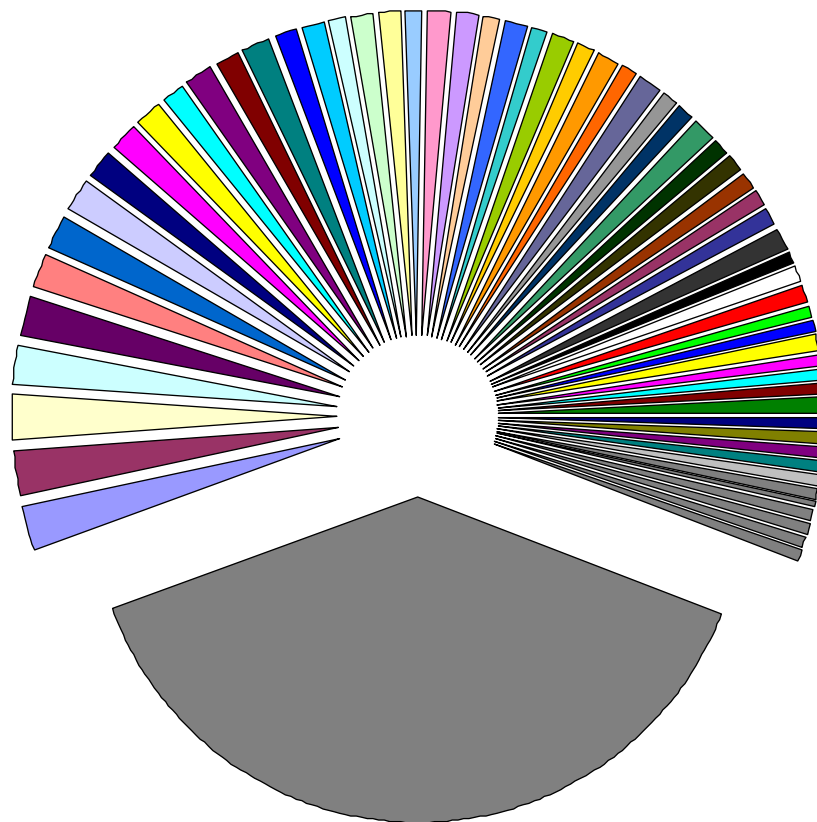
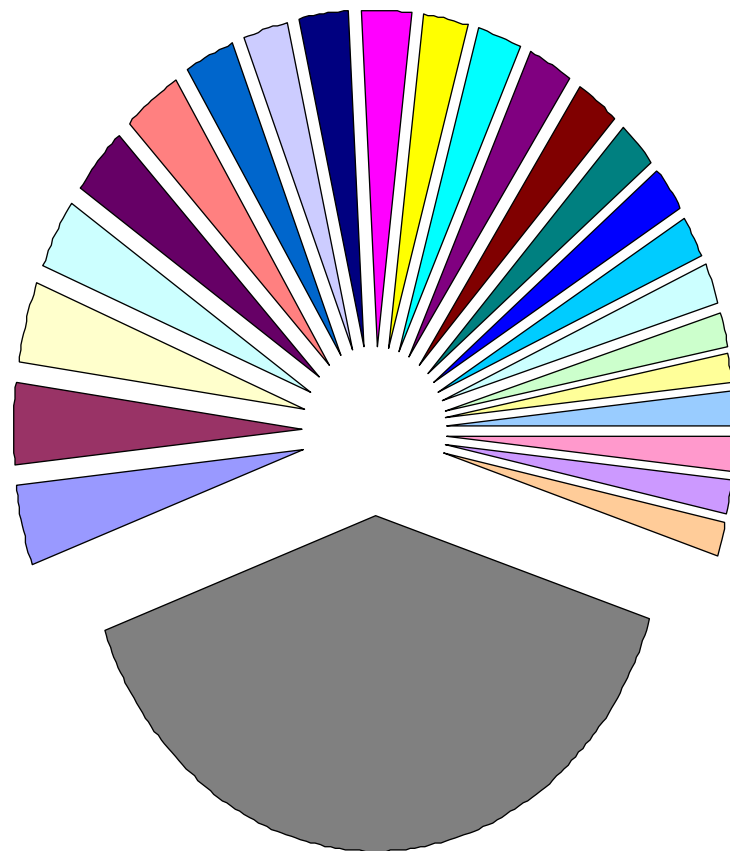


Supplemental Figure 1. Original pie-chart and legend to Up-regulated GO families



- ▣ cell organization and biogenesis
- ▣ biopolymer modification
- transferase activity
- purine nucleotide binding
- ATP binding
- intracellular non-membrane-bound organelle
- kinase activity
- organelle organization and biogenesis
- response to stress
- ▣ phosphate metabolism
- ▣ cellular localization
- ▣ phosphotransferase activity, alcohol group as acceptor
- morphogenesis
- negative regulation of biological process
- cell differentiation
- protein kinase activity
- phosphorylation
- ubiquitin cycle
- enzyme regulator activity
- negative regulation of physiological process
- protein serine/threonine kinase activity
- cell cycle
- RNA binding
- intracellular protein transport
- nervous system development
- ligase activity
- positive regulation of physiological process
- ▣ transcription from RNA polymerase II promoter
- cell motility
- locomotion
- regulation of cell cycle
- development
- ▣ protein modification
- protein complex
- ▣ adenylyl nucleotide binding
- non-membrane-bound organelle
- transferase activity, transferring phosphorus-containing groups
- transcription regulator activity
- intracellular signaling cascade
- ▣ phosphorus metabolism
- establishment of cellular localization
- intracellular transport
- protein localization
- establishment of protein localization
- cytoskeleton
- protein transport
- negative regulation of cellular process
- organ development
- negative regulation of cellular physiological process
- protein amino acid phosphorylation
- transcription factor activity
- ▣ DNA metabolism
- cytoskeleton organization and biogenesis
- system development
- cellular morphogenesis
- positive regulation of cellular process
- positive regulation of cellular physiological process
- cytoskeletal protein binding
- response to abiotic stimulus
- localization of cell
- regulation of progression through cell cycle
- other

Supplemental Figure 2. Original pie-chart and legend to Down-regulated GO families



- system development
- morphogenesis
- nervous system development
- extracellular matrix (sensu Metazoa)
- organ morphogenesis
- transcription factor activity
- peptidase activity
- positive regulation of cellular process
- neuron differentiation
- angiogenesis
- blood vessel development
- enzyme linked receptor protein signaling pathway
- other
- cell differentiation
- transcription regulator activity
- cellular biosynthesis
- extracellular matrix
- calcium ion binding
- cell adhesion
- positive regulation of biological process
- macromolecule biosynthesis
- neurogenesis
- blood vessel morphogenesis
- vasculature development
- sequence-specific DNA binding

Table 1. Up-regulated GO gene sets in co-cultured cholangiocytes

GO Family ^A	Gene Count ^B	Percentage ^C	EASE Score ^D
Cell organization and biogenesis	88	9.95%	7.92000E-07
Development	88	9.95%	7.77000E-05
Biopolymer modification	86	9.73%	5.94000E-09
Protein modification	82	9.28%	2.39000E-08
Transferase activity	73	8.26%	2.03398E-03
Protein complex	72	8.14%	1.74018E-02
Purine nucleotide binding	70	7.92%	2.46000E-04
Adenyl nucleotide binding	59	6.67%	3.15000E-04
ATP binding	58	6.56%	2.39000E-04
Non-membrane-bound organelle	55	6.22%	6.66968E-03
Intracellular non-membrane-bound organelle	55	6.22%	6.66968E-03
Transferase activity, transferring phosphorus-containing groups	52	5.88%	1.53000E-04
Kinase activity	50	5.66%	1.94000E-05
Transcription regulator activity	49	5.54%	4.94198E-03
Organelle organization and biogenesis	47	5.32%	5.51000E-04
Intracellular signaling cascade	45	5.09%	1.05950E-03
Response to stress	44	4.98%	3.15927E-02
Phosphorus metabolism	42	4.75%	6.09000E-04
Phosphate metabolism	42	4.75%	6.09000E-04
Establishment of cellular localization	41	4.64%	1.24000E-05
Cellular localization	41	4.64%	1.52000E-05
Intracellular transport	40	4.52%	2.50000E-05
Phosphotransferase activity, alcohol group as acceptor	40	4.52%	5.50000E-05
Protein localization	40	4.52%	5.51000E-05
Morphogenesis	40	4.52%	1.09671E-02
Establishment of protein localization	39	4.41%	4.70000E-05
Negative regulation of biological process	39	4.41%	4.83000E-04
Cytoskeleton	39	4.41%	1.27384E-03
Cell differentiation	39	4.41%	1.17331E-02
Protein transport	38	4.30%	4.71000E-05
Protein kinase activity	37	4.19%	2.62000E-05
Negative regulation of cellular process	37	4.19%	4.09000E-04
Phosphorylation	37	4.19%	5.42000E-04
Organ development	37	4.19%	5.72216E-02
Ubiquitin cycle	36	4.07%	1.44000E-07
Negative regulation of cellular physiological process	35	3.96%	8.22000E-05
Enzyme regulator activity	35	3.96%	1.02000E-04
Protein amino acid phosphorylation	35	3.96%	1.43000E-04
Negative regulation of physiological process	35	3.96%	1.97000E-04
Transcription factor activity	34	3.85%	4.51600E-02
Protein serine/threonine kinase activity	30	3.39%	1.42000E-04
DNA metabolism	30	3.39%	1.09557E-03
Cell cycle	29	3.28%	1.60859E-02
Cytoskeleton organization and biogenesis	28	3.17%	2.70000E-04
RNA binding	27	3.05%	1.15740E-03
System development	27	3.05%	8.53358E-03

Intracellular protein transport	25	2.83%	2.16018E-03
Cellular morphogenesis	24	2.71%	1.02468E-03
Nervous system development	24	2.71%	1.96118E-02
Positive regulation of cellular process	24	2.71%	7.63648E-02
Ligase activity	23	2.60%	4.28579E-03
Positive regulation of cellular physiological process	23	2.60%	5.68016E-02
Positive regulation of physiological process	23	2.60%	8.58718E-02
Cytoskeletal protein binding	22	2.49%	3.47000E-04
Transcription from RNA polymerase II promoter	22	2.49%	1.45890E-03
Response to abiotic stimulus	21	2.38%	7.34388E-02
Cell motility	20	2.26%	3.89000E-04
Localization of cell	20	2.26%	3.89000E-04
Locomotion	20	2.26%	4.67000E-04
Regulation of progression through cell cycle	20	2.26%	7.20075E-03
Regulation of cell cycle	20	2.26%	7.43645E-03

^A List of GO sets that have been up-regulated in co-cultured cholangiocytes compared to mono-cultures. ^B Absolute number of genes altered. ^C Percentage of genes altered within the GO family. ^D EASE score.

Probe Set ID Symbol	Average Expression Coculture/Average Expression Monoculture	Gene Title	Gene
1415670_at	0.95061	"coatomer protein complex, subunit gamma"	Copg
1415671_at	0.94545	"ATPase, H ⁺ transporting, lysosomal V0 subunit D1"	Atp6v0d1
1415672_at	0.93091	"golgi autoantigen, golgin subfamily a, 7"	Golga7
1415673_at	1.02658	phosphoserine phosphatase	Psph
1415674_a_at	1.09169	trafficking protein particle complex 4	Trappc4
1415675_at	1.05584	dolichol-phosphate (beta-D) mannosyltransferase 2	Dpm2
1415676_a_at	1.06288	"proteasome (prosome, macropain) subunit, beta type 5"	Psmb5
1415677_at	0.97995	dehydrogenase/reductase (SDR family) member 1	Dhrs1
1415678_at	0.95973	"protein phosphatase 1A, magnesium dependent, alpha isoform"	Ppm1a
1415679_at	1.05311	presenilin enhancer 2 homolog (C. elegans)	Psenen
1415680_at	1.12135	anaphase promoting complex subunit 1	Anapc1
1415681_at	1.07250	mitochondrial ribosomal protein L43	Mrpl43
1415682_at	0.99309	exportin 7	Xpo7
1415683_at	1.04942	N-myristoyltransferase 1	Nmt1
1415684_at	1.07304	autophagy-related 5 (yeast)	Atg5
1415685_at	1.12205	mitochondrial translational initiation factor 2	Mtif2
1415686_at	0.95201	"RAB14, member RAS oncogene family"	Rab14
1415687_a_at	0.87781	prosaposin	Psap
1415688_at	0.85738	"ubiquitin-conjugating enzyme E2G 1 (UBC7 homolog, C. elegans)"	Ube2g1
1415689_s_at	0.74390	zinc finger protein 306	Zfp306
1415690_at	1.17290	mitochondrial ribosomal protein L27	Mrpl27
1415691_at	1.07939	"discs, large homolog 1 (Drosophila)"	Dlgh1
1415692_s_at	1.10304	calnexin	Canx
1415693_at	1.00621	"Der1-like domain family, member 1"	Der1l
1415694_at	1.10393	tryptophanyl-tRNA synthetase	Wars
1415695_at	1.02342	"proteasome (prosome, macropain) subunit, alpha type 1"	Psma1
1415696_at	0.88722	SAR1 gene homolog A (S. cerevisiae)	Sar1a
1415697_at	1.12479	RIKEN cDNA E430034L04 gene	E430034L04Rik
1415698_at	0.96860	golgi phosphoprotein 2	Golph2
1415699_a_at	0.97535	G protein pathway suppressor 1	Gps1
1415700_a_at	0.98348	"signal sequence receptor, gamma"	Ssr3
1415701_x_at	1.01856	ribosomal protein L23	Rpl23
1415702_a_at	0.87204	C-terminal binding protein 1	Ctbp1
1415703_at	1.01601	"HECT, UBA and WWE domain containing 1"	Huwe1
1415704_a_at	1.20063	carnitine deficiency-associated gene expressed in ventricle 3	Cdv3
1415705_at	0.87415	RIKEN cDNA 9130011J15 gene	9130011J15Rik
1415706_at	0.91394	coatomer protein complex subunit alpha	Copa
1415707_at	0.98701	anaphase promoting complex subunit 2	Anapc2
1415708_at	0.95803	taurine upregulated gene 1	Tug1
1415709_s_at	0.99431	golgi-specific brefeldin A-resistance factor 1	Gbf1

1415710_at 1.01728 COX18 cytochrome c oxidase assembly homolog (*S. cerevisiae*) Cox18
 1415711_at 0.92829 ADP-ribosylation factor guanine nucleotide-exchange factor 1(brefeldin A-inhibited) Arfgef1
 1415712_at 0.86270 "zinc finger, RAN-binding domain containing 1" Zranb1
 1415713_a_at 0.97428 DEAD (Asp-Glu-Ala-Asp) box polypeptide 24 Ddx24
 1415714_a_at 1.05008 RIKEN cDNA 2610209M04 gene 2610209M04Rik
 1415715_at 0.94432 transmembrane protein 129 Tmem129
 1415716_a_at 0.91185 ribosomal protein S27 Rps27
 1415717_at 0.91605 RIKEN cDNA 4931406I20 gene 4931406I20Rik
 1415718_at 1.00544 SAP30-like Sap30l
 1415719_s_at 0.92885 armadillo repeat containing 1 Armc1
 1415720_s_at 0.96774 MAD2L1 binding protein Mad2l1bp
 1415721_a_at 0.95385 RIKEN cDNA 1200013P24 gene 1200013P24Rik
 1415722_a_at 0.97559 RIKEN cDNA 1110059P08 gene 1110059P08Rik
 1415723_at 1.07928 eukaryotic translation initiation factor 5 Eif5
 1415724_a_at 1.01143 cell division cycle 42 homolog (*S. cerevisiae*) Cdc42
 1415725_at 1.14175 RRN3 RNA polymerase I transcription factor homolog (yeast) Rrn3
 1415726_at 0.97237 ankyrin repeat domain 17 Ankrd17
 1415727_at 1.01865 apolipoprotein A-I binding protein Apoalbp
 1415728_at 1.19108 "poly(A) binding protein, nuclear 1" Pabpn1
 1415729_at 0.94445 3-phosphoinositide dependent protein kinase-1 Pdkp1
 1415730_at 0.99416 RIKEN cDNA 5730453I16 gene 5730453I16Rik
 1415731_at 0.99396 angel homolog 2 (*Drosophila*) Angel2
 1415732_at 0.88899 HLA-B associated transcript 5 Bat5
 1415733_a_at 1.11468 RIKEN cDNA 1110019J04 gene 1110019J04Rik
 1415734_at 1.00006 "RAB7, member RAS oncogene family" Rab7
 1415735_at 1.01757 damage specific DNA binding protein 1 Ddb1
 1415736_at 0.98003 prefoldin 5 Pfdn5
 1415737_at 0.97661 riboflavin kinase Rfk
 1415738_at 0.88480 thioredoxin domain containing 12 (endoplasmic reticulum) Txndc12
 1415739_at 0.91974 RIKEN cDNA 3100004P22 gene 3100004P22Rik
 1415740_at 1.15203 "protease (prosome, macropain) 26S subunit, ATPase 5" Psmc5
 1415741_at 0.88485 transmembrane protein 165 Tmem165
 1415742_at 0.99604 ancient ubiquitous protein Aup1
 1415743_at 0.82408 histone deacetylase 5 Hdac5
 1415744_at 1.15763 H2-K region expressed gene 2 H2-Ke2
 1415745_a_at 1.02067 Down syndrome critical region gene 3 Dscr3
 1415746_at 0.94838 capicua homolog (*Drosophila*) Cic
 1415747_s_at 0.99108 RIO kinase 3 (yeast) Riok3
 1415748_a_at 1.05675 dynactin 5 Dctn5
 1415749_a_at 1.13763 Ras-related GTP binding C Rragc
 1415750_at 1.01870 transducin (beta)-like 3 Tbl3
 1415751_at 0.95825 "heterochromatin protein 1, binding protein 3" Hp1bp3

1415752_at 0.99537 cDNA sequence BC031181 BC031181
 1415753_at 0.92153 "DNA segment, Chr 10, Brigham & Women's Genetics 1364 expressed"
 D10Bwg1364e
 1415754_at 0.96349 polymerase (RNA) II (DNA directed) polypeptide F Polr2f
 1415755_a_at 0.96021 ubiquitin-conjugating enzyme E2 variant 1 /// similar to ubiquitin-conjugating
 enzyme E2 variant 1 Ube2v1 /// LOC666853
 1415756_a_at 0.94673 SNAP-associated protein Snapap
 1415757_at 0.98763 golgi-specific brefeldin A-resistance factor 1 Gbf1
 1415758_at 0.99324 furry homolog-like (Drosophila) Fryl
 1415759_a_at 1.07680 hepatitis B virus x interacting protein Hbxip
 1415760_s_at 0.98276 ATX1 (antioxidant protein 1) homolog 1 (yeast) Atox1
 1415761_at 1.02689 mitochondrial ribosomal protein L52 Mrpl52
 1415762_x_at 1.00904 mitochondrial ribosomal protein L52 Mrpl52
 1415763_a_at 1.04934 --- ---
 1415764_at 1.07036 zinc finger CCCH type containing 11A Zc3h11a
 1415765_at 0.98615 heterogeneous nuclear ribonucleoprotein U-like 2 Hnrpul2
 1415766_at 0.92974 SEC22 vesicle trafficking protein homolog B (S. cerevisiae) Sec22b
 1415767_at 0.97333 YTH domain family 1 Ythdf1
 1415768_a_at 0.89896 ubiquitin-conjugating enzyme E2R 2 Ube2r2
 1415769_at 0.86374 itchy Itch
 1415770_at 0.87368 WD repeat domain 6 Wdr6
 1415771_at 1.31355 nucleolin Ncl
 1415772_at 1.19104 nucleolin Ncl
 1415773_at 1.25252 nucleolin Ncl
 1415774_at 1.00502 signal transducer and activator of transcription interacting protein 1 Statip1
 1415775_at 0.98471 retinoblastoma binding protein 7 Rbbp7
 1415776_at 1.04301 "aldehyde dehydrogenase family 3, subfamily A2" Aldh3a2
 1415777_at 0.95752 pancreatic lipase related protein 1 Pnliprp1
 1415778_at 0.97861 mortality factor 4 like 2 Morf4l2
 1415779_s_at 0.93269 "actin, gamma, cytoplasmic 1" Actg1
 1415780_a_at 0.96980 "armadillo repeat containing, X-linked 2" Armcx2
 1415781_a_at 0.97614 SMT3 suppressor of mif two 3 homolog 2 (yeast) /// similar to SMT3
 suppressor of mif two 3 homolog 2 /// similar to SMT3 suppressor of mif two 3 homolog 2 Sumo2 ///
 LOC672381 /// LOC675440
 1415782_at 0.97112 SMT3 suppressor of mif two 3 homolog 2 (yeast) Sumo2
 1415783_at 0.98985 vacuolar protein sorting 35 Vps35
 1415784_at 0.95225 vacuolar protein sorting 35 Vps35
 1415785_a_at 1.09653 chaperonin subunit 8 (theta) Cct8
 1415786_at 0.85776 progastricsin (pepsinogen C) Pgc
 1415787_at 0.97904 alpha glucosidase 2 alpha neutral subunit Ganab
 1415788_at 0.99339 ubiquitin-like domain containing CTD phosphatase 1 Ublcp1
 1415789_a_at 1.06763 ubiquitin-like domain containing CTD phosphatase 1 /// similar to ubiquitin-
 like domain containing CTD phosphatase 1 Ublcp1 /// LOC665689

1415790_at	1.06106	ubiquitin-like domain containing CTD phosphatase 1	Ublcp1
1415791_at	0.92732	ring finger protein 34	Rnf34
1415792_at	1.00830	RanBP-type and C3HC4-type zinc finger containing 1	Rbck1
1415793_at	1.00955	pyridoxine 5'-phosphate oxidase	Pnp0
1415794_a_at	0.92303	spindlin	Spin
1415795_at	0.99445	spindlin	Spin
1415796_at	0.98173	DAZ associated protein 2	Dazap2
1415797_at	0.86284	"discoidin domain receptor family, member 1"	Ddr1
1415798_at	0.88710	"discoidin domain receptor family, member 1"	Ddr1
1415799_at	1.02262	WW domain binding protein 11	Wbp11
1415800_at	0.91939	gap junction membrane channel protein alpha 1	Gja1
1415801_at	0.87100	gap junction membrane channel protein alpha 1	Gja1
1415802_at	1.03586	"solute carrier family 16 (monocarboxylic acid transporters), member 1"	Slc16a1
1415803_at	1.04938	chemokine (C-X3-C motif) ligand 1	Cx3cl1
1415804_at	1.04757	chemokine (C-X3-C motif) ligand 1	Cx3cl1
1415805_at	0.90602	"colipase, pancreatic"	Clps
1415806_at	0.92149	"plasminogen activator, tissue"	Plat
1415807_s_at	0.99381	"splicing factor, arginine/serine-rich 2 (SC-35)"	Sfrs2
1415808_at	1.04800	trophoblast specific protein alpha	Tpbpa
1415809_at	1.06848	trophoblast specific protein alpha	Tpbpa
1415810_at	1.06940	"ubiquitin-like, containing PHD and RING finger domains, 1"	Uhrf1
1415811_at	1.04151	"ubiquitin-like, containing PHD and RING finger domains, 1"	Uhrf1
1415812_at	0.89135	gelsolin	Gsn
1415813_at	1.05977	apoptosis inhibitor 5	Api5
1415814_at	0.98662	"ATPase, H+ transporting, lysosomal V1 subunit B2"	Atp6v1b2
1415815_at	0.94270	tripartite motif protein 27	Trim27
1415816_at	1.16392	chaperonin subunit 7 (eta)	Cct7
1415817_s_at	1.01732	chaperonin subunit 7 (eta)	Cct7
1415818_at	0.79656	annexin A6	Anxa6
1415819_a_at	0.95664	"protein phosphatase 2 (formerly 2A), regulatory subunit A (PR 65), alpha isoform"	Ppp2r1a
1415820_x_at	0.98573	"non-POU-domain-containing, octamer binding protein"	Nono
1415821_at	0.95726	neuroplastin	Nptn
1415822_at	1.05457	stearoyl-Coenzyme A desaturase 2	Scd2
1415823_at	1.14659	stearoyl-Coenzyme A desaturase 2	Scd2
1415824_at	1.02222	stearoyl-Coenzyme A desaturase 2	Scd2
1415825_s_at	0.93359	"casein kinase 1, delta"	Csnk1d
1415826_at	1.12403	"ATPase, H+ transporting, lysosomal V1 subunit H"	Atp6v1h
1415827_a_at	0.92140	"DNA segment, Chr 3, University of California at Los Angeles 1"	D3Ucla1
1415828_a_at	0.94854	"DNA segment, Chr 3, University of California at Los Angeles 1"	D3Ucla1
1415829_at	1.11509	lamin B receptor	Lbr
1415830_at	0.97824	"origin recognition complex, subunit 5-like (<i>S. cerevisiae</i>)"	Orc5l
1415831_at	1.08753	"proteasome (prosome, macropain) 26S subunit, non-ATPase, 2"	Psmd2

1415832_at 0.92633 "angiotensin II receptor, type 2" Agtr2
 1415833_x_at 0.89011 ribosomal protein S29 /// similar to 40S ribosomal protein S29 /// similar to 40S ribosomal protein S29 Rps29 /// LOC629957 /// LOC637599
 1415834_at 0.93085 dual specificity phosphatase 6 Dusp6
 1415835_at 1.12113 chorionic somatomammotropin hormone 2 Csh2
 1415836_at 1.08310 "aldehyde dehydrogenase 18 family, member A1" Aldh18a1
 1415837_at 0.95275 kallikrein 1 Klk1
 1415838_at 0.92171 serine incorporator 1 Serinc1
 1415839_a_at 1.16155 nucleophosmin 1 Npm1
 1415840_at 1.01054 "ELOVL family member 5, elongation of long chain fatty acids (yeast)" Elovl5
 1415841_at 0.91448 dynein cytoplasmic 1 intermediate chain 2 Dync1i2
 1415842_at 0.91587 G protein beta subunit-like Gbl
 1415843_at 1.10030 G protein beta subunit-like Gbl
 1415844_at 1.06500 synaptotagmin IV Syt4
 1415845_at 1.00003 synaptotagmin IV Syt4
 1415846_a_at 0.82866 lactate dehydrogenase C Ldhc
 1415847_at 1.16015 lactate dehydrogenase C Ldhc
 1415848_at 1.05468 chorionic somatomammotropin hormone 1 Csh1
 1415849_s_at 0.86249 stathmin 1 Stmn1
 1415850_at 1.10982 RAS p21 protein activator 3 Rasa3
 1415851_a_at 1.06399 inosine 5'-phosphate dehydrogenase 2 Impdh2
 1415852_at 1.07864 inosine 5'-phosphate dehydrogenase 2 Impdh2
 1415853_at 1.02658 differentially expressed in FDCP 8 Def8
 1415854_at 0.76649 kit ligand Kitl
 1415855_at 0.82832 kit ligand Kitl
 1415856_at 0.94556 embigin Emb
 1415857_at 1.00515 embigin Emb
 1415858_at 1.10628 "eukaryotic translation initiation factor 3, subunit 8" Eif3s8
 1415859_at 1.01663 "eukaryotic translation initiation factor 3, subunit 8" Eif3s8
 1415860_at 1.09785 karyopherin (importin) alpha 2 /// similar to Importin alpha-2 subunit (Karyopherin alpha-2 subunit) (SRP1-alpha) (RAG cohort protein 1) (Pendulin) (Pore targeting complex 58 kDa subunit) (PTAC58) (Importin alpha P1) /// similar to Importin alpha-2 subunit (Karyopherin alpha-2 subunit) (SRP1-alpha) (RAG cohort protein 1) (Pendulin) (Pore targeting complex 58 kDa subunit) (PTAC58) (Importin alpha P1) Kpna2 /// LOC670551 /// LOC671639
 1415861_at 1.10492 tyrosinase-related protein 1 Tyrp1
 1415862_at 1.04405 tyrosinase-related protein 1 Tyrp1
 1415863_at 1.03167 "eukaryotic translation initiation factor 4, gamma 2" Eif4g2
 1415864_at 0.96637 "2,3-bisphosphoglycerate mutase" Bpgm
 1415865_s_at 1.10339 "2,3-bisphosphoglycerate mutase" Bpgm
 1415866_at 1.12592 unc-45 homolog A (C. elegans) Unc45a
 1415867_at 1.03903 chaperonin subunit 4 (delta) Cct4
 1415868_at 1.11574 chaperonin subunit 4 (delta) Cct4
 1415869_a_at 0.95705 tripartite motif protein 28 Trim28

1415870_at	0.88942	calumenin	Calu	
1415871_at	0.88675	"transforming growth factor, beta induced"	Tgfb1	
1415872_at	1.04502	heterogeneous nuclear ribonucleoprotein H1	Hnrph1	
1415873_a_at	0.95042	ARP1 actin-related protein 1 homolog A (yeast)	Actr1a	
1415874_at	0.83124	sprouty homolog 1 (Drosophila)	Spry1	
1415875_at	0.90589	RIKEN cDNA 3010003L21 gene	3010003L21Rik	
1415876_a_at	1.03884	ribosomal protein S26	Rps26	
1415877_at	0.40085	dihydropyrimidinase-like 3	Dpysl3	
1415878_at	1.07404	ribonucleotide reductase M1	Rrm1	
1415879_a_at	1.06398	"ribosomal protein, large P2"	Rplp2	
1415880_a_at	0.99849	lysosomal membrane glycoprotein 1	Lamp1	
1415881_at	1.02437	growth hormone inducible transmembrane protein	Ghitm	
1415882_at	1.01929	growth hormone inducible transmembrane protein	Ghitm	
1415883_a_at	1.03841	"elastase 3, pancreatic"	Ela3	
1415884_at	0.97289	"elastase 3, pancreatic"	Ela3	
1415885_at	1.02161	chromogranin B	Chgb	
1415886_at	0.99392	SH2 domain containing 3C	Sh2d3c	
1415887_at	0.97795	Trk-fused gene	Tfg	
1415888_at	1.08371	hepatoma-derived growth factor	Hdgf	
1415889_a_at	0.97353	"heat shock protein 90kDa beta (Grp94), member 1"	Hsp90b1	
1415890_at	0.88366	3'-phosphoadenosine 5'-phosphosulfate synthase 1	Papss1	
1415891_at	1.05873	"succinate-CoA ligase, GDP-forming, alpha subunit"	Suc1g1	
1415892_at	0.98659	sphingosine phosphate lyase 1	Sgpl1	
1415893_at	0.93696	sphingosine phosphate lyase 1	Sgpl1	
1415894_at	1.07266	ectonucleotide pyrophosphatase/phosphodiesterase 2	Enpp2	
1415895_at	0.96408	small nuclear ribonucleoprotein N	Snrpn	
1415896_x_at	0.97529	small nuclear ribonucleoprotein N	Snrpn	
1415897_a_at	1.18015	microsomal glutathione S-transferase 1	Mgst1	
1415898_at	1.03431	microsomal glutathione S-transferase 1	Mgst1	
1415899_at	0.82573	Jun-B oncogene	Junb	
1415900_a_at	1.03759	kit oncogene	Kit	
1415901_at	0.87565	"procollagen-lysine, 2-oxoglutarate 5-dioxygenase 3"	Plod3	
1415902_at	0.96009	"aldehyde dehydrogenase family 7, member A1"	Aldh7a1	
1415903_at	1.07623	"solute carrier family 38, member 1"	Slc38a1	
1415904_at	0.64291	lipoprotein lipase	Lpl	
1415905_at	0.89810	regenerating islet-derived 1	Reg1	
1415906_at	0.87979	"thymosin, beta 4, X chromosome"	Tmsb4x	
1415907_at	0.95668	cyclin D3	Ccnd3	
1415908_at	0.85584	"testis-specific protein, Y-encoded-like 1"	Tspyl1	
1415909_at	1.11368	stress-induced phosphoprotein 1	Stip1	
1415910_s_at	0.99490	cytokine induced apoptosis inhibitor 1	Ciapin1	
1415911_at	1.18125	imprinted and ancient	Impact	
1415912_a_at	1.10500	ribosomal protein S13 /// similar to ribosomal protein S13 /// similar to		

ribosomal protein S13 /// similar to ribosomal protein S13 /// similar to ribosomal protein S13 /// similar to ribosomal protein S13
 to ribosomal protein S13 Rps13 /// LOC625298 /// LOC628061 /// LOC633920 /// LOC637251 /// LOC671723

1415913_at 1.10956 ribosomal protein S13 /// similar to ribosomal protein S13 /// similar to ribosomal protein S13 /// similar to ribosomal protein S13 /// similar to ribosomal protein S13 /// similar to ribosomal protein S13 /// similar to ribosomal protein S13
 to ribosomal protein S13 /// similar to ribosomal protein S13 /// similar to ribosomal protein S13
 Rps13 /// LOC625298 /// LOC628061 /// LOC633920 /// LOC637251 /// LOC668239 /// LOC671723 /// LOC671887

1415914_at 1.04724 heterogeneous nuclear ribonucleoprotein A/B Hnrpab

1415915_at 1.03425 DEAD (Asp-Glu-Ala-Asp) box polypeptide 1 Ddx1

1415916_a_at 1.04265 "methylenetetrahydrofolate dehydrogenase (NADP+ dependent), methenyltetrahydrofolate cyclohydrolase, formyltetrahydrofolate synthase" Mthfd1

1415917_at 1.03169 "methylenetetrahydrofolate dehydrogenase (NADP+ dependent), methenyltetrahydrofolate cyclohydrolase, formyltetrahydrofolate synthase" Mthfd1

1415918_a_at 1.17086 triosephosphate isomerase 1 Tpi1

1415919_at 0.94002 "neural proliferation, differentiation and control gene 1" Npdc1

1415920_at 0.77838 "cleavage stimulation factor, 3' pre-RNA subunit 2, tau" Cstf2t

1415921_a_at 0.99104 "tumor necrosis factor receptor superfamily, member 19" Tnfrsf19

1415922_s_at 0.80933 MARCKS-like 1 Marcksl1

1415923_at 0.98202 necdin Ndn

1415924_at 1.00981 transition protein 1 Tnp1

1415925_a_at 0.90608 nucleoporin 62 Nup62

1415926_at 0.94634 nucleoporin 62 Nup62

1415927_at 1.05778 "actin, alpha, cardiac" Actc1

1415928_a_at 0.87793 microtubule-associated protein 1 light chain 3 beta Map1lc3b

1415929_at 0.89484 microtubule-associated protein 1 light chain 3 beta Map1lc3b

1415930_a_at 1.01147 microtubule-associated protein 1 light chain 3 beta Map1lc3b

1415931_at 0.79602 insulin-like growth factor 2 Igf2

1415932_x_at 0.90247 "ATPase, class II, type 9A" Atp9a

1415933_a_at 0.99413 "cytochrome c oxidase, subunit Va" Cox5a

1415934_at 0.94509 "COP9 (constitutive photomorphogenic) homolog, subunit 8 (Arabidopsis thaliana)" Cops8

1415935_at 0.95941 SPARC related modular calcium binding 2 Smoc2

1415936_at 0.89077 breast cancer anti-estrogen resistance 3 Bcar3

1415937_s_at 0.91150 programmed cell death 6 interacting protein Pcd6ip

1415938_at 0.92625 "serine peptidase inhibitor, Kazal type 3" Spink3

1415939_at 0.66901 fibromodulin Fmod

1415940_at 0.98303 "zinc finger, AN1-type domain 2A" Zfand2a

1415941_s_at 0.98476 "zinc finger, AN1-type domain 2A" Zfand2a

1415942_at 1.03210 ribosomal protein 10 /// similar to ribosomal protein L10 /// similar to ribosomal protein L10
 Rpl10 /// LOC234703 /// LOC638133

1415943_at 0.83362 syndecan 1 Sdc1

1415944_at 0.84329 syndecan 1 Sdc1

1415945_at 0.93153 "minichromosome maintenance deficient 5, cell division cycle 46 (S. cerevisiae)" Mcm5
 1415946_at 0.87425 "phosphatidylinositol glycan anchor biosynthesis, class Q" Pigq
 1415947_at 1.06145 cellular repressor of E1A-stimulated genes 1 Creg1
 1415948_at 1.09902 cellular repressor of E1A-stimulated genes 1 Creg1
 1415949_at 0.83458 carboxypeptidase E Cpe
 1415950_a_at 0.99872 phosphatidylethanolamine binding protein 1 Pebp1
 1415951_at 0.80485 FK506 binding protein 10 Fkbp10
 1415952_at 1.05065 MAP/microtubule affinity-regulating kinase 2 Mark2
 1415953_s_at 0.99158 MAP/microtubule affinity-regulating kinase 2 Mark2
 1415954_at 1.08081 "trypsin 4 /// protease, serine, 1 (trypsin 1) /// RIKEN cDNA 1810049H19 gene /// trypsinogen 10" Try4 /// Prss1 /// 1810049H19Rik /// LOC436522
 1415955_x_at 1.08008 protamine 1 Prm1
 1415956_a_at 1.01046 PCTAIRE-motif protein kinase 1 Pctk1
 1415957_a_at 0.97883 novel nuclear protein 1 Nnp1
 1415958_at 0.84053 "solute carrier family 2 (facilitated glucose transporter), member 4" Slc2a4
 1415959_at 0.89012 "solute carrier family 2 (facilitated glucose transporter), member 4" Slc2a4
 1415960_at 0.90126 myeloperoxidase Mpo
 1415961_at 0.81868 integral membrane protein 2C Itm2c
 1415962_at 0.99048 "eukaryotic translation initiation factor 3, subunit 3 (gamma)" Eif3s3
 1415963_at 1.11676 heterogeneous nuclear ribonucleoprotein H2 Hnrph2
 1415964_at 1.40388 stearoyl-Coenzyme A desaturase 1 Scd1
 1415965_at 1.40597 stearoyl-Coenzyme A desaturase 1 Scd1
 1415966_a_at 0.94437 NADH dehydrogenase (ubiquinone) flavoprotein 1 Ndufv1
 1415967_at 0.96119 NADH dehydrogenase (ubiquinone) flavoprotein 1 Ndufv1
 1415968_a_at 0.88147 kidney androgen regulated protein Kap
 1415969_s_at 1.04805 kidney androgen regulated protein Kap
 1415970_at 0.94893 "cytochrome c oxidase, subunit VIc" Cox6c
 1415971_at 1.00054 myristoylated alanine rich protein kinase C substrate Marcks
 1415972_at 1.08693 myristoylated alanine rich protein kinase C substrate Marcks
 1415973_at 0.97335 "0 day neonate cerebellum cDNA, RIKEN full-length enriched library, clone: C230070N19 product:myristoylated alanine rich protein kinase C substrate, full insert sequence" ---
 1415974_at 0.95207 mitogen activated protein kinase kinase 2 Map2k2
 1415975_at 0.85695 calcium regulated heat stable protein 1 Carhsp1
 1415976_a_at 1.00103 calcium regulated heat stable protein 1 Carhsp1
 1415977_at 0.92435 myo-inositol 1-phosphate synthase A1 Isynal
 1415978_at 0.86024 "tubulin, beta 3" Tubb3
 1415979_x_at 1.05233 ribosomal protein L7 Rpl7
 1415980_at 0.99204 "ATP synthase, H+ transporting, mitochondrial F0 complex, subunit c (subunit 9), isoform 2" Atp5g2
 1415981_at 0.98809 HERPUD family member 2 Herpud2
 1415982_at 0.95041 HERPUD family member 2 Herpud2
 1415983_at 1.09232 lymphocyte cytosolic protein 1 Lcp1

1415984_at 0.83823 "acyl-Coenzyme A dehydrogenase, medium chain" Acadm
 1415985_at 1.02914 "splicing factor 3b, subunit 3" Sf3b3
 1415986_at 0.92538 chloride channel 4-2 Clcn4-2
 1415987_at 1.10127 high density lipoprotein (HDL) binding protein Hdlbp
 1415988_at 1.03539 high density lipoprotein (HDL) binding protein Hdlbp
 1415989_at 1.19129 vascular cell adhesion molecule 1 Vcam1
 1415990_at 1.03612 voltage-dependent anion channel 2 Vdac2
 1415991_a_at 0.95531 kelch domain containing 3 Klhdc3
 1415992_at 0.88820 "phosphatidylinositol glycan anchor biosynthesis, class O" Pigo
 1415993_at 1.46319 squalene epoxidase Sqle
 1415994_at 0.88601 "cytochrome P450, family 2, subfamily e, polypeptide 1" Cyp2e1
 1415995_at 0.89945 caspase 6 Casp6
 1415996_at 0.89052 thioredoxin interacting protein Txnip
 1415997_at 0.74160 thioredoxin interacting protein Txnip
 1415998_at 0.98788 voltage-dependent anion channel 1 Vdac1
 1415999_at 0.78428 hairy/enhancer-of-split related with YRPW motif 1 Hey1
 1416000_a_at 1.01835 peroxiredoxin 1 Prdx1
 1416001_a_at 0.91014 coactosin-like 1 (Dictyostelium) Cot11
 1416002_x_at 0.78957 coactosin-like 1 (Dictyostelium) Cot11
 1416003_at 0.84815 claudin 11 Cldn11
 1416004_at 0.93530 "tyrosine 3-monooxygenase/tryptophan 5-monooxygenase activation protein, eta polypeptide" Ywhah
 1416005_at 1.13207 "protease (prosome, macropain) 26S subunit, ATPase 1" Psmc1
 1416006_at 1.01438 midkine Mdk
 1416007_at 0.56739 special AT-rich sequence binding protein 1 Satb1
 1416008_at 0.50955 special AT-rich sequence binding protein 1 Satb1
 1416009_at 0.85239 tetraspanin 3 Tspan3
 1416010_a_at 1.10991 EH-domain containing 1 Ehd1
 1416011_x_at 1.15038 EH-domain containing 1 Ehd1
 1416012_at 1.09124 EH-domain containing 1 Ehd1
 1416013_at 0.76586 "phospholipase D family, member 3" Pld3
 1416014_at 1.00371 "ATP-binding cassette, sub-family E (OABP), member 1" Abce1
 1416015_s_at 1.07786 "ATP-binding cassette, sub-family E (OABP), member 1" Abce1
 1416016_at 0.91215 "transporter 1, ATP-binding cassette, sub-family B (MDR/TAP)" Tap1
 1416017_at 0.93001 "coatamer protein complex, subunit gamma" Copg
 1416018_at 0.95008 down-regulator of transcription 1 Dr1
 1416019_at 0.87504 down-regulator of transcription 1 Dr1
 1416020_a_at 1.04382 "ATP synthase, H+ transporting, mitochondrial F0 complex, subunit c (subunit 9), isoform 1" Atp5g1
 1416021_a_at 1.07194 "fatty acid binding protein 5, epidermal /// similar to Fatty acid-binding protein, epidermal (E-FABP) (Psoriasis-associated fatty acid-binding protein homolog) (PA-FABP) (Keratinocyte lipid-binding protein) /// similar to Fatty acid-binding protein, epidermal (E-FABP) (Psoriasis-associated fatty acid-binding protein homolog) (PA-FABP) (Keratinocyte lipid-binding

protein) /// similar to Fatty acid-binding protein, epidermal (E-FABP) (Psoriasis-associated fatty acid-binding protein homolog) (PA-FABP) (Keratinocyte lipid-binding protein) /// similar to Fatty acid-binding protein, epidermal (E-FABP) (Psoriasis-associated fatty acid-binding protein homolog) (PA-FABP) (Keratinocyte lipid-binding protein)" Fabp5 /// LOC628298 /// LOC637129 /// LOC669559 /// LOC672321

1416022_at 1.10051 "fatty acid binding protein 5, epidermal" Fabp5

1416023_at 0.93678 "fatty acid binding protein 3, muscle and heart" Fabp3

1416024_x_at 1.06047 chaperonin subunit 3 (gamma) Cct3

1416025_at 1.00835 "fibrinogen, gamma polypeptide" Fgg

1416026_a_at 1.08444 ribosomal protein L12 Rpl12

1416027_at 1.00673 programmed cell death 6 Pcd6

1416028_a_at 1.02459 hematological and neurological expressed sequence 1 Hn1

1416029_at 0.75072 Kruppel-like factor 10 Klf10

1416030_a_at 0.87528 minichromosome maintenance deficient 7 (*S. cerevisiae*) Mcm7

1416031_s_at 0.93101 minichromosome maintenance deficient 7 (*S. cerevisiae*) Mcm7

1416032_at 0.90329 transmembrane protein 109 Tmem109

1416033_at 0.98667 transmembrane protein 109 Tmem109

1416034_at 0.85646 CD24a antigen Cd24a

1416035_at 1.06711 --- ---

1416036_at 0.99517 FK506 binding protein 1a Fkbp1a

1416037_a_at 1.01144 chaperonin subunit 2 (beta) Cct2

1416038_at 1.03413 expressed sequence AL033314 AL033314

1416039_x_at 0.70266 cysteine rich protein 61 Cyr61

1416040_at 1.07544 "lipase, gastric" Lipf

1416041_at 0.80530 serum/glucocorticoid regulated kinase Sgk

1416042_s_at 1.01984 nuclear autoantigenic sperm protein (histone-binding) Nasp

1416043_at 1.02684 nuclear autoantigenic sperm protein (histone-binding) Nasp

1416044_at 0.96006 flightless I homolog (*Drosophila*) Fliih

1416045_a_at 0.96813 "SWI/SNF related, matrix associated, actin dependent regulator of chromatin, subfamily b, member 1" Smarcb1

1416046_a_at 1.06369 "fucosidase, alpha-L- 2, plasma" Fuca2

1416047_at 0.92620 "fucosidase, alpha-L- 2, plasma" Fuca2

1416048_at 1.02124 polyhomeotic-like 2 (*Drosophila*) Phc2

1416049_at 1.00693 glycine decarboxylase Glc

1416050_a_at 0.96486 "scavenger receptor class B, member 1" Scarb1

1416051_at 1.08252 complement component 2 (within H-2S) C2

1416052_at 1.00092 phosphoribosyl pyrophosphate synthetase 1 Prps1

1416053_at 1.01560 "leucine rich repeat protein 1, neuronal" Lrrn1

1416054_at 1.04883 ribosomal protein S5 Rps5

1416055_at 0.97089 "amylase 2, pancreatic /// RIKEN cDNA 1810008N23 gene" Amy2 /// 1810008N23Rik

1416056_a_at 0.88312 "NADH dehydrogenase (ubiquinone) 1 beta subcomplex, 11" Ndufb11

1416057_at 0.91955 "NADH dehydrogenase (ubiquinone) 1 beta subcomplex, 11" Ndufb11

1416058_s_at 0.93974 "ATP synthase, H⁺ transporting, mitochondrial F1 complex, gamma polypeptide 1" Atp5c1
 1416059_at 0.91878 SEC23B (*S. cerevisiae*) Sec23b
 1416060_at 1.02558 "TBC1 domain family, member 15" Tbc1d15
 1416061_at 1.08737 "TBC1 domain family, member 15" Tbc1d15
 1416062_at 1.01759 "TBC1 domain family, member 15" Tbc1d15
 1416063_x_at 1.01823 CEA-related cell adhesion molecule 11 Ceacam11
 1416064_a_at 0.97708 heat shock 70kD protein 5 (glucose-regulated protein) Hspa5
 1416065_a_at 1.07026 ankyrin repeat domain 10 Ankrd10
 1416066_at 0.97082 CD9 antigen Cd9
 1416067_at 1.17273 interferon-related developmental regulator 1 Ifrd1
 1416068_at 1.07517 lysyl-tRNA synthetase /// similar to lysyl-tRNA synthetase Kars ///
 LOC631033
 1416069_at 0.99112 "phosphofructokinase, platelet" Pfkp
 1416070_a_at 1.15744 DEAD (Asp-Glu-Ala-Asp) box polypeptide 18 Ddx18
 1416071_at 1.14761 DEAD (Asp-Glu-Ala-Asp) box polypeptide 18 Ddx18
 1416072_at 0.86938 CD34 antigen Cd34
 1416073_a_at 0.98002 nucleoporin 85 Nup85
 1416074_a_at 0.97483 ribosomal protein L28 Rpl28
 1416075_at 1.07563 salvador homolog 1 (*Drosophila*) Sav1
 1416076_at 1.09770 "cyclin B1, related sequence 1 /// cyclin B1 /// similar to G2/mitotic-specific cyclin-B1 /// similar to G2/mitotic-specific cyclin-B1" Ccnb1-rs1 /// Ccnb1 /// LOC434175 ///
 LOC672788
 1416077_at 1.30351 adrenomedullin Adm
 1416078_s_at 0.95535 v-raf-leukemia viral oncogene 1 Raf1
 1416079_a_at 0.99266 "actin related protein 2/3 complex, subunit 1A" Arpc1a
 1416080_at 0.85509 a disintegrin and metallopeptidase domain 15 (metargidin) Adam15
 1416081_at 0.98116 MAD homolog 1 (*Drosophila*) Smad1
 1416082_at 0.97657 "RAB1, member RAS oncogene family" Rab1
 1416083_at 0.91552 "zinc finger, AN1-type domain 5" Zfand5
 1416084_at 0.93110 "zinc finger, AN1-type domain 5" Zfand5
 1416085_s_at 0.91674 "zinc finger, AN1-type domain 5" Zfand5
 1416086_at 0.92378 protein-tyrosine sulfotransferase 2 Tpst2
 1416087_at 0.98801 "adaptor protein complex AP-1, sigma 1" Ap1s1
 1416088_a_at 1.07240 ribosomal protein S15 Rps15
 1416089_at 1.06129 ribosomal protein S15 Rps15
 1416090_at 0.93791 pyruvate dehydrogenase (lipoamide) beta Pdhb
 1416091_at 0.94550 microtubule-associated protein 4 Mtap4
 1416092_a_at 1.01313 microtubule-associated protein 4 Mtap4
 1416093_a_at 1.05778 mitochondrial ribosomal protein L20 Mrpl20
 1416094_at 1.01017 a disintegrin and metallopeptidase domain 9 (meltrin gamma) Adam9
 1416095_x_at 1.00064 RIKEN cDNA 1110005A23 gene 1110005A23Rik
 1416096_at 0.82892 expressed sequence AI413782 AI413782

1416097_at 0.81474 leucine rich repeat containing 4 Lrrc4
1416098_at 0.87467 synaptogyrin 3 Syngr3
1416099_at 1.08280 ribosomal protein L27 /// similar to ribosomal protein L27 /// similar to
ribosomal protein L27 /// similar to ribosomal protein L27 Rpl27 /// LOC621100 /// LOC669376 ///
LOC675088
1416100_at 1.01548 "eukaryotic translation initiation factor 3, subunit 7 (zeta)" Eif3s7
1416101_a_at 1.45734 "histone 1, H1c" Hist1h1c
1416102_at 1.03868 "tyrosine 3-monooxygenase/tryptophan 5-monooxygenase activation protein,
zeta polypeptide" Ywhaz
1416103_at 0.94086 "tyrosine 3-monooxygenase/tryptophan 5-monooxygenase activation protein,
zeta polypeptide" Ywhaz
1416104_at 1.09495 mannose-P-dolichol utilization defect 1 Mpdu1
1416105_at 0.87860 nicotinamide nucleotide transhydrogenase Nnt
1416106_at 1.06621 "KTI12 homolog, chromatin associated (S. cerevisiae)" Kti12
1416107_at 1.06020 neuron specific gene family member 2 Nsg2
1416108_a_at 0.87234 transmembrane emp24 domain containing 3 Tmed3
1416109_at 0.99094 "fucosidase, alpha-L- 1, tissue" Fucal
1416110_at 0.94480 "solute carrier family 35, member A4" Slc35a4
1416111_at 1.09100 CD83 antigen Cd83
1416112_at 0.95158 "cytochrome c oxidase, subunit VIIIa" Cox8a
1416113_at 0.92207 FK506 binding protein 8 Fkbp8
1416114_at 0.93761 "SPARC-like 1 (mast9, hevin)" Sparcl1
1416115_at 0.96965 "origin recognition complex, subunit 3-like (S. cerevisiae)" Orc3l
1416116_at 1.00748 "origin recognition complex, subunit 3-like (S. cerevisiae)" Orc3l
1416117_at 0.88554 "origin recognition complex, subunit 3-like (S. cerevisiae)" Orc3l
1416118_at 0.90415 similar to mouse RING finger 1 LOC630539
1416119_at 1.07842 thioredoxin 1 Txn1
1416120_at 0.92009 ribonucleotide reductase M2 Rrm2
1416121_at 0.92703 lysyl oxidase Lox
1416122_at 1.00978 cyclin D2 Ccnd2
1416123_at 1.26487 cyclin D2 Ccnd2
1416124_at 0.93315 cyclin D2 Ccnd2
1416125_at 1.09269 FK506 binding protein 5 Fkbp5
1416126_at 1.16469 similar to DNA-directed RNA polymerase I 135 kDa polypeptide (RNA
polymerase I subunit 2) (RPA135) LOC671429
1416127_a_at 0.92440 aspartyl aminopeptidase Dnpep
1416128_at 0.97614 "tubulin, alpha 6 /// similar to Tubulin alpha-6 chain (Alpha-tubulin 6)"
Tuba6 /// LOC626534
1416129_at 0.98611 ERBB receptor feedback inhibitor 1 Errfi1
1416130_at 0.83682 prion protein Prnp
1416131_s_at 0.89997 RIKEN cDNA C920006C10 gene C920006C10Rik
1416132_at 0.97578 RIKEN cDNA C920006C10 gene C920006C10Rik
1416133_at 0.94686 RIKEN cDNA C920006C10 gene C920006C10Rik

1416134_at 0.80465 amyloid beta (A4) precursor-like protein 1 Aplp1
 1416135_at 1.24173 apurinic/aprimidinic endonuclease 1 Apex1
 1416136_at 0.87170 matrix metalloproteinase 2 Mmp2
 1416137_at 0.99082 annexin A7 Anxa7
 1416138_at 1.04951 annexin A7 Anxa7
 1416139_at 0.96747 regenerating islet-derived 2 Reg2
 1416140_a_at 1.00153 DEAH (Asp-Glu-Ala-His) box polypeptide 30 Dhx30
 1416141_a_at 1.02854 ribosomal protein S6 /// similar to 40S ribosomal protein S6 /// similar to 40S
 ribosomal protein S6 /// similar to 40S ribosomal protein S6 /// similar to 40S ribosomal protein S6
 Rps6 /// LOC666577 /// LOC667739 /// LOC671306 /// LOC672620
 1416142_at 0.99252 ribosomal protein S6 /// similar to 40S ribosomal protein S6 /// similar to 40S
 ribosomal protein S6 /// similar to 40S ribosomal protein S6 /// similar to 40S ribosomal protein S6 ///
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 ribosomal protein S6 Rps6 /// LOC214738 /// LOC384383 /// LOC545640 /// LOC626259 ///
 LOC629575 /// LOC638497 /// LOC639593 /// LOC639750 /// LOC666577 /// LOC667739 ///
 LOC667922 /// LOC671306 /// LOC671432 /// LOC672620 /// LOC675985
 1416143_at 0.92147 "ATP synthase, H⁺ transporting, mitochondrial F₀ complex, subunit F" Atp5j
 1416144_a_at 0.97593 DEAH (Asp-Glu-Ala-His) box polypeptide 15 Dhx15
 1416145_at 0.98794 DEAH (Asp-Glu-Ala-His) box polypeptide 15 Dhx15
 1416146_at 1.21220 heat shock protein 4 Hspa4
 1416147_at 0.77617 heat shock protein 4 Hspa4
 1416148_at 0.89898 lysosomal-associated protein transmembrane 4B Laptm4b
 1416149_at 0.89536 oligodendrocyte transcription factor 1 Olig1
 1416150_a_at 1.02496 "splicing factor, arginine/serine-rich 3 (SRp20)" Sfrs3
 1416151_at 0.81073 "splicing factor, arginine/serine-rich 3 (SRp20)" Sfrs3
 1416152_a_at 0.85609 "splicing factor, arginine/serine-rich 3 (SRp20)" Sfrs3
 1416153_at 0.96994 signal recognition particle 54 /// similar to signal recognition particle 54
 Srp54 /// LOC665155
 1416154_at 0.88600 signal recognition particle 54 /// similar to signal recognition particle 54
 Srp54 /// LOC665155
 1416155_at 1.06168 high mobility group box 3 Hmgb3
 1416156_at 0.97932 vinculin Vcl
 1416157_at 1.06980 vinculin Vcl
 1416158_at 0.81758 "nuclear receptor subfamily 2, group F, member 2" Nr2f2
 1416159_at 0.86312 "nuclear receptor subfamily 2, group F, member 2" Nr2f2
 1416160_at 0.72729 "nuclear receptor subfamily 2, group F, member 2" Nr2f2
 1416161_at 1.02763 RAD21 homolog (*S. pombe*) Rad21
 1416162_at 0.99156 RAD21 homolog (*S. pombe*) Rad21
 1416163_at 1.01366 "COP9 (constitutive photomorphogenic) homolog, subunit 4 (*Arabidopsis
 thaliana*)" Cops4

1416164_at	0.86994	fibulin 5	Fbln5	
1416165_at	0.96258	"RAB31, member RAS oncogene family"	Rab31	
1416166_a_at	0.88484	peroxiredoxin 4	Prdx4	
1416167_at	0.95316	peroxiredoxin 4	Prdx4	
1416168_at	0.96949	"serine (or cysteine) peptidase inhibitor, clade F, member 1"	Serpinf1	
1416169_at	0.91226	trophoblast specific protein beta	Tpbpb	
1416170_at	0.99033	TNF receptor-associated protein 1	Trap1	
1416171_at	1.03027	RIKEN cDNA 2310037I24 gene	2310037I24Rik	
1416172_at	1.14496	"pescadillo homolog 1, containing BRCT domain (zebrafish)"	Pes1	
1416173_at	0.91860	"pescadillo homolog 1, containing BRCT domain (zebrafish)"	Pes1	
1416174_at	0.72802	retinoblastoma binding protein 9	Rbbp9	
1416175_a_at	0.92261	voltage-dependent anion channel 3	Vdac3	
1416176_at	0.95440	high mobility group box 1	Hmgb1	
1416177_at	1.05308	"RNA binding motif protein, X chromosome retrogene"	Rbmxt	
1416178_a_at	0.90555	"pleckstrin homology domain containing, family B (evectins) member 1"	Plekha1	
1416179_a_at	0.90916	radixin	Rdx	
1416180_a_at	0.86517	radixin	Rdx	
1416181_at	0.89304	mesoderm development candiate 2	Mesdc2	
1416182_at	0.87703	"amyloid beta (A4) precursor protein-binding, family A, member 3"	Apba3	
1416183_a_at	0.96896	lactate dehydrogenase B	Ldhb	
1416184_s_at	1.34221	high mobility group AT-hook 1	Hmga1	
1416185_a_at	1.00643	"alcohol dehydrogenase 5 (class III), chi polypeptide"	Adh5	
1416186_at	0.95572	proline-rich nuclear receptor coactivator 2	Pnrc2	
1416187_s_at	0.88446	proline-rich nuclear receptor coactivator 2	Pnrc2	
1416188_at	0.98931	GM2 ganglioside activator protein	Gm2a	
1416189_a_at	0.92835	Sec61 alpha 1 subunit (<i>S. cerevisiae</i>)	Sec61a1	
1416190_a_at	0.92609	Sec61 alpha 1 subunit (<i>S. cerevisiae</i>)	Sec61a1	
1416191_at	0.91205	Sec61 alpha 1 subunit (<i>S. cerevisiae</i>)	Sec61a1	
1416192_at	0.88513	N-ethylmaleimide sensitive fusion protein attachment protein alpha	Napa	
1416193_at	0.96605	carbonic anhydrase 1	Car1	
1416194_at	0.99130	"cytochrome P450, family 4, subfamily b, polypeptide 1"	Cyp4b1	
1416195_at	1.09258	putative phosphatase	RP23-136K12.4	
1416196_at	0.96390	ribosomal protein SA	Rpsa	
1416197_at	0.99299	U1 small nuclear ribonucleoprotein C	Snrpc	
1416198_at	0.99129	TH1-like homolog (<i>Drosophila</i>)	Th11	
1416199_at	0.94901	kinesin family member C3	Kifc3	
1416200_at	1.10855	RIKEN cDNA 9230117N10 gene	9230117N10Rik	
1416201_at	0.98053	v-crk sarcoma virus CT10 oncogene homolog (avian)	Crk	
1416202_at	1.09773	prohibitin 2	Phb2	
1416203_at	0.91208	aquaporin 1	Aqp1	
1416204_at	0.84436	glycerol-3-phosphate dehydrogenase 1 (soluble)	Gpd1	
1416205_at	0.84241	"galactosidase, beta 1"	Glb1	

1416206_at 0.97811 signal-induced proliferation associated gene 1 Sipal1
1416207_at 0.92043 tafazzin Taz
1416208_at 1.10705 ubiquitin specific peptidase 14 Usp14
1416209_at 1.05988 glutamate dehydrogenase 1 Glud1
1416210_at 1.10355 "IMP3, U3 small nucleolar ribonucleoprotein, homolog (yeast)" Imp3
1416211_a_at 0.92987 pleiotrophin Ptn
1416212_at 1.04955 "mago-nashi homolog, proliferation-associated (Drosophila)" Magoh
1416213_x_at 0.97927 surfeit gene 4 Surf4
1416214_at 0.96488 minichromosome maintenance deficient 4 homolog (S. cerevisiae) Mcm4
1416215_at 0.89205 golgi SNAP receptor complex member 1 Gosr1
1416216_at 1.10470 RalBP1 associated Eps domain containing protein Repls1
1416217_a_at 0.98780 ribosomal protein L37a Rpl37a
1416218_x_at 0.98688 ribosomal protein L37a /// similar to 60S ribosomal protein L37a /// similar to
60S ribosomal protein L37a /// similar to 60S ribosomal protein L37a /// similar to 60S ribosomal
protein L37a Rpl37a /// LOC629949 /// LOC640991 /// LOC668935 /// LOC674834
1416219_at 1.04591 ribosomal protein L19 Rpl19
1416220_at 1.01054 signal peptidase complex subunit 1 homolog (S. cerevisiae) Spcs1
1416221_at 0.74506 follistatin-like 1 Fstl1
1416222_at 1.48491 NAD(P) dependent steroid dehydrogenase-like Nsdhl
1416223_at 0.89669 SH3 binding domain protein 5 like Sh3bp5l
1416224_at 0.99291 zinc finger and BTB domain containing 17 Zbtb17
1416225_at 0.73128 alcohol dehydrogenase 1 (class I) Adh1
1416226_at 1.18326 "actin related protein 2/3 complex, subunit 1B" Arpc1b
1416227_at 1.08340 "actin related protein 2/3 complex, subunit 1B /// similar to Actin-related protein
2/3 complex subunit 1B (ARP2/3 complex 41 kDa subunit) (p41-ARC) /// similar to Actin-related
protein 2/3 complex subunit 1B (ARP2/3 complex 41 kDa subunit) (p41-ARC)" Arpc1b ///
LOC434782 /// LOC671552
1416228_at 1.08119 protein (peptidyl-prolyl cis/trans isomerase) NIMA-interacting 1 Pin1
1416229_at 0.98596 riboflavin kinase Rfk
1416230_at 0.98781 riboflavin kinase Rfk
1416231_at 1.09806 Vac14 homolog (S. cerevisiae) Vac14
1416232_at 1.02742 oligodendrocyte transcription factor 2 Olig2
1416233_at 1.11291 "eukaryotic translation initiation factor 3, subunit 2 (beta)" Eif3s2
1416234_at 1.10591 leucine rich repeat containing 59 Lrrc59
1416235_at 0.99419 leucine rich repeat containing 59 Lrrc59
1416236_a_at 0.85599 epithelial V-like antigen 1 Eva1
1416237_at 0.92988 epithelial V-like antigen 1 Eva1
1416238_at 1.12227 tyrosine kinase receptor 1 Tie1
1416239_at 0.41578 argininosuccinate synthetase 1 Ass1
1416240_at 1.17063 "proteasome (prosome, macropain) subunit, beta type 7" Psmb7
1416241_at 1.04216 SEC13-like 1 (S. cerevisiae) Sec13l1
1416242_at 0.97322 kelch-like 13 (Drosophila) Khl13
1416243_a_at 1.07996 ribosomal protein L35 Rpl35

1416244_a_at 1.10804 cellular nucleic acid binding protein Cnbp
1416245_at 1.01037 aurora kinase A interacting protein 1 Aurkaip1
1416246_a_at 1.09781 "coronin, actin binding protein 1A" Coro1a
1416247_at 1.03763 dynactin 3 Dctn3
1416248_at 1.07606 NAD kinase Nadk
1416249_at 1.04109 NAD kinase Nadk
1416250_at 0.79602 "B-cell translocation gene 2, anti-proliferative" Btg2
1416251_at 0.87763 "minichromosome maintenance deficient 6 (MIS5 homolog, S. pombe) (S. cerevisiae)" Mcm6
1416252_at 1.11419 serine/threonine kinase 38 Stk38
1416253_at 1.07887 "cyclin-dependent kinase inhibitor 2D (p19, inhibits CDK4)" Cdkn2d
1416254_a_at 0.96270 vacuolar protein sorting 16 (yeast) Vps16
1416255_at 0.87833 gap junction membrane channel protein alpha 4 Gja4
1416256_a_at 0.93787 "tubulin, beta 5" Tubb5
1416257_at 0.98767 calpain 2 Capn2
1416258_at 1.08359 thymidine kinase 1 Tk1
1416259_at 1.04910 peroxisomal biogenesis factor 12 Pex12
1416260_a_at 1.02885 sorting nexin 1 Snx1
1416261_at 0.99088 transmembrane protein 19 Tmem19
1416262_at 0.94687 transmembrane protein 19 Tmem19
1416263_at 0.99710 "ATP-binding cassette, sub-family B (MDR/TAP), member 9" Abcb9
1416264_at 0.94004 "ATP-binding cassette, sub-family B (MDR/TAP), member 9" Abcb9
1416265_at 1.05137 calpain 10 Capn10
1416266_at 1.02284 prodynorphin Pdyn
1416267_at 0.99321 short coiled-coil protein Scoc
1416268_at 1.00177 "E26 avian leukemia oncogene 2, 3' domain" Ets2
1416269_at 0.97164 "ATP synthase, H+ transporting, mitochondrial F0 complex, subunit f, isoform 2" Atp5j2
1416270_at 1.00277 polymerase (RNA) II (DNA directed) polypeptide G Polr2g
1416271_at 0.81231 "PERP, TP53 apoptosis effector" Perp
1416272_at 1.02000 mitogen-activated protein kinase kinase 1 interacting protein 1 Map2k1ip1
1416273_at 1.04496 "tumor necrosis factor, alpha-induced protein 2" Tnfaip2
1416274_at 0.91552 "cystinosis, nephropathic" Ctns
1416275_at 0.82794 "solute carrier family 26, member 6" Slc26a6
1416276_a_at 1.01962 "ribosomal protein S4, X-linked" Rps4x
1416277_a_at 1.03822 "ribosomal protein, large, P1" Rplp1
1416278_a_at 0.96879 "ATP synthase, H+ transporting, mitochondrial F1 complex, O subunit" Atp5o
1416279_at 0.98862 "adaptor protein complex AP-1, beta 1 subunit" Ap1b1
1416280_at 1.00888 ubiquitin-like 1 (sentrin) activating enzyme E1B Uble1b
1416281_at 0.96273 Wdr45 like Wdr45l
1416282_at 1.03421 "proteasome (prosome, macropain) 26S subunit, ATPase 3" Psmc3
1416283_at 1.08828 phosphoribosylglycinamide formyltransferase Gart

1416284_at	1.04986	mitochondrial ribosomal protein L28	Mrpl28	
1416285_at	0.91207	"NADH dehydrogenase (ubiquinone) 1, subcomplex unknown, 1"	Ndufc1	
1416286_at	1.00871	regulator of G-protein signaling 4	Rgs4	
1416287_at	0.86961	regulator of G-protein signaling 4	Rgs4	
1416288_at	1.02291	"DnaJ (Hsp40) homolog, subfamily A, member 1"	Dnaja1	
1416289_at	0.82882	"procollagen-lysine, 2-oxoglutarate 5-dioxygenase 1"	Plod1	
1416290_a_at	1.03812	"proteasome (prosome, macropain) 26S subunit, ATPase, 4"	Psmc4	
1416291_at	1.14696	"proteasome (prosome, macropain) 26S subunit, ATPase, 4"	Psmc4	
1416292_at	1.03437	peroxiredoxin 3	Prdx3	
1416293_at	0.89365	nuclear factor I/B	Nfib	
1416294_at	1.00637	secretory carrier membrane protein 3	Scamp3	
1416295_a_at	0.94375	"interleukin 2 receptor, gamma chain"	Il2rg	
1416296_at	1.17229	"interleukin 2 receptor, gamma chain"	Il2rg	
1416297_s_at	0.88660	pancreatitis-associated protein	Pap	
1416298_at	1.04079	matrix metalloproteinase 9	Mmp9	
1416299_at	0.99283	Shc SH2-domain binding protein 1	Shcbp1	
1416300_a_at	0.98619	"solute carrier family 25 (mitochondrial carrier, phosphate carrier), member 3"	Slc25a3	
1416301_a_at	1.00021	early B-cell factor 1	Ebf1	
1416302_at	0.90449	early B-cell factor 1	Ebf1	
1416303_at	1.13145	LPS-induced TN factor	Litaf	
1416304_at	1.06914	LPS-induced TN factor	Litaf	
1416305_at	1.03448	SH3-domain binding protein 4	Sh3bp4	
1416306_at	1.00949	chloride channel calcium activated 3	Clca3	
1416307_at	0.92068	"adaptor-related protein complex AP-1, mu subunit 1"	Ap1m1	
1416308_at	1.10612	UDP-glucose dehydrogenase	Ugdh	
1416309_at	1.04627	nucleolar and spindle associated protein 1	Nusap1	
1416310_at	0.83236	nucleolar and spindle associated protein 1	Nusap1	
1416311_s_at	0.97872	"tubulin, alpha 3 /// tubulin, alpha 7 /// similar to tubulin, alpha 1"	Tuba3 /// Tuba7 /// LOC384954	
1416312_at	1.12364	arginyl-tRNA synthetase	Rars	
1416313_at	1.03889	"myeloid/lymphoid or mixed-lineage leukemia (trithorax homolog, Drosophila); translocated to, 11"	Mllt11	
1416314_at	0.93593	---	---	
1416315_at	0.88522	abhydrolase domain containing 4	Abhd4	
1416316_at	1.13752	"solute carrier family 27 (fatty acid transporter), member 2"	Slc27a2	
1416317_a_at	0.88775	Stam binding protein	Stamp	
1416318_at	1.14764	"serine (or cysteine) peptidase inhibitor, clade B, member 1a"	Serpib1a	
1416319_at	1.00873	adenosine kinase	Adk	
1416320_at	0.85750	SEC22 vesicle trafficking protein-like A (S. cerevisiae)	Sec22a	
1416321_s_at	0.59518	proline arginine-rich end leucine-rich repeat	Prelp	
1416322_at	0.77549	proline arginine-rich end leucine-rich repeat	Prelp	
1416323_at	1.00720	potassium channel tetramerisation domain containing 20	Kctd20	

1416324_s_at 1.01149 potassium channel tetramerisation domain containing 20 Kctd20
1416325_at 1.06040 cysteine-rich secretory protein 1 Crisp1
1416326_at 0.86145 cysteine-rich protein 1 (intestinal) Crip1
1416327_at 1.02121 ubiquitin-fold modifier conjugating enzyme 1 Ufc1
1416328_a_at 0.94643 "ATPase, H+ transporting, lysosomal V0 subunit E" Atp6v0e
1416329_at 1.02315 cytoplasmic FMR1 interacting protein 1 Cyfip1
1416330_at 0.88142 CD 81 antigen Cd81
1416331_a_at 1.02178 "nuclear factor, erythroid derived 2,-like 1" Nfe2l1
1416332_at 0.80769 cold inducible RNA binding protein Cirbp
1416333_at 1.01796 docking protein 2 Dok2
1416334_at 0.79182 WW domain-containing oxidoreductase Wwox
1416335_at 1.25528 macrophage migration inhibitory factor Mif
1416336_s_at 1.14845 small nuclear ribonucleoprotein D1 Snrpd1
1416337_at 0.98571 ubiquinol-cytochrome c reductase binding protein Uqcrb
1416338_at 0.94805 SH3-domain GRB2-like 1 Sh3gl1
1416339_a_at 0.94129 protein kinase C substrate 80K-H PrkcsH
1416340_a_at 0.90368 "mannosidase 2, alpha B1" Man2b1
1416341_at 1.00045 polymerase (RNA) II (DNA directed) polypeptide C Polr2c
1416342_at 0.98939 tenascin C Tnc
1416343_a_at 0.81424 lysosomal membrane glycoprotein 2 Lamp2
1416344_at 0.84626 lysosomal membrane glycoprotein 2 Lamp2
1416345_at 1.10302 translocase of inner mitochondrial membrane 8 homolog a1 (yeast)
Timm8a1
1416346_at 1.22749 translocase of inner mitochondrial membrane 8 homolog a1 (yeast)
Timm8a1
1416347_at 1.05395 multiple endocrine neoplasia 1 Men1
1416348_at 1.04936 multiple endocrine neoplasia 1 Men1
1416349_at 1.01825 mitochondrial ribosomal protein L34 Mrpl34
1416350_at 1.01203 Kruppel-like factor 16 Klf16
1416351_at 1.06905 mitogen activated protein kinase kinase 1 Map2k1
1416352_s_at 1.11885 "glycoprotein, synaptic 2" Gpsn2
1416353_at 0.95077 "nuclear receptor subfamily 1, group H, member 2" Nr1h2
1416354_at 1.00889 "RNA binding motif protein, X chromosome" Rbmx
1416355_at 0.95185 "RNA binding motif protein, X chromosome" Rbmx
1416356_at 1.04101 guanosine monophosphate reductase 2 Gmpr2
1416357_a_at 0.52523 melanoma cell adhesion molecule Mcam
1416358_at 0.98938 RIKEN cDNA 0610009O03 gene 0610009O03Rik
1416359_at 0.97515 sorting nexin associated golgi protein 1 Snag1
1416360_at 0.88164 sorting nexin associated golgi protein 1 Snag1
1416361_a_at 1.07156 dynein cytoplasmic 1 intermediate chain 1 Dync1i1
1416362_a_at 0.93889 FK506 binding protein 4 Fkbp4
1416363_at 1.09409 FK506 binding protein 4 Fkbp4
1416364_at 1.04513 "heat shock protein 90kDa alpha (cytosolic), class B member 1" Hsp90ab1

1416365_at 0.96688 "heat shock protein 90kDa alpha (cytosolic), class B member 1" Hsp90ab1
1416366_at 1.00314 "similar to NADH dehydrogenase (ubiquinone) 1, subcomplex unknown, 2"
LOC675851
1416367_at 0.98889 RIKEN cDNA 1110001J03 gene 1110001J03Rik
1416368_at 1.03616 "glutathione S-transferase, alpha 4" Gsta4
1416369_at 0.96597 hippocampus abundant transcript-like 1 Hiatl1
1416370_at 0.90081 zinc finger proliferation 1 Zipro1
1416371_at 1.02338 apolipoprotein D Apod
1416372_at 0.99518 phosphatidylserine synthase 1 Ptdss1
1416373_at 0.93435 nitrogen fixation gene 1 (S. cerevisiae) Nfs1
1416374_at 1.02921 "adaptor-related protein complex 3, mu 1 subunit /// similar to AP-3 complex
subunit mu-1 (Adapter-related protein complex 3 mu-1 subunit) (Mu-adaptin 3A) (AP-3 adapter
complex mu3A subunit)" Ap3m1 /// LOC671800
1416375_at 0.91771 "adaptor-related protein complex 3, mu 1 subunit" Ap3m1
1416376_at 1.37028 transmembrane protein 97 Tmem97
1416377_at 1.07184 programmed cell death protein 7 Pcd7
1416378_at 0.92109 polynucleotide kinase 3'-phosphatase Pnkp
1416379_at 0.91273 pannexin 1 Panx1
1416380_at 0.93766 Moloney leukemia virus 10 Mov10
1416381_a_at 1.08466 peroxiredoxin 5 Prdx5
1416382_at 1.07202 cathepsin C Ctsc
1416383_a_at 1.00516 pyruvate carboxylase Pcx
1416384_a_at 1.08373 "coatamer protein complex, subunit epsilon" Cope
1416385_a_at 0.95796 "mannose-6-phosphate receptor, cation dependent" M6pr
1416386_a_at 0.95448 "mannose-6-phosphate receptor, cation dependent" M6pr
1416387_at 0.93803 "phosphatidylinositol-4-phosphate 5-kinase, type II, gamma" Pip5k2c
1416388_at 1.01502 "phosphatidylinositol-4-phosphate 5-kinase, type II, gamma" Pip5k2c
1416389_a_at 0.93089 regulator of chromosome condensation (RCC1) and BTB (POZ) domain
containing protein 2 Rcbtb2
1416390_at 0.74556 regulator of chromosome condensation (RCC1) and BTB (POZ) domain
containing protein 2 Rcbtb2
1416391_at 0.78992 pentatricopeptide repeat domain 1 Ptdc1
1416392_a_at 1.06728 "ATPase, H+ transporting, lysosomal V0 subunit C" Atp6v0c
1416393_at 1.18707 EMG1 nucleolar protein homolog (S. cerevisiae) Emg1
1416394_at 0.99970 Bcl2-associated athanogene 1 Bag1
1416395_at 1.01594 guanylate kinase 1 Guk1
1416396_at 0.95893 --- ---
1416397_at 0.89055 mesoderm development candidate 1 Mesdc1
1416398_at 1.02785 mesoderm development candidate 1 Mesdc1
1416399_a_at 0.93752 heme oxygenase (decycling) 2 Hmox2
1416400_at 0.98677 pyrroline-5-carboxylate reductase-like Pycrl
1416401_at 1.13154 CD82 antigen Cd82
1416402_at 0.92549 "ATP-binding cassette, sub-family B (MDR/TAP), member 10" Abcb10

1416403_at	0.95994	"ATP-binding cassette, sub-family B (MDR/TAP), member 10"	Abcb10
1416404_s_at	1.04365	ribosomal protein S16	Rps16
1416405_at	0.81875	biglycan	Bgn
1416406_at	0.96304	phosphoprotein enriched in astrocytes 15	Pea15
1416407_at	0.98715	phosphoprotein enriched in astrocytes 15	Pea15
1416408_at	1.11020	"acyl-Coenzyme A oxidase 1, palmitoyl"	Acox1
1416409_at	1.12796	"acyl-Coenzyme A oxidase 1, palmitoyl"	Acox1
1416410_at	1.09152	"platelet-activating factor acetylhydrolase, isoform 1b, alpha1 subunit"	Pafah1b3
1416411_at	0.89157	"glutathione S-transferase, mu 2"	Gstm2
1416412_at	0.82962	neutral sphingomyelinase (N-SMase) activation associated factor	Nsmaf
1416413_at	0.91539	cathepsin J	Ctsj
1416414_at	0.95550	elastin microfibril interfacier 1	Emilin1
1416415_a_at	1.00416	"H2A histone family, member Z /// similar to H2A histone family, member Z /// similar to H2A histone family, member Z"	H2afz /// LOC666634 /// LOC669445
1416416_x_at	0.83799	"glutathione S-transferase, mu 1"	Gstm1
1416417_a_at	0.95134	"NADH dehydrogenase (ubiquinone) 1 beta subcomplex, 7"	Ndufb7
1416418_at	0.99349	gamma-aminobutyric acid (GABA(A)) receptor-associated protein-like 1	Gabarapl1
1416419_s_at	0.98745	gamma-aminobutyric acid (GABA(A)) receptor-associated protein-like 1	Gabarapl1
1416420_a_at	1.03566	ribosomal protein L9 /// similar to ribosomal protein L9	Rpl9 /// LOC637379
1416421_a_at	1.13279	Sjogren syndrome antigen B	Ssb
1416422_a_at	1.00583	Sjogren syndrome antigen B	Ssb
1416423_x_at	1.08813	Sjogren syndrome antigen B	Ssb
1416424_at	0.94790	mannose-6-phosphate receptor binding protein 1	M6prbp1
1416425_at	1.00141	peroxisome biogenesis factor 19	Pex19
1416426_at	0.92956	"RAB5A, member RAS oncogene family"	Rab5a
1416427_at	0.88148	cyclin I	Ccni
1416428_at	0.93179	THAP domain containing 11	Thap11
1416429_a_at	0.81538	catalase	Cat
1416430_at	0.83613	catalase	Cat
1416431_at	0.98635	"tubulin, beta 6"	Tubb6
1416432_at	1.17522	"6-phosphofructo-2-kinase/fructose-2,6-biphosphatase 3"	Pfkfb3
1416433_at	0.93164	replication protein A2	Rpa2
1416434_at	0.96778	Bcl2-like 10	Bcl2l10
1416435_at	0.95215	lymphotoxin B receptor	Ltbr
1416436_a_at	0.98623	RIKEN cDNA 2410003P15 gene	2410003P15Rik
1416437_a_at	1.00370	mitogen-activated protein kinase 8 interacting protein 3	Mapk8ip3
1416438_at	0.93607	RIKEN cDNA 2410104I19 gene	2410104I19Rik
1416439_at	1.12551	RIKEN cDNA 2410015N17 gene	2410015N17Rik
1416440_at	0.82622	CD164 antigen	Cd164

1416441_at 1.06814 plasma glutamate carboxypeptidase Pgcg
 1416442_at 0.63882 immediate early response 2 Ier2
 1416443_a_at 1.00363 ubiquitin-like 1 (sentrin) activating enzyme E1A Uble1a
 1416444_at 0.84979 "elongation of very long chain fatty acids (FEN1/Elo2, SUR4/Elo3, yeast)-like 2" Elovl2
 1416445_at 0.96943 RIKEN cDNA 2810405J04 gene 2810405J04Rik
 1416446_at 1.00813 transmembrane protein 30A Tmem30a
 1416447_at 0.94352 transmembrane protein 30A Tmem30a
 1416448_at 0.99091 inosine triphosphatase (nucleoside triphosphate pyrophosphatase) Itpa
 1416449_x_at 1.02086 syntaxin binding protein 2 Stxbp2
 1416450_at 1.03262 taube nuss Tbn
 1416451_s_at 1.08058 taube nuss Tbn
 1416452_at 0.91379 ornithine aminotransferase Oat
 1416453_x_at 1.08466 ribosomal protein S12 /// similar to ribosomal protein S12 /// similar to ribosomal protein S12 /// similar to ribosomal protein S12 /// similar to ribosomal protein S12 /// similar to 40S ribosomal protein S12 /// similar to ribosomal protein S12 /// similar to ribosomal protein S12 Rps12 /// LOC432865 /// LOC627835 /// LOC668354 /// LOC670751 /// LOC670832 /// LOC671641 /// LOC672008 /// LOC672415 /// LOC676277
 1416454_s_at 0.61457 "actin, alpha 2, smooth muscle, aorta" Acta2
 1416455_a_at 0.91827 "crystallin, alpha B" Cryab
 1416456_a_at 0.91016 "chitinase, acidic" Chia
 1416457_at 0.90948 dimethylarginine dimethylaminohydrolase 2 Ddah2
 1416458_at 0.91022 ADP-ribosylation factor 2 Arf2
 1416459_at 0.82357 ADP-ribosylation factor 2 Arf2
 1416460_at 1.12455 myo-inositol oxygenase Miox
 1416461_at 0.99380 GPI-anchored membrane protein 1 Gpiap1
 1416462_at 0.97661 GPI-anchored membrane protein 1 Gpiap1
 1416463_at 1.03833 GPI-anchored membrane protein 1 Gpiap1
 1416464_at 0.94583 "solute carrier family 4 (anion exchanger), member 1" Slc4a1
 1416465_a_at 0.98448 "vesicle-associated membrane protein, associated protein A" Vapa
 1416466_at 0.99208 "vesicle-associated membrane protein, associated protein A" Vapa
 1416467_at 1.04646 "DEAD/H (Asp-Glu-Ala-Asp/His) box polypeptide 3, X-linked" Ddx3x
 1416468_at 0.89587 "aldehyde dehydrogenase family 1, subfamily A1" Aldh1a1
 1416469_at 0.95123 leucine zipper protein 1 Luzp1
 1416470_a_at 0.94391 ribophorin I Rpn1
 1416471_at 0.88665 choline/ethanolaminophosphotransferase 1 Cept1
 1416472_at 0.84509 synapse associated protein 1 Syap1
 1416473_a_at 0.86569 neighbor of Punc E11 Nope
 1416474_at 0.91376 neighbor of Punc E11 Nope
 1416475_at 0.98758 ubiquitin-conjugating enzyme E2D 2 Ube2d2
 1416476_a_at 0.95693 ubiquitin-conjugating enzyme E2D 2 Ube2d2
 1416477_at 0.90024 ubiquitin-conjugating enzyme E2D 2 Ube2d2

1416478_a_at 1.02834 "malate dehydrogenase 2, NAD (mitochondrial)" Mdh2
1416479_a_at 1.01952 transmembrane protein 14C Tmem14c
1416480_a_at 1.45071 "HIG1 domain family, member 1A" Higd1a
1416481_s_at 1.40610 "HIG1 domain family, member 1A" Higd1a
1416482_at 0.91468 tetratricopeptide repeat domain 3 Ttc3
1416483_at 0.94588 tetratricopeptide repeat domain 3 Ttc3
1416484_at 0.85326 tetratricopeptide repeat domain 3 Ttc3
1416485_at 1.02917 translocase of inner mitochondrial membrane 23 homolog (yeast) Timm23
1416486_at 1.04783 "small inducible cytokine subfamily E, member 1" Scye1
1416487_a_at 1.04403 yes-associated protein 1 Yap1
1416488_at 1.34148 cyclin G2 Ccng2
1416489_at 1.22085 phosphatidylinositol 4-kinase type 2 beta Pi4k2b
1416490_at 1.00874 transmembrane emp24 protein transport domain containing 6 Tmed6
1416491_at 0.83217 numb-like Numbl
1416492_at 0.95255 cyclin E1 Ccne1
1416493_at 0.92689 dolichyl-di-phosphooligosaccharide-protein glycotransferase Ddost
1416494_at 1.05169 NADH dehydrogenase (ubiquinone) Fe-S protein 5 Ndufs5
1416495_s_at 1.03084 NADH dehydrogenase (ubiquinone) Fe-S protein 5 Ndufs5
1416496_at 0.93254 Morf4 family associated protein 1 Mrfap1
1416497_at 0.81407 protein disulfide isomerase associated 4 Pdia4
1416498_at 1.07059 peptidylprolyl isomerase C Ppic
1416499_a_at 0.92206 dynactin 6 Dctn6
1416500_at 0.98750 "SAC1 (suppressor of actin mutations 1, homolog)-like (S. cerevisiae)" Sacm11
1416501_at 0.96566 3-phosphoinositide dependent protein kinase-1 Pdpk1
1416502_a_at 0.99099 prolactin regulatory element binding Preb
1416503_at 0.94745 latexin Lxn
1416504_at 0.73921 Unc-51 like kinase 1 (C. elegans) Ulk1
1416505_at 0.44269 "nuclear receptor subfamily 4, group A, member 1" Nr4a1
1416506_at 1.09266 "proteasome (prosome, macropain) subunit, alpha type 6" Psma6
1416507_at 1.01548 transmembrane protein 4 Tmem4
1416508_at 1.02050 "mediator of RNA polymerase II transcription, subunit 28 homolog (yeast)"
Med28
1416509_at 0.99728 transmembrane 9 superfamily member 3 Tm9sf3
1416510_at 0.98652 mitochondrial ribosomal protein L4 Mrpl4
1416511_a_at 0.84138 CDC42 effector protein (Rho GTPase binding) 4 Cdc42ep4
1416512_at 0.99409 nucleotide binding protein 2 Nubp2
1416513_at 0.86860 "laminin, beta 2" Lamb2
1416514_a_at 1.05666 "fascin homolog 1, actin bundling protein (Strongylocentrotus purpuratus)"
Fscn1
1416515_at 1.05721 "fascin homolog 1, actin bundling protein (Strongylocentrotus purpuratus)"
Fscn1
1416516_at 1.01914 "fascin homolog 1, actin bundling protein (Strongylocentrotus purpuratus)"
Fscn1

1416517_at 0.98434 patatin-like phospholipase domain containing 6 Pnpla6
1416518_at 0.91352 "H1 histone family, member O, oocyte-specific" H1foo
1416519_at 1.05474 similar to ribosomal protein L36 LOC622534
1416520_x_at 1.11528 similar to ribosomal protein L36 LOC622534
1416521_at 0.73403 "selenoprotein W, muscle 1" Sepw1
1416522_a_at 0.99776 "gene rich cluster, C10 gene" Grcc10
1416523_at 1.07423 "ribonuclease, RNase A family, 1 (pancreatic)" Rnase1
1416524_at 0.97918 speckle-type POZ protein Spop
1416525_at 1.04116 speckle-type POZ protein Spop
1416526_a_at 1.05380 "Parkinson disease (autosomal recessive, early onset) 7" Park7
1416527_at 1.68743 "RAB32, member RAS oncogene family" Rab32
1416528_at 0.89710 SH3 domain binding glutamic acid-rich protein-like 3 Sh3bgrl3
1416529_at 1.02152 epithelial membrane protein 1 Emp1
1416530_a_at 1.05853 purine-nucleoside phosphorylase Pnp
1416531_at 1.26952 glutathione S-transferase omega 1 Gsto1
1416532_at 1.00672 transformation/transcription domain-associated protein Trrap
1416533_at 1.00682 EGL nine homolog 2 (C. elegans) Egl2
1416534_at 0.92527 "D4, zinc and double PHD fingers family 2" Dpf2
1416535_at 0.93320 microspherule protein 1 /// similar to Microspherule protein 1 (58 kDa
microspherule protein) /// similar to Microspherule protein 1 (58 kDa microspherule protein) Mcrs1 ///
LOC433780 /// LOC666215
1416536_at 1.08951 melanoma associated antigen (mutated) 1 Mum1
1416537_at 0.90199 cysteine-rich with EGF-like domains 1 Creld1
1416538_at 0.73769 sialic acid acetyltransferase Sia
1416539_at 0.80629 sialic acid acetyltransferase Sia
1416540_at 0.97459 HGF-regulated tyrosine kinase substrate Hgs
1416541_at 0.96040 ClpB caseinolytic peptidase B homolog (E. coli) Clpb
1416542_at 0.83026 PHD finger protein 1 Phf1
1416543_at 0.97779 "nuclear factor, erythroid derived 2, like 2" Nfe2l2
1416544_at 0.97828 enhancer of zeste homolog 2 (Drosophila) Ezh2
1416545_at 0.94870 "zinc finger, DHHC domain containing 7" Zdhc7
1416546_a_at 1.05017 ribosomal protein L6 Rpl6
1416547_at 1.05058 NADH dehydrogenase (ubiquinone) 1 beta subcomplex 3 Ndufb3
1416548_at 0.97502 "solute carrier family 35, member B4" Slc35b4
1416549_at 0.92552 "solute carrier family 35, member B4" Slc35b4
1416550_at 0.95607 "solute carrier family 35, member B4" Slc35b4
1416551_at 0.90160 "ATPase, Ca⁺⁺ transporting, cardiac muscle, slow twitch 2" Atp2a2
1416552_at 1.04872 developmental pluripotency associated 5 Dppa5
1416553_at 1.05667 stimulated by retinoic acid 13 Stra13
1416554_at 0.94834 PDZ and LIM domain 1 (elfin) Pdlim1
1416555_at 1.00928 etoposide induced 2.4 mRNA Ei24
1416556_at 0.98025 tetraspanin 31 Tspan31
1416557_a_at 0.98509 elongation factor Tu GTP binding domain containing 2 Eftud2

1416558_at 1.03271 maternal embryonic leucine zipper kinase Melk
1416559_at 1.05411 RIKEN cDNA 1500003O22 gene 1500003O22Rik
1416560_at 0.90434 "solute carrier family 13 (sodium-dependent dicarboxylate transporter), member 3" Slc13a3
1416561_at 1.07628 glutamic acid decarboxylase 1 Gad1
1416562_at 1.02334 glutamic acid decarboxylase 1 Gad1
1416563_at 1.28908 cytidine 5'-triphosphate synthase Ctps
1416564_at 0.95013 SRY-box containing gene 7 Sox7
1416565_at 0.91237 "cytochrome c oxidase, subunit VIb polypeptide 1" Cox6b1
1416566_at 1.04473 serine/threonine kinase receptor associated protein Strap
1416567_s_at 0.98083 "ATP synthase, H+ transporting, mitochondrial F1 complex, epsilon subunit" Atp5e
1416568_a_at 1.06827 apoptotic chromatin condensation inducer 1 Acin1
1416569_at 1.01311 actin-like 6A Actl6a
1416570_s_at 0.99761 "G elongation factor, mitochondrial 1" Gfm1
1416571_at 0.94324 Tnf receptor associated factor 4 Traf4
1416572_at 0.82774 matrix metalloproteinase 14 (membrane-inserted) Mmp14
1416573_at 0.83634 protein O-fucosyltransferase 2 Pofut2
1416574_at 0.89993 RIKEN cDNA 5730589K01 gene 5730589K01Rik
1416575_at 0.97095 cell division cycle 45 homolog (S. cerevisiae)-like Cdc45l
1416576_at 0.98502 suppressor of cytokine signaling 3 Socs3
1416577_a_at 0.94395 ring-box 1 Rbx1
1416578_at 0.97288 ring-box 1 Rbx1
1416579_a_at 0.73870 tumor-associated calcium signal transducer 1 Tacstd1
1416580_a_at 1.05904 STIP1 homology and U-Box containing protein 1 Stub1
1416581_at 1.05016 WD repeat domain 5 Wdr5
1416582_a_at 0.94600 Bcl-associated death promoter Bad
1416583_at 0.99878 Bcl-associated death promoter Bad
1416584_at 0.90818 "mannosidase 2, alpha B2" Man2b2
1416585_at 1.10890 RuvB-like protein 1 Ruvbl1
1416586_at 0.81881 zinc finger protein 239 Zfp239
1416587_a_at 0.90698 X-ray repair complementing defective repair in Chinese hamster cells 1 Xrcc1
1416588_at 1.01442 "protein tyrosine phosphatase, receptor type, N /// similar to Receptor-type tyrosine-protein phosphatase-like N precursor (R-PTP-N) (PTP IA-2)" Ptprn /// LOC669060
1416589_at 0.88887 secreted acidic cysteine rich glycoprotein Sparc
1416590_a_at 1.07828 "RAB34, member of RAS oncogene family" Rab34
1416591_at 1.08336 "RAB34, member of RAS oncogene family" Rab34
1416592_at 0.64154 glutaredoxin Glrx
1416593_at 0.66405 glutaredoxin Glrx
1416594_at 1.08896 secreted frizzled-related sequence protein 1 Sfrp1
1416595_at 1.09657 mitochondrial ribosomal protein S22 Mrps22
1416596_at 1.04891 "solute carrier family 44, member 4" Slc44a4
1416597_at 1.02895 "hepatoma-derived growth factor, related protein 2" Hdgfrp2

1416598_at 0.85452 GLIS family zinc finger 2 Glis2
 1416599_at 0.93889 GLIS family zinc finger 2 Glis2
 1416600_a_at 0.82677 Down syndrome critical region homolog 1 (human) Dscr1
 1416601_a_at 0.81090 Down syndrome critical region homolog 1 (human) Dscr1
 1416602_a_at 1.02953 RAD52 homolog (S. cerevisiae) Rad52
 1416603_at 1.03195 ribosomal protein L22 /// similar to 60S ribosomal protein L22 (Heparin-binding protein HBp15) /// similar to 60S ribosomal protein L22 (Heparin-binding protein HBp15) /// similar to 60S ribosomal protein L22 (Heparin binding protein HBp15) /// similar to 60S ribosomal protein L22 (Heparin binding protein HBp15) Rpl22 /// LOC623483 /// LOC627737 /// LOC632026 /// LOC634567
 1416604_at 0.99629 cytochrome c-1 Cyc1
 1416605_at 1.16122 "nucleolar protein family A, member 2" Nola2
 1416606_s_at 1.10235 "nucleolar protein family A, member 2" Nola2
 1416607_at 0.95028 RIKEN cDNA 4931406C07 gene 4931406C07Rik
 1416608_a_at 0.91555 cDNA sequence BC004004 BC004004
 1416609_at 0.90755 cDNA sequence BC004004 BC004004
 1416610_a_at 1.08475 chloride channel 3 Clcn3
 1416611_at 0.95541 secretory carrier membrane protein 2 Scamp2
 1416612_at 0.62304 "cytochrome P450, family 1, subfamily b, polypeptide 1" Cyp1b1
 1416613_at 0.85195 "cytochrome P450, family 1, subfamily b, polypeptide 1" Cyp1b1
 1416614_at 0.94822 CREBBP/EP300 inhibitory protein 1 Cri1
 1416615_at 1.04241 "caseinolytic peptidase, ATP-dependent, proteolytic subunit homolog (E. coli)" Clpp
 1416616_s_at 1.02782 "caseinolytic peptidase, ATP-dependent, proteolytic subunit homolog (E. coli)" Clpp
 1416617_at 0.93314 acyl-CoA synthetase short-chain family member 1 Acss1
 1416618_at 0.91677 protoporphyrinogen oxidase Ppox
 1416619_at 0.90966 RIKEN cDNA 4632428N05 gene 4632428N05Rik
 1416620_at 0.95716 "Swi/SNF related matrix associated, actin dependent regulator of chromatin, subfamily a-like 1" Smarcal1
 1416621_at 1.01255 lethal giant larvae homolog 1 (Drosophila) Llg1
 1416622_at 1.04436 Williams-Beuren syndrome chromosome region 16 homolog (human) Wbscr16
 1416623_at 0.78509 thrombospondin 3 Thbs3
 1416624_a_at 0.96933 ubiquitin A-52 residue ribosomal protein fusion product 1 /// similar to ubiquitin A-52 residue ribosomal protein fusion product 1 /// similar to ubiquitin A-52 residue ribosomal protein fusion product 1 Uba52 /// LOC666586 /// LOC669193 /// LOC676687
 1416625_at 1.07558 "serine (or cysteine) peptidase inhibitor, clade G, member 1" Serping1
 1416626_at 1.06514 "phospholipase A2, group IB, pancreas" Pla2g1b
 1416627_at 0.90011 "serine protease inhibitor, Kunitz type 1" Spint1
 1416628_at 1.15112 RIKEN cDNA 0610006I08 gene 0610006I08Rik
 1416629_at 1.19165 "solute carrier family 1 (neutral amino acid transporter), member 5" Slc1a5
 1416630_at 0.35327 inhibitor of DNA binding 3 Id3
 1416631_at 1.11790 "adaptor-related protein complex AP-4, beta 1" Ap4b1

1416632_at 1.14614 "malic enzyme, supernatant /// similar to NADP-dependent malic enzyme (NADP-ME) (Malic enzyme 1) /// similar to NADP-dependent malic enzyme (NADP-ME) (Malic enzyme 1)" Mod1 /// LOC624892 /// LOC677317
 1416633_a_at 0.93579 RIKEN cDNA 5730536A07 gene 5730536A07Rik
 1416634_at 1.09858 RIKEN cDNA 5730536A07 gene 5730536A07Rik
 1416635_at 1.05073 "sphingomyelin phosphodiesterase, acid-like 3A" Smpdl3a
 1416636_at 1.06985 RAS-homolog enriched in brain Rheb
 1416637_at 0.92163 "solute carrier family 4 (anion exchanger), member 2" Slc4a2
 1416638_at 0.76280 sal-like 2 (Drosophila) Sall2
 1416639_at 0.89079 "solute carrier family 2 (facilitated glucose transporter), member 5" Slc2a5
 1416640_at 1.00154 "potassium voltage-gated channel, Isk-related family, member 1-like" Kcne11
 1416641_at 0.97450 "ligase I, DNA, ATP-dependent" Lig1
 1416642_a_at 0.99540 "tumor protein, translationally-controlled 1" Tpt1
 1416643_at 1.04954 "tumor protein, translationally-controlled 1" Tpt1
 1416644_a_at 0.86655 "sema domain, immunoglobulin domain (Ig), short basic domain, secreted, (semaphorin) 3B" Sema3b
 1416645_a_at 1.31487 alpha fetoprotein Afp
 1416646_at 1.18711 alpha fetoprotein Afp
 1416647_at 0.84100 "branched chain ketoacid dehydrogenase E1, alpha polypeptide" Bckdha
 1416648_at 1.00055 dynein cytoplasmic 1 heavy chain 1 Dync1h1
 1416649_at 1.02246 alpha 1 microglobulin/bikunin Ambp
 1416650_at 1.00728 ret finger protein-like 4 Rfp14
 1416651_at 1.05219 "zinc finger, HIT domain containing 2" Znhit2
 1416652_at 0.93734 asporin Asp
 1416653_at 1.06963 syntaxin binding protein 3A Stxbp3a
 1416654_at 1.05772 "solute carrier family 31, member 2" Slc31a2
 1416655_at 0.94959 C1GALT1-specific chaperone 1 C1galt1c1
 1416656_at 1.01584 chloride intracellular channel 1 Clic1
 1416657_at 0.98023 thymoma viral proto-oncogene 1 Akt1
 1416658_at 0.98776 frizzled-related protein Frzb
 1416659_at 0.99862 "eukaryotic translation initiation factor 3, subunit 10 (theta)" Eif3s10
 1416660_at 1.01806 "eukaryotic translation initiation factor 3, subunit 10 (theta)" Eif3s10
 1416661_at 1.06766 "eukaryotic translation initiation factor 3, subunit 10 (theta)" Eif3s10
 1416662_at 0.80424 sarcosine dehydrogenase Sardh
 1416663_at 1.03088 "NADH dehydrogenase (ubiquinone) 1 alpha subcomplex, 9" Ndufa9
 1416664_at 1.06229 cell division cycle 20 homolog (S. cerevisiae) Cdc20
 1416665_at 1.04143 demethyl-Q 7 Coq7
 1416666_at 1.06737 "serine (or cysteine) peptidase inhibitor, clade E, member 2" Serpine2
 1416667_at 1.11835 phenylalkylamine Ca²⁺ antagonist (emopamil) binding protein Ebp
 1416668_at 0.99523 RIKEN cDNA 4921531G14 gene 4921531G14Rik
 1416669_s_at 1.14385 nascent polypeptide-associated complex alpha polypeptide Naca
 1416670_at 0.86539 "SET domain, bifurcated 1" Setdb1
 1416671_a_at 0.83821 mucolipin 1 Mcoln1

1416672_s_at 1.04333 mucolipin 1 Mcoln1
1416673_at 0.97168 beta-site APP-cleaving enzyme 2 Bace2
1416674_at 0.91298 "protein tyrosine phosphatase, receptor type, U" Ptpnu
1416675_s_at 0.63014 "phospholipase C, delta 1" Plcd1
1416676_at 0.90949 kininogen 1 Kng1
1416677_at 0.87623 apolipoprotein H Apoh
1416678_at 0.99544 "COP9 (constitutive photomorphogenic) homolog, subunit 3 (Arabidopsis thaliana)" Cops3
1416679_at 0.92463 "ATP-binding cassette, sub-family D (ALD), member 3" Abcd3
1416680_at 0.95093 ubiquitin protein ligase E3A Ube3a
1416681_at 0.88834 ubiquitin protein ligase E3A Ube3a
1416682_at 1.01651 ubiquitin protein ligase E3A Ube3a
1416683_at 1.01536 plexin B2 Plxnb2
1416684_at 1.24200 fibrillarlin Fbl
1416685_s_at 1.28945 fibrillarlin Fbl
1416686_at 0.97087 "procollagen lysine, 2-oxoglutarate 5-dioxygenase 2" Plod2
1416687_at 0.94244 "procollagen lysine, 2-oxoglutarate 5-dioxygenase 2" Plod2
1416688_at 1.03779 synaptosomal-associated protein 91 Snap91
1416689_at 1.01663 tuftelin 1 Tuft1
1416690_at 1.07668 GTP binding protein 2 Gtpbp2
1416691_at 0.98984 GTP binding protein 2 Gtpbp2
1416692_at 0.94615 coilin Coil
1416693_at 1.01803 forkhead box C2 Foxc2
1416694_at 1.12658 glycerol kinase 2 Gk2
1416695_at 0.98808 translocator protein Tspo
1416696_at 0.98184 "DNA segment, Chr 17, Wayne State University 104, expressed" D17Wsu104e
1416697_at 0.98076 dipeptidylpeptidase 4 Dpp4
1416698_a_at 1.00276 CDC28 protein kinase 1b Cks1b
1416699_at 0.91448 RIKEN cDNA 1110008F13 gene 1110008F13Rik
1416700_at 0.97012 Rho family GTPase 3 Rnd3
1416701_at 0.99765 Rho family GTPase 3 Rnd3
1416702_at 0.96176 "serine (or cysteine) peptidase inhibitor, clade I, member 1" Serpini1
1416703_at 1.02684 mitogen activated protein kinase 14 Mapk14
1416704_at 0.99573 mitogen activated protein kinase 14 Mapk14
1416705_at 0.95461 ribulose-5-phosphate-3-epimerase Rpe
1416706_at 1.05844 ribulose-5-phosphate-3-epimerase Rpe
1416707_a_at 1.09101 polyamine-modulated factor 1 Pmf1
1416708_a_at 0.93916 GRAM domain containing 1A Gramd1a
1416709_a_at 1.00493 "neugrin, neurite outgrowth associated" Ngrn
1416710_at 1.02823 transmembrane protein 35 Tmem35
1416711_at 0.92387 T-box brain gene 1 Tbr1
1416712_at 0.92553 peptidase D Pepd
1416713_at 0.84989 RIKEN cDNA 2700055K07 gene 2700055K07Rik

1416714_at 0.98201 interferon regulatory factor 8 Irf8
1416715_at 1.11765 gap junction membrane channel protein beta 3 Gjb3
1416716_at 0.99921 embryonal Fyn-associated substrate Efs
1416717_at 1.10834 cysteine-rich secretory protein 2 Crisp2
1416718_at 0.93582 brevican Bcan
1416719_a_at 1.04957 ribosomal protein S10 /// similar to ribosomal protein S10 /// similar to
ribosomal protein S10 /// similar to ribosomal protein S10 /// similar to ribosomal protein S10 Rps10 ///
LOC621772 /// LOC636539 /// LOC666274 /// LOC668457
1416720_at 1.01935 "splicing factor, arginine/serine-rich 6" Sfrs6
1416721_s_at 0.99860 "splicing factor, arginine/serine-rich 6" Sfrs6
1416722_at 0.95921 high mobility group 20A Hmg20a
1416723_at 0.83891 transcription factor 4 Tcf4
1416724_x_at 0.87046 transcription factor 4 Tcf4
1416725_at 0.91444 transcription factor 4 Tcf4
1416726_s_at 1.03743 ubiquitin-conjugating enzyme E2S Ube2s
1416727_a_at 0.97593 cytochrome b-5 Cyb5
1416728_at 1.07815 "casein kinase 2, beta polypeptide" Csnk2b
1416729_at 1.25957 plasminogen Plg
1416730_at 1.09753 RNA terminal phosphate cyclase-like 1 Rcl1
1416731_at 1.07248 topoisomerase (DNA) II beta Top2b
1416732_at 1.00255 topoisomerase (DNA) II beta Top2b
1416733_at 1.02686 "muskelin 1, intracellular mediator containing kelch motifs" Mkl1
1416734_at 0.92267 "muskelin 1, intracellular mediator containing kelch motifs" Mkl1
1416735_at 1.00443 N-acylsphingosine amidohydrolase 1 Asah1
1416736_at 1.01116 cancer susceptibility candidate 3 Casc3
1416737_at 1.04108 --- ---
1416738_at 0.98622 BRCA1 associated protein Brap
1416739_a_at 0.86902 BRCA1 associated protein Brap
1416740_at 0.82789 "procollagen, type V, alpha 1" Col5a1
1416741_at 1.08124 "Procollagen, type V, alpha 1" Col5a1
1416742_at 1.04242 craniofacial development protein 1 Cfdp1
1416743_at 1.06508 UDP-N-acetylglucosamine pyrophosphorylase 1 /// similar to UDP-N-
acetylhexosamine pyrophosphorylase Uap1 /// LOC640502
1416744_at 0.98751 UDP-N-acetylglucosamine pyrophosphorylase 1 Uap1
1416745_x_at 1.00070 UDP-N-acetylglucosamine pyrophosphorylase 1 Uap1
1416746_at 1.00626 "H2A histone family, member X" H2afx
1416747_at 1.02266 "DNA Segment, Chr 15, Mouse Genome Informatics 27" D15Mgi27
1416748_a_at 1.01523 meiotic recombination 11 homolog A (S. cerevisiae) Mre11a
1416749_at 0.70129 HtrA serine peptidase 1 Htra1
1416750_at 1.36854 "opioid receptor, sigma 1" Oprs1
1416751_a_at 1.02417 DEAD (Asp-Glu-Ala-Asp) box polypeptide 20 Ddx20
1416752_at 1.05716 LIM domain binding 3 Ldb3
1416753_at 0.94733 "protein kinase, cAMP dependent regulatory, type I beta" Prkar1b

1416754_at 0.97027 "protein kinase, cAMP dependent regulatory, type I beta" Prkar1b
1416755_at 0.94739 "DnaJ (Hsp40) homolog, subfamily B, member 1" Dnajb1
1416756_at 0.89780 "DnaJ (Hsp40) homolog, subfamily B, member 1" Dnajb1
1416757_at 1.11857 "Zwilch, kinetochore associated, homolog (Drosophila)" Zwilch
1416758_at 1.02026 Rho GTPase activating protein 27 Arhgap27
1416759_at 0.87089 "microtubule associated monooxygenase, calponin and LIM domain containing 1" Mical1
1416760_at 1.02165 UDP-N-acetyl-alpha-D-galactosamine:polypeptide N-acetylgalactosaminyltransferase-like 1 Galnt11
1416761_at 0.97148 --- ---
1416762_at 1.11428 S100 calcium binding protein A10 (calpactin) S100a10
1416763_at 1.08441 RIKEN cDNA G431001I09 gene G431001I09Rik
1416764_at 1.02666 fission 1 (mitochondrial outer membrane) homolog (yeast) Fis1
1416765_s_at 1.02309 mitochondria-associated protein involved in granulocyte-macrophage colony-stimulating factor signal transduction Magmas
1416766_at 1.02848 MOCO sulphurase C-terminal domain containing 2 Mosc2
1416767_a_at 1.00023 RIKEN cDNA 1110003E01 gene 1110003E01Rik
1416768_at 0.96525 RIKEN cDNA 1110003E01 gene 1110003E01Rik
1416769_s_at 0.98629 "ATPase, H+ transporting, lysosomal V0 subunit B" Atp6v0b
1416770_at 0.89332 serine/threonine kinase 25 (yeast) Stk25
1416771_at 1.11310 trafficking protein particle complex 3 Trappc3
1416772_at 0.84761 carnitine palmitoyltransferase 2 Cpt2
1416773_at 0.89011 wee 1 homolog (S. pombe) Wee1
1416774_at 0.82291 wee 1 homolog (S. pombe) Wee1
1416775_at 0.85484 RIKEN cDNA 2310004L02 gene 2310004L02Rik
1416776_at 0.99687 "crystallin, mu" Crym
1416777_at 1.05565 CEA-related cell adhesion molecule 12 Ceacam12
1416778_at 0.83101 serum deprivation response Sdpr
1416779_at 0.85289 serum deprivation response Sdpr
1416780_at 0.83601 "phosphofructokinase, muscle" Pfkf
1416781_at 0.95155 PRA1 domain family 2 Praf2
1416782_s_at 0.92330 PRA1 domain family 2 Praf2
1416783_at 1.06036 tachykinin 1 Tac1
1416784_at 0.96815 phosphoglycerate kinase 2 Pgk2
1416785_at 0.97702 Kv channel-interacting protein 1 Kcnip1
1416786_at 0.82715 "activin A receptor, type 1" Acvr1
1416787_at 0.85847 "activin A receptor, type 1" Acvr1
1416788_a_at 1.00066 "isocitrate dehydrogenase 3 (NAD+), gamma" Idh3g
1416789_at 1.08669 "isocitrate dehydrogenase 3 (NAD+), gamma" Idh3g
1416790_a_at 1.09350 thymine DNA glycosylase /// similar to thymine DNA glycosylase isoform 2 /// similar to G/T mismatch-specific thymine DNA glycosylase (C-JUN leucine zipper interactive protein JZA-3) /// similar to thymine DNA glycosylase isoform 2 /// similar to G/T mismatch-specific thymine DNA glycosylase (C-JUN leucine zipper interactive protein JZA-3) Tdg /// LOC545124 ///

LOC624784 /// LOC670176 /// LOC672659

1416791_a_at 1.02181 nuclear RNA export factor 1 homolog (*S. cerevisiae*) Nxf1
1416792_at 0.97602 "protein phosphatase 1G (formerly 2C), magnesium-dependent, gamma isoform" Ppm1g
1416793_at 1.03986 ADP-ribosylation factor-like 6 interacting protein 2 Arl6ip2
1416794_at 0.98574 ADP-ribosylation factor-like 6 interacting protein 2 Arl6ip2
1416795_at 0.89625 "crystallin, lamda 1" Cry11
1416796_at 0.97636 non-catalytic region of tyrosine kinase adaptor protein 2 Nck2
1416797_at 0.80755 non-catalytic region of tyrosine kinase adaptor protein 2 Nck2
1416798_a_at 0.97632 "expressed in non-metastatic cells 4, protein" Nme4
1416799_at 0.98634 "transient receptor potential cation channel, subfamily M, member 7" Trpm7
1416800_at 1.04744 "transient receptor potential cation channel, subfamily M, member 7" Trpm7
1416801_at 1.10823 "transient receptor potential cation channel, subfamily M, member 7" Trpm7
1416802_a_at 0.98810 cell division cycle associated 5 Cdc5
1416803_at 0.89674 FK506 binding protein 7 Fkbp7
1416804_at 1.23284 RIKEN cDNA G430002G23 gene G430002G23Rik
1416805_at 1.00494 RIKEN cDNA 1110032E23 gene 1110032E23Rik
1416806_at 1.01049 ferredoxin reductase Fdxr
1416807_at 1.19304 ribosomal protein L36a Rpl36a
1416808_at 1.09554 nidogen 1 Nid1
1416809_at 1.00262 "cytochrome P450, family 3, subfamily a, polypeptide 11" Cyp3a11
1416810_at 1.15467 male enhanced antigen 1 Mea1
1416811_s_at 1.03685 cytotoxic T lymphocyte-associated protein 2 alpha /// cytotoxic T lymphocyte-associated protein 2 beta Ctla2a /// Ctla2b
1416812_at 1.02316 cytotoxic granule-associated RNA binding protein 1 Tia1
1416813_at 1.00388 cytotoxic granule-associated RNA binding protein 1 Tia1
1416814_at 0.97669 cytotoxic granule-associated RNA binding protein 1 Tia1
1416815_s_at 1.01494 budding uninhibited by benzimidazoles 3 homolog (*S. cerevisiae*) Bub3
1416816_at 1.01197 NIMA (never in mitosis gene a)-related expressed kinase 7 Nek7
1416817_at 1.11463 NIMA (never in mitosis gene a)-related expressed kinase 7 Nek7
1416818_at 1.00052 "parvin, alpha" Parva
1416819_at 1.08534 cell division cycle 37 homolog (*S. cerevisiae*) Cdc37
1416820_at 0.94690 cell division cycle 37 homolog (*S. cerevisiae*) Cdc37
1416821_at 0.99375 expressed sequence 2 embryonic lethal Es2el
1416822_at 0.89974 expressed sequence 2 embryonic lethal Es2el
1416823_a_at 0.74918 oxysterol binding protein-like 1A Osbpl1a
1416824_at 1.01969 RIKEN cDNA B230118H07 gene B230118H07Rik
1416825_at 0.90557 "syntrophin, acidic 1" Snta1
1416826_a_at 0.96894 Trf (TATA binding protein-related factor)-proximal protein homolog (*Drosophila*) Trfp
1416827_at 1.00246 "thromboxane A synthase 1, platelet" Tbxas1
1416828_at 1.00198 synaptosomal-associated protein 25 Snap25
1416829_at 0.91470 "ATP synthase, H⁺ transporting mitochondrial F1 complex, beta subunit" Atp5b

1416830_at 1.02899 RIKEN cDNA 0610031J06 gene 0610031J06Rik
1416831_at 1.02659 neuraminidase 1 Neu1
1416832_at 0.70950 "solute carrier family 39 (metal ion transporter), member 8" Slc39a8
1416833_at 1.07824 kidney expressed gene 1 Keg1
1416834_x_at 0.97566 "NADH dehydrogenase (ubiquinone) 1 beta subcomplex, 2" Ndufb2
1416835_s_at 1.21369 S-adenosylmethionine decarboxylase 1 /// S-adenosylmethionine decarboxylase
2 Amd1 /// Amd2
1416836_at 0.95754 low-density lipoprotein receptor-related protein 10 Lrp10
1416837_at 1.29242 Bcl2-associated X protein Bax
1416838_at 0.95433 methylmalonyl-Coenzyme A mutase Mut
1416839_at 1.03424 methylmalonyl-Coenzyme A mutase Mut
1416840_at 0.83629 Mid1 interacting protein 1 (gastrulation specific G12-like (zebrafish)) Mid1ip1
1416841_at 0.96036 RIKEN cDNA 1110059E24 gene 1110059E24Rik
1416842_at 0.99976 "glutathione S-transferase, mu 5" Gstm5
1416843_at 0.96618 "phosphodiesterase 6D, cGMP-specific, rod, delta" Pde6d
1416844_at 0.92217 protein arginine N-methyltransferase 2 Prmt2
1416845_at 0.86278 transmembrane protein 132A Tmem132a
1416846_a_at 0.75816 PDZ domain containing RING finger 3 Pdzn3
1416847_s_at 1.01385 2'-5' oligoadenylate synthetase 1D /// 2'-5' oligoadenylate synthetase 1E
Oas1d /// Oas1e
1416848_at 1.04207 ubiquitin-like 5 Ubl5
1416849_at 0.94036 "zinc finger, CDGSH-type domain 1" Zcd1
1416850_s_at 0.90633 "zinc finger, CDGSH-type domain 1" Zcd1
1416851_at 0.95070 Suppression of tumorigenicity 13 St13
1416852_a_at 0.86136 neurochondrin Ncdn
1416853_at 1.00312 neurochondrin Ncdn
1416854_at 0.91906 "solute carrier family 34 (sodium phosphate), member 2" Slc34a2
1416855_at 1.04679 growth arrest specific 1 Gas1
1416856_at 0.99718 RIKEN cDNA 3230401D17 gene 3230401D17Rik
1416857_at 1.00772 stromal cell derived factor 2 Sdf2
1416858_a_at 1.01783 FK506 binding protein 3 Fkbp3
1416859_at 1.04532 FK506 binding protein 3 Fkbp3
1416860_s_at 0.96737 "inhibitor of growth family, member 1" Ing1
1416861_at 1.03338 signal transducing adaptor molecule (SH3 domain and ITAM motif) 1 Stam
1416862_at 0.90065 signal transducing adaptor molecule (SH3 domain and ITAM motif) 1 Stam
1416863_at 0.84775 abhydrolase domain containing 8 Abhd8
1416864_at 1.02234 surfeit gene 6 Surf6
1416865_at 0.94308 "FYVE, RhoGEF and PH domain containing 1" Fgd1
1416866_at 0.82146 blocked early in transport 1 homolog (S. cerevisiae) Bet1
1416867_at 0.91702 blocked early in transport 1 homolog (S. cerevisiae) Bet1
1416868_at 0.78180 "cyclin-dependent kinase inhibitor 2C (p18, inhibits CDK4)" Cdkn2c
1416869_x_at 1.05328 Lck interacting transmembrane adaptor 1 Lime1
1416870_at 0.81887 "dynein, axonemal, light chain 4" Dnalc4

1416871_at 1.14690 a disintegrin and metallopeptidase domain 8 Adam8
1416872_at 1.08142 tetraspanin 6 Tspan6
1416873_a_at 0.96235 cyclin-dependent kinase 2 Cdk2
1416874_a_at 0.98873 "Paf1, RNA polymerase II associated factor, homolog (S. cerevisiae)" Paf1
1416875_at 0.92709 "parvin, gamma" Parvg
1416876_at 0.84861 "parvin, gamma" Parvg
1416877_a_at 0.97652 mitochondrial ribosomal protein L51 Mrpl51
1416878_at 1.02338 mitochondrial ribosomal protein L51 Mrpl51
1416879_at 1.03455 mitochondrial ribosomal protein L51 Mrpl51
1416880_at 0.86039 myeloid cell leukemia sequence 1 Mcl1
1416881_at 0.93346 myeloid cell leukemia sequence 1 Mcl1
1416882_at 0.83310 regulator of G-protein signalling 10 Rgs10
1416883_at 0.92175 cleft lip and palate associated transmembrane protein 1 Clptm1
1416884_at 0.99460 chromobox homolog 3 (Drosophila HP1 gamma) /// similar to chromobox homolog 3 /// similar to chromobox homolog 3 Cbx3 /// LOC629578 /// LOC672750
1416885_at 0.95094 RIKEN cDNA 1110038F14 gene 1110038F14Rik
1416886_at 1.11418 nuclear DNA binding protein C1d
1416887_at 0.94350 nuclear DNA binding protein C1d
1416888_at 1.02172 Fas (TNFRSF6)-associated via death domain Fadd
1416889_at 0.91713 "troponin I, skeletal, fast 2" Tnni2
1416890_at 1.05584 WD repeat domain 74 Wdr74
1416891_at 0.98704 numb gene homolog (Drosophila) Numb
1416892_s_at 1.06051 RIKEN cDNA 3110001A13 gene 3110001A13Rik
1416893_at 0.95746 RIKEN cDNA 3110001A13 gene 3110001A13Rik
1416894_at 0.91819 RIKEN cDNA 1500019G21 gene 1500019G21Rik
1416895_at 1.14136 ephrin A1 Efna1
1416896_at 0.88359 ribosomal protein S6 kinase polypeptide 1 Rps6ka1
1416897_at 1.18071 "poly (ADP-ribose) polymerase family, member 9" Parp9
1416898_a_at 0.97517 interferon regulatory factor 3 Irf3
1416899_at 0.93247 undifferentiated embryonic cell transcription factor 1 Utf1
1416900_s_at 0.95626 growth differentiation factor 1 /// longevity assurance homolog 1 (S. cerevisiae) Gdf1 /// Lass1
1416901_at 1.11139 Niemann Pick type C2 Npc2
1416902_a_at 0.95923 "cytochrome c oxidase, subunit Vb" Cox5b
1416903_at 0.83601 nucleobindin 1 Nucb1
1416904_at 0.89105 muscleblind-like 1 (Drosophila) Mbnl1
1416905_at 0.73845 guanylate cyclase activator 2a (guanylin) Guca2a
1416906_at 0.91797 anaphase-promoting complex subunit 5 Anapc5
1416907_at 1.00594 translin Tsn
1416908_s_at 0.97231 translin Tsn
1416909_at 0.93686 RIKEN cDNA 1810008A14 gene 1810008A14Rik
1416910_at 1.02667 "DnaJ (Hsp40) homolog, subfamily C, member 15" Dnajc15
1416911_a_at 0.98253 RIKEN cDNA 6330407G11 gene 6330407G11Rik

1416912_at 1.00036 RIKEN cDNA 6330407G11 gene 6330407G11Rik
1416913_at 1.05754 esterase 1 Es1
1416914_s_at 0.84362 mammary tumor virus receptor 2 Mtvr2
1416915_at 1.03420 mutS homolog 6 (E. coli) Msh6
1416916_at 1.46524 E74-like factor 3 Elf3
1416917_at 0.93571 RIKEN cDNA 1700123O20 gene 1700123O20Rik
1416918_at 0.79264 "discs, large homolog 3 (Drosophila)" Dlgh3
1416919_a_at 0.92827 nephronophthisis 1 (juvenile) homolog (human) Nphp1
1416920_at 1.08215 RNA binding motif protein 4 Rbm4
1416921_x_at 1.11776 "aldolase 1, A isoform" Aldoa
1416922_a_at 1.22290 BCL2/adenovirus E1B interacting protein 3-like Bnip3l
1416923_a_at 1.14484 BCL2/adenovirus E1B interacting protein 3-like Bnip3l
1416924_at 0.99505 brain protein I3 Bri3
1416925_at 1.14399 karyopherin (importin) beta 1 Kpnb1
1416926_at 0.88702 transformation related protein 53 inducible nuclear protein 1 Trp53inp1
1416927_at 0.97579 transformation related protein 53 inducible nuclear protein 1 Trp53inp1
1416928_at 1.00191 RNA binding motif protein 12 Rbm12
1416929_at 1.02085 RNA binding motif protein 12 Rbm12
1416930_at 1.20245 "lymphocyte antigen 6 complex, locus D" Ly6d
1416931_at 1.03057 Ngg1 interacting factor 3-like 1 (S. pombe) Nif3l1
1416932_at 0.98060 component of oligomeric golgi complex 2 Cog2
1416933_at 0.97414 P450 (cytochrome) oxidoreductase Por
1416934_at 0.90958 X-linked myotubular myopathy gene 1 Mtm1
1416935_at 0.91906 "transient receptor potential cation channel, subfamily V, member 2" Trpv2
1416936_at 0.86390 apoptosis-associated tyrosine kinase Aatk
1416937_at 0.95585 gamma-aminobutyric acid receptor associated protein Gabarap
1416938_at 1.00522 coiled-coil-helix-coiled-coil-helix domain containing 1 Chchd1
1416939_at 1.04761 pyrophosphatase (inorganic) 1 Ppa1
1416940_at 1.04017 peptidylprolyl isomerase F (cyclophilin F) Ppif
1416941_s_at 0.96303 Williams-Beuren syndrome chromosome region 1 homolog (human) Wbscr1
1416942_at 1.08674 type 1 tumor necrosis factor receptor shedding aminopeptidase regulator Arts1
1416943_at 1.12063 "ubiquitin-conjugating enzyme E2E 1, UBC4/5 homolog (yeast) /// similar to Ubiquitin-conjugating enzyme E2 E1 (Ubiquitin-protein ligase E1) (Ubiquitin carrier protein E1) (UbcM3)" Ube2e1 /// LOC631407
1416944_a_at 0.98966 tousled-like kinase 2 (Arabidopsis) Tlk2
1416945_at 0.93943 prostate tumor over expressed gene 1 Ptov1
1416946_a_at 0.83161 acetyl-Coenzyme A acyltransferase 1A /// acetyl-Coenzyme A acyltransferase 1B Acaa1a /// Acaa1b
1416947_s_at 0.95906 acetyl-Coenzyme A acyltransferase 1A /// acetyl-Coenzyme A acyltransferase 1B Acaa1a /// Acaa1b
1416948_at 1.02267 mitochondrial ribosomal protein L23 Mrpl23
1416949_s_at 0.99247 "solute carrier family 39 (zinc transporter), member 7" Slc39a7
1416950_at 1.02579 "tumor necrosis factor, alpha-induced protein 8" Tnfaip8

1416951_a_at	1.04920	"ATPase, H ⁺ transporting, lysosomal V1 subunit D"	Atp6v1d
1416952_at	1.10003	"ATPase, H ⁺ transporting, lysosomal V1 subunit D"	Atp6v1d
1416953_at	0.72628	connective tissue growth factor	Ctgf
1416954_at	1.02612	"solute carrier family 25 (mitochondrial carrier, dicarboxylate transporter), member 10"	Slc25a10
1416955_at	1.00329	"solute carrier family 25 (mitochondrial carrier, dicarboxylate transporter), member 10"	Slc25a10
1416956_at	0.91779	"potassium voltage-gated channel, shaker-related subfamily, beta member 2"	Kcnab2
1416957_at	0.94587	"POU domain, class 2, associating factor 1"	Pou2af1
1416958_at	0.89899	"nuclear receptor subfamily 1, group D, member 2"	Nr1d2
1416959_at	0.84209	"nuclear receptor subfamily 1, group D, member 2"	Nr1d2
1416960_at	0.97753	"beta-1,3-glucuronyltransferase 3 (glucuronosyltransferase I)"	B3gat3
1416961_at	0.92311	"budding uninhibited by benzimidazoles 1 homolog, beta (S. cerevisiae)"	Bub1b
1416962_at	1.05217	regulator of chromosome condensation 1	Rcc1
1416963_at	0.94231	ubiquitin associated domain containing 1	Ubadc1
1416964_at	0.96286	"eukaryotic elongation factor, selenocysteine-tRNA-specific"	Eefsec
1416965_at	0.95382	proprotein convertase subtilisin/kexin type 1 inhibitor	Pcsk1n
1416966_at	0.84985	"solute carrier family 22 (organic anion transporter), member 8"	Slc22a8
1416967_at	1.01355	SRY-box containing gene 2	Sox2
1416968_a_at	0.98523	"hydroxy-delta-5-steroid dehydrogenase, 3 beta- and steroid delta-isomerase 7"	Hsd3b7
1416969_at	1.06044	G two S phase expressed protein 1	Gtse1
1416970_a_at	0.92075	"cytochrome c oxidase, subunit VIIa 2"	Cox7a2
1416971_at	0.96989	"cytochrome c oxidase, subunit VIIa 2"	Cox7a2
1416972_at	1.03865	NHP2 non-histone chromosome protein 2-like 1 (S. cerevisiae) /// similar to Nhp2 non-histone chromosome protein 2-like 1 /// similar to NHP2-like protein 1 (High mobility group-like nuclear protein 2 homolog 1) ([U4/U6.U5] tri-snRNP 15.5 kDa protein) (Sperm specific antigen 1) (Fertilization antigen 1) (FA-1)	Nhp211 /// LOC434401 /// LOC637555
1416973_at	1.10895	NHP2 non-histone chromosome protein 2-like 1 (S. cerevisiae)	Nhp211
1416974_at	0.97165	signal transducing adaptor molecule (SH3 domain and ITAM motif) 2	Stam2
1416975_at	0.99770	signal transducing adaptor molecule (SH3 domain and ITAM motif) 2	Stam2
1416976_at	0.88449	signal transducing adaptor molecule (SH3 domain and ITAM motif) 2	Stam2
1416977_at	1.17914	signal transducing adaptor molecule (SH3 domain and ITAM motif) 2	Stam2
1416978_at	0.62751	"Fc receptor, IgG, alpha chain transporter"	Fcgrt
1416979_at	1.06282	proteasome maturation protein	Pomp
1416980_at	1.23981	methyltransferase like 7B	Mettl7b
1416981_at	0.80669	forkhead box O1	Foxo1
1416982_at	0.76186	forkhead box O1	Foxo1
1416983_s_at	0.76468	forkhead box O1	Foxo1
1416984_at	0.97059	mitochondrial ribosomal protein S18A	Mrps18a
1416985_at	0.85156	signal-regulatory protein alpha	Sirpa
1416986_a_at	0.96188	signal-regulatory protein alpha	Sirpa

1416987_at 1.06256 elongation protein 4 homolog (S. cerevisiae) Elp4
1416988_at 0.94243 mutS homolog 2 (E. coli) Msh2
1416989_at 0.93921 vacuolar protein sorting 53 (yeast) Vps53
1416990_at 0.99088 retinoid X receptor beta Rxrb
1416991_at 0.95611 mitochondrial translation optimization 1 homolog (S. cerevisiae) Mto1
1416992_at 0.89363 manic fringe homolog (Drosophila) Mfng
1416993_at 0.89816 component of oligomeric golgi complex 4 Cog4
1416994_at 0.99012 tetratricopeptide repeat domain 1 Ttc1
1416995_at 0.90980 protein kinase C and casein kinase substrate in neurons 3 Pacsin3
1416996_at 0.95654 "TBC1 domain family, member 8" Tbc1d8
1416997_a_at 0.89801 huntingtin-associated protein 1 Hap1
1416998_at 1.09021 RRS1 ribosome biogenesis regulator homolog (S. cerevisiae) Rrs1
1416999_at 0.94789 "sphingomyelin phosphodiesterase 2, neutral" Smpd2
1417000_at 0.61385 ankyrin repeat and BTB (POZ) domain containing 1 Abtb1
1417001_a_at 1.06898 "DNA segment, Chr 4, Wayne State University 53, expressed" D4Wsu53e
1417002_at 0.96104 RIKEN cDNA 0610012G03 gene 0610012G03Rik
1417003_at 1.13587 RIKEN cDNA 0610012G03 gene /// hypothetical protein LOC638521
0610012G03Rik /// LOC638521
1417004_at 0.96929 tumor susceptibility gene 101 Tsg101
1417005_at 1.09930 kinesin 2 Kns2
1417006_at 1.14155 COMM domain containing 4 Commd4
1417007_a_at 0.99093 vacuolar protein sorting 4b (yeast) Vps4b
1417008_at 0.89886 carnitine acetyltransferase Crat
1417009_at 1.31110 "complement component 1, r subcomponent" C1r
1417010_at 1.00700 zinc finger protein 238 Zfp238
1417011_at 0.76459 syndecan 2 Sdc2
1417012_at 0.85134 syndecan 2 Sdc2
1417013_at 0.76005 heat shock protein 8 Hspb8
1417014_at 0.77810 heat shock protein 8 Hspb8
1417015_at 0.98438 Ras association (RalGDS/AF-6) domain family 3 Rassf3
1417016_at 0.99974 MAP kinase-activated protein kinase 5 Mapkapk5
1417017_at 0.97016 "cytochrome P450, family 17, subfamily a, polypeptide 1" Cyp17a1
1417018_at 0.88887 epidermal growth factor-containing fibulin-like extracellular matrix protein 2
Efemp2
1417019_a_at 1.01243 cell division cycle 6 homolog (S. cerevisiae) Cdc6
1417020_at 0.96508 spermatogenesis associated 4 Spata4
1417021_a_at 0.99851 "sporulation protein, meiosis-specific, SPO11 homolog (S. cerevisiae) ///
similar to Meiotic recombination protein SPO11" Spo11 /// LOC675982
1417022_at 0.95514 "solute carrier family 7 (cationic amino acid transporter, y+ system), member
3" Slc7a3
1417023_a_at 0.97944 "fatty acid binding protein 4, adipocyte" Fabp4
1417024_at 1.12587 histidyl-tRNA synthetase Hars
1417025_at 1.02054 "histocompatibility 2, class II antigen E beta" H2-Eb1

1417026_at 1.01313 prefoldin 1 Pfdn1
1417027_at 0.80583 tripartite motif protein 2 Trim2
1417028_a_at 0.77590 tripartite motif protein 2 Trim2
1417029_a_at 0.75009 tripartite motif protein 2 Trim2
1417030_at 0.98464 RIKEN cDNA 2310028N02 gene 2310028N02Rik
1417031_at 1.04584 RIKEN cDNA 2310028N02 gene 2310028N02Rik
1417032_at 0.99189 ubiquitin-conjugating enzyme E2G 2 Ube2g2
1417033_at 0.93968 ubiquitin-conjugating enzyme E2G 2 Ube2g2
1417034_at 1.09161 trafficking protein particle complex 6A Trappc6a
1417035_at 1.05758 SAC3 domain containing 1 Sac3d1
1417036_at 0.90431 SAC3 domain containing 1 Sac3d1
1417037_at 1.11534 "origin recognition complex, subunit 6-like (*S. cerevisiae*)" Orc6l
1417038_at 0.97498 septin 9 9-Sep
1417039_a_at 1.00904 cullin 7 Cul7
1417040_a_at 0.87859 Bcl-2-related ovarian killer protein Bok
1417041_at 1.14963 RNA polymerase 1-1 Rpo1-1
1417042_at 1.18417 "solute carrier family 37 (glycerol-6-phosphate transporter), member 4" Slc37a4
1417043_at 1.05738 lecithin cholesterol acyltransferase Lcat
1417044_at 0.89081 leucine carboxyl methyltransferase 1 Lcmt1
1417045_at 1.42615 BH3 interacting domain death agonist Bid
1417046_at 0.99179 tumor suppressor candidate 4 Tusc4
1417047_at 1.06024 prominin 2 Prom2
1417048_at 0.92857 prominin 2 Prom2
1417049_at 0.91138 "Rh blood group, D antigen" Rhd
1417050_at 0.94949 C1q and tumor necrosis factor related protein 4 C1qtnf4
1417051_at 1.02550 protocadherin 8 Pcdh8
1417052_at 1.08888 "proteasome (prosome, macropain) subunit, beta type 3" Psmb3
1417053_at 1.09140 prohibitin /// similar to prohibitin /// similar to prohibitin /// similar to
prohibitin Phb /// LOC665511 /// LOC669197 /// LOC674314
1417054_a_at 1.01082 RIKEN cDNA 0610009D07 gene 0610009D07Rik
1417055_at 1.02668 RIKEN cDNA 0610009D07 gene 0610009D07Rik
1417056_at 1.13365 "proteasome (prosome, macropain) 28 subunit, alpha" Psmel
1417057_a_at 1.06849 peptidylprolyl isomerase D (cyclophilin D) /// lysosomal-associated membrane
protein 3 /// similar to peptidylprolyl isomerase D Ppid /// Lamp3 /// LOC671770
1417058_a_at 1.04418 keratinocyte associated protein 2 Krtcap2
1417059_at 1.06626 keratinocyte associated protein 2 Krtcap2
1417060_at 1.00651 "protein phosphatase 1, regulatory (inhibitor) subunit 11" Ppp1r11
1417061_at 0.65488 "solute carrier family 40 (iron-regulated transporter), member 1" Slc40a1
1417062_at 0.93323 RIKEN cDNA 2810037C14 gene 2810037C14Rik
1417063_at 1.08170 "complement component 1, q subcomponent, beta polypeptide" C1qb
1417064_at 1.08118 jagunal homolog 1 (*Drosophila*) Jagn1
1417065_at 0.41795 early growth response 1 Egr1
1417066_at 0.84945 "chaperone, ABC1 activity of bc1 complex like (*S. pombe*)" Cabcl

1417067_s_at 1.01069 "chaperone, ABC1 activity of bc1 complex like (S. pombe)" Cabcl
1417068_a_at 0.96966 "protein tyrosine phosphatase, non-receptor type 1" Ptpn1
1417069_a_at 0.98054 "glia maturation factor, beta" Gmfb
1417070_at 0.92826 "cytochrome P450, family 4, subfamily v, polypeptide 3" Cyp4v3
1417071_s_at 0.79261 "cytochrome P450, family 4, subfamily v, polypeptide 3" Cyp4v3
1417072_at 0.98709 "solute carrier family 22 (organic anion transporter), member 6" Slc22a6
1417073_a_at 0.97526 quaking Qk
1417074_at 1.02746 CEA-related cell adhesion molecule 10 Ceacam10
1417075_at 1.16367 RIKEN cDNA 2010309E21 gene 2010309E21Rik
1417076_at 1.03551 "fatty acid binding protein 9, testis" Fabp9
1417077_at 0.96387 B-cell receptor-associated protein 29 Bcap29
1417078_at 1.02202 "lectin, galactose-binding, soluble 2" Lgals2
1417079_s_at 1.17751 "lectin, galactose-binding, soluble 2" Lgals2
1417080_a_at 1.01648 ECSIT homolog (Drosophila) Ecsit
1417081_a_at 1.05681 synaptogyrin 2 Syngr2
1417082_at 1.05072 "acidic nuclear phosphoprotein 32 family, member B" Anp32b
1417083_at 1.11344 Sec61 beta subunit Sec61b
1417084_at 1.01686 eukaryotic translation initiation factor 4E binding protein 2 Eif4ebp2
1417085_at 0.96706 "aldo-keto reductase family 1, member C6" Akr1c6
1417086_at 0.93276 "platelet-activating factor acetylhydrolase, isoform 1b, beta1 subunit" Pafah1b1
1417087_at 0.88180 golgi apparatus protein 1 Glg1
1417088_at 0.99221 zinc finger protein 346 Zfp346
1417089_a_at 0.96710 "creatine kinase, mitochondrial 1, ubiquitous" Ckmt1
1417090_at 0.92743 reticulocalbin 1 Rcn1
1417091_at 1.03333 conserved helix-loop-helix ubiquitous kinase Chuk
1417092_at 0.91547 parathyroid hormone receptor 1 Pthr1
1417093_a_at 0.96082 "general transcription factor II H, polypeptide 4" Gtf2h4
1417094_at 1.04540 acyl-CoA thioesterase 7 Acot7
1417095_a_at 1.02068 heat shock protein 14 Hspa14
1417096_at 1.26180 RIKEN cDNA 2810430M08 gene 2810430M08Rik
1417097_at 1.07843 mitochondrial trans-2-enoyl-CoA reductase Mecr
1417098_s_at 1.15375 mitochondrial trans-2-enoyl-CoA reductase Mecr
1417099_at 0.96775 Ftsj homolog 1 (E. coli) Ftsj1
1417100_at 0.95613 CD320 antigen Cd320
1417101_at 0.79212 heat shock protein 2 Hspa2
1417102_a_at 0.90593 "NADH dehydrogenase (ubiquinone) 1 beta subcomplex, 5" Ndufb5
1417103_at 1.10563 D-dopachrome tautomerase Ddt
1417104_at 1.06995 epithelial membrane protein 3 Emp3
1417105_at 1.05237 trafficking protein particle complex 2-like Trappc2l
1417106_at 0.93320 tumor protein D52-like 2 Tpd52l2
1417107_at 0.91189 tumor protein D52-like 2 Tpd52l2
1417108_at 0.83877 kinesin light chain 4 Klc4
1417109_at 0.96789 tubulointerstitial nephritis antigen-like Tinagl

1417110_at	0.61379	"mannosidase 1, alpha"	Man1a
1417111_at	0.69133	"mannosidase 1, alpha"	Man1a
1417112_at	0.91362	ADP-ribosylation factor-like 2 binding protein	Arl2bp
1417113_at	0.85913	germ cell-less homolog 1 (Drosophila)	Gmcl1
1417114_at	0.90148	germ cell-less homolog 1 (Drosophila)	Gmcl1
1417115_at	0.85461	mitogen activated protein kinase kinase kinase 12	Map3k12
1417116_at	0.93430	"solute carrier family 6 (neurotransmitter transporter, creatine), member 8"	Slc6a8
1417117_at	1.10811	"cleavage stimulation factor, 3' pre-RNA, subunit 1"	Cstf1
1417118_a_at	1.14429	N-acetyltransferase ARD1 homolog (S. cerevisiae)	Ard1
1417119_at	0.97368	zinc finger like protein 1	Zfp11
1417120_at	1.00449	"DNA segment, Chr 4, Wayne State University 114, expressed"	D4Wsu114e
1417121_at	0.99607	"gamma-aminobutyric acid (GABA-A) receptor, subunit alpha 6"	Gabra6
1417122_at	0.94956	vav 3 oncogene	Vav3
1417123_at	0.87883	vav 3 oncogene	Vav3
1417124_at	0.91923	destrin	Dstn
1417125_at	1.07038	S-adenosylhomocysteine hydrolase	Ahcy
1417126_a_at	1.01342	ribosomal protein L22 like 1	Rpl22i1
1417127_at	0.82753	"homeo box, msh-like 1"	Msx1
1417128_at	0.91318	"pleckstrin homology domain containing, family O member 1"	Plekho1
1417129_a_at	0.99442	myeloid ecotropic viral integration site-related gene 1	Mrg1
1417130_s_at	1.16747	angiopoietin-like 4	Angptl4
1417131_at	0.94829	cell division cycle 25 homolog A (S. cerevisiae)	Cdc25a
1417132_at	0.93716	cell division cycle 25 homolog A (S. cerevisiae)	Cdc25a
1417133_at	0.74125	peripheral myelin protein	Pmp22
1417134_at	0.98241	serine/arginine-rich protein specific kinase 2	Srpk2
1417135_at	0.96342	serine/arginine-rich protein specific kinase 2	Srpk2
1417136_s_at	0.96954	serine/arginine-rich protein specific kinase 2	Srpk2
1417137_at	1.16105	uridine-cytidine kinase 2	Uck2
1417138_s_at	1.10457	polymerase (RNA) II (DNA directed) polypeptide E	Polr2e
1417139_at	0.93502	RIKEN cDNA 1700022L09 gene	1700022L09Rik
1417140_a_at	1.15488	"protein tyrosine phosphatase, non-receptor type 2"	Ptpn2
1417141_at	1.05588	interferon gamma induced GTPase	Igtp
1417142_at	0.91196	RIKEN cDNA 4932442K08 gene	4932442K08Rik
1417143_at	0.94328	"endothelial differentiation, lysophosphatidic acid G-protein-coupled receptor, 2"	Edg2
1417144_at	0.93713	"tubulin, gamma 1"	Tubg1
1417145_at	0.92986	"nuclear transcription factor, X-box binding-like 1"	Nfxl1
1417146_at	1.20669	RIKEN cDNA 2410018C20 gene	2410018C20Rik
1417147_at	0.84073	RIKEN cDNA B230317C12 gene	B230317C12Rik
1417148_at	0.85824	"platelet derived growth factor receptor, beta polypeptide"	Pdgfrb
1417149_at	0.93171	"procollagen-proline, 2-oxoglutarate 4-dioxygenase (proline 4-hydroxylase), alpha II polypeptide"	P4ha2

1417150_at 1.10302 "solute carrier family 6 (neurotransmitter transporter, serotonin), member 4"
Slc6a4

1417151_a_at 0.91952 neurotensin receptor 2 Ntsr2

1417152_at 0.85594 BTB (POZ) domain containing 14A Btbd14a

1417153_at 0.94506 BTB (POZ) domain containing 14A Btbd14a

1417154_at 0.93889 "solute carrier family 25 (mitochondrial carrier, brain), member 14" Slc25a14

1417155_at 0.77211 "v-myc myelocytomatosis viral related oncogene, neuroblastoma derived
(avian)" Mycn

1417156_at 1.11357 keratin 19 Krt19

1417157_at 1.01524 ARP10 actin-related protein 10 homolog (*S. cerevisiae*) Actr10

1417158_at 1.00289 ZXD family zinc finger C Zxdc

1417159_at 1.11710 --- ---

1417160_s_at 1.14408 extracellular proteinase inhibitor Expi

1417161_at 0.96398 CDK2-associated protein 2 Cdk2ap2

1417162_at 1.04082 transmembrane BAX inhibitor motif containing 1 Tmbim1

1417163_at 1.09287 dual specificity phosphatase 10 Dusp10

1417164_at 1.01507 dual specificity phosphatase 10 Dusp10

1417165_at 1.01072 methyl-CpG binding domain protein 2 Mbd2

1417166_at 1.01778 PC4 and SFRS1 interacting protein 1 Psip1

1417167_at 1.03317 exosome component 5 Exosc5

1417168_a_at 0.52594 ubiquitin specific peptidase 2 Usp2

1417169_at 0.67345 ubiquitin specific peptidase 2 Usp2

1417170_at 0.96257 leucine zipper transcription factor-like 1 Lztf11

1417171_at 0.85775 IL2-inducible T-cell kinase Itk

1417172_at 1.13782 ubiquitin-conjugating enzyme E2L 6 Ube2l6

1417173_at 0.94076 cAMP responsive element binding protein-like 1 Creb11

1417174_at 0.93698 RIKEN cDNA 1810021J13 gene 1810021J13Rik

1417175_at 0.87628 "casein kinase 1, epsilon" Csnk1e

1417176_at 0.91063 "casein kinase 1, epsilon" Csnk1e

1417177_at 1.27653 galactokinase 1 Galk1

1417178_at 0.93573 "GIPC PDZ domain containing family, member 2" Gipc2

1417179_at 0.82271 tetraspanin 5 Tspan5

1417180_at 0.94469 proprotein convertase subtilisin/kexin type 7 Pcsk7

1417181_a_at 0.89599 kinesin-associated protein 3 Kifap3

1417182_at 0.92988 "DnaJ (Hsp40) homolog, subfamily A, member 2" Dnaja2

1417183_at 0.98009 "DnaJ (Hsp40) homolog, subfamily A, member 2" Dnaja2

1417184_s_at 0.96683 "hemoglobin, beta adult major chain /// hemoglobin, beta adult minor chain"
Hbb-b1 /// Hbb-b2

1417185_at 1.09157 "lymphocyte antigen 6 complex, locus A" Ly6a

1417186_at 1.06668 huntingtin interacting protein 2 Hip2

1417187_at 0.98828 huntingtin interacting protein 2 Hip2

1417188_s_at 0.96442 huntingtin interacting protein 2 Hip2

1417189_at 1.19405 "proteasome (prosome, macropain) 28 subunit, beta" Psme2

1417190_at 1.27562 pre-B-cell colony-enhancing factor 1 Pbef1
 1417191_at 0.78152 "DnaJ (Hsp40) homolog, subfamily B, member 9" Dnajb9
 1417192_at 1.11930 translocase of outer mitochondrial membrane 70 homolog A (yeast)
 Tomm70a
 1417193_at 0.99827 "superoxide dismutase 2, mitochondrial" Sod2
 1417194_at 0.96617 "superoxide dismutase 2, mitochondrial" Sod2
 1417195_at 0.94857 "WW, C2 and coiled-coil domain containing 2" Wwc2
 1417196_s_at 0.94795 "WW, C2 and coiled-coil domain containing 2" Wwc2
 1417197_at 0.97949 "WW, C2 and coiled-coil domain containing 2" Wwc2
 1417198_at 0.94115 "WW, C2 and coiled-coil domain containing 2" Wwc2
 1417199_at 0.98637 RIKEN cDNA 1300007B12 gene 1300007B12Rik
 1417200_at 1.03172 RIKEN cDNA 1300007B12 gene 1300007B12Rik
 1417201_at 0.91134 "5'-nucleotidase, cytosolic II" Nt5c2
 1417202_s_at 0.93927 ubiquitin-activating enzyme E1C Ube1c
 1417203_at 1.08829 ethylmalonic encephalopathy 1 Ethel
 1417204_at 0.98451 KDEL (Lys-Asp-Glu-Leu) endoplasmic reticulum protein retention receptor
 2 Kdelr2
 1417205_at 0.93193 KDEL (Lys-Asp-Glu-Leu) endoplasmic reticulum protein retention receptor
 2 Kdelr2
 1417206_at 1.08928 uroporphyrinogen decarboxylase Urod
 1417207_at 1.07443 "dishevelled 2, dsh homolog (Drosophila)" Dvl2
 1417208_at 0.95050 alpha-methylacyl-CoA racemase Amacr
 1417209_at 1.01309 SERTA domain containing 2 Sertad2
 1417210_at 1.10990 "eukaryotic translation initiation factor 2, subunit 3, structural gene Y-
 linked" Eif2s3y
 1417211_a_at 0.89900 RIKEN cDNA 1110032A03 gene 1110032A03Rik
 1417212_at 0.93000 RIKEN cDNA 9530058B02 gene 9530058B02Rik
 1417213_a_at 0.93679 RNA binding motif protein 6 Rbm6
 1417214_at 0.76138 "RAB27b, member RAS oncogene family" Rab27b
 1417215_at 0.82291 "RAB27b, member RAS oncogene family" Rab27b
 1417216_at 1.09746 proviral integration site 2 Pim2
 1417217_at 0.94014 "melanoma antigen, family L, 2" Magel2
 1417218_at 0.88083 RIKEN cDNA 2810048G17 gene 2810048G17Rik
 1417219_s_at 1.09901 "thymosin, beta 10" Tmsb10
 1417220_at 1.02640 fumarylacetoacetate hydrolase Fah
 1417221_at 0.87566 --- ---
 1417222_a_at 0.88608 transmembrane protein 123 Tmem123
 1417223_at 0.97153 CD2 antigen (cytoplasmic tail) binding protein 2 Cd2bp2
 1417224_a_at 0.87092 CD2 antigen (cytoplasmic tail) binding protein 2 Cd2bp2
 1417225_at 1.07670 ADP-ribosylation factor-like 6 interacting protein 5 Arl6ip5
 1417226_at 0.97991 F-box and WD-40 domain protein 4 Fbxw4
 1417227_at 1.04273 methylcrotonoyl-Coenzyme A carboxylase 1 (alpha) Mccc1
 1417228_at 0.96425 calpain 1 Capn1

1417229_at	1.02276	calpain 1	Capn1
1417230_at	1.18296	Ral GEF with PH domain and SH3 binding motif 2	Ralgps2
1417231_at	1.20249	claudin 2	Cldn2
1417232_at	1.13888	claudin 2	Cldn2
1417233_at	1.01868	coiled-coil-helix-coiled-coil-helix domain containing 4	Chchd4
1417234_at	0.68750	matrix metalloproteinase 11	Mmp11
1417235_at	0.86141	EH-domain containing 3	Ehd3
1417236_at	0.93856	EH-domain containing 3	Ehd3
1417237_at	0.99761	phospholipase D2	Pld2
1417238_at	1.01614	Ewing sarcoma breakpoint region 1	Ewsr1
1417239_at	1.06402	centrin 3	Cetn3
1417240_at	0.86872	zyxin	Zyx
1417241_at	0.90643	EST X83328	X83328
1417242_at	1.00341	DEAD (Asp-Glu-Ala-Asp) box polypeptide 48	Ddx48
1417243_at	0.95318	RIKEN cDNA 2310065K24 gene	2310065K24Rik
1417244_a_at	1.03279	interferon regulatory factor 7	Irf7
1417245_at	0.99497	G protein-coupled receptor 180	Gpr180
1417246_at	1.01460	pregnancy zone protein	Pzp
1417247_at	0.93240	expressed sequence AI597479	AI597479
1417248_at	1.06120	ralA binding protein 1	Ralbp1
1417249_at	1.14092	"polymerase (DNA directed), mu"	Polm
1417250_at	1.11782	ring finger protein 12	Rnf12
1417251_at	1.14835	palmdelphin	Palmd
1417252_at	1.21615	"5',3'-nucleotidase, cytosolic"	Nt5c
1417253_at	1.00340	FSHD region gene 1	Frg1
1417254_at	1.08168	spermatogenesis associated 5	Spata5
1417255_at	0.78749	intraflagellar transport 140 homolog (Chlamydomonas)	Ift140
1417256_at	0.96038	matrix metalloproteinase 13	Mmp13
1417257_at	0.89696	carboxyl ester lipase	Cel
1417258_at	1.06158	chaperonin subunit 5 (epsilon)	Cct5
1417259_a_at	1.01419	"capping protein (actin filament) muscle Z-line, beta"	Capzb
1417260_at	0.96188	U2 small nuclear ribonucleoprotein auxiliary factor (U2AF) 2	U2af2
1417261_at	0.93696	mbt domain containing 1	Mbtd1
1417262_at	1.09336	prostaglandin-endoperoxide synthase 2	Ptgs2
1417263_at	1.24440	prostaglandin-endoperoxide synthase 2	Ptgs2
1417264_at	0.87241	"coenzyme Q5 homolog, methyltransferase (yeast)"	Coq5
1417265_s_at	0.98244	"coenzyme Q5 homolog, methyltransferase (yeast)"	Coq5
1417266_at	1.02135	chemokine (C-C motif) ligand 6	Ccl6
1417267_s_at	1.17349	FK506 binding protein 11	Fkbp11
1417268_at	1.45882	CD14 antigen	Cd14
1417269_at	1.03316	cyclin-dependent kinase 9 (CDC2-related kinase)	Cdk9
1417270_at	1.20914	WD repeat domain 12	Wdr12
1417271_a_at	0.99485	endoglin	Eng

1417272_at 0.75434 RIKEN cDNA 9130005N14 gene 9130005N14Rik
1417273_at 0.77880 "pyruvate dehydrogenase kinase, isoenzyme 4" Pdk4
1417274_at 1.08655 small nuclear ribonucleoprotein polypeptide A Snrpa
1417275_at 1.00181 "myelin and lymphocyte protein, T-cell differentiation protein" Mal
1417276_at 0.86732 tubby-like protein 2 Tulp2
1417277_at 0.92550 "cytochrome P450, family 4, subfamily f, polypeptide 16" Cyp4f16
1417278_a_at 0.98806 naked cuticle 1 homolog (Drosophila) /// similar to naked cuticle 1 homolog ///
similar to naked cuticle 1 homolog Nkd1 /// LOC634379 /// LOC671838
1417279_at 0.71255 "inositol 1,4,5-triphosphate receptor 1" Itpr1
1417280_at 1.15011 "solute carrier family 17 (sodium phosphate), member 1" Slc17a1
1417281_a_at 0.98620 matrix metalloproteinase 23 Mmp23
1417282_at 1.05930 matrix metalloproteinase 23 Mmp23
1417283_at 0.77308 Ly6/neurotoxin 1 Lynx1
1417284_at 0.92796 mitogen-activated protein kinase associated protein 1 Mapkap1
1417285_a_at 0.91675 "NADH dehydrogenase (ubiquinone) 1 alpha subcomplex, 5" Ndufa5
1417286_at 0.94089 "NADH dehydrogenase (ubiquinone) 1 alpha subcomplex, 5" Ndufa5
1417287_at 0.94451 histocompatibility 13 H13
1417288_at 1.14815 "pleckstrin homology domain-containing, family A (phosphoinositide binding
specific) member 2" Plekha2
1417289_at 1.06316 "pleckstrin homology domain-containing, family A (phosphoinositide binding
specific) member 2" Plekha2
1417290_at 0.93216 leucine-rich alpha-2-glycoprotein 1 Lrg1
1417291_at 1.01050 "tumor necrosis factor receptor superfamily, member 1a" Tnfrsf1a
1417292_at 1.08552 interferon gamma inducible protein 47 Ifi47
1417293_at 0.88217 heparan sulfate 6-O-sulfotransferase 1 Hs6st1
1417294_at 0.89341 "aldo-keto reductase family 7, member A5 (aflatoxin aldehyde reductase)"
Akr7a5
1417295_at 1.09610 metastasis associated 1 Mta1
1417296_at 1.02234 activating transcription factor 1 Atf1
1417297_at 1.03043 "inositol 1,4,5-triphosphate receptor 3" Itpr3
1417298_at 1.01554 emopamil binding protein-like Ebpl
1417299_at 1.19431 NIMA (never in mitosis gene a)-related expressed kinase 2 Nek2
1417300_at 1.10560 "sphingomyelin phosphodiesterase, acid-like 3B" Smpdl3b
1417301_at 0.79249 frizzled homolog 6 (Drosophila) Fzd6
1417302_at 1.16353 REST corepressor 2 Rcor2
1417303_at 1.52229 mevalonate (diphospho) decarboxylase Mvd
1417304_at 0.94354 chordin Chrd
1417305_at 0.76976 SPEG complex locus Speg
1417306_at 1.14477 tyrosine kinase 2 Tyk2
1417307_at 0.93475 "dystrophin, muscular dystrophy" Dmd
1417308_at 1.04527 "pyruvate kinase, muscle" Pkm2
1417309_at 0.99176 "transducer of ERBB2, 2" Tob2
1417310_at 0.87896 "transducer of ERBB2, 2" Tob2

1417311_at 0.82916 cysteine rich protein 2 Crip2
1417312_at 1.02060 dickkopf homolog 3 (Xenopus laevis) Dkk3
1417313_at 1.07047 "LSM7 homolog, U6 small nuclear RNA associated (S. cerevisiae)" Lsm7
1417314_at 0.92738 complement factor B Cfb
1417315_at 0.92120 GRIP1 associated protein 1 Gripap1
1417316_at 1.02592 thioesterase superfamily member 2 Them2
1417317_s_at 1.01072 ribosomal protein L35a Rpl35a
1417318_at 0.95392 deleted in bladder cancer 1 (human) Dbc1
1417319_at 0.79120 poliovirus receptor-related 3 Pvr13
1417320_at 1.07464 "GrpE-like 1, mitochondrial" Grpel1
1417321_at 0.95817 "DNA segment, Chr 4, Wayne State University 132, expressed" D4Wsu132e
1417322_at 1.03085 deoxyhypusine hydroxylase/monooxygenase Dohh
1417323_at 1.02608 proline/serine-rich coiled-coil 1 Psrc1
1417324_at 0.97621 microtubule associated serine/threonine kinase 2 Mast2
1417325_at 0.97057 beta-transducin repeat containing protein Btrc
1417326_a_at 0.94631 anaphase promoting complex subunit 11 homolog (yeast) Anapc11
1417327_at 0.92789 caveolin 2 Cav2
1417328_at 0.97977 "excision repair cross-complementing rodent repair deficiency, complementation group 1" Ercc1
1417329_at 0.91471 "solute carrier family 23 (nucleobase transporters), member 2" Slc23a2
1417330_at 0.95548 "solute carrier family 23 (nucleobase transporters), member 2" Slc23a2
1417331_a_at 0.91101 ADP-ribosylation factor-like 6 Arl6
1417332_at 0.88273 "regulatory factor X, 2 (influences HLA class II expression)" Rfx2
1417333_at 0.99946 RAS p21 protein activator 4 Rasa4
1417334_at 0.99548 serine/threonine kinase 19 Stk19
1417335_at 0.88248 "sulfotransferase family, cytosolic, 2B, member 1" Sult2b1
1417336_a_at 0.88069 synaptotagmin-like 4 Sytl4
1417337_at 0.95853 erythrocyte protein band 4.2 Epb4.2
1417338_at 1.11908 erythrocyte protein band 4.2 Epb4.2
1417339_a_at 0.96066 "dynein light chain LC8-type 1 /// similar to dynein, cytoplasmic, light peptide /// similar to dynein, cytoplasmic, light peptide /// similar to dynein, cytoplasmic, light peptide" Dynll1 /// LOC627788 /// LOC637840 /// LOC672375
1417340_at 1.04784 thioredoxin-like 2 Txnl2
1417341_a_at 1.02132 "protein phosphatase 1, regulatory (inhibitor) subunit 2" Ppp1r2
1417342_at 1.02336 "protein phosphatase 1, regulatory (inhibitor) subunit 2" Ppp1r2
1417343_at 1.02118 FXYD domain-containing ion transport regulator 6 Fxyd6
1417344_at 0.98554 RIKEN cDNA 2900064A13 gene 2900064A13Rik
1417345_at 1.06515 RIKEN cDNA 2900064A13 gene 2900064A13Rik
1417346_at 0.96692 PYD and CARD domain containing Pycard
1417347_at 0.92608 PYD and CARD domain containing Pycard
1417348_at 1.04776 RIKEN cDNA 2310039H08 gene 2310039H08Rik
1417349_at 1.01651 pallidin Pldn
1417350_at 1.03282 pallidin Pldn

1417351_a_at 1.14762 small nuclear ribonucleoprotein polypeptide A' Snrpa1
1417352_s_at 1.24863 small nuclear ribonucleoprotein polypeptide A' Snrpa1
1417353_x_at 1.11259 small nuclear ribonucleoprotein polypeptide A' Snrpa1
1417354_at 0.96325 endoplasmic reticulum chaperone SIL1 homolog (S. cerevisiae) Sil1
1417355_at 0.96069 paternally expressed 3 Peg3
1417356_at 0.90423 paternally expressed 3 Peg3
1417357_at 1.00768 emerlin Emd
1417358_s_at 0.74323 sorbin and SH3 domain containing 1 Sorbs1
1417359_at 1.09867 microfibrillar-associated protein 2 Mfap2
1417360_at 1.01930 mutL homolog 1 (E. coli) Mlh1
1417361_at 0.99013 ankyrin repeat and SOCS box-containing protein 3 Asb3
1417362_at 0.92871 Rhesus blood group-associated C glycoprotein Rhcg
1417363_at 0.96095 zinc finger protein 61 Zfp61
1417364_at 1.03786 eukaryotic translation elongation factor 1 gamma Eef1g
1417365_a_at 1.06049 calmodulin 1 Calm1
1417366_s_at 1.14226 calmodulin 1 Calm1
1417367_at 0.97182 "protein phosphatase 2 (formerly 2A), catalytic subunit, alpha isoform" Ppp2ca
1417368_s_at 1.04521 "NADH dehydrogenase (ubiquinone) 1 alpha subcomplex, 2" Ndufa2
1417369_at 0.95611 hydroxysteroid (17-beta) dehydrogenase 4 Hsd17b4
1417370_at 0.96654 "trefoil factor 3, intestinal" Tff3
1417371_at 1.06963 pellino 1 Peli1
1417372_a_at 0.97971 pellino 1 Peli1
1417373_a_at 1.01068 "tubulin, alpha 4" Tuba4
1417374_at 1.02877 "tubulin, alpha 4" Tuba4
1417375_at 0.93934 "tubulin, alpha 4" Tuba4
1417376_a_at 0.82472 "immunoglobulin superfamily, member 4A" Igsf4a
1417377_at 0.90262 "immunoglobulin superfamily, member 4A" Igsf4a
1417378_at 0.89497 "immunoglobulin superfamily, member 4A" Igsf4a
1417379_at 1.02261 IQ motif containing GTPase activating protein 1 Iqgap1
1417380_at 1.02316 IQ motif containing GTPase activating protein 1 Iqgap1
1417381_at 0.90026 "complement component 1, q subcomponent, alpha polypeptide" C1qa
1417382_at 0.90213 ectonucleoside triphosphate diphosphohydrolase 5 Entpd5
1417383_at 0.94699 ectonucleoside triphosphate diphosphohydrolase 5 Entpd5
1417384_at 0.83728 ectonucleoside triphosphate diphosphohydrolase 5 Entpd5
1417385_at 1.05674 aminopeptidase puromycin sensitive Npepps
1417386_at 1.13858 aminopeptidase puromycin sensitive Npepps
1417387_at 1.01368 "mediator of RNA polymerase II transcription, subunit 31 homolog (yeast)"
Med31
1417388_at 0.91041 brain expressed X-linked 2 Bex2
1417389_at 0.83436 glypican 1 Gpc1
1417390_at 1.02376 XPA binding protein 1 Xab1
1417391_a_at 1.07215 interleukin 16 Il16
1417392_a_at 0.57932 "solute carrier family 7 (cationic amino acid transporter, y+ system), member

7" Slc7a7

1417393_a_at	0.94534	C1q domain containing 2	C1qdc2
1417394_at	1.01146	Kruppel-like factor 4 (gut)	Klf4
1417395_at	0.85248	Kruppel-like factor 4 (gut)	Klf4
1417396_at	0.86996	podocalyxin-like	Podxl
1417397_at	0.93772	"solute carrier family 9 (sodium/hydrogen exchanger), member 1"	Slc9a1
1417398_at	0.99914	related RAS viral (r-ras) oncogene homolog 2	Rras2
1417399_at	0.85753	growth arrest specific 6	Gas6
1417400_at	1.19874	retinoic acid induced 14	Rai14
1417401_at	1.17654	retinoic acid induced 14	Rai14
1417402_at	0.96078	RIKEN cDNA 1190017O12 gene	1190017O12Rik
1417403_at	1.49220	"ELOVL family member 6, elongation of long chain fatty acids (yeast)"	Elovl6
1417404_at	1.52747	"ELOVL family member 6, elongation of long chain fatty acids (yeast)"	Elovl6
1417405_at	0.85255	START domain containing 3	Stard3
1417406_at	1.08882	SERTA domain containing 1	Sertad1
1417407_at	1.02281	F-box and leucine-rich repeat protein 14	Fbxl14
1417408_at	1.12619	coagulation factor III	F3
1417409_at	0.69016	Jun oncogene	Jun
1417410_s_at	1.00937	"protein kinase C, iota"	Prkci
1417411_at	1.09540	nucleosome assembly protein 1-like 5	Nap115
1417412_at	0.92054	factor 8-associated gene A	F8a
1417413_at	0.89159	CUB and zona pellucida-like domains 1	Cuzd1
1417414_at	1.10586	septin 3	3-Sep
1417415_at	0.88485	"solute carrier family 6 (neurotransmitter transporter, dopamine), member 3"	Slc6a3
1417416_at	0.91322	"potassium voltage-gated channel, shaker-related subfamily, member 1"	Kcna1
1417417_a_at	0.97493	"cytochrome c oxidase, subunit VI a, polypeptide 1"	Cox6a1
1417418_s_at	0.96376	"cytochrome c oxidase, subunit VI a, polypeptide 1 /// similar to Cytochrome c oxidase polypeptide VIa-liver, mitochondrial precursor /// similar to Cytochrome c oxidase polypeptide VIa-liver, mitochondrial precursor"	Cox6a1 /// LOC665806 /// LOC673054
1417419_at	1.56313	cyclin D1	Ccnd1
1417420_at	1.37958	cyclin D1	Ccnd1
1417421_at	0.85954	S100 calcium binding protein A1	S100a1
1417422_at	0.94605	glycine N-methyltransferase	Gnmt
1417423_at	0.94560	"glutamate receptor, ionotropic, N-methyl D-aspartate-associated protein 1 (glutamate binding)"	Grina
1417424_at	0.95709	immediate early response 3 interacting protein 1	Ier3ip1
1417425_at	1.05255	Prkr interacting protein 1 (IL11 inducible)	Prkrip1
1417426_at	1.02987	"proteoglycan 1, secretory granule"	Prg1
1417427_at	1.01535	RIKEN cDNA 1500026D16 gene	1500026D16Rik
1417428_at	1.01514	"guanine nucleotide binding protein (G protein), gamma 3 subunit"	Gng3
1417429_at	1.05388	flavin containing monooxygenase 1	Fmo1
1417430_at	0.71039	cerebellar degeneration-related 2	Cdr2

1417431_a_at 0.95858 sphingosine kinase 2 Sphk2
1417432_a_at 1.01535 "guanine nucleotide binding protein, beta 1" Gnb1
1417433_at 0.92203 lysophospholipase 2 Lypla2
1417434_at 1.26026 "glycerol phosphate dehydrogenase 2, mitochondrial" Gpd2
1417435_at 0.88645 like-glycosyltransferase Large
1417436_at 1.01367 like-glycosyltransferase Large
1417437_at 0.95330 X-ray repair complementing defective repair in Chinese hamster cells 6 Xrcc6
1417438_at 0.89859 retinol dehydrogenase 14 (all-trans and 9-cis) Rdh14
1417439_at 0.68583 "CD248 antigen, endosialin" Cd248
1417440_at 0.91035 AT rich interactive domain 1A (Swi1 like) /// similar to AT rich interactive domain 1A isoform a Arid1a /// LOC675933
1417441_at 0.98643 "DnaJ (Hsp40) homolog, subfamily C, member 12" Dnajc12
1417442_a_at 0.95172 peroxisomal biogenesis factor 3 Pex3
1417443_at 0.74180 cDNA sequence BC026682 BC026682
1417444_at 1.04530 E2F transcription factor 5 E2f5
1417445_at 1.04369 kinetochore associated 2 Kntc2
1417446_at 1.04474 "solute carrier family 12, member 4" Slc12a4
1417447_at 0.96839 transcription factor 21 Tcf21
1417448_at 0.95902 RIKEN cDNA 1810008A18 gene 1810008A18Rik
1417449_at 1.00673 acyl-CoA thioesterase 8 Acot8
1417450_a_at 1.03419 "transforming, acidic coiled-coil containing protein 3" Tacc3
1417451_a_at 0.98428 peptidylprolyl isomerase A Ppia
1417452_a_at 1.06281 Finkel-Biskis-Reilly murine sarcoma virus (FBR-MuSV) ubiquitously expressed (fox derived) Fau
1417453_at 1.12566 cullin 4B Cul4b
1417454_at 0.98780 cullin 4B Cul4b
1417455_at 0.80670 "transforming growth factor, beta 3" Tgfb3
1417456_at 0.95499 glyceronephosphate O-acyltransferase Gnpat
1417457_at 0.98765 CDC28 protein kinase regulatory subunit 2 Cks2
1417458_s_at 0.96240 CDC28 protein kinase regulatory subunit 2 Cks2
1417459_at 0.93565 demilune cell and parotid protein Dcpp
1417460_at 0.95662 interferon induced transmembrane protein 2 Ifitm2
1417461_at 0.96829 "CAP, adenylate cyclase-associated protein 1 (yeast)" Cap1
1417462_at 0.94467 "CAP, adenylate cyclase-associated protein 1 (yeast)" Cap1
1417463_a_at 1.00256 RIKEN cDNA 2400001E08 gene /// hypothetical LOC433216 2400001E08Rik /// LOC433216
1417464_at 0.93767 "troponin C2, fast" Tnnc2
1417465_at 0.96479 "farnesyltransferase, CAAX box, alpha" Fnta
1417466_at 1.03892 --- ---
1417467_a_at 0.95898 "transcriptional adaptor 3 (NGG1 homolog, yeast)-like" Tada3l
1417468_at 0.93644 nitrilase 1 Nit1
1417469_at 0.99479 nitrilase 1 Nit1
1417470_at 1.02142 apolipoprotein B editing complex 3 Apobec3

1417471_s_at 1.03663 "DNA segment, Chr 1, ERATO Doi 622, expressed" D1Ert622e
1417472_at 0.92200 "myosin, heavy polypeptide 9, non-muscle" Myh9
1417473_a_at 0.98709 phosphopantothenoylcysteine synthetase Ppcs
1417474_at 1.04006 RIKEN cDNA 1500035H01 gene 1500035H01Rik
1417475_at 0.99709 ATPase type 13A1 Atp13a1
1417476_at 0.91559 F-box and WD-40 domain protein 5 Fbxw5
1417477_at 0.87397 gene trap locus F3b Gtlf3b
1417478_a_at 0.95301 RIKEN cDNA 4930511A21 gene 4930511A21Rik
1417479_at 0.89946 RIKEN cDNA 4930511A21 gene 4930511A21Rik
1417480_at 0.89316 f-box only protein 9 Fbxo9
1417481_at 0.65511 receptor (calcitonin) activity modifying protein 1 Ramp1
1417482_at 1.03483 testis expressed gene 19 Tex19
1417483_at 1.18481 "nuclear factor of kappa light polypeptide gene enhancer in B-cells inhibitor,
zeta" Nfkbiz
1417484_at 1.01317 integrin binding sialoprotein Ibsp
1417485_at 1.06844 integrin binding sialoprotein Ibsp
1417486_at 1.01323 src homology 2 domain-containing transforming protein D Shd
1417487_at 1.64743 fos-like antigen 1 Fosl1
1417488_at 1.60200 fos-like antigen 1 Fosl1
1417489_at 1.04435 neuropeptide Y receptor Y2 Npy2r
1417490_at 0.92910 cathepsin B Ctsb
1417491_at 0.79648 cathepsin B Ctsb
1417492_at 0.72150 cathepsin B Ctsb
1417493_at 0.91078 B lymphoma Mo-MLV insertion region 1 Bmi1
1417494_a_at 0.90846 ceruloplasmin Cp
1417495_x_at 0.95846 ceruloplasmin Cp
1417496_at 0.98932 ceruloplasmin Cp
1417497_at 0.72817 ceruloplasmin Cp
1417498_at 0.89097 "serine (or cysteine) peptidase inhibitor, clade F, member 2" Serpinf2
1417499_at 1.10354 translocase of inner mitochondrial membrane 10 homolog (yeast) Timm10
1417500_a_at 0.95144 "transglutaminase 2, C polypeptide" Tgm2
1417501_at 1.25918 F-box only protein 6b Fbxo6b
1417502_at 0.79077 tetraspanin 7 Tspan7
1417503_at 0.92979 replication factor C (activator 1) 2 Rfc2
1417504_at 0.92849 calbindin-28K Calb1
1417505_s_at 1.02361 "interleukin 11 receptor, alpha chain 1 /// interleukin 11 receptor, alpha chain
2 /// similar to interleukin 11 receptor, alpha chain 2" Il11ra1 /// Il11ra2 /// LOC621887
1417506_at 1.06236 geminin Gmnn
1417507_at 0.87295 cytochrome b-561 Cyb561
1417508_at 1.03664 ring finger protein (C3HC4 type) 19 Rnf19
1417509_at 1.14711 ring finger protein (C3HC4 type) 19 Rnf19
1417510_at 0.91163 vacuolar protein sorting 4a (yeast) Vps4a
1417511_at 1.14273 Ly1 antibody reactive clone Lyar

1417512_at	0.98208	ecotropic viral integration site 5	Evi5
1417513_at	0.94150	ecotropic viral integration site 5	Evi5
1417514_at	1.03026	"synovial sarcoma, X breakpoint 2 interacting protein"	Ssx2ip
1417515_at	0.97662	U7 snRNP-specific Sm-like protein LSM10	Lsm10
1417516_at	1.21066	DNA-damage inducible transcript 3	Ddit3
1417517_at	0.98652	pleiomorphic adenoma gene-like 2	Plagl2
1417518_at	1.03552	pleiomorphic adenoma gene-like 2	Plagl2
1417519_at	0.99598	pleiomorphic adenoma gene-like 2	Plagl2
1417520_at	0.67652	"nuclear factor, erythroid derived 2, like 3"	Nfe2l3
1417521_at	0.85686	ephrin A2	Efna2
1417522_at	0.83423	F-box only protein 32	Fbxo32
1417523_at	1.01635	pleckstrin	Plek
1417524_at	0.60007	cornichon homolog 2 (Drosophila)	Cnih2
1417525_at	1.21613	heart and neural crest derivatives expressed transcript 1	Hand1
1417526_at	0.83420	poly(rC) binding protein 3	Pcbp3
1417527_at	0.83521	"adaptor-related protein complex 3, mu 2 subunit"	Ap3m2
1417528_at	1.17943	sperm associated antigen 6	Spag6
1417529_at	0.98391	"RAB33A, member of RAS oncogene family"	Rab33a
1417530_a_at	1.13107	signal recognition particle 9	Srp9
1417531_at	0.71164	"cytochrome P450, family 2, subfamily j, polypeptide 5"	Cyp2j5
1417532_at	1.14296	"cytochrome P450, family 2, subfamily j, polypeptide 5"	Cyp2j5
1417533_a_at	0.71244	integrin beta 5	Itgb5
1417534_at	0.68920	integrin beta 5	Itgb5
1417535_at	0.92725	F-box only protein 25	Fbxo25
1417536_at	1.06612	"zinc finger, matrin type 2"	Zmat2
1417537_at	0.96747	"zinc finger, matrin type 2"	Zmat2
1417538_at	0.98252	"solute carrier family 35 (CMP-sialic acid transporter), member 1"	Slc35a1
1417539_at	0.93646	"solute carrier family 35 (CMP-sialic acid transporter), member 1"	Slc35a1
1417540_at	1.07094	E74-like factor 1	Elf1
1417541_at	1.02714	"helicase, lymphoid specific"	Hells
1417542_at	0.87377	"ribosomal protein S6 kinase, polypeptide 2"	Rps6ka2
1417543_at	0.96257	"ribosomal protein S6 kinase, polypeptide 2"	Rps6ka2
1417544_a_at	0.94737	flotillin 2	Flot2
1417545_at	0.88269	"transient receptor potential cation channel, subfamily V, member 4"	Trpv4
1417546_at	0.97443	"interleukin 2 receptor, beta chain"	Il2rb
1417547_at	0.98146	squamous cell carcinoma antigen recognized by T-cells 3	Sart3
1417548_at	1.00292	squamous cell carcinoma antigen recognized by T-cells 3	Sart3
1417549_at	0.74690	zinc finger protein 68	Zfp68
1417550_a_at	0.98486	germ cell-specific gene 1	Gsg1
1417551_at	1.12898	"ceroid lipofuscinosis, neuronal 3, juvenile (Batten, Spielmeier-Vogt disease)"	Cln3
1417552_at	0.88451	fibroblast activation protein	Fap
1417553_at	1.05028	placental specific protein 1	Plac1

1417554_at 0.90917 "hydroxy-delta-5-steroid dehydrogenase, 3 beta- and steroid delta-isomerase 4"
Hsd3b4

1417555_at 0.94654 "ATPase family, AAA domain containing 1" Atad1

1417556_at 1.00821 "fatty acid binding protein 1, liver" Fabp1

1417557_at 1.01218 UBX domain containing 1 Ubx1

1417558_at 0.92321 Fyn proto-oncogene Fyn

1417559_at 1.08925 sideroflexin 1 Sfxn1

1417560_at 1.02476 sideroflexin 1 Sfxn1

1417561_at 1.00815 apolipoprotein C-I Apoc1

1417562_at 1.21948 eukaryotic translation initiation factor 4E binding protein 1 Eif4ebp1

1417563_at 1.02088 eukaryotic translation initiation factor 4E binding protein 1 Eif4ebp1

1417564_at 0.98865 "cofactor required for Sp1 transcriptional activation, subunit 9" Crsp9

1417565_at 1.07866 abhydrolase domain containing 5 Abhd5

1417566_at 1.06931 abhydrolase domain containing 5 Abhd5

1417567_at 0.90933 catenin beta interacting protein 1 Ctnnb1

1417568_at 0.82173 neurocalcin delta Ncald

1417569_at 0.84546 neurocalcin delta Ncald

1417570_at 0.90986 anaphase promoting complex subunit 1 Anapc1

1417571_at 1.03300 N-methylpurine-DNA glycosylase Mpg

1417572_at 0.80416 N-methylpurine-DNA glycosylase Mpg

1417573_at 0.97781 RIKEN cDNA 2010311D03 gene 2010311D03Rik

1417574_at 0.73532 chemokine (C-X-C motif) ligand 12 Cxcl12

1417575_at 0.88933 "OTU domain, ubiquitin aldehyde binding 2" Otub2

1417576_a_at 0.84572 "OTU domain, ubiquitin aldehyde binding 2" Otub2

1417577_at 0.87543 "transient receptor potential cation channel, subfamily C, member 3" Trpc3

1417578_a_at 0.95053 GDP-mannose pyrophosphorylase A Gmppa

1417579_x_at 0.95555 GDP-mannose pyrophosphorylase A Gmppa

1417580_s_at 0.73845 selenium binding protein 1 Selenbp1

1417581_at 0.90270 dihydroorotate dehydrogenase Dhodh

1417582_s_at 1.03730 dihydroorotate dehydrogenase Dhodh

1417583_a_at 1.03292 charged amino acid rich leucine zipper 1 Crlz1

1417584_at 0.96626 "solute carrier family 11 (proton-coupled divalent metal ion transporters),
member 2" Slc11a2

1417585_at 0.83283 nucleoporin 210 Nup210

1417586_at 0.99147 timeless homolog (Drosophila) Timeless

1417587_at 0.91030 timeless homolog (Drosophila) Timeless

1417588_at 1.08690 UDP-N-acetyl-alpha-D-galactosamine:polypeptide N-
acetylgalactosaminyltransferase 3 Galnt3

1417589_at 1.06788 UDP-N-acetyl-alpha-D-galactosamine:polypeptide N-
acetylgalactosaminyltransferase 3 /// hypothetical protein LOC620893 Galnt3 /// LOC620893

1417590_at 1.00704 "cytochrome P450, family 27, subfamily a, polypeptide 1" Cyp27a1

1417591_at 1.03294 prostaglandin E synthase 2 Ptges2

1417592_at 0.98066 FK506 binding protein 12-rapamycin associated protein 1 Frap1

1417593_at 0.91075 tumor suppressor candidate 2 Tusc2
1417594_at 0.87146 G kinase anchoring protein 1 Gkap1
1417595_at 1.07829 mesenchyme homeobox 1 Meox1
1417596_at 1.01092 endothelial precursor protein B9 Eppb9
1417597_at 1.00713 CD28 antigen Cd28
1417598_a_at 1.04340 "fragile X mental retardation gene 1, autosomal homolog" Fxr1h
1417599_at 0.87034 CD276 antigen Cd276
1417600_at 0.94168 "solute carrier family 15 (H+/peptide transporter), member 2" Slc15a2
1417601_at 1.07196 regulator of G-protein signaling 1 Rgs1
1417602_at 0.76418 period homolog 2 (Drosophila) Per2
1417603_at 1.09339 period homolog 2 (Drosophila) Per2
1417604_at 0.91132 calcium/calmodulin-dependent protein kinase I Camk1
1417605_s_at 0.89578 calcium/calmodulin-dependent protein kinase I /// similar to calcium/
calmodulin-dependent protein kinase I Camk1 /// LOC669267
1417606_a_at 0.93248 calreticulin Calr
1417607_at 1.02096 "cytochrome c oxidase, subunit VI a, polypeptide 2" Cox6a2
1417608_a_at 1.05007 ribosomal protein L13a Rpl13a
1417609_at 0.85829 "ubiquitin-conjugating enzyme E2A, RAD6 homolog (S. cerevisiae)" Ube2a
1417610_at 0.97923 apolipoprotein A-V Apoa5
1417611_at 0.86964 transmembrane protein 37 Tmem37
1417612_at 0.81048 immediate early response 5 Ier5
1417613_at 0.87293 immediate early response 5 Ier5
1417614_at 1.00160 "creatine kinase, muscle" Ckm
1417615_a_at 1.11410 ribosomal protein L11 Rpl11
1417616_at 0.74881 "ST6 (alpha-N-acetyl-neuraminy-2,3-beta-galactosyl-1,3)-N-
acetylgalactosaminide alpha-2,6-sialyltransferase 2" St6galnac2
1417617_at 1.06229 "ST6 (alpha-N-acetyl-neuraminy-2,3-beta-galactosyl-1,3)-N-
acetylgalactosaminide alpha-2,6-sialyltransferase 2" St6galnac2
1417618_at 0.92386 "inter-alpha trypsin inhibitor, heavy chain 2" Itih2
1417619_at 1.02337 "growth arrest and DNA-damage-inducible, gamma interacting protein 1"
Gadd45gip1
1417620_at 0.86929 RAS-related C3 botulinum substrate 2 Rac2
1417621_at 1.20110 "nuclear factor of activated T-cells, cytoplasmic, calcineurin-dependent 1"
Nfatc1
1417622_at 1.09443 "solute carrier family 12, member 2" Slc12a2
1417623_at 1.12377 "solute carrier family 12, member 2" Slc12a2
1417624_at 1.05657 Ngfi-A binding protein 1 Nab1
1417625_s_at 0.85012 chemokine orphan receptor 1 Cmkor1
1417626_at 0.67085 phosphodiesterase 4D interacting protein (myomegalin) Pde4dip
1417627_a_at 0.86340 "LIM-domain containing, protein kinase" Limk1
1417628_at 0.92733 suppressor of Ty 6 homolog (S. cerevisiae) Supt6h
1417629_at 1.06106 "proline dehydrogenase /// similar to Proline oxidase, mitochondrial precursor
(Proline dehydrogenase)" Prodh /// LOC671160

1417630_at	0.90092	MAP kinase-interacting serine/threonine kinase 1	Mknk1
1417631_at	0.86181	MAP kinase-interacting serine/threonine kinase 1	Mknk1
1417632_at	1.04259	"ATPase, H ⁺ transporting, lysosomal V0 subunit A1"	Atp6v0a1
1417633_at	0.84354	"superoxide dismutase 3, extracellular"	Sod3
1417634_at	0.92216	"superoxide dismutase 3, extracellular"	Sod3
1417635_at	0.55658	sperm autoantigenic protein 17	Spa17
1417636_at	1.11200	"solute carrier family 6 (neurotransmitter transporter, glycine), member 9"	Slc6a9
1417637_a_at	0.94773	high mobility group 20 B	Hmg20b
1417638_at	0.85014	left right determination factor 1	Lefty1
1417639_at	0.79494	"solute carrier family 22 (organic cation transporter), member 4"	Slc22a4
1417640_at	0.95981	CD79B antigen	Cd79b
1417641_at	1.04680	UDP-N-acetyl-alpha-D-galactosamine:polypeptide N-acetylgalactosaminyltransferase-like 5	Galntl5
1417642_at	0.96408	"aldehyde dehydrogenase family 1, subfamily A3"	Aldh1a3
1417643_at	1.10657	testis specific gene A2	Tsga2
1417644_at	0.85280	sarcospan	Sspn
1417645_at	0.85410	sarcospan	Sspn
1417646_a_at	1.08505	sorting nexin 5	Snx5
1417647_at	0.98055	sorting nexin 5	Snx5
1417648_s_at	0.98861	sorting nexin 5	Snx5
1417649_at	0.89878	cyclin-dependent kinase inhibitor 1C (P57)	Cdkn1c
1417650_at	1.13075	L-amino acid oxidase 1	Lao1
1417651_at	1.06326	"cytochrome P450, family 2, subfamily c, polypeptide 29"	Cyp2c29
1417652_a_at	1.04051	tubulin cofactor a	Tbca
1417653_at	1.11626	parvalbumin	Pvalb
1417654_at	1.12873	syndecan 4	Sdc4
1417655_a_at	1.01455	arsenate resistance protein 2	Ars2
1417656_at	1.03065	myeloblastosis oncogene-like 2	Mybl2
1417657_s_at	1.17388	"DnaJ (Hsp40) homolog, subfamily C, member 2"	Dnajc2
1417658_at	1.08542	transforming growth factor beta regulated gene 4	Tbrg4
1417659_at	1.10299	vacuolar protein sorting 29 (S. pombe)	Vps29
1417660_s_at	1.06758	vacuolar protein sorting 29 (S. pombe)	Vps29
1417661_at	0.82841	RAD52 motif 1	Rdm1
1417662_at	1.12238	"ELK3, member of ETS oncogene family"	Elk3
1417663_a_at	0.97037	N-myc downstream regulated gene 3	Ndr3
1417664_a_at	0.84364	N-myc downstream regulated gene 3	Ndr3
1417665_a_at	1.05733	cleavage and polyadenylation specific factor 1	Cpsf1
1417666_at	0.98669	"deoxynucleotidyltransferase, terminal, interacting protein 1"	Dnttip1
1417667_a_at	0.92387	phosphotriesterase related	Pter
1417668_at	0.92631	reticulon 4 interacting protein 1	Rtn4ip1
1417669_at	0.89064	abhydrolase domain containing 12	Abhd12
1417670_at	0.97523	translocase of inner mitochondrial membrane 44	Timm44

1417671_at 0.97005 selenocysteine lyase Scly
1417672_at 1.00315 "solute carrier family 4, sodium bicarbonate cotransporter-like, member 10" Slc4a10
1417673_at 0.65093 growth factor receptor bound protein 14 Grb14
1417674_s_at 1.01981 "golgi autoantigen, golgin subfamily a, 4" Golga4
1417675_a_at 1.40014 midasin homolog (yeast) Mdn1
1417676_a_at 1.06986 "protein tyrosine phosphatase, receptor type, O" Ptpro
1417677_at 1.09161 opsin (encephalopsin) Opn3
1417678_at 0.76799 matrix metalloproteinase 24 Mmp24
1417679_at 1.07539 growth factor independent 1 Gfi1
1417680_at 1.03520 "potassium voltage-gated channel, shaker-related subfamily, member 5" Kcna5
1417681_at 0.99661 nudix (nucleoside diphosphate linked moiety X)-type motif 21 Nudt21
1417682_a_at 0.96975 "protease, serine, 2" Prss2
1417683_at 0.86170 diablo homolog (Drosophila) Diablo
1417684_at 1.01514 THUMP domain containing 3 Thumpd3
1417685_at 0.97326 ankyrin repeat and FYVE domain containing 1 Ankfy1
1417686_at 1.00269 "lectin, galactose binding, soluble 12" Lgals12
1417687_at 0.80451 "lectin, galactose binding, soluble 12" Lgals12
1417688_at 0.81999 cDNA sequence BC004044 BC004044
1417689_a_at 1.07287 PDZK1 interacting protein 1 Pdzk1ip1
1417690_at 0.87938 "protein kinase, AMP-activated, gamma 1 non-catalytic subunit" Prkag1
1417691_at 0.93656 bridging integrator 3 Bin3
1417692_at 0.98326 developmentally regulated GTP binding protein 2 Drg2
1417693_a_at 0.79932 growth factor receptor bound protein 2-associated protein 1 Gab1
1417694_at 0.88669 growth factor receptor bound protein 2-associated protein 1 Gab1
1417695_a_at 0.82375 sterol O-acyltransferase 1 Soat1
1417696_at 0.98772 sterol O-acyltransferase 1 Soat1
1417697_at 0.99673 sterol O-acyltransferase 1 Soat1
1417698_at 1.07730 "general transcription factor IIF, polypeptide 1" Gtf2f1
1417699_at 1.03269 "general transcription factor IIF, polypeptide 1" Gtf2f1
1417700_at 0.76080 "Rab38, member of RAS oncogene family" Rab38
1417701_at 0.78410 "protein phosphatase 1, regulatory (inhibitor) subunit 14c" Ppp1r14c
1417702_a_at 0.87766 histamine N-methyltransferase Hnmt
1417703_at 0.99533 poliovirus receptor-related 2 Pvr12
1417704_a_at 1.08964 Rho GTPase activating protein 6 Arhgap6
1417705_at 0.95493 "OTU domain, ubiquitin aldehyde binding 1" Otub1
1417706_at 0.83616 alpha-N-acetylglucosaminidase (Sanfilippo disease IIIB) Naglu
1417707_at 0.55964 RIKEN cDNA B230342M21 gene B230342M21Rik
1417708_at 0.96653 synaptotagmin III Syt3
1417709_at 0.86783 "cytochrome P450, family 46, subfamily a, polypeptide 1" Cyp46a1
1417710_at 0.98116 methyltransferase like 9 Mettl9
1417711_at 0.94645 RIKEN cDNA 0610012D09 gene 0610012D09Rik
1417712_at 1.12108 "eukaryotic translation initiation factor 2, subunit 2 (beta)" Eif2s2

1417713_at 1.19986 "eukaryotic translation initiation factor 2, subunit 2 (beta)" Eif2s2
1417714_x_at 0.94398 "hemoglobin alpha, adult chain 1" Hba-a1
1417715_a_at 0.94088 "glutamate oxaloacetate transaminase 2, mitochondrial" Got2
1417716_at 0.95675 "glutamate oxaloacetate transaminase 2, mitochondrial" Got2
1417717_a_at 1.10375 tyrosinase Tyr
1417718_at 1.10312 "eukaryotic translation initiation factor 3, subunit 4 (delta)" Eif3s4
1417719_at 1.21874 sin3 associated polypeptide Sap30
1417720_at 1.02665 polymerase (RNA) II (DNA directed) polypeptide J Polr2j
1417721_s_at 0.89291 lysosomal-associated protein transmembrane 5 /// similar to Lysosomal-associated multitransmembrane protein (Retinoic acid-inducible E3 protein) Laptm5 /// LOC669058
1417722_at 1.03348 6-phosphogluconolactonase Pgl
1417723_at 1.06097 "ubiquitin-conjugating enzyme E2, J1" Ube2j1
1417724_at 1.06620 THO complex 4 Thoc4
1417725_a_at 1.02749 Sjogren's syndrome/scleroderma autoantigen 1 homolog (human) Sssca1
1417726_at 1.15957 Sjogren's syndrome/scleroderma autoantigen 1 homolog (human) Sssca1
1417727_at 1.00849 "splicing factor, arginine/serine rich 9" Sfrs9
1417728_at 1.14975 methyl-CpG binding domain protein 3 Mbd3
1417729_at 0.98089 "myosin, heavy polypeptide 6, cardiac muscle, alpha" Myh6
1417730_at 0.84961 exostoses (multiple) 1 Ext1
1417731_at 1.05635 polyglutamine binding protein 1 Pqbp1
1417732_at 1.34093 annexin A8 Anxa8
1417733_at 0.95456 ring finger protein 146 Rnf146
1417734_at 0.85044 A kinase (PRKA) anchor protein 8-like Akap8l
1417735_at 0.91852 RIKEN cDNA 1810030J14 gene /// similar to Serum amyloid P-component precursor (SAP) 1810030J14Rik /// LOC630754
1417736_at 0.94866 structural maintenance of chromosomes 6 Smc6
1417737_at 0.96206 mitochondrial ribosomal protein S31 Mrps31
1417738_at 0.96465 "RAB25, member RAS oncogene family" Rab25
1417739_at 1.09416 "mediator of RNA polymerase II transcription, subunit 11 homolog (S. cerevisiae)" Med11
1417740_at 1.07120 cell division cycle 37 homolog (S. cerevisiae)-like 1 Cdc37l1
1417741_at 0.95478 liver glycogen phosphorylase Pygl
1417742_a_at 0.99374 DNA methyltransferase 1-associated protein 1 Dmap1
1417743_at 1.03647 CDC-like kinase 2 Clk2
1417744_a_at 1.00496 v-ral simian leukemia viral oncogene homolog B (ras related) Ralb
1417745_at 0.87841 "carboxypeptidase N, polypeptide 1" Cpn1
1417746_at 1.01785 complexin 1 Cplx1
1417747_at 0.82133 complexin 1 Cplx1
1417748_x_at 1.05171 forkhead box M1 Foxm1
1417749_a_at 1.01539 tight junction protein 1 Tjp1
1417750_a_at 1.56601 "solute carrier family 25, member 37" Slc25a37
1417751_at 1.08913 serine/threonine kinase 10 Stk10
1417752_at 1.02886 "coronin, actin binding protein 1C" Coro1c

1417753_at 0.86567 polycystic kidney disease 2 Pkd2
1417754_at 0.94708 "topoisomerase I binding, arginine/serine-rich" Topors
1417755_at 0.94079 "topoisomerase I binding, arginine/serine-rich" Topors
1417756_a_at 0.87633 lymphocyte specific 1 Lsp1
1417757_at 1.02162 unc-13 homolog B (C. elegans) Unc13b
1417758_at 0.99455 integrin alpha 2b Itga2b
1417759_at 0.86907 pore forming protein-like Pfp1
1417760_at 0.99535 "nuclear receptor subfamily 0, group B, member 1" Nr0b1
1417761_at 1.01002 apolipoprotein A-IV Apoa4
1417762_a_at 1.05889 ribosomal protein L8 Rpl8
1417763_at 1.03634 "signal sequence receptor, alpha /// similar to signal sequence receptor, alpha" Ssr1 /// LOC636537
1417764_at 0.99497 "signal sequence receptor, alpha /// similar to signal sequence receptor, alpha" Ssr1 /// LOC636537
1417765_a_at 0.87354 "amylase 1, salivary" Amy1
1417766_at 1.17976 cytochrome b5 type B Cyb5b
1417767_at 1.21166 cytochrome b5 type B Cyb5b
1417768_at 0.93029 patatin-like phospholipase domain containing 8 Pnpla8
1417769_at 0.79068 --- ---
1417770_s_at 0.92755 --- ---
1417771_a_at 1.06847 "proteasome (prosome, macropain) 26S subunit, ATPase, 6" Psmc6
1417772_at 1.26842 glyoxylate reductase/hydroxypyruvate reductase Grhpr
1417773_at 1.02208 N-acetylneuraminic acid synthase (sialic acid synthase) Nans
1417774_at 0.99469 N-acetylneuraminic acid synthase (sialic acid synthase) Nans
1417775_at 1.15512 RNA polymerase 1-4 Rpo1-4
1417776_at 0.93792 "alpha-2-glycoprotein 1, zinc" Azgp1
1417777_at 0.84322 leukotriene B4 12-hydroxydehydrogenase Ltb4dh
1417778_at 0.94124 zinc finger protein 35 Zfp35
1417779_at 0.84058 RIKEN cDNA 2310079N02 gene 2310079N02Rik
1417780_at 0.85505 longevity assurance homolog 4 (S. cerevisiae) Lass4
1417781_at 1.10720 longevity assurance homolog 4 (S. cerevisiae) Lass4
1417782_at 1.02963 longevity assurance homolog 4 (S. cerevisiae) Lass4
1417783_at 1.01526 amyotrophic lateral sclerosis 2 (juvenile) homolog (human) Als2
1417784_at 0.95018 amyotrophic lateral sclerosis 2 (juvenile) homolog (human) Als2
1417785_at 1.36871 phospholipase A1 member A Pla1a
1417786_a_at 1.03597 regulator of G-protein signaling 19 Rgs19
1417787_at 0.95262 dickkopf-like 1 Dkk1
1417788_at 1.06433 "synuclein, gamma" Sncg
1417789_at 1.07103 small chemokine (C-C motif) ligand 11 Ccl11
1417790_at 0.99414 docking protein 1 Dok1
1417791_a_at 0.95669 "zinc finger, matrin-like" Zfml
1417792_at 0.94453 "zinc finger, matrin-like" Zfml
1417793_at 1.07889 interferon inducible GTPase 2 Iigp2

1417794_at 0.85512 "zinc finger, MYM-type 3" Zmym3
1417795_at 0.94636 cell adhesion molecule with homology to L1CAM Chl1
1417796_at 0.94844 G protein pathway suppressor 2 Gps2
1417797_a_at 0.97108 RIKEN cDNA 1810019J16 gene 1810019J16Rik
1417798_at 1.02111 RIKEN cDNA 1810019J16 gene 1810019J16Rik
1417799_at 0.94613 "ATPase, H+ transporting, lysosomal V1 subunit G2" Atp6v1g2
1417800_at 0.90499 "poly (ADP-ribose) polymerase family, member 2" Parp2
1417801_a_at 0.97341 "protein tyrosine phosphatase, receptor-type, F interacting protein, binding protein 2" Ppfibp2
1417802_at 0.63353 RIKEN cDNA 1110032A04 gene 1110032A04Rik
1417803_at 0.71940 RIKEN cDNA 1110032A04 gene 1110032A04Rik
1417804_at 1.14406 "RAS, guanyl releasing protein 2" Rasgrp2
1417805_at 1.03471 "X-prolyl aminopeptidase (aminopeptidase P) 2, membrane-bound" Xpnpep2
1417806_at 0.95114 popeye domain containing 2 Popdc2
1417807_at 1.00640 RIKEN cDNA 2700038N03 gene 2700038N03Rik
1417808_at 1.05532 RIKEN cDNA 2310050C09 gene 2310050C09Rik
1417809_at 1.07685 "solute carrier family 22 (organic cation transporter), member 18" Slc22a18
1417810_a_at 0.92313 "potassium voltage gated channel, Shab-related subfamily, member 1 /// protein kinase C and casein kinase substrate in neurons 2" Kcnb1 /// Pacsin2
1417811_at 1.11538 "solute carrier family 24 (sodium/potassium/calcium exchanger), member 6" Slc24a6
1417812_a_at 2.84457 "laminin, beta 3" Lamb3
1417813_at 1.07976 inhibitor of kappaB kinase epsilon Ikbke
1417814_at 0.78960 "phospholipase A2, group V" Pla2g5
1417815_a_at 0.82967 serine incorporator 3 Serinc3
1417816_s_at 0.87731 serine incorporator 3 Serinc3
1417817_a_at 0.91184 WW domain containing transcription regulator 1 Wwtr1
1417818_at 0.86433 WW domain containing transcription regulator 1 Wwtr1
1417819_at 0.94028 "torsin family 1, member B" Tor1b
1417820_at 1.02549 "torsin family 1, member B" Tor1b
1417821_at 1.27607 "DNA segment, Chr 17, human D6S56E 5" D17H6S56E-5
1417822_at 1.24939 "DNA segment, Chr 17, human D6S56E 5" D17H6S56E-5
1417823_at 1.33610 glycine C-acetyltransferase (2-amino-3-ketobutyrate-coenzyme A ligase) Gcat
1417824_at 1.00776 glycine C-acetyltransferase (2-amino-3-ketobutyrate-coenzyme A ligase) Gcat
1417825_at 1.12800 esterase D/formylglutathione hydrolase Esd
1417826_at 0.87806 "aldo-keto reductase family 1, member E1" Akr1e1
1417827_at 0.87923 N-glycanase 1 Ngly1
1417828_at 0.87747 aquaporin 8 Aqp8
1417829_a_at 0.96052 "RAB15, member RAS oncogene family" Rab15
1417830_at 1.02682 structural maintenance of chromosomes 1A Smc1a
1417831_at 0.91836 structural maintenance of chromosomes 1A Smc1a
1417832_at 0.91850 structural maintenance of chromosomes 1A Smc1a
1417833_at 0.96430 zinc finger CCCH type containing 10 Zc3h10

1417834_at 1.01169 synaptojanin 2 binding protein Synj2bp
1417835_at 0.88789 murinoglobulin 1 /// similar to murinoglobulin 1 Mug1 /// LOC669343
1417836_at 1.00095 glutathione peroxidase 7 Gpx7
1417837_at 0.94722 "pleckstrin homology-like domain, family A, member 2" Phlda2
1417838_at 1.05296 spermiogenesis specific transcript on the Y 2 /// similar to RIKEN cDNA
1700029H17 /// similar to spermiogenesis specific transcript on the Y 2 /// similar to spermiogenesis
specific transcript on the Y 2 /// similar to spermiogenesis specific transcript on the Y 2 /// similar to
Spindlin-like protein 2 (SPIN-2) Ssty2 /// LOC382133 /// MGC107098 /// LOC434960 ///
LOC435023 /// LOC666126
1417839_at 0.99714 claudin 5 Cldn5
1417840_at 0.88295 RIKEN cDNA 1500031L02 gene 1500031L02Rik
1417841_at 0.98512 peroxisomal membrane protein 2 Pxmp2
1417842_at 0.96192 calcium modulating ligand Caml
1417843_s_at 0.91490 EPS8-like 2 Eps8l2
1417844_at 1.07040 "mediator of RNA polymerase II transcription, subunit 4 homolog (yeast)"
Med4
1417845_at 1.11477 claudin 6 Cldn6
1417846_at 0.88480 Unc-51 like kinase 2 (C. elegans) Ulk2
1417847_at 0.75750 Unc-51 like kinase 2 (C. elegans) Ulk2
1417848_at 0.76773 zinc finger protein 704 Zfp704
1417849_at 0.83715 zinc finger protein 704 Zfp704
1417850_at 0.89033 retinoblastoma 1 Rb1
1417851_at 0.95343 chemokine (C-X-C motif) ligand 13 Cxcl13
1417852_x_at 1.57536 chloride channel calcium activated 1 Clca1
1417853_at 1.19399 chloride channel calcium activated 1 Clca1
1417854_at 1.02090 mitogen activated protein kinase kinase 5 Map2k5
1417855_at 1.07605 F-box and leucine-rich repeat protein 15 Fbxl15
1417856_at 1.28754 avian reticuloendotheliosis viral (v-rel) oncogene related B Relb
1417857_at 0.86684 methylmalonic aciduria (cobalamin deficiency) type A Mmaa
1417858_at 1.21568 RAS protein activator like 1 (GAP1 like) Rasal1
1417859_at 1.16055 growth arrest specific 7 Gas7
1417860_a_at 0.40565 "spondin 2, extracellular matrix protein" Spon2
1417861_at 0.99463 cyclin C Ccnc
1417862_at 0.95405 RIKEN cDNA A830059I20 gene A830059I20Rik
1417863_at 1.09326 spermatogenesis associated 19 Spata19
1417864_at 1.28497 phosphoglycerate kinase 1 /// phosphoglycerate kinase 1-like ///
phosphoglycerate kinase 1-like Pgk1 /// LOC433594 /// LOC668435
1417865_at 1.00840 "tumor necrosis factor, alpha-induced protein 1 (endothelial)" Tnfaip1
1417866_at 1.10293 "tumor necrosis factor, alpha-induced protein 1 (endothelial)" Tnfaip1
1417867_at 1.13216 complement factor D (adipsin) Cfd
1417868_a_at 0.95475 cathepsin Z Ctsz
1417869_s_at 1.02929 cathepsin Z Ctsz
1417870_x_at 1.03158 cathepsin Z Ctsz

1417871_at 1.35922 hydroxysteroid (17-beta) dehydrogenase 7 Hsd17b7
1417872_at 0.62267 four and a half LIM domains 1 Fhl1
1417873_at 1.01973 PWP1 homolog (S. cerevisiae) Pwp1
1417874_at 0.96069 "TMEM9 domain family, member B" Tmem9b
1417875_at 0.98488 DEAD (Asp-Glu-Ala-Asp) box polypeptide 50 Ddx50
1417876_at 0.89900 "Fc receptor, IgG, high affinity I" Fcgr1
1417877_at 0.69834 RIKEN cDNA 2310005P05 gene 2310005P05Rik
1417878_at 1.01801 E2F transcription factor 1 E2f1
1417879_at 1.03350 neuron derived neurotrophic factor Nenf
1417880_at 1.03680 "glucose-6-phosphatase, catalytic" G6pc
1417881_at 0.77602 "solute carrier family 39 (zinc transporter), member 3" Slc39a3
1417882_at 1.01731 "solute carrier family 39 (zinc transporter), member 3" Slc39a3
1417883_at 0.97136 "glutathione S-transferase, theta 2" Gstt2
1417884_at 0.79989 "solute carrier family 16 (monocarboxylic acid transporters), member 6" Slc16a6
1417885_at 0.96175 microtubule-associated protein tau Mapt
1417886_at 1.12470 RIKEN cDNA 1810009A15 gene 1810009A15Rik
1417887_at 0.98708 integrator complex subunit 5 Ints5
1417888_at 0.85598 tripartite motif protein 13 Trim13
1417889_at 1.02252 apolipoprotein B editing complex 2 Apobec2
1417890_at 1.21027 "pyridoxal (pyridoxine, vitamin B6) phosphatase" Pdxp
1417891_at 0.90407 splA/ryanodine receptor domain and SOCS box containing 3 Spsb3
1417892_a_at 0.92210 "sirtuin 3 (silent mating type information regulation 2, homolog) 3 (S. cerevisiae)" Sirt3
1417893_at 0.99118 sideroflexin 3 Sfxn3
1417894_at 0.88743 G protein-coupled receptor 97 Gpr97
1417895_a_at 1.07257 transmembrane protein 54 Tmem54
1417896_at 1.09135 tight junction protein 3 Tjp3
1417897_at 1.02274 breast cancer metastasis-suppressor 1 Brms1
1417898_a_at 0.98415 granzyme A Gzma
1417899_at 0.88862 zona pellucida 3 receptor Zp3r
1417900_a_at 1.02711 very low density lipoprotein receptor Vldlr
1417901_a_at 0.97836 islet cell autoantigen 1 Ica1
1417902_at 1.37062 "solute carrier family 19 (thiamine transporter), member 2" Slc19a2
1417903_at 1.02101 "deafness, autosomal dominant 5 homolog (human)" Dfna5h
1417904_at 1.00548 "DNA cross-link repair 1A, PSO2 homolog (S. cerevisiae)" Dclre1a
1417905_at 1.04788 prolactin-like protein F Prlpf
1417906_at 0.98655 RIKEN cDNA 1700001F09 gene 1700001F09Rik
1417907_at 0.94764 ubiquitin-conjugating enzyme E2L 3 /// ribosomal protein S28 Ube213 /// Rps28
1417908_s_at 0.98703 ubiquitin-conjugating enzyme E2L 3 /// ribosomal protein S28 Ube213 /// Rps28
1417909_at 1.03349 "serine (or cysteine) peptidase inhibitor, clade C (antithrombin), member 1" Serpincl1

1417910_at	0.99070 cyclin A2	Ccna2	
1417911_at	1.01006 cyclin A2	Ccna2	
1417912_at	1.04663 transmembrane protein 93	Tmem93	
1417913_at	1.02832 GINS complex subunit 4 (Sld5 homolog)	Gins4	
1417914_at	1.09867 "RAP2B, member of RAS oncogene family"	Rap2b	
1417915_at	1.12389 "RAP2B, member of RAS oncogene family"	Rap2b	
1417916_a_at	1.06135 fractured callus expressed transcript 1	Fxc1	
1417917_at	0.79353 calponin 1	Cnn1	
1417918_at	1.05530 mitochondrial ribosomal protein L11	Mrpl11	
1417919_at	1.09856 "protein phosphatase 1, regulatory (inhibitor) subunit 7"		Ppp1r7
1417920_at	1.09737 amnionless	Amn	
1417921_at	1.11137 RIKEN cDNA 2610029G23 gene	2610029G23Rik	
1417922_at	0.94526 kelch repeat and BTB (POZ) domain containing 4	Kbtbd4	
1417923_at	0.95076 p21 (CDKN1A)-activated kinase 3	Pak3	
1417924_at	1.04005 p21 (CDKN1A)-activated kinase 3	Pak3	
1417925_at	0.95256 chemokine (C-C motif) ligand 22	Ccl22	
1417926_at	0.96624 leucine zipper protein 5	Luzp5	
1417927_at	0.98815 DEAD (Asp-Glu-Ala-Asp) box polypeptide 19a		Ddx19a
1417928_at	0.86884 PDZ and LIM domain 4	Pdlim4	
1417929_at	0.96560 "solute carrier family 7 (cationic amino acid transporter, y+ system), member 8"		Slc7a8
1417930_at	0.98625 Ngfi-A binding protein 2	Nab2	
1417931_at	0.90337 N-deacetylase/N-sulfotransferase (heparan glucosaminyl) 2		Ndst2
1417932_at	1.17634 interleukin 18	Il18	
1417933_at	0.82741 insulin-like growth factor binding protein 6	Igfbp6	
1417934_at	0.98286 "DnaJ (Hsp40) homolog, subfamily C, member 4"	Dnajc4	
1417935_at	0.85409 "makorin, ring finger protein, 2"	Mkrn2	
1417936_at	1.06849 chemokine (C-C motif) ligand 9	Ccl9	
1417937_at	1.05583 "dapper homolog 1, antagonist of beta-catenin (xenopus)"		Dact1
1417938_at	0.94311 RAD51 associated protein 1	Rad51ap1	
1417939_at	1.07755 RAD51 associated protein 1	Rad51ap1	
1417940_s_at	1.06734 RAD51 associated protein 1	Rad51ap1	
1417941_at	0.94186 N-acetylneuraminic acid phosphatase	Nanp	
1417942_at	1.12077 Ly6/Plaur domain containing 3	Lypd3	
1417943_at	0.96017 "guanine nucleotide binding protein (G protein), gamma 4 subunit"		Gng4
1417944_at	1.00339 "guanine nucleotide binding protein (G protein), gamma 4 subunit"		Gng4
1417945_at	1.00058 "POU domain, class 5, transcription factor 1"	Pou5f1	
1417946_at	0.88625 abhydrolase domain containing 3	Abhd3	
1417947_at	1.00464 proliferating cell nuclear antigen	Pcna	
1417948_s_at	0.87972 interleukin enhancer binding factor 2	Ilf2	
1417949_at	0.96070 interleukin enhancer binding factor 2	Ilf2	
1417950_a_at	0.91159 apolipoprotein A-II	Apoa2	
1417951_at	0.96820 "enolase 3, beta muscle"	Eno3	

1417952_at 1.04168 "cytochrome P450, family 2, subfamily j, polypeptide 6" Cyp2j6
 1417953_at 0.73023 "DNA segment, Chr 6, Wayne State University 176, expressed" D6Wsu176e
 1417954_at 0.94865 somatostatin Sst
 1417955_at 0.90201 coiled-coil domain containing 71 Ccdc71
 1417956_at 0.93385 "cell death-inducing DNA fragmentation factor, alpha subunit-like effector A"
 Cidea
 1417957_a_at 1.07471 tetraspanin 1 Tspan1
 1417958_at 0.86494 tetraspanin 1 Tspan1
 1417959_at 0.79904 PDZ and LIM domain 7 Pdlim7
 1417960_at 1.18421 cytoplasmic polyadenylation element binding protein 1 Cpeb1
 1417961_a_at 1.00885 tripartite motif protein 30 Trim30
 1417962_s_at 0.53859 growth hormone receptor Ghr
 1417963_at 0.87487 phospholipid transfer protein Pltp
 1417964_at 1.05054 "adaptor-related protein complex 3, delta 1 subunit" Ap3d1
 1417965_at 0.90664 "pleckstrin homology domain containing, family A (phosphoinositide binding
 specific) member 1" Plekha1
 1417966_at 1.03766 mitochondrial ribosomal protein L39 Mrpl39
 1417967_at 1.07562 MMS19 (MET18 *S. cerevisiae*)-like Mms19l
 1417968_a_at 0.90452 methyl-CpG binding domain protein 1 Mbd1
 1417969_at 0.88317 F-box only protein 31 Fbxo31
 1417970_at 0.94513 "ATP synthase, H⁺ transporting, mitochondrial F0 complex, subunit s" Atp5s
 1417971_at 0.83518 nurim (nuclear envelope membrane protein) Nrm
 1417972_s_at 0.97548 "processing of precursor 5, ribonuclease P/MRP family (*S. cerevisiae*)" Pop5
 1417973_at 1.17955 "inter-alpha trypsin inhibitor, heavy chain 1" Itih1
 1417974_at 1.00425 karyopherin (importin) alpha 4 Kpna4
 1417975_at 1.03056 karyopherin (importin) alpha 4 Kpna4
 1417976_at 0.99701 adenosine deaminase Ada
 1417977_at 0.60174 eukaryotic translation initiation factor 4E member 3 Eif4e3
 1417978_at 0.73109 eukaryotic translation initiation factor 4E member 3 Eif4e3
 1417979_at 0.88067 tenomodulin Tnmd
 1417980_a_at 1.09204 insulin induced gene 2 Insig2
 1417981_at 1.10784 insulin induced gene 2 Insig2
 1417982_at 1.06362 insulin induced gene 2 Insig2
 1417983_a_at 1.08230 ubiquitin-conjugating enzyme E2 variant 2 Ube2v2
 1417984_at 0.95546 ubiquitin-conjugating enzyme E2 variant 2 Ube2v2
 1417985_at 0.98495 Notch-regulated ankyrin repeat protein Nrarp
 1417986_at 0.94155 Notch-regulated ankyrin repeat protein Nrarp
 1417987_at 0.93951 biotinidase Btd
 1417988_at 0.82019 regulated endocrine-specific protein 18 Resp18
 1417989_at 1.07781 testis-specific serine kinase 1 Tssk1
 1417990_at 1.03221 "protein phosphatase 1, regulatory (inhibitor) subunit 14D" Ppp1r14d
 1417991_at 0.91852 "deiodinase, iodothyronine, type I" Dio1
 1417992_a_at 1.01249 seminal vesicle secretion 3 /// RIKEN cDNA 9530004A22 Svs3 ///

9530004A22Rik

1417993_at 1.04123 seminal vesicle secretion 3 Svs3
 1417994_a_at 0.97698 "amiloride-sensitive cation channel 1, neuronal (degenerin)" Accn1
 1417995_at 0.99895 "protein tyrosine phosphatase, non-receptor type 22 (lymphoid)" Ptpn22
 1417996_at 0.96047 neuroglobin Ngb
 1417997_at 0.96113 neuroglobin Ngb
 1417998_at 0.99327 prostaglandin E synthase 3 (cytosolic) Ptges3
 1417999_at 0.85101 integral membrane protein 2B Itm2b
 1418000_a_at 0.92444 integral membrane protein 2B Itm2b
 1418001_at 0.95150 integral membrane protein 2B Itm2b
 1418002_at 0.95535 "HIG1 domain family, member 2A" Higd2a
 1418003_at 0.92455 RIKEN cDNA 1190002H23 gene 1190002H23Rik
 1418004_a_at 1.16945 RIKEN cDNA 1810009M01 gene 1810009M01Rik
 1418005_at 0.92616 "succinate dehydrogenase complex, subunit B, iron sulfur (Ip)" Sdhb
 1418006_at 1.08561 RIKEN cDNA 5830416A07 gene 5830416A07Rik
 1418007_at 0.99837 RIKEN cDNA 1810007M14 gene 1810007M14Rik
 1418008_at 1.17711 RIKEN cDNA 1810007M14 gene 1810007M14Rik
 1418009_at 0.96661 RIKEN cDNA 1810007M14 gene 1810007M14Rik
 1418010_a_at 1.10224 SH3-domain GRB2-like B1 (endophilin) Sh3glb1
 1418011_a_at 0.98900 SH3-domain GRB2-like B1 (endophilin) Sh3glb1
 1418012_at 0.89263 SH3-domain GRB2-like B1 (endophilin) Sh3glb1
 1418013_at 0.95409 camello-like 1 Cml1
 1418014_a_at 0.95815 "UDP-Gal:betaGlcNAc beta 1,4- galactosyltransferase, polypeptide 1"
 B4galt1
 1418015_at 1.03963 pumilio 2 (Drosophila) Pum2
 1418016_at 1.02600 pumilio 2 (Drosophila) Pum2
 1418017_at 0.94078 pumilio 2 (Drosophila) Pum2
 1418018_at 0.98809 carboxypeptidase D Cpd
 1418019_at 0.90784 carboxypeptidase D Cpd
 1418020_s_at 0.92703 carboxypeptidase D Cpd
 1418021_at 0.78389 complement component 4B (Childo blood group) /// similar to Complement C4
 precursor C4b /// LOC675521
 1418022_at 1.16188 NMDA receptor-regulated gene 1 Narg1
 1418023_at 1.22904 NMDA receptor-regulated gene 1 Narg1
 1418024_at 1.17614 NMDA receptor-regulated gene 1 Narg1
 1418025_at 1.40296 "basic helix-loop-helix domain containing, class B2" Bhlhb2
 1418026_at 1.06307 exonuclease 1 Exo1
 1418027_at 1.04014 exonuclease 1 Exo1
 1418028_at 0.95253 dopachrome tautomerase Dct
 1418029_at 1.14920 Fas apoptotic inhibitory molecule Faim
 1418030_at 1.02086 "solute carrier organic anion transporter family, member 3a1" Slco3a1
 1418031_at 0.90709 myosin IXb Myo9b
 1418032_at 0.97819 integrin alpha FG-GAP repeat containing 2 Itfg2

1418033_s_at 0.96753 zinc finger protein 535 Zfp535
1418034_at 1.09175 mitochondrial ribosomal protein S9 Mrps9
1418035_a_at 0.91551 "DNA primase, p58 subunit" Prim2
1418036_at 0.96506 "DNA primase, p58 subunit" Prim2
1418037_at 0.61867 complement component 4 binding protein C4bp
1418038_s_at 1.13734 dual specificity phosphatase 19 Dusp19
1418039_at 0.91151 "RAB, member of RAS oncogene family-like 3" Rabl3
1418040_at 0.94935 RIKEN cDNA 4432406C05 gene 4432406C05Rik
1418041_at 0.97044 RIKEN cDNA 4432406C05 gene 4432406C05Rik
1418042_a_at 1.07270 "ATP-binding cassette, sub-family C (CFTR/MRP), member 5" Abcc5
1418043_at 0.96876 "ATP-binding cassette, sub-family C (CFTR/MRP), member 5" Abcc5
1418044_at 1.07786 RIKEN cDNA A630042L21 gene A630042L21Rik
1418045_at 1.00570 inositol polyphosphate-1-phosphatase Inpp1
1418046_at 0.90269 nucleosome assembly protein 1-like 2 Nap1l2
1418047_at 0.99155 neurogenic differentiation 6 Neurod6
1418048_at 0.99994 RIKEN cDNA 1110059G10 gene 1110059G10Rik
1418049_at 0.74271 latent transforming growth factor beta binding protein 3 Ltbp3
1418050_at 1.00046 glycosylphosphatidylinositol specific phospholipase D1 Gpld1
1418051_at 1.07052 Eph receptor B6 Ephb6
1418052_at 1.23049 mevalonate kinase Mvk
1418053_at 0.92221 "synuclein, beta" Sncb
1418054_at 0.96857 neurogenic differentiation 4 Neurod4
1418055_at 0.93873 neurogenic differentiation 4 Neurod4
1418056_at 0.88691 IQ motif containing F3 Iqcf3
1418057_at 0.98356 T-cell lymphoma invasion and metastasis 1 Tiam1
1418058_at 1.09774 "EGF, latrophilin seven transmembrane domain containing 1" Eltd1
1418059_at 1.12943 "EGF, latrophilin seven transmembrane domain containing 1" Eltd1
1418060_a_at 1.01179 mitogen activated protein kinase 7 Mapk7
1418061_at 0.99764 latent transforming growth factor beta binding protein 2 Ltbp2
1418062_at 1.02918 eukaryotic translation elongation factor 1 alpha 2 Eef1a2
1418063_at 1.11111 keratocan Kera
1418064_at 1.00745 TCF3 (E2A) fusion partner Tfpt
1418065_at 0.98598 recombination activating gene 2 Rag2
1418066_at 1.02413 "cofilin 2, muscle" Cfl2
1418067_at 0.96735 "cofilin 2, muscle" Cfl2
1418068_at 0.93466 NADH dehydrogenase (ubiquinone) 1 alpha subcomplex 10 Ndufa10
1418069_at 1.01659 apolipoprotein C-II Apoc2
1418070_at 0.79735 "chromodomain protein, Y chromosome-like" Cdyl
1418071_s_at 0.96785 "chromodomain protein, Y chromosome-like" Cdyl
1418072_at 0.85060 "histone 1, H2bc" Hist1h2bc
1418073_at 1.15689 acyl-CoA thioesterase 9 Acot9
1418074_at 0.85209 "ST6 (alpha-N-acetyl-neuraminyl-2,3-beta-galactosyl-1,3)-N-acetylgalactosaminide alpha-2,6-sialyltransferase 4" St6galnac4

1418075_at 0.87154 "ST6 (alpha-N-acetyl-neuraminyl-2,3-beta-galactosyl-1,3)-N-acetylgalactosaminide alpha-2,6-sialyltransferase 4" St6galnac4
 1418076_at 1.00795 suppression of tumorigenicity 14 (colon carcinoma) St14
 1418077_at 0.97398 tripartite motif protein 21 Trim21
 1418078_at 1.00083 "proteasome (prosome, macropain) 28 subunit, 3" Psme3
 1418079_at 1.07329 "proteasome (prosome, macropain) 28 subunit, 3" Psme3
 1418080_at 0.95462 "UDP-Gal:betaGlcNAc beta 1,4- galactosyltransferase, polypeptide 2" B4galt2
 1418081_at 0.98768 Williams-Beuren syndrome chromosome region 18 homolog (human) Wbscr18
 1418082_at 1.05083 N-myristoyltransferase 1 Nmt1
 1418083_at 0.97923 RIKEN cDNA 0610009B22 gene 0610009B22Rik
 1418084_at 0.53265 neuropilin 1 Nrp1
 1418085_at 0.83192 "protein kinase C, zeta" Prkcz
 1418086_at 1.11804 "protein phosphatase 1, regulatory (inhibitor) subunit 14A" Ppp1r14a
 1418087_at 0.98440 ubiquitin fusion degradation 1 like Ufd1l
 1418088_a_at 1.03883 syntaxin 8 Stx8
 1418089_at 0.99884 syntaxin 8 Stx8
 1418090_at 0.89722 plasmalemma vesicle associated protein Plvap
 1418091_at 1.05527 transcription factor CP2-like 1 Tcfcp2l1
 1418092_s_at 0.98423 thyroid hormone receptor interactor 10 Trip10
 1418093_a_at 0.93632 epidermal growth factor Egf
 1418094_s_at 1.05338 carbonic anhydrase 4 Car4
 1418095_at 1.03407 "small muscle protein, X-linked" Smpx
 1418096_at 0.99878 docking protein 3 Dok3
 1418097_a_at 0.86173 transmembrane phosphoinositide 3-phosphatase and tensin homolog 2 Tpte2
 1418098_at 1.04089 adenylate cyclase 4 Adcy4
 1418099_at 1.35003 "tumor necrosis factor receptor superfamily, member 1b" Tnfrsf1b
 1418100_at 0.77640 RIKEN cDNA A030009H04 gene A030009H04Rik
 1418101_a_at 0.86977 reticulon 3 Rtn3
 1418102_at 0.61345 hairy and enhancer of split 1 (Drosophila) Hes1
 1418103_at 0.89938 "solute carrier family 12 (potassium/chloride transporters), member 9" Slc12a9
 1418104_at 0.93698 nuclear receptor interacting protein 3 Nrip3
 1418105_at 0.81646 stathmin-like 4 Stmn4
 1418106_at 1.02971 hairy/enhancer-of-split related with YRPW motif 2 Hey2
 1418107_at 1.08740 "transcription elongation factor A (SII), 2" Tcea2
 1418108_at 0.89929 "pleckstrin homology domain containing, family K member 1" Plekhk1
 1418109_at 1.06519 G1 to S phase transition 2 Gspt2
 1418110_a_at 0.97610 inositol polyphosphate-5-phosphatase D Inpp5d
 1418111_at 0.91070 RIKEN cDNA 1810008K16 gene 1810008K16Rik
 1418112_at 0.97988 mitochondrial ribosomal protein L10 Mrpl10
 1418113_at 0.91507 "cytochrome P450, family 2, subfamily d, polypeptide 10" Cyp2d10
 1418114_at 0.98325 recombining binding protein suppressor of hairless (Drosophila) Rbpsuh

1418115_s_at 1.02529 interferon alpha responsive gene Ifrg15
1418116_at 1.06647 interferon alpha responsive gene Ifrg15
1418117_at 0.77753 NADH dehydrogenase (ubiquinone) Fe-S protein 4 Ndufs4
1418118_at 0.87742 "solute carrier family 22 (organic cation transporter), member 1" Slc22a1
1418119_at 1.14761 RNA binding motif protein 8a Rbm8a
1418120_at 0.95470 RNA binding motif protein 8a Rbm8a
1418121_at 0.96169 vaccinia related kinase 3 Vrk3
1418122_at 0.96547 "myosin, heavy polypeptide 11, smooth muscle" Myh11
1418123_at 0.98364 unc-119 homolog (C. elegans) Unc119
1418124_at 1.05534 transmembrane protein 85 Tmem85
1418125_at 0.89590 INO80 complex homolog 1 (S. cerevisiae) Inoc1
1418126_at 1.51512 chemokine (C-C motif) ligand 5 Ccl5
1418127_a_at 0.99988 programmed cell death 8 Pcdcd8
1418128_at 1.06002 adenylate cyclase 6 Adcy6
1418129_at 1.25496 24-dehydrocholesterol reductase Dhcr24
1418130_at 1.00916 --- ---
1418131_at 1.11840 "SAM domain and HD domain, 1" Samhd1
1418132_a_at 0.98321 "DNA segment, Chr 7, Wayne State University 128, expressed" D7Wsu128e
1418133_at 1.23037 B-cell leukemia/lymphoma 3 Bcl3
1418134_at 0.96384 RIKEN cDNA 1200007B05 gene 1200007B05Rik
1418135_at 0.91487 "AF4/FMR2 family, member 1" Aff1
1418136_at 0.75019 transforming growth factor beta 1 induced transcript 1 Tgfb1i1
1418137_at 1.02001 mitochondrial ribosomal protein 63 Mrp63
1418138_at 1.07777 "sulfotransferase family 1D, member 1" Sult1d1
1418139_at 0.86034 doublecortin Dcx
1418140_at 0.94991 doublecortin Dcx
1418141_at 0.87884 doublecortin Dcx
1418142_at 0.88936 "potassium inwardly-rectifying channel, subfamily J, member 8" Kcnj8
1418143_at 0.94967 vacuolar protein sorting 45 (yeast) Vps45
1418144_a_at 1.09624 "phosphatidylinositol-4-phosphate 5-kinase, type 1 beta" Pip5k1b
1418145_at 0.95571 tuftelin interacting protein 11 Tfip11
1418146_a_at 0.93790 retinoblastoma-like 2 /// similar to retinoblastoma-like 2 Rbl2 ///
LOC635075
1418147_at 0.92121 "transcription factor AP-2, gamma" Tcfap2c
1418148_at 1.12614 abhydrolase domain containing 1 Abhd1
1418149_at 1.12299 chromogranin A Chga
1418150_at 0.84602 myotubularin related protein 4 Mtmr4
1418151_at 0.94960 myotubularin related protein 4 Mtmr4
1418152_at 1.05585 nucleosome binding protein 1 Nsbp1
1418153_at 1.10541 "laminin, alpha 1" Lama1
1418154_at 1.11484 cDNA sequence BC004022 BC004022
1418155_at 0.95911 myotilin Myot
1418156_at 0.99436 "potassium voltage-gated channel, Isk-related subfamily, gene 4" Kcne4

1418157_at	0.90146	"nuclear receptor subfamily 2, group F, member 1"	Nr2f1
1418158_at	0.91391	transformation related protein 63	Trp63
1418159_at	0.91102	transcription factor CP2	Tcfcp2
1418160_at	0.98316	"makorin, ring finger protein, 3"	Mkrm3
1418161_at	0.90665	junctophilin 3	Jph3
1418162_at	1.16751	toll-like receptor 4	Tlr4
1418163_at	0.98418	toll-like receptor 4	Tlr4
1418164_at	0.85931	syntaxin 2	Stx2
1418165_at	0.90554	intelectin a	Itlna
1418166_at	1.05565	"interleukin 12 receptor, beta 1"	Il12rb1
1418167_at	1.06235	transcription factor AP4	Tcfap4
1418168_at	1.01998	"zinc finger, CCHC domain containing 14"	Zcchc14
1418169_at	0.98442	"zinc finger, CCHC domain containing 14"	Zcchc14
1418170_a_at	0.98968	"zinc finger, CCHC domain containing 14"	Zcchc14
1418171_at	0.86657	transcription elongation factor A (SII)-like 8	Tceal8
1418172_at	0.93908	heme binding protein 1	Hebp1
1418173_at	1.02251	keratin 25	Krt25
1418174_at	0.77592	D site albumin promoter binding protein	Dbp
1418175_at	0.86176	vitamin D receptor	Vdr
1418176_at	0.91094	vitamin D receptor	Vdr
1418177_at	1.06887	"gamma-aminobutyric acid (GABA-A) receptor, subunit gamma 2"	Gabrg2
1418178_at	0.88153	"internexin neuronal intermediate filament protein, alpha"	Ina
1418179_at	0.84074	autophagy-related 3 (yeast)	Atg3
1418180_at	0.92965	trans-acting transcription factor 1	Sp1
1418181_at	0.65415	protein tyrosine phosphatase 4a3	Ptp4a3
1418182_at	0.88831	protein tyrosine phosphatase 4a3	Ptp4a3
1418183_a_at	0.87633	"pleckstrin homology, Sec7 and coiled-coil domains 1"	Pscd1
1418184_at	1.06037	centromere protein M	Cenpm
1418185_at	1.02630	RIKEN cDNA 4733401H18 gene	4733401H18Rik
1418186_at	0.87018	"glutathione S-transferase, theta 1"	Gstt1
1418187_at	1.17684	receptor (calcitonin) activity modifying protein 2	Ramp2
1418188_a_at	1.81155	Metastasis associated lung adenocarcinoma transcript 1 (non-coding RNA)	Malat1
1418189_s_at	1.86723	Metastasis associated lung adenocarcinoma transcript 1 (non-coding RNA)	Malat1
1418190_at	1.08157	paraoxonase 1	Pon1
1418191_at	0.99325	ubiquitin specific peptidase 18	Usp18
1418192_at	1.05475	max binding protein	Mnt
1418193_at	0.97499	similar to max binding protein	LOC672705
1418194_at	0.93270	UDP-N-acetyl-alpha-D-galactosamine:polypeptide N-acetylgalactosaminyltransferase 10	Galnt10
1418195_at	0.85001	UDP-N-acetyl-alpha-D-galactosamine:polypeptide N-acetylgalactosaminyltransferase 10	Galnt10

1418196_at 1.10093 telomerase associated protein 1 Tep1
1418197_at 0.98517 "uncoupling protein 1 (mitochondrial, proton carrier)" Ucp1
1418198_a_at 0.87048 transmembrane 9 superfamily member 1 Tm9sf1
1418199_at 1.05126 hemogen Hemgn
1418200_at 0.96979 GLI-Kruppel family member HKR3 Hkr3
1418201_at 1.00873 "pleckstrin homology domain containing, family G (with RhoGef domain) member 2" Plekhg2
1418202_a_at 1.03920 widely-interspaced zinc finger motifs Wiz
1418203_at 1.21457 phorbol-12-myristate-13-acetate-induced protein 1 Pmaip1
1418204_s_at 1.22016 allograft inflammatory factor 1 Aif1
1418205_at 0.77050 --- ---
1418206_at 0.96384 stromal cell-derived factor 2-like 1 Sdf2l1
1418207_at 0.97560 FXYD domain-containing ion transport regulator 4 Fxyd4
1418208_at 1.08617 paired box gene 8 Pax8
1418209_a_at 0.72224 profilin 2 Pfn2
1418210_at 0.74180 profilin 2 Pfn2
1418211_at 0.90887 pink-eyed dilution p
1418212_at 0.99814 oligodendrocyte myelin glycoprotein Omg
1418213_at 0.98975 keratin 23 Krt23
1418214_at 0.91987 kinesin light chain 2 Klc2
1418215_at 1.00295 meprin 1 beta Mep1b
1418216_at 0.99592 gamma-glutamyltransferase-like activity 1 Ggta1
1418217_at 0.92420 "non-metastatic cells 7, protein expressed in" Nme7
1418218_at 1.03025 CEA-related cell adhesion molecule 9 Ceacam9
1418219_at 0.81493 interleukin 15 Il15
1418220_at 0.81340 forkhead box F2 Foxf2
1418221_at 0.99883 casein alpha s2-like B Csn1s2b
1418222_at 1.14041 RIKEN cDNA 2610024G14 gene 2610024G14Rik
1418223_at 1.02653 Sec11-like 1 (S. cerevisiae) Sec11l1
1418224_at 0.98960 Sec11-like 1 (S. cerevisiae) Sec11l1
1418225_at 1.08148 "origin recognition complex, subunit 2-like (S. cerevisiae)" Orc2l
1418226_at 1.05266 "origin recognition complex, subunit 2-like (S. cerevisiae)" Orc2l
1418227_at 1.10050 "origin recognition complex, subunit 2-like (S. cerevisiae)" Orc2l
1418228_at 1.01806 histone cell cycle regulation defective interacting protein 5 Hirip5
1418229_s_at 1.05930 histone cell cycle regulation defective interacting protein 5 Hirip5
1418230_a_at 1.01058 LIM and senescent cell antigen-like domains 1 Lims1
1418231_at 1.01843 LIM and senescent cell antigen-like domains 1 Lims1
1418232_s_at 1.06847 LIM and senescent cell antigen-like domains 1 Lims1
1418233_a_at 1.07653 trafficking protein particle complex 5 Trappc5
1418234_s_at 0.94445 breast carcinoma amplified sequence 2 Bcas2
1418235_at 1.01458 autophagy-related 5 (yeast) Atg5
1418236_s_at 1.01838 autophagy-related 5 (yeast) Atg5
1418237_s_at 1.00791 "procollagen, type XVIII, alpha 1" Col18a1

1418238_at 0.95753 isovaleryl coenzyme A dehydrogenase Ivd
1418239_at 0.89778 apolipoprotein F Apof
1418240_at 1.38599 guanylate nucleotide binding protein 2 Gbp2
1418241_at 0.98878 upstream transcription factor 2 Usf2
1418242_at 1.02596 Fas-associated factor 1 Faf1
1418243_at 0.84868 ficolin A Fcna
1418244_at 1.01507 "N-acetyltransferase 5 (ARD1 homolog, *S. cerevisiae*)" Nat5
1418245_a_at 0.94846 RNA binding motif protein 9 Rbm9
1418246_at 1.01239 RNA binding motif protein 9 Rbm9
1418247_s_at 1.01169 RNA binding motif protein 9 Rbm9
1418248_at 1.13355 "galactosidase, alpha" Gla
1418249_at 1.03093 calcitonin gene-related peptide-receptor component protein Crcp
1418250_at 0.73351 ADP-ribosylation factor 4-like Arfl4
1418251_at 0.89958 tubby-like protein 3 Tulp3
1418252_at 0.87931 "peptidyl arginine deiminase, type II /// similar to peptidyl arginine deiminase, type II" Padi2 /// LOC638935
1418253_a_at 1.06891 heat shock protein 4 like Hspa4l
1418254_at 0.91675 APAF1 interacting protein Apip
1418255_s_at 0.75916 serum response factor Srf
1418256_at 0.89397 serum response factor Srf
1418257_at 0.79408 "solute carrier family 12, member 7" Slc12a7
1418258_s_at 0.97139 dynein light chain LC8-type 2 Dynll2
1418259_a_at 0.88792 ectonucleoside triphosphate diphosphohydrolase 2 Entpd2
1418260_at 0.95497 hormonally upregulated Neu-associated kinase /// similar to hormonally upregulated Neu-associated kinase Hunk /// LOC630567
1418261_at 0.96973 spleen tyrosine kinase Syk
1418262_at 1.06117 spleen tyrosine kinase Syk
1418263_at 1.01921 DEAD (Asp-Glu-Ala-Asp) box polypeptide 25 Ddx25
1418264_at 0.95232 centromere protein K Cenpk
1418265_s_at 0.99860 interferon regulatory factor 2 Irf2
1418266_at 1.01553 "arachidonate 12-lipoxygenase, 12R type" Alox12b
1418267_at 0.96355 macrophage stimulating 1 (hepatocyte growth factor-like) Mst1
1418268_at 0.98104 5-hydroxytryptamine (serotonin) receptor 3A Htr3a
1418269_at 0.86697 lysyl oxidase-like 3 Loxl3
1418270_at 1.00525 "a disintegrin-like and metallopeptidase (reprolysin type) with thrombospondin type 1 motif, 8" Adamts8
1418271_at 1.01506 "basic helix-loop-helix domain containing, class B5" Bhlhb5
1418272_at 0.86561 3-oxoacid CoA transferase 2A Oxct2a
1418273_a_at 1.06138 ribosomal protein L30 /// similar to ribosomal protein L30 /// similar to ribosomal protein L30 /// similar to ribosomal protein L30 Rpl30 /// LOC666899 /// LOC670014 /// LOC675333 /// LOC677010
1418274_at 1.04703 nuclear transport factor 2 /// similar to Nuclear transport factor 2 (NTF-2) /// similar to Nuclear transport factor 2 (NTF-2) /// similar to Nuclear transport factor 2 (NTF-2) /// similar

to Nuclear transport factor 2 (NTF-2) Nutf2 /// LOC621832 /// LOC668830 /// LOC670887 ///
LOC675580

1418275_a_at 0.92759 E74-like factor 2 Elf2
1418276_at 0.88729 E74-like factor 2 Elf2
1418277_at 0.96666 retinitis pigmentosa 9 homolog (human) Rp9h
1418278_at 0.90745 apolipoprotein C-III Apoc3
1418279_a_at 0.99456 A kinase (PRKA) anchor protein 1 Akap1
1418280_at 1.03551 Kruppel-like factor 6 Klf6
1418281_at 1.07177 RAD51 homolog (S. cerevisiae) Rad51
1418282_x_at 0.84477 "serine (or cysteine) preptidase inhibitor, clade A, member 1b" Serpina1b
1418283_at 1.04069 claudin 4 Cldn4
1418284_at 0.92815 vacuolar protein sorting 72 (yeast) Vps72
1418285_at 0.79557 ephrin B1 Efnb1
1418286_a_at 0.82225 ephrin B1 Efnb1
1418287_a_at 0.99057 deleted in malignant brain tumors 1 Dmbt1
1418288_at 1.24649 lipin 1 Lpin1
1418289_at 0.99323 nestin Nes
1418290_a_at 0.98952 enhancer of zeste homolog 1 (Drosophila) Ezh1
1418291_at 0.85480 zinc finger protein 87 Zfp87
1418292_at 1.03293 "arsA (bacterial) arsenite transporter, ATP-binding, homolog 1" Asna1
1418293_at 0.93522 interferon-induced protein with tetratricopeptide repeats 2 Ifit2
1418294_at 0.53490 erythrocyte protein band 4.1-like 4b Epb4.114b
1418295_s_at 0.93854 diacylglycerol O-acyltransferase 1 Dgat1
1418296_at 0.85146 FXYD domain-containing ion transport regulator 5 Fxyd5
1418297_at 0.78048 dihydropyrimidinase-like 4 Dpysl4
1418298_s_at 1.05494 dihydropyrimidinase-like 4 Dpysl4
1418299_at 0.94905 dihydropyrimidinase-like 4 Dpysl4
1418300_a_at 0.90000 MAP kinase-interacting serine/threonine kinase 2 Mknk2
1418301_at 0.92887 interferon regulatory factor 6 Irf6
1418302_at 0.93447 palmitoyl-protein thioesterase 2 Ppt2
1418303_at 1.03831 RIKEN cDNA 1700113O17 gene 1700113O17Rik
1418304_at 1.06557 protocadherin 21 Pcdh21
1418305_s_at 1.13839 "nucleolar protein family A, member 1 (H/ACA small nucleolar RNPs)"
Nola1
1418306_at 0.88951 "crystallin, beta B1" Crybb1
1418307_a_at 0.96117 spermine binding protein Sbp
1418308_at 1.06118 Hus1 homolog (S. pombe) Hus1
1418309_at 0.99392 "tumor necrosis factor receptor superfamily, member 11b (osteoprotegerin)"
Tnfrsf11b
1418310_a_at 0.96949 retinaldehyde binding protein 1 Rlbp1
1418311_at 0.76561 fructosamine 3 kinase Fn3k
1418312_at 0.91769 zinc finger protein (C2H2 type) 276 Zfp276
1418313_at 0.97895 zinc finger protein (C2H2 type) 276 Zfp276

1418314_a_at 1.00514 ataxin 2 binding protein 1 A2bp1
1418315_at 0.96435 "nuclear receptor subfamily 5, group A, member 1" Nr5a1
1418316_a_at 0.93947 MAP/microtubule affinity-regulating kinase 3 Mark3
1418317_at 0.90837 LIM homeobox protein 2 Lhx2
1418318_at 0.89038 ring finger protein 128 Rnf128
1418319_at 1.12075 RIKEN cDNA 1810047C23 gene 1810047C23Rik
1418320_at 1.12857 "protease, serine, 8 (prostasin)" Prss8
1418321_at 1.02157 "dodecenoyl-Coenzyme A delta isomerase (3,2 trans-enoyl-Coenzyme A isomerase)" Dci
1418322_at 0.84393 cAMP responsive element modulator Crem
1418323_at 0.89792 feminization 1 homolog b (C. elegans) Fem1b
1418324_at 1.02288 feminization 1 homolog b (C. elegans) Fem1b
1418325_at 0.99033 selenophosphate synthetase 2 Sephs2
1418326_at 1.11613 "solute carrier family 7 (cationic amino acid transporter, y+ system), member 5" Slc7a5
1418327_at 0.96580 RIKEN cDNA 1110058L19 gene 1110058L19Rik
1418328_at 0.98021 "carnitine palmitoyltransferase 1b, muscle" Cpt1b
1418329_at 0.99956 pyroglutamyl-peptidase I Pgpep1
1418330_at 0.99973 CCCTC-binding factor Ctf
1418331_at 0.96078 RIKEN cDNA 1110031I02 gene 1110031I02Rik
1418332_a_at 0.90009 ATP/GTP binding protein 1 /// similar to ATP/GTP binding protein 1 Agtpbp1 /// LOC671356
1418333_at 1.13955 metal response element binding transcription factor 1 Mtf1
1418334_at 1.04712 DBF4 homolog (S. cerevisiae) Dbf4
1418335_a_at 0.99928 DPH1 homolog (S. cerevisiae) /// candidate tumor suppressor OVCA2 Dph1 /// RP23-143A14.5
1418336_at 1.08333 AFG3(ATPase family gene 3)-like 1 (yeast) Afg311
1418337_at 1.24679 ribose 5-phosphate isomerase A Rpia
1418338_at 0.93266 WD repeat domain 33 Wdr33
1418339_at 0.97703 RecQ protein-like Recql
1418340_at 0.95394 "Fc receptor, IgE, high affinity I, gamma polypeptide" Fcer1g
1418341_at 0.91784 "RAB4A, member RAS oncogene family" Rab4a
1418342_at 1.01042 replication factor C 1 Recc1
1418343_at 0.89972 GTF2I repeat domain containing 2 Gtf2ird2
1418344_at 1.05758 transmembrane protein 8 (five membrane-spanning domains) Tmem8
1418345_at 1.28481 "tumor necrosis factor (ligand) superfamily, member 13 /// tumor necrosis factor (ligand) superfamily, member 12-member 13" Tnfsf13 /// Tnfsf12-tnfsf13
1418346_at 0.75898 insulin-like 6 Insl6
1418347_at 1.08817 coiled-coil domain containing 22 Ccdc22
1418348_a_at 1.15720 activating signal cointegrator 1 complex subunit 2 Ascc2
1418349_at 0.98060 heparin-binding EGF-like growth factor Hbegf
1418350_at 0.95156 heparin-binding EGF-like growth factor Hbegf
1418351_a_at 1.00331 DNA methyltransferase 3B Dnmt3b

1418352_at 0.98300 hydroxysteroid (17-beta) dehydrogenase 2 Hsd17b2
1418353_at 0.93290 CD5 antigen Cd5
1418354_at 0.90564 component of oligomeric golgi complex 1 Cog1
1418355_at 1.08799 nucleobindin 2 Nucb2
1418356_at 0.93919 mercaptopyruvate sulfurtransferase Mpst
1418357_at 0.78096 forkhead box G1 Foxg1
1418358_at 0.95681 sperm mitochondria-associated cysteine-rich protein Smcp
1418359_at 1.23892 Williams Beuren syndrome chromosome region 27 (human) Wbscr27
1418360_at 1.09876 zinc finger protein 179 Zfp179
1418361_at 0.92995 growth arrest specific 8 Gas8
1418362_at 0.91485 zinc finger protein 42 Zfp42
1418363_at 0.97290 "lactalbumin, alpha" Lalba
1418364_a_at 1.15670 ferritin light chain 1 Ft11
1418365_at 0.87319 cathepsin H Ctsh
1418366_at 0.98419 "histone 2, H3c1 /// histone 2, H2aa1 /// histone 1, H2ad /// histone 1, H2an ///
histone 2, H2ab /// histone 2, H2ac /// histone 2, H2aa2" Hist2h3c1 /// Hist2h2aa1 /// Hist1h2ad ///
Hist1h2an /// Hist2h2ab /// Hist2h2ac /// Hist2h2aa2
1418367_x_at 0.95200 "histone 2, H3c1 /// histone 2, H2aa1 /// histone 1, H2ad /// histone 1, H2an ///
histone 2, H2ab /// histone 2, H2ac /// histone 2, H2aa2" Hist2h3c1 /// Hist2h2aa1 /// Hist1h2ad ///
Hist1h2an /// Hist2h2ab /// Hist2h2ac /// Hist2h2aa2
1418368_at 0.98288 resistin like beta Retnlb
1418369_at 0.94726 "DNA primase, p49 subunit" Prim1
1418370_at 0.95711 "troponin C, cardiac/slow skeletal" Tnnc1
1418371_at 1.00879 dynein light chain LC8-type 2 Dynll2
1418372_at 1.06314 adenylosuccinate lyase Adsl
1418373_at 0.94470 phosphoglycerate mutase 2 Pgam2
1418374_at 0.66440 FXYD domain-containing ion transport regulator 3 Fxyd3
1418375_at 1.04079 methyl-CpG binding domain protein 6 Mbd6
1418376_at 1.05547 fibroblast growth factor 15 Fgf15
1418377_a_at 1.03602 Cd27 binding protein (Hindu God of destruction) Siva
1418378_at 0.83331 bactericidal/permeability-increasing protein-like 1 Bpil1
1418379_s_at 0.84513 G protein-coupled receptor 124 Gpr124
1418380_at 1.05788 telomeric repeat binding factor 1 Terf1
1418381_at 0.93888 zinc finger protein 148 Zfp148
1418382_at 0.53481 adenomatosis polyposis coli down-regulated 1 Apcdd1
1418383_at 0.89221 adenomatosis polyposis coli down-regulated 1 Apcdd1
1418384_at 0.99618 RIKEN cDNA 9430083G14 gene 9430083G14Rik
1418385_at 0.91350 CUE domain containing 2 Cuedc2
1418386_at 1.13854 RIKEN cDNA 2510005D08 gene 2510005D08Rik
1418387_at 1.18616 RIKEN cDNA 4930548G07 gene 4930548G07Rik
1418388_s_at 1.10589 RIKEN cDNA 4930548G07 gene 4930548G07Rik
1418389_at 1.06523 RIKEN cDNA 2810453I06 gene 2810453I06Rik
1418390_at 0.93418 PHD finger protein 21A Phf21a

1418391_at 0.94795 PHD finger protein 21A Phf21a
1418392_a_at 1.68492 guanylate nucleotide binding protein 4 Gbp4
1418393_a_at 0.79204 integrin alpha 7 Itga7
1418394_a_at 0.94930 CD97 antigen Cd97
1418395_at 0.89657 RIKEN cDNA 1300013J15 gene 1300013J15Rik
1418396_at 1.09835 "G-protein signalling modulator 3 (AGS3-like, C. elegans)" Gpsm3
1418397_at 0.96630 zinc finger protein 275 Zfp275
1418398_a_at 0.89783 tetraspanin 32 Tspan32
1418399_at 1.07994 potassium channel tetramerisation domain containing 9 Kctd9
1418400_at 0.95491 "La ribonucleoprotein domain family, member 6" Larp6
1418401_a_at 1.02534 dual specificity phosphatase 16 Dusp16
1418402_at 1.03990 a disintegrin and metallopeptidase domain 19 (meltrin beta) Adam19
1418403_at 0.94578 a disintegrin and metallopeptidase domain 19 (meltrin beta) Adam19
1418404_at 0.96623 RAD9 homolog (S. pombe) Rad9
1418405_at 0.91674 hepatocyte growth factor activator Hgfac
1418406_at 0.85768 phosphodiesterase 8A Pde8a
1418407_at 0.93727 phosphodiesterase 8A Pde8a
1418408_at 0.80720 "zinc finger, AN1-type domain 1" Zfand1
1418409_at 0.84062 jerky Jrk
1418410_at 0.95875 jerky Jrk
1418411_at 0.92402 F-box and leucine-rich repeat protein 8 Fbxl8
1418412_at 0.95414 tumor protein D52-like 1 Tpd52l1
1418413_at 1.02578 caveolin 3 Cav3
1418414_at 0.90202 "potassium voltage-gated channel, subfamily H (eag-related), member 1" Kcnh1
1418415_at 1.09204 homeo box B5 Hoxb5
1418416_x_at 1.02575 pregnancy-specific glycoprotein 23 Psg23
1418417_at 1.01342 musculin Msc
1418418_a_at 1.05708 prolactin-like protein C 1 Prlpc1
1418419_at 0.98152 F-box and leucine-rich repeat protein 20 Fbxl20
1418420_at 0.89915 myogenic differentiation 1 Myod1
1418421_at 1.12667 "B-cell CLL/lymphoma 6, member B" Bcl6b
1418422_at 0.97944 "serine (or cysteine) peptidase inhibitor, clade B, member 9g" Serpinb9g
1418423_s_at 0.98699 "serine (or cysteine) peptidase inhibitor, clade B, member 9f /// serine (or cysteine) peptidase inhibitor, clade B, member 9e /// serine (or cysteine) peptidase inhibitor, clade B, member 9g /// similar to serine (or cysteine) proteinase inhibitor, clade B, member 9f" Serpinb9f /// Serpinb9e /// Serpinb9g /// LOC544923
1418424_at 1.14669 tumor necrosis factor alpha induced protein 6 Tnfaip6
1418425_at 0.85288 trans-acting transcription factor 7 Sp7
1418426_at 0.92272 "general transcription factor II A, 1-like factor" Gtf2a1lf
1418427_at 1.06323 kinesin family member 5B Kif5b
1418428_at 0.99238 kinesin family member 5B Kif5b
1418429_at 1.02757 kinesin family member 5B Kif5b
1418430_at 1.04446 kinesin family member 5B Kif5b

1418431_at	1.04119	kinesin family member 5B	Kif5b
1418432_at	0.96502	calcium binding protein 39	Cab39
1418433_at	0.99707	calcium binding protein 39	Cab39
1418434_at	0.93451	"makorin, ring finger protein, 1"	Mkrn1
1418435_at	0.97373	"makorin, ring finger protein, 1"	Mkrn1
1418436_at	1.03524	syntaxin 7	Stx7
1418437_a_at	0.94276	MAX-like protein X	Mlx
1418438_at	0.87473	"fatty acid binding protein 2, intestinal"	Fabp2
1418439_at	1.00158	"DNA segment, Chr 10, ERATO Doi 322, expressed"	D10Ert322e
1418440_at	1.06235	"procollagen, type VIII, alpha 1"	Col8a1
1418441_at	1.07135	"procollagen, type VIII, alpha 1"	Col8a1
1418442_at	1.03276	"exportin 1, CRM1 homolog (yeast)"	Xpo1
1418443_at	1.11813	"exportin 1, CRM1 homolog (yeast)"	Xpo1
1418444_a_at	1.12695	membrane interacting protein of RGS16	Mir16
1418445_at	1.12115	---	---
1418446_at	0.97412	"solute carrier family 16 (monocarboxylic acid transporters), member 2"	Slc16a2
1418447_at	0.93014	"golgi autoantigen, golgin subfamily a, 5"	Golga5
1418448_at	1.16793	"Harvey rat sarcoma oncogene, subgroup R"	Rras
1418449_at	0.61308	ladinin	Lad1
1418450_at	0.54241	immunoglobulin superfamily containing leucine-rich repeat	Islr
1418451_at	0.86047	"guanine nucleotide binding protein (G protein), gamma 2 subunit"	Gng2
1418452_at	1.08397	"guanine nucleotide binding protein (G protein), gamma 2 subunit"	Gng2
1418453_a_at	0.71759	"ATPase, Na ⁺ /K ⁺ transporting, beta 1 polypeptide"	Atp1b1
1418454_at	1.10746	microfibrillar associated protein 5	Mfap5
1418455_at	0.72607	"coatamer protein complex, subunit zeta 2"	Copz2
1418456_a_at	1.02705	chemokine (C-X-C motif) ligand 14	Cxcl14
1418457_at	1.22893	chemokine (C-X-C motif) ligand 14	Cxcl14
1418458_at	1.02351	anaphase promoting complex subunit 7	Anapc7
1418459_at	1.16688	coiled-coil domain containing 91	Ccdc91
1418460_at	0.98116	SH3 domain protein D19	Sh3d19
1418461_at	1.02626	SH3 domain protein D19	Sh3d19
1418462_at	1.04527	exosome component 9	Exosc9
1418463_at	0.92669	"phosphatidylinositol 3-kinase, regulatory subunit, polypeptide 2 (p85 beta)"	Pik3r2
1418464_at	0.82588	matrilin 4	Matn4
1418465_at	1.20924	neutrophil cytosolic factor 4	Ncf4
1418466_at	1.07680	ring finger protein 141	Rnf141
1418467_at	0.74982	"SWI/SNF related, matrix associated, actin dependent regulator of chromatin, subfamily d, member 3"	Smardc3
1418468_at	0.93156	annexin A11	Anxa11
1418469_at	0.93519	nuclear receptor interacting protein 1	Nrip1
1418470_at	0.86578	Yamaguchi sarcoma viral (v-yes) oncogene homolog 1	Yes1
1418471_at	1.05193	placental growth factor	Pgf

1418472_at	0.93422	aspartoacylase (aminoacylase) 2	Aspa
1418473_at	0.93152	cutC copper transporter homolog (E.coli)	Cutc
1418474_at	1.07525	RIKEN cDNA 1500005A01 gene	1500005A01Rik
1418475_at	0.99160	"sodium channel, nonvoltage-gated 1 beta"	Scnn1b
1418476_at	0.63286	cytokine receptor-like factor 1	Crlf1
1418477_at	1.19457	"matrilin 1, cartilage matrix protein 1"	Matn1
1418478_at	0.99242	LIM domain only 1	Lmo1
1418479_at	1.04409	vacuolar protein sorting 54 (yeast)	Vps54
1418480_at	0.98785	chemokine (C-X-C motif) ligand 7	Cxcl7
1418481_at	1.02892	"protein kinase, membrane associated tyrosine/threonine 1"	Pkmyt1
1418482_at	0.95306	cytochrome b-561 domain containing 2	Cyb561d2
1418483_a_at	1.13568	"glycoprotein galactosyltransferase alpha 1, 3"	Ggta1
1418484_at	0.96601	tektin 2	Tekt2
1418485_at	0.80040	"solute carrier family 4 (anion exchanger), member 3"	Slc4a3
1418486_at	1.29981	vanin 1	Vnn1
1418487_at	0.87629	receptor-interacting serine-threonine kinase 4	Ripk4
1418488_s_at	0.89726	receptor-interacting serine-threonine kinase 4	Ripk4
1418489_a_at	0.94159	calcitonin receptor-like	Calcr1
1418490_at	0.90477	serine dehydratase-like	Sdsl
1418491_a_at	1.04680	pseudouridine synthase 3	Pus3
1418492_at	0.93587	"gremlin 2 homolog, cysteine knot superfamily (Xenopus laevis)"	Grem2
1418493_a_at	0.87696	"synuclein, alpha"	Snca
1418494_at	0.96649	early B-cell factor 2	Ebf2
1418495_at	1.17527	zinc finger CCCH type containing 8	Zc3h8
1418496_at	0.63891	forkhead box A1	Foxa1
1418497_at	0.87350	fibroblast growth factor 13	Fgf13
1418498_at	0.98653	fibroblast growth factor 13	Fgf13
1418499_a_at	0.96914	"potassium voltage-gated channel, Isk-related subfamily, gene 3"	Kcne3
1418500_at	0.88596	nucleosome assembly protein 1-like 3	Nap113
1418501_a_at	0.90977	oxidation resistance 1	Oxr1
1418502_a_at	0.95502	oxidation resistance 1	Oxr1
1418503_at	1.09365	heat shock protein 9A	Hspa9a
1418504_at	1.11541	heat shock protein 9A	Hspa9a
1418505_at	0.86267	nudix (nucleoside diphosphate linked moiety X)-type motif 4	Nudt4
1418506_a_at	0.91993	peroxiredoxin 2	Prdx2
1418507_s_at	0.38764	suppressor of cytokine signaling 2	Socs2
1418508_a_at	0.95953	growth factor receptor bound protein 2	Grb2
1418509_at	0.43678	carbonyl reductase 2	Cbr2
1418510_s_at	0.86629	F-box only protein 8	Fbxo8
1418511_at	1.15444	dermatopontin	Dpt
1418512_at	0.95252	"serine/threonine kinase 3 (Ste20, yeast homolog)"	Stk3
1418513_at	0.96346	"serine/threonine kinase 3 (Ste20, yeast homolog)"	Stk3
1418514_at	1.01234	metal response element binding transcription factor 2	Mtf2

1418515_at	1.01251	metal response element binding transcription factor 2	Mtf2
1418516_at	1.00766	metal response element binding transcription factor 2	Mtf2
1418517_at	0.69596	Iroquois related homeobox 3 (Drosophila)	Irx3
1418518_at	0.86213	furin (paired basic amino acid cleaving enzyme)	Furin
1418519_at	1.04276	aminoadipate aminotransferase	Aadat
1418520_at	0.94368	trans-golgi network protein	Tgoln1
1418521_a_at	1.02503	metaxin 1	Mtx1
1418522_at	1.00966	metaxin 1	Mtx1
1418523_at	1.20720	ariadne homolog 2 (Drosophila)	Arih2
1418524_at	0.96109	pericentriolar material 1	Pcm1
1418525_at	0.90067	pericentriolar material 1	Pcm1
1418526_at	1.02400	FUS interacting protein (serine-arginine rich) 1	Fusip1
1418527_a_at	1.03783	FUS interacting protein (serine-arginine rich) 1	Fusip1
1418528_a_at	1.01182	defender against cell death 1	Dad1
1418529_at	1.09979	O-sialoglycoprotein endopeptidase	Osgep
1418530_at	1.03479	nucleoporin 160	Nup160
1418531_at	1.04532	oocyte secreted protein 1	Oosp1
1418532_at	0.89441	frizzled homolog 2 (Drosophila)	Fzd2
1418533_s_at	0.90332	frizzled homolog 2 (Drosophila)	Fzd2
1418534_at	0.81222	---	---
1418535_at	0.80119	"ral guanine nucleotide dissociation stimulator,-like 1"	Rgl1
1418536_at	1.36802	"similar to H-2 class I histocompatibility antigen, Q7 alpha chain precursor (QA-2 antigen) /// region containing histocompatibility 2, Q region locus 9; histocompatibility 2, Q region locus 7"	LOC630509 /// LOC674192
1418537_at	1.08209	RIKEN cDNA 0610042E07 gene	0610042E07Rik
1418538_at	0.79600	KDEL (Lys-Asp-Glu-Leu) endoplasmic reticulum protein retention receptor 3	Kdelr3
1418539_a_at	1.83979	"protein tyrosine phosphatase, receptor type, E"	Ptpre
1418540_a_at	1.40858	"protein tyrosine phosphatase, receptor type, E"	Ptpre
1418541_at	1.08075	centromere protein O	Cenpo
1418542_s_at	1.02984	centromere protein O	Cenpo
1418543_s_at	0.99372	coiled-coil domain containing 43	Ccdc43
1418544_at	0.93292	"Kv channel interacting protein 3, calsenilin"	Kcnip3
1418545_at	1.00479	WASP family 1	Wasf1
1418546_a_at	0.97144	Stam binding protein like 1	Stamp1l
1418547_at	0.99039	tissue factor pathway inhibitor 2	Tfpi2
1418548_at	0.92550	seminal vesicle secretion 6	Svs6
1418549_at	0.96127	"glycoprotein hormones, alpha subunit"	Cga
1418550_x_at	0.99435	defensin related sequence cryptdin peptide (paneth cells)	Defcr-rs1
1418551_at	0.63284	"myosin binding protein C, cardiac"	Mybpc3
1418552_at	0.79740	"opsin 1 (cone pigments), short-wave-sensitive (color blindness, tritan)"	Opn1sw
1418553_at	0.95475	rho/rac guanine nucleotide exchange factor (GEF) 18	Arhgef18

1418554_at	1.07644	adrenomedullin receptor	Admr	
1418555_x_at	0.90244	Spi-C transcription factor (Spi-1/PU.1 related)	Spic	
1418556_at	0.93481	"capping protein (actin filament) muscle Z-line, alpha 3"	Capza3	Capza3
1418557_s_at	0.89175	testis-specific serine kinase 3	Tssk3	
1418558_at	0.99530	retina and anterior neural fold homeobox	Rax	Rax
1418559_at	1.03243	testicular serine protease 1	Tesp1	
1418560_at	1.03830	pyruvate dehydrogenase E1 alpha 1	Pdha1	Pdha1
1418561_at	0.92662	"splicing factor 3b, subunit 1"	Sf3b1	
1418562_at	1.11664	"splicing factor 3b, subunit 1"	Sf3b1	
1418563_at	0.97998	Serpine1 mRNA binding protein 1	Serbp1	
1418564_s_at	1.04356	Serpine1 mRNA binding protein 1	Serbp1	
1418565_at	0.80384	Serpine1 mRNA binding protein 1	Serbp1	
1418566_s_at	1.02658	NudC domain containing 2	Nudcd2	
1418567_a_at	1.02854	signal recognition particle 14	Srp14	
1418568_x_at	1.03045	signal recognition particle 14	Srp14	
1418569_at	0.94052	filamin binding LIM protein 1	Fblim1	
1418570_at	0.96972	nicastrin	Ncstn	
1418571_at	1.07855	"tumor necrosis factor receptor superfamily, member 12a"	Tnfrsf12a	Tnfrsf12a
1418572_x_at	1.05725	"tumor necrosis factor receptor superfamily, member 12a"	Tnfrsf12a	Tnfrsf12a
1418573_a_at	0.99860	hnRNP-associated with lethal yellow	Raly	
1418574_a_at	0.93562	split hand/foot malformation (ectrodactyly) type 1	Shfm1	Shfm1
1418575_at	0.96039	split hand/foot malformation (ectrodactyly) type 1	Shfm1	Shfm1
1418576_at	0.85637	"Yip1 domain family, member 5"	Yipf5	
1418577_at	1.05068	tripartite motif protein 8	Trim8	
1418578_at	0.88457	"diacylglycerol kinase, alpha"	Dgka	
1418579_at	0.92731	centrin 2	Cetn2	
1418580_at	1.34205	receptor transporter protein 4	Rtp4	
1418581_a_at	1.02110	LIM motif-containing protein kinase 2	Limk2	
1418582_at	0.75397	"core-binding factor, runt domain, alpha subunit 2, translocated to, 3 homolog (human)"	Cbfa2t3h	
1418583_at	1.02643	histidine triad nucleotide binding protein 3	Hint3	
1418584_at	1.11291	cyclin H /// similar to cyclin H	Ccnh /// LOC672655	
1418585_at	0.91345	cyclin H	Ccnh	
1418586_at	1.00290	adenylate cyclase 9	Adcy9	
1418587_at	1.21231	Tnf receptor-associated factor 3	Traf3	
1418588_at	0.88895	neurensin 1	Nrsn1	
1418589_a_at	0.88434	myeloid leukemia factor 1	Mlf1	
1418590_at	0.99493	karyopherin (importin) alpha 6	Kpna6	
1418591_at	0.89408	"DnaJ (Hsp40) homolog, subfamily A, member 4"	Dnaja4	
1418592_at	0.99297	"DnaJ (Hsp40) homolog, subfamily A, member 4"	Dnaja4	
1418593_at	0.92918	"TAF6 RNA polymerase II, TATA box binding protein (TBP)-associated factor"	Taf6	
1418594_a_at	0.74046	nuclear receptor coactivator 1	Ncoa1	

dehydrogenase (GAPDH) /// similar to Glyceraldehyde-3-phosphate dehydrogenase (GAPDH) /// similar
 to Glyceraldehyde-3-phosphate dehydrogenase (GAPDH) /// similar to glyceraldehyde-3-phosphate
 dehydrogenase /// glyceraldehyde-3-phosphate dehydrogenase pseudogene /// similar to Glyceraldehyde-
 3-phosphate dehydrogenase (GAPDH) /// similar to Glyceraldehyde-3-phosphate dehydrogenase
 (GAPDH) /// similar to Glyceraldehyde-3-phosphate dehydrogenase (GAPDH) /// similar to
 Glyceraldehyde-3-phosphate dehydrogenase (GAPDH) /// similar to Glyceraldehyde-3-phosphate
 dehydrogenase (GAPDH) /// similar to Glyceraldehyde-3-phosphate dehydrogenase (GAPDH) /// similar
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 dehydrogenase (GAPDH) /// similar to Glyceraldehyde-3-phosphate dehydrogenase (GAPDH) /// similar
 to Glyceraldehyde-3-phosphate dehydrogenase (GAPDH) /// similar to Glyceraldehyde-3-phosphate
 dehydrogenase (GAPDH) /// similar to Glyceraldehyde-3-phosphate dehydrogenase (GAPDH)
 LOC14433 /// LOC219102 /// LOC224508 /// LOC230044 /// LOC242823 /// 3000002C10Rik ///
 LOC380687 /// LOC384643 /// Gapdh /// LOC432919 /// LOC433184 /// LOC433273 /// LOC435292 ///
 LOC545984 /// LOC622365 /// LOC630967 /// LOC632720 /// LOC633247 /// RP23-227C11.3 ///
 LOC666342 /// LOC666488 /// LOC667170 /// LOC668426 /// LOC669993 /// LOC670403 ///
 LOC671133 /// LOC672308 /// LOC675199 /// LOC675385 /// LOC675515 /// LOC676292
 1418626_a_at 0.82082 clusterin Clu
 1418627_at 0.88306 "glutamate-cysteine ligase , modifier subunit" Gclm
 1418628_at 1.01435 "KH domain containing, RNA binding, signal transduction associated 1"
 Khdrbs1
 1418629_a_at 0.94308 --- ---
 1418630_at 1.18952 "KH domain containing, RNA binding, signal transduction associated 1"
 Khdrbs1
 1418631_at 0.92321 ubiquitin-conjugating enzyme E2H Ube2h
 1418632_at 0.83984 ubiquitin-conjugating enzyme E2H Ube2h
 1418633_at 0.89857 Notch gene homolog 1 (Drosophila) Notch1
 1418634_at 0.89011 Notch gene homolog 1 (Drosophila) Notch1
 1418635_at 1.09532 ets variant gene 3 Etv3
 1418636_at 1.06219 ets variant gene 3 Etv3
 1418637_at 1.06572 ets variant gene 3 Etv3
 1418638_at 1.06367 "histocompatibility 2, class II, locus Mb1" H2-DMb1
 1418639_at 0.87351 surfactant associated protein C Sftpc
 1418640_at 0.94735 "sirtuin 1 ((silent mating type information regulation 2, homolog) 1 (S.
 cerevisiae)" Sirt1
 1418641_at 0.97639 lymphocyte cytosolic protein 2 Lcp2
 1418642_at 0.94096 lymphocyte cytosolic protein 2 Lcp2
 1418643_at 0.88433 tetraspanin 13 Tspan13
 1418644_a_at 0.97509 serine/threonine kinase 11 Stk11
 1418645_at 0.92893 histidine ammonia lyase /// similar to Histidine ammonia-lyase (Histidase)
 Hal /// LOC638196
 1418646_at 1.03798 "guanine nucleotide binding protein, related sequence 1" Gna-rs1
 1418647_at 0.94874 --- ---
 1418648_at 1.23358 EGL nine homolog 3 (C. elegans) EglN3

1418649_at 1.33106 EGL nine homolog 3 (C. elegans) Egl3
1418650_at 0.73874 spermatogenesis associated 6 Spata6
1418651_at 0.76492 spermatogenesis associated 6 Spata6
1418652_at 0.99055 chemokine (C-X-C motif) ligand 9 Cxcl9
1418653_at 0.91068 "cytochrome P450, family 2. subfamily c, polypeptide 37 /// cytochrome P450, family 2, subfamily c, polypeptide 50 /// cytochrome P450 /// similar to cytochrome P450" Cyp2c37 /// Cyp2c50 /// Cyp2c54 /// LOC639023
1418654_at 1.05666 hydroxyacid oxidase (glycolate oxidase) 3 Hao3
1418655_at 0.75000 "beta-1,4-N-acetyl-galactosaminyl transferase 1" B4galnt1
1418656_at 1.09236 "LSM5 homolog, U6 small nuclear RNA associated (S. cerevisiae)" Lsm5
1418657_at 0.98152 "zinc finger, HIT type 4" Znhit4
1418658_at 0.91912 RIKEN cDNA 2410005O16 gene 2410005O16Rik
1418659_at 1.06955 circadian locomoter output cycles kaput Clock
1418660_at 1.01228 circadian locomoter output cycles kaput /// hypothetical protein LOC620729 Clock /// LOC620729
1418661_at 0.98451 abhydrolase domain containing 2 Abhd2
1418662_at 0.92170 RIKEN cDNA 2210012G02 gene 2210012G02Rik
1418663_at 0.97065 multiple PDZ domain protein Mpdz
1418664_at 0.90968 multiple PDZ domain protein Mpdz
1418665_at 1.00462 inositol (myo)-1(or 4)-monophosphatase 2 Impa2
1418666_at 1.00705 pentraxin related gene Ptx3
1418667_at 1.17427 RIKEN cDNA 2410002O22 gene 2410002O22Rik
1418668_at 1.06842 acyl-CoA synthetase medium-chain family member 1 Acsm1
1418669_at 0.97561 perlecan (heparan sulfate proteoglycan 2) Hspg2
1418670_s_at 0.93413 perlecan (heparan sulfate proteoglycan 2) Hspg2
1418671_at 0.93885 calpain 5 Capn5
1418672_at 0.90039 "aldo-keto reductase family 1, member C13" Akr1c13
1418673_at 1.00661 snail homolog 2 (Drosophila) Snai2
1418674_at 1.07435 oncostatin M receptor Osmr
1418675_at 0.96663 oncostatin M receptor Osmr
1418676_at 1.02224 insulin related protein 2 (islet 2) Isl2
1418677_at 0.88509 actinin alpha 3 Actn3
1418678_at 0.90708 hyaluronan synthase 2 Has2
1418679_at 1.11436 granzyme F Gzmf
1418680_at 0.89702 "serine (or cysteine) peptidase inhibitor, clade D, member 1" Serpind1
1418681_at 1.12409 glycosyltransferase 28 domain containing 1 Glt28d1
1418682_at 1.05989 testis nuclear RNA binding protein Tenr
1418683_at 1.05658 lin-7 homolog B (C. elegans) Lin7b
1418684_at 1.02022 RIKEN cDNA 2310012P17 gene 2310012P17Rik
1418685_at 1.16093 toll-interleukin 1 receptor (TIR) domain-containing adaptor protein Tirap
1418686_at 0.95592 2'-5' oligoadenylate synthetase 1C Oas1c
1418687_at 0.69073 activity regulated cytoskeletal-associated protein Arc
1418688_at 1.04404 calcitonin receptor Calcr

1418689_at 0.96565 cathepsin 6 Cts6
 1418690_at 1.02203 "protein tyrosine phosphatase, receptor type Z, polypeptide 1" Ptporz1
 1418691_at 0.98299 regulator of G-protein signaling 9 Rgs9
 1418692_at 1.02943 "RAB8A, member RAS oncogene family" Rab8a
 1418693_at 0.97263 heterogeneous nuclear ribonucleoprotein C /// similar to heterogeneous nuclear ribonucleoprotein C /// similar to heterogeneous nuclear ribonucleoprotein C Hnrpc /// LOC620883 /// LOC676606
 1418694_at 0.98885 potassium channel modulatory factor 1 Kcmf1
 1418695_a_at 1.05143 potassium channel modulatory factor 1 Kcmf1
 1418696_at 0.91996 cDNA sequence BC021608 BC021608
 1418697_at 0.44263 indolethylamine N-methyltransferase Inmt
 1418698_a_at 0.99631 ferrocyclase Fech
 1418699_s_at 0.96986 ferrocyclase Fech
 1418700_at 1.00407 lipoic acid synthetase Lias
 1418701_at 1.00672 catechol-O-methyltransferase Comt
 1418702_a_at 0.99202 RIKEN cDNA 2810428I15 gene 2810428I15Rik
 1418703_at 1.01480 "RNA binding motif, single stranded interacting protein 1" Rbms1
 1418704_at 0.93663 S100 calcium binding protein A13 S100a13
 1418705_at 1.01780 cone-rod homeobox containing gene Crx
 1418706_at 0.94497 "solute carrier family 38, member 3" Slc38a3
 1418707_at 0.89770 BCL2-associated athanogene 4 Bag4
 1418708_at 1.04840 apolipoprotein C-IV Apoc4
 1418709_at 0.98088 "cytochrome c oxidase, subunit VIIa 1" Cox7a1
 1418710_at 0.83730 CD59a antigen Cd59a
 1418711_at 0.91409 "platelet derived growth factor, alpha" Pdgfa
 1418712_at 1.02926 CDC42 effector protein (Rho GTPase binding) 5 Cdc42ep5
 1418713_at 1.00097 pterin 4 alpha carbinolamine dehydratase/dimerization cofactor of hepatocyte nuclear factor 1 alpha (TCF1) 1 Pcbd1
 1418714_at 0.92597 dual specificity phosphatase 8 Dusp8
 1418715_at 1.02322 pantothenate kinase 1 Pank1
 1418716_at 0.96071 mitochondrial ribosomal protein S25 Mrps25
 1418717_at 0.90773 mitochondrial ribosomal protein S25 Mrps25
 1418718_at 1.32894 chemokine (C-X-C motif) ligand 16 Cxcl16
 1418719_at 0.82885 RIKEN cDNA 2410004L22 gene 2410004L22Rik
 1418720_at 0.91030 "COP9 (constitutive photomorphogenic) homolog, subunit 7b (Arabidopsis thaliana)" Cops7b
 1418721_at 0.89595 "COP9 (constitutive photomorphogenic) homolog, subunit 7b (Arabidopsis thaliana)" Cops7b
 1418722_at 0.85309 neutrophilic granule protein Ngp
 1418723_at 1.11056 "endothelial differentiation, lysophosphatidic acid G-protein-coupled receptor 7" Edg7
 1418724_at 0.93374 complement component factor i Cfi
 1418725_at 0.97499 "DnaJ (Hsp40) homolog, subfamily C, member 5 beta" Dnajc5b

1418726_a_at 0.74020 "troponin T2, cardiac" Tnnt2
1418727_at 1.16491 nucleoporin 155 Nup155
1418728_at 1.11367 steroidogenic acute regulatory protein Star
1418729_at 0.95150 steroidogenic acute regulatory protein Star
1418730_at 1.00246 ring finger protein 12 Rnf12
1418731_at 1.02188 ring finger protein 12 Rnf12
1418732_s_at 1.09328 RIKEN cDNA 1500041N16 gene 1500041N16Rik
1418733_at 0.98014 twist gene homolog 1 (Drosophila) Twist1
1418734_at 0.99864 expressed sequence BE136769 BE136769
1418735_at 0.97775 keratin 4 Krt4
1418736_at 0.96763 "UDP-GalNAc:betaGlcNAc beta 1,3-galactosaminyltransferase, polypeptide 1" B3galnt1
1418737_at 1.10975 nudix (nucleoside diphosphate linked moiety X)-type motif 2 Nudt2
1418738_at 1.00983 "sodium channel, voltage-gated, type I, beta" Scn1b
1418739_at 1.05180 serum/glucocorticoid regulated kinase 2 Sgk2
1418740_at 1.11956 spermatid associated Spert
1418741_at 1.05822 integrin beta 7 Itgb7
1418742_at 0.90577 keratin 34 Krt34
1418743_a_at 0.80531 tescalcin Tesc
1418744_s_at 0.72234 tescalcin Tesc
1418745_at 0.67605 osteomodulin Omd
1418746_at 0.91483 paroxysmal nonkinesigenic dyskinesia Pnkd
1418747_at 0.95714 SFFV proviral integration 1 Sfp1
1418748_at 0.94512 caspase 14 Casp14
1418749_at 0.84195 pleckstrin and Sec7 domain containing 3 Psd3
1418750_at 0.93029 plexin B3 Plxnb3
1418751_at 0.93590 suppression inducing transmembrane adaptor 1 Sit1
1418752_at 1.81220 "aldehyde dehydrogenase family 3, subfamily A1" Aldh3a1
1418753_at 0.89762 glutamine fructose-6-phosphate transaminase 2 Gfpt2
1418754_at 1.01297 adenylate cyclase 8 Adcy8
1418755_at 0.91723 T-box 15 Tbx15
1418756_at 0.89501 thyrotropin releasing hormone Trh
1418757_at 0.85698 ring finger protein 36 Rnf36
1418758_a_at 0.99624 "pleckstrin homology, Sec7 and coiled-coil domains 3" Pscd3
1418759_at 1.06456 "protein tyrosine phosphatase, non-receptor type 20" Ptpn20
1418760_at 1.25807 retinol dehydrogenase 11 Rdh11
1418761_at 1.18038 insulin-like growth factor 2 mRNA binding protein 1 Igf2bp1
1418762_at 1.01597 CD55 antigen Cd55
1418763_at 0.89354 "nitrilase family, member 2" Nit2
1418764_a_at 1.14449 bisphosphate 3'-nucleotidase 1 Bpnt1
1418765_at 1.11611 T-cell immunoglobulin and mucin domain containing 2 Timd2
1418766_s_at 0.96899 T-cell immunoglobulin and mucin domain containing 2 Timd2
1418767_at 0.89547 "cytochrome P450, family 4, subfamily f, polypeptide 13" Cyp4f13

1418768_at 0.87312 optic atrophy 1 homolog (human) Opa1
1418769_at 0.99034 myozenin 2 Myoz2
1418770_at 1.01565 CD2 antigen Cd2
1418771_a_at 0.96610 carboxypeptidase B2 (plasma) Cpb2
1418772_at 1.12255 cDNA sequence BC016423 BC016423
1418773_at 0.87182 fatty acid desaturase 3 Fads3
1418774_a_at 0.99809 "ATPase, Cu++ transporting, alpha polypeptide" Atp7a
1418775_at 0.88489 expressed sequence AI837181 AI837181
1418776_at 1.04550 RIKEN cDNA 5830443L24 gene /// similar to macrophage activation 2 like
5830443L24Rik /// LOC626578
1418777_at 1.05959 chemokine (C-C motif) ligand 25 Ccl25
1418778_at 1.05337 coiled-coil domain containing 109B Ccdc109b
1418779_at 0.95617 "RCE1 homolog, prenyl protein peptidase (S. cerevisiae)" Rce1
1418780_at 0.76192 "cytochrome P450, family 39, subfamily a, polypeptide 1" Cyp39a1
1418781_at 0.89541 NAT9 Nat9
1418782_at 0.94458 retinoid X receptor gamma Rxrg
1418783_at 1.08351 "transient receptor potential cation channel, subfamily M, member 5" Trpm5
1418784_at 0.94362 "glutamate receptor, ionotropic, kainate 5 (gamma 2)" Grik5
1418785_at 0.99819 mitogen-activated protein kinase 8 interacting protein 2 Mapk8ip2
1418786_at 0.95952 mitogen-activated protein kinase 8 interacting protein 2 Mapk8ip2
1418787_at 0.96690 mannose binding lectin (C) Mbl2
1418788_at 0.89338 endothelial-specific receptor tyrosine kinase Tek
1418789_at 1.00860 "syntrophin, gamma 2" Sntg2
1418790_at 1.00489 Fez family zinc finger 2 Fezf2
1418791_at 1.00273 SH3-domain GRB2-like 2 Sh3gl2
1418792_at 0.98277 SH3-domain GRB2-like 2 Sh3gl2
1418793_at 1.04878 "iduronidase, alpha-L-" Idua
1418794_at 1.01967 CDP-diacylglycerol synthase (phosphatidate cytidyltransferase) 2 Cds2
1418795_at 0.86957 CDP-diacylglycerol synthase (phosphatidate cytidyltransferase) 2 Cds2
1418796_at 1.07805 "C-type lectin domain family 11, member a" Clec11a
1418797_at 0.97323 "membrane-spanning 4-domains, subfamily A, member 8A" Ms4a8a
1418798_s_at 0.93906 serine/arginine-rich protein specific kinase 3 Srpk3
1418799_a_at 0.96224 "procollagen, type XVII, alpha 1" Col17a1
1418800_at 0.84691 "basic helix-loop-helix domain containing, class B, 8" Bhlhb8
1418801_at 0.78334 zinc finger with KRAB and SCAN domains 1 Zkscan1
1418802_at 0.92824 expressed sequence R74862 R74862
1418803_a_at 1.13485 "Fas ligand (TNF superfamily, member 6)" FasL
1418804_at 1.09101 succinate receptor 1 Sucnr1
1418805_at 0.97897 secretin Sct
1418806_at 0.95878 colony stimulating factor 3 receptor (granulocyte) Csf3r
1418807_at 1.08394 RIKEN cDNA 3110070M22 gene 3110070M22Rik
1418808_at 1.05261 retinol dehydrogenase 5 Rdh5
1418809_at 0.94760 paired-Ig-like receptor A2 Pira2

1418810_at 0.96941 corticotropin releasing hormone receptor 1 Crhr1
1418811_at 0.97710 BarH-like 1 (Drosophila) Barhl1
1418812_a_at 0.92372 BarH-like 1 (Drosophila) Barhl1
1418813_at 1.00688 "serine (or cysteine) peptidase inhibitor, clade A, member 5" Serpina5
1418814_s_at 0.98800 "NADH dehydrogenase (ubiquinone) 1 alpha subcomplex, 12" Ndufa12
1418815_at 1.00095 cadherin 2 Cdh2
1418816_at 0.91330 chromatin modifying protein 1B Chmp1b
1418817_at 0.87997 chromatin modifying protein 1B Chmp1b
1418818_at 1.13081 aquaporin 5 Aqp5
1418819_at 1.00247 ADP-ribosylation factor-like 8B Arl8b
1418820_s_at 1.05526 "zinc finger, CCHC domain containing 10" Zcchc10
1418821_at 1.09432 "cytochrome P450, family 2, subfamily a, polypeptide 12" Cyp2a12
1418822_a_at 0.91223 ADP-ribosylation factor 6 Arf6
1418823_at 0.89346 ADP-ribosylation factor 6 Arf6
1418824_at 0.67210 ADP-ribosylation factor 6 Arf6
1418825_at 0.87946 "immunity-related GTPase family, M" Irgm
1418826_at 0.96736 "membrane-spanning 4-domains, subfamily A, member 6B" Ms4a6b
1418827_at 0.96316 three prime histone mRNA exonuclease 1 Thex1
1418828_at 0.93226 three prime histone mRNA exonuclease 1 Thex1
1418829_a_at 1.98889 "enolase 2, gamma neuronal" Eno2
1418830_at 0.94191 CD79A antigen (immunoglobulin-associated alpha) Cd79a
1418831_at 0.93862 plakophilin 3 Pkp3
1418832_at 0.91870 plakophilin 3 Pkp3
1418833_at 1.00701 alanine-glyoxylate aminotransferase Agxt
1418834_at 0.92085 cappuccino Cno
1418835_at 1.16165 "pleckstrin homology-like domain, family A, member 1" Phlda1
1418836_at 0.90811 quinolinate phosphoribosyltransferase Qprt
1418837_at 1.01044 quinolinate phosphoribosyltransferase Qprt
1418838_at 0.93398 "ATP-binding cassette, sub-family D (ALD), member 1" Abcd1
1418839_at 1.10367 "glomulin, FKBP associated protein" Glmn
1418840_at 0.82091 programmed cell death 4 /// similar to programmed cell death 4 Pdcd4 ///
LOC670861
1418841_s_at 0.96534 cell division cycle 2-like 1 Cdc2l1
1418842_at 0.86267 hematopoietic cell specific Lyn substrate 1 Hcls1
1418843_at 1.13983 "solute carrier family 30 (zinc transporter), member 4" Slc30a4
1418844_at 1.07198 "asparagine-linked glycosylation 9 homolog (yeast, alpha 1,2
mannosyltransferase)" Alg9
1418845_at 0.83120 protein C Proc
1418846_at 0.98031 "adaptor-related protein complex AP-4, mu 1" Ap4m1
1418847_at 1.07608 arginase type II Arg2
1418848_at 1.03812 aquaporin 7 Aqp7
1418849_x_at 1.00510 aquaporin 7 Aqp7
1418850_at 0.98238 enhancer of polycomb homolog 1 (Drosophila) Epc1

1418851_at 1.13672 tripartite motif protein 39 Trim39
1418852_at 0.91336 "cholinergic receptor, nicotinic, alpha polypeptide 1 (muscle)" Chrna1
1418853_at 0.98792 apolipoprotein N Apon
1418854_at 0.92723 baculoviral IAP repeat-containing 2 Birc2
1418855_at 0.95565 RIKEN cDNA 1110008K04 gene 1110008K04Rik
1418856_a_at 1.02473 "Fanconi anemia, complementation group A" Fanca
1418857_at 0.91555 "solute carrier family 13 (sodium-dependent dicarboxylate transporter), member 2" Slc13a2
1418858_at 0.98011 aldehyde oxidase 3 Aox3
1418859_at 0.98778 regulatory factor X-associated protein Rfxap
1418860_a_at 0.79078 LETM1 domain containing 1 Letmd1
1418861_at 0.97603 protein inhibitor of activated STAT 4 Pias4
1418862_at 1.13049 enoyl Coenzyme A hydratase domain containing 3 Echdc3
1418863_at 1.16166 GATA binding protein 4 Gata4
1418864_at 0.91674 GATA binding protein 4 Gata4
1418865_at 0.87221 zinc finger protein 385 Zfp385
1418866_at 1.02067 "cytochrome P450, family 24, subfamily a, polypeptide 1" Cyp24a1
1418867_at 1.04698 "cytochrome P450, family 24, subfamily a, polypeptide 1" Cyp24a1
1418868_at 1.06090 engrailed 2 En2
1418869_a_at 1.07133 pseudouridine synthase 1 Pus1
1418870_at 0.95011 RIKEN cDNA 4930579J09 gene 4930579J09Rik
1418871_a_at 1.13215 RIKEN cDNA 1810007E14 gene 1810007E14Rik
1418872_at 1.11194 "ATP-binding cassette, sub-family B (MDR/TAP), member 1B" Abcb1b
1418873_at 0.75907 sideroflexin 4 Sfxn4
1418874_a_at 1.07209 "proteasome (prosome, macropain) 26S subunit, non-ATPase, 4" Psmd4
1418875_at 0.87347 synaptogyrin 4 Syngr4
1418876_at 0.91783 forkhead box D1 Foxd1
1418877_at 1.32283 Forkhead box D1 Foxd1
1418878_at 0.86361 acrosomal vesicle protein 1 Acrv1
1418879_at 0.88201 RIKEN cDNA 9030611O19 gene 9030611O19Rik
1418880_at 0.92885 glial cell line derived neurotrophic factor family receptor alpha 3 Gfra3
1418881_at 0.94058 EF hand calcium binding protein 2 Efcbp2
1418882_at 0.97224 "NACHT, leucine rich repeat and PYD containing 5" Nalp5
1418883_a_at 1.05070 "poly A binding protein, cytoplasmic 1" Pabpc1
1418884_x_at 0.95630 "tubulin, alpha 1" Tuba1
1418885_a_at 0.95041 isocitrate dehydrogenase 3 (NAD+) beta Idh3b
1418886_s_at 1.01983 isocitrate dehydrogenase 3 (NAD+) beta Idh3b
1418887_a_at 0.85258 "DNA segment, Chr 11, Wayne State University 99, expressed" D11Wsu99e
1418888_a_at 1.03140 selenoprotein X 1 Sepx1
1418889_a_at 1.00489 "casein kinase 1, delta" Csnk1d
1418890_a_at 0.84535 "RAB3D, member RAS oncogene family" Rab3d
1418891_a_at 0.74424 "RAB3D, member RAS oncogene family" Rab3d
1418892_at 1.00012 "ras homolog gene family, member J" Rhoj

1418893_at 0.95079 pre B-cell leukemia transcription factor 2 Pbx2
1418894_s_at 0.85984 pre B-cell leukemia transcription factor 2 Pbx2
1418895_at 1.03993 src family associated phosphoprotein 2 Scap2
1418896_a_at 0.92452 ribophorin II Rpn2
1418897_at 0.87168 coagulation factor II F2
1418898_at 0.97312 lin-7 homolog C (C. elegans) Lin7c
1418899_at 0.96741 ubiquitin-fold modifier 1 Ufm1
1418900_at 0.98424 ubiquitin-fold modifier 1 Ufm1
1418901_at 0.94248 "CCAAT/enhancer binding protein (C/EBP), beta" Cebp
1418902_at 0.81513 RIKEN cDNA 3110023E09 gene 3110023E09Rik
1418903_at 1.02565 aquaporin 2 Aqp2
1418904_at 0.96842 --- ---
1418905_at 1.19826 nucleotide binding protein 1 Nubp1
1418906_at 1.06438 nucleotide binding protein 1 Nubp1
1418907_at 0.95915 --- ---
1418908_at 0.98772 peptidylglycine alpha-amidating monooxygenase Pam
1418909_at 1.07910 erythroblast membrane-associated protein Ermap
1418910_at 0.55126 bone morphogenetic protein 7 Bmp7
1418911_s_at 1.39198 acyl-CoA synthetase long-chain family member 4 Acsl4
1418912_at 1.12600 plexin domain containing 2 Plxdc2
1418913_at 1.07797 betaine-homocysteine methyltransferase 2 Bhmt2
1418914_s_at 1.04071 betaine-homocysteine methyltransferase 2 Bhmt2
1418915_at 1.07833 "methylmalonic aciduria cblC type, with homocystinuria" Mmachc
1418916_a_at 1.06388 secreted phosphoprotein 2 Spp2
1418917_at 0.81112 heme binding protein 2 Hebp2
1418918_at 0.95028 insulin-like growth factor binding protein 1 Igfbp1
1418919_at 1.02278 shugoshin-like 1 (S. pombe) Sgol1
1418920_at 0.95143 claudin 15 Cldn15
1418921_at 0.94991 "immunoglobulin superfamily, member 4B" Igsf4b
1418922_at 1.08189 "immunoglobulin superfamily, member 4B" Igsf4b
1418923_at 0.93604 "solute carrier family 17 (sodium phosphate), member 3" Slc17a3
1418924_at 1.08197 Ras association (RalGDS/AF-6) domain family 7 Rassf7
1418925_at 0.75929 cadherin EGF LAG seven-pass G-type receptor 1 Celsr1
1418926_at 0.99248 zinc finger homeobox 1a Zfhx1a
1418927_a_at 0.81273 similar to hyaluronic acid binding protein 4 LOC671769
1418928_a_at 0.95272 RIKEN cDNA 2310038H17 gene 2310038H17Rik
1418929_at 0.93017 intraflagellar transport 57 homolog (Chlamydomonas) Ift57
1418930_at 1.52223 chemokine (C-X-C motif) ligand 10 Cxcl10
1418931_at 0.99484 "regenerating islet-derived family, member 4" Reg4
1418932_at 0.91585 "nuclear factor, interleukin 3, regulated" Nfil3
1418933_at 0.91280 "solute carrier family 1 (high affinity aspartate/glutamate transporter), member 6" Slc1a6
1418934_at 0.90801 mab-21-like 2 (C. elegans) Mab21l2

1418935_at 0.90764 "transient receptor potential cation channel, subfamily M, member 1" Trpm1
1418936_at 1.26513 "v-maf musculoaponeurotic fibrosarcoma oncogene family, protein F (avian)" Maff
1418937_at 0.89076 "deiodinase, iodothyronine, type II" Dio2
1418938_at 0.97832 "deiodinase, iodothyronine, type II" Dio2
1418939_at 0.95279 H2.0-like homeo box 1 (Drosophila) Hlx1
1418940_at 0.97035 "sulfotransferase family 1B, member 1" Sult1b1
1418941_at 0.95242 protocadherin beta 22 Pcdhb22
1418942_at 1.00262 intraflagellar transport 74 homolog (Chlamydomonas) Ift74
1418943_at 0.85763 RIKEN cDNA B230120H23 gene B230120H23Rik
1418944_at 0.78195 cysteinyl leukotriene receptor 1 Cysltr1
1418945_at 1.01436 matrix metalloproteinase 3 Mmp3
1418946_at 1.05471 "ST3 beta-galactoside alpha-2,3-sialyltransferase 1" St3gal1
1418947_at 0.99395 NIMA (never in mitosis gene a)-related expressed kinase 3 Nek3
1418948_at 0.98403 "fascin homolog 3, actin-bundling protein, testicular (Strongylocentrotus purpuratus)" Fscn3
1418949_at 0.96307 growth differentiation factor 15 Gdf15
1418950_at 0.93500 dopamine receptor 2 Drd2
1418951_at 1.04343 taxilin beta Txlnb
1418952_at 0.89436 taxilin beta Txlnb
1418953_at 0.93130 F-box protein 16 Fbxo16
1418954_at 0.94649 "calcium/calmodulin-dependent protein kinase kinase 1, alpha" Camkk1
1418955_at 0.88855 zinc finger protein 93 Zfp93
1418956_at 1.18877 testis-specific serine kinase 6 Tssk6
1418957_at 0.87576 src homology three (SH3) and cysteine rich domain Stac
1418958_at 1.05718 acyl-malonyl condensing enzyme 1 Amac1
1418959_at 1.04620 "transmembrane protease, serine 5 (spinesin)" Tmprss5
1418960_at 1.07026 PHD finger protein 20-like 1 Phf20l1
1418961_at 0.95496 NECAP endocytosis associated 2 Necap2
1418962_at 0.98724 NECAP endocytosis associated 2 Necap2
1418963_at 0.90185 RIKEN cDNA 2310047O13 gene 2310047O13Rik
1418964_at 0.83289 "phosphatidylinositol glycan anchor biosynthesis, class M" Pigm
1418965_at 0.90095 nitric oxide synthase interacting protein Nosip
1418966_a_at 0.98490 "discoidin, CUB and LCCL domain containing 1" Dcbl1
1418967_a_at 0.99891 Suppression of tumorigenicity 7 St7
1418968_at 0.78661 RB1-inducible coiled-coil 1 Rb1cc1
1418969_at 1.06795 S-phase kinase-associated protein 2 (p45) Skp2
1418970_a_at 1.08235 B-cell leukemia/lymphoma 10 Bcl10
1418971_x_at 1.07027 B-cell leukemia/lymphoma 10 Bcl10
1418972_at 1.06461 B-cell leukemia/lymphoma 10 Bcl10
1418973_at 0.90587 basic leucine zipper nuclear factor 1 Blzf1
1418974_at 1.02781 basic leucine zipper nuclear factor 1 Blzf1
1418975_at 1.14385 NCK interacting protein with SH3 domain Nckipsd

1418976_s_at 0.84591 "cell death-inducing DNA fragmentation factor, alpha subunit-like effector B"
Cideb

1418977_at 0.90010 SAP30 binding protein Sap30bp

1418978_at 0.88391 SAP30 binding protein /// similar to transcriptional regulator protein
Sap30bp /// LOC639905

1418979_at 0.59823 "aldo-keto reductase family 1, member C14" Akr1c14

1418980_a_at 0.80916 cyclic nucleotide phosphodiesterase 1 Cnp1

1418981_at 1.04943 caspase 12 Casp12

1418982_at 0.89686 "CCAAT/enhancer binding protein (C/EBP), alpha" Cebpa

1418983_at 0.71909 InaD-like (Drosophila) Inadl

1418984_at 0.81535 InaD-like (Drosophila) Inadl

1418985_at 0.95469 CTTNBP2 N-terminal like Ctnnbp2nl

1418986_a_at 1.16387 ubiquitously expressed transcript Uxt

1418987_at 1.03458 "phospholipase A2, group IID" Pla2g2d

1418988_at 0.95731 peroxisome biogenesis factor 7 Pex7

1418989_at 0.83527 cathepsin E Ctse

1418990_at 1.04175 "membrane-spanning 4-domains, subfamily A, member 4D" Ms4a4d

1418991_at 1.05571 BCL2-antagonist/killer 1 Bak1

1418992_at 0.92912 coagulation factor X F10

1418993_s_at 0.92271 coagulation factor X F10

1418994_at 1.00288 RIKEN cDNA 2410116G06 gene 2410116G06Rik

1418995_at 1.01836 --- ---

1418996_a_at 0.94662 RIKEN cDNA 4930469P12 gene 4930469P12Rik

1418997_at 1.00699 --- ---

1418998_at 1.23911 kynurenine 3-monooxygenase (kynurenine 3-hydroxylase) Kmo

1418999_at 1.03935 RIKEN cDNA 2310033P09 gene 2310033P09Rik

1419000_at 0.88547 carboxypeptidase X 2 (M14 family) Cpxm2

1419001_at 1.08183 bile acid-Coenzyme A: amino acid N-acyltransferase Baat

1419002_s_at 0.90832 bile acid-Coenzyme A: amino acid N-acyltransferase Baat

1419003_at 1.07690 blood vessel epicardial substance Bves

1419004_s_at 0.97285 B-cell leukemia/lymphoma 2 related protein A1a /// B-cell leukemia/lymphoma
2 related protein A1b /// B-cell leukemia/lymphoma 2 related protein A1d Bcl2a1a /// Bcl2a1b ///
Bcl2a1d

1419005_at 0.85921 "crystallin, beta B3" Crybb3

1419006_s_at 0.95529 pellino 2 Peli2

1419007_at 0.97206 zona pellucida glycoprotein 3 Zp3

1419008_at 1.04701 neuropeptide Y receptor Y5 Npy5r

1419009_at 0.96975 actin-like 7a Actl7a

1419010_x_at 0.98769 kallikrein 1-related peptidase b5 Klk1b5

1419011_at 1.03988 "crystallin, beta A2" Cryba2

1419012_at 1.03922 "zinc finger protein, multitype 2" Zfpm2

1419013_at 0.91009 G patch domain containing 1 Gpatc1

1419014_at 0.92469 Rhesus blood group-associated A glycoprotein Rhag

1419015_at 1.03716 WNT1 inducible signaling pathway protein 2 Wisp2
1419016_at 0.96332 RIKEN cDNA 1700034I23 gene 1700034I23Rik
1419017_at 0.92414 corin Corin
1419018_at 0.81666 reproductive homeobox 6 Rhox6
1419019_a_at 0.93853 A kinase (PRKA) anchor protein 4 Akap4
1419020_at 0.99125 gastric intrinsic factor Gif
1419021_at 1.08524 mcf.2 transforming sequence Mcf2
1419022_a_at 1.04822 "enolase 1, alpha non-neuron /// similar to enolase 1, alpha non-neuron /// similar to Alpha-enolase (2-phospho-D-glycerate hydro-lyase) (Non-neural enolase) (NNE) (Enolase 1)" Eno1 /// LOC433182 /// LOC672232
1419023_x_at 1.05505 "enolase 1, alpha non-neuron" Eno1
1419024_at 1.07066 protein tyrosine phosphatase 4a1 Ptp4a1
1419025_at 1.14422 retinal S-antigen Sag
1419026_at 0.99254 Fas death domain-associated protein Daxx
1419027_s_at 0.90154 glycolipid transfer protein Gltf
1419028_at 0.93715 "cyclic AMP-regulated phosphoprotein, 21" Arpp21
1419029_at 1.42131 ERO1-like (S. cerevisiae) Ero1l
1419030_at 1.29477 ERO1-like (S. cerevisiae) Ero1l
1419031_at 1.14331 fatty acid desaturase 2 Fads2
1419032_at 1.12372 RIKEN cDNA 2610018G03 gene 2610018G03Rik
1419033_at 0.95047 --- ---
1419034_at 1.08681 "casein kinase 2, alpha 1 polypeptide" Csnk2a1
1419035_s_at 1.01851 "casein kinase 2, alpha 1 polypeptide" Csnk2a1
1419036_at 1.09983 "casein kinase 2, alpha 1 polypeptide" Csnk2a1
1419037_at 1.01032 "casein kinase 2, alpha 1 polypeptide" Csnk2a1
1419038_a_at 1.21260 "casein kinase 2, alpha 1 polypeptide" Csnk2a1
1419039_at 0.98611 "cytochrome P450, family 2, subfamily d, polypeptide 22" Cyp2d22
1419040_at 0.95565 "cytochrome P450, family 2, subfamily d, polypeptide 22" Cyp2d22
1419041_at 0.85546 integrin alpha FG-GAP repeat containing 1 Itfg1
1419042_at 0.99876 interferon inducible GTPase 1 Iigp1
1419043_a_at 1.33152 interferon inducible GTPase 1 Iigp1
1419044_at 1.16250 contactin associated protein 4 Cntnap4
1419045_at 0.68900 "solute carrier family 25 (mitochondrial carrier; phosphate carrier), member 23" Slc25a23
1419046_at 1.21353 brain protein 16 Brp16
1419047_at 0.93269 pecanex homolog (Drosophila) Pcnx
1419048_at 0.98747 pecanex homolog (Drosophila) Pcnx
1419049_at 0.78875 pecanex homolog (Drosophila) Pcnx
1419050_at 1.16248 RIKEN cDNA 1110002H13 gene 1110002H13Rik
1419051_at 0.98045 OVO homolog-like 1 (Drosophila) Ovol1
1419052_at 0.96487 OVO homolog-like 1 (Drosophila) Ovol1
1419053_at 0.85596 peroxisomal biogenesis factor 14 Pex14
1419054_a_at 1.00605 "protein tyrosine phosphatase, non-receptor type 21" Ptpn21

1419055_a_at 1.00760 "protein tyrosine phosphatase, non-receptor type 21" Ptpn21
1419056_at 0.79722 reticulon 2 (Z-band associated protein) Rtn2
1419057_at 0.79970 "solute carrier family 5 (sodium/glucose cotransporter), member 1" Slc5a1
1419058_at 1.17693 polymerase (RNA) I polypeptide E Polr1e
1419059_at 0.99560 serum amyloid P-component Apcs
1419060_at 0.95991 granzyme B Gzmb
1419061_at 0.95240 "ras homolog gene family, member D" Rhod
1419062_at 1.01865 erythrocyte protein band 4.1-like 3 Epb4.113
1419063_at 0.99185 UDP galactosyltransferase 8A Ugt8a
1419064_a_at 0.95602 UDP galactosyltransferase 8A Ugt8a
1419065_at 1.03800 RIKEN cDNA 5730521E12 gene 5730521E12Rik
1419066_at 0.79341 immediate early response 5-like Ier5l
1419067_a_at 1.03619 RAB guanine nucleotide exchange factor (GEF) 1 Rabgef1
1419068_at 0.98572 RAB guanine nucleotide exchange factor (GEF) 1 Rabgef1
1419069_at 0.91064 RAB guanine nucleotide exchange factor (GEF) 1 Rabgef1
1419070_at 0.74643 cystin 1 Cys1
1419071_at 0.78803 cystin 1 Cys1
1419072_at 0.90784 "glutathione S-transferase, mu 7" Gstm7
1419073_at 0.83240 transmembrane protein with EGF-like and two follistatin-like domains 2 Tmeff2
1419074_at 1.17020 RIKEN cDNA 2510006C20 gene 2510006C20Rik
1419075_s_at 0.82192 serum amyloid A 1 Saal
1419076_a_at 0.96171 breast cancer 2 Brca2
1419077_at 0.92046 "membrane protein, palmitoylated 3 (MAGUK p55 subfamily member 3)"
Mpp3
1419078_at 1.01784 ninein Nin
1419079_at 0.95398 "sodium channel, nonvoltage-gated 1 gamma" Scnn1g
1419080_at 0.99556 glial cell line derived neurotrophic factor Gdnf
1419081_at 0.79003 autophagy-related 10 (yeast) Atg10
1419082_at 1.10410 "serine (or cysteine) peptidase inhibitor, clade B, member 2" Serpinb2
1419083_at 0.97366 "tumor necrosis factor (ligand) superfamily, member 11" Tnfsf11
1419084_a_at 1.11578 Purkinje cell protein 2 (L7) Pcp2
1419085_at 1.00011 Purkinje cell protein 2 (L7) Pcp2
1419086_at 1.43209 fibroblast growth factor binding protein 1 Fgfbp1
1419087_s_at 0.85496 "splicing factor 3a, subunit 1" Sf3a1
1419088_at 0.70944 tissue inhibitor of metalloproteinase 3 Timp3
1419089_at 0.71220 tissue inhibitor of metalloproteinase 3 Timp3
1419090_x_at 1.01104 kallikrein 1-related peptidase b26 Klk1b26
1419091_a_at 0.98555 annexin A2 Anxa2
1419092_a_at 0.98005 STE20-like kinase (yeast) Slk
1419093_at 0.95167 "tryptophan 2,3-dioxygenase" Tdo2
1419094_at 0.99160 "cytochrome P450, family 2, subfamily c, polypeptide 37 /// cytochrome P450,
family 2, subfamily c, polypeptide 50 /// cytochrome P450 /// similar to cytochrome P450"
Cyp2c37 /// Cyp2c50 /// Cyp2c54 /// LOC639023

1419095_a_at	1.35646	apolipoprotein M	Apom
1419096_at	0.93473	apolipoprotein M	Apom
1419097_a_at	1.01558	stomatin	Stom
1419098_at	0.88803	stomatin	Stom
1419099_x_at	0.88576	stomatin	Stom
1419100_at	0.99816	"serine (or cysteine) peptidase inhibitor, clade A, member 3N"	Serpina3n
1419101_at	1.09174	"transcriptional regulator, SIN3A (yeast)"	Sin3a
1419102_at	1.09082	"transcriptional regulator, SIN3A (yeast)"	Sin3a
1419103_a_at	0.97709	abhydrolase domain containing 6	Abhd6
1419104_at	0.91965	abhydrolase domain containing 6	Abhd6
1419105_at	0.92538	"nuclear receptor subfamily 1, group H, member 4"	Nr1h4
1419106_at	1.07434	RIKEN cDNA 2210409E12 gene	2210409E12Rik
1419107_at	0.89657	oligophrenin 1	Ophn1
1419108_at	0.95640	oligophrenin 1	Ophn1
1419109_at	0.91481	histidine rich calcium binding protein	Hrc
1419110_at	0.89480	RIO kinase 1 (yeast)	Riok1
1419111_at	0.98956	"inhibitor of growth family, member 2"	Ing2
1419112_at	1.09895	nemo like kinase	Nlk
1419113_at	1.11376	"adaptor protein complex AP-1, gamma 2 subunit"	Ap1g2
1419114_at	0.93253	asparagine-linked glycosylation 14 homolog (yeast)	Alg14
1419115_at	0.92094	asparagine-linked glycosylation 14 homolog (yeast)	Alg14
1419116_at	1.02890	asparagine-linked glycosylation 14 homolog (yeast)	Alg14
1419117_at	0.85807	"solute carrier family 22 (organic cation transporter), member 2"	Slc22a2
1419118_at	1.00072	RIKEN cDNA 2900093B09 gene	2900093B09Rik
1419119_at	1.00663	hematopoietic cell signal transducer	Hcst
1419120_at	1.00102	lymphoblastomic leukemia	Lyl1
1419121_at	0.99752	thiopurine methyltransferase	Tpmt
1419122_at	1.15383	methyltransferase-like 1	Mettl1
1419123_a_at	0.67458	"platelet-derived growth factor, C polypeptide"	Pdgfc
1419124_at	0.82950	RIKEN cDNA 2210010L05 gene	2210010L05Rik
1419125_at	0.86618	"protein tyrosine phosphatase, non-receptor type 18"	Ptpn18
1419126_at	1.03048	homeo box D9	Hoxd9
1419127_at	0.91487	neuropeptide Y	Npy
1419128_at	0.90957	integrin alpha X	Itgax
1419129_at	0.93923	"solute carrier family 22 (organic cation transporter), member 13"	Slc22a13
1419130_at	1.21864	deaminase domain containing 1	Deadc1
1419131_at	1.10993	"coagulation factor XIII, beta subunit"	F13b
1419132_at	1.16331	toll-like receptor 2	Tlr2
1419133_at	0.87650	Envoplakin	Evpl
1419134_at	1.00965	Rhesus blood group-associated B glycoprotein	Rhbg
1419135_at	1.06824	lymphotoxin B	Ltb
1419136_at	0.95527	"aldo-keto reductase family 1, member C18"	Akr1c18
1419137_at	0.92936	SH3/ankyrin domain gene 3	Shank3

1419138_at 0.94782 "UDP-Gal:betaGalNAc beta 1,3-galactosyltransferase, polypeptide 4"
 B3galt4
 1419139_at 0.96583 growth differentiation factor 5 Gdf5
 1419140_at 1.01460 activin receptor IIB Acvr2b
 1419141_at 0.96686 "crystallin, gamma D" Crygd
 1419142_at 0.97966 cathepsin R Ctsr
 1419143_at 0.86345 cysteine-rich perinuclear theca 5 Cyp5
 1419144_at 0.93907 CD163 antigen Cd163
 1419145_at 0.90867 smoothelin-like 1 Smtnl1
 1419146_a_at 0.95925 glucokinase Gck
 1419147_at 1.00120 REC8-like 1 (yeast) Rec8L1
 1419148_at 0.90209 advillin Avil
 1419149_at 1.23910 "serine (or cysteine) peptidase inhibitor, clade E, member 1" Serpine1
 1419150_at 1.03097 myogenic factor 6 Myf6
 1419151_at 1.04694 RIKEN cDNA 4930449I24 gene 4930449I24Rik
 1419152_at 1.12060 RIKEN cDNA 2810417H13 gene 2810417H13Rik
 1419153_at 1.10126 RIKEN cDNA 2810417H13 gene 2810417H13Rik
 1419154_at 0.88767 "transmembrane protease, serine 2" Tmprs2
 1419155_a_at 0.73414 SRY-box containing gene 4 /// similar to Transcription factor SOX-4
 Sox4 /// LOC672274
 1419156_at 0.67671 SRY-box containing gene 4 /// similar to Transcription factor SOX-4 Sox4 ///
 LOC672274
 1419157_at 0.69594 SRY-box containing gene 4 /// similar to Transcription factor SOX-4 Sox4 ///
 LOC672274
 1419158_a_at 1.00165 histidyl-tRNA synthetase-like Harsl
 1419159_at 0.91581 "golgi autoantigen, golgin subfamily a, 3" Golga3
 1419160_at 0.81354 "golgi autoantigen, golgin subfamily a, 3" Golga3
 1419161_a_at 0.99265 NADPH oxidase 4 Nox4
 1419162_s_at 0.85702 "DnaJ (Hsp40) homolog, subfamily C, member 3" Dnajc3
 1419163_s_at 0.86276 "DnaJ (Hsp40) homolog, subfamily C, member 3" Dnajc3
 1419164_at 0.86178 zinc finger protein 260 Zfp260
 1419165_at 0.89636 zinc finger protein 260 /// similar to Zinc finger protein OZF (Zinc finger
 protein 260) (Zfp-260) Zfp260 /// LOC635007
 1419166_at 1.02782 "solute carrier family 5 (sodium/glucose cotransporter), member 2" Slc5a2
 1419167_at 0.92795 proline-rich acidic protein 1 Prap1
 1419168_at 1.13062 mitogen-activated protein kinase 6 Mapk6
 1419169_at 1.07750 mitogen-activated protein kinase 6 Mapk6
 1419170_at 0.88536 transmembrane protein 157 Tmem157
 1419171_at 0.95809 transmembrane protein 157 Tmem157
 1419172_at 0.89032 dihydrofolate reductase Dhfr
 1419173_at 1.10320 aminoacylase 1 Acyl1
 1419174_at 0.94032 RIKEN cDNA 2410004B18 gene 2410004B18Rik
 1419175_a_at 1.00915 "butyrophilin, subfamily 1, member A1" Btn1a1

1419176_at	1.04650	vacuolar protein sorting 37A (yeast)	Vps37a
1419177_at	1.01738	vacuolar protein sorting 37A (yeast)	Vps37a
1419178_at	1.04032	"CD3 antigen, gamma polypeptide"	Cd3g
1419179_at	0.94374	thioredoxin-like 4	Txn14
1419180_at	1.03249	B-cell CLL/lymphoma 9-like	Bcl9l
1419181_at	0.98125	zinc finger protein 326	Zfp326
1419182_at	0.98126	"sushi, von Willebrand factor type A, EGF and pentraxin domain containing 1"	Svep1
1419183_at	1.10091	PAP associated domain containing 4	Papd4
1419184_a_at	0.95078	four and a half LIM domains 2	Fhl2
1419185_a_at	1.05445	MLX interacting protein-like	Mlxip1
1419186_a_at	0.89322	"ST8 alpha-N-acetyl-neuraminide alpha-2,8-sialyltransferase 4"	St8sia4
1419187_at	1.01151	RIKEN cDNA 1700012L04 gene /// similar to Histone H2A.x (H2a/x) 1700012L04Rik /// LOC547154	
1419188_s_at	1.05556	chemokine (C-C motif) ligand 27	Ccl27
1419189_at	0.99206	vesicle transport through interaction with t-SNAREs homolog 1A (yeast)	Vti1a
1419190_at	1.03105	vesicle transport through interaction with t-SNAREs homolog 1A (yeast)	Vti1a
1419191_at	0.83959	homeodomain interacting protein kinase 3	Hipk3
1419192_at	0.80923	interleukin 4 induced 1	Il4i1
1419193_a_at	0.94602	"glia maturation factor, gamma"	Gmfg
1419194_s_at	0.93090	"glia maturation factor, gamma"	Gmfg
1419195_at	0.95916	WAP four-disulfide core domain 15	Wfdc15
1419196_at	1.02622	hepcidin antimicrobial peptide 1	Hamp1
1419197_x_at	0.96159	hepcidin antimicrobial peptide 1	Hamp1
1419198_at	0.93988	chromobox homolog 8 (Drosophila Pc class)	Cbx8
1419199_at	1.00834	chromobox homolog 8 (Drosophila Pc class)	Cbx8
1419200_at	0.94827	FXDY domain-containing ion transport regulator 7	Fxyd7
1419201_at	0.91752	kallikrein 11	Klk11
1419202_at	1.13524	cystatin F (leukocystatin)	Cst7
1419203_at	1.08939	gene trap locus F3a	Gtlf3a
1419204_at	1.02319	delta-like 1 (Drosophila)	Dll1
1419205_x_at	1.37464	G patch domain containing 4	Gpatc4
1419206_at	1.03249	CD37 antigen	Cd37
1419207_at	0.95166	zinc finger protein 37	Zfp37
1419208_at	0.89182	mitogen activated protein kinase kinase kinase 8	Map3k8
1419209_at	8.15299	chemokine (C-X-C motif) ligand 1	Cxcl1
1419210_at	1.17361	histamine receptor H 1	Hrh1
1419211_s_at	0.95587	RIKEN cDNA 4933424B01 gene	4933424B01Rik
1419212_at	1.31876	icos ligand	Icosl
1419213_at	1.02725	N-acetyltransferase 6	Nat6
1419214_at	0.87803	"tumor necrosis factor receptor superfamily, member 11a"	Tnfrsf11a
1419215_at	0.84548	aldehyde oxidase 4	Aox4
1419216_at	0.87884	5-azacytidine induced gene 1	Azil

1419217_at 0.91293 secretion regulating guanine nucleotide exchange factor Sergef
 1419218_at 0.84564 "vang-like 2 (van gogh, Drosophila)" Vangl2
 1419219_at 0.98207 "cytochrome P450, family 4, subfamily f, polypeptide 18" Cyp4f18
 1419220_at 0.89822 cardiomyopathy associated 1 Cmya1
 1419221_a_at 1.01031 regulator of G-protein signaling 14 Rgs14
 1419222_at 0.87057 thromboxane A2 receptor Tbx2r
 1419223_a_at 0.97651 dystrobrevin alpha Dtna
 1419224_at 1.00542 "cat eye syndrome chromosome region, candidate 6 homolog (human)"
 Cacr6
 1419225_at 1.10957 "calcium channel, voltage-dependent, alpha2/delta subunit 3" Cacna2d3
 1419226_at 0.93691 CD96 antigen Cd96
 1419227_at 1.07890 chaperonin subunit 6b (zeta) Cct6b
 1419228_at 1.07241 elacC homolog 1 (E. coli) Elac1
 1419229_at 0.96586 reproductive homeobox 4B /// reproductive homeobox 4E /// reproductive
 homeobox 4C /// reproductive homeobox 4H /// reproductive homeobox 4G /// reproductive homeobox
 4A /// reproductive homeobox 4D /// similar to reproductive homeobox 4E Rhox4b /// Rhox4e ///
 Rhox4c /// Rhox4h /// Rhox4g /// Rhox4a /// Rhox4d /// LOC665169
 1419230_at 1.05130 keratin 12 Krt12
 1419231_s_at 1.03808 keratin 12 Krt12
 1419232_a_at 0.96795 apolipoprotein A-I Apo1
 1419233_x_at 0.98021 apolipoprotein A-I Apo1
 1419234_at 1.18537 helicase (DNA) B Helb
 1419235_s_at 1.03971 helicase (DNA) B Helb
 1419236_at 1.05868 helicase (DNA) B Helb
 1419237_at 1.02358 ubiquitin specific peptidase 29 Usp29
 1419238_at 1.01950 "ATP-binding cassette, sub-family A (ABC1), member 7" Abca7
 1419239_at 0.95234 zinc finger protein 54 Zfp54
 1419240_at 0.98965 testis expressed gene 14 Tex14
 1419241_a_at 1.07161 autoimmune regulator (autoimmune polyendocrinopathy candidiasis
 ectodermal dystrophy) Aire
 1419242_at 0.94788 RIKEN cDNA 9530002B09 gene 9530002B09Rik
 1419243_at 0.97859 "RAB14, member RAS oncogene family" Rab14
 1419244_a_at 0.94482 "RAB14, member RAS oncogene family" Rab14
 1419245_at 0.75875 "RAB14, member RAS oncogene family" Rab14
 1419246_s_at 0.94471 "RAB14, member RAS oncogene family" Rab14
 1419247_at 0.80963 regulator of G-protein signaling 2 Rgs2
 1419248_at 0.84777 regulator of G-protein signaling 2 Rgs2
 1419249_at 0.85239 PFTAIRES protein kinase 1 Pftk1
 1419250_a_at 0.81681 PFTAIRES protein kinase 1 Pftk1
 1419251_at 1.07202 epidermal growth factor receptor pathway substrate 15 Eps15
 1419252_at 0.99614 --- ---
 1419253_at 0.97224 "methylenetetrahydrofolate dehydrogenase (NAD+ dependent),
 methenyltetrahydrofolate cyclohydrolase" Mthfd2

1419254_at	1.06571	"methylenetetrahydrofolate dehydrogenase (NAD+ dependent), methenyltetrahydrofolate cyclohydrolase"	Mthfd2	
1419255_at	1.51739	spectrin beta 2	Spnb2	
1419256_at	0.99570	spectrin beta 2	Spnb2	
1419257_at	1.15257	transcription elongation factor A (SII) 1	Tcea1	
1419258_at	1.11832	transcription elongation factor A (SII) 1	Tcea1	
1419259_at	0.87142	Ras suppressor protein 1	Rsu1	
1419260_a_at	1.13606	small nuclear ribonucleoprotein B	Snrpb	
1419261_at	0.94400	"acyl-Coenzyme A dehydrogenase family, member 8"		Acad8
1419262_at	0.95356	"acyl-Coenzyme A dehydrogenase family, member 8"		Acad8
1419263_a_at	1.03983	adhesion regulating molecule 1	Adrm1	
1419264_at	1.02117	RIKEN cDNA 1200014M14 gene	1200014M14Rik	
1419265_at	0.95489	RIKEN cDNA 1200014M14 gene	1200014M14Rik	
1419266_at	1.03593	nuclear transcription factor-Y beta	Nfyb	
1419267_at	0.94393	nuclear transcription factor-Y beta	Nfyb	
1419268_at	1.02572	anterior gradient 2 (Xenopus laevis)	Agr2	
1419269_at	0.75474	deoxyuridine triphosphatase	Dut	
1419270_a_at	1.00582	deoxyuridine triphosphatase	Dut	
1419271_at	1.15105	paired box gene 6	Pax6	
1419272_at	1.02065	myeloid differentiation primary response gene 88		Myd88
1419273_at	1.07734	expressed sequence C80913	C80913	
1419274_at	1.10736	expressed sequence C80913	C80913	
1419275_at	1.04350	DAZ associated protein 1	Dazap1	
1419276_at	0.68422	ectonucleotide pyrophosphatase/phosphodiesterase 1	Enpp1	
1419277_at	0.96914	ubiquitin specific peptidase 48	Usp48	
1419278_at	0.97995	ubiquitin specific peptidase 48	Usp48	
1419279_at	0.69683	"phosphatidylinositol-4-phosphate 5-kinase, type II, alpha"		Pip5k2a
1419280_at	0.59692	"phosphatidylinositol-4-phosphate 5-kinase, type II, alpha"		Pip5k2a
1419281_a_at	1.08361	zinc finger protein 259	Zfp259	
1419282_at	0.98257	chemokine (C-C motif) ligand 12	Ccl12	
1419283_s_at	0.81207	tensin 1	Tns1	
1419284_at	0.97868	cysteine and histidine rich 1	Cyhr1	
1419285_s_at	0.96105	cysteine and histidine rich 1	Cyhr1	
1419286_s_at	0.92441	intraflagellar transport 81 homolog (Chlamydomonas)	Ift81	
1419287_at	0.96733	HSPC171 protein	Hspc171	
1419288_at	0.99582	junction adhesion molecule 2	Jam2	
1419289_a_at	0.91733	synaptogyrin 1	Syngr1	
1419290_at	1.10972	methyltransferase like 6	Mettl6	
1419291_x_at	1.14625	growth arrest specific 5	Gas5	
1419292_at	0.81119	HtrA serine peptidase 3	Htra3	
1419293_at	0.91228	Down syndrome cell adhesion molecule	Dscam	
1419294_at	1.06061	RIKEN cDNA 1700011H14 gene	1700011H14Rik	
1419295_at	0.98525	cAMP responsive element binding protein 3-like 1		Creb3l1

1419296_at	0.95669	Rho GTPase activating protein 4	Arhgap4	
1419297_at	1.02710	"histocompatibility 2, O region alpha locus"	H2-Oa	
1419298_at	0.84226	paraoxonase 3	Pon3	
1419299_at	0.96153	RIKEN cDNA 2010012O05 gene	2010012O05Rik	
1419300_at	0.94367	FMS-like tyrosine kinase 1	Flt1	
1419301_at	0.71424	frizzled homolog 4 (Drosophila)	Fzd4	
1419302_at	1.05882	hairy/enhancer-of-split related with YRPW motif-like	Heyl	
1419303_at	1.02284	hairy/enhancer-of-split related with YRPW motif-like	Heyl	
1419304_at	0.98754	brachyury	T	
1419305_a_at	0.96483	F-box only protein 36	Fbxo36	
1419306_at	0.74834	coiled-coil domain containing 96	Ccdc96	
1419307_at	0.89380	"tumor necrosis factor receptor superfamily, member 13c"	Tnfrsf13c	
1419308_at	0.89860	inversin	Invs	
1419309_at	1.14401	podoplanin	Pdpn	
1419310_s_at	0.89542	regulatory factor X-associated ankyrin-containing protein	Rfxank	
1419311_at	1.12936	tripartite motif protein 10	Trim10	
1419312_at	0.98042	"ATPase, Ca ⁺⁺ transporting, cardiac muscle, fast twitch 1"	Atp2a1	
1419313_at	1.05082	cyclin T1	Ccnt1	
1419314_at	0.49222	tubulointerstitial nephritis antigen	Tinag	
1419315_at	0.94486	SLAM family member 9	Slamf9	
1419316_s_at	1.00569	"guanine nucleotide binding protein (G protein), beta polypeptide 1-like"	Gnb1l	
1419317_x_at	0.87717	RIKEN cDNA 2310002A05 gene /// small proline rich-like 1 /// similar to late cornified envelope protein /// hypothetical protein LOC630971 /// similar to late cornified envelope protein 2310002A05Rik /// Sprr11 /// LOC545548 /// LOC630971 /// LOC630994		
1419318_at	0.91067	serum amyloid A 4	Saa4	
1419319_at	0.89212	serum amyloid A 4	Saa4	
1419320_at	1.01245	carbohydrate (N-acetylglucosamine 6-O) sulfotransferase 5	Chst5	
1419321_at	0.93464	coagulation factor VII	F7	
1419322_at	0.94418	"FYVE, RhoGEF and PH domain containing 6"	Fgd6	
1419323_at	1.01893	"peptidyl arginine deiminase, type I"	Padi1	
1419324_at	0.85071	LIM homeobox protein 9	Lhx9	
1419325_at	0.87045	neuromedin U	Nmu	
1419326_at	1.01137	RIKEN cDNA 1700028P14 gene	1700028P14Rik	
1419327_at	0.92381	RIKEN cDNA 2210010A19 gene	2210010A19Rik	
1419328_at	0.88148	"sema domain, immunoglobulin domain (Ig), TM domain, and short cytoplasmic domain"	Sema4f	
1419329_at	0.94298	sorbin and SH3 domain containing 3	Sorbs3	
1419330_a_at	1.04660	glycoprotein A33 (transmembrane)	Gpa33	
1419331_at	0.95173	cadherin 17	Cdh17	
1419332_at	0.99322	"EGF-like-domain, multiple 6"	Egfl6	
1419333_at	1.03092	RIKEN cDNA 1110008J03 gene	1110008J03Rik	
1419334_at	1.02714	cytotoxic T-lymphocyte-associated protein 4	Ctla4	

1419335_at 0.93717 RIKEN cDNA 4930441O14 gene 4930441O14Rik
1419336_at 0.92747 lipocalin 5 Lcn5
1419337_at 0.92613 RIKEN cDNA 1700080E11 gene 1700080E11Rik
1419338_at 0.91878 ICRFP703B1614Q5.5 gene ICRFP703B1614Q5.5
1419339_at 0.93565 neuraminidase 3 Neu3
1419340_at 1.11028 Moloney leukemia virus 10-like 1 Mov10l1
1419341_at 0.99832 Eph receptor A8 EphA8
1419342_at 0.94194 translocating chain-associating membrane protein 2 Tram2
1419343_at 1.09628 "solute carrier family 15 (oligopeptide transporter), member 1" Slc15a1
1419344_at 1.12792 t-complex-associated testis expressed 1 Tcte1
1419345_at 1.09527 parathyroid hormone Pth
1419346_a_at 0.99136 seminal vesicle secretion 5 Svs5
1419347_x_at 0.96614 seminal vesicle secretion 5 Svs5
1419348_at 0.95012 parotid secretory protein Psp
1419349_a_at 0.87250 "cytochrome P450, family 2, subfamily d, polypeptide 9" Cyp2d9
1419350_at 1.11085 hook homolog 2 (Drosophila) Hook2
1419351_a_at 1.10329 RIKEN cDNA 0610007P06 gene 0610007P06Rik
1419352_at 1.09866 RIKEN cDNA 0610007P06 gene 0610007P06Rik
1419353_at 0.93706 dolichol-phosphate (beta-D) mannosyltransferase 1 Dpm1
1419354_at 0.83277 Kruppel-like factor 7 (ubiquitous) Klf7
1419355_at 0.96870 Kruppel-like factor 7 (ubiquitous) Klf7
1419356_at 0.90601 Kruppel-like factor 7 (ubiquitous) Klf7
1419357_at 1.04757 RIKEN cDNA 5830446M03 gene 5830446M03Rik
1419358_at 0.76647 sortilin-related VPS10 domain containing receptor 2 Sorcs2
1419359_at 1.02117 hexamethylene bis-acetamide inducible 1 Hexim1
1419360_a_at 0.91339 "synovial sarcoma translocation, Chromosome 18" Ss18
1419361_at 0.94448 "synovial sarcoma translocation, Chromosome 18" Ss18
1419362_at 0.92989 mitochondrial ribosomal protein L35 Mrpl35
1419363_a_at 1.07807 mitochondrial ribosomal protein L35 Mrpl35
1419364_a_at 1.00764 ribosomal protein S7 /// similar to 40S ribosomal protein S7 (S8) /// similar to
40S ribosomal protein S7 (S8) /// similar to 40S ribosomal protein S7 (S8) /// similar to 40S ribosomal
protein S7 (S8) Rps7 /// LOC621699 /// LOC624124 /// LOC634044 /// LOC670565
1419365_at 0.96524 peroxisomal biogenesis factor 11a Pex11a
1419366_at 1.06388 "zinc finger, matrin type 5" Zmat5
1419367_at 1.01163 "2,4-dienoyl CoA reductase 1, mitochondrial" Decr1
1419368_a_at 1.03480 ring finger protein 138 Rnf138
1419369_at 0.86515 ring finger protein 138 Rnf138
1419370_a_at 0.99718 microfibrillar-associated protein 1 Mfap1
1419371_s_at 0.93712 golgi SNAP receptor complex member 2 Gosr2
1419372_at 1.03194 golgi SNAP receptor complex member 2 Gosr2
1419373_at 0.95093 "ATPase, H+ transporting, lysosomal V1 subunit B1" Atp6v1b1
1419374_at 1.10385 WW domain binding protein 4 Wbp4
1419375_at 0.91924 WW domain binding protein 4 Wbp4

1419376_at 0.74819 RIKEN cDNA 1110018M03 gene 1110018M03Rik
 1419377_at 1.02209 "mediator of RNA polymerase II transcription, subunit 9 homolog (yeast)"
 Med9
 1419378_a_at 0.96580 FXYP domain-containing ion transport regulator 2 Fxyd2
 1419379_x_at 1.00783 FXYP domain-containing ion transport regulator 2 Fxyd2
 1419380_at 1.01134 zinc finger protein 423 Zfp423
 1419381_at 0.93982 "telomeric repeat binding factor 2, interacting protein" Terf2ip
 1419382_a_at 1.04815 dehydrogenase/reductase (SDR family) member 4 Dhrs4
 1419383_at 0.88965 "S100 protein, beta polypeptide, neural" S100b
 1419384_at 1.00773 protein interacting with C kinase 1 Pick1
 1419385_a_at 1.03260 ubiquilin 1 Ubqln1
 1419386_at 0.98380 "mucin 13, epithelial transmembrane" Muc13
 1419387_s_at 0.86109 "mucin 13, epithelial transmembrane" Muc13
 1419388_at 1.01879 transmembrane 4 L six family member 20 Tm4sf20
 1419389_at 1.08663 phosphodiesterase 10A Pde10a
 1419390_at 1.10786 phosphodiesterase 10A Pde10a
 1419391_at 0.96259 myogenin Myog
 1419392_at 0.97613 piccolo (presynaptic cytomatrix protein) Pclo
 1419393_at 0.89184 "ATP-binding cassette, sub-family G (WHITE), member 5" Abcg5
 1419394_s_at 0.97088 S100 calcium binding protein A8 (calgranulin A) S100a8
 1419395_at 0.99038 acyl-CoA thioesterase 12 Acot12
 1419396_at 1.07854 AT rich interactive domain 3A (Bright like) Arid3a
 1419397_at 0.96928 "polymerase (DNA directed), alpha 1" Pola1
 1419398_a_at 0.96080 receptor accessory protein 5 Reep5
 1419399_at 1.08683 microsomal triglyceride transfer protein Mttp
 1419400_at 1.07935 microsomal triglyceride transfer protein Mttp
 1419401_at 0.94376 ankyrin repeat and SOCS box-containing protein 13 Asb13
 1419402_at 0.94986 meiosis-specific nuclear structural protein 1 Mns1
 1419403_at 1.02827 cDNA sequence BC017612 BC017612
 1419404_s_at 0.82161 seven in absentia 1A /// seven in absentia 1B Siah1a /// Siah1b
 1419405_at 0.98097 neuromedin B Nmb
 1419406_a_at 1.08882 B-cell CLL/lymphoma 11A (zinc finger protein) Bcl11a
 1419407_at 1.09224 hemolytic complement Hc
 1419408_at 0.85188 sine oculis-related homeobox 6 homolog (Drosophila) Six6
 1419409_at 0.99682 small proline rich-like 5 Sprrl5
 1419410_at 1.03071 "basic leucine zipper transcription factor, ATF-like" Batf
 1419411_at 1.02092 tachykinin 2 Tac2
 1419412_at 0.98846 chemokine (C motif) ligand 1 Xcl1
 1419413_at 1.26455 chemokine (C-C motif) ligand 17 Ccl17
 1419414_at 1.03281 "guanine nucleotide binding protein 13, gamma" Gng13
 1419415_a_at 0.94269 "retinoic acid receptor, gamma" Rarg
 1419416_a_at 1.01532 "retinoic acid receptor, gamma" Rarg
 1419417_at 0.75396 vascular endothelial growth factor C Vegfc

1419418_a_at 1.15242 microrchidia 1 Morc1
1419419_at 1.08651 "DNA segment, human D6S2654E" D0H6S2654E
1419420_at 0.86449 "ST6 (alpha-N-acetyl-neuraminy1-2,3-beta-galactosyl-1,3)-N-acetylgalactosaminide alpha-2,6-sialyltransferase 5" St6galnac5
1419421_at 0.88770 "ankyrin 1, erythroid" Ank1
1419422_at 0.98919 polycystic kidney disease 2-like 2 Pkd2l2
1419423_at 0.98023 stabilin 2 Stab2
1419424_at 0.96868 "pancreas specific transcription factor, 1a" Ptf1a
1419425_at 0.99035 cannabinoid receptor 1 (brain) Cnr1
1419426_s_at 0.95679 chemokine (C-C motif) ligand 21b /// chemokine (C-C motif) ligand 21a /// chemokine (C-C motif) ligand 21c (leucine) Ccl21b /// Ccl21a /// Ccl21c
1419427_at 1.28720 colony stimulating factor 3 (granulocyte) Csf3
1419428_a_at 0.90348 "glucosidase, alpha, acid" Gaa
1419429_at 1.03689 ciliary neurotrophic factor receptor Cntfr
1419430_at 0.96450 "cytochrome P450, family 26, subfamily a, polypeptide 1" Cyp26a1
1419431_at 0.87500 epiregulin Ereg
1419432_at 1.02544 sperm adhesion molecule 1 Spam1
1419433_at 1.06270 nth (endonuclease III)-like 1 (E.coli) Nth1
1419434_at 0.95997 "solute carrier family 2 (facilitated glucose transporter), member 10" Slc2a10
1419435_at 0.89119 aldehyde oxidase 1 Aox1
1419436_at 0.98392 complement factor H-related 1 Cfhr1
1419437_at 0.98063 single-minded homolog 2 (Drosophila) Sim2
1419438_at 1.17413 single-minded homolog 2 (Drosophila) Sim2
1419439_at 1.00857 serine/threonine kinase 22 substrate 1 /// similar to serine/threonine kinase 22 substrate 1 Stk22s1 /// LOC672873
1419440_at 1.03274 tripartite motif-containing 54 Trim54
1419441_at 1.02805 acidic ribosomal phosphoprotein P0 Arbp
1419442_at 0.86344 matrilin 2 Matn2
1419443_at 1.05311 Sin3-associated polypeptide 18 Sap18
1419444_at 1.13434 Sin3-associated polypeptide 18 Sap18
1419445_s_at 1.07232 Sin3-associated polypeptide 18 Sap18
1419446_at 0.83686 "TBC1 domain family, member 1" Tbc1d1
1419447_s_at 0.92220 "TBC1 domain family, member 1" Tbc1d1
1419448_at 0.92452 "TBC1 domain family, member 1" Tbc1d1
1419449_a_at 0.96191 "guanine nucleotide binding protein, alpha inhibiting 2" Gnai2
1419450_at 1.03739 ORM1-like 3 (S. cerevisiae) Ormdl3
1419451_at 0.94650 fizzy/cell division cycle 20 related 1 (Drosophila) Fzr1
1419452_at 1.06388 ubiquitin carboxyl-terminal esterase L5 Uchl5
1419453_at 1.09238 ubiquitin carboxyl-terminal esterase L5 Uchl5
1419454_x_at 0.91919 protein inhibitor of activated STAT 2 Pias2
1419455_at 1.13132 "interleukin 10 receptor, beta" Il10rb
1419456_at 0.91391 dicarbonyl L-xylulose reductase Dcxr
1419457_at 0.98686 Rho-guanine nucleotide exchange factor Rgnef

1419458_at	1.01788	Rho-guanine nucleotide exchange factor	Rgnef
1419459_a_at	0.97371	RIKEN cDNA 2610529C04 gene	2610529C04Rik
1419460_at	1.03691	ribonuclease P 14 subunit (human)	Rpp14
1419461_at	1.08310	ribonuclease P 14 subunit (human)	Rpp14
1419462_s_at	1.02853	gene trap locus 3	Gtl3
1419463_at	2.04763	chloride channel calcium activated 2	Clca2
1419464_at	0.99949	RIKEN cDNA 2810046M22 gene	2810046M22Rik
1419465_at	0.98087	naked cuticle 2 homolog (Drosophila)	Nkd2
1419466_at	0.98356	naked cuticle 2 homolog (Drosophila)	Nkd2
1419467_at	1.10987	"C-type lectin domain family 14, member a"	Clec14a
1419468_at	1.02658	"C-type lectin domain family 14, member a"	Clec14a
1419469_at	0.73713	"guanine nucleotide binding protein, beta 4"	Gnb4
1419470_at	0.76323	"guanine nucleotide binding protein, beta 4"	Gnb4
1419471_a_at	1.11955	nuclear distribution gene C homolog (Aspergillus)	Nudc
1419472_s_at	1.08025	"nuclear distribution gene C homolog (Aspergillus) /// nuclear distribution gene C homolog (Aspergillus), pseudogene 1"	Nudc /// Nudc-ps1
1419473_a_at	0.98468	cholecystokinin	Cck
1419474_a_at	1.01629	ets homologous factor	Ehf
1419475_a_at	1.00647	ets homologous factor	Ehf
1419476_at	0.94488	"ADAM-like, decysin 1"	Adamdec1
1419477_at	1.04456	"C-type lectin domain family 2, member d"	Clec2d
1419478_at	1.07255	secreted and transmembrane 1	Sectm1
1419479_at	1.02519	secreted and transmembrane 1	Sectm1
1419480_at	0.97827	"selectin, lymphocyte"	Sell
1419481_at	0.99259	"selectin, lymphocyte"	Sell
1419482_at	1.14830	complement component 3a receptor 1	C3ar1
1419483_at	1.17715	complement component 3a receptor 1	C3ar1
1419484_a_at	0.86042	glioblastoma amplified sequence	Gbas
1419485_at	0.76069	forkhead box C1	Foxc1
1419486_at	0.68906	forkhead box C1	Foxc1
1419487_at	0.88515	myosin binding protein H	Mybph
1419488_at	1.03498	TNFAIP3 interacting protein 2	Tnip2
1419489_at	0.97881	expressed sequence AW049604	AW049604
1419490_at	0.94228	expressed sequence AW049604	AW049604
1419491_at	0.90415	defensin beta 1	Defb1
1419492_s_at	0.97287	defensin beta 1	Defb1
1419493_a_at	0.97467	tumor protein D52	Tpd52
1419494_a_at	1.02403	tumor protein D52	Tpd52
1419495_at	1.07617	IMP2 inner mitochondrial membrane peptidase-like (S. cerevisiae)	Imp21
1419496_at	1.01185	"solute carrier organic anion transporter family, member 1a6"	Slco1a6
1419497_at	0.81809	cyclin-dependent kinase inhibitor 1B	Cdkn1b
1419498_at	0.99194	transmembrane and immunoglobulin domain containing 1	Tmigd1
1419499_at	0.92926	"glycerol-3-phosphate acyltransferase, mitochondrial"	Gpam

1419500_at 0.89757 "poly A binding protein, cytoplasmic 2" Pabpc2
1419501_at 1.18028 "polymerase (DNA directed), kappa" Polk
1419502_at 0.94475 GH3 domain containing Ghdc
1419503_at 0.99073 stanniocalcin 2 Stc2
1419504_at 1.20253 monoacylglycerol O-acyltransferase 1 Mogat1
1419505_a_at 0.96397 geranylgeranyl diphosphate synthase 1 Ggps1
1419506_at 0.95768 geranylgeranyl diphosphate synthase 1 Ggps1
1419507_at 1.00626 keratin associated protein 15 Krtap15
1419508_at 0.93220 receptor (TNFRSF)-interacting serine-threonine kinase 1 Ripk1
1419509_a_at 1.04620 N-acetylglucosamine kinase Nagk
1419510_at 0.93627 esterase 22 Es22
1419511_at 0.96101 mutS homolog 4 (E. coli) Msh4
1419512_at 0.77368 PRP40 pre-mRNA processing factor 40 homolog B (yeast) Prpf40b
1419513_a_at 0.97491 ect2 oncogene Ect2
1419514_at 0.98707 paired-like homeodomain transcription factor 1 Pitx1
1419515_at 0.99863 "FYVE, RhoGEF and PH domain containing 2" Fgd2
1419516_at 1.09765 "DNA segment, human DXS9928E" D0HXS9928E
1419517_at 0.98151 cornichon homolog 3 (Drosophila) Cnih3
1419518_at 0.96761 "tubulin, alpha 8" Tuba8
1419519_at 0.69357 insulin-like growth factor 1 Igf1
1419520_at 0.99033 camello-like 4 Cml4
1419521_at 1.12495 zinc finger protein 94 Zfp94
1419522_at 1.14369 "zinc finger, MYND domain containing 19" Zmynd19
1419523_at 1.04517 "cytochrome P450, family 3, subfamily a, polypeptide 13" Cyp3a13
1419524_at 1.08662 tryptophan hydroxylase 1 Tph1
1419525_at 1.10218 "carbonic anhydrase 5a, mitochondrial" Car5a
1419526_at 0.84660 Gardner-Rasheed feline sarcoma viral (Fgr) oncogene homolog Fgr
1419527_at 0.75984 cartilage oligomeric matrix protein Comp
1419528_at 0.92422 "sulfotransferase family 2A, dehydroepiandrosterone (DHEA)-preferring, member 2" Sult2a2
1419529_at 0.95172 "interleukin 23, alpha subunit p19" Il23a
1419530_at 1.03911 interleukin 12b Il12b
1419531_at 0.93696 RIKEN cDNA 1700102P08 gene 1700102P08Rik
1419532_at 1.05634 "interleukin 1 receptor, type II" Il1r2
1419533_at 1.07490 nescient helix loop helix 1 Nhlh1
1419534_at 1.11611 oxidized low density lipoprotein (lectin-like) receptor 1 Olr1
1419535_at 0.86188 "solute carrier organic anion transporter family, member 6b1" Slco6b1
1419536_a_at 0.94349 v-rel reticuloendotheliosis viral oncogene homolog A (avian) Rela
1419537_at 0.99572 transcription factor EC Tcfec
1419538_at 0.99307 FMS-like tyrosine kinase 3 Flt3
1419539_at 0.89846 Iroquois related homeobox 4 (Drosophila) Irx4
1419540_at 1.03867 "ferritin, heavy polypeptide-like 17 /// similar to ferritin, heavy polypeptide-like 17 /// similar to ferritin, heavy polypeptide-like 17 /// similar to ferritin, heavy polypeptide-like 17"

Fthl17 /// LOC434726 /// LOC434727 /// LOC434728

1419541_at	0.98722	RIKEN cDNA 4930403L05 gene	4930403L05Rik
1419542_at	0.93429	deleted in azoospermia-like	Dazl
1419543_a_at	1.06504	"splicing factor, arginine/serine-rich 10 (transformer 2 homolog, Drosophila)"	Sfrs10
1419544_at	1.07331	"ATPase, H+ transporting, lysosomal V1 subunit C1"	Atp6v1c1
1419545_a_at	1.07527	"ATPase, H+ transporting, lysosomal V1 subunit C1"	Atp6v1c1
1419546_at	1.06271	"ATPase, H+ transporting, lysosomal V1 subunit C1"	Atp6v1c1
1419547_at	1.20378	fumarylacetoacetate hydrolase domain containing 1	Fahd1
1419548_at	1.06155	karyopherin (importin) alpha 1	Kpna1
1419549_at	1.00034	"arginase 1, liver"	Arg1
1419550_a_at	0.88131	"serine/threonine kinase 39, STE20/SPS1 homolog (yeast)"	Stk39
1419551_s_at	0.90393	"serine/threonine kinase 39, STE20/SPS1 homolog (yeast)"	Stk39
1419552_at	1.08624	enoyl Coenzyme A hydratase domain containing 1	Echdc1
1419553_a_at	1.04473	"RAB geranylgeranyl transferase, b subunit"	Rabggtb
1419554_at	1.23286	"CD47 antigen (Rh-related antigen, integrin-associated signal transducer)"	Cd47
1419555_at	0.99217	E74-like factor 5	Elf5
1419556_at	0.92753	E74-like factor 5	Elf5
1419557_a_at	0.81672	transmembrane protein 9	Tmem9
1419558_at	1.02193	transformed mouse 3T3 cell double minute 4	Mdm4
1419559_at	0.90362	"cytochrome P450, family 4, subfamily f, polypeptide 14"	Cyp4f14
1419560_at	0.75762	"lipase, hepatic"	Lipc
1419561_at	0.88731	chemokine (C-C motif) ligand 3	Ccl3
1419562_at	1.02439	baculoviral IAP repeat-containing 6	Birc6
1419563_at	1.02446	baculoviral IAP repeat-containing 6	Birc6
1419564_at	0.86968	zinc finger protein 467	Zfp467
1419565_a_at	0.87361	zinc finger protein X-linked	Zfx
1419566_at	0.88730	fibronectin type 3 and ankyrin repeat domains 1	Fank1
1419567_at	1.01334	fibronectin type 3 and ankyrin repeat domains 1	Fank1
1419568_at	1.06818	mitogen activated protein kinase 1	Mapk1
1419569_a_at	0.99418	interferon-stimulated protein	Isg20
1419570_at	1.04513	"solute carrier family 28 (sodium-coupled nucleoside transporter), member 3"	Slc28a3
1419571_at	1.03754	"solute carrier family 28 (sodium-coupled nucleoside transporter), member 3"	Slc28a3
1419572_a_at	1.14186	"ATP-binding cassette, sub-family D (ALD), member 4"	Abcd4
1419573_a_at	1.00211	"lectin, galactose binding, soluble 1"	Lgals1
1419574_at	0.94135	zinc finger protein 292	Zfp292
1419575_s_at	0.91093	zinc finger protein 292	Zfp292
1419576_at	1.04566	homeo box B13	Hoxb13
1419577_at	0.83196	RIKEN cDNA A530089I17 gene	A530089I17Rik
1419578_at	1.01525	mannose binding lectin (A)	Mbl1

1419579_at 1.03354 "solute carrier family 7 (cationic amino acid transporter, y+ system), member 12" Slc7a12

1419580_at 0.93384 --- ---

1419581_at 0.77775 "discs, large homolog 4 (Drosophila)" Dlgh4

1419582_at 1.00805 "cytochrome P450, family 2, subfamily c, polypeptide 55" Cyp2c55

1419583_at 0.79359 chromobox homolog 4 (Drosophila Pc class) Cbx4

1419584_at 0.74050 tetratricopeptide repeat domain 28 Ttc28

1419585_at 1.10238 retinitis pigmentosa 2 homolog (human) Rp2h

1419586_at 1.00528 retinitis pigmentosa 2 homolog (human) Rp2h

1419587_s_at 1.01457 retinitis pigmentosa 2 homolog (human) Rp2h

1419588_at 0.92606 sperm associated antigen 1 Spag1

1419589_at 0.90483 CD93 antigen Cd93

1419590_at 0.88949 "cytochrome P450, family 2, subfamily b, polypeptide 9" Cyp2b9

1419591_at 0.94153 "melanoma-derived leucine zipper, extra-nuclear factor" Mlze

1419592_at 0.90638 unc-5 homolog C (C. elegans) Unc5c

1419593_at 0.85992 gene regulated by estrogen in breast cancer protein Greb1

1419594_at 0.89711 cathepsin G Ctsg

1419595_a_at 0.86261 gamma-glutamyl hydrolase /// similar to Gamma-glutamyl hydrolase precursor (Gamma-Glu-x carboxypeptidase) (Conjugase) (GH) (Folylpolyglutamate hydrolase) (FGPH) Ggh /// LOC667301

1419596_at 0.94694 ectodysplasin-A /// similar to Ectodysplasin-A (EDA protein homolog) (Tabby protein) Eda /// LOC675952

1419597_at 0.97506 ectodysplasin-A Eda

1419598_at 0.93933 "membrane-spanning 4-domains, subfamily A, member 6D" Ms4a6d

1419599_s_at 1.04870 "membrane-spanning 4-domains, subfamily A, member 11" Ms4a11

1419600_at 0.96046 defensin beta 4 Defb4

1419601_at 1.01092 "potassium inwardly-rectifying channel, subfamily J, member 10" Kcnj10

1419602_at 0.94302 homeo box A2 Hoxa2

1419603_at 1.11061 similar to Interferon-activatable protein 204 (Ifi-204) (Interferon-inducible protein p204) LOC672547

1419604_at 0.97391 Z-DNA binding protein 1 Zbp1

1419605_at 0.88862 macrophage galactose N-acetyl-galactosamine specific lectin 1 Mgl1

1419606_a_at 0.98331 "troponin T1, skeletal, slow" Tnnt1

1419607_at 0.84808 tumor necrosis factor Tnf

1419608_a_at 0.94271 melanoma inhibitory activity 1 Mia1

1419609_at 1.01230 chemokine (C-C motif) receptor 1 Ccr1

1419610_at 0.99913 chemokine (C-C motif) receptor 1 Ccr1

1419611_at 0.97151 --- ---

1419612_at 0.98301 RIKEN cDNA 4632415L05 gene 4632415L05Rik

1419613_at 1.32142 "procollagen, type VII, alpha 1" Col7a1

1419614_at 1.06522 "phospholipase A2, group XIIB" Pla2g12b

1419615_at 0.88687 "transient receptor potential cation channel, subfamily V, member 6" Trpv6

1419616_at 1.16928 "bone morphogenic protein receptor, type II (serine/threonine kinase)" Bmpr2

1419617_at 0.88696 "potassium intermediate/small conductance calcium-activated channel, subfamily N, member 1" Kcnn1
 1419618_at 1.07816 "butyrobetaine (gamma), 2-oxoglutarate dioxygenase 1 (gamma-butyrobetaine hydroxylase)" Bbox1
 1419619_at 1.05821 keratin 80 Krt80
 1419620_at 1.25014 pituitary tumor-transforming 1 Pttg1
 1419621_at 0.99071 ankyrin repeat domain 2 (stretch responsive muscle) Ankrd2
 1419622_at 0.94405 "UDP glucuronosyltransferase 2 family, polypeptide B5" Ugt2b5
 1419623_at 0.95956 "protease, serine, 21" Prss21
 1419624_a_at 1.06255 RIKEN cDNA 1700010I14 gene 1700010I14Rik
 1419625_at 1.03462 heat shock protein 1-like Hspa11
 1419626_at 1.06358 similar to ADAM 25 precursor (A disintegrin and metalloproteinase domain 25) (Testase 2) LOC382006
 1419627_s_at 0.91904 "C-type lectin domain family 4, member n" Clec4n
 1419628_at 1.01791 C. elegans ceh-10 homeo domain containing homolog Chx10
 1419629_at 1.06439 mesoderm posterior 2 Mesp2
 1419630_a_at 1.11923 tripartite motif protein 11 Trim11
 1419631_at 1.02710 Wiskott-Aldrich syndrome homolog (human) Was
 1419632_at 0.86415 tectorin alpha Tecta
 1419633_at 0.97724 Unc4.1 homeobox (C. elegans) Uncx4.1
 1419634_a_at 1.00561 growth hormone releasing hormone Ghrh
 1419635_at 0.83912 RIKEN cDNA 4833420G17 gene 4833420G17Rik
 1419636_at 1.06416 RIKEN cDNA 4833420G17 gene 4833420G17Rik
 1419637_s_at 0.96587 RIKEN cDNA 4833420G17 gene 4833420G17Rik
 1419638_at 0.80671 ephrin B2 Efnb2
 1419639_at 0.83160 ephrin B2 Efnb2
 1419640_at 1.04011 purine rich element binding protein B Purb
 1419641_at 1.13962 purine rich element binding protein B Purb
 1419642_at 1.11953 purine rich element binding protein B Purb
 1419643_s_at 1.11481 RIKEN cDNA 2310057J18 gene /// RIKEN cDNA 2310065H11 gene 2310057J18Rik /// 2310065H11Rik
 1419644_at 1.03914 "cleavage stimulation factor, 3' pre-RNA subunit 2" Cstf2
 1419645_at 0.99290 "cleavage stimulation factor, 3' pre-RNA subunit 2" Cstf2
 1419646_a_at 0.86459 myelin basic protein Mbp
 1419647_a_at 1.50754 immediate early response 3 Ier3
 1419648_at 1.03201 myosin IC Myo1c
 1419649_s_at 1.04336 myosin IC Myo1c
 1419650_at 1.01112 zinc finger RNA binding protein Zfr
 1419651_at 1.02488 RIKEN cDNA 2610200G18 gene 2610200G18Rik
 1419652_s_at 1.01018 RIKEN cDNA 2610200G18 gene 2610200G18Rik
 1419653_a_at 0.96857 DEAD (Asp-Glu-Ala-Asp) box polypeptide 5 Ddx5
 1419654_at 0.88204 "transducin-like enhancer of split 3, homolog of Drosophila E(spl)" Tle3
 1419655_at 0.85765 "transducin-like enhancer of split 3, homolog of Drosophila E(spl)" Tle3

1419656_at 0.71948 "solute carrier family 25, member 36" Slc25a36
1419657_a_at 0.91411 "solute carrier family 25, member 36" Slc25a36
1419658_at 1.09012 RIKEN cDNA C920025E04 gene C920025E04Rik
1419659_s_at 0.95516 cysteine-rich hydrophobic domain 2 Chic2
1419660_at 0.93906 RIKEN cDNA 1600012F09 gene 1600012F09Rik
1419661_at 0.96760 RIKEN cDNA 1600012F09 gene /// hypothetical LOC624446
1600012F09Rik /// LOC624446
1419662_at 0.69936 osteoglycin Ogn
1419663_at 0.68411 osteoglycin Ogn
1419664_at 0.93444 serine racemase Srr
1419665_a_at 1.37603 nuclear protein 1 Nupr1
1419666_x_at 1.32508 nuclear protein 1 Nupr1
1419667_at 0.75772 "sarcoglycan, beta (dystrophin-associated glycoprotein)" Sgcb
1419668_at 0.77042 "sarcoglycan, beta (dystrophin-associated glycoprotein)" Sgcb
1419669_at 0.79882 proteinase 3 Prtn3
1419670_at 1.09996 formiminotransferase cyclodeaminase Ftcd
1419671_a_at 0.86187 interleukin 17 receptor C Il17rc
1419672_at 0.97096 "sparc/osteonectin, cwcv and kazal-like domains proteoglycan 1" Spock1
1419673_at 1.11908 "sparc/osteonectin, cwcv and kazal-like domains proteoglycan 1 /// similar to
sparc/osteonectin, cwcv and kazal-like domains proteoglycan 1" Spock1 /// LOC671214
1419674_a_at 0.99226 dipeptidase 1 (renal) Dpep1
1419675_at 0.79456 "nerve growth factor, beta" Ngfb
1419676_at 1.10638 myxovirus (influenza virus) resistance 2 Mx2
1419677_at 0.92009 mannan-binding lectin serine peptidase 1 Masp1
1419678_at 0.95110 large tumor suppressor 2 Lats2
1419679_at 1.10990 large tumor suppressor 2 Lats2
1419680_a_at 1.07060 elaC homolog 2 (E. coli) Elac2
1419681_a_at 0.99805 prokineticin 2 Prok2
1419682_a_at 1.20231 Trp53 regulating kinase /// RIKEN cDNA 2810408M09 gene Trp53rk ///
2810408M09Rik
1419683_at 1.18019 Trp53 regulating kinase Trp53rk
1419684_at 1.25065 chemokine (C-C motif) ligand 8 Ccl8
1419685_at 0.98857 UPF1 regulator of nonsense transcripts homolog (yeast) Upf1
1419686_at 1.02039 testis specific gene A14 Tsga14
1419687_at 0.91822 RIKEN cDNA D930010J01 gene D930010J01Rik
1419688_at 0.75811 glypican 6 Gpc6
1419689_at 0.93952 glypican 6 Gpc6
1419690_at 0.97612 RIKEN cDNA 2610002M06 gene 2610002M06Rik
1419691_at 1.05188 cathelicidin antimicrobial peptide Camp
1419692_a_at 1.16251 leukotriene C4 synthase Ltc4s
1419693_at 1.00867 collectin sub-family member 12 Colec12
1419694_at 0.92441 "ST8 alpha-N-acetyl-neuraminide alpha-2,8-sialyltransferase 1" St8sia1
1419695_at 0.90807 "ST8 alpha-N-acetyl-neuraminide alpha-2,8-sialyltransferase 1" St8sia1

1419696_at 0.97559 CD4 antigen Cd4
1419697_at 2.56569 chemokine (C-X-C motif) ligand 11 Cxcl11
1419698_at 1.35578 chemokine (C-X-C motif) ligand 11 Cxcl11
1419699_at 1.01413 "secretoglobin, family 3A, member 1" Scgb3a1
1419700_a_at 1.01971 prominin 1 Prom1
1419701_s_at 1.09487 ankyrin repeat and sterile alpha motif domain containing 3 Anks3
1419702_at 1.14406 "TATA box binding protein (Tbp)-associated factor, RNA polymerase I, A" Taf1a
1419703_at 1.12225 "procollagen, type V, alpha 3" Col5a3
1419704_at 1.07089 "cytochrome P450, family 3, subfamily a, polypeptide 41" Cyp3a41
1419705_at 0.79570 "carbonic anhydrase 5b, mitochondrial" Car5b
1419706_a_at 0.69399 A kinase (PRKA) anchor protein (gravin) 12 Akap12
1419707_at 0.90315 keratin associated protein 14 Krtap14
1419708_at 0.95111 wingless-related MMTV integration site 6 Wnt6
1419709_at 0.93228 stefin A3 Stfa3
1419710_at 0.91664 neurexophilin 3 Nxph3
1419711_at 0.99655 CD7 antigen Cd7
1419712_at 0.94614 "interleukin 3 receptor, alpha chain" Il3ra
1419713_at 0.90154 "ATPase, Class I, type 8B, member 3" Atp8b3
1419714_at 1.14861 CD274 antigen Cd274
1419715_at 1.02623 RIKEN cDNA 1700029F12 gene 1700029F12Rik
1419716_a_at 1.04128 "POU domain, class 2, transcription factor 1" Pou2f1
1419717_at 0.99344 "sema domain, immunoglobulin domain (Ig), short basic domain, secreted, (semaphorin) 3E" Sema3e
1419718_at 0.98360 RIKEN cDNA 4921530L21 gene 4921530L21Rik
1419719_at 1.04581 "gamma-aminobutyric acid (GABA-A) receptor, subunit beta 1 /// RIKEN cDNA B230208N19 gene" Gabrb1 /// B230208N19Rik
1419720_at 0.87688 glycoprotein 9 (platelet) Gp9
1419721_at 0.89210 G protein-coupled receptor 109A Gpr109a
1419722_at 0.96102 kallikrein 8 Klk8
1419723_at 0.94384 "opsin 1 (cone pigments), medium-wave-sensitive (color blindness, deutan)" Opn1mw
1419724_at 0.99782 ectodysplasin-A receptor /// similar to Tumor necrosis factor receptor superfamily member EDAR precursor (Anhidrotic ectodysplasin receptor 1) (Ectodysplasin-A receptor) (Ectodermal dysplasia receptor) (Downless) Edar /// LOC669001
1419725_at 0.86571 "solute carrier family 26, member 4" Slc26a4
1419726_at 1.04702 a disintegrin and metallopeptidase domain 18 Adam18
1419727_at 0.95086 decay accelerating factor 2 Daf2
1419728_at 17.96972 chemokine (C-X-C motif) ligand 5 Cxcl5
1419729_at 0.92451 testis expressed gene 11 Tex11
1419730_at 1.27952 RIKEN cDNA 4631427C17 gene 4631427C17Rik
1419731_at 1.09461 "cytochrome P450, family 2, subfamily b, polypeptide 19" Cyp2b19
1419732_at 0.94276 reticulon 4 receptor Rtn4r

1419733_at	0.94612	GLI pathogenesis-related 1 like 2	Glipr112
1419734_at	1.27109	"actin, beta, cytoplasmic"	Actb
1419735_at	1.17759	casein kappa	Csn3
1419736_a_at	1.02736	"eukaryotic translation initiation factor 1A, Y-linked"	Eif1ay
1419737_a_at	1.22801	lactate dehydrogenase A	Ldha
1419738_a_at	0.87767	"tropomyosin 2, beta"	Tpm2
1419739_at	0.84181	"tropomyosin 2, beta"	Tpm2
1419740_at	0.94269	"phosphodiesterase 6B, cGMP, rod receptor, beta polypeptide"	Pde6b
1419741_at	1.06494	suppressor of Ty 16 homolog (<i>S. cerevisiae</i>)	Supt16h
1419742_at	1.03071	RIKEN cDNA 1700037H04 gene	1700037H04Rik
1419743_s_at	0.98727	coactivator-associated arginine methyltransferase 1	Carm1
1419744_at	0.95985	"histocompatibility 2, class II, locus Mb2"	H2-DMb2
1419745_at	0.99590	Rho GTPase activating protein 23	Arhgap23
1419746_at	1.11729	Rho GTPase activating protein 23	Arhgap23
1419747_at	1.06167	asialoglycoprotein receptor 2	Asgr2
1419748_at	0.94544	"ATP-binding cassette, sub-family D (ALD), member 2"	Abcd2
1419749_at	1.01441	DNA methyltransferase 2	Dnmt2
1419750_at	1.16193	DNA methyltransferase 2	Dnmt2
1419751_x_at	0.94285	cDNA sequence AB056442	AB056442
1419752_at	1.03136	"nuclear transcription factor, X-box binding 1"	Nfx1
1419753_at	0.98284	"nuclear transcription factor, X-box binding 1"	Nfx1
1419754_at	1.25580	myosin Va	Myo5a
1419755_at	1.00050	antigen p97 (melanoma associated) identified by monoclonal antibodies 133.2 and 96.5	Mfi2
1419756_at	1.04356	"diacylglycerol kinase, gamma"	Dgkg
1419757_at	0.92134	"phosphatidylinositol transfer protein, membrane-associated 2"	Pitpmn2
1419758_at	1.01937	"ATP-binding cassette, sub-family B (MDR/TAP), member 1A"	Abcb1a
1419759_at	1.05133	"ATP-binding cassette, sub-family B (MDR/TAP), member 1A"	Abcb1a
1419760_a_at	0.99591	open reading frame 5	ORF5
1419761_a_at	0.84408	"GA repeat binding protein, beta 1"	Gabpb1
1419762_at	1.73338	ubiquitin D	Ubd
1419763_at	0.90075	RIKEN cDNA 2610020O08 gene	2610020O08Rik
1419764_at	1.02081	chitinase 3-like 3	Chi3l3
1419765_at	0.96260	cullin 2	Cul2
1419766_at	0.98792	SNF1-like kinase	Snf1lk
1419767_at	1.23128	"peptidyl arginine deiminase, type III"	Padi3
1419768_at	0.96434	CD22 antigen	Cd22
1419769_at	0.90422	CD22 antigen	Cd22
1419770_at	0.97220	---	---
1419771_at	1.02149	"PREDICTED: Mus musculus similar to dachsous 2 isoform 2 (LOC627094), mRNA"	---
1419772_at	1.04075	Myeloid/lymphoid or mixed lineage-leukemia translocation to 10 homolog (<i>Drosophila</i>)	Mllt10

1419773_at	0.88437	RIKEN cDNA 2010301N04 gene	2010301N04Rik
1419774_at	0.95070	"DNA segment, Chr 2, ERATO Doi 239, expressed"	D2Ertd239e
1419775_at	0.86830	---	---
1419776_at	1.06304	---	---
1419777_at	0.91749	Serine/threonine kinase 25 (yeast)	Stk25
1419778_at	1.04186	---	---
1419779_at	1.07262	---	---
1419780_at	0.92369	---	---
1419781_at	0.87161	RIKEN cDNA D630004N19 gene	D630004N19Rik
1419782_at	0.87136	expressed sequence AA517864	AA517864
1419783_at	0.94503	---	---
1419784_x_at	0.90961	---	---
1419785_at	0.92911	expressed sequence AA516738	AA516738
1419786_at	1.07447	latent transforming growth factor beta binding protein 1	Ltbp1
1419787_a_at	0.89633	zinc finger protein 628 Zfp628	
1419788_at	0.87665	Transcribed locus	---
1419789_at	0.92141	"Centaurin, gamma 1"	Centg1
1419790_at	0.94800	---	---
1419791_at	0.93771	zinc fingers and homeoboxes 3	Zhx3
1419792_at	1.04450	M-phase phosphoprotein 1	Mphosph1
1419793_at	1.08255	"DNA segment, Chr 5, ERATO Doi 615, expressed"	D5Ertd615e
1419794_at	0.90845	Transcribed locus	---
1419795_at	0.93933	expressed sequence AA517631	AA517631
1419796_at	1.05776	---	---
1419797_at	1.09253	expressed sequence AA672641	AA672641
1419798_at	1.17728	RIKEN cDNA 2610019E17 gene	2610019E17Rik
1419799_at	1.03669	"Proteasome (prosome, macropain) 26S subunit, non-ATPase, 1"	Psmd1
1419800_at	1.08248	Transcribed locus	---
1419801_x_at	1.03028	Transcribed locus	---
1419802_at	0.97584	coiled-coil domain containing 12	Ccdc12
1419803_s_at	1.03111	coiled-coil domain containing 12	Ccdc12
1419804_at	1.07112	RIKEN cDNA 1700051A21 gene	1700051A21Rik
1419805_s_at	0.96752	geranylgeranyl diphosphate synthase 1	Ggps1
1419806_at	0.92992	high density lipoprotein (HDL) binding protein	Hdlbp
1419807_at	0.91922	"DNA segment, Chr 4, ERATO Doi 335, expressed"	D4Ertd335e
1419808_at	1.02632	component of oligomeric golgi complex 4	Cog4
1419809_s_at	0.94442	component of oligomeric golgi complex 4	Cog4
1419810_x_at	0.85933	Rho GTPase activating protein 9	Arhgap9
1419811_at	0.76345	"DNA segment, Chr 16, Wayne State University 65, expressed"	D16Wsu65e
1419812_s_at	1.06271	coiled-coil domain containing 56	Ccdc56
1419813_at	0.93069	Transcribed locus	---
1419814_s_at	0.76958	S100 calcium binding protein A1	S100a1
1419815_at	0.94454	methyltransferase 11 domain containing 1	Mett11d1

1419816_s_at 0.88847 ERBB receptor feedback inhibitor 1 Errf1
1419817_s_at 1.16231 "DNA segment, Chr 1, ERATO Doi 161, expressed" D1Ert161e
1419818_x_at 0.98427 "DNA segment, Chr 1, ERATO Doi 161, expressed" D1Ert161e
1419819_s_at 1.08114 SEC63-like (*S. cerevisiae*) Sec63
1419820_at 0.50608 polycystic kidney and hepatic disease 1 Pkhd1
1419821_s_at 1.27236 "isocitrate dehydrogenase 1 (NADP+), soluble" Idh1
1419822_at 0.92388 "Eukaryotic translation initiation factor 3, subunit 6" Eif3s6
1419823_s_at 1.06531 kinase suppressor of ras 1 Ksr1
1419824_a_at 1.11117 RIKEN cDNA A230062G08 gene A230062G08Rik
1419825_at 0.98657 Neurexin III Nrxn3
1419826_at 0.90187 kinesin family member 17 Kif17
1419827_s_at 0.94979 kinesin family member 17 Kif17
1419828_at 1.04896 ubiquitin specific peptidase 31 Usp31
1419829_a_at 0.97709 Growth factor receptor bound protein 2-associated protein 2 Gab2
1419830_at 0.84745 --- ---
1419831_at 0.92427 expressed sequence AA416453 AA416453
1419832_s_at 0.92168 "acid phosphatase, prostate" AcpP
1419833_s_at 1.19455 "centaurin, delta 3" Centd3
1419834_x_at 1.03064 MAP/microtubule affinity-regulating kinase 1 Mark1
1419835_s_at 0.97366 plectin 1 Plec1
1419836_at 0.95401 --- ---
1419837_at 1.02771 triggering receptor expressed on myeloid cells-like 2 Trem12
1419838_s_at 0.95988 polo-like kinase 4 (*Drosophila*) Plk4
1419839_x_at 1.02520 PRP19/PSO4 pre-mRNA processing factor 19 homolog (*S. cerevisiae*)
Prpf19
1419840_at 1.12688 keratin 72 Krt72
1419842_at 0.94771 Transcribed locus ---
1419843_at 0.89313 Calreticulin 4 Calr4
1419844_a_at 1.03236 hypothetical protein LOC545086 LOC545086
1419845_at 0.97893 "distal-less homeobox 1, antisense" Dlx1as
1419846_at 0.98044 "TAF15 RNA polymerase II, TATA box binding protein (TBP)-associated
factor" Taf15
1419847_at 0.88084 Protein serine kinase H1 Pskh1
1419848_x_at 1.12954 toll-like receptor 7 Tlr7
1419849_at 1.09472 Ankyrin repeat domain 44 Ankrd44
1419851_at 1.03406 --- ---
1419853_a_at 0.95885 "purinergic receptor P2X, ligand-gated ion channel, 7" P2rx7
1419857_at 0.97737 Transcribed locus ---
1419858_at 0.82994 Interferon activated gene 203 Ifi203
1419859_at 0.94879 Transcribed locus ---
1419861_at 0.95617 --- ---
1419863_at 0.90808 Tropomodulin 1 Tmod1
1419864_x_at 0.94649 Transcribed locus ---

1419866_s_at 1.00764 ataxin 2 Atxn2
1419867_a_at 1.13072 ankyrin repeat and KH domain containing 1 Ankhd1
1419869_s_at 0.92407 high density lipoprotein (HDL) binding protein Hdlbp
1419871_at 1.07784 "Protein phosphatase 2 (formerly 2A), regulatory subunit A (PR 65), beta isoform" Ppp2r1b
1419872_at 0.65649 colony stimulating factor 1 receptor Csf1r
1419873_s_at 0.79954 colony stimulating factor 1 receptor Csf1r
1419874_x_at 1.01727 zinc finger and BTB domain containing 16 Zbtb16
1419875_at 1.07874 Transcribed locus ---
1419876_at 0.98605 RIKEN cDNA 2810449G22 gene 2810449G22Rik
1419877_x_at 0.91470 Transcribed locus ---
1419879_s_at 1.06142 tripartite motif protein 25 Trim25
1419880_x_at 0.99290 Transcribed locus ---
1419881_x_at 0.98277 --- ---
1419882_at 0.88833 "6-phosphofructo-2-kinase/fructose-2,6-biphosphatase 3" Pfkfb3
1419883_s_at 1.08259 "ATPase, H+ transporting, lysosomal V1 subunit B2" Atp6v1b2
1419884_at 1.07242 Ethanolamine kinase 1 Etnk1
1419885_at 0.93465 "DNA segment, Chr X, ERATO Doi 223, expressed" DXErtd223e
1419886_at 1.58098 "DNA segment, Chr X, ERATO Doi 223, expressed" DXErtd223e
1419887_at 0.97460 NudC domain containing 1 Nudcd1
1419888_at 1.22880 RIKEN cDNA 1110033M05 gene 1110033M05Rik
1419889_at 1.11670 RIKEN cDNA 1110033M05 gene 1110033M05Rik
1419890_at 1.04110 --- ---
1419891_s_at 0.86400 expressed sequence C77545 C77545
1419892_at 0.92392 RIKEN cDNA 1110021J02 gene 1110021J02Rik
1419893_at 0.96729 RIKEN cDNA D930048N14 gene D930048N14Rik
1419894_at 0.96189 "Similar to proteasome (prosome, macropain) subunit, beta type 7" MGC107702
1419895_at 1.11041 expressed sequence AA536748 AA536748
1419896_at 1.10181 expressed sequence BB001228 BB001228
1419897_at 1.12614 expressed sequence BB001228 BB001228
1419898_s_at 1.20280 zinc finger CCCH type containing 7 A Zc3h7a
1419899_at 0.94831 Zinc finger CCCH type containing 7 A Zc3h7a
1419900_at 0.95686 --- ---
1419901_at 0.93916 "NOD-derived CD11c +ve dendritic cells cDNA, RIKEN full-length enriched library, clone:F630006H17 product:hypothetical protein, full insert sequence" ---
1419902_at 0.99335 "NOD-derived CD11c +ve dendritic cells cDNA, RIKEN full-length enriched library, clone:F630006H17 product:hypothetical protein, full insert sequence" ---
1419903_at 0.94488 "Adult male corpora quadrigemina cDNA, RIKEN full-length enriched library, clone:B230215D24 product:unclassifiable, full insert sequence" ---
1419904_at 0.89313 "Adult male corpora quadrigemina cDNA, RIKEN full-length enriched library, clone:B230215D24 product:unclassifiable, full insert sequence" ---
1419905_s_at 1.01722 hydroxyprostaglandin dehydrogenase 15 (NAD) Hpgd

1419906_at	0.99380	Hydroxyprostaglandin dehydrogenase 15 (NAD)	Hpgd
1419907_s_at	1.13910	Fc receptor-like A	Fcrla
1419908_at	0.99767	Fc receptor-like A	Fcrla
1419909_at	1.08367	M-phase phosphoprotein 9	Mphosph9
1419910_at	0.74871	M-phase phosphoprotein 9	Mphosph9
1419911_at	0.90950	"Coronin, actin binding protein 1C"	Coro1c
1419912_s_at	1.01554	Serine/threonine kinase receptor associated protein	Strap
1419913_at	1.04027	Serine/threonine kinase receptor associated protein	Strap
1419914_s_at	1.05992	"DNA segment, Chr 10, ERATO Doi 438, expressed"	D10Ertd438e
1419915_at	0.72123	"DNA segment, Chr 10, ERATO Doi 438, expressed"	D10Ertd438e
1419916_at	0.87469	Ring finger protein 20	Rnf20
1419917_s_at	0.93150	transmembrane emp24 protein transport domain containing 7	Tmed7
1419918_at	0.81393	transmembrane emp24 protein transport domain containing 7	Tmed7
1419919_at	0.88010	RIKEN cDNA 4833422M21 gene	4833422M21Rik
1419920_s_at	1.08178	ubiquitin specific peptidase 7	Usp7
1419921_s_at	0.91947	Ubiquitin specific peptidase 7	Usp7
1419922_s_at	0.73120	attractin like 1	Atrn1
1419923_at	0.89426	Similar to Zinc finger protein 124 (HZF-16)	LOC626391
1419924_at	0.90533	RIKEN cDNA A730024A03 gene	A730024A03Rik
1419925_s_at	0.93965	---	---
1419926_at	0.86511	Thioredoxin domain containing 10	Txndc10
1419927_s_at	1.04745	RAB interacting factor	Rabif
1419928_at	1.01694	leucine-rich repeats and immunoglobulin-like domains 1	Lrig1
1419929_at	1.00773	"DNA segment, Chr 15, ERATO Doi 55, expressed"	D15Ertd55e
1419930_at	0.92218	"DNA segment, Chr 15, ERATO Doi 55, expressed"	D15Ertd55e
1419931_at	0.97584	"ATP-binding cassette, sub-family B (MDR/TAP), member 7"	Abcb7
1419932_s_at	1.04278	---	---
1419933_at	1.02277	p53 and DNA damage regulated 1	Pdrg1
1419934_at	0.95869	---	---
1419935_s_at	0.97136	"casein kinase 2, alpha prime polypeptide"	Csnk2a2
1419936_at	0.84347	"Casein kinase II, alpha 2, polypeptide"	Csnk2a2
1419937_at	0.88588	expressed sequence AA536749	AA536749
1419938_s_at	0.88559	Rho guanine nucleotide exchange factor (GEF) 17	Arhgef17
1419939_at	0.90464	Rho guanine nucleotide exchange factor (GEF) 17	Arhgef17
1419940_at	1.19334	RIKEN cDNA C030018P15 gene	C030018P15Rik
1419941_at	0.90913	RIKEN cDNA C030018P15 gene	C030018P15Rik
1419942_at	1.06424	Sulfiredoxin 1 homolog (S. cerevisiae)	Srxn1
1419943_s_at	1.07897	cyclin B1	Ccnb1
1419944_at	0.84215	cyclin B1	Ccnb1
1419945_s_at	0.88444	"RAB2, member RAS oncogene family"	Rab2
1419946_s_at	0.85773	"RAB2, member RAS oncogene family"	Rab2
1419947_at	1.02052	"DNA segment, Chr 4, ERATO Doi 117, expressed"	D4Ertd117e
1419948_at	1.01839	"DNA segment, Chr 4, ERATO Doi 117, expressed"	D4Ertd117e

1419949_at	1.02770	Transmembrane protein 38B	Tmem38b
1419950_s_at	1.07563	transportin 3	Tnpo3
1419951_at	0.98756	"Lectin, mannose-binding, 1"	Lman1
1419952_at	0.88963	Syntaxin 5A	Stx5a
1419953_at	1.01077	expressed sequence C80678	C80678
1419954_s_at	0.92201	"zinc finger, AN1-type domain 3"	Zfand3
1419955_at	0.95422	"Zinc finger, AN1-type domain 3"	Zfand3
1419956_at	1.08028	expressed sequence C77144	C77144
1419957_at	0.94464	expressed sequence C77144	C77144
1419958_at	0.94165	RIKEN cDNA C330003B14 gene	C330003B14Rik
1419959_s_at	0.94977	RIKEN cDNA C330003B14 gene	C330003B14Rik
1419960_at	0.96647	RIKEN cDNA C330003B14 gene	C330003B14Rik
1419961_s_at	1.04358	RIKEN cDNA C330003B14 gene	C330003B14Rik
1419962_at	1.00224	Transcribed locus	---
1419963_at	0.97374	DEP domain containing 6	Depdc6
1419964_s_at	1.08251	hepatoma-derived growth factor	Hdgf
1419965_at	0.94936	expressed sequence C79445	C79445
1419966_at	1.03254	"tubulin, beta 2b"	Tubb2b
1419967_at	1.09949	SEH1-like (<i>S. cerevisiae</i>)	Seh1
1419968_at	0.93570	expressed sequence C77370	C77370
1419969_at	1.06658	expressed sequence C77370	C77370
1419970_at	0.81119	"solute carrier family 35, member A5"	Slc35a5
1419971_s_at	0.88651	"solute carrier family 35, member A5"	Slc35a5
1419972_at	0.99876	"Solute carrier family 35, member A5"	Slc35a5
1419973_at	0.94501	"3-oxoacyl-ACP synthase, mitochondrial"	Oxsm
1419974_at	0.96614	"sterol carrier protein 2, liver"	Scp2
1419975_at	0.94180	---	---
1419976_s_at	0.96925	"nuclear factor of activated T-cells, cytoplasmic, calcineurin-dependent 3"	Nfatc3
1419977_s_at	0.76183	RNA binding motif protein 35b	Rbm35b
1419978_s_at	1.02474	"DNA segment, Chr 10, ERATO Doi 610, expressed"	D10Ertd610e
1419979_s_at	0.88685	cAMP responsive element binding protein 3	Creb3
1419980_at	0.92148	glucosidase beta 2	Gba2
1419981_at	0.87590	RIO kinase 3 (yeast)	Riok3
1419982_s_at	1.00543	---	---
1419983_at	0.87522	zinc finger protein 644	Zfp644
1419984_s_at	0.87491	zinc finger protein 644	Zfp644
1419985_s_at	0.83916	coiled-coil domain containing 69	Ccdc69
1419986_at	1.01772	Coiled-coil domain containing 69	Ccdc69
1419987_at	0.88820	---	---
1419988_at	1.00999	mitogen activated protein kinase kinase kinase 7	Map3k7
1419989_at	0.93699	COMM domain containing 5	Commd5
1419990_at	1.12054	RIKEN cDNA A830082K12 gene	A830082K12Rik

1419991_at	0.81344	RAB3A interacting protein (rabin3)-like 1	Rab3il1
1419992_x_at	0.87556	RAB3A interacting protein (rabin3)-like 1	Rab3il1
1419993_at	0.88531	RAB3A interacting protein (rabin3)-like 1	Rab3il1
1419994_s_at	0.97380	"DNA segment, Chr 10, ERATO Doi 641, expressed"	D10Ertd641e
1419995_at	0.92973	"DNA segment, Chr 10, ERATO Doi 641, expressed"	D10Ertd641e
1419996_s_at	0.97840	aryl hydrocarbon receptor nuclear translocator	Arnt
1419997_at	0.96531	---	---
1419998_at	0.89880	Transcribed locus	---
1419999_at	1.03305	immunoglobulin (CD79A) binding protein 1	Igbp1
1420000_s_at	1.07629	immunoglobulin (CD79A) binding protein 1	Igbp1
1420001_at	0.98452	"DNA segment, Chr 7, ERATO Doi 1, expressed"	D7Ertd1e
1420002_at	1.09302	"DNA segment, Chr 7, ERATO Doi 1, expressed"	D7Ertd1e
1420003_at	1.03818	"NOL1/NOP2/Sun domain family, member 5"	Nsun5
1420004_s_at	0.97814	nuclear pore membrane protein 121	Pom121
1420005_s_at	0.93487	bone morphogenetic protein 15	Bmp15
1420006_at	0.89355	Bone morphogenetic protein 15	Bmp15
1420007_at	0.98292	---	---
1420008_s_at	0.92874	---	---
1420009_at	1.00927	"WW, C2 and coiled-coil domain containing 1"	Wwc1
1420010_at	0.82959	hypothetical LOC574418	LOC574418
1420011_s_at	0.85972	X-box binding protein 1	Xbp1
1420012_at	0.96467	X-box binding protein 1	Xbp1
1420013_s_at	1.60090	lanosterol synthase	Lss
1420014_at	1.09274	Lanosterol synthase	Lss
1420015_s_at	1.13869	palmitoyl-protein thioesterase 1	Ppt1
1420016_at	0.86117	Palmitoyl-protein thioesterase 1	Ppt1
1420017_at	0.97249	tetraspanin 8	Tspan8
1420018_s_at	0.94401	tetraspanin 8	Tspan8
1420019_at	0.90182	Tetraspanin 8	Tspan8
1420020_at	0.95185	suppressor of zeste 12 homolog (Drosophila)	Suz12
1420021_s_at	1.05767	suppressor of zeste 12 homolog (Drosophila)	Suz12
1420022_s_at	0.93531	Suppressor of zeste 12 homolog (Drosophila)	Suz12
1420023_at	1.01304	eukaryotic translation termination factor 1	Etf1
1420024_s_at	1.11670	eukaryotic translation termination factor 1	Etf1
1420025_s_at	1.15374	hypothetical protein LOC637741	LOC637741
1420026_at	0.88800	AP2 associated kinase 1	Aak1
1420027_at	0.93960	expressed sequence C80171	C80171
1420028_s_at	1.05444	minichromosome maintenance deficient 3 (S. cerevisiae) /// similar to DNA replication licensing factor MCM3 (DNA polymerase alpha holoenzyme-associated protein P1) (P1-MCM3)	Mcm3 /// LOC671904
1420029_at	0.85627	Minichromosome maintenance deficient 3 (S. cerevisiae)	Mcm3
1420030_at	1.01780	"DNA segment, Chr 11, ERATO Doi 730, expressed"	D11Ertd730e
1420031_at	1.02374	checkpoint kinase 1 homolog (S. pombe)	Chek1

1420032_at 0.98334 Checkpoint kinase 1 homolog (*S. pombe*) Chek1
1420033_s_at 1.01201 "protein phosphatase 2, regulatory subunit B, delta isoform" Ppp2r2d
1420034_at 0.92245 "Protein phosphatase 2, regulatory subunit B, delta isoform" Ppp2r2d
1420035_at 1.03288 Aquaporin 6 Aqp6
1420036_at 0.93068 Arginine-tRNA-protein transferase 1 Ate1
1420037_at 0.88662 "ATP synthase, H⁺ transporting, mitochondrial F1 complex, alpha subunit, isoform 1" Atp5a1
1420038_at 1.02208 "VATPase, H⁺ transporting, lysosomal V1 subunit E1" Atp6v1e1
1420039_s_at 1.03508 chromobox homolog 7 Cbx7
1420040_at 0.94295 RIKEN cDNA 5730472N09 gene 5730472N09Rik
1420041_at 1.06840 --- ---
1420042_at 1.00331 THO complex 1 Thoc1
1420043_s_at 0.87204 THO complex 1 Thoc1
1420044_at 0.92950 --- ---
1420045_at 1.04590 --- ---
1420046_s_at 0.89142 MAF1 homolog (*S. cerevisiae*) Maf1
1420047_at 1.04208 MAF1 homolog (*S. cerevisiae*) Maf1
1420048_at 1.04149 --- ---
1420049_at 0.89721 expressed sequence C78859 C78859
1420050_at 0.98030 Transcribed locus ---
1420051_at 1.08036 Transcribed locus ---
1420052_x_at 1.10820 "proteasome (prosome, macropain) subunit, beta type 1" Psmb1
1420053_at 0.94000 "Proteasome (prosome, macropain) subunit, beta type 1" Psmb1
1420054_s_at 1.01499 "solute carrier family 35, member C2" Slc35c2
1420055_at 0.85445 "solute carrier family 35, member C2" Slc35c2
1420056_s_at 1.36470 phosphatidylserine receptor Ptdsr
1420057_at 0.98872 Phosphatidylserine receptor Ptdsr
1420058_s_at 0.91643 RIKEN cDNA 2410166I05 gene 2410166I05Rik
1420059_at 0.97948 RIKEN cDNA 2410166I05 gene 2410166I05Rik
1420060_s_at 0.93387 expressed sequence AA675344 AA675344
1420061_s_at 1.06636 expressed sequence AA675344 AA675344
1420062_at 1.02867 Transcribed locus ---
1420063_at 1.06257 Transcribed locus ---
1420064_s_at 0.86892 transketolase-like 1 Tktl1
1420065_at 0.97449 Transketolase-like 1 Tktl1
1420066_s_at 0.96675 CD160 antigen Cd160
1420067_at 0.94636 CD160 antigen Cd160
1420068_at 1.12576 --- ---
1420069_at 0.95136 "PREDICTED: *Mus musculus* hypothetical protein LOC628408 (LOC628408), mRNA" ---
1420070_a_at 0.96217 "solute carrier family 9 (sodium/hydrogen exchanger), member 5" Slc9a5
1420071_at 0.94003 "serine (or cysteine) peptidase inhibitor, clade B, member 6e" Serpinb6e
1420072_s_at 1.07598 --- ---

1420073_s_at 1.00692 --- ---
1420074_at 0.95578 "DNA segment, Chr 8, ERATO Doi 738, expressed" D8Ertd738e
1420075_at 1.16418 "ATPase, class I, type 8B, member 1" Atp8b1
1420076_at 0.87385 "ATPase, class I, type 8B, member 1" Atp8b1
1420077_at 0.95669 expressed sequence AI452102 AI452102
1420078_at 0.98487 "Transcribed locus, moderately similar to XP_218017.3 PREDICTED: similar to putative pheromone receptor [Rattus norvegicus]" ---
1420079_at 1.23441 similar to putative pheromone receptor (Go-VN5) /// similar to putative pheromone receptor (Go-VN5) /// similar to putative pheromone receptor (Go-VN5) /// similar to putative pheromone receptor (Go-VN5) /// hypothetical LOC627111 /// similar to putative pheromone receptor (Go-VN5) /// similar to putative pheromone receptor (Go-VN5) /// similar to putative pheromone receptor (Go-VN5) /// similar to putative pheromone receptor (Go-VN5) /// similar to putative pheromone receptor (Go-VN5) /// similar to putative pheromone receptor (Go-VN5)
LOC195691 /// LOC224552 /// LOC328759 /// LOC433070 /// LOC627111 /// LOC627367 /// LOC627476 /// LOC637382 /// LOC665253 /// LOC665376 /// LOC670484
1420080_a_at 0.92902 glutamine fructose-6-phosphate transaminase 1 Gfpt1
1420081_s_at 1.06187 "DNA segment, Chr 2, ERATO Doi 750, expressed" D2Ertd750e
1420082_at 0.91452 "DNA segment, Chr 2, ERATO Doi 750, expressed" D2Ertd750e
1420083_at 0.81857 expressed sequence AU016916 AU016916
1420084_at 0.93452 Transcribed locus ---
1420085_at 0.89899 fibroblast growth factor 4 Fgf4
1420086_x_at 0.95402 fibroblast growth factor 4 Fgf4
1420087_at 1.00328 Frizzled homolog 3 (Drosophila) Fzd3
1420088_at 1.36582 "nuclear factor of kappa light chain gene enhancer in B-cells inhibitor, alpha" Nfkbia
1420089_at 0.99115 "Nuclear factor of kappa light chain gene enhancer in B-cells inhibitor, alpha" Nfkbia
1420090_at 1.05548 --- ---
1420091_s_at 1.01215 microrchidia 3 Morc3
1420092_at 0.98161 microrchidia 3 Morc3
1420093_s_at 0.92837 heterogeneous nuclear ribonucleoprotein D-like Hnrpdl
1420094_at 0.98235 Heterogeneous nuclear ribonucleoprotein D-like Hnrpdl
1420095_s_at 0.88865 zinc finger proliferation 1 Zipr1
1420096_at 0.82397 Zinc finger proliferation 1 Zipr1
1420097_at 0.94389 "DNA segment, Chr 13, ERATO Doi 787, expressed" D13Ertd787e
1420098_s_at 1.24434 "DNA segment, Chr 13, ERATO Doi 787, expressed" D13Ertd787e
1420099_at 0.91532 "DNA segment, Chr 13, ERATO Doi 787, expressed" D13Ertd787e
1420100_s_at 0.96834 "DNA segment, Chr 13, ERATO Doi 787, expressed" D13Ertd787e
1420101_at 0.93736 RIKEN cDNA 2610020H08 gene 2610020H08Rik
1420102_at 0.89175 --- ---
1420103_at 1.05707 --- ---
1420104_at 1.03525 --- ---
1420105_at 0.91454 "DNA segment, Chr 13, ERATO Doi 205, expressed" D13Ertd205e

1420106_at	0.94696	"DNA Segment, Chr 9, Mouse Genome Informatics 7"	D9Mgi7
1420107_at	0.89763	Bardet-Biedl syndrome 4 homolog (human) Bbs4	
1420108_at	1.14676	---	---
1420109_at	1.03025	"Phosphatidylinositol-4-phosphate 5-kinase, type II, alpha"	Pip5k2a
1420110_s_at	0.99616	zinc finger protein 334 Zfp334	
1420111_at	0.96621	Zinc finger protein 334 Zfp334	
1420112_at	1.13246	Phosphofurin acidic cluster sorting protein 1	Pacs1
1420113_s_at	1.27939	RIKEN cDNA 2410022L05 gene	2410022L05Rik
1420114_s_at	0.91348	RIKEN cDNA 2410022L05 gene	2410022L05Rik
1420115_at	0.81251	exosome component 8	Exosc8
1420116_s_at	0.97802	golgi phosphoprotein 3	Golph3
1420117_at	0.80711	Transcribed locus	---
1420118_s_at	1.12409	---	---
1420119_s_at	1.13395	PHD finger protein 3	Phf3
1420120_at	0.87971	expressed sequence AU020177	AU020177
1420121_at	1.03860	expressed sequence AA517858	AA517858
1420122_at	0.90200	expressed sequence AA517858	AA517858
1420123_at	0.90932	T-cell leukemia translocation altered gene	Tcta
1420124_s_at	0.91991	T-cell leukemia translocation altered gene	Tcta
1420125_at	1.05342	T-cell leukemia translocation altered gene	Tcta
1420126_at	0.99285	expressed sequence AA522020	AA522020
1420127_s_at	0.88915	cell cycle progression 1	Ccpg1
1420128_s_at	0.86560	"phosphatidylinositol glycan anchor biosynthesis, class B"	Pigb
1420129_s_at	1.04127	"DNA segment, Chr 10, Wayne State University 52, expressed"	D10Wsu52e
1420130_s_at	1.00693	"DNA segment, Chr 10, Wayne State University 52, expressed"	D10Wsu52e
1420131_s_at	1.00135	pituitary tumor-transforming 1 interacting protein	Pttg1ip
1420132_s_at	1.08212	Pituitary tumor-transforming 1 interacting protein	Pttg1ip
1420133_at	0.93123	hypothetical protein LOC433719	RP23-289L21.1
1420134_at	1.08511	Hypothetical protein LOC433719	RP23-289L21.1
1420135_at	0.90318	CDNA sequence BC021438	BC021438
1420136_a_at	1.66527	---	---
1420137_at	1.09727	---	---
1420138_at	1.07688	"solute carrier family 19 (sodium/hydrogen exchanger), member 1"	Slc19a1
1420139_s_at	0.99173	"KRR1, small subunit (SSU) processome component, homolog (yeast)"	
		Krr1	
1420140_at	1.02996	"bin3, bicoid-interacting 3, homolog (Drosophila)"	Bcdin3
1420141_at	1.03562	expressed sequence AA517023	AA517023
1420142_s_at	1.12368	proliferation-associated 2G4	Pa2g4
1420143_at	1.08273	membrane associated DNA binding protein	Mnab
1420144_x_at	0.95469	membrane associated DNA binding protein	Mnab
1420145_at	0.97359	Transcribed locus	---
1420146_at	1.01271	---	---
1420147_at	1.10036	expressed sequence AA407331	AA407331

1420148_at	1.08424	"solute carrier family 6 (neurotransmitter transporter, taurine), member 6"	
Slc6a6			
1420149_at	1.02003	Pericentrin (kendrin)	Pcnt
1420150_at	0.73751	splA/ryanodine receptor domain and SOCS box containing 1	Spsb1
1420151_at	0.94241	---	---
1420152_at	0.86139	---	---
1420153_at	1.01750	---	---
1420154_at	1.12135	---	---
1420155_at	1.02910	expressed sequence AA408954	AA408954
1420156_at	0.91531	expressed sequence AA408954	AA408954
1420157_s_at	1.01904	"ATP-binding cassette, sub-family F (GCN20), member 1"	Abcf1
1420158_s_at	0.97512	"ATP-binding cassette, sub-family F (GCN20), member 1"	Abcf1
1420159_at	1.32887	myosin IE	Myo1e
1420160_s_at	0.98064	myosin IE	Myo1e
1420161_at	1.46855	expressed sequence AA409749	AA409749
1420162_at	1.07640	expressed sequence AA409749	AA409749
1420163_at	0.98500	expressed sequence AA511261	AA511261
1420164_at	1.12157	"DNA segment, Chr 7, ERATO Doi 183, expressed"	D7Ertd183e
1420165_s_at	1.15450	"DnaJ (Hsp40) homolog, subfamily C, member 17"	Dnajc17
1420166_at	0.93785	"Deoxynucleotidyltransferase, terminal"	Dntt
1420167_at	1.05574	---	---
1420168_at	1.01222	expressed sequence AA517841	AA517841
1420169_at	1.08563	expressed sequence AA517841	AA517841
1420170_at	0.83862	"myosin, heavy polypeptide 9, non-muscle"	Myh9
1420171_s_at	0.91246	"myosin, heavy polypeptide 9, non-muscle"	Myh9
1420172_at	0.87110	"Myosin, heavy polypeptide 9, non-muscle"	Myh9
1420173_at	1.00623	"DNA segment, Chr 11, ERATO Doi 326, expressed"	D11Ertd326e
1420174_s_at	0.94169	---	---
1420175_at	0.81568	Tax1 (human T-cell leukemia virus type I) binding protein 1	Tax1bp1
1420176_x_at	0.99074	immunoglobulin lambda-like polypeptide 1	Igl11
1420177_at	0.97226	immunoglobulin lambda-like polypeptide 1	Igl11
1420178_at	1.13172	Transcribed locus	---
1420179_at	0.99492	dolichyl-di-phosphooligosaccharide-protein glycotransferase	Ddost
1420180_at	1.00116	---	---
1420181_at	0.94182	RIKEN cDNA 1700011M02 gene /// hypothetical protein LOC668339	1700011M02Rik /// LOC668339
1420182_x_at	0.85111	doublesex and mab-3 related transcription factor 3	Dmrt3
1420183_at	0.90491	Loricrin	Lor
1420184_at	0.97067	---	---
1420185_at	0.90819	"synovial sarcoma, X breakpoint 2 interacting protein"	Ssx2ip
1420186_at	0.93984	---	---
1420187_at	0.91264	expressed sequence C76628	C76628
1420188_at	1.03882	synaptotagmin VI	Syt6

1420189_at	0.89771	---	---	
1420190_at	1.03531	---	---	
1420191_s_at	1.18310	"DNA segment, Chr 16, Brigham & Women's Genetics 1494 expressed"		
D16Bwg1494e				
1420192_at	0.91652	"DNA segment, Chr 16, Brigham & Women's Genetics 1494 expressed"		
D16Bwg1494e				
1420193_at	1.02601	keratin 17	Krt17	
1420194_at	1.01047	---	---	
1420195_at	0.98818	---	---	
1420196_s_at	0.91282	"TBC1 domain family, member 14" Tbc1d14		
1420197_at	0.90791	Growth arrest and DNA-damage-inducible 45 beta	Gadd45b	
1420198_at	0.89024	---	---	
1420199_at	1.00080	"Solute carrier family 35, member A4" Slc35a4		
1420200_at	1.02890	---	---	
1420201_at	0.95321	---	---	
1420202_at	0.98571	---	---	
1420203_at	0.93582	---	---	
1420204_at	1.07885	---	---	
1420205_at	0.89013	---	---	
1420206_at	0.94963	---	---	
1420207_at	0.96003	---	---	
1420208_at	1.04141	---	---	
1420209_at	0.98768	Dihydrouridine synthase 4-like (<i>S. cerevisiae</i>)	Dus4l	
1420210_at	0.94665	Expressed sequence AV344025	AV344025	
1420211_at	1.03277	---	---	
1420212_at	0.92199	---	---	
1420213_x_at	0.94801	---	---	
1420214_at	0.99608	RIKEN cDNA 1810012K16 gene	1810012K16Rik	
1420215_x_at	0.97794	RIKEN cDNA 1810012K16 gene	1810012K16Rik	
1420216_at	0.86167	RIKEN cDNA 1810012K16 gene	1810012K16Rik	
1420217_x_at	0.78395	---	---	
1420218_at	0.91024	---	---	
1420219_at	0.93402	RIKEN cDNA 4930461P20 gene	4930461P20Rik	
1420220_x_at	0.98146	RIKEN cDNA 4930461P20 gene	4930461P20Rik	
1420221_at	0.87244	RIKEN cDNA 4930461P20 gene	4930461P20Rik	
1420222_at	1.04423	---	---	
1420223_at	0.86382	---	---	
1420224_at	0.93837	---	---	
1420225_at	0.90385	---	---	
1420226_x_at	1.00123	homeo box A4	Hoxa4	
1420227_at	1.01657	Transcribed locus	---	
1420228_at	0.93395	---	---	
1420229_at	0.99671	---	---	

1420230_at	1.08817	expressed sequence AA414993	AA414993
1420231_at	0.92222	---	---
1420232_at	0.90943	---	---
1420233_at	1.02457	---	---
1420234_at	0.96157	---	---
1420235_at	0.98319	"biphenyl hydrolase-like (serine hydrolase, breast epithelial mucin-associated antigen)"	Bphl
1420236_at	1.16319	Transcribed locus	---
1420237_at	1.06096	Transcribed locus	---
1420238_at	1.10002	---	---
1420239_x_at	0.89529	membrane-associated ring finger (C3HC4) 2	2-Mar
1420240_at	0.89307	---	---
1420241_at	0.95055	---	---
1420242_at	1.00550	---	---
1420243_at	0.90599	zinc finger homeobox 1a	Zfx1a
1420244_at	1.17678	---	---
1420245_x_at	1.11963	transgelin 2 /// similar to transgelin 2	Tagln2 /// LOC672466
1420246_at	1.14822	"DNA segment, Chr 1, ERATO Doi 161, expressed"	D1Ertd161e
1420247_at	0.93662	"galactosidase, beta 1-like"	Glb1l
1420248_at	1.00649	"tubulin, gamma 2"	Tubg2
1420249_s_at	1.09542	chemokine (C-C motif) ligand 6	Ccl6
1420250_at	0.87582	---	---
1420251_at	0.96476	---	---
1420252_at	1.09318	---	---
1420253_at	0.87218	"DNA segment, Chr 2, ERATO Doi 63, expressed"	D2Ertd63e
1420254_at	0.96493	---	---
1420255_at	1.06543	---	---
1420256_x_at	1.01662	Peripherin 1	Prph1
1420257_at	1.01457	RIKEN cDNA 1700073E17 gene	1700073E17Rik
1420258_at	0.96519	RIKEN cDNA 1700073E17 gene	1700073E17Rik
1420259_at	1.04027	Plakophilin 2	Pkp2
1420260_at	0.99743	PHD finger protein 7	Phf7
1420261_at	0.99926	presenilin 1	Psen1
1420262_at	0.89663	---	---
1420263_at	0.98605	---	---
1420264_at	1.06758	---	---
1420265_x_at	0.98681	component of oligomeric golgi complex 8	Cog8
1420266_at	0.98141	---	---
1420267_at	0.85386	Nucleophosmin 1	Npm1
1420268_x_at	0.93410	Nucleophosmin 1	Npm1
1420269_at	1.01320	Nucleophosmin 1	Npm1
1420270_at	0.97505	---	---
1420271_at	1.02383	"3 days neonate thymus cDNA, RIKEN full-length enriched library, clone:	

A630030O14 product:unclassifiable, full insert sequence" ---

1420272_at	0.84483	---	---	
1420273_x_at	0.84759	---	---	
1420274_at	1.01324	"transporter 1, ATP-binding cassette, sub-family B (MDR/TAP)"	Tap1	
1420275_at	1.08449	---	---	
1420276_x_at	0.89826	---	---	
1420277_at	1.12590	---	---	
1420278_at	0.90485	cDNA sequence BC048599	BC048599	
1420279_at	1.03753	CDNA sequence BC048599	BC048599	
1420280_x_at	1.02797	"procollagen, type IX, alpha 3"	Col9a3	
1420281_at	0.88037	---	---	
1420282_s_at	1.00624	"protease, serine, 29"	Prss29	
1420283_at	1.03008	---	---	
1420284_at	0.93983	HLA-B associated transcript 4	Bat4	
1420285_at	0.90045	---	---	
1420286_at	0.85547	Transcribed locus	---	
1420287_at	0.93748	Transcribed locus	---	
1420288_at	0.90800	expressed sequence T25656	T25656	
1420289_at	0.91653	expressed sequence T25656	T25656	
1420290_at	0.94318	Transcribed locus	---	
1420291_at	1.04469	---	---	
1420292_x_at	0.99919	---	---	
1420293_at	0.84774	acyltransferase like 2	Aytl2	
1420294_at	0.88760	Acytransferase like 2	Aytl2	
1420295_x_at	1.06168	chloride channel 5	Clcn5	
1420296_at	0.88080	chloride channel 5	Clcn5	
1420297_at	1.04971	---	---	
1420298_at	0.95666	---	---	
1420299_at	0.99250	"gamma-aminobutyric acid (GABA-A) receptor, subunit alpha 2"	Gabra2	
1420300_at	1.10506	---	---	
1420301_at	0.99989	expressed sequence AA414903	AA414903	
1420302_at	0.81649	expressed sequence AA414903	AA414903	
1420303_x_at	0.93743	"Transcribed locus, strongly similar to XP_537399.1 PREDICTED: similar to 40S ribosomal protein S4, X isoform [Canis familiaris]"	---	
1420304_x_at	0.98462	"Transcribed locus, strongly similar to XP_537399.1 PREDICTED: similar to 40S ribosomal protein S4, X isoform [Canis familiaris]"	---	
1420305_at	0.94941	Transcribed locus	---	
1420307_a_at	1.02041	"phosphatidylinositol transfer protein, beta"	Pitpnb	
1420310_at	1.45167	---	---	
1420311_s_at	0.92474	---	---	
1420312_s_at	1.06280	---	---	
1420313_x_at	1.05151	---	---	
1420318_at	0.99737	Similar to zinc finger protein 709	LOC626877	

1420321_at	1.09890	Radical S-adenosyl methionine domain containing 1	Rsad1
1420325_at	1.09913	"Crm, cramped-like (Drosophila)"	Cramp11
1420326_s_at	0.98791	"Crm, cramped-like (Drosophila)"	Cramp11
1420328_at	1.30438	RIKEN cDNA 2210417D09 gene	2210417D09Rik
1420329_at	0.89315	RIKEN cDNA 4930455C21 gene	4930455C21Rik
1420330_at	0.88639	"C-type lectin domain family 4, member e"	Clec4e
1420331_at	1.06359	"C-type lectin domain family 4, member e"	Clec4e
1420332_x_at	0.98623	small proline rich-like 7	Sprrl7
1420333_at	1.02819	thioredoxin domain containing 8	Txndc8
1420334_at	1.02789	"solute carrier family 12 (potassium/chloride transporters), member 8"	Slc12a8
1420335_at	0.99580	disrupted meiotic cDNA 1 homolog	Dmc1h
1420336_at	1.00757	ADAMTS-like 5	Adamts15
1420337_at	0.90533	gastrulation brain homeobox 2	Gbx2
1420338_at	1.04427	arachidonate 15-lipoxygenase	Alox15
1420339_at	0.98671	RIKEN cDNA 0610016J10 gene	0610016J10Rik
1420340_at	1.03687	RIKEN cDNA 4930413O22 gene	4930413O22Rik
1420341_at	0.93387	transmembrane and coiled-coil domains 5	Tmco5
1420342_at	2.50815	ganglioside-induced differentiation-associated-protein 10	Gdap10
1420343_at	0.98285	granzyme D	Gzmd
1420344_x_at	0.99664	granzyme D	Gzmd
1420345_at	0.89090	claudin 14	Cldn14
1420346_at	0.97092	ankyrin repeat and SOCS box-containing protein 12	Asb12
1420347_at	1.01876	"palate, lung, and nasal epithelium carcinoma associated"	Plunc
1420348_at	0.98061	LIM homeobox protein 5	Lhx5
1420349_at	0.86281	prostaglandin F receptor	Ptgfr
1420350_at	1.07795	small proline rich-like 2	Sprrl2
1420351_at	1.08180	"tumor necrosis factor receptor superfamily, member 4"	Tnfrsf4
1420352_at	1.04732	"protease, serine, 22"	Prss22
1420353_at	0.96656	lymphotoxin A	Lta
1420354_at	1.05631	cyclin M1	Cnm1
1420355_at	0.89985	"G protein-coupled receptor kinase 2, groucho gene related (Drosophila)"	Gprk21
1420356_at	1.00040	ninjurin 2	Ninj2
1420357_s_at	0.83469	X-linked lymphocyte-regulated 3A /// X-linked lymphocyte-regulated 3B /// hypothetical protein LOC574437	Xlr3a /// Xlr3b /// MGC76689
1420358_at	0.96731	keratin associated protein 13	Krtap13
1420359_at	1.03918	seminal vesicle antigen Sva	
1420360_at	1.04597	dickkopf homolog 1 (Xenopus laevis)	Dkk1
1420361_at	1.25263	"solute carrier family 11 (proton-coupled divalent metal ion transporters), member 1"	Slc11a1
1420362_a_at	0.99536	Bcl2-interacting killer	Bik
1420363_at	0.95889	Bcl2-interacting killer	Bik
1420364_at	0.97669	G protein-coupled receptor 87	Gpr87

1420365_a_at 1.04234 heterogeneous nuclear ribonucleoprotein A2/B1 Hnrpa2b1
1420366_at 0.94171 sperm equatorial segment protein 1 Spesp1
1420367_at 0.88797 density-regulated protein Denr
1420368_at 1.14003 density-regulated protein Denr
1420369_a_at 0.97767 casein beta Csn2
1420370_s_at 0.97146 casein beta Csn2
1420371_at 0.95951 "syntrophin, basic 2" Sntb2
1420372_at 1.10294 "syntrophin, basic 2" Sntb2
1420373_at 0.80352 forkhead box J2 Foxj2
1420374_at 0.71465 forkhead box J2 Foxj2
1420375_at 0.97456 kinesin family member 3A Kif3a
1420376_a_at 0.93936 "H3 histone, family 3B" H3f3b
1420377_at 0.86930 "ST8 alpha-N-acetyl-neuraminide alpha-2,8-sialyltransferase 2" St8sia2
1420378_at 1.20170 surfactant associated protein D Sftpd
1420379_at 0.86858 "solute carrier organic anion transporter family, member 1a1" Slco1a1
1420380_at 3.37928 chemokine (C-C motif) ligand 2 Ccl2
1420381_a_at 1.04070 ribosomal protein L31 Rpl31
1420382_at 0.90697 apolipoprotein B48 receptor Apob48r
1420383_a_at 1.08436 "procollagen, type IV, alpha 3 (Goodpasture antigen) binding protein"
Col4a3bp
1420384_at 0.98492 "procollagen, type IV, alpha 3 (Goodpasture antigen) binding protein"
Col4a3bp
1420385_at 0.95050 "guanine nucleotide binding protein, alpha 14" Gna14
1420386_at 1.00354 SEH1-like (S. cerevisiae) Seh1
1420387_at 1.07385 "Mpv17 transgene, kidney disease mutant" Mpv17
1420388_at 1.09072 "protease, serine, 12 neurotrypsin (motopsin)" Prss12
1420389_at 1.03723 paired box gene 3 Pax3
1420390_s_at 0.88978 zinc finger protein 354A Zfp354a
1420391_at 0.79757 par-3 (partitioning defective 3) homolog (C. elegans) Pard3
1420392_at 1.04617 ubiquitin-like 4B Ubl4b
1420393_at 1.08462 "nitric oxide synthase 2, inducible, macrophage" Nos2
1420394_s_at 1.07614 "glycoprotein 49 A /// leukocyte immunoglobulin-like receptor, subfamily B,
member 4" Gp49a /// Lilrb4
1420395_a_at 0.87540 kinesin family member 9 Kif9
1420396_at 0.94075 CD160 antigen Cd160
1420397_a_at 0.96605 "SPEN homolog, transcriptional regulator (Drosophila)" Spen
1420398_at 0.98838 regulator of G-protein signaling 18 Rgs18
1420399_at 0.84005 growth factor independent 1B Gfi1b
1420400_at 0.99386 seminal vesicle antigen-like 1 Sval1
1420401_a_at 1.03308 receptor (calcitonin) activity modifying protein 3 Ramp3
1420402_at 0.92088 "ATPase, Ca++ transporting, plasma membrane 2" Atp2b2
1420403_at 1.00652 "ATPase, Ca++ transporting, plasma membrane 2" Atp2b2
1420404_at 1.01450 CD86 antigen Cd86

1420405_at 0.95897 "solute carrier organic anion transporter family, member 1a4" Slco1a4
 1420406_at 0.90279 paternally expressed 12 Peg12
 1420407_at 0.99266 leukotriene B4 receptor 1 Ltb4r1
 1420408_a_at 0.88645 "ATP-binding cassette, sub-family C (CFTR/MRP), member 9" Abcc9
 1420409_at 0.95454 keratin 35 Krt35
 1420410_at 0.93761 "nuclear receptor subfamily 5, group A, member 2" Nr5a2
 1420411_a_at 1.15354 phosphatidylinositol 4-kinase type 2 beta Pi4k2b
 1420412_at 0.99105 "tumor necrosis factor (ligand) superfamily, member 10" Tnfsf10
 1420413_at 1.19021 "solute carrier family 7 (cationic amino acid transporter, y+ system), member 11" Slc7a11
 1420414_at 1.11022 homeo box A11 Hoxa11
 1420415_at 0.84210 cDNA sequence AY026312 AY026312
 1420416_at 0.63470 "sema domain, immunoglobulin domain (Ig), short basic domain, secreted, (semaphorin) 3A" Sema3a
 1420417_at 0.97239 "sema domain, immunoglobulin domain (Ig), short basic domain, secreted, (semaphorin) 3A" Sema3a
 1420418_at 1.12596 synaptotagmin II Syt2
 1420419_a_at 1.00557 otoferlin Otof
 1420420_at 1.00274 "hydroxyacid oxidase 1, liver" Hao1
 1420421_s_at 1.11944 killer cell lectin-like receptor subfamily B member 1B Klrb1b
 1420422_at 0.88968 protocadherin beta 21 Pcdhb21
 1420423_at 1.00076 "T-cell leukemia/lymphoma 1B, 4" Tcl1b4
 1420424_at 0.96318 RIKEN cDNA 1700054O13 gene 1700054O13Rik
 1420425_at 0.92673 "PR domain containing 1, with ZNF domain" Prdm1
 1420426_at 0.85849 myosin VIIb Myo7b
 1420427_a_at 0.82497 DEAH (Asp-Glu-Ala-His) box polypeptide 32 Dhx32
 1420428_at 0.79236 advanced glycosylation end product-specific receptor Ager
 1420429_at 0.91971 protocadherin beta 3 Pcdhb3
 1420430_a_at 0.94092 RIKEN cDNA 2510048L02 gene 2510048L02Rik
 1420431_at 0.95728 repetin Rptn
 1420432_at 1.01981 "taste receptor, type 1, member 1" Tas1r1
 1420433_at 0.96881 "TAF7-like RNA polymerase II, TATA box binding protein (TBP)-associated factor" Taf7l
 1420434_at 0.92093 claudin 16 Cldn16
 1420435_at 1.05983 RIKEN cDNA 1700011A15 gene 1700011A15Rik
 1420436_x_at 1.04850 testis expressed gene 21 Tex21
 1420437_at 1.19248 "indoleamine-pyrrole 2,3 dioxygenase" Indo
 1420438_at 1.07593 orosomucoid 2 Orm2
 1420439_at 0.99237 testis specific X-linked gene Tsx
 1420440_at 0.92880 pancreatic polypeptide Ppy
 1420441_at 1.03431 centromere protein C1 Cenpc1
 1420442_at 1.06770 "calcium channel, voltage-dependent, L type, alpha 1S subunit" Cacna1s
 1420443_at 0.87509 protocadherin beta 19 Pcdhb19

1420444_at 0.92996 "solute carrier family 22 (organic cation transporter), member 3" Slc22a3
 1420445_at 0.93910 "solute carrier family 16 (monocarboxylic acid transporters), member 8" Slc16a8
 1420446_at 0.95814 outer dense fiber of sperm tails 3 Odf3
 1420447_at 1.11325 "sulfotransferase family 1E, member 1" Sult1e1
 1420448_at 1.17667 reproductive homeobox 2 Rhox2
 1420449_at 1.03951 RIKEN cDNA A230048G03 gene A230048G03Rik
 1420450_at 0.94298 matrix metalloproteinase 10 Mmp10
 1420451_at 1.08313 "amiloride-sensitive cation channel 5, intestinal" Accn5
 1420452_at 0.94372 keratin associated protein 5-2 Krtap5-2
 1420453_at 1.03144 "crystallin, gamma S" Crygs
 1420454_at 1.26259 retinoic acid induced 1 Rai1
 1420455_at 1.11291 glial cells missing homolog 2 (Drosophila) Gcm2
 1420456_at 1.05953 glial cells missing homolog 2 (Drosophila) Gcm2
 1420457_at 1.06371 "dishevelled 3, dsh homolog (Drosophila)" Dvl3
 1420458_at 1.06365 tachykinin 4 Tac4
 1420459_at 1.05290 Down syndrome critical region homolog 6 (human) Dscr6
 1420460_a_at 0.90373 peroxisomal biogenesis factor 11b Pex11b
 1420461_at 0.84347 macrophage stimulating 1 receptor (c-met-related tyrosine kinase) Mst1r
 1420462_at 0.97978 interleukin 1 receptor accessory protein-like 2 Il1rapl2
 1420463_at 1.06527 cytokine-dependent hematopoietic cell linker Clnk
 1420464_s_at 1.00794 "paired-Ig-like receptor A1 /// paired-Ig-like receptor A11 /// paired-Ig-like
 receptor A2 /// paired-Ig-like receptor A3 /// paired-Ig-like receptor A4 /// paired-Ig-like receptor A6 ///
 leukocyte immunoglobulin-like receptor, subfamily B (with TM and ITIM domains), member 3 ///
 hypothetical LOC619608 /// similar to paired-Ig-like receptor A11 /// similar to paired-Ig-like receptor
 A11 /// similar to paired-Ig-like receptor A1" Pira1 /// Pira11 /// Pira2 /// Pira3 /// Pira4 /// Pira6 ///
 Lilrb3 /// LOC619608 /// LOC669449 /// LOC669458 /// LOC675749
 1420465_s_at 1.01288 major urinary protein 1 /// major urinary protein 2 Mup1 /// Mup2
 1420466_at 0.92473 salivary protein 2 Spt2
 1420467_at 1.01554 psoriasis susceptibility 1 candidate 2 (human) Psors1c2
 1420468_at 0.99694 ankyrin repeat and SOCS box-containing protein 17 Asb17
 1420469_at 0.97295 RIKEN cDNA 4933411G06 gene 4933411G06Rik
 1420470_at 1.05694 "sulfotransferase family, cytosolic, 1C, member 1" Sult1c1
 1420471_at 0.88311 hypocretin Hcrt
 1420472_at 1.01089 myotrophin Mtpn
 1420473_at 1.00011 myotrophin Mtpn
 1420474_at 1.06860 myotrophin Mtpn
 1420475_at 0.97677 myotrophin Mtpn
 1420476_a_at 1.07790 nucleosome assembly protein 1-like 1 Nap111
 1420477_at 1.14491 nucleosome assembly protein 1-like 1 Nap111
 1420478_at 1.02867 nucleosome assembly protein 1-like 1 Nap111
 1420479_a_at 1.04303 nucleosome assembly protein 1-like 1 Nap111
 1420480_at 0.78010 cyclin M3 Cnm3
 1420481_at 0.81343 cyclin M3 Cnm3

1420482_at	0.94063 cyclin M3	Cnm3
1420483_at	1.08422 cyclin M3	Cnm3
1420484_a_at	1.01275 vitronectin	Vtn
1420485_at	1.08574 nucleolar protein 7	Nol7
1420486_at	1.14809 nucleolar protein 7	Nol7
1420487_at	0.94462 nucleolar protein 7	Nol7
1420488_at	0.98406 mitochondrial ribosomal protein S14	Mrps14
1420489_at	1.00408 mitochondrial ribosomal protein S14	Mrps14
1420490_at	0.94569 kallikrein 1-related peptidase b16	Klk1b16
1420491_at	1.11603 "eukaryotic translation initiation factor 2, subunit 1 alpha"	Eif2s1
1420492_s_at	0.90651 submaxillary gland androgen regulated protein 1 /// submaxillary gland androgen regulated protein 3	Smr1 /// Smr3
1420493_a_at	1.24317 "phosphate cytidyltransferase 2, ethanolamine"	Pcyt2
1420494_x_at	0.94185 ubiquitin C	Ubc
1420495_a_at	1.03822 vacuolar protein sorting 26 homolog A (yeast)	Vps26a
1420496_at	0.96343 coagulation factor XII (Hageman factor) F12	
1420497_a_at	1.05058 CCAAT/enhancer binding protein zeta	Cebpz
1420498_a_at	0.91474 disabled homolog 2 (Drosophila) Dab2	
1420499_at	0.96353 GTP cyclohydrolase 1	Gch1
1420500_at	0.98699 "DnaJ (Hsp40) homolog, subfamily C, member 1"	Dnajc1
1420501_at	0.94136 "DnaJ (Hsp40) homolog, subfamily C, member 1"	Dnajc1
1420502_at	1.03779 spermidine/spermine N1-acetyl transferase 1	Sat1
1420503_at	1.00668 "solute carrier family 6 (neurotransmitter transporter), member 14"	Slc6a14
1420504_at	0.95744 "solute carrier family 6 (neurotransmitter transporter), member 14"	Slc6a14
1420505_a_at	1.14599 syntaxin binding protein 1	Stxbp1
1420506_a_at	1.02031 syntaxin binding protein 1	Stxbp1
1420507_a_at	0.83509 RIKEN cDNA 3110031B13 gene	3110031B13Rik
1420508_at	0.88627 "sema domain, immunoglobulin domain (Ig), short basic domain, secreted, (semaphorin) 3 F"	Sema3f
1420509_at	1.02833 serum response factor binding protein 1	Srfbp1
1420510_at	1.08121 --- ---	
1420511_at	1.08089 "retinal degeneration, slow (retinitis pigmentosa 7)"	Rds
1420512_at	1.01809 dickkopf homolog 2 (Xenopus laevis)	Dkk2
1420513_at	1.12326 EF-hand calcium binding domain 2	Efcab2
1420514_at	1.01609 Transmembrane protein 47	Tmem47
1420515_a_at	1.15206 peptidoglycan recognition protein 2	Pglyrp2
1420516_at	1.00651 nonagouti	a
1420517_at	0.86282 RIKEN cDNA 2310010I16 gene	2310010I16Rik
1420518_a_at	0.93263 "immunoglobulin superfamily, member 9"	Igsf9
1420519_a_at	0.99501 Era (G-protein)-like 1 (E. coli)	Eral1
1420520_x_at	0.98234 Era (G-protein)-like 1 (E. coli)	Eral1
1420521_at	0.99586 "papilin, proteoglycan-like sulfated glycoprotein"	Papln
1420522_at	0.91117 coiled-coil domain containing 50	Ccdc50

1420523_at 0.91411 coiled-coil domain containing 50 Ccdc50
 1420524_a_at 0.86036 mannan-binding lectin serine peptidase 2 Masp2
 1420525_a_at 1.09404 ornithine transcarbamylase Otc
 1420526_at 0.95796 "discoidin, CUB and LCCL domain containing 2" Dcbld2
 1420527_s_at 1.01160 t-complex protein 10a /// t-complex protein 10b /// t-complex protein 10c
 Tcp10a /// Tcp10b /// Tcp10c
 1420528_at 0.95789 G patch domain containing 2 Gpatc2
 1420529_at 0.98303 neuronal d4 domain family member Neud4
 1420530_at 0.93083 neuronal d4 domain family member Neud4
 1420531_at 0.97650 "hydroxy-delta-5-steroid dehydrogenase, 3 beta- and steroid delta-isomerase 5"
 Hsd3b5
 1420532_at 0.88542 "membrane associated guanylate kinase, WW and PDZ domain containing 2"
 Magi2
 1420533_at 0.92064 "guanylate cyclase 1, soluble, alpha 3" Gucy1a3
 1420534_at 0.93080 "guanylate cyclase 1, soluble, alpha 3" Gucy1a3
 1420535_a_at 0.99452 negative regulator of ubiquitin-like proteins 1 Nub1
 1420536_at 0.94807 "crystallin, beta B2" Crybb2
 1420537_at 0.90729 potassium channel tetramerisation domain containing 4 Kctd4
 1420538_at 0.90595 "G protein-coupled receptor, family C, group 5, member D" Gprc5d
 1420539_a_at 0.95258 chordin-like 2 Chrdl2
 1420540_a_at 1.00462 Ras-like without CAAX 1 Rit1
 1420541_at 1.04034 retinol dehydrogenase 16 Rdh16
 1420542_at 1.03169 open reading frame 28 ORF28
 1420543_at 0.89509 open reading frame 28 ORF28
 1420544_at 0.95578 germinal center expressed transcript 2 Gcet2
 1420545_a_at 0.96580 chimerin (chimaerin) 1 Chn1
 1420546_at 1.00823 tyrosine hydroxylase Th
 1420547_at 1.12306 galactosylceramidase Galc
 1420548_a_at 1.22697 RIKEN cDNA 2310008H09 gene 2310008H09Rik
 1420549_at 1.32318 guanylate nucleotide binding protein 1 Gbp1
 1420550_at 0.93221 RIKEN cDNA 1110055J05 gene 1110055J05Rik
 1420551_at 0.98826 RIKEN cDNA 2310039E09 gene 2310039E09Rik
 1420552_s_at 1.02032 testis specific gene A8 Tsga8
 1420553_x_at 0.98110 "serine (or cysteine) peptidase inhibitor, clade A, member 1a" Serpina1a
 1420554_a_at 0.99027 RAS-related C3 botulinum substrate 3 Rac3
 1420555_at 0.92350 aristaless 3 Alx3
 1420556_at 0.99735 oxytocin Oxt
 1420557_at 1.12246 Eph receptor A5 Epha5
 1420558_at 1.23025 "selectin, platelet" Selp
 1420559_a_at 0.94704 short stature homeobox 2 Shox2
 1420560_at 1.04707 "cholinergic receptor, nicotinic, epsilon polypeptide" Chrne
 1420561_at 0.93216 "transient receptor potential cation channel, subfamily C, member 7" Trpc7
 1420562_at 0.95114 secreted Ly6/Plaur domain containing 1 Slurp1

1420563_at	1.09899	"glutamate receptor, ionotropic, AMPA3 (alpha 3)"	Gria3
1420564_at	0.96918	insulin receptor-related receptor	Insr
1420565_at	0.88002	homeo box A1	Hoxa1
1420566_at	0.99810	---	---
1420567_at	0.95901	"protein kinase C, nu"	Prkc
1420568_at	0.85498	stimulated by retinoic acid gene 8	Stra8
1420569_at	0.96587	chondroadherin	Chad
1420570_x_at	0.96702	"T-cell leukemia/lymphoma 1B, 3"	Tcl1b3
1420571_at	1.11484	prolactin-like protein B	Prlpb
1420572_at	1.05171	"membrane-spanning 4-domains, subfamily A, member 3"	Ms4a3
1420573_at	0.95650	homeo box D1	Hoxd1
1420574_at	0.94084	RIKEN cDNA 1700065I17 gene	1700065I17Rik
1420575_at	0.92149	metallothionein 3	Mt3
1420576_at	0.94809	"endothelial differentiation, lysophosphatidic acid G-protein-coupled receptor 4"	Edg4
1420577_at	0.88022	activation-induced cytidine deaminase	Aicda
1420578_at	0.92773	opticin	Optc
1420579_s_at	0.95944	cystic fibrosis transmembrane conductance regulator homolog	Cftr
1420580_at	1.03029	RIKEN cDNA 4930429B21 gene	4930429B21Rik
1420581_at	0.96408	G protein-coupled receptor 143	Gpr143
1420582_at	1.02329	CD209e antigen	Cd209e
1420583_a_at	0.85468	RAR-related orphan receptor alpha	Rora
1420584_at	1.00969	"phospholipase A2, group IIC"	Pla2g2c
1420585_a_at	0.95763	nuclear RNA export factor 2	Nxf2
1420586_at	1.00501	RIKEN cDNA 2900054J07 gene	2900054J07Rik
1420587_at	1.02183	testis specific gene A13	Tsga13
1420588_at	0.93172	"serine protease inhibitor-like, with Kunitz and WAP domains 1 (eppin)"	Spinlw1
1420589_at	0.86213	hyaluronan synthase 3	Has3
1420590_at	0.92714	hyaluronan synthase1	Has1
1420591_at	1.02032	G protein-coupled receptor 84	Gpr84
1420592_a_at	1.11345	"acidic (leucine-rich) nuclear phosphoprotein 32 family, member E"	Anp32e
1420593_a_at	0.95881	TEA domain family member 3	Tead3
1420594_at	0.97578	BRCA1 associated RING domain 1	Bard1
1420595_at	0.73223	distal-less homeobox 4	Dlx4
1420596_at	0.93889	"calcium channel, voltage-dependent, gamma subunit 2"	Cacng2
1420597_a_at	0.96296	calcium binding protein 2	Cabp2
1420598_x_at	1.04175	"defensin related cryptdin, related sequence 2"	Defcr-rs2
1420599_at	1.00365	RIKEN cDNA 2300006N05 gene	/// hypothetical protein LOC670472
		hypothetical protein LOC670482	2300006N05Rik
		/// LOC670472	/// LOC670482
1420600_at	0.98704	thyroid peroxidase	Tpo
1420601_at	1.04326	glial cells missing homolog 1 (Drosophila)	Gcm1
1420602_a_at	0.90200	"extraembryonic, spermatogenesis, homeobox 1"	Esx1

1420603_s_at 0.91981 "retinoic acid early transcript 1, alpha /// retinoic acid early transcript beta ///
retinoic acid early transcript gamma /// retinoic acid early transcript delta /// retinoic acid early transcript
1E" Raet1a /// Raet1b /// Raet1c /// Raet1d /// Raet1e

1420604_at 0.96877 homeo box gene expressed in ES cells Hesx1

1420605_at 1.03446 metastasis associated gene 2 Mtag2

1420606_at 1.06390 neuropeptide FF-amide peptide precursor Npff

1420607_at 0.95015 RNA binding motif protein 18 Rbm18

1420608_at 0.92833 RNA binding motif protein 18 Rbm18

1420609_at 0.87759 membrane-associated ring finger (C3HC4) 7 7-Mar

1420610_at 0.83466 "protein kinase, cAMP dependent, catalytic, beta" Prkacb

1420611_at 0.89267 "protein kinase, cAMP dependent, catalytic, beta" Prkacb

1420612_s_at 1.13899 protein tyrosine phosphatase 4a2 Ptp4a2

1420613_at 1.19703 protein tyrosine phosphatase 4a2 Ptp4a2

1420614_at 0.84874 dynein light chain Tctex-type 3 Dynlt3

1420615_at 0.80211 "ash2 (absent, small, or homeotic)-like (Drosophila)" Ash2l

1420616_at 0.98416 "ash2 (absent, small, or homeotic)-like (Drosophila)" Ash2l

1420617_at 0.89481 cytoplasmic polyadenylation element binding protein 4 Cpeb4

1420618_at 0.77197 cytoplasmic polyadenylation element binding protein 4 Cpeb4

1420619_a_at 0.83789 amino-terminal enhancer of split Aes

1420620_a_at 0.89689 ring finger protein 13 Rnf13

1420621_a_at 0.81069 amyloid beta (A4) precursor protein App

1420622_a_at 0.97792 heat shock protein 8 /// similar to heat shock protein 8 /// heat shock protein 8
pseudogene /// similar to heat shock protein 8 Hspa8 /// LOC621284 /// LOC666904 /// LOC669493

1420623_x_at 0.97471 heat shock protein 8 /// similar to heat shock protein 8 /// heat shock protein 8
pseudogene Hspa8 /// LOC621284 /// LOC666904

1420624_a_at 1.03840 vesicle-associated membrane protein 8 Vamp8

1420625_at 0.98096 RIKEN cDNA 2410016F19 gene 2410016F19Rik

1420626_at 1.05873 RIKEN cDNA 2410016F19 gene 2410016F19Rik

1420627_a_at 0.97334 casein alpha s1 Csn1s1

1420628_at 1.14468 purine rich element binding protein A Pura

1420629_a_at 1.12709 "DnaJ (Hsp40) homolog, subfamily A, member 3" Dnaja3

1420630_at 0.86290 RIKEN cDNA 8430419L09 gene 8430419L09Rik

1420631_a_at 1.00894 bladder cancer associated protein homolog (human) Blcap

1420632_a_at 0.93732 Bernardinelli-Seip congenital lipodystrophy 2 homolog (human) Bslc2

1420633_a_at 0.93235 casein alpha s2-like A Csn1s2a

1420634_a_at 1.05670 MAD homolog 2 (Drosophila) Smad2

1420635_a_at 1.01707 "T-cell, immune regulator 1, ATPase, H+ transporting, lysosomal V0 protein
A3" Tcirg1

1420636_a_at 1.01117 dual specificity phosphatase 12 Dusp12

1420637_at 0.92076 phosphoribosyl pyrophosphate synthetase 2 Prps2

1420638_at 1.00582 phosphoribosyl pyrophosphate synthetase 2 Prps2

1420639_at 0.87687 junction-mediating and regulatory protein Jmy

1420640_at 0.96781 junction-mediating and regulatory protein Jmy

1420641_a_at 0.86802 sulfide quinone reductase-like (yeast) Sqrdl
1420642_a_at 0.88247 RIKEN cDNA 2010100O12 gene 2010100O12Rik
1420643_at 0.75393 lunatic fringe gene homolog (Drosophila) Lfng
1420644_a_at 1.12670 "Sec61, alpha subunit 2 (S. cerevisiae)" Sec61a2
1420645_at 1.01440 polycomb group ring finger 2 Pcgf2
1420646_at 1.12312 "DNA segment, Chr 10, ERATO Doi 438, expressed" D10Ert438e
1420647_a_at 0.91617 "keratin 8 /// similar to Keratin, type II cytoskeletal 8 (Cytokeratin 8) (Cytokeratin endo A) /// similar to Keratin, type II cytoskeletal 8 (Cytokeratin-8) (CK-8) (Keratin-8) (K8) (Cytokeratin endo A)" Krt8 /// LOC434261 /// LOC675884
1420648_at 1.13398 tripartite motif protein 12 Trim12
1420649_at 0.95328 AT motif binding factor 1 Atbf1
1420650_at 0.92736 AT motif binding factor 1 Atbf1
1420651_at 0.96877 arginine-tRNA-protein transferase 1 Ate1
1420652_at 0.95359 arginine-tRNA-protein transferase 1 Ate1
1420653_at 0.99570 "transforming growth factor, beta 1" Tgfb1
1420654_a_at 1.12658 "glucan (1,4-alpha-), branching enzyme 1" Gbe1
1420655_at 0.86215 patatin-like phospholipase domain containing 3 Pnpla3
1420656_at 0.80675 "ATP-binding cassette, sub-family G (WHITE), member 8" Abcg8
1420657_at 1.09789 "uncoupling protein 3 (mitochondrial, proton carrier)" Ucp3
1420658_at 0.99103 "uncoupling protein 3 (mitochondrial, proton carrier)" Ucp3
1420659_at 0.99404 SLAM family member 6 Slamf6
1420660_at 1.01480 leucine rich repeat containing 6 (testis) Lrrc6
1420661_a_at 0.87461 RIKEN cDNA 4933439F18 gene 4933439F18Rik
1420662_at 1.06622 RIKEN cDNA 4933439F18 gene 4933439F18Rik
1420663_at 1.06974 Zinc finger and BTB domain containing 7B Zbtb7b
1420664_s_at 1.12527 "protein C receptor, endothelial" Procr
1420665_at 0.93105 integrin beta 3 binding protein (beta3-endonexin) Itgb3bp
1420666_at 0.96766 "double C2, beta" Doc2b
1420667_at 0.99388 "double C2, beta" Doc2b
1420668_a_at 0.94301 "Yip1 domain family, member 2" Yipf2
1420669_at 0.94474 aryl hydrocarbon receptor nuclear translocator 2 Arnt2
1420670_at 1.12150 aryl hydrocarbon receptor nuclear translocator 2 Arnt2
1420671_x_at 0.96436 "membrane-spanning 4-domains, subfamily A, member 4C" Ms4a4c
1420672_at 0.98106 "potassium voltage-gated channel, Isk-related subfamily, member 1" Kcne1
1420673_a_at 0.89856 "acyl-Coenzyme A oxidase 2, branched chain" Acox2
1420674_at 1.08746 "phospholipase A2, group IIE" Pla2g2e
1420675_at 0.90682 zinc finger protein 113 Zfp113
1420676_at 1.06954 small proline rich-like 3 Sprl3
1420677_x_at 0.97958 small proline rich-like 3 Sprl3
1420678_a_at 1.07567 interleukin 17 receptor B Il17rb
1420679_a_at 0.82168 androgen-induced 1 Aig1
1420680_at 0.92941 cathepsin 8 Cts8
1420681_at 1.00704 RIKEN cDNA 1700120K04 gene 1700120K04Rik

1420682_at 1.18456 "cholinergic receptor, nicotinic, beta polypeptide 1 (muscle)" Chrb1
 1420683_at 1.14992 BCL2/adenovirus E1B 19kD interacting protein like Bnpl
 1420684_at 0.84405 "acyl-Coenzyme A oxidase 3, pristanoyl" Acox3
 1420685_at 0.82832 GRB2-related adaptor protein 2 /// similar to GRB2-related adaptor protein 2
 (GADS protein) (Growth factor receptor binding protein) (GRBLG) (GRB-2-like protein) (GRB2L)
 (Hematopoietic cell-associated adaptor protein GrpL) (GRB-2-related monocytic adapter protein)
 (Monocytic adapter... Grap2 /// LOC674050
 1420686_at 0.95459 "crystallin, beta A4" Cryba4
 1420687_at 0.96062 RIKEN cDNA 4932438H23 gene 4932438H23Rik
 1420688_a_at 0.86498 "sarcoglycan, epsilon" Sgce
 1420689_at 0.97586 RIKEN cDNA 4933415F23 gene 4933415F23Rik
 1420690_at 0.88806 fibroblast growth factor 10 Fgf10
 1420691_at 0.97919 "interleukin 2 receptor, alpha chain" Il2ra
 1420692_at 0.93259 "interleukin 2 receptor, alpha chain" Il2ra
 1420693_at 0.89809 myomesin 1 Myom1
 1420694_a_at 1.05165 dachshund 1 (Drosophila) Dach1
 1420695_at 0.94146 dachshund 1 (Drosophila) Dach1
 1420696_at 0.94569 "sema domain, immunoglobulin domain (Ig), short basic domain, secreted,
 (semaphorin) 3C" Sema3c
 1420697_at 0.91166 "solute carrier family 15, member 3" Slc15a3
 1420698_at 1.08587 glutathione peroxidase 5 Gpx5
 1420699_at 1.00232 "C-type lectin domain family 7, member a" Clec7a
 1420700_s_at 1.05449 folate receptor 4 (delta) Folr4
 1420701_at 1.00996 kallikrein 1-related peptidase b1 Klk1b1
 1420702_at 1.08727 RIKEN cDNA 1700093K21 gene 1700093K21Rik
 1420703_at 1.04832 "colony stimulating factor 2 receptor, alpha, low-affinity (granulocyte-
 macrophage)" Csf2ra
 1420704_at 0.86633 "colony stimulating factor 2 receptor, alpha, low-affinity (granulocyte-
 macrophage) /// similar to Granulocyte-macrophage colony-stimulating factor receptor alpha chain
 precursor (GM-CSF-R-alpha) (GMR) (CD116 antigen)" Csf2ra /// LOC671366
 1420705_at 0.94281 forkhead box B1 Foxb1
 1420706_at 1.00913 P140 gene P140
 1420707_a_at 1.23079 TRAF-interacting protein Traip
 1420708_at 0.91157 TRAF-interacting protein Traip
 1420709_s_at 0.97797 D-amino acid oxidase 1 Dao1
 1420710_at 1.17098 reticuloendotheliosis oncogene Rel
 1420711_a_at 0.92764 peroxisomal membrane protein 3 Pxmp3
 1420712_a_at 1.36636 hepsin Hpn
 1420713_a_at 0.97833 MyoD family inhibitor Mdf1
 1420714_at 0.94155 ladybird homeobox homolog 2 (Drosophila) Lbx2
 1420715_a_at 0.93175 peroxisome proliferator activated receptor gamma Pparg
 1420716_at 0.79570 CD247 antigen Cd247
 1420717_at 0.98586 RIKEN cDNA A930018P22 gene A930018P22Rik

1420718_at 1.03048 odd Oz/ten-m homolog 2 (Drosophila) /// similar to odd Oz/ten-m homolog
 2 Odz2 /// LOC669490
 1420719_at 0.88220 testis expressed gene 15 Tex15
 1420720_at 1.11656 neuronal pentraxin 2 Nptx2
 1420721_at 1.08943 RIKEN cDNA 4921536K21 gene 4921536K21Rik
 1420722_at 1.00397 "elongation of very long chain fatty acids (FEN1/Elo2, SUR4/Elo3, yeast)-like
 3" Elovl3
 1420723_at 1.00668 vanin 3 Vnn3
 1420724_at 0.92883 RIKEN cDNA 1700067P10 gene 1700067P10Rik
 1420725_at 0.97229 "trimethyllysine hydroxylase, epsilon" Tmlhe
 1420726_x_at 1.10285 "trimethyllysine hydroxylase, epsilon" Tmlhe
 1420727_a_at 1.07937 "trimethyllysine hydroxylase, epsilon" Tmlhe
 1420728_at 1.03314 keratin 32 Krt32
 1420729_at 0.93465 "2-cell-stage, variable group, member 1" Tcstv1
 1420730_a_at 1.08648 t-complex protein 11 Tcp11
 1420731_a_at 0.77282 cysteine and glycine-rich protein 2 Csrp2
 1420732_at 0.97874 RIKEN cDNA 2510027J23 gene 2510027J23Rik
 1420733_at 0.93974 calcium binding protein 39-like Cab39l
 1420734_at 1.01299 "protein phosphatase 1, regulatory (inhibitor) subunit 3F" Ppp1r3f
 1420735_at 1.00755 "gamma-aminobutyric acid (GABA-C) receptor, subunit rho 2" Gabrr2
 1420736_at 0.90574 single-minded homolog 1 (Drosophila) Sim1
 1420737_at 0.96538 polyamine modulated factor 1 binding protein 1 Pmfbp1
 1420738_at 1.09254 myeloid zinc finger 1 Mzf1
 1420739_at 0.92348 contactin 3 Cntn3
 1420740_at 1.04033 interleukin 25 Il25
 1420741_x_at 0.99437 RIKEN cDNA 2310069N01 gene 2310069N01Rik
 1420742_at 0.98465 a disintegrin and metallopeptidase domain 7 Adam7
 1420743_a_at 0.99367 "protein phosphatase 3, catalytic subunit, gamma isoform" Ppp3cc
 1420744_at 1.01820 "cholinergic receptor, nicotinic, beta polypeptide 2 (neuronal)" Chrn2
 1420745_a_at 0.88142 cyclin D-type binding-protein 1 Ccndbp1
 1420746_at 0.99190 prolactin-like protein I Prlpi
 1420747_at 2.34942 per-pentamer repeat gene Ppnr
 1420748_a_at 1.05881 "adenosine deaminase, tRNA-specific 1" Adat1
 1420749_a_at 0.93094 "POU domain, class 6, transcription factor 1" Pou6f1
 1420750_at 1.06160 RIKEN cDNA 1110006O24 gene 1110006O24Rik
 1420751_at 0.99885 keratin associated protein 6-1 Krtap6-1
 1420752_at 0.83271 deltex 3 homolog (Drosophila) Dtx3
 1420753_at 1.00301 tolloid-like Tll1
 1420754_at 1.03309 transcription termination factor 1 Ttf1
 1420755_a_at 1.11980 parkin Park2
 1420756_at 1.02062 fucosyltransferase 7 Fut7
 1420757_at 0.99681 myogenic factor 5 Myf5
 1420758_at 0.99003 RIKEN cDNA 583041J07 gene 583041J07Rik

1420759_s_at 1.01886 "zinc finger protein 1, Y linked /// zinc finger protein 2, Y linked" Zfy1 ///
 Zfy2
 1420760_s_at 1.08424 N-myc downstream regulated-like Ndr1
 1420761_at 0.97825 "cholinergic receptor, nicotinic, delta polypeptide" Chrnd
 1420762_a_at 0.86020 Y box protein 2 Ybx2
 1420763_at 0.95666 "kallikrein 4 (prostase, enamel matrix, prostate)" Klk4
 1420764_at 0.94209 scrapie responsive gene 1 Scrg1
 1420765_a_at 0.90726 forkhead box P3 Foxp3
 1420766_at 0.97756 suppressor of cytokine signaling 7 Socs7
 1420767_at 0.95823 RIKEN cDNA 1700019G17 gene 1700019G17Rik
 1420768_a_at 0.92876 "DNA segment, Chr 11, Lothar Hennighausen 2, expressed" D11Lgp2e
 1420769_at 0.99975 "DNA segment, Chr 11, Lothar Hennighausen 2, expressed" D11Lgp2e
 1420770_at 1.05633 kallikrein 1-related peptidase b24 Klk1b24
 1420771_at 1.00066 small proline-rich protein 2D Sprr2d
 1420772_a_at 0.71944 TSC22 domain family 3 Tsc22d3
 1420773_at 0.84839 deubiquitinating enzyme 1 Dub1
 1420774_a_at 1.11303 RIKEN cDNA 4930583H14 gene 4930583H14Rik
 1420775_at 0.98224 haloacid dehalogenase-like hydrolase domain Hdhd1a
 1420776_a_at 0.83973 AU RNA binding protein/enoyl-coenzyme A hydratase Auh
 1420777_a_at 1.00434 RIKEN cDNA 4933400A11 gene 4933400A11Rik
 1420778_at 0.99178 "taste receptor, type 1, member 3" Tas1r3
 1420779_at 1.05728 four and a half LIM domains 3 Fhl3
 1420780_at 0.99696 achaete-scute complex homolog-like 3 (Drosophila) Ascl3
 1420781_at 1.20562 ectopic ossification 1 Etos1
 1420782_at 1.08268 "tumor necrosis factor receptor superfamily, member 17" Tnfrsf17
 1420783_at 0.94434 tumor protein D52-like 3 Trpd52l3
 1420784_at 0.96981 "sodium channel, voltage-gated, type XI, alpha" Scn11a
 1420785_at 1.18373 growth factor receptor bound protein 2-associated protein 2 Gab2
 1420786_a_at 0.97899 "RNA binding motif protein, Y chromosome, family 1, member A1" Rbmy1a1
 1420787_at 1.09162 RIKEN cDNA 4933436I01 gene 4933436I01Rik
 1420788_at 1.13949 "killer cell lectin-like receptor subfamily G, member 1" Klr1g1
 1420789_at 1.13837 "killer cell lectin-like receptor, subfamily A, member 5" Klra5
 1420790_x_at 1.11555 "killer cell lectin-like receptor, subfamily A, member 16" Klra16
 1420791_x_at 1.03340 spermatogenesis associated glutamate (E)-rich protein 4f Speer4f
 1420792_at 1.06637 RIKEN cDNA 4930433N12 gene 4930433N12Rik
 1420793_at 0.93058 major urinary protein 4 Mup4
 1420794_at 0.98393 ADP-ribosyltransferase 2b Art2b
 1420795_at 1.07925 fibroblast growth factor 9 Fgf9
 1420796_at 1.01909 aryl-hydrocarbon receptor repressor Ahrr
 1420797_at 0.95782 otogelin Otog
 1420798_s_at 0.82033 "protocadherin alpha 4 /// protocadherin alpha 6 /// protocadherin alpha 7 ///
 protocadherin alpha 5 /// protocadherin alpha 11 /// protocadherin alpha 10 /// protocadherin alpha 1 ///
 protocadherin alpha 9 /// protocadherin alpha 3 /// protocadherin alpha 12 /// protocadherin alpha 2 ///

protocadherin alpha 8 /// protocadherin alpha subfamily C, 1 /// protocadherin alpha subfamily C, 2"
 Pcdha4 /// Pcdha6 /// Pcdha7 /// Pcdha5 /// Pcdha11 /// Pcdha10 /// Pcdha1 /// Pcdha9 /// Pcdha3 ///
 Pcdha12 /// Pcdha2 /// Pcdha8 /// Pcdhac1 /// Pcdhac2
 1420799_at 1.00355 neurotensin receptor 1 Ntsr1
 1420800_a_at 1.16036 "potassium voltage-gated channel, subfamily Q, member 2" Kcnq2
 1420801_at 0.89593 neuronal PAS domain protein 1 Npas1
 1420802_at 1.05941 interleukin 13 Il13
 1420803_at 1.05125 testis expressed gene 18 Tex18
 1420804_s_at 1.08230 "C-type lectin domain family 4, member d" Clec4d
 1420805_at 0.98258 "myosin light chain 2, precursor lymphocyte-specific" Mylc2pl
 1420806_at 1.04543 fibroblast growth factor 16 Fgf16
 1420807_a_at 1.12154 "EGF-like-domain, multiple 9" Egf19
 1420808_at 0.95859 nuclear receptor coactivator 4 /// similar to nuclear receptor coactivator 4
 Ncoa4 /// LOC627557
 1420809_a_at 0.94974 RIKEN cDNA 1500003O03 gene 1500003O03Rik
 1420810_at 0.99943 RIKEN cDNA 1500003O03 gene 1500003O03Rik
 1420811_a_at 1.02418 "catenin (cadherin associated protein), beta 1" Ctnnb1
 1420812_at 0.98766 histone deacetylase 7A Hdac7a
 1420813_at 0.98448 histone deacetylase 7A Hdac7a
 1420814_at 0.98187 guanosine diphosphate (GDP) dissociation inhibitor 2 Gdi2
 1420815_at 0.95050 guanosine diphosphate (GDP) dissociation inhibitor 2 Gdi2
 1420816_at 1.03854 "3-monooxygenase/tryptophan 5-monooxygenase activation protein, gamma
 polypeptide" Ywhag
 1420817_at 1.01022 "3-monooxygenase/tryptophan 5-monooxygenase activation protein, gamma
 polypeptide" Ywhag
 1420818_at 0.87266 src-like adaptor Sla
 1420819_at 0.84659 src-like adaptor Sla
 1420820_at 0.98420 RIKEN cDNA 2900073G15 gene 2900073G15Rik
 1420821_at 1.09303 sphingosine-1-phosphate phosphatase 1 Sgpp1
 1420822_s_at 1.07028 sphingosine-1-phosphate phosphatase 1 Sgpp1
 1420823_at 0.86625 "sema domain, immunoglobulin domain (Ig), transmembrane domain (TM) and
 short cytoplasmic domain, (semaphorin) 4D" Sema4d
 1420824_at 0.89478 "sema domain, immunoglobulin domain (Ig), transmembrane domain (TM) and
 short cytoplasmic domain, (semaphorin) 4D" Sema4d
 1420825_at 0.97330 leucine zipper-EF-hand containing transmembrane protein 1 Letm1
 1420826_at 1.01848 leucine zipper-EF-hand containing transmembrane protein 1 Letm1
 1420827_a_at 1.13854 cyclin G1 Ccng1
 1420828_s_at 0.97580 "tyrosine 3-monooxygenase/tryptophan 5-monooxygenase activation protein,
 theta polypeptide" Ywhaq
 1420829_a_at 0.93118 "tyrosine 3-monooxygenase/tryptophan 5-monooxygenase activation protein,
 theta polypeptide" Ywhaq
 1420830_x_at 0.88458 "tyrosine 3-monooxygenase/tryptophan 5-monooxygenase activation protein,
 theta polypeptide" Ywhaq

1420831_at	0.94286	quiescin Q6	Qscn6	
1420832_at	1.00199	quiescin Q6	Qscn6	
1420833_at	0.85761	vesicle-associated membrane protein 2	Vamp2	
1420834_at	0.92322	vesicle-associated membrane protein 2	Vamp2	
1420835_at	0.94754	"solute carrier family 25, member 30"	Slc25a30	
1420836_at	0.94990	"solute carrier family 25, member 30"	Slc25a30	
1420837_at	1.04204	"neurotrophic tyrosine kinase, receptor, type 2"	Ntrk2	
1420838_at	1.12562	"neurotrophic tyrosine kinase, receptor, type 2"	Ntrk2	
1420839_at	0.99631	"pleckstrin homology domain-containing, family A (phosphoinositide binding specific) member 3"	Plekha3	
1420840_at	0.97214	"pleckstrin homology domain-containing, family A (phosphoinositide binding specific) member 3"	Plekha3	
1420841_at	0.95828	"protein tyrosine phosphatase, receptor type, F"	Ptprf	
1420842_at	0.93359	"protein tyrosine phosphatase, receptor type, F"	Ptprf	
1420843_at	0.91314	"protein tyrosine phosphatase, receptor type, F"	Ptprf	
1420844_at	0.99078	ubiquilin 2	Ubqln2	
1420845_at	0.95986	mitochondrial ribosomal protein S2	Mrps2	
1420846_at	0.97358	mitochondrial ribosomal protein S2	Mrps2	
1420847_a_at	0.56703	fibroblast growth factor receptor 2	Fgfr2	
1420848_at	0.90520	suppressor of fused homolog (Drosophila)	Sufu	
1420849_at	1.04332	"Crn, crooked neck-like 1 (Drosophila)"	Crnk11	
1420850_at	0.97569	"Crn, crooked neck-like 1 (Drosophila)"	Crnk11	
1420851_at	0.87232	par-6 partitioning defective 6 homolog gamma (C. elegans)	Pard6g	
1420852_a_at	0.96264	"UDP-GlcNAc:betaGal beta-1,3-N-acetylglucosaminyltransferase 2"	B3gnt2	
1420853_at	1.01098	syndecan 3	Sdc3	
1420854_at	1.06557	elastin	Eln	
1420855_at	1.03183	elastin	Eln	
1420856_a_at	0.94760	LanC (bacterial lantibiotic synthetase component C)-like 2	Lancl2	
1420857_at	1.01063	LanC (bacterial lantibiotic synthetase component C)-like 2	Lancl2	
1420858_at	0.79175	"protein kinase inhibitor, alpha"	Pkia	
1420859_at	0.75745	"protein kinase inhibitor, alpha"	Pkia	
1420860_at	0.87016	integrin alpha 9	Itga9	
1420861_at	0.90042	dynactin 4	Dctn4	
1420862_at	0.93959	dynactin 4	Dctn4	
1420863_at	0.89153	dynactin 4	Dctn4	
1420864_at	0.82678	zinc finger protein 161	Zfp161	
1420865_at	0.92611	zinc finger protein 161	Zfp161	
1420866_at	0.80675	zinc finger protein 161	Zfp161	
1420867_at	0.83464	transmembrane emp24 domain trafficking protein 2	Tmed2	
1420868_s_at	0.90044	transmembrane emp24 domain trafficking protein 2	Tmed2	
1420869_at	0.99586	myeloid/lymphoid or mixed lineage-leukemia translocation to 10 homolog (Drosophila)	Mllt10	
1420870_at	1.07046	myeloid/lymphoid or mixed lineage-leukemia translocation to 10 homolog		

(Drosophila) Mllt10

1420871_at 1.13354 "guanylate cyclase 1, soluble, beta 3" Gucy1b3

1420872_at 1.14404 "guanylate cyclase 1, soluble, beta 3" Gucy1b3

1420873_at 1.08597 protein tyrosine kinase 9 Ptk9

1420874_at 0.93300 protein tyrosine kinase 9 Ptk9

1420875_at 1.03224 protein tyrosine kinase 9 Ptk9

1420876_a_at 0.95704 septin 6 6-Sep

1420877_at 0.80914 septin 6 6-Sep

1420878_a_at 0.97172 "tyrosine 3-monooxygenase/tryptophan 5-monooxygenase activation protein, beta polypeptide" Ywhab

1420879_a_at 0.88713 "tyrosine 3-monooxygenase/tryptophan 5-monooxygenase activation protein, beta polypeptide" Ywhab

1420880_a_at 0.88969 "tyrosine 3-monooxygenase/tryptophan 5-monooxygenase activation protein, beta polypeptide" Ywhab

1420881_at 0.85786 nuclear receptor-binding SET-domain protein 1 Nsd1

1420882_a_at 0.97073 nuclear receptor-binding SET-domain protein 1 /// adrenocortical dysplasia Nsd1 /// Acd

1420883_at 0.92759 sarcolipin Sln

1420884_at 0.92208 sarcolipin Sln

1420885_a_at 1.09806 seizure related gene 6 Sez6

1420886_a_at 0.83042 X-box binding protein 1 Xbp1

1420887_a_at 0.89753 Bcl2-like 1 Bcl2l1

1420888_at 1.02580 Bcl2-like 1 Bcl2l1

1420889_at 0.90404 holocytochrome c synthetase Hccs

1420890_at 0.91128 holocytochrome c synthetase Hccs

1420891_at 1.11765 wingless-related MMTV integration site 7B Wnt7b

1420892_at 1.20304 wingless-related MMTV integration site 7B Wnt7b

1420893_a_at 0.95614 "transforming growth factor, beta receptor I" Tgfbr1

1420894_at 0.82972 "transforming growth factor, beta receptor I" Tgfbr1

1420895_at 0.87068 "transforming growth factor, beta receptor I" Tgfbr1

1420896_at 0.97300 synaptosomal-associated protein 23 Snap23

1420897_at 0.93582 synaptosomal-associated protein 23 Snap23

1420898_at 1.02065 synaptosomal-associated protein 23 Snap23

1420899_at 0.95076 "RAB18, member RAS oncogene family" Rab18

1420900_a_at 0.96081 "RAB18, member RAS oncogene family" Rab18

1420901_a_at 1.02740 hexokinase 1 Hk1

1420902_at 1.07994 "ST6 (alpha-N-acetyl-neuraminy-2,3-beta-galactosyl-1,3)-N-acetylgalactosaminide alpha-2,6-sialyltransferase 3" St6galnac3

1420903_at 0.88776 "ST6 (alpha-N-acetyl-neuraminy-2,3-beta-galactosyl-1,3)-N-acetylgalactosaminide alpha-2,6-sialyltransferase 3" St6galnac3

1420904_at 0.96155 interleukin 17 receptor A Il17ra

1420905_at 1.31580 interleukin 17 receptor A Il17ra

1420906_at 1.15706 CD2-associated protein Cd2ap

1420907_at	1.19999	CD2-associated protein	Cd2ap
1420908_at	1.15694	CD2-associated protein	Cd2ap
1420909_at	1.03136	vascular endothelial growth factor A	Vegfa
1420910_at	0.86458	phosphatidic acid phosphatase type 2c	Ppap2c
1420911_a_at	0.82143	milk fat globule-EGF factor 8 protein	Mfge8
1420912_at	1.65249	"AF4/FMR2 family, member 4"	Aff4
1420913_at	0.97706	"solute carrier organic anion transporter family, member 2a1"	Slco2a1
1420914_at	1.10944	---	---
1420915_at	0.99210	signal transducer and activator of transcription 1	Stat1
1420916_at	1.09834	PRP40 pre-mRNA processing factor 40 homolog A (yeast)	Prpf40a
1420917_at	1.03257	PRP40 pre-mRNA processing factor 40 homolog A (yeast)	Prpf40a
1420918_at	0.77922	serum/glucocorticoid regulated kinase 3	Sgk3
1420919_at	0.72781	serum/glucocorticoid regulated kinase 3	Sgk3
1420920_a_at	0.97394	ADP-ribosylation factor 1	Arf1
1420921_at	1.05338	CD151 antigen	Cd151
1420922_at	0.96384	"ubiquitin specific peptidase 9, X chromosome"	Usp9x
1420923_at	1.17624	"ubiquitin specific peptidase 9, X chromosome"	Usp9x
1420924_at	0.80993	tissue inhibitor of metalloproteinase 2	Timp2
1420925_at	0.90042	tubby candidate gene	Tub
1420926_at	1.06070	aristaless related homeobox gene (Drosophila)	Arx
1420927_at	0.88079	"beta galactoside alpha 2,6 sialyltransferase 1"	St6gal1
1420928_at	0.88620	"beta galactoside alpha 2,6 sialyltransferase 1"	St6gal1
1420929_at	0.87285	"catenin (cadherin associated protein), alpha-like 1"	Ctnnal1
1420930_s_at	0.87699	"catenin (cadherin associated protein), alpha-like 1"	Ctnnal1
1420931_at	1.06712	mitogen activated protein kinase 8	Mapk8
1420932_at	1.06900	mitogen activated protein kinase 8	Mapk8
1420933_a_at	1.09490	eyes absent 3 homolog (Drosophila)	Eya3
1420934_a_at	1.07536	serine/arginine repetitive matrix 1	Srrm1
1420935_a_at	1.13489	serine/arginine repetitive matrix 1	Srrm1
1420936_s_at	1.07742	cleavage and polyadenylation specific factor 2	Cpsf2
1420937_at	1.03963	cleavage and polyadenylation specific factor 2	Cpsf2
1420938_at	0.89702	heparan sulfate 6-O-sulfotransferase 2	Hs6st2
1420939_at	0.98079	heparan sulfate 6-O-sulfotransferase 2	Hs6st2
1420940_x_at	0.93962	regulator of G-protein signaling 5	Rgs5
1420941_at	0.96711	regulator of G-protein signaling 5	Rgs5
1420942_s_at	0.93722	---	---
1420943_at	0.88805	zinc finger protein 185	Zfp185
1420944_at	0.79373	zinc finger protein 185	Zfp185
1420945_at	0.89667	alpha thalassemia/mental retardation syndrome X-linked homolog (human)	Atrx
1420946_at	0.91715	alpha thalassemia/mental retardation syndrome X-linked homolog (human)	Atrx
1420947_at	1.07058	alpha thalassemia/mental retardation syndrome X-linked homolog (human)	Atrx
1420948_s_at	0.87014	RIKEN cDNA 4833408C14 gene	4833408C14Rik
1420949_at	0.79631	zinc and ring finger 1	Znrf1

1420950_at	0.98093	zinc and ring finger 1	Znrf1
1420951_a_at	0.90079	Son cell proliferation protein	Son
1420952_at	0.92706	Son cell proliferation protein	Son
1420953_at	0.89042	adducin 1 (alpha)	Add1
1420954_a_at	0.91909	adducin 1 (alpha)	Add1
1420955_at	1.10386	visinin-like 1	Vsnl1
1420956_at	1.01601	adenomatosis polyposis coli	Apc
1420957_at	0.86561	adenomatosis polyposis coli	Apc
1420958_at	1.10744	---	---
1420959_at	0.75283	aspartate-beta-hydroxylase	Asph
1420960_at	0.85952	"Fanconi anemia, complementation group G"	Fancg
1420961_a_at	0.65129	influenza virus NS1A binding protein	Ivns1abp
1420962_at	1.02888	hyaluronan and proteoglycan link protein 2	Hapln2
1420963_at	0.89712	hyaluronan and proteoglycan link protein 2	Hapln2
1420964_at	1.04391	ectodermal-neural cortex 1	Enc1
1420965_a_at	0.98595	ectodermal-neural cortex 1	Enc1
1420966_at	0.94397	"solute carrier family 25 (mitochondrial carrier ornithine transporter), member 15"	Slc25a15
1420967_at	1.03218	"solute carrier family 25 (mitochondrial carrier ornithine transporter), member 15"	Slc25a15
1420968_at	0.98596	BTB (POZ) domain containing 14B	Btbd14b
1420969_at	0.97924	BTB (POZ) domain containing 14B	Btbd14b
1420970_at	0.96748	adenylate cyclase 7	Adcy7
1420971_at	0.98895	ubiquitin protein ligase E3 component n-recognin 1	Ubr1
1420972_at	0.92040	AT rich interactive domain 5B (Mrf1 like)	Arid5b
1420973_at	0.76241	AT rich interactive domain 5B (Mrf1 like)	Arid5b
1420974_at	1.76011	"SET domain, bifurcated 1"	Setdb1
1420975_at	1.04652	"bromodomain adjacent to zinc finger domain, 1B"	Baz1b
1420976_at	1.03011	"mannosidase, alpha, class 1A, member 2"	Man1a2
1420977_at	1.00739	"mannosidase, alpha, class 1A, member 2"	Man1a2
1420978_at	1.00587	nuclear respiratory factor 1	Nrf1
1420979_at	0.99340	p21 (CDKN1A)-activated kinase 1	Pak1
1420980_at	0.95104	p21 (CDKN1A)-activated kinase 1	Pak1
1420981_a_at	0.91617	LIM domain only 4	Lmo4
1420982_at	0.96376	RNA binding motif protein 39	Rbm39
1420983_at	1.00650	phosphatidylcholine transfer protein	Pctp
1420984_at	1.11142	phosphatidylcholine transfer protein	Pctp
1420985_at	1.16252	"ash1 (absent, small, or homeotic)-like (Drosophila)"	Ash11
1420986_s_at	1.00934	---	---
1420987_at	0.91203	kinesin family member 3B	Kif3b
1420988_at	1.05406	"polymerase (DNA directed), eta (RAD 30 related)"	Polh
1420989_at	0.82483	RIKEN cDNA 4933411K20 gene	4933411K20Rik
1420990_at	1.15187	chromodomain helicase DNA binding protein 1	Chd1

1420991_at 1.18914 ankyrin repeat domain 1 (cardiac muscle) Ankrd1
1420992_at 1.24399 ankyrin repeat domain 1 (cardiac muscle) Ankrd1
1420993_at 0.99897 "UDP-GlcNAc:betaGal beta-1,3-N-acetylglucosaminyltransferase 5" B3gnt5
1420994_at 0.97896 "UDP-GlcNAc:betaGal beta-1,3-N-acetylglucosaminyltransferase 5" B3gnt5
1420995_at 0.87135 plexin A3 Plxna3
1420996_at 0.83726 plexin A3 Plxna3
1420997_a_at 1.03975 glucose phosphate isomerase 1 /// similar to Glucose-6-phosphate isomerase (GPI) (Phosphoglucose isomerase) (PGI) (Phosphohexose isomerase) (PHI) (Neuroleukin) (NLK) Gpi1 /// LOC676596
1420998_at 0.95618 ets variant gene 5 Etv5
1420999_at 1.17964 "CCR4-NOT transcription complex, subunit 4" Cnot4
1421000_at 1.33805 "CCR4-NOT transcription complex, subunit 4" Cnot4
1421001_a_at 0.97460 carbonic anhydrase 6 Car6
1421002_at 0.81116 angiopoietin-like 2 Angptl2
1421003_at 0.93910 glucocorticoid modulatory element binding protein 1 Gmeb1
1421004_at 0.84310 centrosomal protein 110 Cep110
1421005_at 0.98896 centrosomal protein 110 Cep110
1421006_at 0.90507 "procollagen, type IV, alpha 6" Col4a6
1421007_at 0.88375 "procollagen, type IV, alpha 6" Col4a6
1421008_at 0.97717 radical S-adenosyl methionine domain containing 2 Rsad2
1421009_at 1.12333 radical S-adenosyl methionine domain containing 2 Rsad2
1421010_at 1.08223 myelin-associated oligodendrocytic basic protein Mobp
1421011_at 0.80898 dehydrogenase/reductase (SDR family) member 8 Dhhrs8
1421012_at 0.86428 "signal recognition particle receptor, B subunit" Srprb
1421013_at 1.00454 "phosphatidylinositol transfer protein, beta" Pitpnb
1421014_a_at 0.64337 citrate lyase beta like Clybl
1421015_s_at 1.05345 "polymerase (DNA directed), epsilon 3 (p17 subunit)" Pole3
1421016_at 0.97277 immunoglobulin mu binding protein 2 Ighmbp2
1421017_at 0.99567 neuregulin 3 Nrg3
1421018_at 1.00680 RIKEN cDNA 1110018J18 gene 1110018J18Rik
1421019_at 0.92500 RIKEN cDNA 1700021F05 gene 1700021F05Rik
1421020_at 0.88365 RAN binding protein 17 Ranbp17
1421021_at 0.96807 --- ---
1421022_x_at 0.86712 "acylphosphatase 1, erythrocyte (common) type" Acyp1
1421023_at 0.99951 "phosphatidylinositol 3-kinase, C2 domain containing, alpha polypeptide" Pik3c2a
1421024_at 0.99301 "1-acylglycerol-3-phosphate O-acyltransferase 1 (lysophosphatidic acid acyltransferase, alpha)" Agpat1
1421025_at 0.88708 "1-acylglycerol-3-phosphate O-acyltransferase 1 (lysophosphatidic acid acyltransferase, alpha)" Agpat1
1421026_at 0.88029 "guanine nucleotide binding protein, alpha 12" Gna12
1421027_a_at 0.98572 myocyte enhancer factor 2C Mef2c
1421028_a_at 0.95104 myocyte enhancer factor 2C Mef2c

1421029_a_at 0.92355 Hbs1-like (*S. cerevisiae*) Hbs11
 1421030_at 0.93337 zinc finger protein 64 Zfp64
 1421031_a_at 1.52452 RIKEN cDNA 2310016C08 gene 2310016C08Rik
 1421032_a_at 0.96622 "DnaJ (Hsp40) homolog, subfamily B, member 12" Dnajb12
 1421033_a_at 1.14302 transcription elongation regulator 1 (CA150) Tcerg1
 1421034_a_at 0.96299 "interleukin 4 receptor, alpha" Il4ra
 1421035_a_at 0.95303 "membrane associated guanylate kinase, WW and PDZ domain containing 3"
 Magi3
 1421036_at 0.90333 neuronal PAS domain protein 2 Npas2
 1421037_at 1.00875 neuronal PAS domain protein 2 Npas2
 1421038_a_at 1.04912 "potassium intermediate/small conductance calcium-activated channel,
 subfamily N, member 4" Kcnn4
 1421039_at 1.10553 major intrinsic protein of eye lens fiber Mip
 1421040_a_at 0.97022 "glutathione S-transferase, alpha 2 (Yc2)" Gsta2
 1421041_s_at 1.85571 "glutathione S-transferase, alpha 1 (Ya) /// glutathione S-transferase, alpha 2
 (Yc2)" Gsta1 /// Gsta2
 1421042_at 1.01596 rho/rac guanine nucleotide exchange factor (GEF) 2 Arhgef2
 1421043_s_at 0.97924 rho/rac guanine nucleotide exchange factor (GEF) 2 Arhgef2
 1421044_at 1.09571 "mannose receptor, C type 2" Mrc2
 1421045_at 1.01801 "mannose receptor, C type 2" Mrc2
 1421046_a_at 1.03217 "poly A binding protein, cytoplasmic 4 /// similar to Poly(A) binding protein,
 cytoplasmic 4, isoform 1" Pabpc4 /// LOC432881
 1421047_at 0.96505 MAD homolog 5 (*Drosophila*) Smad5
 1421048_a_at 0.62127 yippee-like 1 (*Drosophila*) Ypell
 1421049_at 1.11461 RIKEN cDNA 0610009O20 gene 0610009O20Rik
 1421050_at 1.07091 vacuolar protein sorting 25 (yeast) Vps25
 1421051_s_at 1.03935 vacuolar protein sorting 25 (yeast) Vps25
 1421052_a_at 1.09902 spermine synthase /// similar to Spermine synthase (Spermidine
 aminopropyltransferase) (SPMSY) Sms /// LOC671878
 1421053_at 0.96887 kinesin family member 1A Kif1a
 1421054_at 1.28294 exportin 4 Xpo4
 1421055_at 1.29048 large tumor suppressor 2 /// exportin 4 Lats2 /// Xpo4
 1421056_at 1.07677 deoxyribonuclease 1-like 3 Dnase113
 1421057_at 0.99846 deoxyribonuclease 1-like 3 Dnase113
 1421058_at 0.85104 "alcohol dehydrogenase 7 (class IV), mu or sigma polypeptide" Adh7
 1421059_a_at 0.89838 "asparagine-linked glycosylation 2 homolog (yeast, alpha-1,3-
 mannosyltransferase)" Alg2
 1421060_at 0.87567 myeloid/lymphoid or mixed lineage-leukemia translocation to 1 homolog
 (*Drosophila*) Mllt1
 1421061_at 1.21937 guanylate cyclase activator 1a (retina) Guca1a
 1421062_s_at 0.99243 "clathrin, light polypeptide (Lca)" Clta
 1421063_s_at 0.99554 small nuclear ribonucleoprotein N /// SNRPN upstream reading frame
 Snrpn /// Snurf

1421064_at 1.04500 "membrane protein, palmitoylated 5 (MAGUK p55 subfamily member 5)"
Mpp5

1421065_at 0.96349 Janus kinase 2 Jak2

1421066_at 0.95607 --- ---

1421067_a_at 0.94884 PHD finger protein 2 Phf2

1421068_at 1.11014 PHD finger protein 2 Phf2

1421069_at 0.79616 PHD finger protein 2 Phf2

1421070_at 0.73941 "DNA segment, Chr 3, ERATO Doi 300, expressed" D3ErtD300e

1421071_at 1.09989 von Hippel-Lindau syndrome homolog Vhlh

1421072_at 0.95723 Iroquois related homeobox 5 (Drosophila) Irx5

1421073_a_at 0.94826 prostaglandin E receptor 4 (subtype EP4) Ptger4

1421074_at 0.92259 "cytochrome P450, family 7, subfamily b, polypeptide 1" Cyp7b1

1421075_s_at 0.95262 "cytochrome P450, family 7, subfamily b, polypeptide 1" Cyp7b1

1421076_at 0.85555 SERTA domain containing 3 Sertad3

1421077_at 0.89813 SERTA domain containing 3 Sertad3

1421078_at 0.95055 transcription factor 23 Tcf23

1421079_at 1.04581 "nuclear receptor subfamily 4, group A, member 3" Nr4a3

1421080_at 0.91301 "nuclear receptor subfamily 4, group A, member 3" Nr4a3

1421081_a_at 1.11880 barrier to autointegration factor 1 Banf1

1421082_s_at 1.05804 barrier to autointegration factor 1 Banf1

1421083_x_at 1.10646 barrier to autointegration factor 1 Banf1

1421084_at 1.00739 retinoschisis 1 homolog (human) Rs1h

1421085_at 1.01275 retinoschisis 1 homolog (human) Rs1h

1421086_at 0.95937 period homolog 3 (Drosophila) Per3

1421087_at 0.90740 period homolog 3 (Drosophila) Per3

1421088_at 0.75616 glypican 4 Gpc4

1421089_a_at 1.05540 RIKEN cDNA 2610028A01 gene 2610028A01Rik

1421090_at 0.94986 erythrocyte protein band 4.1-like 1 Epb4.111

1421091_at 1.09259 "serine (or cysteine) peptidase inhibitor, clade A (alpha-1 antiproteinase, antitrypsin), member 12" Serpina12

1421092_at 1.06675 "serine (or cysteine) peptidase inhibitor, clade A (alpha-1 antiproteinase, antitrypsin), member 12" Serpina12

1421093_at 0.86869 "solute carrier family 7 (cationic amino acid transporter, y+ system), member 10" Slc7a10

1421094_at 0.96433 zinc finger and BTB domain containing 33 Zbtb33

1421095_a_at 0.98323 "transient receptor potential cation channel, subfamily C, member 1" Trpc1

1421096_at 1.01206 "transient receptor potential cation channel, subfamily C, member 1" Trpc1

1421097_at 0.94303 endonuclease G Endog

1421098_at 0.94867 expressed sequence AI586015 AI586015

1421099_at 0.98819 "basic helix-loop-helix domain containing, class B3" Bhlhb3

1421100_a_at 0.97518 disabled homolog 1 (Drosophila) Dab1

1421101_a_at 0.94666 LIM domain binding 2 Ldb2

1421102_a_at 0.99705 vesicle-associated membrane protein 3 Vamp3

1421103_at 0.94201 BMP2 inducible kinase Bmp2k
 1421104_at 0.89039 "macrophage activation 2 /// macrophage activation 2 like /// similar to
 macrophage activation 2 /// similar to macrophage activation 2 like /// similar to macrophage activation
 2 /// similar to guanylate binding protein family, member 6 /// similar to guanylate binding protein
 family, member 6" Mpa2 /// Mpa2l /// LOC547126 /// LOC626578 /// LOC634650 /// LOC671219 ///
 LOC673195
 1421105_at 1.06581 jagged 1 Jag1
 1421106_at 0.88644 jagged 1 Jag1
 1421107_at 1.08877 serine/threonine kinase 4 Stk4
 1421108_at 0.90777 camello-like 2 Cml2
 1421109_at 1.05822 camello-like 2 Cml2
 1421110_at 1.07358 Metadherin Mtdh
 1421111_at 0.87824 RING1 and YY1 binding protein Rybp
 1421112_at 1.00964 "NK2 transcription factor related, locus 2 (Drosophila)" Nkx2-2
 1421113_at 0.99835 "pepsinogen 5, group I" Pga5
 1421114_a_at 1.03884 dermatan sulphate proteoglycan 3 Dspg3
 1421115_a_at 0.96054 "zinc finger, DHHC domain containing 16" Zdhhc16
 1421116_a_at 0.93395 reticulon 4 Rtn4
 1421117_at 0.97084 dystonin Dst
 1421118_a_at 0.51927 G protein-coupled receptor 56 Gpr56
 1421119_at 0.93703 kinesin family member 21B Kif21b
 1421120_at 1.07459 myosin VI Myo6
 1421121_at 1.01212 A kinase (PRKA) anchor protein 10 Akap10
 1421122_at 0.99461 Casitas B-lineage lymphoma-like 1 Cbl1
 1421123_at 0.91366 "cyclin-dependent kinase 5, regulatory subunit (p35) 1" Cdk5r1
 1421124_at 1.01315 "cyclin-dependent kinase 5, regulatory subunit (p35) 1" Cdk5r1
 1421125_at 1.16951 spermatid perinuclear RNA binding protein Strbp
 1421126_at 0.95126 "ryanodine receptor 2, cardiac" Ryr2
 1421127_at 0.94835 transmembrane protein 42 Tmem42
 1421128_at 1.03011 "U2 small nuclear ribonucleoprotein auxiliary factor (U2AF) 1, related sequence
 2" U2af1-rs2
 1421129_a_at 0.92008 "ATPase, Ca++ transporting, ubiquitous" Atp2a3
 1421130_at 0.90806 zinc finger protein 111 Zfp111
 1421131_a_at 0.76347 zinc finger protein 111 Zfp111
 1421132_at 0.78391 poliovirus receptor-related 3 Pvr13
 1421133_at 1.13443 poliovirus receptor-related 3 Pvr13
 1421134_at 1.01317 amphiregulin Areg
 1421135_a_at 1.03076 "CCR4-NOT transcription complex, subunit 8" Cnot8
 1421136_at 0.80140 endothelin 3 Edn3
 1421137_a_at 1.11963 "protein kinase inhibitor beta, cAMP dependent, testis specific" Pkib
 1421138_a_at 1.11031 "protein kinase inhibitor beta, cAMP dependent, testis specific" Pkib
 1421139_a_at 1.03560 zinc finger protein 386 (Kruppel-like) Zfp386
 1421140_a_at 0.88106 forkhead box P1 Foxp1

1421141_a_at 0.89077 forkhead box P1 Foxp1
1421142_s_at 0.96846 forkhead box P1 Foxp1
1421143_at 1.07714 diaphanous homolog 1 (Drosophila) Diap1
1421144_at 0.99710 retinitis pigmentosa GTPase regulator interacting protein 1 Rpgrip1
1421145_at 0.99554 "solute carrier family 26 (sulfate transporter), member 2" Slc26a2
1421146_at 1.24541 Rap guanine nucleotide exchange factor (GEF) 1 Rapgef1
1421147_at 0.84601 telomeric repeat binding factor 2 Terf2
1421148_a_at 1.10993 Tial1 cytotoxic granule-associated RNA binding protein-like 1 Tial1
1421149_a_at 0.97856 atrophin 1 Atn1
1421150_at 1.00195 human immunodeficiency virus type I enhancer binding protein 3 Hivep3
1421151_a_at 1.12402 Eph receptor A2 Epha2
1421152_a_at 0.68773 "guanine nucleotide binding protein, alpha o" Gnao1
1421153_at 1.16225 lysyl oxidase-like 4 Loxl4
1421154_at 0.91784 "hyperpolarization-activated, cyclic nucleotide-gated K+ 2" Hcn2
1421155_at 1.03971 "UDP-Gal:betaGal beta 1,3-galactosyltransferase, polypeptide 6" B3galt6
1421156_a_at 1.00202 desmocollin 2 Dsc2
1421157_at 1.14533 frizzled homolog 3 (Drosophila) Fzd3
1421158_at 1.01182 cingulin-like 1 Cgnl1
1421159_at 1.00981 cingulin-like 1 /// similar to cingulin-like 1 isoform 1 Cgnl1 /// LOC677485
1421160_a_at 0.98744 radical fringe gene homolog (Drosophila) Rfng
1421161_at 1.03351 "betacellulin, epidermal growth factor family member" Btc
1421162_a_at 0.93080 nuclear factor I/A Nfia
1421163_a_at 0.83825 nuclear factor I/A Nfia
1421164_a_at 0.88139 Rho guanine nucleotide exchange factor (GEF) 1 Arhgef1
1421165_at 0.85542 c-myc binding protein Mycbp
1421166_at 0.95084 attractin Atrn
1421167_at 0.99383 "ATPase, class VI, type 11A" Atp11a
1421168_at 0.98700 "ATP-binding cassette, sub-family G (WHITE), member 3" Abcg3
1421169_at 1.11432 zinc finger protein 46 Zfp46
1421170_a_at 1.01289 "phospholipase C, beta 1" Plcb1
1421171_at 0.95901 a disintegrin and metallopeptidase domain 12 (meltrin alpha) Adam12
1421172_at 0.77663 a disintegrin and metallopeptidase domain 12 (meltrin alpha) Adam12
1421173_at 1.08202 interferon regulatory factor 4 Irf4
1421174_at 0.93544 interferon regulatory factor 4 Irf4
1421175_at 1.00119 myelin transcription factor 1-like Myt1l
1421176_at 0.83306 RAS guanyl releasing protein 1 Rasgrp1
1421177_at 0.90559 "THAP domain containing, apoptosis associated protein 2" Thap2
1421178_at 0.96569 "mannosyl (alpha-1,3-)-glycoprotein beta-1,4-N-acetylglucosaminyltransferase, isozyme C (putative)" Mgat4c
1421179_at 1.29155 tweety homolog 2 (Drosophila) Ttyh2
1421180_at 0.92461 limb expression 1 homolog (chicken) Lix1
1421181_at 1.20846 neuronal pentraxin receptor Nptxr
1421182_at 1.06111 "C-type lectin domain family 1, member b" Clec1b

1421183_at 0.82470 testis expressed gene 12 Tex12
1421184_a_at 0.95669 methyltransferase like 7A /// UbiE-YGHL1 fusion protein /// UbiE-YGHL1 fusion protein Mettl7a /// Ubie /// LOC554292
1421185_at 0.81547 methyltransferase like 7A Mettl7a
1421186_at 1.05732 chemokine (C-C motif) receptor 2 Ccr2
1421187_at 0.95737 chemokine (C-C motif) receptor 2 Ccr2
1421188_at 0.98504 chemokine (C-C motif) receptor 2 Ccr2
1421189_at 1.08836 "gamma-aminobutyric acid (GABA-A) receptor, subunit beta 3" Gabrb3
1421190_at 0.92749 "gamma-aminobutyric acid (GABA-A) receptor, subunit beta 3" Gabrb3
1421191_s_at 1.11354 golgi associated PDZ and coiled-coil motif containing Gopc
1421192_a_at 0.98889 intersectin 1 (SH3 domain protein 1A) Itsn1
1421193_a_at 0.99366 pre B-cell leukemia transcription factor 3 Pbx3
1421194_at 0.92744 integrin alpha 4 Itga4
1421195_at 0.98720 cholecystokinin A receptor Cckar
1421196_at 0.98871 "protein tyrosine phosphatase, non-receptor type 11" Ptpn11
1421197_a_at 1.05151 apoptotic chromatin condensation inducer 1 Acin1
1421198_at 0.88899 integrin alpha V Itgav
1421199_at 1.11967 "discs, large homolog 2 (Drosophila)" Dlgh2
1421200_at 0.94456 "discs, large homolog 2 (Drosophila)" Dlgh2
1421201_a_at 0.89566 trophinin Tro
1421202_at 0.97628 "cholinergic receptor, nicotinic, alpha polypeptide 4" Chrna4
1421203_at 1.01436 "cholinergic receptor, nicotinic, alpha polypeptide 4" Chrna4
1421204_a_at 0.86430 nudix (nucleoside diphosphate linked moiety X)-type motif 16 Nudt16
1421205_at 1.37856 ataxia telangiectasia mutated homolog (human) Atm
1421206_at 0.89041 leukemia inhibitory factor Lif
1421207_at 1.48106 leukemia inhibitory factor Lif
1421208_at 0.94360 inhibitor of kappaB kinase gamma Ikbkg
1421209_s_at 0.93812 inhibitor of kappaB kinase gamma Ikbkg
1421210_at 0.99602 class II transactivator C2ta
1421211_a_at 1.07158 class II transactivator C2ta
1421212_at 0.91271 "ATP-binding cassette, sub-family C (CFTR/MRP), member 6" Abcc6
1421213_at 1.05555 EDAR (ectodysplasin-A receptor)-associated death domain Edaradd
1421214_at 0.94727 cytidine monophospho-N-acetylneuraminic acid hydroxylase Cmah
1421215_a_at 1.06027 sarcolemma associated protein Smap
1421216_a_at 0.81675 iduronate 2-sulfatase Ids
1421217_a_at 1.08587 "lectin, galactose binding, soluble 9" Lgals9
1421218_at 1.06479 butyrylcholinesterase Bche
1421219_at 0.96319 "tubulin tyrosine ligase-like family, member 3" Ttl3
1421220_at 0.98637 ankyrin repeat domain 17 Ankrd17
1421221_at 0.86658 "beta-carotene 9', 10'-dioxygenase 2" Bcdo2
1421222_at 0.94031 FIP1 like 1 (S. cerevisiae) Fip111
1421223_a_at 1.05763 annexin A4 Anxa4
1421224_a_at 0.94655 transcription factor 2 Tcf2

1421225_a_at 1.13830 "solute carrier family 4 (anion exchanger), member 4" Slc4a4
1421226_at 0.96356 triggering receptor expressed on myeloid cells 2 Trem2
1421227_at 1.20754 granzyme E Gzme
1421228_at 1.17972 chemokine (C-C motif) ligand 7 Ccl7
1421229_at 1.02325 "DNA fragmentation factor, beta subunit" Dffb
1421230_a_at 0.97711 Musashi homolog 2 (Drosophila) Msi2
1421231_at 0.90764 Musashi homolog 2 (Drosophila) Msi2
1421232_at 0.98395 plexin A1 Plxna1
1421233_at 0.87315 Pbx/knotted 1 homeobox Pknox1
1421234_at 0.91125 transcription factor 1 Tcf1
1421235_s_at 0.85220 RecQ protein-like 5 Recql5
1421236_at 1.12725 receptor (TNFRSF)-interacting serine-threonine kinase 2 Ripk2
1421237_at 0.83984 thymopoietin Tmpo
1421238_a_at 0.84454 "protein tyrosine phosphatase, receptor type, T" Ptprt
1421239_at 0.91015 interleukin 6 signal transducer Il6st
1421240_at 0.98254 endoplasmic reticulum (ER) to nucleus signalling 1 Ern1
1421241_at 1.08104 "nerve growth factor receptor (TNFR superfamily, member 16)" Ngfr
1421242_at 1.13499 ring finger protein 144 Rnf144
1421243_at 0.92698 ring finger protein 144 Rnf144
1421244_at 1.06545 estrogen receptor 1 (alpha) Esr1
1421245_at 0.93400 sclerostin Sost
1421246_at 0.98404 paired box gene 9 Pax9
1421247_at 1.10879 paired box gene 9 Pax9
1421248_at 0.95686 synapsin III Syn3
1421249_at 1.10415 "similar to solute carrier family 39 (zinc transporter), member 9" LOC674308
1421250_at 0.99302 "similar to solute carrier family 39 (zinc transporter), member 9" LOC674308
1421251_at 0.93853 zinc finger protein 40 Zfp40
1421252_a_at 0.81696 myocyte enhancer factor 2A Mef2a
1421253_at 1.11011 nebulin-related anchoring protein Nrap
1421254_a_at 0.98435 "sarcoglycan, gamma (dystrophin-associated glycoprotein)" Sgcg
1421255_a_at 0.98904 calcium binding protein 1 Cabp1
1421256_at 1.04587 granzyme C Gzmc
1421257_at 1.08024 "phosphatidylinositol glycan anchor biosynthesis, class B" Pigb
1421258_a_at 0.91699 pyruvate kinase liver and red blood cell Pklr
1421259_at 0.92953 pyruvate kinase liver and red blood cell Pklr
1421260_a_at 1.04523 spermidine synthase Srm
1421261_at 0.94329 "lipase, endothelial" Lipg
1421262_at 1.12562 "lipase, endothelial" Lipg
1421263_at 0.99301 "gamma-aminobutyric acid (GABA-A) receptor, subunit alpha 3" Gabra3
1421264_at 0.97541 butyrophilin related 1 Butr1
1421265_a_at 1.06472 RNA binding motif protein 38 Rbm38
1421266_s_at 1.09531 "nuclear factor of kappa light chain gene enhancer in B-cells inhibitor, beta"
Nfkbib

1421267_a_at 0.90660 "Cbp/p300-interacting transactivator, with Glu/Asp-rich carboxy-terminal domain, 2" Cited2

1421268_at 1.00546 UDP-glucose ceramide glucosyltransferase Ugcg

1421269_at 1.11828 UDP-glucose ceramide glucosyltransferase Ugcg

1421270_at 1.10341 SH3 domain containing ring finger 1 Sh3rf1

1421271_at 1.09247 SH3 domain containing ring finger 1 Sh3rf1

1421272_at 0.94884 Vav2 oncogene Vav2

1421273_at 0.95423 suppressor of cytokine signaling 4 Socs4

1421274_at 0.99918 suppressor of cytokine signaling 4 Socs4

1421275_s_at 0.92498 suppressor of cytokine signaling 4 Socs4

1421276_a_at 0.97921 dystonin Dst

1421277_at 0.64078 spectrin alpha 1 /// similar to spectrin alpha 1 Spna1 /// LOC630963

1421278_s_at 0.94835 similar to spectrin alpha 1 LOC630963

1421279_at 1.22320 "laminin, gamma 2" Lamc2

1421280_at 1.03673 "gamma-aminobutyric acid (GABA-A) receptor, subunit alpha 1" Gabra1

1421281_at 1.10628 "gamma-aminobutyric acid (GABA-A) receptor, subunit alpha 1" Gabra1

1421282_at 1.06589 bone morphogenetic protein 5 Bmp5

1421283_at 0.97084 bone morphogenetic protein 5 Bmp5

1421284_at 0.92452 "phosphatidylinositol glycan anchor biosynthesis, class N" Pign

1421285_at 1.06677 phosphoinositide-3-kinase adaptor protein 1 Pik3ap1

1421286_a_at 0.87113 "ATPase, H+/K+ exchanging, gastric, alpha polypeptide" Atp4a

1421287_a_at 0.95087 platelet/endothelial cell adhesion molecule 1 Pecam1

1421288_at 0.90899 coagulation factor II (thrombin) receptor-like 3 F2rl3

1421289_at 1.05674 "heat shock protein family, member 7 (cardiovascular)" Hspb7

1421290_at 0.97679 "heat shock protein family, member 7 (cardiovascular)" Hspb7

1421291_at 1.01125 interleukin 18 receptor accessory protein Il18rap

1421292_a_at 0.89268 RIKEN cDNA A730008L03 gene A730008L03Rik

1421293_at 1.01567 hepatoma derived growth factor-like 1 Hdgfl1

1421294_at 0.81060 hepatoma derived growth factor-like 1 Hdgfl1

1421295_at 0.92653 chordin-like 1 Chrdl1

1421296_at 1.22070 "tumor necrosis factor receptor superfamily, member 10b" Tnfrsf10b

1421297_a_at 0.84397 "calcium channel, voltage-dependent, L type, alpha 1C subunit" Cacna1c

1421298_a_at 0.87302 homeodomain interacting protein kinase 1 Hipk1

1421299_a_at 0.92304 lymphoid enhancer binding factor 1 Lef1

1421300_at 0.92907 "adenosine deaminase, RNA-specific, B2" Adarb2

1421301_at 1.03593 Zic finger protein of the cerebellum 2 Zic2

1421302_a_at 0.91592 "guanine nucleotide binding protein, alpha 15" Gna15

1421303_at 1.01516 "zinc finger protein, subfamily 1A, 1 (Ikaros)" Zfpn1a1

1421304_at 1.06277 "killer cell lectin-like receptor, subfamily A, member 2" Klra2

1421305_x_at 1.00364 "rabaptin, RAB GTPase binding effector protein 1" Rabep1

1421306_a_at 1.02067 histone deacetylase 9 Hdac9

1421307_at 1.13040 carbonic anhydrase 13 Car13

1421308_at 1.04355 carbonic anhydrase 13 Car13

1421309_at 0.98440 O-6-methylguanine-DNA methyltransferase Mgmt
 1421310_at 0.94718 RIKEN cDNA 4921517D21 gene 4921517D21Rik
 1421311_at 0.90649 protocadherin 10 Pcdh10
 1421312_a_at 0.88987 kinesin family member C2 Kifc2
 1421313_s_at 1.02774 cortactin Ctn
 1421314_at 1.10364 cortactin Ctn
 1421315_s_at 0.97751 cortactin Ctn
 1421316_at 0.87496 RIKEN cDNA 1110058A15 gene 1110058A15Rik
 1421317_x_at 0.95994 myeloblastosis oncogene Myb
 1421318_at 0.92133 N-deacetylase/N-sulfotransferase (heparin glucosaminyl) 4 Ndst4
 1421319_at 0.87912 prostaglandin F2 receptor negative regulator Ptgfrn
 1421320_a_at 1.06435 queuine tRNA-ribosyltransferase domain containing 1 Qtrtd1
 1421321_a_at 0.93480 neuroepithelial cell transforming gene 1 Net1
 1421322_a_at 0.99704 interferon dependent positive acting transcription factor 3 gamma Isgf3g
 1421323_a_at 1.06981 RIKEN cDNA E430034L04 gene E430034L04Rik
 1421324_a_at 0.93485 thymoma viral proto-oncogene 2 Akt2
 1421325_at 1.10247 "phospholipase A2, group IIF" Pla2g2f
 1421326_at 1.03267 "colony stimulating factor 2 receptor, beta 1, low-affinity (granulocyte-macrophage)" Csf2rb1
 1421327_at 0.82817 microtubule-associated protein 2 /// hypothetical protein A730034C02
 Mtap2 /// A730034C02
 1421328_at 0.86946 microtubule-associated protein 2 /// hypothetical protein A730034C02
 Mtap2 /// A730034C02
 1421329_a_at 0.92987 SET and MYND domain containing 1 Smyd1
 1421330_at 1.08398 "protein tyrosine phosphatase, non-receptor type 4" Ptpn4
 1421331_at 0.83832 heparan sulfate (glucosamine) 3-O-sulfotransferase 3B1 Hs3st3b1
 1421332_at 0.94561 heparan sulfate (glucosamine) 3-O-sulfotransferase 3B1 Hs3st3b1
 1421333_a_at 0.86967 myoneurin Mynn
 1421334_x_at 0.85859 myoneurin Mynn
 1421335_a_at 0.99247 EGF-like domain 7 Egfl7
 1421336_at 0.98797 prospero-related homeobox 1 Prox1
 1421337_at 0.90607 E74-like factor 4 (ets domain transcription factor) Elf4
 1421338_at 1.01589 E74-like factor 4 (ets domain transcription factor) /// hypothetical protein
 LOC676319 Elf4 /// LOC676319
 1421339_at 1.07934 exostoses (multiple)-like 3 Extl3
 1421340_at 0.66197 mitogen activated protein kinase kinase kinase 5 /// similar to mitogen activated
 protein kinase kinase kinase 5 Map3k5 /// LOC675366
 1421341_at 0.91566 axin2 Axin2
 1421342_at 1.09981 "K+ voltage-gated channel, subfamily S, 2" Kcns2
 1421343_at 1.02720 BTB and CNC homology 2 Bach2
 1421344_a_at 1.03301 ajuba Jub
 1421345_at 0.96473 lecithin-retinol acyltransferase (phosphatidylcholine-retinol-O-acyltransferase) Lrat

1421346_a_at 1.20363 "solute carrier family 6 (neurotransmitter transporter, taurine), member 6" Slc6a6

1421347_at 0.94149 ferritin mitochondrial Ftmt

1421348_a_at 1.05217 cell cycle exit and neuronal differentiation 1 Cend1

1421349_x_at 1.00323 cell cycle exit and neuronal differentiation 1 Cend1

1421350_a_at 0.85527 glutamate receptor interacting protein 1 Grip1

1421351_at 1.08615 "glutamate receptor, ionotropic, AMPA4 (alpha 4)" Gria4

1421352_at 0.93919 toll-like receptor 6 Tlr6

1421353_at 1.03917 phosphodiesterase 7B Pde7b

1421354_at 0.92236 "protein kinase, cGMP-dependent, type II" Prkg2

1421355_at 0.97651 "transglutaminase 3, E polypeptide" Tgm3

1421356_at 1.06655 testis expressed gene 9 Tex9

1421357_at 1.02364 "general transcription factor II A, 1" Gtf2a1

1421358_at 1.10300 "histocompatibility 2, M region locus 3" H2-M3

1421359_at 1.02397 ret proto-oncogene Ret

1421360_at 0.92122 "inositol polyphosphate-4-phosphatase, type I" Inpp4a

1421361_at 0.89687 G protein-coupled receptor kinase 1 Grk1

1421362_a_at 1.07128 fyn-related kinase Frk

1421363_at 0.95058 "cytochrome P450, family 2, subfamily c, polypeptide 39" Cyp2c39

1421364_at 0.84329 leucine rich repeat and fibronectin type III domain containing 1 Lfn1

1421365_at 0.93067 follistatin Fst

1421366_at 0.90620 "C-type lectin domain family 5, member a" Clec5a

1421367_at 0.87841 RIKEN cDNA 4930549C01 gene 4930549C01Rik

1421368_at 1.04626 "scratch homolog 1, zinc finger protein (Drosophila)" Sqrt1

1421369_a_at 0.88891 mab-21-like 1 (C. elegans) Mab21l1

1421370_a_at 0.82083 "interleukin 1 family, member 5 (delta)" Il1f5

1421371_at 0.95571 --- ---

1421372_at 1.03799 kallikrein 1-related peptidase b4 Klk1b4

1421373_at 1.00951 cytochrome c oxidase subunit IV isoform 2 Cox4i2

1421374_a_at 0.86174 FXFD domain-containing ion transport regulator 1 Fxyd1

1421375_a_at 1.12175 S100 calcium binding protein A6 (calcylin) S100a6

1421376_at 1.13966 Tnf receptor-associated factor 6 Traf6

1421377_at 0.76199 Tnf receptor-associated factor 6 Traf6

1421378_s_at 1.14766 "ATP-binding cassette, sub-family C (CFTR/MRP), member 1" Abcc1

1421379_at 0.96383 zinc finger protein 354B Zfp354b

1421380_at 0.91513 insulin receptor Insr

1421381_a_at 0.98103 "procollagen, type IX, alpha 1" Col9a1

1421382_at 0.93795 prolactin receptor Prlr

1421383_at 1.16744 coiled-coil domain containing 82 Ccdc82

1421384_at 0.66355 lysosomal trafficking regulator Lyst

1421385_a_at 0.63885 myosin VIIa Myo7a

1421386_at 1.18450 ankyrin repeat domain 6 Ankrd6

1421387_at 1.02880 kringle containing transmembrane protein 1 Kremen1

1421388_at 0.92872 myocyte enhancer factor 2D Mef2d
1421389_a_at 0.96116 eukaryotic translation initiation factor 2 alpha kinase 4 Eif2ak4
1421390_at 0.99451 "solute carrier family 12, member 1" Slc12a1
1421391_at 1.07612 vasoactive intestinal peptide receptor 2 Vipr2
1421392_a_at 1.20644 baculoviral IAP repeat-containing 3 Birc3
1421393_at 0.99553 "glutamate receptor, ionotropic, NMDA2D (epsilon 4)" Grin2d
1421394_a_at 0.87611 baculoviral IAP repeat-containing 4 Birc4
1421395_at 0.92524 zinc finger protein interacting with K protein 1 Zik1
1421396_at 0.84357 proprotein convertase subtilisin/kexin type 1 Pcsk1
1421397_a_at 1.16216 leucine-rich and death domain containing Lrdd
1421398_at 1.01581 tripartite motif protein 7 Trim7
1421399_at 0.99758 insulinoma-associated 1 Insm1
1421400_at 1.11572 "potassium large conductance calcium-activated channel, subfamily M, beta member 1" Kcnmb1
1421401_at 0.96040 "Potassium large conductance calcium-activated channel, subfamily M, beta member 1" Kcnmb1
1421402_at 0.84516 metastasis associated 3 Mta3
1421403_at 1.16836 peptidase inhibitor 15 Pi15
1421404_at 1.08856 chemokine (C-X-C motif) ligand 15 Cxcl15
1421405_at 0.99303 "zinc finger, imprinted 1" Zim1
1421406_at 0.95686 --- ---
1421407_at 1.09950 coagulation factor II (thrombin) receptor-like 2 F2rl2
1421408_at 1.07983 "immunoglobulin superfamily, member 6" Igsf6
1421409_at 1.04654 Musashi homolog 1(Drosophila) Msi1
1421410_a_at 0.93336 proline-serine-threonine phosphatase-interacting protein 2 Pstpip2
1421411_at 1.14571 proline-serine-threonine phosphatase-interacting protein 2 Pstpip2
1421412_at 0.98243 gooseoid Gsc
1421413_a_at 0.86799 PDZ and LIM domain 5 /// similar to PDZ and LIM domain protein 5 (Enigma homolog) (Enigma-like PDZ and LIM domains protein) Pdlim5 /// LOC669660
1421414_a_at 0.92486 "sema domain, transmembrane domain (TM), and cytoplasmic domain, (semaphorin) 6A" Sema6a
1421415_s_at 0.89229 "glucosaminyl (N-acetyl) transferase 2, I-branching enzyme" Gcnt2
1421416_at 0.96849 mitogen activated protein kinase kinase 7 Map2k7
1421417_s_at 1.13297 elaC homolog 1 (E. coli) Elac1
1421418_a_at 0.91789 pregnancy specific glycoprotein 19 Psg19
1421419_at 0.90384 "potassium channel, subfamily K, member 4" Kcnk4
1421420_at 1.02336 chemokine (C-C motif) receptor 10 Ccr10
1421421_at 0.95468 angiopoietin-like 1 Angpt1
1421422_at 0.89174 cDNA sequence AF397014 AF397014
1421423_at 0.88512 prolactin-like protein O Prlpo
1421424_a_at 0.82477 alanyl (membrane) aminopeptidase Anpep
1421425_a_at 0.92231 Down syndrome critical region gene 1-like 1 Dscr111
1421426_at 0.99152 Hedgehog-interacting protein Hhip

1421427_at	0.98670	"procollagen, type XXV, alpha 1"	Col25a1
1421428_at	1.09780	"solute carrier family 5 (choline transporter), member 7"	Slc5a7
1421429_a_at	0.71617	nephronectin	Npnt
1421430_at	0.96375	RAD51-like 1 (S. cerevisiae)	Rad5111
1421431_at	1.07243	polymerase I and transcript release factor	Ptrf
1421432_at	0.93698	ADP-ribosylation factor-like 10	Arl10
1421433_at	0.89336	zinc finger homeodomain 4	Zfhx4
1421434_at	1.01933	"dynein, axonemal, heavy chain 5"	Dnahc5
1421435_at	1.06025	"glutamate receptor, ionotropic, delta 2"	Grid2
1421436_at	0.94296	"glutamate receptor, ionotropic, delta 2"	Grid2
1421437_x_at	0.90448	protocadherin beta 14	Pcdhb14
1421438_at	1.03269	---	---
1421439_at	1.08513	wingless related MMTV integration site 8b	Wnt8b
1421440_at	0.91275	similar to wingless related MMTV integration site 8b	LOC677060
1421441_at	0.91840	angiopoietin 1	Angpt1
1421442_at	1.28954	FMS-like tyrosine kinase 4	Flt4
1421443_at	0.99144	G protein-coupled receptor 110	Gpr110
1421444_at	0.89884	progesterone receptor	Pgr
1421445_at	1.01042	"solute carrier family 26, member 3"	Slc26a3
1421446_at	1.11121	"protein kinase C, gamma"	Prkcc
1421447_at	1.02377	"one cut domain, family member 1"	Onecut1
1421448_at	0.79597	GTPase activating RANGAP domain-like 1	Garnl1
1421449_at	1.06072	CUB and Sushi multiple domains 1	Csmd1
1421450_a_at	0.97383	mitogen activated protein kinase kinase kinase 4	Map3k4
1421451_at	0.96751	crumbs homolog 1 (Drosophila)	Crb1
1421452_at	0.85401	cell adhesion molecule-related/down-regulated by oncogenes	Cdon
1421453_at	0.98780	junctionophilin 2	Jph2
1421454_at	0.89322	zinc finger protein 108	Zfp108
1421455_at	0.84985	"syntrophin, basic 1"	Sntb1
1421456_at	0.80503	"purinergic receptor P2Y, G-protein coupled 1"	P2ry1
1421457_a_at	0.90882	"SAM domain, SH3 domain and nuclear localization signals, 1"	Samsn1
1421458_at	0.90301	zinc finger protein 112	Zfp112
1421459_a_at	0.97470	"low density lipoprotein receptor-related protein 8, apolipoprotein e receptor"	Lrp8
1421460_at	0.92742	desmocollin 1	Dsc1
1421461_at	0.99956	myeloproliferative leukemia virus oncogene	Mpl
1421462_a_at	0.83733	leprecan 1	Lepre1
1421463_at	1.05715	sialic acid binding Ig-like lectin E	Siglece
1421464_at	0.95539	bagpipe homeobox gene 1 homolog (Drosophila)	Bapx1
1421465_at	0.68140	wingless related MMTV integration site 2b	Wnt2b
1421466_at	1.02936	ankyrin repeat and SOCS box-containing protein 10	Asb10
1421467_at	0.88612	runt related transcription factor 3	Runx3
1421468_at	0.86445	"potassium inwardly-rectifying channel, subfamily J, member 3"	Kcnj3

1421469_a_at 0.98031 signal transducer and activator of transcription 5A Stat5a
1421470_at 1.02532 gastrin releasing peptide receptor Grpr
1421471_at 1.04320 neuropeptide Y receptor Y1 Npy1r
1421472_at 1.02741 RIKEN cDNA 2900083I11 gene 2900083I11Rik
1421473_at 0.89519 interleukin 1 alpha Il1a
1421474_a_at 0.90407 splA/ryanodine receptor domain and SOCS box containing 1 Spsb1
1421475_at 0.96578 "membrane-spanning 4-domains, subfamily A, member 2" Ms4a2
1421476_a_at 0.91465 calcium activated nucleotidase 1 Cant1
1421477_at 0.92550 complexin 2 Cplx2
1421478_a_at 1.00316 zinc finger protein 318 Zfp318
1421479_at 0.89881 zinc finger protein 318 Zfp318
1421480_a_at 1.02000 "adenosine deaminase, RNA-specific, B1" Adarb1
1421481_at 0.96306 "tumor necrosis factor receptor superfamily, member 9" Tnfrsf9
1421482_at 0.89942 "Bartter syndrome, infantile, with sensorineural deafness (Barttin)" Bsnd
1421483_at 1.09527 LIM homeobox protein 4 Lhx4
1421484_at 0.98725 "UDP glucuronosyltransferase 2 family, polypeptide A1 /// UDP
glucuronosyltransferase 2 family, polypeptide A2" Ugt2a1 /// Ugt2a2
1421485_at 1.05055 tau tubulin kinase 2 Ttbk2
1421486_at 0.92898 early growth response 3 Egr3
1421487_a_at 1.07438 non-catalytic region of tyrosine kinase adaptor protein 1 Nck1
1421488_at 1.00012 RAB GTPase activating protein 1-like Rabgap11
1421489_a_at 1.00430 RIKEN cDNA 2010106E10 gene 2010106E10Rik
1421490_at 0.92928 proline-rich protein MP5 /// cDNA sequence BC061494 Prpmp5 /// BC061494
1421491_a_at 0.92121 transmembrane protein 49 Tmem49
1421492_at 1.06059 "prostaglandin D2 synthase 2, hematopoietic" Ptgds2
1421493_a_at 0.96118 regulator of G-protein signaling 20 Rgs20
1421494_at 0.90217 RIKEN cDNA 1110033F04 gene 1110033F04Rik
1421495_a_at 1.04619 RIKEN cDNA 1700052N19 gene 1700052N19Rik
1421496_at 1.01723 N-acylsphingosine amidohydrolase 3-like Asah3l
1421497_at 0.89454 glycoprotein hormone alpha 2 Gpha2
1421498_a_at 1.31511 RIKEN cDNA 2010204K13 gene 2010204K13Rik
1421499_a_at 1.06987 "protein tyrosine phosphatase, non-receptor type 14" Ptpn14
1421500_at 1.00457 steroid sulfatase Sts
1421501_a_at 0.86818 ELKS/RAB6-interacting/CAST family member 1 Erc1
1421502_at 0.96738 ubiquitin specific peptidase 26 Usp26
1421503_at 0.79616 protocadherin 15 Pcdh15
1421504_at 0.98478 trans-acting transcription factor 4 Sp4
1421505_at 0.96142 Mix1 homeobox-like 1 (Xenopus laevis) Mixl1
1421506_at 0.98589 olfactory receptor 78 Olfr78
1421507_at 1.01057 olfactory receptor 78 Olfr78
1421508_at 0.91355 odd Oz/ten-m homolog 1 (Drosophila) Odz1
1421509_at 0.96094 VPS10 domain receptor protein SORCS 1 Sorcs1
1421510_at 0.96959 "protease, serine, 7 (enterokinase)" Prss7

1421511_at 0.94241 integrin beta 3 Itgb3
1421512_at 0.97700 centrosomal protein 250 Cep250
1421513_at 0.95908 tudor domain containing 1 Tdrd1
1421514_a_at 0.96087 sex comb on midleg-like 2 (Drosophila) Scml2
1421515_at 0.90303 "nuclear receptor subfamily 6, group A, member 1" Nr6a1
1421516_at 0.98290 "nuclear receptor subfamily 6, group A, member 1" Nr6a1
1421517_at 1.01228 "ST6 (alpha-N-acetyl-neuraminyl-2,3-beta-galactosyl-1,3)-N-acetylgalactosaminide alpha-2,6-sialyltransferase 1" St6galnac1
1421518_at 1.11829 "K+ voltage-gated channel, subfamily S, 1" Kcns1
1421519_a_at 0.89156 zinc finger protein 120 Zfp120
1421520_at 0.95472 junctophilin 1 Jph1
1421521_at 0.99035 RIKEN cDNA 4930430A15 gene 4930430A15Rik
1421522_at 0.99608 "beta-1,4-N-acetyl-galactosaminyl transferase 2" B4galnt2
1421523_at 1.07270 fibroblast growth factor 17 Fgf17
1421524_at 0.95151 "cripto, FRL-1, cryptic family 1" Cfc1
1421525_a_at 1.03583 baculoviral IAP repeat-containing 1e Birc1e
1421526_at 1.03119 tolloid-like 2 Tll2
1421527_at 0.92504 Eph receptor A6 Epha6
1421528_a_at 0.92118 surfeit gene 5 Surf5
1421529_a_at 0.97644 thioredoxin reductase 1 Txnrd1
1421530_a_at 1.11235 "glutamate receptor, metabotropic 8" Grm8
1421531_at 1.03261 A kinase (PRKA) anchor protein 3 Akap3
1421532_at 0.94099 relaxin/insulin-like family peptide receptor 2 Rxfp2
1421533_at 1.02881 "solute carrier family 7 (cationic amino acid transporter, y+ system), member 1" Slc7a1
1421534_at 0.89665 "deafness, autosomal dominant 5 homolog (human)" Dfna5h
1421535_a_at 1.03689 "phosphodiesterase 4A, cAMP specific" Pde4a
1421536_at 0.97630 "gamma-aminobutyric acid (GABA-A) receptor, subunit theta" Gabrq
1421537_at 0.85807 homeo box D3 Hoxd3
1421538_at 1.00332 "potassium voltage-gated channel, Shal-related family, member 1" Kcnd1
1421539_at 1.16747 zinc finger protein of the cerebellum 4 Zic4
1421540_at 0.93126 testicular cell adhesion molecule 1 Tcam1
1421541_a_at 0.86606 myocyte enhancer factor 2B Mef2b
1421542_at 0.91983 DAZ interacting protein 1 Dzip1
1421543_at 0.96621 F-box only protein 4 Fbxo4
1421544_at 0.92101 "dynein, axonemal, heavy chain 11" Dnahc11
1421545_a_at 0.88049 synaptic nuclear envelope 1 Syne1
1421546_a_at 0.92722 Rac GTPase-activating protein 1 Racgap1
1421547_at 0.97122 CD180 antigen Cd180
1421548_at 0.97844 protocadherin beta 2 Pcdhb2
1421549_at 1.02584 protocadherin beta 15 Pcdhb15
1421550_a_at 0.80623 tripartite motif protein 34 /// similar to Tripartite motif protein 34 /// similar to Tripartite motif protein 34 Trim34 /// LOC434218 /// LOC640746

1421551_s_at 1.33871 interferon activated gene 202B Ifi202b
1421552_at 0.94369 caudal type homeo box 4 Cdx4
1421553_at 0.99939 "epilepsy, progressive myoclonic epilepsy, type 2 gene alpha" Epm2a
1421554_at 1.07450 LIM homeobox transcription factor 1 alpha Lmx1a
1421555_at 0.91395 "adrenergic receptor, beta 3" Adrb3
1421556_at 0.95405 "serine (or cysteine) peptidase inhibitor, clade A, member 3A" Serpina3a
1421557_x_at 0.98528 thioredoxin 2 Txn2
1421558_at 0.91263 brachyury 2 T2
1421559_at 1.02803 "ABO blood group (transferase A, alpha 1-3-N-acetylgalactosaminyltransferase, transferase B, alpha 1-3-galactosyltransferase)" Abo
1421560_at 0.86946 snail homolog 3 (Drosophila) Snai3
1421561_at 1.02137 "basic helix-loop-helix domain containing, class B4" Bhlhb4
1421562_at 0.98093 CD209c antigen Cd209c
1421563_at 0.94470 "ATPase, Cu++ transporting, beta polypeptide" Atp7b
1421564_at 1.15721 "serine (or cysteine) peptidase inhibitor, clade A, member 3C" Serpina3c
1421565_at 1.08787 roundabout homolog 3 (Drosophila) Robo3
1421566_at 0.88339 plasmacytoma expressed transcript 2 Pet2
1421567_at 0.93532 neuronal PAS domain protein 3 Npas3
1421568_at 1.05815 "potassium voltage-gated channel, shaker-related, subfamily, member 6" Kcna6
1421569_at 1.10023 "glutamate receptor, ionotropic, delta 1" Grid1
1421570_at 0.89822 interleukin 9 receptor Il9r
1421571_a_at 1.17632 "lymphocyte antigen 6 complex, locus C" Ly6c
1421572_at 0.99554 "hypoxia inducible factor 3, alpha subunit" Hif3a
1421573_at 1.03075 RIKEN cDNA 4930533N22 gene 4930533N22Rik
1421574_at 0.92945 RAS related protein 2a Rap2a
1421575_at 1.01192 RIKEN cDNA 2310057N15 gene 2310057N15Rik
1421576_at 1.00779 RIKEN cDNA 4930524B15 gene 4930524B15Rik
1421577_at 0.85831 even skipped homeotic gene 2 homolog Evx2
1421578_at 0.92071 chemokine (C-C motif) ligand 4 Ccl4
1421579_at 1.01954 homeo box A9 Hoxa9
1421580_at 1.05725 contactin associated protein 1 Cntnap1
1421581_at 1.10592 "potassium inwardly-rectifying channel, subfamily J, member 6" Kcnj6
1421582_a_at 0.90124 cAMP responsive element binding protein 1 Creb1
1421583_at 1.06861 cAMP responsive element binding protein 1 Creb1
1421584_at 0.96899 opsin 4 (melanopsin) Opn4
1421585_at 0.96794 RIKEN cDNA 4933439F11 gene 4933439F11Rik
1421586_a_at 0.93257 "CD46 antigen, complement regulatory protein" Cd46
1421587_at 1.05012 kallikrein 1-related peptidase b27 Klk1b27
1421588_at 0.96824 "tumor necrosis factor (ligand) superfamily, member 14" Tnfsf14
1421589_at 0.98118 keratin 31 Krt31
1421590_at 0.95952 "MAS-related GPR, member H" Mrgprh
1421591_at 0.92132 "cylicin, basic protein of sperm head cytoskeleton 1" Cylc1
1421592_at 1.06722 neural cell adhesion molecule 2 Ncam2

1421593_at 0.96237 neural cell adhesion molecule 2 Ncam2
1421594_a_at 0.70155 synaptotagmin-like 2 Syt12
1421595_at 0.93554 RIKEN cDNA 9630031F12 gene 9630031F12Rik
1421596_s_at 0.96403 histocompatibility 28 H28
1421597_a_at 0.86082 "homeo box, msh-like 3" Msx3
1421598_at 0.96947 --- ---
1421599_at 0.93800 heparan sulfate 6-O-sulfotransferase 3 Hs6st3
1421600_a_at 0.98784 tripartite motif protein 26 Trim26
1421601_at 0.95775 genomic screened homeo box 2 Gsh2
1421602_at 1.01094 sex hormone binding globulin Shbg
1421603_a_at 1.04215 CEA-related cell adhesion molecule 2 Ceacam2
1421604_a_at 0.90788 Kruppel-like factor 3 (basic) Klf3
1421605_a_at 1.00470 aquaporin 9 Aqp9
1421606_a_at 1.18348 "sulfotransferase family 4A, member 1" Sult4a1
1421607_at 0.99798 vestigial like 1 homolog (Drosophila) Vgl1
1421608_at 1.20476 interleukin 20 Il20
1421609_a_at 0.94407 camello-like 3 Cml3
1421610_at 1.03928 chemokine (C-X3-C motif) ligand 1 Cx3cl1
1421611_at 0.89206 "hypothetical protein, MNCb-2457" AB041806
1421612_a_at 1.07465 "H2A histone family, member Y3 /// H2A histone family, member Y2"
H2afy3 /// H2afy2
1421613_at 1.06673 "H2A histone family, member Y3" H2afy3
1421614_at 0.86951 zonadhesin Zan
1421615_at 1.12543 myosin XV Myo15
1421616_at 1.05471 "glutamate receptor, ionotropic, NMDA2A (epsilon 1)" Grin2a
1421617_at 1.13247 "transient receptor potential cation channel, subfamily M, member 8" Trpm8
1421618_at 0.91156 myosin IF Myo1f
1421619_at 0.93308 "potassium voltage-gated channel, subfamily H (eag-related), member 3" Kcnh3
1421620_at 1.02197 "interleukin 5 receptor, alpha" Il5ra
1421621_at 0.82207 RAS protein-specific guanine nucleotide-releasing factor 2 Rasgrf2
1421622_a_at 0.97439 Rap guanine nucleotide exchange factor (GEF) 4 Rapgef4
1421623_at 1.10999 "interleukin 12 receptor, beta 2" Il12rb2
1421624_a_at 1.05420 enabled homolog (Drosophila) Enah
1421625_a_at 0.96000 transmembrane channel-like gene family 1 Tmc1
1421626_at 0.88357 transmembrane channel-like gene family 1 Tmc1
1421627_at 0.94192 even skipped homeotic gene 1 homolog Evx1
1421628_at 1.01347 interleukin 18 receptor 1 Il18r1
1421629_at 0.92921 "gamma-aminobutyric acid (GABA-A) receptor, subunit epsilon" Gabre
1421630_at 1.00255 "zinc finger protein 1, Y linked /// zinc finger protein 2, Y linked" Zfy1 /// Zfy2
1421631_at 0.97959 protocadherin beta 1 Pcdhb1
1421632_at 0.87911 "potassium intermediate/small conductance calcium-activated channel,
subfamily N, member 3" Kcnn3
1421633_a_at 0.96046 hyaluronan and proteoglycan link protein 1 Hapln1

1421634_at 1.00209 calpain 8 Capn8
1421635_at 0.83850 cyclin M4 Cnnm4
1421636_at 0.97058 RIKEN cDNA 9030605E16 gene 9030605E16Rik
1421637_at 0.95172 "solute carrier family 5, member 4a" Slc5a4a
1421638_at 1.01787 pregnancy specific glycoprotein 17 Psg17
1421639_at 0.93634 RIKEN cDNA 1700082M22 gene 1700082M22Rik
1421640_a_at 1.03486 TRAF family member-associated Nf-kappa B activator Tank
1421641_at 0.88386 "solute carrier family 6 (neurotransmitter transporter, noradrenalin), member 2" Slc6a2
1421642_a_at 1.07208 cysteinyl leukotriene receptor 2 Cysltr2
1421643_at 0.98372 "zinc finger protein, subfamily 1A, 2 (Helios)" Zfpn1a2
1421644_at 1.09518 schlafen 3 Slfn3
1421645_at 1.02963 RIKEN cDNA 9930013L23 gene 9930013L23Rik
1421646_a_at 0.88531 protein inhibitor of activated STAT 3 Pias3
1421647_at 0.98512 CD1d2 antigen Cd1d2
1421648_at 1.01143 neuroligin 1 Nlgn1
1421649_at 0.93587 "tumor necrosis factor receptor superfamily, member 8" Tnfrsf8
1421650_at 0.96901 matrix extracellular phosphoglycoprotein with ASARM motif (bone) Mepe
1421651_at 1.07557 ankyrin repeat domain 22 Ankrd22
1421652_at 1.00458 5-hydroxytryptamine (serotonin) receptor 3B Htr3b
1421653_a_at 0.96986 immunoglobulin heavy chain (J558 family) /// similar to immunoglobulin heavy chain variable region /// similar to immunoglobulin mu-chain /// similar to anti-poly(dC) monoclonal antibody heavy chain /// similar to Ig heavy chain V region 23 precursor /// similar to Ig heavy chain V region 102 precursor /// similar to Ig heavy chain V region VH558 A1/A4 precursor Igh-VJ558 /// LOC238447 /// LOC544903 /// LOC544907 /// LOC633774 /// LOC676478 /// LOC677563
1421654_a_at 1.00245 lamin A Lmna
1421655_a_at 0.92023 chemokine (C-C motif) receptor 4 Ccr4
1421656_at 0.98460 sprouty homolog 2 (Drosophila) Spry2
1421657_a_at 0.90766 SRY-box containing gene 17 Sox17
1421658_x_at 0.99818 pregnancy specific glycoprotein 18 Psg18
1421659_at 0.92976 "adrenergic receptor, alpha 1a" Adra1a
1421660_at 0.99390 "sodium channel, voltage-gated, type IX, alpha /// similar to sodium channel 25" Scn9a /// LOC671835
1421661_at 1.02473 formyl peptide receptor-like 1 Fprl1
1421662_a_at 0.91403 tumor suppressor candidate 3 Tusc3
1421663_at 1.00475 dehydrogenase/reductase member 2 Dhrs2
1421664_a_at 1.10419 phosphoserine/threonine/tyrosine interaction protein Styx
1421665_a_at 1.01437 gonadotropin releasing hormone receptor Gnhr
1421666_a_at 0.94616 arylalkylamine N-acetyltransferase Aanat
1421667_at 0.82737 neuromedin U receptor 1 Nmur1
1421668_x_at 0.99319 spermatogenesis associated glutamate (E)-rich protein 3 Speer3
1421669_at 0.97586 "sulfotransferase family 3A, member 1" Sult3a1
1421670_a_at 0.89546 interleukin-1 receptor-associated kinase 4 Irak4

1421671_at 1.10243 hydroxysteroid (17-beta) dehydrogenase 3 Hsd17b3
1421672_at 0.97678 interleukin 17A Il17a
1421673_s_at 0.94741 syntaxin 1B2 /// syntaxin 1B1 Stx1b2 /// Stx1b1
1421674_at 1.01144 --- ---
1421675_at 0.97277 RIKEN cDNA 1700123K08 gene 1700123K08Rik
1421676_at 1.08237 RIKEN cDNA 4930542N07 gene 4930542N07Rik
1421677_at 1.06302 fibroblast growth factor 20 Fgf20
1421678_at 1.00512 "inositol 1,4,5-triphosphate receptor 2" Itpr2
1421679_a_at 1.13529 cyclin-dependent kinase inhibitor 1A (P21) Cdkn1a
1421680_at 1.04455 RIKEN cDNA A030005L19 gene A030005L19Rik
1421681_at 0.83868 neuregulin 4 Nrg4
1421682_a_at 1.02663 t-complex-associated testis expressed 3 Tcte3
1421683_at 1.08288 t-complex-associated testis expressed 3 Tcte3
1421684_at 1.01418 RIKEN cDNA 2310051M13 gene 2310051M13Rik
1421685_at 1.00403 "C-type lectin domain family 4, member b1" Clec4b1
1421686_at 0.93030 RFamide-related peptide Rfrp
1421687_at 1.12162 beta-microseminoprotein Msmb
1421688_a_at 0.95225 chemokine (C-C motif) ligand 1 Ccl1
1421689_at 0.95626 keratin associated protein 8-2 Krtap8-2
1421690_s_at 1.02172 agouti related protein Agrp
1421691_at 0.93669 keratin associated protein 16-7 Krtap16-7
1421692_at 1.14777 "calcium channel, voltage-dependent, R type, alpha 1E subunit" Cacna1e
1421693_a_at 0.99505 G protein-coupled receptor 98 Gpr98
1421694_a_at 0.91884 chondroitin sulfate proteoglycan 2 Cspg2
1421695_at 0.83334 stereocilin Strc
1421696_at 1.04922 polycystic kidney and hepatic disease 1-like 1 Pkhd11l
1421697_at 0.98472 "Usher syndrome 2A (autosomal recessive, mild) homolog (human)" Ush2a
1421698_a_at 0.93224 "procollagen, type XIX, alpha 1" Col19a1
1421699_at 1.03527 enamel Enam
1421700_at 0.97809 testis expressed gene 16 Tex16
1421701_at 0.93941 "vomeronasal 2, receptor, 16" V2r16
1421702_at 1.05438 retinol dehydrogenase 1 (all trans) Rdh1
1421703_at 0.91203 zinc finger protein 59 Zfp59
1421704_a_at 1.04936 "phosphatidylinositol 3-kinase, C2 domain containing, gamma polypeptide" Pik3c2g
1421705_at 0.96537 "sodium channel, voltage-gated, type III, alpha" Scn3a
1421706_at 0.92131 matrix metalloproteinase 20 (enamelysin) Mmp20
1421707_at 0.88659 transmembrane channel-like gene family 2 Tmc2
1421708_a_at 1.02382 signal transducer and activator of transcription 6 Stat6
1421709_a_at 0.89300 flavin containing monooxygenase 5 Fmo5
1421710_at 0.92001 zinc finger protein 92 Zfp92
1421711_at 0.94578 zinc finger protein 109 Zfp109
1421712_at 1.01038 "selectin, endothelial cell" Sele

1421713_at	1.03425	preferentially expressed antigen in melanoma-like 1	Pramell1
1421714_at	0.94636	"transmembrane protease, serine 3"	Tmprss3
1421715_at	1.02240	pheromone receptor V2Rx	V2rx
1421716_at	1.04923	"vomeronasal 1 receptor, D6"	V1rd6
1421717_at	1.13943	"vomeronasal 1 receptor, D4"	V1rd4
1421718_at	1.24079	striamin	Strm
1421719_at	1.05369	"vomeronasal 2, receptor, 1b"	V2r1b
1421720_a_at	0.99423	deltex 2 homolog (Drosophila)	Dtx2
1421721_a_at	0.90250	aryl hydrocarbon receptor nuclear translocator	Arnt
1421722_at	0.87394	---	---
1421723_at	0.97670	protocadherin beta 18	Pcdhb18
1421724_at	0.97721	"vomeronasal 1 receptor, D7"	V1rd7
1421725_at	0.97054	"solute carrier family 26, member 5"	Slc26a5
1421726_at	0.94685	"adaptor-related protein complex AP-4, beta 1"	Ap4b1
1421727_at	0.89001	eyes absent 1 homolog (Drosophila)	Eya1
1421728_at	0.85956	oligodendrocyte transcription factor 3	Olig3
1421729_a_at	1.06102	"fer (fms/fps related) protein kinase, testis specific 2"	Fert2
1421730_at	1.02000	"nuclear receptor subfamily 5, group A, member 1"	Nr5a1
1421731_a_at	0.95451	flap structure specific endonuclease 1	Fen1
1421732_at	1.11618	glutamine repeat protein 1	Glrp1
1421733_a_at	0.96213	protein-tyrosine sulfotransferase 1	Tpst1
1421734_at	1.00378	---	---
1421735_a_at	0.91731	"ST8 alpha-N-acetyl-neuraminide alpha-2,8-sialyltransferase 5"	St8sia5
1421736_at	0.87637	"vomeronasal 1 receptor, D3"	V1rd3
1421737_at	0.93278	aristaless 4	Alx4
1421738_at	1.01815	"gamma-aminobutyric acid (GABA-A) receptor, subunit alpha 2"	Gabra2
1421739_a_at	1.05299	megakaryocyte-associated tyrosine kinase	Matk
1421740_at	1.04881	"GNAS (guanine nucleotide binding protein, alpha stimulating) complex locus"	Gnas
1421741_at	0.96535	"cytochrome P450, family 3, subfamily a, polypeptide 16"	Cyp3a16
1421742_at	0.88386	guanosine diphosphate (GDP) dissociation inhibitor 2	Gdi2
1421743_a_at	0.91062	poly(rC) binding protein 2	Pcbp2
1421744_at	0.90678	"tumor necrosis factor (ligand) superfamily, member 4"	Tnfsf4
1421745_at	0.95312	pleiomorphic adenoma gene 1	Plag1
1421746_a_at	1.12308	F-box protein 17	Fbxo17
1421747_at	0.93877	estrogen-related receptor gamma	Esrrg
1421748_a_at	1.07656	"tubulin, delta 1"	Tubd1
1421749_at	0.94186	---	---
1421750_a_at	1.00697	von Hippel-Lindau binding protein 1	Vbp1
1421751_a_at	1.11809	"proteasome (prosome, macropain) 26S subunit, non-ATPase, 14"	Psmd14
1421752_a_at	0.94049	"serine (or cysteine) peptidase inhibitor, clade B, member 5"	Serpib5
1421753_a_at	0.99089	LIM homeobox protein 3	Lhx3
1421754_at	0.88327	cDNA sequence AY036118	AY036118

1421755_at 0.89308 G protein-coupled receptor 132 Gpr132
1421756_a_at 0.75255 G protein-coupled receptor 19 Gpr19
1421757_at 1.03439 5-hydroxytryptamine (serotonin) receptor 6 Htr6
1421758_at 1.07675 N-acetyltransferase 1 (arylamine N-acetyltransferase) Nat1
1421759_a_at 0.96213 forkhead box K1 Foxk1
1421760_at 1.21206 pre T-cell antigen receptor alpha Ptcra
1421761_a_at 0.96075 BarH-like homeobox 2 Barx2
1421762_at 1.06534 "potassium inwardly-rectifying channel, subfamily J, member 5" Kcnj5
1421763_at 1.05226 bone morphogenetic protein 10 Bmp10
1421764_at 1.08447 "vomeronasal 1 receptor, D9" V1rd9
1421765_at 0.95399 paired box gene 5 Pax5
1421766_at 1.13378 ubiquitin carboxyl-terminal esterase L4 Uchl4
1421767_at 0.95567 Adenosine kinase Adk
1421768_a_at 1.07327 homer homolog 1 (Drosophila) Homer1
1421769_at 0.93265 LIM homeobox transcription factor 1 beta Lmx1b
1421770_a_at 0.90418 "T-cell leukemia, homeobox 2" Tlx2
1421771_a_at 0.97189 IAP promoted placental gene Ipp
1421772_a_at 0.93853 cytochrome c oxidase subunit VIIa polypeptide 2-like Cox7a2l
1421773_at 1.05329 ets variant gene 2 Etv2
1421774_at 0.87412 ventral anterior homeobox containing gene 1 Vax1
1421775_at 0.95428 "Fc receptor, IgE, high affinity I, alpha polypeptide" Fcer1a
1421776_at 0.96384 olfactory receptor 74 Olfr74
1421777_at 1.06026 olfactory receptor 73 Olfr73
1421778_at 0.96550 "vomeronasal 1, receptor B2" V1rb2
1421779_at 1.00968 Ras-like without CAAX 2 Rit2
1421780_a_at 0.90812 calcium binding protein 5 Cabp5
1421781_at 0.82474 uroplakin 2 Upk2
1421782_a_at 1.00257 submaxillary gland androgen regulated protein 2 Smr2
1421783_a_at 1.07748 Kv channel-interacting protein 2 Kcnip2
1421784_a_at 0.82562 ephrin A4 Efna4
1421785_at 0.91355 UEV and lactate/malate dehydrogenase domains Uevld
1421786_at 1.02378 "protein phosphatase 3, regulatory subunit B, alpha isoform (calcineurin B, type I)" Ppp3r1
1421787_at 0.94294 --- ---
1421788_x_at 0.87533 epidermal growth factor binding protein type B Egfbp2
1421789_s_at 1.05192 ADP-ribosylation factor 3 Arf3
1421790_a_at 0.88159 "potassium voltage-gated channel, shaker-related subfamily, beta member 3" Kcnab3
1421791_at 1.00556 "angiogenin, ribonuclease A family, member 3" Ang3
1421792_s_at 1.05646 triggering receptor expressed on myeloid cells 2 Trem2
1421793_at 0.97252 fibroblast growth factor 11 Fgf11
1421794_at 1.09724 "killer cell lectin-like receptor subfamily C, member 3" Klrc3
1421795_s_at 0.95427 "killer cell lectin-like receptor subfamily C, member 2 /// killer cell lectin-like

receptor subfamily C, member 3" Klrc2 /// Klrc3

1421796_a_at 0.87462 ephrin A5 Efna5
 1421797_a_at 0.99536 sorting nexin 12 Snx12
 1421798_at 1.02026 mesogenin 1 Msgn1
 1421799_at 1.02746 urotensin 2 Uts2
 1421800_at 1.35281 per-hexamer repeat gene 1 Phxr1
 1421801_at 1.04121 --- ---
 1421802_at 0.94432 "eosinophil-associated, ribonuclease A family, member 1" Ear1
 1421803_at 0.93868 androgen-binding protein eta Apbh
 1421804_at 0.93062 "fatty acid binding protein 9, testis" Fabp9
 1421805_at 0.97961 --- ---
 1421806_at 0.99058 defensin beta 3 Defb3
 1421807_at 0.96684 defensin beta 6 Defb6
 1421808_at 1.06336 defensin beta 5 Defb5
 1421809_at 0.96746 DiGeorge syndrome critical region gene 2 Dgcr2
 1421810_at 1.01604 DiGeorge syndrome critical region gene 2 Dgcr2
 1421811_at 0.68873 thrombospondin 1 /// similar to thrombospondin 1 Thbs1 /// LOC640441
 1421812_at 1.26214 TAP binding protein Tapbp
 1421813_a_at 0.80694 prosaposin Psap
 1421814_at 1.03349 moesin Msn
 1421815_at 0.77499 ependymin related protein 2 (zebrafish) Epdr2
 1421816_at 0.97191 "similar to Glutathione reductase, mitochondrial precursor (GR) (GRase)"
 LOC630729
 1421817_at 0.93235 glutathione reductase 1 Gsr
 1421818_at 0.99575 B-cell leukemia/lymphoma 6 Bcl6
 1421819_a_at 1.19659 SET translocation /// similar to SET protein (Phosphatase 2A inhibitor I2PP2A) (I-2PP2A) (Template-activating factor I) (TAF-I) (Liver regeneration-related protein LRRGR00002) Set /// LOC671392
 1421820_a_at 0.93107 neurofibromatosis 2 Nf2
 1421821_at 1.45311 low density lipoprotein receptor Ldlr
 1421822_at 0.97204 "protein phosphatase 2 (formerly 2A), catalytic subunit, beta isoform" Ppp2cb
 1421823_a_at 0.97664 "protein phosphatase 2 (formerly 2A), catalytic subunit, beta isoform" Ppp2cb
 1421824_at 1.04091 beta-site APP cleaving enzyme 1 Bace1
 1421825_at 0.92734 beta-site APP cleaving enzyme 1 Bace1
 1421826_at 1.00036 delta-like 4 (Drosophila) Dll4
 1421827_at 0.76524 delta-like 4 (Drosophila) Dll4
 1421828_at 1.08412 karyopherin (importin) alpha 3 Kpna3
 1421829_at 1.11696 adenylate kinase 3 alpha-like 1 Ak311
 1421830_at 1.48040 adenylate kinase 3 alpha-like 1 Ak311
 1421831_at 1.07299 --- ---
 1421832_at 0.79265 twisted gastrulation homolog 1 (Drosophila) Twsg1
 1421833_at 0.93886 "phosphatidylinositol-4-phosphate 5-kinase, type 1 alpha" Pip5k1a
 1421834_at 0.96266 "phosphatidylinositol-4-phosphate 5-kinase, type 1 alpha" Pip5k1a

1421835_at	0.97077	microtubule-associated protein 7	Mtap7
1421836_at	0.95588	---	---
1421837_at	1.07022	ribosomal protein S18	Rps18
1421838_at	1.09357	ribosomal protein S18	Rps18
1421839_at	0.52932	"ATP-binding cassette, sub-family A (ABC1), member 1"	Abca1
1421840_at	0.25567	"ATP-binding cassette, sub-family A (ABC1), member 1"	Abca1
1421841_at	0.73813	fibroblast growth factor receptor 3	Fgfr3
1421842_a_at	0.89342	secretory carrier membrane protein 4	Scamp4
1421843_at	0.89242	interleukin 1 receptor accessory protein	Il1rap
1421844_at	1.23110	---	---
1421845_at	0.89332	golgi phosphoprotein 3	Golph3
1421846_at	1.00945	WD repeat and SOCS box-containing 2	Wsb2
1421847_at	0.98409	WD repeat and SOCS box-containing 2	Wsb2
1421848_at	0.97408	"solute carrier family 22 (organic cation transporter), member 5"	Slc22a5
1421849_at	0.91267	stromal antigen 2	Stag2
1421850_at	1.08473	microtubule-associated protein 1 B	Mtap1b
1421851_at	1.11920	microtubule-associated protein 1 B	Mtap1b
1421852_at	1.02349	"potassium channel, subfamily K, member 5"	Kcnk5
1421853_at	0.94879	presenilin 1	Psen1
1421854_at	1.05270	fibrinogen-like protein 2	Fgl2
1421855_at	1.00599	fibrinogen-like protein 2	Fgl2
1421856_at	1.35877	S100 calcium binding protein A3	S100a3
1421857_at	0.94745	a disintegrin and metallopeptidase domain 17	Adam17
1421858_at	0.98915	a disintegrin and metallopeptidase domain 17	Adam17
1421859_at	1.03727	a disintegrin and metallopeptidase domain 17	Adam17
1421860_at	0.96010	calsyntenin 1	Clstn1
1421861_at	0.94977	calsyntenin 1	Clstn1
1421862_a_at	1.01704	vesicle-associated membrane protein 1	Vamp1
1421863_at	0.84691	vesicle-associated membrane protein 1	Vamp1
1421864_at	1.00306	diazepam binding inhibitor-like 5	Dbil5
1421865_at	0.97893	diazepam binding inhibitor-like 5	Dbil5
1421866_at	0.89714	"nuclear receptor subfamily 3, group C, member 1"	Nr3c1
1421867_at	1.04664	"nuclear receptor subfamily 3, group C, member 1"	Nr3c1
1421868_a_at	0.94537	pancreatic lipase	Pnlip
1421869_at	1.12215	tripartite motif-containing 44	Trim44
1421870_at	0.97716	tripartite motif-containing 44	Trim44
1421871_at	0.79209	SH3-binding domain glutamic acid-rich protein like	Sh3bgrl
1421872_at	1.15240	"RAB24, member RAS oncogene family"	Rab24
1421873_s_at	1.13757	"RAB24, member RAS oncogene family"	Rab24
1421874_a_at	1.00700	mitochondrial ribosomal protein S23	Mrps23
1421875_a_at	0.98037	mitochondrial ribosomal protein S23	Mrps23
1421876_at	0.95597	mitogen activated protein kinase 9	Mapk9
1421877_at	0.86462	mitogen activated protein kinase 9	Mapk9

1421878_at 0.91576 mitogen activated protein kinase 9 Mapk9
1421879_at 0.95574 myotubularin related protein 1 Mtmr1
1421880_at 0.87773 myotubularin related protein 1 Mtmr1
1421881_a_at 0.95054 "ELAV (embryonic lethal, abnormal vision, Drosophila)-like 2 (Hu antigen B)" Elavl2
1421882_a_at 1.02149 "ELAV (embryonic lethal, abnormal vision, Drosophila)-like 2 (Hu antigen B)" Elavl2
1421883_at 1.03097 "ELAV (embryonic lethal, abnormal vision, Drosophila)-like 2 (Hu antigen B)" Elavl2
1421884_at 1.02719 Son of sevenless homolog 1 (Drosophila) Sos1
1421885_at 1.02397 Son of sevenless homolog 1 (Drosophila) Sos1
1421886_at 1.13381 Son of sevenless homolog 1 (Drosophila) Sos1
1421887_a_at 0.90928 amyloid beta (A4) precursor-like protein 2 Aplp2
1421888_x_at 0.94037 amyloid beta (A4) precursor-like protein 2 Aplp2
1421889_a_at 0.86890 amyloid beta (A4) precursor-like protein 2 Aplp2
1421890_at 0.82770 "ST3 beta-galactoside alpha-2,3-sialyltransferase 2" St3gal2
1421891_at 0.82819 "ST3 beta-galactoside alpha-2,3-sialyltransferase 2" St3gal2
1421892_at 0.87857 --- ---
1421893_a_at 1.04670 tripeptidyl peptidase II Tpp2
1421894_a_at 1.09390 tripeptidyl peptidase II Tpp2
1421895_at 1.12130 "eukaryotic translation initiation factor 2, subunit 3, structural gene X-linked /// similar to eukaryotic translation initiation factor 2, subunit 3, structural gene X-linked" Eif2s3x /// LOC673373
1421896_at 0.98618 "ELK1, member of ETS oncogene family" Elk1
1421897_at 1.00888 "ELK1, member of ETS oncogene family" Elk1
1421898_a_at 0.91217 "major histocompatibility complex, class I-related" Mr1
1421899_a_at 0.92830 "major histocompatibility complex, class I-related" Mr1
1421900_at 0.98909 eukaryotic translation initiation factor 2 alpha kinase 1 Eif2ak1
1421901_at 0.93396 eukaryotic translation initiation factor 2 alpha kinase 1 Eif2ak1
1421902_at 0.98656 intersex-like (Drosophila) Ixl
1421903_at 0.98110 intersex-like (Drosophila) Ixl
1421904_at 1.00090 nuclear receptor coactivator 6 interacting protein Ncoa6ip
1421905_at 1.02551 nuclear receptor coactivator 6 interacting protein Ncoa6ip
1421906_at 1.11024 peroxisome proliferator activated receptor binding protein Pparbp
1421907_at 1.11821 peroxisome proliferator activated receptor binding protein Pparbp
1421908_a_at 0.90963 transcription factor 12 Tcf12
1421909_at 0.97990 transcription factor 20 Tcf20
1421910_at 1.09777 transcription factor 20 Tcf20
1421911_at 0.96549 signal transducer and activator of transcription 2 Stat2
1421912_at 1.01021 "solute carrier family 23 (nucleobase transporters), member 1" Slc23a1
1421913_at 1.08589 mitochondrial ribosomal protein L19 Mrpl19
1421914_s_at 1.07264 mitochondrial ribosomal protein L19 Mrpl19
1421915_a_at 0.61199 "ST3 beta-galactoside alpha-2,3-sialyltransferase 3" St3gal3

1421916_at	0.91248	"platelet derived growth factor receptor, alpha polypeptide"	Pdgfra
1421917_at	1.00173	"platelet derived growth factor receptor, alpha polypeptide"	Pdgfra
1421918_at	0.95759	"acidic (leucine-rich) nuclear phosphoprotein 32 family, member A"	Anp32a
1421919_a_at	0.97475	chemokine (C-C motif) receptor 9	Ccr9
1421920_a_at	0.91692	chemokine (C-C motif) receptor 9	Ccr9
1421921_at	0.94712	"serine (or cysteine) peptidase inhibitor, clade A, member 3M"	Serpina3m
1421922_at	1.24248	SH3-domain binding protein 5 (BTK-associated)	Sh3bp5
1421923_at	1.32441	SH3-domain binding protein 5 (BTK-associated)	Sh3bp5
1421924_at	0.96427	"solute carrier family 2 (facilitated glucose transporter), member 3"	Slc2a3
1421925_at	1.05119	mitogen-activated protein kinase 11	Mapk11
1421926_at	1.40835	mitogen-activated protein kinase 11	Mapk11
1421927_at	1.14119	transducin (beta)-like 2	Tbl2
1421928_at	1.04876	Eph receptor A4	Epha4
1421929_at	1.13585	Eph receptor A4	Epha4
1421930_at	0.99818	inducible T-cell co-stimulator	Icos
1421931_at	0.91253	inducible T-cell co-stimulator	Icos
1421932_at	1.03713	chromobox homolog 5 (Drosophila HP1a)	Cbx5
1421933_at	0.93702	chromobox homolog 5 (Drosophila HP1a)	Cbx5
1421934_at	1.01995	chromobox homolog 5 (Drosophila HP1a)	Cbx5
1421935_at	1.05989	ribosomal protein S20 /// similar to 40S ribosomal protein S20 /// similar to 40S ribosomal protein S20	Rps20 /// LOC245676 /// LOC623568
1421936_at	1.11981	dual adaptor for phosphotyrosine and 3-phosphoinositides 1	Dapp1
1421937_at	1.09205	dual adaptor for phosphotyrosine and 3-phosphoinositides 1	Dapp1
1421938_at	0.84200	serine hydrolase-like	Serhl
1421939_a_at	0.95013	stromal antigen 1	Stag1
1421940_at	1.02215	stromal antigen 1	Stag1
1421941_at	0.95327	calcium/calmodulin-dependent protein kinase IV	Camk4
1421942_s_at	0.99772	transforming growth factor alpha	Tgfa
1421943_at	1.18026	transforming growth factor alpha	Tgfa
1421944_a_at	1.01243	asialoglycoprotein receptor 1	Asgr1
1421945_a_at	1.16080	brix domain containing 1	Bxdc1
1421946_at	0.98720	"C-reactive protein, pentraxin-related"	Crp
1421947_at	1.12967	"guanine nucleotide binding protein (G protein), gamma 12"	Gng12
1421948_a_at	0.97381	RIKEN cDNA 2610507L03 gene	2610507L03Rik
1421949_a_at	1.02394	RIKEN cDNA 2610507L03 gene	2610507L03Rik
1421950_at	1.10836	prefoldin 2	Pfdn2
1421951_at	0.71705	LIM homeobox protein 1	Lhx1
1421952_at	1.17356	calpain 6	Capn6
1421953_at	0.92603	v-crk sarcoma virus CT10 oncogene homolog (avian)-like	Crkl
1421954_at	0.91127	v-crk sarcoma virus CT10 oncogene homolog (avian)-like	Crkl
1421955_a_at	0.90040	"neural precursor cell expressed, developmentally down-regulated gene 4"	Nedd4
1421956_at	1.11437	recombining binding protein suppressor of hairless-like (Drosophila)	Rbpsuhl

1421957_a_at 1.03966 "phosphate cytidyltransferase 1, choline, alpha isoform" Pcyt1a
1421958_at 0.95177 L1 cell adhesion molecule L1cam
1421959_s_at 0.96574 adenylate cyclase 3 Adcy3
1421960_at 0.93843 adenylate cyclase 3 Adcy3
1421961_a_at 0.98662 "DnaJ (Hsp40) homolog, subfamily B, member 5" Dnajb5
1421962_at 0.92277 "DnaJ (Hsp40) homolog, subfamily B, member 5" Dnajb5
1421963_a_at 1.06251 cell division cycle 25 homolog B (S. cerevisiae) Cdc25b
1421964_at 0.69701 Notch gene homolog 3 (Drosophila) Notch3
1421965_s_at 0.56422 Notch gene homolog 3 (Drosophila) Notch3
1421966_at 0.87309 neural cell adhesion molecule 1 Ncam1
1421967_at 1.09085 "UDP-Gal:betaGlcNAc beta 1,4-galactosyltransferase, polypeptide 5"
B4galt5
1421968_a_at 0.96658 non imprinted in Prader-Willi/Angelman syndrome 2 homolog (human)
Nipa2
1421969_a_at 0.88994 fatty acid amide hydrolase Faah
1421970_a_at 0.89454 "glutamate receptor, ionotropic, AMPA2 (alpha 2)" Gria2
1421971_a_at 1.12335 mitochondrial ribosomal protein S34 Mrps34
1421972_s_at 1.01170 host cell factor C1 Hcfc1
1421973_at 0.91864 glial cell line derived neurotrophic factor family receptor alpha 1 Gfra1
1421974_at 0.98083 leucine rich repeat containing 50 Lrrc50
1421975_a_at 1.08992 adducin 2 (beta) Add2
1421976_at 0.89409 matrix metalloproteinase 19 Mmp19
1421977_at 1.04228 matrix metalloproteinase 19 Mmp19
1421978_at 0.99840 glutamic acid decarboxylase 2 Gad2
1421979_at 0.89391 "phosphate regulating gene with homologies to endopeptidases on the X
chromosome (hypophosphatemia, vitamin D resistant rickets)" Phex
1421980_at 1.04893 "potassium voltage gated channel, Shaw-related subfamily, member 3" Kcnc3
1421981_at 0.93527 "potassium voltage gated channel, Shaw-related subfamily, member 3" Kcnc3
1421982_a_at 1.04950 unc-50 homolog (C. elegans) Unc50
1421983_s_at 0.96391 "hepatic nuclear factor 4, alpha" Hnf4a
1421984_at 1.06618 stanniocalcin 1 Stc1
1421985_a_at 1.10347 eukaryotic translation initiation factor 4E member 2 Eif4e2
1421986_at 1.02839 eukaryotic translation initiation factor 4E member 2 Eif4e2
1421987_at 0.98691 3'-phosphoadenosine 5'-phosphosulfate synthase 2 Papss2
1421988_at 1.00895 3'-phosphoadenosine 5'-phosphosulfate synthase 2 Papss2
1421989_s_at 1.00173 3'-phosphoadenosine 5'-phosphosulfate synthase 2 Papss2
1421990_at 1.01130 synaptotagmin I Syt1
1421991_a_at 0.90082 insulin-like growth factor binding protein 4 Igfbp4
1421992_a_at 0.67232 insulin-like growth factor binding protein 4 Igfbp4
1421993_a_at 1.10783 transmembrane protein 134 Tmem134
1421994_a_at 0.95342 HCLS1 binding protein 3 Hs1bp3
1421995_at 1.01484 "transcription factor AP-2, alpha" Tcfap2a
1421996_at 1.02100 "transcription factor AP-2, alpha" Tcfap2a

1421997_s_at 1.02701 integrin alpha 3 Itga3
1421998_at 1.36372 "torsin family 3, member A" Tor3a
1421999_at 0.92251 thyroid stimulating hormone receptor Tshr
1422000_at 0.88625 "aldo-keto reductase family 1, member C12" Akr1c12
1422001_at 0.97826 inhibin beta-C Inhbc
1422002_at 0.89731 MAX dimerization protein 1 Mxd1
1422003_at 0.95025 Burkitt lymphoma receptor 1 Blr1
1422004_at 0.96135 RIKEN cDNA D730048I06 gene D730048I06Rik
1422005_at 1.13052 eukaryotic translation initiation factor 2-alpha kinase 2 Eif2ak2
1422006_at 1.19911 eukaryotic translation initiation factor 2-alpha kinase 2 Eif2ak2
1422007_at 1.11478 aquaporin 3 Aqp3
1422008_a_at 1.07332 aquaporin 3 Aqp3
1422009_at 0.86687 "ATPase, Na+/K+ transporting, beta 2 polypeptide" Atp1b2
1422010_at 0.89966 toll-like receptor 7 Tlr7
1422011_s_at 1.03985 X-linked lymphocyte-regulated complex /// RIKEN cDNA 3830403N18 gene Xlr /// 3830403N18Rik
1422012_at 1.04612 corticotropin releasing hormone receptor 2 Crhr2
1422013_at 1.22288 "C-type lectin domain family 4, member a2" Clec4a2
1422014_at 0.76901 forkhead box P2 Foxp2
1422015_a_at 0.96413 "ATP-binding cassette, sub-family B (MDR/TAP), member 8" Abcb8
1422016_a_at 0.97228 centromere protein H Cenph
1422017_s_at 0.88913 RIKEN cDNA 4833439L19 gene 4833439L19Rik
1422018_at 0.80733 human immunodeficiency virus type I enhancer binding protein 2 Hivep2
1422019_at 0.96254 "transforming growth factor, beta receptor II" Tgfbr2
1422020_at 0.91069 sprouty homolog 4 (Drosophila) Spry4
1422021_at 0.92081 sprouty homolog 4 (Drosophila) Spry4
1422022_at 0.82289 zinc finger protein 319 Zfp319
1422023_at 1.00162 bassoon Bsn
1422024_at 1.04102 Friend leukemia integration 1 Fli1
1422025_at 0.99488 microphthalmia-associated transcription factor Mitf
1422026_at 0.93608 peptidase inhibitor 16 Pi16
1422027_a_at 1.03981 "E26 avian leukemia oncogene 1, 5' domain" Ets1
1422028_a_at 1.20961 "E26 avian leukemia oncogene 1, 5' domain" Ets1
1422029_at 2.23945 chemokine (C-C motif) ligand 20 Ccl20
1422030_at 1.03947 "ATPase, H+ transporting, lysosomal V0 subunit A4" Atp6v0a4
1422031_a_at 0.89570 "zinc finger, AN1-type domain 6" Zfand6
1422032_a_at 0.94531 "zinc finger, AN1-type domain 6" Zfand6
1422033_a_at 1.15115 ciliary neurotrophic factor /// zinc finger protein 91 Cntf /// Zfp91
1422034_a_at 0.85147 paralemmin Palm
1422035_at 1.05334 "serine (or cysteine) peptidase inhibitor, clade B, member 9c" Serpinb9c
1422036_at 0.99051 "striatin, calmodulin binding protein" Strn
1422037_at 1.07345 distal-less homeobox 3 Dlx3
1422038_a_at 1.09677 "tumor necrosis factor receptor superfamily, member 22" Tnfrsf22

1422039_at 0.97789 "tumor necrosis factor receptor superfamily, member 22" Tnfrsf22
1422040_at 0.88159 "sema domain, immunoglobulin domain (Ig), and GPI membrane anchor, (semaphorin) 7A" Sema7a
1422041_at 1.01227 paired immunoglobulin-like type 2 receptor beta 1 Pilrb1
1422042_at 1.10444 gap junction membrane channel protein epsilon 1 Gje1
1422043_at 1.09640 tuberous sclerosis 1 Tsc1
1422044_at 0.86751 N-deacetylase/N-sulfotransferase (heparan glucosaminyl) 1 Ndst1
1422045_a_at 1.04787 "protein tyrosine phosphatase, non-receptor type 12" Ptpn12
1422046_at 1.02543 integrin alpha M Itgam
1422047_at 0.98717 cadherin 5 Cdh5
1422048_at 1.12505 "transient receptor potential cation channel, subfamily C, member 5" Trpc5
1422049_at 1.00591 "NK1 transcription factor related, locus 2 (Drosophila)" Nkx1-2
1422050_at 1.05658 "NK1 transcription factor related, locus 2 (Drosophila)" Nkx1-2
1422051_a_at 1.21578 "gamma-aminobutyric acid (GABA-B) receptor, 1" Gabbr1
1422052_at 0.91100 cadherin 8 Cdh8
1422053_at 0.79646 inhibin beta-A Inhba
1422054_a_at 0.89291 SKI-like Skil
1422055_at 1.21504 midline 1 Mid1
1422056_at 0.90351 netrin 2-like (chicken) Ntn2l
1422057_at 1.05905 nodal Nodal
1422058_at 0.99336 --- ---
1422059_at 1.02390 chromobox homolog 2 (Drosophila Pc class) Cbx2
1422060_at 0.95694 "parvin, beta /// similar to parvin, beta" Parvb /// LOC676024
1422061_at 0.91806 "aldo-keto reductase family 1, member C20" Akr1c20
1422062_at 1.04022 macrophage scavenger receptor 1 Msr1
1422063_a_at 0.81969 peroxisome biogenesis factor 5 Pex5
1422064_a_at 0.88096 zinc finger and BTB domain containing 20 Zbtb20
1422065_at 1.01746 killer cell lectin-like receptor subfamily B member 1D Klrbl1d
1422066_x_at 0.95770 killer cell lectin-like receptor subfamily B member 1D Klrbl1d
1422067_at 0.96673 killer cell lectin-like receptor subfamily B member 1D Klrbl1d
1422068_at 1.06767 "POU domain, class 3, transcription factor 1" Pou3f1
1422069_at 1.10179 melanocortin 1 receptor Mc1r
1422070_at 0.98141 "alcohol dehydrogenase 4 (class II), pi polypeptide" Adh4
1422071_at 0.94488 "lectin, galactose binding, soluble 6" Lgals6
1422072_a_at 0.89593 "glutathione S-transferase, mu 6" Gstm6
1422073_a_at 0.73748 cadherin EGF LAG seven-pass G-type receptor 2 Celsr2
1422074_at 0.99555 caudal type homeo box 2 Cdx2
1422075_at 0.87008 caudal type homeo box 2 Cdx2
1422076_at 0.95803 acyl-CoA thioesterase 4 Acot4
1422077_at 1.09610 acyl-CoA thioesterase 4 Acot4
1422078_at 0.79681 thymoma viral proto-oncogene 3 Akt3
1422079_at 0.85508 "protein kinase C, eta" Prkch
1422080_at 1.06633 interleukin 7 Il7

1422081_at 1.02213 kyphoscoliosis peptidase Ky
1422082_a_at 1.00349 nuclear transcription factor-Y alpha Nfya
1422083_at 0.96456 toll-like receptor 9 Tlr9
1422084_at 0.93548 BMX non-receptor tyrosine kinase Bmx
1422085_at 0.87330 T-box 19 Tbx19
1422086_at 0.83875 T-box 19 Tbx19
1422087_at 0.87374 "v-myc myelocytomatosis viral oncogene homolog 1, lung carcinoma derived (avian)" Mycl1
1422088_at 0.98742 "v-myc myelocytomatosis viral oncogene homolog 1, lung carcinoma derived (avian)" Mycl1
1422089_at 0.96700 natural cytotoxicity triggering receptor 1 Ncr1
1422090_a_at 0.84604 "6-phosphofructo-2-kinase/fructose-2,6-biphosphatase 2" Pfkfb2
1422091_at 0.97818 "6-phosphofructo-2-kinase/fructose-2,6-biphosphatase 2" Pfkfb2
1422092_at 0.92946 "6-phosphofructo-2-kinase/fructose-2,6-biphosphatase 2" Pfkfb2
1422093_at 1.07032 wingless-related MMTV integration site 3A Wnt3a
1422094_a_at 1.39290 zinc finger protein 329 Zfp329
1422095_a_at 1.05887 thymidylate kinase family LPS-inducible member Tyki
1422096_at 0.92275 RIKEN cDNA 4933434I20 gene 4933434I20Rik
1422097_at 0.87734 RIKEN cDNA 4933434I20 gene 4933434I20Rik
1422098_at 1.04609 "activin A receptor, type 1B" Acvr1b
1422099_a_at 1.11354 opioid receptor-like 1 Oprl1
1422100_at 1.00702 "cytochrome P450, family 7, subfamily a, polypeptide 1" Cyp7a1
1422101_at 1.14531 "tumor necrosis factor receptor superfamily, member 23" Tnfrsf23
1422102_a_at 0.88462 signal transducer and activator of transcription 5B Stat5b
1422103_a_at 0.93987 signal transducer and activator of transcription 5B Stat5b
1422104_at 0.98624 visual system homeobox 1 homolog (zebrafish) Vsx1
1422105_at 1.04289 "CD3 antigen, epsilon polypeptide" Cd3e
1422106_a_at 0.98247 splA/ryanodine receptor domain and SOCS box containing 2 Spsb2
1422107_at 0.87245 RIKEN cDNA 2410066E13 gene 2410066E13Rik
1422108_at 0.92553 "protein phosphatase 1, regulatory (inhibitor) subunit 3A" Ppp1r3a
1422109_at 1.01811 "regulatory factor X, 1 (influences HLA class II expression)" Rfx1
1422110_at 0.92048 hairless Hr
1422111_at 0.98744 chemokine binding protein 2 Ccbp2
1422112_at 1.00152 chemokine binding protein 2 Ccbp2
1422113_at 0.89888 "solute carrier family 5 (sodium iodide symporter), member 5" Slc5a5
1422114_at 1.02776 Ets2 repressor factor Erf
1422115_a_at 1.05427 FMS-like tyrosine kinase 3 ligand Flt3l
1422116_at 0.94639 "AF4/FMR2 family, member 2" Aff2
1422117_s_at 0.98996 "KH domain containing, RNA binding, signal transduction associated 2" Khdrbs2
1422118_at 1.03246 syncoilin Sync
1422119_at 1.10249 "RAB5B, member RAS oncogene family" Rab5b
1422120_at 1.00250 ELL associated factor 2 Eaf2

1422121_at 1.07588 "opioid receptor, delta 1" Oprd1
1422122_at 0.99201 "Fc receptor, IgE, low affinity II, alpha polypeptide" Fcgr2a
1422123_s_at 0.94468 CEA-related cell adhesion molecule 1 /// CEA-related cell adhesion molecule
2 Ceacam1 /// Ceacam2
1422124_a_at 1.01086 "protein tyrosine phosphatase, receptor type, C" Ptpcr
1422125_at 0.92403 5-hydroxytryptamine (serotonin) receptor 2B Htr2b
1422126_a_at 1.03625 nudix (nucleoside diphosphate linked moiety X)-type motif 13 Nudt13
1422127_at 1.05276 desert hedgehog Dhh
1422128_at 1.09024 ribosomal protein L14 Rpl14
1422129_at 0.95053 adenomatous polyposis coli 2 Apc2
1422130_at 1.00320 neuronal pentraxin 1 Nptx1
1422131_at 0.95031 decapping enzyme Dcp1a
1422132_at 0.92896 "5,10-methylenetetrahydrofolate reductase" Mthfr
1422133_at 0.99568 sialophorin Spn
1422134_at 0.56161 FBJ osteosarcoma oncogene B Fosb
1422135_at 1.05398 similar to zinc finger protein 146 LOC677248
1422136_at 1.12867 U2AF homology motif (UHM) kinase 1 /// similar to U2AF homology motif
(UHM) kinase 1 Uhmk1 /// LOC677213
1422137_at 0.93599 RIKEN cDNA 9030623N16 gene 9030623N16Rik
1422138_at 0.94315 "plasminogen activator, urokinase" Plau
1422139_at 0.84730 "plasminogen activator, urokinase" Plau
1422140_at 0.96519 similar to component of Sp100-rs /// similar to component of Sp100-rs ///
similar to component of Sp100-rs /// similar to component of Sp100-rs /// similar to component of Sp100-
rs /// similar to component of Sp100-rs /// similar to component of Sp100-rs /// similar to component of
Sp100-rs /// similar to component of Sp100-rs /// similar to component of Sp100-rs Sp100-rs ///
LOC620113 /// LOC665338 /// LOC665378 /// LOC673083 /// LOC673135 /// LOC673206 ///
LOC673671 /// LOC673849 /// LOC673997
1422141_s_at 0.99968 component of Sp100-rs Csprs
1422142_at 1.03863 "nephrosis 1 homolog, nephrin (human)" Nphs1
1422143_at 0.95824 A kinase (PKA) anchor protein 7 Akap7
1422144_at 0.98659 inhibin beta E Inhbe
1422145_at 1.02896 mannoside acetylglucosaminyltransferase 3 Mgat3
1422146_at 0.87925 "sema domain, seven thrombospondin repeats (type 1 and type 1-like),
transmembrane domain (TM) and short cytoplasmic domain, (semaphorin) 5B" Sema5b
1422147_a_at 1.08325 "phospholipase A2, group VI" Pla2g6
1422148_at 0.89499 matrilin 3 Matn3
1422149_at 0.95255 myeloid/lymphoid or mixed lineage-leukemia translocation to 7 homolog
(Drosophila) Mllt7
1422150_at 0.94855 H6 homeo box 3 Hmx3
1422151_at 1.05013 opposite strand transcription unit to Stag3 Gats
1422152_at 0.82182 H6 homeo box 1 Hmx1
1422153_a_at 0.97081 ankyrin repeat and SOCS box-containing protein 11 Asb11
1422154_at 0.84525 G protein-coupled receptor 27 Gpr27

1422155_at 1.01345 "histone 2, H3c2" Hist2h3c2
 1422156_a_at 1.02494 ribosomal protein S2 /// similar to 40S ribosomal protein S2 /// similar to 40S
 ribosomal protein S2 /// similar to 40S ribosomal protein S2 /// similar to 40S ribosomal protein S2 ///
 similar to 40S ribosomal protein S2 /// similar to 40S ribosomal protein S2 /// similar to 40S ribosomal
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 protein S2 /// similar to 40S ribosomal protein S2 /// similar to 40S ribosomal protein S2 Rps2 ///
 LOC544872 /// LOC546164 /// LOC546298 /// LOC620248 /// LOC620559 /// LOC623242 ///
 LOC628962 /// LOC637606 /// LOC664975 /// LOC665951 /// LOC667279 /// LOC667846 ///
 LOC667847 /// LOC669206 /// LOC669347 /// LOC670479 /// LOC671929 /// LOC672594 ///
 LOC674755 /// LOC675043 /// LOC676775 /// LOC677135
 1422157_a_at 1.19510 integrin beta 1 binding protein 1 Itgb1bp1
 1422158_at 1.01172 calsyntenin 2 Clstn2
 1422159_at 1.03789 "protein phosphatase, EF hand calcium-binding domain 2" Ppef2
 1422160_at 1.10980 "histocompatibility 2, T region locus 24" H2-T24
 1422161_at 1.05310 "sialic acid binding Ig-like lectin 1, sialoadhesin" Siglec1
 1422162_at 0.87774 deleted in colorectal carcinoma Dcc
 1422163_at 0.90273 SH3 and PX domains 2A Sh3pxd2a
 1422164_at 1.04403 "POU domain, class 3, transcription factor 4" Pou3f4
 1422165_at 0.89677 "POU domain, class 3, transcription factor 4" Pou3f4
 1422166_at 0.99370 "C-type lectin domain family 2, member i" Clec2i
 1422167_at 1.03423 "sema domain, seven thrombospondin repeats (type 1 and type 1-like),
 transmembrane domain (TM) and short cytoplasmic domain, (semaphorin) 5A" Sema5a
 1422168_a_at 0.88026 brain derived neurotrophic factor Bdnf
 1422169_a_at 0.87686 brain derived neurotrophic factor Bdnf
 1422170_at 1.18834 "solute carrier family 5 (inositol transporters), member 3" Slc5a3
 1422171_at 0.89254 prostaglandin D receptor Ptgdr
 1422172_x_at 0.89511 RIKEN cDNA 2810408B13 gene 2810408B13Rik
 1422173_at 0.98825 "insulin promoter factor 1, homeodomain transcription factor" Ipf1
 1422174_at 1.02559 "insulin promoter factor 1, homeodomain transcription factor" Ipf1
 1422175_at 1.06220 matrix metalloproteinase 1a (interstitial collagenase) /// similar to matrix
 metalloproteinase 1a (interstitial collagenase) Mmp1a /// LOC672109
 1422176_at 1.02914 fibroblast growth factor 23 Fgf23
 1422177_at 1.09138 "interleukin 13 receptor, alpha 2" Il13ra2
 1422178_a_at 0.87058 "RAB17, member RAS oncogene family" Rab17
 1422179_at 1.24857 gap junction membrane channel protein beta 4 Gjb4
 1422180_a_at 0.90359 mast cell protease 6 Mcpt6
 1422181_at 1.04421 cyclic nucleotide gated channel alpha 2 Cnga2
 1422182_at 1.01612 "hepatocyte nuclear factor 4, gamma" Hnf4g
 1422183_a_at 0.82987 "adrenergic receptor, alpha 1b" Adra1b

1422184_a_at 0.77704 adenylate kinase 1 Ak1
1422185_a_at 1.00601 cytochrome b5 reductase 3 Cyb5r3
1422186_s_at 0.98698 cytochrome b5 reductase 3 Cyb5r3
1422187_at 0.86198 "gamma-aminobutyric acid (GABA-A) receptor, subunit gamma 3" Gabrg3
1422188_s_at 0.96188 "T-cell receptor gamma chain /// T cell receptor gamma chain /// region containing T-cell receptor gamma chain; T-cell receptor gamma, variable 4" Tcrg /// LOC670271
1422189_x_at 1.06588 "T-cell receptor gamma chain /// region containing T-cell receptor gamma chain; T-cell receptor gamma, variable 4" Tcrg /// LOC670271
1422190_at 0.96201 complement component 5a receptor 1 C5ar1
1422191_at 0.99479 CD200 receptor 1 Cd200r1
1422192_at 1.00079 gap junction membrane channel protein alpha 5 Gja5
1422193_at 1.04765 guanylate cyclase 2e Gucy2e
1422194_at 1.00717 "sodium channel, voltage-gated, type V, alpha" Scn5a
1422195_s_at 0.79393 T-box 15 Tbx15
1422196_at 1.00468 5-hydroxytryptamine (serotonin) receptor 5B Htr5b
1422197_at 0.93716 "potassium voltage-gated channel, shaker-related subfamily, member 2" Kcna2
1422198_a_at 1.14878 serine hydroxymethyl transferase 1 (soluble) Shmt1
1422199_at 0.94633 olfactory marker protein Omp
1422200_at 0.97545 olfactory marker protein Omp
1422201_at 1.09416 --- ---
1422202_at 0.92913 thyroid hormone receptor beta Thrb
1422203_at 0.92239 "solute carrier family 18 (vesicular monoamine), member 3" Slc18a3
1422204_at 0.94170 arginine vasopressin receptor 1B Avpr1b
1422205_at 0.97753 SRY-box containing gene 1 Sox1
1422206_at 0.97886 "UDP-Gal:betaGlcNAc beta 1,3-galactosyltransferase, polypeptide 1" B3galt1
1422207_at 1.09572 5-hydroxytryptamine (serotonin) receptor 5A Htr5a
1422208_a_at 0.78333 "guanine nucleotide binding protein, beta 5" Gnb5
1422209_s_at 0.99705 keratin associated protein 9-1 /// similar to keratin associated protein 9-1 /// similar to keratin associated protein 9-1 /// similar to keratin associated protein 9-1 Krtap9-1 /// LOC432600 /// LOC546515 /// LOC670533
1422210_at 1.05256 forkhead box D3 Foxd3
1422211_a_at 0.89466 "UDP-GlcNAc:betaGal beta-1,3-N-acetylglucosaminyltransferase 3" B3gnt3
1422212_at 0.89625 forkhead box H1 Foxh1
1422213_s_at 0.88695 forkhead box H1 Foxh1
1422214_at 0.97070 neuropeptide FF receptor 2 Npffr2
1422215_at 1.06346 RIKEN cDNA 4930542C12 gene 4930542C12Rik
1422216_at 0.82226 midline 2 Mid2
1422217_a_at 0.44727 "cytochrome P450, family 1, subfamily a, polypeptide 1" Cyp1a1
1422218_at 0.94056 "purinergic receptor P2X, ligand-gated ion channel, 7" P2rx7
1422219_a_at 0.94517 T-box 20 Tbx20
1422220_at 1.00087 pituitary specific transcription factor 1 Pit1
1422221_at 0.92561 heart and neural crest derivatives expressed transcript 2 Hand2

1422222_at 0.82475 involucrin Iv1
1422223_at 1.12226 "glutamate receptor, ionotropic, NMDA2B (epsilon 2)" Grin2b
1422224_at 1.10119 t-complex protein 10a /// t-complex protein 10b /// t-complex protein 10c
Tcp10a /// Tcp10b /// Tcp10c
1422225_s_at 0.93059 t-complex protein 10a /// t-complex protein 10b /// t-complex protein 10c
Tcp10a /// Tcp10b /// Tcp10c
1422226_at 0.72486 --- ---
1422227_at 1.04668 Kruppel-like factor 12 /// RIKEN cDNA D530033K05 gene Klf12 ///
D530033K05Rik
1422228_at 1.04771 wingless-related MMTV integration site 8A Wnt8a
1422229_at 0.86017 deubiquitinating enzyme 2 Dub2
1422230_s_at 1.02425 "cytochrome P450, family 2, subfamily a, polypeptide 4 /// cytochrome P450,
family 2, subfamily a, polypeptide 5" Cyp2a4 /// Cyp2a5
1422231_a_at 1.01466 "tumor necrosis factor receptor superfamily, member 25" Tnfrsf25
1422232_at 0.88280 paired-like homeobox 2b Phox2b
1422233_at 0.93451 "ELK4, member of ETS oncogene family" Elk4
1422234_at 0.93533 tachykinin receptor 2 Tacr2
1422235_at 0.92600 5-hydroxytryptamine (serotonin) receptor 7 Htr7
1422236_at 1.00113 RIKEN cDNA 4930560E09 gene 4930560E09Rik
1422237_at 0.89155 melanocortin 3 receptor Mc3r
1422238_at 0.97477 upregulated during skeletal muscle growth 1 Usmg1
1422239_at 0.73206 homeo box D13 Hoxd13
1422240_s_at 1.00084 small proline-rich protein 2H Sprr2h
1422241_a_at 0.95360 "NADH dehydrogenase (ubiquinone) 1 alpha subcomplex, 1" Ndufa1
1422242_at 0.92372 "defensin related cryptdin, related sequence 10" Defcr-rs10
1422243_at 1.11509 fibroblast growth factor 7 Fgf7
1422244_at 0.94826 "polycystic kidney disease (polycystin) and REJ (sperm receptor for egg jelly,
sea urchin homolog)-like" Pkdrej
1422245_a_at 1.04665 MRV integration site 1 Mrvi1
1422246_at 1.07816 dentin sialophosphoprotein Dspp
1422247_a_at 1.12670 "ubiquitously transcribed tetratricopeptide repeat gene, Y chromosome" Uty
1422248_at 0.83123 insulin receptor substrate 4 /// similar to insulin receptor substrate 4 Irs4 ///
LOC669982
1422249_s_at 0.78528 "zinc finger protein, autosomal /// zinc finger protein X-linked" Zfa /// Zfx
1422250_at 0.87330 mitogen activated protein kinase kinase kinase 2 Map3k2
1422251_at 1.05418 neuralized-like homolog (Drosophila) Neurl
1422252_a_at 1.02318 cell division cycle 25 homolog C (S. cerevisiae) Cdc25c
1422253_at 0.81023 "procollagen, type X, alpha 1" Col10a1
1422254_a_at 0.92333 dual-specificity tyrosine-(Y)-phosphorylation regulated kinase 1b Dyrk1b
1422255_at 0.96161 "potassium voltage-gated channel, shaker-related subfamily, member 4" Kcna4
1422256_at 1.07486 somatostatin receptor 2 Sstr2
1422257_s_at 0.90498 "cytochrome P450, family 2, subfamily b, polypeptide 10" Cyp2b10
1422258_at 0.93191 "cholinergic receptor, muscarinic 3, cardiac" Chrm3

1422259_a_at	1.05293	chemokine (C-C motif) receptor 5	Ccr5
1422260_x_at	1.03593	chemokine (C-C motif) receptor 5	Ccr5
1422261_a_at	0.96116	mutS homolog 3 (E. coli)	Msh3
1422262_a_at	0.85597	LIM homeobox protein 6	Lhx6
1422263_at	1.03179	"bradykinin receptor, beta 2"	Bdkrb2
1422264_s_at	0.87984	Kruppel-like factor 9	Klf9
1422265_at	1.21806	bombesin-like receptor 3	Brs3
1422266_at	0.75085	"myc-like oncogene, s-myc protein"	Mycs
1422267_at	0.91282	forkhead box B2	Foxb2
1422268_a_at	0.99043	"ribosomal protein S6 kinase, polypeptide 2"	Rps6kb2
1422269_at	1.06175	insulinoma-associated 2	Insm2
1422270_a_at	0.97359	"interleukin 6 receptor, alpha"	Il6ra
1422271_at	0.86442	pancreatic polypeptide receptor 1	Ppyr1
1422272_at	2.91765	per-hexamer repeat gene 4	Phxr4
1422273_at	0.94575	matrix metalloproteinase 1b (interstitial collagenase)	Mmp1b
1422274_at	0.88308	gap junction membrane channel protein alpha 8	Gja8
1422275_at	1.01838	G protein-coupled receptor 44	Gpr44
1422276_at	0.98819	"pyrimidinergic receptor P2Y, G-protein coupled, 4"	P2ry4
1422277_at	0.98008	"glycine receptor, alpha 1 subunit"	Gla1
1422278_at	0.99945	dopamine receptor 3	Drd3
1422279_at	1.16205	Friend virus susceptibility 1	Fv1
1422280_at	1.05601	granzyme K	Gzmk
1422281_at	0.87981	somatostatin receptor 4	Sstr4
1422282_at	0.97280	tachykinin receptor 1	Tacr1
1422283_at	1.03152	CD40 ligand	Cd40lg
1422284_at	1.03924	"NK2 transcription factor related, locus 9 (Drosophila)"	Nkx2-9
1422285_at	1.01359	orthopedia homolog (Drosophila)	Otp
1422286_a_at	0.98171	TG interacting factor	Tgif
1422287_at	0.99936	per-hexamer repeat gene 2	Phxr2
1422288_at	0.82925	5-hydroxytryptamine (serotonin) receptor 1B	Htr1b
1422289_a_at	0.94370	cathepsin Q	Ctsq
1422290_at	1.00707	5-hydroxytryptamine (serotonin) receptor 1D	Htr1d
1422291_at	1.11825	chemokine (C-C motif) receptor 8	Ccr8
1422292_at	0.83746	RIKEN cDNA A030005K14 gene	A030005K14Rik
1422293_a_at	0.70652	potassium channel tetramerisation domain containing 1	Kctd1
1422294_at	0.92419	chemokine (C motif) receptor 1	Xcr1
1422295_at	0.93542	myelodysplasia syndrome 1 homolog (human)	Mds1
1422296_at	1.09392	RIKEN cDNA 5730470L24 gene	5730470L24Rik
1422297_at	1.06098	prefoldin 5	Pfdn5
1422298_at	0.99671	RIKEN cDNA 5530401N06 gene	5530401N06Rik
1422299_a_at	1.05493	serine/threonine/tyrosine interacting-like 1	Styx11
1422300_at	1.11800	noggin	Nog
1422301_at	1.11054	ferritin light chain 2	Ftl2

1422302_s_at 1.12357 ferritin light chain 1 /// ferritin light chain 2 /// similar to Ferritin light chain 1 (Ferritin L subunit 1) /// similar to Ferritin light chain 1 (Ferritin L subunit 1) /// similar to Ferritin light chain 1 (Ferritin L subunit 1) /// similar to Ferritin light chain 1 (Ferritin L subunit 1) /// similar to Ferritin light chain 1 (Ferritin L subunit 1) /// similar to Ferritin light chain 1 (Ferritin L subunit 1) /// similar to Ferritin light chain 1 (Ferritin L subunit 1) /// Ftl1 /// Ftl2 /// LOC544951 /// LOC634386 /// LOC665937 /// LOC666673 /// LOC669088 /// LOC675170
 1422303_a_at 0.92579 "tumor necrosis factor receptor superfamily, member 18" Tnfrsf18
 1422304_at 1.01129 lipocalin 4 Lcn4
 1422305_at 1.01487 "interferon beta 1, fibroblast" Ifnb1
 1422306_at 0.90379 lipocalin 3 Lcn3
 1422307_at 0.82436 --- ---
 1422308_a_at 1.11189 "lectin, galactose binding, soluble 7" Lgals7
 1422309_a_at 0.86042 lens epithelial protein Lenep
 1422310_at 1.00958 stannin Snn
 1422311_a_at 0.88744 polymerase (RNA) II (DNA directed) polypeptide A Polr2a
 1422312_a_at 0.88703 neurogenin 3 Neurog3
 1422313_a_at 0.93480 insulin-like growth factor binding protein 5 Igfbp5
 1422314_at 0.95498 chloride channel 6 Clcn6
 1422315_x_at 0.97470 phosphorylase kinase gamma 1 Phkg1
 1422316_at 1.05624 "glycoprotein 1b, alpha polypeptide" Gp1ba
 1422317_a_at 0.91569 interleukin 1 receptor-like 1 Il1rl1
 1422318_at 1.09397 forkhead box D4 Foxd4
 1422319_at 1.01978 UDP-N-acetyl-alpha-D-galactosamine:polypeptide N-acetylgalactosaminyltransferase 11 Galnt11
 1422320_x_at 0.95222 per-hexamer repeat gene 5 Phxr5
 1422321_a_at 1.06633 splicing factor 1 Sf1
 1422322_at 0.95703 choroideremia-like Chml
 1422323_a_at 0.90171 ladybird homeobox homolog 1 (Drosophila) Lbx1
 1422324_a_at 1.14241 parathyroid hormone-like peptide Pthlh
 1422325_at 1.02136 "melanoma antigen, family A, 5" Magea5
 1422326_at 1.00225 glucose-6-phosphate dehydrogenase 2 G6pd2
 1422327_s_at 1.10152 glucose-6-phosphate dehydrogenase 2 /// glucose-6-phosphate dehydrogenase X-linked G6pd2 /// G6pdx
 1422328_at 0.92952 gap junction membrane channel protein alpha 10 Gja10
 1422329_a_at 0.96457 "neurotrophic tyrosine kinase, receptor, type 3" Ntrk3
 1422330_at 0.93714 glucagon-like peptide 1 receptor Glp1r
 1422331_at 0.99655 "POU domain, class 3, transcription factor 3" Pou3f3
 1422332_at 0.95830 interferon alpha 11 Ifna11
 1422333_at 0.99851 "cytochrome P450, family 21, subfamily a, polypeptide 1" Cyp21a1
 1422334_a_at 1.07210 surfactant associated protein A1 Sftpa1
 1422335_at 0.99825 "adrenergic receptor, alpha 2c" Adra2c
 1422336_at 1.08717 homeo box A13 Hoxa13
 1422337_at 0.99288 olfactory receptor 156 Olfr156

1422338_at 1.04537 G protein-coupled receptor 63 Gpr63
1422339_at 0.98649 gap junction membrane channel protein alpha 3 Gja3
1422340_a_at 0.97221 "actin, gamma 2, smooth muscle, enteric" Actg2
1422341_s_at 0.89995 lysophospholipase 3 Lypla3
1422342_at 0.92134 neuromedin B receptor Nmbr
1422343_at 1.02270 olfactory receptor 155 Olfr155
1422344_s_at 1.01680 "tumor necrosis factor receptor superfamily, member 10b" Tnfrsf10b
1422345_s_at 1.09435 "melanoma antigen, family B, 1 /// melanoma antigen, family B, 2 ///
melanoma antigen, family B, 3" Mageb1 /// Mageb2 /// Mageb3
1422346_at 0.92574 thyroid transcription factor 1 Titf1
1422347_at 0.95248 neuropeptide Y receptor Y6 Npy6r
1422348_at 1.12060 "similar to H-2 class I histocompatibility antigen, L-D alpha chain precursor"
LOC676708
1422349_at 1.05841 chemokine (C-C motif) receptor 1-like 1 Ccr111
1422350_at 0.86948 "formyl peptide receptor, related sequence 3" Fpr-rs3
1422351_at 0.98944 olfactory receptor 480 Olfr480
1422352_at 0.94687 mast cell protease 1 Mcpt1
1422353_at 0.95806 "POU domain, class 4, transcription factor 3" Pou4f3
1422354_at 0.85117 olfactory receptor 544 Olfr544
1422355_at 0.95743 trace amine-associated receptor 1 Taar1
1422356_at 0.96611 "vomeronasal 1 receptor, B10" V1rb10
1422357_at 1.02880 "melanoma antigen, family A, 6" Magea6
1422358_at 0.91891 "formyl peptide receptor, related sequence 4" Fpr-rs4
1422359_at 1.17030 olfactory receptor 690 Olfr690
1422360_at 1.09093 olfactory receptor 672 Olfr672
1422361_at 1.00213 T lymphoma oncogene Tlm
1422362_s_at 0.97167 olfactory receptor 68 /// olfactory receptor 69 Olfr68 /// Olfr69
1422363_at 1.01375 olfactory receptor 65 Olfr64
1422364_at 0.97469 olfactory receptor 68 Olfr68
1422365_at 0.99588 olfactory receptor 17 Olfr17
1422366_at 0.91541 "vomeronasal 1 receptor, A9" V1ra9
1422367_at 0.97691 olfactory receptor 70 Olfr70
1422368_at 0.90726 "vomeronasal 1 receptor, A5" V1ra5
1422369_at 1.03791 "vomeronasal 1 receptor, A6" V1ra6
1422370_at 0.80420 olfactory receptor 49 Olfr49
1422371_at 0.96665 olfactory receptor 1507 Olfr1507
1422372_at 0.97087 olfactory receptor 15 Olfr15
1422373_at 0.81318 olfactory receptor 71 Olfr71
1422374_s_at 1.19116 olfactory receptor 64 /// olfactory receptor 66 Olfr65 /// Olfr66
1422375_a_at 0.90127 ADP-ribosyltransferase 1 Art1
1422376_at 0.97525 "vomeronasal 1 receptor, B1" V1rb1
1422377_at 1.07962 "vomeronasal 1 receptor, B4" V1rb4
1422378_at 1.08881 "vomeronasal 1 receptor, B7" V1rb7

1422379_x_at 1.14473 "vomeronasal 1 receptor, B9" V1rb9
1422380_at 1.04415 "vomeronasal 1 receptor, B5" V1rb5
1422381_at 0.87099 olfactory receptor 160 Olfr160
1422382_at 1.08059 olfactory receptor 16 Olfr16
1422383_at 0.94036 "vomeronasal 1 receptor, B6" V1rb6
1422384_at 0.99022 olfactory receptor 1509 Olfr1509
1422385_at 0.79631 olfactory receptor 1264 Olfr1264
1422386_at 1.11265 olfactory receptor 64 Olfr65
1422387_at 0.98928 "vomeronasal 1 receptor, C2" V1rc2
1422388_at 1.02178 "vomeronasal 1 receptor, C8" V1rc8
1422389_at 1.09756 "taste receptor, type 2, member 105" Tas2r105
1422390_at 0.97765 "vomeronasal 1 receptor, C5" V1rc5
1422391_at 0.96114 "vomeronasal 1 receptor, C3" V1rc3
1422392_at 1.06153 "vomeronasal 1 receptor, C6" V1rc6
1422393_at 0.88522 forkhead box E3 Foxe3
1422394_at 1.06107 "vomeronasal 1 receptor, C4" V1rc4
1422395_at 0.97669 "vomeronasal 1 receptor, C8 /// vomeronasal 1 receptor, C22" V1rc8 ///
V1rc22
1422396_s_at 0.87184 achaete-scute complex homolog-like 2 (Drosophila) Ascl2
1422397_a_at 1.00191 "interleukin 15 receptor, alpha chain" Il15ra
1422398_at 0.95688 "histone 1, H1e" Hist1h1e
1422399_a_at 1.08767 "RAB23, member RAS oncogene family" Rab23
1422400_a_at 0.97096 hematopoietic cell transcript 1 Hemt1
1422401_at 0.98338 small proline-rich protein 3 Sprr3
1422402_a_at 0.94006 "ankyrin repeat, SAM and basic leucine zipper domain containing 1" Asz1
1422403_at 0.90602 interferon alpha 1 /// interferon alpha 2 /// interferon alpha 4 /// interferon alpha
5 /// interferon alpha 6 /// interferon alpha 7 /// interferon alpha 9 /// interferon alpha B /// interferon
alpha 6T /// alpha-interferon /// interferon alpha 12 /// interferon alpha 8/6 precursor; IFNa8/6 ///
interferon alpha 14 Ifna1 /// Ifna2 /// Ifna4 /// Ifna5 /// Ifna6 /// Ifna7 /// Ifna9 /// Ifnab /// Ifna6T ///
LOC242517 /// Ifna12 /// LOC384100 /// Ifna14
1422404_x_at 1.04806 interferon alpha 1 /// interferon alpha 2 /// interferon alpha 4 /// interferon alpha
5 /// interferon alpha 6 /// interferon alpha 7 /// interferon alpha 9 /// interferon alpha B /// interferon
alpha 6T /// alpha-interferon /// interferon alpha 12 /// interferon alpha 8/6 precursor; IFNa8/6 ///
interferon alpha 14 Ifna1 /// Ifna2 /// Ifna4 /// Ifna5 /// Ifna6 /// Ifna7 /// Ifna9 /// Ifnab /// Ifna6T ///
LOC242517 /// Ifna12 /// LOC384100 /// Ifna14
1422405_at 0.94848 "histocompatibility 2, D region locus 4" H2-D4
1422406_at 0.96014 interferon alpha 9 Ifna9
1422407_s_at 0.99311 Harvey rat sarcoma virus oncogene 1 Hras1
1422408_at 0.79137 interferon alpha 4 Ifna4
1422409_at 0.94106 hairy and enhancer of split 3 (Drosophila) Hes3
1422410_at 1.11453 Fer3-like (Drosophila) Ferd3l
1422411_s_at 0.97308 "eosinophil-associated, ribonuclease A family, member 1 /// eosinophil-
associated, ribonuclease A family, member 2 /// eosinophil-associated, ribonuclease A family, member

3 /// eosinophil-associated, ribonuclease A family, member 12" Ear1 /// Ear2 /// Ear3 /// Ear12
 1422412_x_at 0.96774 "eosinophil-associated, ribonuclease A family, member 3" Ear3
 1422413_at 0.95055 --- ---
 1422414_a_at 1.02939 calmodulin 2 Calm2
 1422415_at 0.92053 "angiogenin, ribonuclease A family, member 2" Ang2
 1422416_s_at 0.97892 pre-B lymphocyte gene 1 /// pre-B lymphocyte gene 2 Vpreb1 /// Vpreb2
 1422417_at 0.97683 --- ---
 1422418_s_at 1.01985 suppressor of Ty 4 homolog 1 (S. cerevisiae) /// suppressor of Ty 4 homolog 2
 (S. cerevisiae) Supt4h1 /// Supt4h2
 1422419_s_at 0.90709 transition protein 2 Tnp2
 1422420_at 0.99216 myoglobin Mb
 1422421_at 0.90675 "defensin related cryptdin, related sequence 12" Defcr-rs12
 1422422_at 1.11956 defensin related cryptdin 4 Defcr4
 1422423_at 0.98301 "melanoma antigen, family A, 7" Magea7
 1422424_at 1.05010 human papillomavirus 18 E5 central sequence motif gene 2 Hpv2
 1422425_at 0.98349 small proline-rich protein 2K Sprr2k
 1422426_at 0.96508 cell matrix adhesion regulator Cmar
 1422427_a_at 1.06659 "seminal vesicle protein, secretion 2" Svs2
 1422428_at 1.18151 acyl-CoA synthetase bubblegum family member 1 Acsbg1
 1422429_at 1.10451 ring finger protein 14 Rnf14
 1422430_at 1.05130 fidgetin-like 1 Figl1
 1422431_at 1.01773 "melanoma antigen, family E, 1" Magee1
 1422432_at 1.27036 diazepam binding inhibitor Dbi
 1422433_s_at 1.30445 "isocitrate dehydrogenase 1 (NADP+), soluble" Idh1
 1422434_a_at 1.10813 RIKEN cDNA 2210010C04 gene 2210010C04Rik
 1422435_at 0.99871 RIKEN cDNA 2210010C04 gene 2210010C04Rik
 1422436_at 1.02849 RIKEN cDNA 2210010C04 gene 2210010C04Rik
 1422437_at 0.94837 "procollagen, type V, alpha 2" Col5a2
 1422438_at 1.10184 "epoxide hydrolase 1, microsomal" Ephx1
 1422439_a_at 1.05126 cyclin-dependent kinase 4 /// similar to Cell division protein kinase 4 (Cyclin-
 dependent kinase 4) (PSK-J3) (CRK3) Cdk4 /// LOC640611
 1422440_at 1.10902 cyclin-dependent kinase 4 /// similar to Cell division protein kinase 4 (Cyclin-
 dependent kinase 4) (PSK-J3) (CRK3) Cdk4 /// LOC640611
 1422441_x_at 1.00594 cyclin-dependent kinase 4 Cdk4
 1422442_at 1.04626 smu-1 suppressor of mec-8 and unc-52 homolog (C. elegans) Smu1
 1422443_at 1.17974 "X-prolyl aminopeptidase (aminopeptidase P) 1, soluble" Xpnpep1
 1422444_at 0.97717 integrin alpha 6 Itga6
 1422445_at 0.87104 integrin alpha 6 Itga6
 1422446_x_at 0.95800 insulin II Ins2
 1422447_at 0.87538 insulin I Ins1
 1422448_at 1.38608 trefoil factor 2 (spasmolytic protein 1) Tff2
 1422449_s_at 0.80342 reticulocalbin 2 Rcn2
 1422450_at 1.05213 "catenin (cadherin associated protein), delta 1" Ctnnd1

1422451_at 0.95582 mitochondrial ribosomal protein S21 Mrps21
1422452_at 0.93908 Bcl2-associated athanogene 3 Bag3
1422453_at 0.98955 pre-mRNA processing factor 8 Prpf8
1422454_at 0.91226 keratin 13 Krt13
1422455_s_at 0.96597 N-ethylmaleimide sensitive fusion protein Nsf
1422456_at 1.03551 N-ethylmaleimide sensitive fusion protein Nsf
1422457_s_at 1.07222 SMT3 suppressor of mif two 3 homolog 3 (yeast) Sumo3
1422458_at 1.04382 T-cell lymphoma breakpoint 1 Tcl1
1422459_a_at 1.06203 "proteasome (prosome, macropain) 26S subunit, non-ATPase, 13" Psmd13
1422460_at 1.04194 "MAD2 (mitotic arrest deficient, homolog)-like 1 (yeast)" Mad2l1
1422461_at 1.08476 "ATPase family, AAA domain containing 3A" Atad3a
1422462_at 1.02593 ubiquitin-conjugating enzyme E2T (putative) Ube2t
1422463_a_at 1.01604 mitochondrial ribosomal protein L3 Mrpl3
1422464_at 0.93946 mitochondrial ribosomal protein L3 Mrpl3
1422465_a_at 0.97182 nucleoredoxin Nxn
1422466_at 1.00696 nucleoredoxin Nxn
1422467_at 0.97757 palmitoyl-protein thioesterase 1 Ppt1
1422468_at 0.97911 palmitoyl-protein thioesterase 1 Ppt1
1422469_at 1.17849 TANK-binding kinase 1 Tbk1
1422470_at 1.35236 "BCL2/adenovirus E1B interacting protein 1, NIP3" Bnip3
1422471_at 0.93392 peroxisomal biogenesis factor 13 Pex13
1422472_at 1.11895 peroxisomal biogenesis factor 13 Pex13
1422473_at 0.80155 "phosphodiesterase 4B, cAMP specific" Pde4b
1422474_at 0.94145 "phosphodiesterase 4B, cAMP specific" Pde4b
1422475_a_at 0.99085 ribosomal protein S3a Rps3a
1422476_at 0.89322 interferon gamma inducible protein 30 Ifi30
1422477_at 0.99634 Cdk5 and Abl enzyme substrate 1 /// similar to Cdk5 and Abl enzyme substrate 1 Cables1 /// LOC635753
1422478_a_at 1.42846 acyl-CoA synthetase short-chain family member 2 Acss2
1422479_at 1.38007 acyl-CoA synthetase short-chain family member 2 Acss2
1422480_at 0.98046 sorting nexin 3 Snx3
1422481_at 0.98644 keratin 1 Krt1
1422482_at 1.14184 RuvB-like protein 2 Ruvbl2
1422483_a_at 1.00247 "cytochrome c, somatic /// similar to Cytochrome c, somatic" Cyps /// LOC672195
1422484_at 0.95740 "cytochrome c, somatic" Cyps
1422485_at 1.00119 MAD homolog 4 (Drosophila) Smad4
1422486_a_at 1.04166 MAD homolog 4 (Drosophila) Smad4
1422487_at 1.00579 MAD homolog 4 (Drosophila) Smad4
1422488_at 1.15968 NTF2-related export protein 1 Nxt1
1422489_at 0.94588 glucosidase 1 Gcs1
1422490_at 1.03511 "BCL2/adenovirus E1B interacting protein 1, NIP2" Bnip2
1422491_a_at 1.00523 "BCL2/adenovirus E1B interacting protein 1, NIP2" Bnip2

1422492_at	1.04045	coproporphyrinogen oxidase	Cpox	
1422493_at	0.96749	coproporphyrinogen oxidase	Cpox	
1422494_s_at	0.97482	cell cycle related kinase	Ccrk	
1422495_a_at	1.01333	high mobility group nucleosomal binding domain 1	Hmgn1	
1422496_at	0.91734	high mobility group nucleosomal binding domain 1	Hmgn1	
1422497_at	0.90509	"solute carrier family 30 (zinc transporter), member 5"	Slc30a5	
1422498_at	0.94569	"melanoma antigen, family H, 1"	Mageh1	
1422499_at	0.93890	LIM domain and actin binding 1	Lima1	
1422500_at	1.01262	isocitrate dehydrogenase 3 (NAD+) alpha	Idh3a	
1422501_s_at	0.87755	isocitrate dehydrogenase 3 (NAD+) alpha	Idh3a	
1422502_at	1.03049	"poly (ADP-ribose) polymerase family, member 1"	Parp1	
1422503_s_at	1.01200	"poly (ADP-ribose) polymerase family, member 1"	Parp1	
1422504_at	1.06552	"glycine receptor, beta subunit"	Glrb	
1422505_at	1.07279	chromatin accessibility complex 1	Chrac1	
1422506_a_at	1.20639	cystatin B	Cstb	
1422507_at	1.29093	cystatin B	Cstb	
1422508_at	1.02333	"ATPase, H+ transporting, lysosomal V1 subunit A"	Atp6v1a	
1422509_at	1.11344	U2 small nuclear ribonucleoprotein auxiliary factor (U2AF) 1	U2af1	
1422510_at	1.06001	"CTD (carboxy-terminal domain, RNA polymerase II, polypeptide A) small phosphatase-like"	Ctdspl	
1422511_a_at	1.01310	opioid growth factor receptor	Ogfr	
1422512_a_at	0.96876	opioid growth factor receptor	Ogfr	
1422513_at	1.10660	cyclin F	Ccnf	
1422514_at	0.88617	AE binding protein 1	Aebp1	
1422515_at	0.98331	"seminal vesicle protein, secretion 7"	Svs7	
1422516_a_at	1.05418	fibroblast growth factor (acidic) intracellular binding protein	Fibp	
1422517_a_at	1.07544	"zinc ribbon domain containing, 1"	Znrd1	
1422518_at	0.93644	calcium/calmodulin-dependent serine protein kinase (MAGUK family)	Cask	
1422519_at	0.95176	calcium/calmodulin-dependent serine protein kinase (MAGUK family)	Cask	
1422520_at	1.02980	"neurofilament 3, medium"	Nef3	
1422521_at	0.86142	dynactin 1	Dctn1	
1422522_at	0.98370	"fragile X mental retardation gene 2, autosomal homolog"	Fxr2h	
1422523_at	0.89862	silver	Si	
1422524_at	1.05505	"ATP-binding cassette, sub-family B (MDR/TAP), member 6"	Abcb6	
1422525_at	0.87017	"ATP synthase, H+ transporting, mitochondrial F1F0 complex, subunit e"	Atp5k	
1422526_at	1.15083	acyl-CoA synthetase long-chain family member 1	Acs11	
1422527_at	1.19324	"histocompatibility 2, class II, locus DMA"	H2-DMA	
1422528_a_at	0.74727	"zinc finger protein 36, C3H type-like 1"	Zfp3611	
1422529_s_at	0.92189	calsequestrin 2	Casq2	
1422530_at	1.09143	similar to peripherin 1 LOC673890		
1422531_at	1.07014	synaptotagmin V	Syt5	
1422532_at	0.81083	"xeroderma pigmentosum, complementation group C"	Xpc	

1422533_at	1.55909	"cytochrome P450, family 51"	Cyp51
1422534_at	1.20441	"cytochrome P450, family 51"	Cyp51
1422535_at	1.01954	cyclin E2	Ccne2
1422536_at	0.91630	"troponin I, cardiac"	Tnni3
1422537_a_at	0.75078	inhibitor of DNA binding 2	Id2
1422538_at	0.96591	exotoses (multiple)-like 2	Extl2
1422539_at	0.95490	exotoses (multiple)-like 2	Extl2
1422540_at	0.86516	fibulin 1	Fbln1
1422541_at	0.72290	"protein tyrosine phosphatase, receptor type, M"	Ptprm
1422542_at	0.95522	G protein-coupled receptor 34	Gpr34
1422543_at	0.77626	"ropporin, rhophilin associated protein 1"	Ropn1
1422544_at	1.03207	myosin X	Myo10
1422545_at	0.92584	---	---
1422546_at	0.93902	interleukin enhancer binding factor 3	Ilf3
1422547_at	1.08531	RAN binding protein 1	Ranbp1
1422548_at	1.05174	dehydrodolichyl diphosphate synthase	Dhdds
1422549_at	1.06653	ADP-ribosylation factor-like 2	Arl2
1422550_a_at	1.06463	microtubule-associated protein 6	Mtap6
1422551_at	0.72845	zinc finger protein 306	Zfp306
1422552_at	1.01995	"reprimin, TP53 dependent G2 arrest mediator candidate"	Rprm
1422553_at	0.95831	phosphatase and tensin homolog	Pten
1422554_at	0.98017	necdin-like 2	Ndnl2
1422555_s_at	1.05466	"guanine nucleotide binding protein, alpha 13"	Gna13
1422556_at	1.08523	"guanine nucleotide binding protein, alpha 13"	Gna13
1422557_s_at	0.98681	metallothionein 1	Mt1
1422558_at	0.78907	guanidinoacetate methyltransferase	Gamt
1422559_at	1.15158	ubiquitin-conjugating enzyme E2N	Ube2n
1422560_at	1.07881	"DNA-damage inducible protein 2 /// regulatory solute carrier protein, family 1, member 1"	Ddi2 /// Rsc1a1
1422561_at	0.68097	"a disintegrin-like and metallopeptidase (reprolysin type) with thrombospondin type 1 motif, 5 (aggrecanase-2)"	Adamts5
1422562_at	0.93735	Ras-related associated with diabetes	Rrad
1422563_at	1.02467	complement receptor related protein	Crry
1422564_at	0.98831	actin-like 6B	Actl6b
1422565_s_at	0.88906	nuclear factor I/C	Nfic
1422566_at	0.88485	transcription factor EB	Tcf7
1422567_at	0.94249	niban protein	Niban
1422568_at	0.96614	nuclear distribution gene E-like homolog 1 (A. nidulans)	Ndel1
1422569_at	1.02320	YY1 transcription factor	Yy1
1422570_at	0.96071	---	---
1422571_at	1.06976	thrombospondin 2	Thbs2
1422572_at	0.98087	"ras homolog gene family, member G"	Rhog
1422573_at	1.24993	AMP deaminase 3	Ampd3

1422574_at 0.89186 Max dimerization protein 4 Mxd4
 1422575_at 0.93345 Max dimerization protein 4 /// similar to Max dimerization protein 4 Mxd4 ///
 LOC635153
 1422576_at 0.92993 ataxin 10 Atxn10
 1422577_at 1.02522 citrate synthase Cs
 1422578_at 1.05509 citrate synthase Cs
 1422579_at 1.12180 "heat shock protein 1 (chaperonin 10) /// similar to 10 kDa heat shock protein,
 mitochondrial (Hsp10) (10 kDa chaperonin) (CPN10) /// similar to 10 kDa heat shock protein,
 mitochondrial (Hsp10) (10 kDa chaperonin) (CPN10)" Hspe1 /// LOC628438 /// LOC636088
 1422580_at 0.82980 "myosin, light polypeptide 4" Myl4
 1422581_at 0.84012 protein inhibitor of activated STAT 1 Pias1
 1422582_at 0.99612 leptin Lep
 1422583_at 1.31085 "RAB3B, member RAS oncogene family" Rab3b
 1422584_at 1.04409 superkiller viralicidic activity 2-like (S. cerevisiae) Skiv21
 1422585_at 1.03562 outer dense fiber of sperm tails 1 Odf1
 1422586_at 0.96078 endothelin converting enzyme-like 1 Ecel1
 1422587_at 0.99891 transmembrane protein 45a Tmem45a
 1422588_at 0.94039 keratin 6B Krt6b
 1422589_at 0.83030 "RAB3A, member RAS oncogene family" Rab3a
 1422590_at 1.03300 cyclin-dependent kinase 5 Cdk5
 1422591_at 1.06344 "transcription elongation factor B (SIII), polypeptide 3" Tceb3
 1422592_at 1.03264 "catenin (cadherin associated protein), delta 2" Ctnnd2
 1422593_at 0.98620 "adaptor-related protein complex 3, sigma 1 subunit" Ap3s1
 1422594_at 1.01077 RIKEN cDNA 5730470L24 gene 5730470L24Rik
 1422595_s_at 0.95973 RIKEN cDNA 5730470L24 gene 5730470L24Rik
 1422596_at 0.84477 RIKEN cDNA C030019F02 gene C030019F02Rik
 1422597_at 0.69120 matrix metalloproteinase 15 Mmp15
 1422598_at 0.94900 calsequestrin 1 Casq1
 1422599_s_at 1.00005 zinc finger protein 143 Zfp143
 1422600_at 1.21460 RAS protein-specific guanine nucleotide-releasing factor 1 Rasgrf1
 1422601_at 1.11958 "serine (or cysteine) peptidase inhibitor, clade B, member 9" Serpinb9
 1422602_a_at 1.00321 wingless-related MMTV integration site 5B Wnt5b
 1422603_at 0.90704 "ribonuclease, RNase A family 4" Rnase4
 1422604_at 0.93051 urate oxidase Uox
 1422605_at 1.00033 "protein phosphatase 1, regulatory (inhibitor) subunit 1A" Ppp1r1a
 1422606_at 0.95862 C1q and tumor necrosis factor related protein 3 C1qtnf3
 1422607_at 0.99214 ets variant gene 1 Etv1
 1422608_at 1.02222 cAMP-regulated phosphoprotein 19 /// RIKEN cDNA 2700024H10 gene
 Arpp19 /// 2700024H10Rik
 1422609_at 0.88978 cAMP-regulated phosphoprotein 19 Arpp19
 1422610_s_at 0.99607 insulin-like growth factor 2 mRNA binding protein 3 Igf2bp3
 1422611_s_at 1.07797 insulin-like growth factor 2 mRNA binding protein 3 Igf2bp3
 1422612_at 1.85998 hexokinase 2 Hk2

1422613_a_at 1.03805 ribosomal protein L7a Rpl7a
 1422614_s_at 1.02646 "biogenesis of lysosome-related organelles complex-1, subunit 1" Bloc1s1
 1422615_at 1.13942 mitogen-activated protein kinase kinase kinase 4 Map4k4
 1422616_s_at 1.00925 WD repeat domain 54 Wdr54
 1422617_at 1.01550 "Xmr protein /// similar to Xlr-related, meiosis regulated /// similar to Xlr-related, meiosis regulated /// similar to Synaptonemal complex protein 3 (SCP-3) /// similar to Xlr-related, meiosis regulated /// similar to Xlr-related, meiosis regulated /// similar to Xlr-related, meiosis regulated /// similar to Xlr-related, meiosis regulated /// similar to Xlr-related, meiosis regulated" Xmr ///
 LOC546272 /// LOC619991 /// LOC664810 /// LOC664861 /// LOC664877 /// LOC664890 ///
 LOC664906 /// LOC664923 /// LOC664944 /// LOC664989
 1422618_x_at 1.10673 "Xmr protein /// similar to Xlr-related, meiosis regulated /// similar to Xlr-related, meiosis regulated /// similar to Synaptonemal complex protein 3 (SCP-3) /// similar to Xlr-related, meiosis regulated /// similar to Xlr-related, meiosis regulated /// similar to Xlr-related, meiosis regulated /// similar to Xlr-related, meiosis regulated /// similar to Xlr-related, meiosis regulated" Xmr ///
 LOC546272 /// LOC619991 /// LOC664810 /// LOC664861 /// LOC664877 /// LOC664890 ///
 LOC664906 /// LOC664923 /// LOC664944 /// LOC664989
 1422619_at 1.15459 phosphatidic acid phosphatase 2a Ppap2a
 1422620_s_at 1.11905 phosphatidic acid phosphatase 2a Ppap2a
 1422621_at 1.04340 RAN binding protein 2 Ranbp2
 1422622_at 1.22229 "nitric oxide synthase 3, endothelial cell" Nos3
 1422623_x_at 1.03418 Ribosomal protein L41 Rpl41
 1422624_at 1.26437 REV1-like (*S. cerevisiae*) Rev1l
 1422625_at 0.98947 "lymphocyte antigen 6 complex, locus H" Ly6h
 1422626_at 0.98850 matrix metalloproteinase 16 Mmp16
 1422627_a_at 1.00341 McKusick-Kaufman syndrome protein Mkks
 1422628_at 0.96568 RIKEN cDNA 4632417K18 gene 4632417K18Rik
 1422629_s_at 0.90321 shroom family member 3 Shroom3
 1422630_at 1.06849 RAD50 homolog (*S. cerevisiae*) Rad50
 1422631_at 0.83170 aryl-hydrocarbon receptor Ahr
 1422632_at 1.03537 cathepsin W Ctsw
 1422633_at 0.97294 RIKEN cDNA 1700020D05 gene 1700020D05Rik
 1422634_a_at 0.94854 V-set and immunoglobulin domain containing 2 Vsig2
 1422635_at 0.95999 acetylcholinesterase Ache
 1422636_at 0.88322 cyclin D binding myb-like transcription factor 1 Dmtf1
 1422637_at 0.81576 Ras association (RalGDS/AF-6) domain family 5 Rassf5
 1422638_s_at 0.85158 Ras association (RalGDS/AF-6) domain family 5 Rassf5
 1422639_at 1.00308 "calcitonin-related polypeptide, beta" Calcb
 1422640_at 0.77191 protocadherin beta 9 Pcdhb9
 1422641_at 0.92168 docking protein 5 Dok5
 1422642_at 0.76516 CDC42 effector protein (Rho GTPase binding) 3 Cdc42ep3
 1422643_at 1.03057 "monooxygenase, DBH-like 1" Moxd1

1422644_at 1.06497 SH3-binding domain glutamic acid-rich protein Sh3bgr
 1422645_at 0.96127 hemochromatosis Hfe
 1422646_at 1.00655 MAX gene associated Mga
 1422647_at 0.95079 ring finger protein 1 Ring1
 1422648_at 1.10487 "solute carrier family 7 (cationic amino acid transporter, y+ system), member 2" Slc7a2
 1422649_at 1.02366 contactin 6 Cntn6
 1422650_a_at 1.10462 RIO kinase 3 (yeast) Riok3
 1422651_at 0.94976 "adiponectin, C1Q and collagen domain containing" Adipoq
 1422652_at 1.05942 "crystallin, gamma A" Cryga
 1422653_at 0.95142 centrosomal protein 70 Cep70
 1422654_at 1.03925 "sarcoglycan, alpha (dystrophin-associated glycoprotein)" Sgca
 1422655_at 0.98698 patched homolog 2 Ptch2
 1422656_at 0.94275 "RAS-like, family 2, locus 9" Rasl2-9
 1422657_at 0.93664 prolactin-like protein L Prlpl
 1422658_at 1.00514 "phospholipase C, zeta 1" Plcz1
 1422659_at 1.09466 "calcium/calmodulin-dependent protein kinase II, delta" Camk2d
 1422660_at 1.16031 similar to Putative RNA-binding protein 3 (RNA-binding motif protein 3)
 LOC671237
 1422661_at 1.04943 "lectin, galactose binding, soluble 8" Lgals8
 1422662_at 1.05354 "lectin, galactose binding, soluble 8" Lgals8
 1422663_at 1.36690 "origin recognition complex, subunit 1-like (S.cerevisiae)" Orc11
 1422664_at 0.99462 "RAB10, member RAS oncogene family" Rab10
 1422665_a_at 0.97020 protein-L-isoaspartate (D-aspartate) O-methyltransferase 1 Pcmt1
 1422666_at 1.02863 Casitas B-lineage lymphoma c Cblc
 1422667_at 0.91282 keratin 15 Krt15
 1422668_at 1.58161 "serine (or cysteine) peptidase inhibitor, clade B, member 9b" Serpinb9b
 1422669_at 0.89880 estrogen receptor-binding fragment-associated gene 9 Ebag9
 1422670_at 0.98104 Rho family GTPase 2 Rnd2
 1422671_s_at 0.86503 N-acetylated alpha-linked acidic dipeptidase 2 Naalad2
 1422672_at 0.93316 small proline-rich protein 1B Sprr1b
 1422673_at 0.86622 "protein kinase C, mu" Prkcm
 1422674_s_at 0.98070 "crystallin, gamma B /// crystallin, gamma C" Crygb /// Crygc
 1422675_at 1.06267 "SWI/SNF related, matrix associated, actin dependent regulator of chromatin, subfamily e, member 1" Smarce1
 1422676_at 0.92850 "SWI/SNF related, matrix associated, actin dependent regulator of chromatin, subfamily e, member 1" Smarce1
 1422677_at 0.96463 diacylglycerol O-acyltransferase 2 Dgat2
 1422678_at 0.97538 diacylglycerol O-acyltransferase 2 Dgat2
 1422679_s_at 0.97180 "Ctr9, Paf1/RNA polymerase II complex component, homolog (S. cerevisiae)" Ctr9
 1422680_at 1.02775 "Ctr9, Paf1/RNA polymerase II complex component, homolog (S. cerevisiae)" Ctr9

1422681_at 0.96512 "Ctr9, Paf1/RNA polymerase II complex component, homolog (S. cerevisiae)" Ctr9
 1422682_s_at 0.90151 "protease, serine, 3 /// protease, serine, 1 (trypsin 1) /// trypsinogen 12" Prss3 /// Prss1 /// LOC436523
 1422683_at 1.07279 interleukin-1 receptor-associated kinase 1 binding protein 1 Irak1bp1
 1422684_a_at 0.90095 exocyst complex component 4 Exoc4
 1422685_at 0.99379 exocyst complex component 4 Exoc4
 1422686_s_at 0.96701 exocyst complex component 4 Exoc4
 1422687_at 1.02437 --- ---
 1422688_a_at 1.04845 neuroblastoma ras oncogene Nras
 1422689_at 0.90047 RIKEN cDNA 4930547C10 gene 4930547C10Rik
 1422690_at 0.91088 "serine palmitoyltransferase, long chain base subunit 1" Sptlc1
 1422691_at 0.96056 "serine palmitoyltransferase, long chain base subunit 1" Sptlc1
 1422692_at 0.89810 SUB1 homolog (S. cerevisia Sub1
 1422693_a_at 1.10476 SUB1 homolog (S. cerevisia Sub1
 1422694_at 0.96410 tweety homolog 1 (Drosophila) Ttyh1
 1422695_at 0.93865 "tweety homolog 1 (Drosophila) /// TAF1 RNA polymerase II, TATA box binding protein (TBP)-associated factor" Ttyh1 /// Taf1
 1422696_at 0.91847 tweety homolog 1 (Drosophila) Ttyh1
 1422697_s_at 0.99285 --- ---
 1422698_s_at 0.94089 "jumonji, AT rich interactive domain 2" Jarid2
 1422699_at 1.02867 arachidonate 12-lipoxygenase Alox12
 1422700_at 1.00340 arachidonate 12-lipoxygenase Alox12
 1422701_at 0.92776 zeta-chain (TCR) associated protein kinase Zap70
 1422702_at 0.94177 antizyme inhibitor 1 Azin1
 1422703_at 0.94171 glycerol kinase Gyk
 1422704_at 1.11499 glycerol kinase Gyk
 1422705_at 0.57493 "transmembrane, prostate androgen induced RNA" Tmepai
 1422706_at 0.73214 similar to Nedd4 WW binding protein 4 LOC637870
 1422707_at 0.95861 "phosphoinositide-3-kinase, catalytic, gamma polypeptide" Pik3cg
 1422708_at 1.08766 "phosphoinositide-3-kinase, catalytic, gamma polypeptide" Pik3cg
 1422709_a_at 1.09719 WD repeat domain 46 Wdr46
 1422710_a_at 0.96168 "calcium channel, voltage-dependent, T type, alpha 1H subunit" Cacna1h
 1422711_a_at 1.23811 pregnancy upregulated non-ubiquitously expressed CaM kinase Pnck
 1422712_a_at 1.04249 ubiquitin-conjugating enzyme E2I Ube2i
 1422713_a_at 0.91996 ubiquitin-conjugating enzyme E2I Ube2i
 1422714_at 0.91530 "ubiquitin-conjugating enzyme E2I /// similar to Chain A, Human Ubiquitin-Conjugating Enzyme Ubc9 /// similar to Ubiquitin-conjugating enzyme E2 I (Ubiquitin-protein ligase I) (Ubiquitin carrier protein I) (SUMO-1-protein ligase) (SUMO-1-conjugating enzyme) (Ubiquitin-conjugating enzyme UbcE2A)" Ube2i /// LOC546265 /// LOC669417
 1422715_s_at 1.01624 "acid phosphatase 1, soluble /// similar to acid phosphatase 1, soluble /// similar to acid phosphatase 1 isoform b /// similar to acid phosphatase 1 isoform b /// similar to acid phosphatase 1 isoform b" Acp1 /// LOC631286 /// LOC668486 /// LOC670469 /// LOC676625

1422716_a_at	1.04701	"acid phosphatase 1, soluble"	Acp1	
1422717_at	0.84951	"acid phosphatase 1, soluble"	Acp1	
1422718_at	0.88485	"adaptor-related protein complex 3, sigma 2 subunit"	Ap3s2	
1422719_s_at	0.98277	nucleoporin 50	Nup50	
1422720_at	1.09066	"ISL1 transcription factor, LIM/homeodomain"	Isl1	
1422721_at	0.88287	"tyrosine kinase, non-receptor, 1"	Tnk1	
1422722_at	1.08770	RIKEN cDNA 1700001K19 gene	1700001K19Rik	
1422723_at	1.07242	stimulated by retinoic acid gene 6	Stra6	
1422724_at	0.92193	male germ cell-associated kinase	Mak	
1422725_at	1.00168	male germ cell-associated kinase	Mak	
1422726_x_at	1.05289	spermatogenesis associated glutamate (E)-rich protein 4a	Speer4a	
1422727_at	0.84085	expressed in non-metastatic cells 5	Nme5	
1422728_at	0.86276	inhibin alpha	Inha	
1422729_at	1.08078	protocadherin beta 10	Pcdhb10	
1422730_at	0.88153	LIM domains containing 1	Limd1	
1422731_at	0.84136	LIM domains containing 1	Limd1	
1422732_at	0.94016	"polymerase (DNA-directed), delta interacting protein 2"	Poldip2	
1422733_at	1.00217	four jointed box 1 (Drosophila)	Fjx1	
1422734_a_at	0.96889	myeloblastosis oncogene	Myb	
1422735_at	0.74873	forkhead box Q1	Foxq1	
1422736_at	0.88636	RAN binding protein 9	Ranbp9	
1422737_at	1.03013	nuclear receptor coactivator 3	Ncoa3	
1422738_at	1.24044	"discoidin domain receptor family, member 2"	Ddr2	
1422739_at	0.90288	Heparan sulfate 2-O-sulfotransferase 1	Hs2st1	
1422740_at	0.97747	"tumor necrosis factor receptor superfamily, member 21"	Tnfrsf21	
1422741_a_at	0.94644	bobby sox homolog (Drosophila)	Bbx	
1422742_at	1.08579	human immunodeficiency virus type I enhancer binding protein 1	Hivep1	
1422743_at	1.13211	phosphorylase kinase alpha 1	Phka1	
1422744_at	0.97099	phosphorylase kinase alpha 1	Phka1	
1422745_at	0.91072	bicaudal D homolog 2 (Drosophila)	Bicd2	
1422746_s_at	1.05194	bicaudal D homolog 2 (Drosophila)	Bicd2	
1422747_at	1.01738	CHK2 checkpoint homolog (S. pombe)	Chek2	
1422748_at	0.94461	zinc finger homeobox 1b	Zfx1b	
1422749_at	0.95519	"lymphocyte antigen 6 complex, locus G6C"	Ly6g6c	
1422750_a_at	0.95165	"zinc finger, MYND domain containing 10"	Zmynd10	
1422751_at	0.84374	"transducin-like enhancer of split 1, homolog of Drosophila E(spl)"	Tle1	
1422752_at	0.93009	polymerase (RNA) III (DNA directed) polypeptide K	Polr3k	
1422753_a_at	1.05748	polymerase (RNA) III (DNA directed) polypeptide K	Polr3k	
1422754_at	0.84219	---	---	
1422755_at	0.86715	Bruton agammaglobulinemia tyrosine kinase	Btk	
1422756_at	1.08443	"solute carrier family 32 (GABA vesicular transporter), member 1"	Slc32a1	
1422757_at	1.07522	"solute carrier family 5 (neutral amino acid transporters, system A), member 4b"	Slc5a4b	

1422758_at 1.00820 carbohydrate sulfotransferase 2 Chst2
 1422759_a_at 0.97176 exportin 6 Xpo6
 1422760_at 1.11143 "peptidyl arginine deiminase, type IV /// similar to Protein-arginine deiminase type IV (Peptidylarginine deiminase IV)" Padi4 /// LOC670543
 1422761_at 0.90473 transducin (beta)-like 1X-linked receptor 1 Tbl1xr1
 1422762_at 0.93667 kinesin family member 17 Kif17
 1422763_at 0.99245 "GIPC PDZ domain containing family, member 1" Gipc1
 1422764_at 1.03032 "microtubule-associated protein, RP/EB family, member 1" Mapre1
 1422765_at 0.94713 "microtubule-associated protein, RP/EB family, member 1" Mapre1
 1422766_at 0.97427 staufen (RNA binding protein) homolog 1 (Drosophila) Stau1
 1422767_at 1.07141 bystin-like Bysl
 1422768_at 0.99051 "synaptotagmin binding, cytoplasmic RNA interacting protein" Syncrip
 1422769_at 1.03680 "synaptotagmin binding, cytoplasmic RNA interacting protein" Syncrip
 1422770_at 1.06693 RAD51-like 3 (S. cerevisiae) Rad51l3
 1422771_at 0.67822 similar to Mothers against decapentaplegic homolog 6 (SMAD 6) (Mothers against DPP homolog 6) (Smad6) (Mad homolog 7) LOC670044
 1422772_at 0.93465 "core 1 UDP-galactose:N-acetylgalactosamine-alpha-R beta 1,3-galactosyltransferase" C1galt1
 1422773_at 0.89436 myelin transcription factor 1 Myt1
 1422774_at 0.75676 RIKEN cDNA 1700088E04 gene 1700088E04Rik
 1422775_at 0.88215 B lymphoid kinase Blk
 1422776_at 0.93530 "serine (or cysteine) peptdiase inhibitor, clade B, member 8" Serpinb8
 1422777_at 0.94838 "complement component 1, q subcomponent-like 1" C1ql1
 1422778_at 1.06020 "TAF9 RNA polymerase II, TATA box binding protein (TBP)-associated factor" Taf9
 1422779_at 0.98362 "sphingomyelin phosphodiesterase 3, neutral" Smpd3
 1422780_at 1.01140 peroxisomal membrane protein 4 Pxmp4
 1422781_at 0.98814 toll-like receptor 3 Tlr3
 1422782_s_at 0.92477 toll-like receptor 3 Tlr3
 1422783_a_at 0.93198 keratin 6A Krt6a
 1422784_at 0.93295 keratin 6A Krt6a
 1422785_at 1.19080 RAS p21 protein activator 2 Rasa2
 1422786_at 1.10714 "solute carrier family 30 (zinc transporter), member 1" Slc30a1
 1422787_at 0.98192 FK506 binding protein-like Fkbpl
 1422788_at 1.10546 "solute carrier family 43, member 3" Slc43a3
 1422789_at 0.95671 "aldehyde dehydrogenase family 1, subfamily A2" Aldh1a2
 1422790_at 0.95565 natriuretic peptide precursor type C Nppc
 1422791_at 1.10896 "platelet-activating factor acetylhydrolase, isoform 1b, alpha2 subunit" Pafah1b2
 1422792_at 1.04317 "platelet-activating factor acetylhydrolase, isoform 1b, alpha2 subunit" Pafah1b2
 1422793_at 0.97416 "platelet-activating factor acetylhydrolase, isoform 1b, alpha2 subunit" Pafah1b2

1422794_at	0.98327	cullin 3	Cul3	
1422795_at	0.86503	cullin 3	Cul3	
1422796_at	1.10517	prolyl endopeptidase	Prep	
1422797_at	1.05939	mitogen activated protein binding protein interacting protein	Mapbpip	
1422798_at	1.16706	contactin associated protein-like 2	Cntnap2	
1422799_at	0.97360	HLA-B associated transcript 2	Bat2	
1422800_at	0.86364	HLA-B associated transcript 2	Bat2	
1422801_at	1.10053	Ras-GTPase-activating protein SH3-domain binding protein		RP23-336J1.4
1422802_at	1.02836	defensin related cryptdin 3	Defcr3	
1422803_at	0.75644	follistatin-like 3	Fstl3	
1422804_at	1.08989	"serine (or cysteine) peptidase inhibitor, clade B, member 6b"	Serpinb6b	
1422805_a_at	0.88283	"inhibitor of growth family, member 3"	Ing3	
1422806_x_at	0.85080	"inhibitor of growth family, member 3"	Ing3	
1422807_at	0.91066	ADP-ribosylation factor 5	Arf5	
1422808_s_at	0.89491	dedicator of cyto-kinesis 2	Dock2	
1422809_at	0.98234	regulating synaptic membrane exocytosis 2	Rims2	
1422810_at	0.76683	zinc finger protein 191	Zfp191	
1422811_at	0.98548	"solute carrier family 27 (fatty acid transporter), member 1"	Slc27a1	
1422812_at	1.02472	chemokine (C-X-C motif) receptor 6	Cxcr6	
1422813_at	1.01748	"calcium channel, voltage-dependent, gamma subunit 1"	Cacng1	
1422814_at	1.03982	"asp (abnormal spindle)-like, microcephaly associated (Drosophila)"	Aspm	
1422815_at	0.97607	complement component 9	C9	
1422816_a_at	1.00355	mutY homolog (E. coli)	Mutyh	
1422817_at	0.98851	glycoprotein 5 (platelet)	Gp5	
1422818_at	0.84361	"neural precursor cell expressed, developmentally down-regulated gene 9"	Nedd9	
1422819_at	0.97965	mitochondrial ribosomal protein L36	Mrpl36	
1422820_at	0.85568	"lipase, hormone sensitive"	Lipe	
1422821_s_at	0.94109	StAR-related lipid transfer (START) domain containing 5	Stard5	
1422822_at	0.95583	StAR-related lipid transfer (START) domain containing 5	Stard5	
1422823_at	1.30166	epidermal growth factor receptor pathway substrate 8	Eps8	
1422824_s_at	1.31315	epidermal growth factor receptor pathway substrate 8	Eps8	
1422825_at	0.91958	cocaine and amphetamine regulated transcript	Cart	
1422826_at	1.02405	"insulin-like growth factor binding protein, acid labile subunit"	Igfals	
1422827_x_at	0.93208	"solute carrier family 35 (UDP-galactose transporter), member 2"		Slc35a2
1422828_at	0.90247	"CD3 antigen, delta polypeptide"	Cd3d	
1422829_at	0.99984	dopamine receptor 4	Drd4	
1422830_s_at	0.97789	dopamine receptor 4	Drd4	
1422831_at	1.03490	fibrillin 2 /// similar to fibrillin 2	Fbn2 ///	LOC639042
1422832_at	0.96679	retinal G protein coupled receptor	Rgr	
1422833_at	0.80633	forkhead box A2	Foxa2	
1422834_at	0.92866	"potassium voltage-gated channel, Shal-related family, member 2"		Kcnd2
1422835_at	1.00976	"potassium voltage-gated channel, Shal-related family, member 2"		Kcnd2

1422836_at 0.89999 muscleblind-like 3 (Drosophila) Mbnl3
1422837_at 1.06894 sciellin Scel
1422838_at 0.94194 "potassium channel, subfamily U, member 1" Kcnu1
1422839_at 0.96931 neurogenin 2 Neurog2
1422840_at 1.05034 RIKEN cDNA 1700013N18 gene 1700013N18Rik
1422841_at 0.92162 "1-acylglycerol-3-phosphate O-acyltransferase 6 (lysophosphatidic acid acyltransferase, zeta)" Agpat6
1422842_at 1.14455 5'-3' exoribonuclease 2 Xrn2
1422843_at 1.09309 --- ---
1422844_a_at 1.03901 WD repeat domain 77 Wdr77
1422845_at 0.98531 calnexin Canx
1422846_at 1.02875 "retinol binding protein 2, cellular" Rbp2
1422847_a_at 0.93359 "protein kinase C, delta" Prkcd
1422848_a_at 1.12178 "poly(A) binding protein, nuclear 1" Pabpn1
1422849_a_at 1.02270 "poly(A) binding protein, nuclear 1" Pabpn1
1422850_at 1.23947 "poly(A) binding protein, nuclear 1" Pabpn1
1422851_at 1.16590 high mobility group AT-hook 2 Hmga2
1422852_at 0.71761 calcium and integrin binding family member 2 Cib2
1422853_at 0.99805 src homology 2 domain-containing transforming protein C1 Shc1
1422854_at 1.02686 src homology 2 domain-containing transforming protein C1 Shc1
1422855_at 1.06746 cleavage and polyadenylation specificity factor 3 Cpsf3
1422856_at 0.73703 "solute carrier family 12, member 3" Slc12a3
1422857_at 1.05202 thyroid hormone receptor interactor 4 Trip4
1422858_at 0.95343 thyroid hormone receptor interactor 4 Trip4
1422859_a_at 1.27977 ribosomal protein L23 Rpl23
1422860_at 0.93075 neurotensin Nts
1422861_s_at 0.85872 similar to PDZ and LIM domain protein 5 (Enigma homolog) (Enigma-like PDZ and LIM domains protein) LOC669660
1422862_at 0.75626 similar to PDZ and LIM domain protein 5 (Enigma homolog) (Enigma-like PDZ and LIM domains protein) LOC669660
1422863_s_at 0.78268 similar to PDZ and LIM domain protein 5 (Enigma homolog) (Enigma-like PDZ and LIM domains protein) LOC669660
1422864_at 1.05777 runt related transcription factor 1 Runx1
1422865_at 1.33967 runt related transcription factor 1 Runx1
1422866_at 0.97547 "procollagen, type XIII, alpha 1" Col13a1
1422867_at 1.11140 granzyme G Gzmg
1422868_s_at 1.00140 guanine deaminase Gda
1422869_at 0.95235 c-mer proto-oncogene tyrosine kinase Mertk
1422870_at 1.00867 homeo box C4 Hoxc4
1422871_at 0.99147 "potassium inwardly-rectifying channel, subfamily J, member 12" Kcnj12
1422872_at 0.91876 "bone morphogenetic protein receptor, type 1B" Bmpr1b
1422873_at 0.96219 "proteoglycan 2, bone marrow" Prg2
1422874_at 0.90153 proacrosin binding protein Acrbp

1422875_at	1.01147	CD84 antigen	Cd84	
1422876_at	1.02142	calpain 9 (nCL-4) /// similar to calpain 9 (nCL-4)	Capn9	LOC639715
1422877_at	0.93691	protocadherin beta 12	Pcdhb12	
1422878_at	0.93643	synaptotagmin XII	Syt12	
1422879_at	0.94720	synaptophysin-like protein	Sypl	
1422880_at	0.99654	synaptophysin-like protein	Sypl	
1422881_s_at	0.98356	synaptophysin-like protein	Sypl	
1422882_at	0.91969	synaptophysin-like protein	Sypl	
1422883_at	0.95513	ubiquitin specific peptidase 8	Usp8	
1422884_at	1.16711	small nuclear ribonucleoprotein D3	Snrpd3	
1422885_at	1.09468	small nuclear ribonucleoprotein D3	Snrpd3	
1422886_a_at	0.99917	CDC like kinase 4	Clk4	
1422887_a_at	0.91881	C-terminal binding protein 2	Ctbp2	
1422888_at	0.92686	ring finger protein 5	Rnf5	
1422889_at	0.77761	protocadherin 18	Pcdh18	
1422890_at	0.76008	protocadherin 18	Pcdh18	
1422891_at	0.96701	"histocompatibility 2, class II antigen E alpha"	H2-Ea	
1422892_s_at	1.06112	"histocompatibility 2, class II antigen E alpha"	H2-Ea	
1422893_at	0.85394	Scm-like with four mbt domains 1	Sfmbt1	
1422894_at	0.91120	Scm-like with four mbt domains 1	Sfmbt1	
1422895_at	0.98456	vesicle-associated membrane protein 4	Vamp4	
1422896_at	1.10956	---	---	
1422897_at	0.97265	"solute carrier family 22 (organic anion/cation transporter), member 12"		
Slc22a12				
1422898_s_at	0.77040	"solute carrier family 22 (organic anion/cation transporter), member 12"		
Slc22a12				
1422899_at	0.90216	"solute carrier family 6 (neurotransmitter transporter), member 20"	Slc6a20	
1422900_at	0.94486	meningioma expressed antigen 5 (hyaluronidase)	Mgea5	
1422901_at	0.92397	meningioma expressed antigen 5 (hyaluronidase)	Mgea5	
1422902_s_at	0.95331	meningioma expressed antigen 5 (hyaluronidase)	Mgea5	
1422903_at	1.10064	lymphocyte antigen 86	Ly86	
1422904_at	0.90174	flavin containing monooxygenase 2	Fmo2	
1422905_s_at	1.00506	flavin containing monooxygenase 2	Fmo2	
1422906_at	0.82496	"ATP-binding cassette, sub-family G (WHITE), member 2"	Abcg2	
1422907_at	0.94940	"guanine nucleotide binding protein, alpha transducing 2"	Gnat2	
1422908_at	0.96804	"ATPase, (Na+)/K+ transporting, beta 4 polypeptide"	Atp1b4	
1422909_at	1.13598	structural maintenance of chromosomes 6	Smc6	
1422910_s_at	0.93766	structural maintenance of chromosomes 6	Smc6	
1422911_at	0.89872	cerebellin 3 precursor protein	Cbln3	
1422912_at	1.10765	bone morphogenetic protein 4	Bmp4	
1422913_at	1.02451	protamine 3	Prm3	
1422914_at	0.95502	trans-acting transcription factor 5	Sp5	
1422915_at	1.09704	gastrin	Gast	

1422916_at 0.96547 fibroblast growth factor 21 Fgf21
1422917_at 0.89612 Eph receptor A1 EphA1
1422918_at 0.93986 RIKEN cDNA 1810009J06 gene 1810009J06Rik
1422919_at 0.84021 HRAS-like suppressor Hrasls
1422920_at 0.96364 claudin 13 Cldn13
1422921_at 0.89655 pre-B lymphocyte gene 3 Vpreb3
1422922_at 1.12410 RecQ protein-like 4 Recql4
1422923_at 0.99613 fibroblast growth factor 3 Fgf3
1422924_at 1.53340 "tumor necrosis factor (ligand) superfamily, member 9" Tnfsf9
1422925_s_at 1.06013 acyl-CoA thioesterase 3 Acot3
1422926_at 1.15268 melanocortin 2 receptor Mc2r
1422927_at 0.99124 "Yip1 domain family, member 7" Yipf7
1422928_at 0.95424 "elastase 2, neutrophil" Ela2
1422929_s_at 0.97741 atonal homolog 7 (Drosophila) Atoh7
1422930_at 0.85792 "intercellular adhesion molecule 4, Landsteiner-Wiener blood group" Icam4
1422931_at 1.13149 fos-like antigen 2 Fosl2
1422932_a_at 0.91997 vav 1 oncogene Vav1
1422933_at 0.90239 X-linked lymphocyte-regulated 5D /// similar to X-linked lymphocyte-regulated 5C Xlr5d /// LOC627081
1422934_x_at 1.08154 "defensin related cryptdin, related sequence 7" Defcr-rs7
1422935_x_at 1.02131 chromobox homolog 1 (Drosophila HP1 beta) Cbx1
1422936_at 0.95156 MAS1 oncogene Mas1
1422937_at 0.91805 frizzled homolog 5 (Drosophila) Fzd5
1422938_at 0.88842 B-cell leukemia/lymphoma 2 Bcl2
1422939_at 0.98373 "serine (or cysteine) peptidase inhibitor, clade B (ovalbumin), member 3B" Serpinb3b
1422940_x_at 1.12278 "serine (or cysteine) peptidase inhibitor, clade B (ovalbumin), member 3B" Serpinb3b
1422941_at 1.03273 wingless-related MMTV integration site 16 Wnt16
1422942_at 0.94599 galanin receptor 2 Galr2
1422943_a_at 0.69132 heat shock protein 1 Hspb1
1422944_a_at 1.09314 diaphanous homolog 3 (Drosophila) Diap3
1422945_a_at 0.95361 kinesin family member 5C Kif5c
1422946_a_at 0.96493 DNA methyltransferase (cytosine-5) 1 Dnmt1
1422947_at 1.07801 "histone 2, H4 /// histone 1, H4f /// histone 1, H4m /// histone 1, H4a /// histone 1, H4b" Hist2h4 /// Hist1h4f /// Hist1h4m /// Hist1h4a /// Hist1h4b
1422948_s_at 1.13518 "histone 1, H4h /// histone 2, H4 /// histone 1, H4c /// histone 1, H4i /// histone 1, H4j /// histone 1, H4k /// histone 1, H4m /// histone 1, H4a /// histone 1, H4b" Hist1h4h /// Hist2h4 /// Hist1h4c /// Hist1h4i /// Hist1h4j /// Hist1h4k /// Hist1h4m /// Hist1h4a /// Hist1h4b
1422949_at 1.01379 "nitric oxide synthase 1, neuronal" Nos1
1422950_at 0.96545 hairy and enhancer of split 7 (Drosophila) Hes7
1422951_at 0.86517 protocadherin beta 4 Pcdhb4
1422952_at 1.03452 Ng23 protein Ng23

1422953_at 1.08069 "formyl peptide receptor, related sequence 2" Fpr-rs2
1422954_at 0.91014 zinc finger protein 60 Zfp60
1422955_at 1.04232 synaptotagmin XVII Syt17
1422956_at 1.01238 "DNA segment, Chr 1, Pasteur Institute 1" D1Pas1
1422957_at 1.02527 chemokine (C-C motif) receptor 3 Ccr3
1422958_at 0.96247 keratin associated protein 5-4 Krtap5-4
1422959_s_at 0.88710 zinc finger protein 313 Zfp313
1422960_at 0.92074 steroid 5 alpha-reductase 2 Srd5a2
1422961_at 1.08446 N-acetyltransferase 3 Nat3
1422962_a_at 1.48513 "proteasome (prosome, macropain) subunit, beta type 8 (large multifunctional peptidase 7)" Psm8
1422963_at 0.91164 small proline-rich protein 2I Sprr2i
1422964_at 1.04236 RAD23a homolog (S. cerevisiae) Rad23a
1422965_at 0.96348 "angiotensin II, type I receptor-associated protein" Agrap
1422966_a_at 1.02573 transferrin receptor Tfrc
1422967_a_at 1.04450 transferrin receptor Tfrc
1422968_at 0.81814 inositol hexaphosphate kinase 1 Ihpk1
1422969_s_at 0.92717 inositol hexaphosphate kinase 1 Ihpk1
1422970_at 0.84110 Max dimerization protein 3 Mxd3
1422971_at 1.06206 GCN5 general control of amino acid synthesis-like 2 (yeast) Gcn512
1422972_s_at 1.15768 GCN5 general control of amino acid synthesis-like 2 (yeast) Gcn512
1422973_a_at 1.04652 thyroid hormone responsive SPOT14 homolog (Rattus) Thrsp
1422974_at 0.82335 "5' nucleotidase, ecto" Nt5e
1422975_at 1.07077 membrane metallo endopeptidase Mme
1422976_x_at 1.00168 "NADH dehydrogenase (ubiquinone) 1 alpha subcomplex, 7 (B14.5a)" Ndufa7
1422977_at 0.98447 "glycoprotein Ib, beta polypeptide" Gp1bb
1422978_at 1.10553 "cytochrome b-245, beta polypeptide" Cybb
1422979_at 1.06284 suppressor of variegation 3-9 homolog 2 (Drosophila) Suv39h2
1422980_a_at 0.99771 blocked early in transport 1 homolog (S. cerevisiae)-like Bet11
1422981_at 0.95417 DiGeorge syndrome critical region gene 8 Dgcr8
1422982_at 1.01361 androgen receptor Ar
1422983_at 0.89364 integrin beta 6 Itgb6
1422984_at 1.13460 cytoplasmic linker 2 Cyln2
1422985_at 0.96733 frizzled homolog 1 (Drosophila) Fzd1
1422986_at 1.05748 "estrogen related receptor, beta" Esrrb
1422987_at 1.01665 netrin 1 Ntn1
1422988_at 1.09103 N-sulfoglucosamine sulfohydrolase (sulfamidase) Sgsh
1422989_a_at 0.93121 dystrophin myotonic-protein kinase Dmpk
1422990_at 1.09576 met proto-oncogene Met
1422991_at 1.02316 phosphatidylinositol 3-kinase catalytic delta polypeptide Pik3cd
1422992_s_at 1.08586 phosphatidylinositol 3-kinase catalytic delta polypeptide Pik3cd
1422993_s_at 1.08660 THO complex 4 /// RNA and export factor binding protein 2 Thoc4 ///

Refbp2

1422994_at	0.99261	"phosphatidylinositol-3-phosphate/phosphatidylinositol 5-kinase, type III"	
Pip5k3			
1422995_at	0.80169	Bcl2 modifying factor	Bmf
1422996_at	1.02003	acyl-CoA thioesterase 2	Acot2
1422997_s_at	0.89538	acyl-CoA thioesterase 1 /// acyl-CoA thioesterase 2	Acot1 /// Acot2
1422998_a_at	0.92776	glutaredoxin 2 (thioltransferase)	Glx2
1422999_at	1.01972	mitogen-activated protein kinase kinase kinase 14	Map3k14
1423000_a_at	0.87422	"diacylglycerol kinase, epsilon"	Dgke
1423001_at	0.92104	"sprouty-related, EVH1 domain containing 2"	Spred2
1423002_at	0.93589	phosphoprotein associated with glycosphingolipid microdomains 1	Pag1
1423003_at	0.92204	SET domain containing (lysine methyltransferase) 7	Setd7
1423004_at	0.91121	vasoactive intestinal peptide receptor 1	Vipr1
1423005_a_at	0.97187	espin	Espn
1423006_at	1.03875	proviral integration site 1	Pim1
1423007_a_at	0.93379	glial cell line derived neurotrophic factor family receptor alpha 2	Gfra2
1423008_at	1.00094	ADP-ribosyltransferase 2a	Art2a
1423009_at	1.07491	secretory blood group 1	Sec1
1423010_at	1.00377	sacsin	Sacs
1423011_at	0.91845	ecotropic viral integration site 1	Evi1
1423012_at	0.97482	synaptotagmin VII	Syt7
1423013_at	0.83345	forkhead box F1a	Foxf1a
1423014_at	1.11165	protocadherin beta 6	Pcdhb6
1423015_at	0.91878	kin of IRRE like 1 (Drosophila)	Kirrel1
1423016_a_at	0.84172	glycophorin A	Gypa
1423017_a_at	1.07801	interleukin 1 receptor antagonist	Il1rn
1423018_at	0.91978	"potassium voltage-gated channel, shaker-related subfamily, member 3"	Kcna3
1423019_at	1.03689	gap junction membrane channel protein alpha 9	Gja9
1423020_at	0.83858	---	---
1423021_s_at	0.98498	insulin-like 3 /// Janus kinase 3	Insl3 /// Jak3
1423022_at	1.05532	"adrenergic receptor, alpha 2a"	Adra2a
1423023_at	0.98858	secreted frizzled-related sequence protein 5	Sfrp5
1423024_at	0.79036	SH2 domain protein 1B1	Sh2d1b1
1423025_a_at	0.89398	schwannomin interacting protein 1	Schip1
1423026_at	0.83060	Rad51 homolog c (S. cerevisiae)	Rad51c
1423027_at	1.19049	forkhead box L1	Foxl1
1423028_at	0.89132	interferon alpha 2	Ifna2
1423029_at	0.90686	hairy and enhancer of split 2 (Drosophila)	Hes2
1423030_at	1.11507	valosin containing protein /// similar to Transitional endoplasmic reticulum ATPase (TER ATPase) (15S Mg(2+)-ATPase p97 subunit) (Valosin-containing protein) (VCP)	Vcp /// LOC675857
1423031_at	0.95649	macrophage erythroblast attacher	Maea
1423032_at	1.06671	ribosomal protein L39	Rpl39

1423033_at 1.04215 "STT3, subunit of the oligosaccharyltransferase complex, homolog A (S. cerevisiae)" Stt3a

1423034_at 1.07116 thioredoxin-like 5 Txnl5

1423035_s_at 1.11030 thioredoxin-like 5 Txnl5

1423036_at 0.87229 thioredoxin-like 5 Txnl5

1423037_at 0.96019 angiotensin receptor-like 1 Agtrl1

1423038_at 0.98223 syntaxin 6 Stx6

1423039_a_at 0.95984 --- ---

1423040_at 1.13482 basic leucine zipper and W2 domains 1 Bzw1

1423041_a_at 1.05545 basic leucine zipper and W2 domains 1 Bzw1

1423042_at 1.00706 "DEAD/H (Asp-Glu-Ala-Asp/His) box polypeptide 3, X-linked" Ddx3x

1423043_s_at 1.01525 "DEAD/H (Asp-Glu-Ala-Asp/His) box polypeptide 3, X-linked" Ddx3x

1423044_at 1.01455 proline synthetase co-transcribed Prosc

1423045_at 0.95862 nuclear cap binding protein subunit 2 Ncbp2

1423046_s_at 0.87246 nuclear cap binding protein subunit 2 Ncbp2

1423047_at 1.00138 toll interacting protein Tollip

1423048_a_at 0.84967 toll interacting protein Tollip

1423049_a_at 0.92873 "tropomyosin 1, alpha" Tpm1

1423050_s_at 0.95211 heterogeneous nuclear ribonucleoprotein U Hnrpu

1423051_at 1.13762 heterogeneous nuclear ribonucleoprotein U Hnrpu

1423052_at 0.98138 ADP-ribosylation factor 4 Arf4

1423053_at 0.92038 ADP-ribosylation factor 4 Arf4

1423054_at 0.96726 WD repeat domain 1 Wdr1

1423055_at 0.92580 neuron specific gene family member 1 Nsg1

1423056_at 0.98407 neuron specific gene family member 1 Nsg1

1423057_at 1.01135 "capping protein (actin filament) muscle Z-line, alpha 2" Capza2

1423058_at 1.07045 "capping protein (actin filament) muscle Z-line, alpha 2" Capza2

1423059_at 1.03853 PTK2 protein tyrosine kinase 2 Ptk2

1423060_at 1.15671 proliferation-associated 2G4 Pa2g4

1423061_at 0.81342 armadillo repeat gene deleted in velo-cardio-facial syndrome Arvcf

1423062_at 0.99824 insulin-like growth factor binding protein 3 Igfbp3

1423063_at 0.74538 DNA methyltransferase 3A Dnmt3a

1423064_at 0.98785 DNA methyltransferase 3A Dnmt3a

1423065_at 0.62061 DNA methyltransferase 3A Dnmt3a

1423066_at 0.87415 DNA methyltransferase 3A Dnmt3a

1423067_at 0.95037 CDK5 regulatory subunit associated protein 3 Cdk5rap3

1423068_at 0.74841 intraflagellar transport 172 homolog (Chlamydomonas) Ift172

1423069_at 0.89451 activity-dependent neuroprotective protein Adnp

1423070_at 0.91802 ribosomal protein L21 Rpl21

1423071_x_at 0.79828 hypothetical gene supported by BC019681; BC027236 /// hypothetical LOC434328 /// hypothetical LOC544983 /// hypothetical LOC545175 /// hypothetical LOC619711 /// hypothetical LOC622782 /// hypothetical protein LOC623857 /// hypothetical LOC624831 /// hypothetical protein LOC665837 /// hypothetical protein LOC666017 /// hypothetical protein

LOC666200 /// hypothetical protein LOC666464 /// hypothetical protein LOC668524 /// hypothetical protein LOC670117 /// hypothetical protein LOC670242 LOC270335 /// LOC434328 /// LOC544983 /// LOC545175 /// LOC619711 /// LOC622782 /// LOC623857 /// LOC624831 /// LOC665837 /// LOC666017 /// LOC666200 /// LOC666464 /// LOC668524 /// LOC670117 /// LOC670242
 1423072_at 0.80948 RIKEN cDNA 6720475J19 gene /// similar to putative retrovirus-related gag protein 6720475J19Rik /// LOC670480
 1423073_at 0.92812 cytidylate kinase Cmpk
 1423074_at 0.80108 "lectin, mannose-binding 2" Lman2
 1423075_at 0.80541 "lectin, mannose-binding 2" Lman2
 1423076_at 1.00038 --- ---
 1423077_at 0.98966 sorting nexin 9 Snx9
 1423078_a_at 1.57335 sterol-C4-methyl oxidase-like Sc4mol
 1423079_a_at 1.10249 translocase of outer mitochondrial membrane 20 homolog (yeast) /// similar to translocase of outer mitochondrial membrane 20 homolog /// similar to translocase of outer mitochondrial membrane 20 homolog Tomm20 /// LOC546321 /// LOC673774
 1423080_at 1.13826 translocase of outer mitochondrial membrane 20 homolog (yeast) Tomm20
 1423081_a_at 1.00563 translocase of outer mitochondrial membrane 20 homolog (yeast) Tomm20
 1423082_at 0.88221 "Der1-like domain family, member 1" Der1
 1423083_at 0.84246 "RAB33B, member of RAS oncogene family" Rab33b
 1423084_at 1.04577 "UDP-Gal:betaGlcNAc beta 1,3-galactosyltransferase, polypeptide 2" B3galt2
 1423085_at 0.86435 ephrin B3 Efnb3
 1423086_at 1.08057 Niemann Pick type C1 Npc1
 1423087_a_at 1.09056 RIKEN cDNA 1110002E23 gene 1110002E23Rik
 1423088_at 0.89639 tropomodulin 3 Tmod3
 1423089_at 0.96552 tropomodulin 3 Tmod3
 1423090_x_at 1.03875 "SEC61, gamma subunit" Sec61g
 1423091_a_at 1.00602 glycoprotein m6b Gpm6b
 1423092_at 1.07117 inner centromere protein Incenp
 1423093_at 0.96387 inner centromere protein Incenp
 1423094_at 0.92392 cereblon Crbn
 1423095_s_at 0.90265 cereblon Crbn
 1423096_at 1.00200 calpain 7 Capn7
 1423097_s_at 1.01121 calpain 7 Capn7
 1423098_at 0.95943 calpain 7 Capn7
 1423099_a_at 0.94531 methyltransferase-like 3 Mettl3
 1423100_at 0.43932 FBJ osteosarcoma oncogene Fos
 1423101_at 0.77203 progesterin and adipoQ receptor family member IV Paqr4
 1423102_a_at 0.94723 ring finger protein 10 Rnf10
 1423103_at 0.97168 "regulatory factor X, 5 (influences HLA class II expression)" Rfx5
 1423104_at 0.76445 insulin receptor substrate 1 Irs1
 1423105_a_at 0.96633 YEATS domain containing 4 Yeats4

1423106_at 0.87717 "ubiquitin-conjugating enzyme E2B, RAD6 homology (S. cerevisiae)" Ube2b
 1423107_at 1.03716 "ubiquitin-conjugating enzyme E2B, RAD6 homology (S. cerevisiae)" Ube2b
 1423108_at 1.03442 "solute carrier family 25 (mitochondrial carnitine/acylcarnitine translocase), member 20" Slc25a20
 1423109_s_at 0.98388 "solute carrier family 25 (mitochondrial carnitine/acylcarnitine translocase), member 20" Slc25a20
 1423110_at 1.02728 "procollagen, type I, alpha 2" Col1a2
 1423111_at 0.95663 "ATP synthase, H⁺ transporting, mitochondrial F1 complex, alpha subunit, isoform 1" Atp5a1
 1423112_at 0.84967 "ubiquitin-conjugating enzyme E2D 3 (UBC4/5 homolog, yeast)" Ube2d3
 1423113_a_at 1.01554 "ubiquitin-conjugating enzyme E2D 3 (UBC4/5 homolog, yeast)" Ube2d3
 1423114_at 0.85573 "ubiquitin-conjugating enzyme E2D 3 (UBC4/5 homolog, yeast)" Ube2d3
 1423115_at 0.82406 "ST6 (alpha-N-acetyl-neuraminy-2,3-beta-galactosyl-1,3)-N-acetylgalactosaminide alpha-2,6-sialyltransferase 6" St6galnac6
 1423116_at 1.02154 DOM-3 homolog Z (C. elegans) Dom3z
 1423117_at 1.07224 pumilio 1 (Drosophila) Pum1
 1423118_at 1.06094 RIKEN cDNA 1200014J11 gene 1200014J11Rik
 1423119_at 0.89115 radial spokehead-like 2A Rshl2a
 1423120_at 1.13448 insulin degrading enzyme Ide
 1423121_at 1.02598 insulin degrading enzyme Ide
 1423122_at 1.19784 arginine vasopressin-induced 1 Avp1
 1423123_at 0.95965 RAD54 like (S. cerevisiae) Rad54l
 1423124_x_at 0.76271 RAD54 like (S. cerevisiae) Rad54l
 1423125_at 0.98201 doublecortin and calcium/calmodulin-dependent protein kinase-like 1 Dcamk1l
 1423126_at 0.94489 "ATPase, Na⁺/K⁺ transporting, beta 3 polypeptide" Atp1b3
 1423127_at 1.06988 inositol (myo)-1(or 4)-monophosphatase 1 Imp1
 1423128_at 1.06351 aryl-hydrocarbon receptor-interacting protein Aip
 1423129_at 0.99131 soc-2 (suppressor of clear) homolog (C. elegans) Shoc2
 1423130_a_at 0.98450 "splicing factor, arginine/serine-rich 5 (SRp40, HRS) /// similar to Splicing factor, arginine/serine-rich 5 (Pre-mRNA splicing factor SRP40) (Insulin-induced growth response protein CL-4) (Delayed-early protein HRS)" Sfrs5 /// LOC622469
 1423131_at 0.99695 similar to TGF beta-inducible nuclear protein 1 (L-name-related protein 42) (LNR42) LOC433230
 1423132_a_at 1.01753 RIKEN cDNA 5730427N09 gene /// similar to TGF beta-inducible nuclear protein 1 (L-name-related protein 42) (LNR42) /// similar to TGF beta-inducible nuclear protein 1 (L-name related LNR42) 5730427N09Rik /// LOC433230 /// LOC636306
 1423133_at 1.08645 RIKEN cDNA 0610040D20 gene 0610040D20Rik
 1423134_at 1.19131 cDNA sequence BC003324 BC003324
 1423135_at 0.97955 "thymus cell antigen 1, theta" Thy1
 1423136_at 0.53920 fibroblast growth factor 1 Fgf1

1423137_at	1.07358	v-ral simian leukemia viral oncogene homolog A (ras related)	Rala
1423138_at	1.11990	WD repeat domain 4	Wdr4
1423139_at	0.92309	WD repeat domain 4	Wdr4
1423140_at	0.86508	lysosomal acid lipase 1	Lip1
1423141_at	0.94130	Lysosomal acid lipase 1	Lip1
1423142_a_at	1.08637	GTP binding protein 4	Gtpbp4
1423143_at	1.03820	GTP binding protein 4	Gtpbp4
1423144_at	1.02544	"phosphatidylinositol 3-kinase, catalytic, alpha polypeptide"	Pik3ca
1423145_a_at	1.02962	titin-cap	Tcap
1423146_at	1.08270	hairy and enhancer of split 5 (Drosophila)	Hes5
1423147_at	1.01876	"methionine adenosyltransferase I, alpha"	Mat1a
1423148_at	0.96179	S-phase kinase-associated protein 1A	Skp1a
1423149_at	0.91904	S-phase kinase-associated protein 1A	Skp1a
1423150_at	1.07247	secretogranin V	Scg5
1423151_at	0.98397	"DnaJ (Hsp40) homolog, subfamily B, member 11"	Dnajb11
1423152_at	0.98513	"vesicle-associated membrane protein, associated protein B and C"	Vapb
1423153_x_at	1.01962	complement component factor h	Cfh
1423154_at	0.99398	cDNA sequence BC005537	BC005537
1423155_at	0.90963	sorcin	Sri
1423156_at	0.97667	glucosamine-phosphate N-acetyltransferase 1	Gnpnat1
1423157_at	0.84401	glucosamine-phosphate N-acetyltransferase 1	Gnpnat1
1423158_at	1.04663	glucosamine-phosphate N-acetyltransferase 1	Gnpnat1
1423159_at	0.96521	dihydrolipoamide dehydrogenase	Dld
1423160_at	1.07517	"sprouty protein with EVH-1 domain 1, related sequence"	Spred1
1423161_s_at	1.11769	"sprouty protein with EVH-1 domain 1, related sequence"	Spred1
1423162_s_at	1.12772	"sprouty protein with EVH-1 domain 1, related sequence"	Spred1
1423163_at	1.06077	HLA-B associated transcript 4	Bat4
1423164_at	0.99582	similar to transmembrane 7 superfamily member 1	LOC664862
1423165_a_at	0.97711	"metastasis-associated gene family, member 2"	Mta2
1423166_at	1.03552	CD36 antigen	Cd36
1423167_at	1.00405	preimplantation protein 3	Prei3
1423168_at	1.07702	preimplantation protein 3	Prei3
1423169_at	0.82015	"TAF7 RNA polymerase II, TATA box binding protein (TBP)-associated factor"	Taf7
1423170_at	0.78655	"TAF7 RNA polymerase II, TATA box binding protein (TBP)-associated factor"	Taf7
1423171_at	0.94438	G-protein coupled receptor 88 /// similar to G-protein coupled receptor 88	Gpr88 /// LOC671461
1423172_at	1.03221	N-ethylmaleimide sensitive fusion protein attachment protein beta	Napb
1423173_at	1.09635	N-ethylmaleimide sensitive fusion protein attachment protein beta	Napb
1423174_a_at	0.99000	par-6 (partitioning defective 6) homolog beta (C. elegans)	Pard6b
1423175_s_at	0.99671	par-6 (partitioning defective 6) homolog beta (C. elegans)	Pard6b
1423176_at	0.91201	transducer of ErbB-2.1	Tob1

1423177_a_at	0.99509	abl-interactor 1	Abi1	
1423178_at	0.87451	abl-interactor 1	Abi1	
1423179_at	0.88794	"potassium voltage gated channel, Shab-related subfamily, member 1"		Kcnb1
1423180_at	0.93103	"potassium voltage gated channel, Shab-related subfamily, member 1"		Kcnb1
1423181_s_at	1.06063	"chloride channel, nucleotide-sensitive, 1A"	Clns1a	
1423182_at	0.95521	"tumor necrosis factor receptor superfamily, member 13b"		Tnfrsf13b
1423183_at	0.99422	"leucine-rich repeat LGI family, member 1"	Lgi1	
1423184_at	1.15685	intersectin 2	Its2	
1423185_a_at	1.01463	ubiquitin-associated protein 1	Ubap1	
1423186_at	1.35928	T-cell lymphoma invasion and metastasis 2	Tiam2	
1423187_at	0.99471	gamma-aminobutyric acid (GABA-A) receptor-associated protein-like 2		Gabarapl2
1423188_a_at	0.97318	RIKEN cDNA 6720456B07 gene	6720456B07Rik	
1423189_at	0.99857	RIKEN cDNA 6720456B07 gene	6720456B07Rik	
1423190_at	0.93242	RIKEN cDNA 1810020G14 gene	1810020G14Rik	
1423191_at	1.07910	formin binding protein 4	Fnbp4	
1423192_at	1.04292	paraspeckle protein 1	Pspc1	
1423193_at	0.99621	paraspeckle protein 1	Pspc1	
1423194_at	0.86457	Rho GTPase activating protein 5	Arhgap5	
1423195_at	1.02824	hippocampus abundant gene transcript 1	Hiat1	
1423196_at	0.95743	"neural precursor cell expressed, developmentally down-regulated gene 1"		Nedd1
1423197_a_at	1.06452	expressed sequence AW011752	AW011752	
1423198_a_at	1.02420	expressed sequence AW011752	AW011752	
1423199_at	1.10316	bromodomain containing 3	Brd3	
1423200_at	0.89411	nuclear receptor co-repressor 1	Ncor1	
1423201_at	0.97123	nuclear receptor co-repressor 1	Ncor1	
1423202_a_at	0.99999	nuclear receptor co-repressor 1	Ncor1	
1423203_a_at	0.92460	centrin 1	Cetn1	
1423204_at	1.06256	transmembrane 9 superfamily protein member 4	Tm9sf4	
1423205_at	0.97779	transmembrane 9 superfamily protein member 4	Tm9sf4	
1423206_s_at	1.03314	RIKEN cDNA 2310003F16 gene	2310003F16Rik	
1423207_at	0.90864	---	---	
1423208_at	0.94448	transmembrane protein 167	Tmem167	
1423209_at	1.05164	---	---	
1423210_a_at	0.96436	"nucleolar protein family A, member 3"	Nola3	
1423211_at	1.10196	"nucleolar protein family A, member 3"	Nola3	
1423212_at	0.95636	polyhomeotic-like 1 (Drosophila)	Phc1	
1423213_at	1.04204	plexin C1	Plxnc1	
1423214_at	0.98333	plexin C1	Plxnc1	
1423215_at	0.98713	signal peptidase complex subunit 2 homolog (S. cerevisiae)		Spcs2
1423216_a_at	0.99668	RIKEN cDNA 2510049I19 gene	2510049I19Rik	
1423217_a_at	0.88633	RIKEN cDNA 2510049I19 gene	2510049I19Rik	

1423218_a_at 0.96667 mitochondrial ribosomal protein L49 Mrpl49
 1423219_a_at 0.99321 mitochondrial ribosomal protein L49 Mrpl49
 1423220_at 1.03794 eukaryotic translation initiation factor 4E /// hypothetical LOC630527 Eif4e ///
 LOC630527
 1423221_at 1.02214 "tubulin, beta 4" Tubb4
 1423222_at 0.98595 "CAP, adenylate cyclase-associated protein, 2 (yeast)" Cap2
 1423223_a_at 1.14280 peroxiredoxin 6 Prdx6
 1423224_at 0.80411 RIKEN cDNA 4432405B04 gene 4432405B04Rik
 1423225_at 1.05521 selenoprotein K Selk
 1423226_at 0.93016 "membrane-spanning 4-domains, subfamily A, member 1" Ms4a1
 1423227_at 0.95573 keratin 17 Krt17
 1423228_at 1.26409 "UDP-Gal:betaGlcNAc beta 1,4-galactosyltransferase, polypeptide 6 /// similar
 to Beta-1,4-galactosyltransferase 6 (Beta-1,4-GalTase 6) (Beta4Gal-T6) (b4Gal-T6) (UDP-galactose:
 beta-N-acetylglucosamine beta-1,4-galactosyltransferase 6) (UDP-Gal:beta-GlcNAc beta-1,4-
 galactosyltransferase 6)" B4galt6 /// LOC675709
 1423229_at 0.96386 inositol polyphosphate-5-phosphatase E Inpp5e
 1423230_at 1.03563 inositol polyphosphate-5-phosphatase E Inpp5e
 1423231_at 0.95821 neurogranin Nrgn
 1423232_at 1.20752 "ets variant gene 4 (E1A enhancer binding protein, E1AF)" Etv4
 1423233_at 3.68779 "CCAAT/enhancer binding protein (C/EBP), delta" Cebpd
 1423234_at 1.00154 "proteasome (prosome, macropain) 26S subunit, non-ATPase, 5" Psmd5
 1423235_at 0.96676 "adaptor-related protein complex 3, beta 1 subunit" Ap3b1
 1423236_at 0.85649 UDP-N-acetyl-alpha-D-galactosamine:polypeptide N-
 acetylgalactosaminyltransferase 1 Galnt1
 1423237_at 0.94695 UDP-N-acetyl-alpha-D-galactosamine:polypeptide N-
 acetylgalactosaminyltransferase 1 Galnt1
 1423238_at 1.25183 integrin beta 1 binding protein 2 Itgb1bp2
 1423239_at 0.98807 inosine 5'-phosphate dehydrogenase 1 Impdh1
 1423240_at 1.06383 Rous sarcoma oncogene Src
 1423241_a_at 0.97968 transcription factor Dp 1 /// similar to Transcription factor Dp-1 (E2F
 dimerization partner 1) (DRTF1-polypeptide 1) /// similar to Transcription factor Dp-1 (E2F
 dimerization partner 1) (DRTF1-polypeptide 1) /// similar to Transcription factor Dp-1 (E2F
 dimerization partner 1) (DRTF1-polypeptide 1) Tfdp1 /// LOC664889 /// LOC672007 /// LOC674691
 1423242_at 1.00793 mitochondrial ribosomal protein S36 Mrps36
 1423243_at 1.09718 "membrane protein, palmitoylated" Mpp1
 1423244_at 1.01033 RIKEN cDNA 9030012A22 gene 9030012A22Rik
 1423245_at 1.11980 "COP9 (constitutive photomorphogenic) homolog, subunit 7a (Arabidopsis
 thaliana)" Cops7a
 1423246_at 0.79335 thioredoxin domain containing 4 (endoplasmic reticulum) Txndc4
 1423247_at 0.91138 thioredoxin domain containing 4 (endoplasmic reticulum) Txndc4
 1423248_at 1.08227 natural killer tumor recognition sequence Nkr
 1423249_at 0.93841 natural killer tumor recognition sequence Nkr
 1423250_a_at 0.85011 "transforming growth factor, beta 2" Tgfb2

1423251_at 1.08718 LUC7-like 2 (*S. cerevisiae*) Luc712
 1423252_at 1.12123 "hepatoma-derived growth factor, related protein 3" Hdgfrp3
 1423253_at 1.06414 myelin protein zero Mpz
 1423254_x_at 1.06983 ribosomal protein S27-like Rps271
 1423255_at 1.17684 "ATPase, H⁺ transporting, lysosomal V1 subunit G1" Atp6v1g1
 1423256_a_at 1.03425 "ATPase, H⁺ transporting, lysosomal V1 subunit G1 /// similar to vacuolar H⁺ ATPase G1 /// similar to vacuolar H⁺ ATPase G1 /// similar to vacuolar H⁺ ATPase G1" Atp6v1g1 ///
 LOC666030 /// LOC671155 /// LOC671196
 1423257_at 0.98427 "cytochrome P450, family 4, subfamily a, polypeptide 14" Cyp4a14
 1423258_at 0.78763 synaptotagmin IX Syt9
 1423259_at 0.65132 inhibitor of DNA binding 4 Id4
 1423260_at 0.72511 inhibitor of DNA binding 4 Id4
 1423261_at 0.79840 RIKEN cDNA 1500015O10 gene 1500015O10Rik
 1423262_a_at 0.92844 "H3 histone, family 3A /// similar to H3 histone, family 3B /// similar to H3 histone, family 3B /// similar to H3 histone, family 3B /// similar to H3 histone, family 3B /// similar to H3 histone, family 3B" H3f3a /// LOC623365 /// LOC624812 ///
 LOC625328 /// LOC633468 /// LOC667250 /// LOC670535
 1423263_at 0.89735 "H3 histone, family 3A" H3f3a
 1423264_at 1.04373 block of proliferation 1 Bop1
 1423265_at 0.95806 multiple inositol polyphosphate histidine phosphatase 1 Minpp1
 1423266_at 0.88696 RIKEN cDNA 2810405K02 gene 2810405K02Rik
 1423267_s_at 0.95327 integrin alpha 5 (fibronectin receptor alpha) Itga5
 1423268_at 0.87035 integrin alpha 5 (fibronectin receptor alpha) Itga5
 1423269_a_at 0.92018 "neural precursor cell expressed, developmentally down-regulated gene 4-like" Nedd4l
 1423270_at 0.88855 "neural precursor cell expressed, developmentally down-regulated gene 4-like" Nedd4l
 1423271_at 0.92356 gap junction membrane channel protein beta 2 Gjb2
 1423272_at 0.96388 "polymerase (DNA directed), gamma" Polg
 1423273_at 1.00092 "polymerase (DNA directed), gamma" Polg
 1423274_at 1.09619 integrator complex subunit 6 Ints6
 1423275_at 1.10839 integrator complex subunit 6 Ints6
 1423276_at 0.92178 immunoglobulin-like domain containing receptor 1 Ildr1
 1423277_at 0.94510 "protein tyrosine phosphatase, receptor type, K" Ptpkr
 1423278_at 0.72619 "protein tyrosine phosphatase, receptor type, K" Ptpkr
 1423279_at 0.92906 "solute carrier family 34 (sodium phosphate), member 1" Slc34a1
 1423280_at 1.02288 stathmin-like 2 Stmn2
 1423281_at 0.90125 stathmin-like 2 Stmn2
 1423282_at 0.98652 "phosphatidylinositol transfer protein, alpha" Pitpna
 1423283_at 1.01204 "phosphatidylinositol transfer protein, alpha" Pitpna
 1423284_at 1.12183 MANS domain containing 1 Mansc1
 1423285_at 0.99630 coagulation factor C homolog (*Limulus polyphemus*) Coch
 1423286_at 0.88480 cerebellin 1 precursor protein Cbln1

1423287_at	0.94441 cerebellin 1 precursor protein	Cbln1	
1423288_s_at	0.85961 cerebellin 1 precursor protein	Cbln1	
1423289_a_at	1.07855 RIKEN cDNA 1810029B16 gene	1810029B16Rik	
1423290_at	0.96903 hypoxia up-regulated 1	Hyou1	
1423291_s_at	0.92966 hypoxia up-regulated 1	Hyou1	
1423292_a_at	0.90006 periaxin	Prx	
1423293_at	1.04899 replication protein A1	Rpa1	
1423294_at	0.95324 mesoderm specific transcript	Mest	
1423295_at	0.95040 transmembrane 9 superfamily member 2	Tm9sf2	
1423296_at	1.06839 "proteasome (prosome, macropain) 26S subunit, non-ATPase, 8"	Psm8	
1423297_at	0.94772 adducin 3 (gamma)	Add3	
1423298_at	0.85254 adducin 3 (gamma)	Add3	
1423299_at	0.98924 thioredoxin-like 1	Txn1	
1423300_at	0.98107 "zinc finger, DHHC domain containing 6"	Zdhhc6	
1423301_at	0.99358 "coatamer protein complex, subunit beta 1"	Cop1	
1423302_a_at	1.02937 PAX interacting (with transcription-activation domain) protein 1		Paxip1
1423303_at	0.95994 PAX interacting (with transcription-activation domain) protein 1		Paxip1
1423304_a_at	1.01950 ring finger 111	Rnf111	
1423305_at	0.95400 exostoses (multiple)-like 1	Ext1	
1423306_at	1.22941 RIKEN cDNA 2010002N04 gene	2010002N04Rik	
1423307_s_at	0.88219 trans-golgi network protein /// trans-golgi network protein 2	Tgoln1 /// Tgoln2	
1423308_at	0.97863 trans-golgi network protein	Tgoln1	
1423309_at	0.96256 trans-golgi network protein	Tgoln1	
1423310_at	0.93915 trophoblast glycoprotein	Tpbg	
1423311_s_at	0.89731 trophoblast glycoprotein	Tpbg	
1423312_at	1.02244 trophoblast glycoprotein	Tpbg	
1423313_at	1.05308 phosphodiesterase 7A	Pde7a	
1423314_s_at	1.20123 phosphodiesterase 7A	Pde7a	
1423315_at	0.99780 Bcl-2 binding component 3	Bbc3	
1423316_at	0.94600 transmembrane protein 39a	Tmem39a	
1423317_at	0.93309 RIKEN cDNA 3110001D03 gene	3110001D03Rik	
1423318_at	0.97608 RAD18 homolog (S. cerevisiae)	Rad18	
1423319_at	0.92610 hematopoietically expressed homeobox	Hhex	
1423320_at	1.22676 deoxyribonuclease 1-like 2	Dnase12	
1423321_at	1.03342 myeloid-associated differentiation marker	Myadm	
1423322_at	0.96850 lin-7 homolog C (C. elegans)	Lin7c	
1423323_at	1.03069 tumor-associated calcium signal transducer 2	Tacstd2	
1423324_at	1.12009 pinin	Pnn	
1423325_at	0.91482 pinin	Pnn	
1423326_at	0.94586 ectonucleoside triphosphate diphosphohydrolase 1	Entpd1	
1423327_at	0.97994 RIKEN cDNA 4930517K11 gene	4930517K11Rik	
1423328_at	0.96189 ganglioside-induced differentiation-associated-protein 1		Gdap1
1423329_at	1.01363 ganglioside-induced differentiation-associated-protein 1		Gdap1

1423330_at	0.89013	endosulfine alpha	Ensa	
1423331_a_at	0.84131	poliovirus receptor-related 3	Pvrl3	
1423332_at	0.97572	syndecan binding protein	Sdcbp	
1423333_at	0.99280	RIKEN cDNA 1200007D18 gene	1200007D18Rik	
1423334_at	1.02514	RIKEN cDNA 1200007D18 gene	1200007D18Rik	
1423335_at	1.04085	RIKEN cDNA 1110004F10 gene	1110004F10Rik	
1423336_at	1.02306	"origin recognition complex, subunit 4-like (<i>S. cerevisiae</i>)"	Orc4l	
1423337_at	1.02774	"origin recognition complex, subunit 4-like (<i>S. cerevisiae</i>)"	Orc4l	
1423338_at	0.97547	coiled-coil domain containing 16	Ccdc16	
1423339_s_at	1.00596	coiled-coil domain containing 16	Ccdc16	
1423340_at	0.97205	transcription factor AP-2 beta	Tcfap2b	
1423341_at	0.74397	chondroitin sulfate proteoglycan 4	Cspg4	
1423342_at	1.07724	BarH-like homeobox 1	Barx1	
1423343_at	1.03952	"solute carrier organic anion transporter family, member 1c1"	Slco1c1	
1423344_at	1.03886	erythropoietin receptor	Epor	
1423345_at	0.98566	degenerative spermatocyte homolog 1 (<i>Drosophila</i>)	Degs1	
1423346_at	0.91524	degenerative spermatocyte homolog 1 (<i>Drosophila</i>)	Degs1	
1423347_at	0.81268	SEC23A (<i>S. cerevisiae</i>)	Sec23a	
1423348_at	0.70832	frizzled homolog 8 (<i>Drosophila</i>)	Fzd8	
1423349_at	1.01336	suppressor of cytokine signaling 5	Socs5	
1423350_at	1.02929	suppressor of cytokine signaling 5	Socs5	
1423351_at	0.87892	mitochondrial ribosomal protein L1	Mrpl1	
1423352_at	0.95904	cysteine-rich secretory protein LCCL domain containing 1	Crispld1	
1423353_at	1.02865	cysteine-rich secretory protein LCCL domain containing 1	Crispld1	
1423354_at	0.93505	synaptosomal-associated protein Snap29		
1423355_at	1.24592	synaptosomal-associated protein Snap29		
1423356_at	0.82396	synaptosomal-associated protein Snap29		
1423357_at	0.86033	RIKEN cDNA 2610209A20 gene	2610209A20Rik	
1423358_at	0.92073	RIKEN cDNA 1810009K13 gene	1810009K13Rik	
1423359_at	1.03688	phospholamban	Pln	
1423360_at	1.05289	YME1-like 1 (<i>S. cerevisiae</i>)	Yme1l1	
1423361_at	1.08017	YME1-like 1 (<i>S. cerevisiae</i>)	Yme1l1	
1423362_at	0.79553	sortilin 1	Sort1	
1423363_at	0.81731	sortilin 1	Sort1	
1423364_a_at	0.90985	fused toes	Fts	
1423365_at	0.73502	"calcium channel, voltage-dependent, T type, alpha 1G subunit"	Cacna1g	
1423366_at	0.91354	stearoyl-coenzyme A desaturase 3	Scd3	
1423367_at	1.17881	wingless-related MMTV integration site 7A	Wnt7a	
1423368_at	0.94776	lysosomal-associated protein transmembrane 4A	Laptm4a	
1423369_at	0.96004	fragile X mental retardation syndrome 1 homolog	Fmr1	
1423370_a_at	0.97592	"casein kinase 1, gamma 2"	Csnk1g2	
1423371_at	0.91974	"polymerase (DNA-directed), epsilon 4 (p12 subunit)"	Pole4	
1423372_at	1.02751	"polymerase (DNA-directed), epsilon 4 (p12 subunit)"	Pole4	

1423373_at 1.09196 ribonuclease P/MRP 30 subunit (human) Rpp30
1423374_at 1.06333 nuclear receptor coactivator 6 Ncoa6
1423375_at 1.18996 RIKEN cDNA 1700023B02 gene 1700023B02Rik
1423376_a_at 0.88264 docking protein 4 Dok4
1423377_at 1.10557 insulin-like growth factor binding protein-like 1 Igfbpl1
1423378_at 0.91446 a disintegrin and metallopeptidase domain 23 Adam23
1423379_at 0.89269 "nuclear factor of activated T-cells, cytoplasmic, calcineurin-dependent 4 /// similar to cytoplasmic nuclear factor of activated T-cells 4" Nfatc4 /// LOC672021
1423380_s_at 0.76808 "nuclear factor of activated T-cells, cytoplasmic, calcineurin-dependent 4" Nfatc4
1423381_at 0.82248 RIKEN cDNA 4921521F21 gene 4921521F21Rik
1423382_a_at 1.00025 heterogeneous nuclear ribonucleoprotein F /// similar to heterogeneous nuclear ribonucleoprotein F Hnrpf /// LOC637008 /// LOC672616
1423383_a_at 0.96895 oxysterol binding protein-like 9 Osbp19
1423384_s_at 0.96445 testis expressed gene 261 Tex261
1423385_at 0.90365 ARP8 actin-related protein 8 homolog (S. cerevisiae) Actr8
1423386_at 1.08987 "proteasome (prosome, macropain) 26S subunit, non-ATPase, 9" Psm9
1423387_at 0.81435 "proteasome (prosome, macropain) 26S subunit, non-ATPase, 9" Psm9
1423388_at 1.03979 "adaptor protein complex AP-1, gamma 1 subunit" Ap1g1
1423389_at 0.76961 MAD homolog 7 (Drosophila) Smad7
1423390_at 0.95199 seven in absentia 1A Siah1a
1423391_at 1.01384 G protein-coupled receptor kinase-interactor 2 Git2
1423392_at 0.98039 chloride intracellular channel 4 (mitochondrial) Clic4
1423393_at 0.97095 chloride intracellular channel 4 (mitochondrial) Clic4
1423394_at 0.91651 prenylcysteine oxidase 1 Pcyox1
1423395_at 1.00847 translin-associated factor X Tsnax
1423396_at 0.73761 "angiotensinogen (serpin peptidase inhibitor, clade A, member 8)" Agt
1423397_at 0.92987 "UDP glucuronosyltransferase 2 family, polypeptide B38" Ugt2b38
1423398_at 1.04324 "TAF12 RNA polymerase II, TATA box binding protein (TBP)-associated factor" Taf12
1423399_a_at 0.93946 YY1 associated factor 2 Yaf2
1423400_at 1.00497 klotho Kl
1423401_at 0.97792 ets variant gene 6 (TEL oncogene) Etv6
1423402_at 1.11107 cAMP responsive element binding protein 1 Creb1
1423403_at 0.81480 mitogen activated protein kinase binding protein 1 Mapkbp1
1423404_at 1.03830 gastroke 1 Gkn1
1423405_at 1.00544 tissue inhibitor of metalloproteinase 4 Timp4
1423406_at 0.81627 synaptic vesicle glycoprotein 2 a Sv2a
1423407_a_at 0.76751 fibulin 2 Fbln2
1423408_a_at 0.97210 RIKEN cDNA 2500003M10 gene 2500003M10Rik
1423409_a_at 0.93617 RIKEN cDNA 2500003M10 gene 2500003M10Rik
1423410_at 0.95944 meiosis expressed gene 1 Meig1

1423411_at 0.91199 cDNA sequence BC013481 BC013481
1423412_at 0.95537 cDNA sequence BC013481 BC013481
1423413_at 1.04805 N-myc downstream regulated gene 1 Ndrp1
1423414_at 0.62171 prostaglandin-endoperoxide synthase 1 Ptgs1
1423415_at 0.89160 G protein-coupled receptor 83 Gpr83
1423416_at 0.97051 "SWI/SNF related, matrix associated, actin dependent regulator of chromatin, subfamily c, member 1" Smarcc1
1423417_at 1.01290 "SWI/SNF related, matrix associated, actin dependent regulator of chromatin, subfamily c, member 1" Smarcc1
1423418_at 1.59456 farnesyl diphosphate synthetase Fdps
1423419_at 1.11087 "ligase III, DNA, ATP-dependent" Lig3
1423420_at 1.10370 "adrenergic receptor, beta 1" Adrb1
1423421_at 0.97543 ankyrin repeat domain 49 Ankrd49
1423422_at 1.03095 ankyrin repeat and SOCS box-containing protein 4 Asb4
1423423_at 0.97977 protein disulfide isomerase associated 3 Pdia3
1423424_at 0.93080 zinc finger protein of the cerebellum 3 Zic3
1423425_at 0.91149 RIKEN cDNA 1300012G16 gene 1300012G16Rik
1423426_at 0.94024 RIKEN cDNA 1300012G16 gene 1300012G16Rik
1423427_at 0.96945 adenylate cyclase activating polypeptide 1 Adcyap1
1423428_at 0.91268 receptor tyrosine kinase-like orphan receptor 2 Ror2
1423429_at 0.94201 reproductive homeobox 5 Rhox5
1423430_at 1.24814 MYB binding protein (P160) 1a Mybbp1a
1423431_a_at 1.24075 MYB binding protein (P160) 1a Mybbp1a
1423432_at 1.12047 pleckstrin homology domain interacting protein Phip
1423433_at 1.07914 "TROVE domain family, member 2" Trove2
1423434_at 1.02186 TEA domain family member 1 Tead1
1423435_at 0.96484 --- ---
1423436_at 1.09022 "glutathione S-transferase, alpha 3" Gsta3
1423437_at 1.01114 --- ---
1423438_at 1.04461 kaptin Kptn
1423439_at 0.99430 "phosphoenolpyruvate carboxykinase 1, cytosolic" Pck1
1423440_at 0.99059 RIKEN cDNA 1110001A07 gene 1110001A07Rik
1423441_at 1.04075 "transcription factor B2, mitochondrial" Tfb2m
1423442_a_at 0.98987 F-box and WD-40 domain protein 2 Fbxw2
1423443_at 0.79563 "solute carrier family 4 (anion exchanger), member 8" Slc4a8
1423444_at 1.06536 Rho-associated coiled-coil containing protein kinase 1 Rock1
1423445_at 0.97341 Rho-associated coiled-coil containing protein kinase 1 Rock1
1423446_at 0.97899 death-associated kinase 3 Dapk3
1423447_at 1.05493 caseinolytic peptidase X (E.coli) Clpx
1423448_at 0.94431 "RAB11B, member RAS oncogene family" Rab11b
1423449_a_at 0.98257 actinin alpha 4 Actn4
1423450_a_at 0.89962 heparan sulfate (glucosamine) 3-O-sulfotransferase 1 Hs3st1
1423451_at 0.87020 progesterone receptor membrane component 1 Pgrmc1

1423452_at 0.93657 serine/threonine kinase 17b (apoptosis-inducing) Stk17b
 1423453_at 1.08403 nucleolar protein 12 Nol12
 1423454_a_at 0.81368 "sema domain, transmembrane domain (TM), and cytoplasmic domain, (semaphorin) 6C" Sema6c
 1423455_at 1.01447 prothymosin alpha Ptma
 1423456_at 1.09269 basic leucine zipper and W2 domains 2 Bzw2
 1423457_at 0.86625 "solute carrier family 35, member A5" Slc35a5
 1423458_at 0.91073 "solute carrier family 35, member A5" Slc35a5
 1423459_at 1.02785 "COP9 (constitutive photomorphogenic) homolog, subunit 2 (Arabidopsis thaliana)" Cops2
 1423460_at 1.30355 "PERQ amino acid rich, with GYF domain 1" Perq1
 1423461_a_at 0.97034 ubiquitin-like 3 Ubl3
 1423462_at 0.97475 mitogen-activated protein kinase kinase kinase 7 interacting protein 2 Map3k7ip2
 1423463_a_at 1.07181 "DNA segment, Chr 2, ERATO Doi 750, expressed" D2Ertd750e
 1423464_at 1.03848 "DNA segment, Chr 2, ERATO Doi 750, expressed" D2Ertd750e
 1423465_at 1.10010 ferric-chelate reductase 1 /// similar to stromal cell derived factor receptor 2 Frrs1 /// LOC635993
 1423466_at 1.00542 chemokine (C-C motif) receptor 7 Ccr7
 1423467_at 0.95151 "membrane-spanning 4-domains, subfamily A, member 4B" Ms4a4b
 1423468_at 0.92362 STEAP family member 3 Steap3
 1423469_at 0.99787 RIKEN cDNA 1700129C05 gene 1700129C05Rik
 1423470_at 0.92069 polypyrimidine tract binding protein 2 Ptbp2
 1423471_at 0.71956 polypyrimidine tract binding protein 2 Ptbp2
 1423472_at 1.07020 septin 2 2-Sep
 1423473_at 1.09205 septin 2 2-Sep
 1423474_at 0.96819 topoisomerase (DNA) I Top1
 1423475_at 0.82408 cyclin M2 Cnm2
 1423476_at 0.86371 thymic stromal cotransporter Tscot
 1423477_at 1.01835 zinc finger protein of the cerebellum 1 Zic1
 1423478_at 0.85651 "protein kinase C, beta 1" Prkcb1
 1423479_at 1.00827 nucleolar protein 11 Nol11
 1423480_at 1.04674 nucleolar protein 11 Nol11
 1423481_at 1.04994 RIO kinase 2 (yeast) Riok2
 1423482_at 0.90965 uroporphyrinogen III synthase Uros
 1423483_s_at 1.02555 "TATA box binding protein (Tbp)-associated factor, RNA polymerase I, C" Taf1c
 1423484_at 0.84879 bicaudal C homolog 1 (Drosophila) Biccl
 1423485_at 0.86618 Rad54 like 2 (S. cerevisiae) Rad54l2
 1423486_at 1.04666 cysteine-rich PDZ-binding protein Cript
 1423487_at 0.93449 cysteine-rich PDZ-binding protein Cript
 1423488_at 0.82204 monocyte to macrophage differentiation-associated Mmd
 1423489_at 0.82427 monocyte to macrophage differentiation-associated Mmd

1423490_at	0.85723	F-box only protein 3	Fbxo3	
1423491_at	0.87771	F-box only protein 3	Fbxo3	
1423492_at	1.14690	mitochondrial ribosomal protein L45	Mrpl45	
1423493_a_at	0.82930	nuclear factor I/X	Nfix	
1423494_at	0.91289	RIKEN cDNA 2310042E22 gene	2310042E22Rik	
1423495_at	0.92146	"2-4-dienoyl-Coenzyme A reductase 2, peroxisomal"	Decr2	
1423496_a_at	0.96333	putative neuronal cell adhesion molecule	Punc	
1423497_at	1.00288	kelch-like 10 (Drosophila)	Klhl10	
1423498_at	1.12000	"aldolase 1, A isoform, pseudogene 1"	Aldoa-ps1	
1423499_at	0.99190	"synuclein, alpha interacting protein (synphilin)"	Sncaip	
1423500_a_at	0.91776	SRY-box containing gene 5	Sox5	
1423501_at	1.01606	Max protein	Max	
1423502_at	0.92130	bromodomain containing 2	Brd2	
1423503_at	0.85295	junction adhesion molecule 3	Jam3	
1423504_at	0.90199	junction adhesion molecule 3	Jam3	
1423505_at	0.64889	transgelin	Tagln	
1423506_a_at	1.04429	neuronatin	Nnat	
1423507_a_at	0.93978	"sirtuin 2 (silent mating type information regulation 2, homolog) 2 (S. cerevisiae)"	Sirt2	
1423508_at	0.95603	MYST histone acetyltransferase monocytic leukemia 4	Myst4	
1423509_a_at	0.98908	islet amyloid polypeptide	Iapp	
1423510_at	1.01596	islet amyloid polypeptide	Iapp	
1423511_at	1.01076	ASF1 anti-silencing function 1 homolog A (S. cerevisiae)	Asf1a	
1423512_at	0.90663	expressed sequence AW209491	AW209491	
1423513_at	1.00100	neogenin /// RIKEN cDNA E030041M21 gene Neo1 /// E030041M21Rik		
1423514_at	0.88456	glycerol kinase-like 1	Gykl1	
1423515_at	0.95272	"sodium channel, voltage-gated, type VIII, alpha"	Scn8a	
1423516_a_at	1.08822	nidogen 2	Nid2	
1423517_at	1.00771	chaperonin subunit 6a (zeta)	Cct6a	
1423518_at	0.96846	c-src tyrosine kinase	Csk	
1423519_at	0.89620	RIKEN cDNA 2210412D01 gene	2210412D01Rik	
1423520_at	1.09728	lamin B1	Lmnb1	
1423521_at	1.00263	lamin B1	Lmnb1	
1423522_at	1.18694	"nucleoplasmin 3 /// nucleoplasmin 3, pseudogene 1 /// similar to Nucleoplasmin-3"	Npm3 /// Npm3-ps1 /// LOC639633	
1423523_at	1.01880	aminoadipate-semialdehyde synthase	Aass	
1423524_at	1.01985	microtubule associated serine/threonine kinase-like	Mastl	
1423525_at	1.00800	microtubule associated serine/threonine kinase-like	Mastl	
1423526_at	1.00570	AT rich interactive domain 3B (Bright like)	Arid3b	
1423527_at	1.02060	RIKEN cDNA 4921510H08 gene	4921510H08Rik	
1423528_at	0.79961	breast carcinoma amplified sequence 3	Bcas3	
1423529_at	0.96934	"glucose-6-phosphatase, catalytic, 2"	G6pc2	
1423530_at	0.85105	serine/threonine kinase 32C	Stk32c	

1423531_a_at 1.06525 heterogeneous nuclear ribonucleoprotein A1 Hnrpa1
1423532_at 0.98460 ring finger protein 44 Rnf44
1423533_a_at 1.01127 "ras homolog gene family, member T1" Rhot1
1423534_at 1.09962 programmed cell death 2 Pcd2
1423535_at 0.96540 "striatin, calmodulin binding protein 3" Strn3
1423536_at 1.03114 "striatin, calmodulin binding protein 3" Strn3
1423537_at 1.10453 growth associated protein 43 Gap43
1423538_at 0.96516 N-terminal Asn amidase Ntan1
1423539_at 1.03542 postmeiotic segregation increased 2 (S. cerevisiae) Pms2
1423540_at 0.98362 "RNA binding motif, single stranded interacting protein 2" Rbms2
1423541_at 1.03215 RIKEN cDNA 4930511I11 gene 4930511I11Rik
1423542_at 1.03022 "kallikrein 7 (chymotryptic, stratum corneum)" Klk7
1423543_at 0.98462 SWA-70 protein Swap70
1423544_at 1.03814 "protein tyrosine phosphatase, non-receptor type 5" Ptpn5
1423545_a_at 1.09111 zinc finger protein 207 Zfp207
1423546_at 1.09895 zinc finger protein 207 Zfp207
1423547_at 1.21895 lysozyme Lyzs
1423548_s_at 0.94257 ERGIC and golgi 3 Ergic3
1423549_at 1.03053 "solute carrier family 1 (glutamate/neutral amino acid transporter), member 4"
Slc1a4
1423550_at 1.01499 "solute carrier family 1 (glutamate/neutral amino acid transporter), member 4"
Slc1a4
1423551_at 1.23782 cadherin 13 Cdh13
1423552_at 1.00670 leptin receptor overlapping transcript-like 1 Leprotl1
1423553_at 1.04287 "DnaJ (Hsp40) homolog, subfamily B, member 3" Dnajb3
1423554_at 0.86040 gamma-glutamyl carboxylase Ggcx
1423555_a_at 1.02445 interferon-induced protein 44 Ifi44
1423556_at 0.78163 "aldo-keto reductase family 1, member B7" Akr1b7
1423557_at 1.27628 interferon gamma receptor 2 Ifngr2
1423558_at 1.22295 interferon gamma receptor 2 Ifngr2
1423559_at 1.02194 "potassium voltage gated channel, Shaw-related subfamily, member 1" Kcnc1
1423560_at 1.16816 NEL-like 2 (chicken) Nell2
1423561_at 0.88926 NEL-like 2 (chicken) Nell2
1423562_at 1.09370 proline-rich transmembrane protein 1 Prrt1
1423563_at 0.87434 proline-rich transmembrane protein 1 Prrt1
1423564_a_at 0.93400 "phosphoribosylaminoimidazole carboxylase,
phosphoribosylaminoribosylaminoimidazole, succinocarboxamide synthetase" Paics
1423565_at 0.95343 "phosphoribosylaminoimidazole carboxylase,
phosphoribosylaminoribosylaminoimidazole, succinocarboxamide synthetase" Paics
1423566_a_at 0.98504 heat shock protein 110 Hsp110
1423567_a_at 1.12658 "proteasome (prosome, macropain) subunit, alpha type 7" Psma7
1423568_at 1.16844 "proteasome (prosome, macropain) subunit, alpha type 7" Psma7
1423569_at 0.94458 glycine amidinotransferase (L-arginine:glycine amidinotransferase) Gatm

1423570_at 1.01809 "ATP-binding cassette, sub-family G (WHITE), member 1" Abcg1
1423571_at 0.93004 endothelial differentiation sphingolipid G-protein-coupled receptor 1 Edg1
1423572_at 1.14467 Bcl2-like 2 Bcl2l2
1423573_at 1.06386 steroid 5 alpha-reductase 2-like Srd5a2l
1423574_s_at 1.06040 steroid 5 alpha-reductase 2-like Srd5a2l
1423575_a_at 0.96094 Williams Beuren syndrome chromosome region 22 Wbscr22
1423576_a_at 0.93231 four and a half LIM domains 4 Fhl4
1423577_at 1.09841 ankyrin repeat domain 32 Ankrd32
1423578_at 1.01219 "procollagen, type XI, alpha 2" Col11a2
1423579_a_at 1.05767 "transient receptor potential cation channel, subfamily C, member 4" Trpc4
1423580_at 0.98320 ankyrin repeat domain 48 /// similar to spermatogenic cell-specific gene 2
Ankrd48 /// LOC633134
1423581_at 1.13142 N-myristoyltransferase 2 Nmt2
1423582_at 1.05548 doublesex and mab-3 related transcription factor 1 Dmrt1
1423583_at 0.98703 feminization 1 homolog a (C. elegans) Fem1a
1423584_at 0.86651 insulin-like growth factor binding protein 7 Igfbp7
1423585_at 0.79656 insulin-like growth factor binding protein 7 /// similar to O-acyltransferase
(membrane bound) domain containing 2 Igfbp7 /// LOC621527
1423586_at 1.11796 AXL receptor tyrosine kinase Axl
1423587_a_at 1.00555 exosome component 10 Exosc10
1423588_at 1.00504 "actin related protein 2/3 complex, subunit 4" Arpc4
1423589_at 0.96059 "actin related protein 2/3 complex, subunit 4" Arpc4
1423590_at 0.92117 napsin A aspartic peptidase Napsa
1423591_at 0.95973 FGFR1 oncogene partner 2 Fgfr1op2
1423592_at 1.19517 Rho-associated coiled-coil containing protein kinase 2 Rock2
1423593_a_at 0.96260 colony stimulating factor 1 receptor Csf1r
1423594_a_at 1.03561 endothelin receptor type B Ednrb
1423595_at 0.92338 myc induced nuclear antigen Mina
1423596_at 1.18543 NIMA (never in mitosis gene a)-related expressed kinase 6 /// similar to NIMA
(never in mitosis gene a)-related expressed kinase 6 Nek6 /// LOC674247
1423597_at 0.93435 "ATPase, aminophospholipid transporter (APLT), class I, type 8A, member
1" Atp8a1
1423598_at 0.83163 "ATPase, aminophospholipid transporter (APLT), class I, type 8A, member
1" Atp8a1
1423599_a_at 0.87538 phosducin-like Pdcl
1423600_a_at 1.14108 "Treacher Collins Franceschetti syndrome 1, homolog" Tcof1
1423601_s_at 1.11197 "Treacher Collins Franceschetti syndrome 1, homolog" Tcof1
1423602_at 0.90906 Tnf receptor-associated factor 1 Traf1
1423603_at 1.19203 "zinc finger protein, multitype 1" Zfp1
1423604_at 0.89330 chondroitin sulfate proteoglycan 3 Cspg3
1423605_a_at 1.06022 transformed mouse 3T3 cell double minute 2 Mdm2
1423606_at 0.86248 "periostin, osteoblast specific factor" Postn
1423607_at 1.04414 lumican Lum

1423608_at 1.00104 integral membrane protein 2A Itm2a
 1423609_a_at 0.88888 mannoside acetylglucosaminyltransferase 1 Mgat1
 1423610_at 1.09171 methionine aminopeptidase 2 Metap2
 1423611_at 0.92356 "alkaline phosphatase 2, liver" Akp2
 1423612_at 0.95250 expressed sequence AI462438 AI462438
 1423613_at 0.98528 --- ---
 1423614_at 0.85490 "leucine rich repeat containing 8 family, member C" Lrrc8c
 1423615_at 0.98208 zinc finger protein 364 Zfp364
 1423616_at 1.01061 TAR (HIV) RNA binding protein 2 Tarbp2
 1423617_at 1.00513 component of oligomeric golgi complex 8 Cog8
 1423618_at 0.99033 bridging integrator 1 Bin1
 1423619_at 0.89622 "RAS, dexamethasone-induced 1" Rasd1
 1423620_at 1.01949 centromere protein Q Cenpq
 1423621_a_at 0.93825 "solute carrier family 33 (acetyl-CoA transporter), member 1" Slc33a1
 1423622_a_at 0.91924 cyclin L1 Ccn1
 1423623_at 0.95654 RIKEN cDNA 2810021B07 gene 2810021B07Rik
 1423624_at 0.94201 "Fanconi anemia, complementation group L" Fanc1
 1423625_a_at 1.07250 "DnaJ (Hsp40) homolog, subfamily C, member 19" Dnajc19
 1423626_at 0.95631 dystonin Dst
 1423627_at 1.52902 "NAD(P)H dehydrogenase, quinone 1" Nqo1
 1423628_s_at 0.94454 "protocadherin gamma subfamily B, 1 /// protocadherin gamma subfamily B, 2 /// protocadherin gamma subfamily B, 4 /// protocadherin gamma subfamily B, 5 /// protocadherin gamma subfamily B, 6 /// protocadherin gamma subfamily B, 7 /// protocadherin gamma subfamily B, 8 /// protocadherin gamma subfamily C, 3 /// protocadherin gamma subfamily C, 4 /// protocadherin gamma subfamily C, 5 /// protocadherin gamma subfamily A, 1 /// protocadherin gamma subfamily A, 2 /// protocadherin gamma subfamily A, 3 /// protocadherin gamma subfamily A, 4 /// protocadherin gamma subfamily A, 5 /// protocadherin gamma subfamily A, 6 /// protocadherin gamma subfamily A, 7 /// protocadherin gamma subfamily A, 8 /// protocadherin gamma subfamily A, 9 /// protocadherin gamma subfamily A, 10 /// protocadherin gamma subfamily A, 11 /// protocadherin gamma subfamily A, 12" Pcdhgb1 /// Pcdhgb2 /// Pcdhgb4 /// Pcdhgb5 /// Pcdhgb6 /// Pcdhgb7 /// Pcdhgb8 /// Pcdhgc3 /// Pcdhgc4 /// Pcdhgc5 /// Pcdhga1 /// Pcdhga2 /// Pcdhga3 /// Pcdhga4 /// Pcdhga5 /// Pcdhga6 /// Pcdhga7 /// Pcdhga8 /// Pcdhga9 /// Pcdhga10 /// Pcdhga11 /// Pcdhga12
 1423629_at 1.05432 dynamin 2 Dnm2
 1423630_at 0.94886 cytoglobin Cygb
 1423631_at 0.98835 "nuclear receptor subfamily 2, group E, member 3" Nr2e3
 1423632_at 1.10816 G protein-coupled receptor 146 Gpr146
 1423633_at 1.09086 RIKEN cDNA 6530403A03 gene 6530403A03Rik
 1423634_at 1.02526 gasdermin 1 Gsdm1
 1423635_at 0.81211 bone morphogenetic protein 2 Bmp2
 1423636_at 0.90661 WD repeat domain 31 Wdr31
 1423637_at 1.10645 UDP-N-acetyl-alpha-D-galactosamine:polypeptide N-acetylglucosaminyltransferase 4 Galnt4
 1423638_at 0.94409 UDP-N-acetyl-alpha-D-galactosamine:polypeptide N-

acetylgalactosaminyltransferase 4	Galnt4	
1423639_at	1.00947 histamine receptor H 2	Hrh2
1423640_at	1.01031 synaptoporin	Synpr
1423641_s_at	0.97490 "CCR4-NOT transcription complex, subunit 7"	Cnot7
1423642_at	1.04776 "tubulin, beta 2c"	Tubb2c
1423643_at	1.13810 DEAD (Asp-Glu-Ala-Asp) box polypeptide 39	Ddx39
1423644_at	1.09000 aconitase 1	Aco1
1423645_a_at	0.88063 DEAD (Asp-Glu-Ala-Asp) box polypeptide 5	Ddx5
1423646_at	0.97541 "zinc finger, DHHC domain containing 3"	Zdhhc3
1423647_a_at	1.04676 "zinc finger, DHHC domain containing 3"	Zdhhc3
1423648_at	0.98069 protein disulfide isomerase associated 6	Pdia6
1423649_at	1.00460 transmembrane protein 68	Tmem68
1423650_at	0.92430 ring finger protein 26	Rnf26
1423651_at	0.91234 HESB like domain containing 2	Hbld2
1423652_at	0.88575 HESB like domain containing 2	Hbld2
1423653_at	0.86120 "ATPase, Na+/K+ transporting, alpha 1 polypeptide"	Atp1a1
1423654_a_at	1.03496 ring finger protein 4	Rnf4
1423655_a_at	0.90616 RIKEN cDNA 1500010J02 gene	1500010J02Rik
1423656_x_at	0.94808 RIKEN cDNA 1500010J02 gene	1500010J02Rik
1423657_at	0.96230 CDP-diacylglycerol--inositol 3-phosphatidyltransferase (phosphatidylinositol synthase)	Cdpt
1423658_at	0.93210 signal peptide peptidase 3	Spp13
1423659_a_at	1.16560 "TBC1 domain family, member 17"	Tbc1d17
1423660_at	1.08851 "CTD (carboxy-terminal domain, RNA polymerase II, polypeptide A) small phosphatase 2"	Ctdsp2
1423661_s_at	1.06043 "CTD (carboxy-terminal domain, RNA polymerase II, polypeptide A) small phosphatase 2"	Ctdsp2
1423662_at	1.00036 "ATPase, H+ transporting, lysosomal accessory protein 2"	Atp6ap2
1423663_at	0.98811 folliculin	Flcn
1423664_at	1.05219 quinoid dihydropteridine reductase	Qdpr
1423665_a_at	1.01183 ribosomal protein L5	Rpl5
1423666_s_at	1.06578 ribosomal protein L5 /// similar to 60S ribosomal protein L5 /// similar to 60S ribosomal protein L5 /// similar to 60S ribosomal protein L5 /// similar to 60S ribosomal protein L5	Rpl5 /// LOC382740 /// LOC665407 /// LOC668032 /// LOC672748 /// LOC675821
1423667_at	0.95989 "methionine adenosyltransferase II, alpha"	Mat2a
1423668_at	0.92642 "zinc finger, DHHC domain containing 14"	Zdhhc14
1423669_at	0.72693 "procollagen, type I, alpha 1"	Coll1a1
1423670_a_at	1.01830 signal recognition particle receptor ('docking protein')	Srpr
1423671_at	0.94138 delta/notch-like EGF-related receptor	Dner
1423672_at	0.73796 RIKEN cDNA 2510042P03 gene	2510042P03Rik
1423673_at	0.95122 "leucine zipper, down-regulated in cancer 1-like"	Ldoc11
1423674_at	1.03018 ubiquitin specific peptidase 1	Usp1

1423675_at	0.93482	ubiquitin specific peptdiase 1	Usp1
1423676_at	1.00793	"ATP synthase, H+ transporting, mitochondrial F0 complex, subunit d"	Atp5h
1423677_at	0.86863	FK506 binding protein 9	Fkbp9
1423678_at	0.95421	cDNA sequence	BC017643 BC017643
1423679_at	0.85752	RIKEN cDNA 2810432L12	gene 2810432L12Rik
1423680_at	1.08579	fatty acid desaturase 1	Fads1
1423681_at	1.05838	RIKEN cDNA 1300018I05	gene 1300018I05Rik
1423682_a_at	0.96544	cell division cycle associated 4	Cdca4
1423683_at	0.96189	cell division cycle associated 4	Cdca4
1423684_at	1.00395	heterogeneous nuclear ribonucleoprotein K	Hnrpk
1423685_at	1.17502	alanyl-tRNA synthetase	Aars
1423686_a_at	1.04152	proline rich 13	Prr13
1423687_a_at	1.07003	"mannosidase, alpha, class 2C, member 1"	Man2c1
1423688_at	0.97627	"mannosidase, alpha, class 2C, member 1"	Man2c1
1423689_a_at	0.81500	"G-protein signalling modulator 1 (AGS3-like, C. elegans)"	Gpsm1
1423690_s_at	0.93562	"G-protein signalling modulator 1 (AGS3-like, C. elegans)"	Gpsm1
1423691_x_at	0.99177	keratin 8	Krt8
1423692_at	1.07262	"NADH dehydrogenase (ubiquinone) 1 alpha subcomplex, 8"	Ndufa8
1423693_at	0.84027	"elastase 1, pancreatic"	Ela1
1423694_at	0.96967	potassium channel tetramerisation domain containing 10	Kctd10
1423695_at	0.91012	"ER degradation enhancer, mannosidase alpha-like 2"	Edem2
1423696_a_at	1.00350	"proteasome (prosome, macropain) 26S subunit, non-ATPase, 6"	Psm6
1423697_at	1.06203	"proteasome (prosome, macropain) 26S subunit, non-ATPase, 6"	Psm6
1423698_at	0.85856	"DNA segment, Chr 15, ERATO Doi 785, expressed"	D15Ertd785e
1423699_at	0.89807	"DNA segment, Chr 15, ERATO Doi 785, expressed"	D15Ertd785e
1423700_at	0.98153	replication factor C (activator 1) 3	Rfc3
1423701_at	0.98128	Coenzyme A synthase	Coasy
1423702_at	0.96053	"H1 histone family, member 0"	H1f0
1423703_at	1.07679	peter pan homolog (Drosophila)	Ppan
1423704_at	0.96106	lysophospholipase 3	Lypla3
1423705_at	1.07295	RIKEN cDNA 2310057D15	gene 2310057D15Rik
1423706_a_at	1.26301	phosphogluconate dehydrogenase	Pgd
1423707_at	0.89394	transmembrane protein 50B	Tmem50b
1423708_a_at	1.05253	"phenylalanine-tRNA synthetase-like, beta subunit"	Farslb
1423709_s_at	1.05535	"phenylalanine-tRNA synthetase-like, beta subunit"	Farslb
1423710_at	1.00511	dihydrolipoamide S-succinyltransferase (E2 component of 2-oxo-glutarate complex)	Dlst
1423711_at	0.95397	"NADH dehydrogenase (ubiquinone) 1 alpha subcomplex, assembly factor 1"	Ndufaf1
1423712_a_at	1.02371	glutaminyl-tRNA synthetase	Qars
1423713_at	0.98329	"ATP-binding cassette, sub-family B (MDR/TAP), member 8"	Abcb8
1423714_at	1.00673	ASF1 anti-silencing function 1 homolog B (S. cerevisiae)	Asf1b
1423715_a_at	1.11057	"neural precursor cell expressed, developmentally down-regulated gene 8"	

Nedd8

1423716_s_at 0.96048 "ATP synthase, H⁺ transporting, mitochondrial F1 complex, delta subunit"

Atp5d

1423717_at 0.93053 adenylate kinase 3 Ak3

1423718_at 0.87184 adenylate kinase 3 Ak3

1423719_at 1.00968 cDNA sequence U46068 U46068

1423720_a_at 1.01247 SAR1 gene homolog A (*S. cerevisiae*) Sar1a

1423721_at 0.90707 "tropomyosin 1, alpha" Tpm1

1423722_at 0.96055 transmembrane protein 49 Tmem49

1423723_s_at 1.05847 TAR DNA binding protein Tardbp

1423724_at 0.90263 ZW10 interactor Zwint

1423725_at 1.02653 plastin 3 (T-isoform) Pls3

1423726_at 1.17301 vesicle amine transport protein 1 homolog (*T. californica*) Vat1

1423727_at 0.97774 cornichon homolog (*Drosophila*) Cnih

1423728_at 0.93037 "eukaryotic translation initiation factor 3, subunit 6 interacting protein"

Eif3s6ip

1423729_a_at 1.02082 RIKEN cDNA 2500002L14 gene 2500002L14Rik

1423730_at 0.98011 CLPTM1-like Clptm1l

1423731_at 1.09477 "aldehyde dehydrogenase 16 family, member A1" Aldh16a1

1423732_at 0.94670 translocating chain-associating membrane protein 1 Tram1

1423733_a_at 0.87103 Flt3 interacting zinc finger protein 1 Fiz1

1423734_at 1.02470 RAS-related C3 botulinum substrate 1 Rac1

1423735_a_at 1.09599 WD repeat domain 36 Wdr36

1423736_a_at 0.99811 dymeclin Dym

1423737_at 0.98623 NADH dehydrogenase (ubiquinone) Fe-S protein 3 Ndufs3

1423738_at 0.96396 oxidase assembly 1-like Oxa1l

1423739_x_at 0.89616 amyloid beta (A4) precursor-like protein 2 Aplp2

1423740_a_at 0.97394 RNA binding motif protein 10 Rbm10

1423741_at 0.95811 RNA binding motif protein 10 Rbm10

1423742_at 1.03939 RNA binding motif protein 10 Rbm10

1423743_at 0.94552 archain 1 Arcn1

1423744_x_at 1.08453 "eukaryotic translation initiation factor 2, subunit 3, structural gene X-linked" Eif2s3x

1423745_at 1.02341 RIKEN cDNA 1110031B06 gene 1110031B06Rik

1423746_at 0.90019 thioredoxin domain containing 5 Txndc5

1423747_a_at 1.15948 "pyruvate dehydrogenase kinase, isoenzyme 1" Pdk1

1423748_at 1.28526 "pyruvate dehydrogenase kinase, isoenzyme 1" Pdk1

1423749_s_at 1.04256 RAN GTPase activating protein 1 Rangap1

1423750_a_at 1.01875 Splicing factor 1 Sf1

1423751_at 1.10573 splicing factor 1 Sf1

1423752_at 1.02271 DEAD (Asp-Glu-Ala-Asp) box polypeptide 47 Ddx47

1423753_at 0.54296 "BMP and activin membrane-bound inhibitor, homolog (*Xenopus laevis*)"

Bambi

1423754_at 1.45086 interferon induced transmembrane protein 3 Ifitm3
1423755_at 0.95277 "zinc finger, CCHC domain containing 8" Zcchc8
1423756_s_at 0.48289 insulin-like growth factor binding protein 4 Igfbp4
1423757_x_at 0.78559 insulin-like growth factor binding protein 4 Igfbp4
1423758_at 1.02389 RIKEN cDNA E430034L04 gene E430034L04Rik
1423759_a_at 1.02016 transmembrane and coiled-coil domains 1 Tmco1
1423760_at 1.19661 CD44 antigen Cd44
1423761_at 1.03279 RIKEN cDNA 5630401D24 gene 5630401D24Rik
1423762_at 0.99354 aarF domain containing kinase 1 Adck1
1423763_x_at 0.92532 Cytotoxic granule-associated RNA binding protein 1 Tia1
1423764_s_at 1.01587 mitochondrial ribosomal protein L37 Mrpl37
1423765_at 0.86750 "ATH1, acid trehalase-like 1 (yeast)" Athl1
1423766_at 1.01565 PAK1 interacting protein 1 Pak1ip1
1423767_at 0.96998 RIKEN cDNA 2810410M20 gene 2810410M20Rik
1423768_at 1.18450 unc-93 homolog B1 (C. elegans) Unc93b1
1423769_at 0.98192 pentatricopeptide repeat domain 2 Ptdc2
1423770_at 0.86623 transmembrane channel-like gene family 6 Tmc6
1423771_at 1.05841 "protein kinase C, delta binding protein" Prkcdbp
1423772_x_at 1.05482 "solute carrier family 25 (mitochondrial carrier, adenine nucleotide translocator), member 5" Slc25a5
1423773_at 0.98170 GC-rich promoter binding protein 1 Gbbp1
1423774_a_at 0.99156 protein regulator of cytokinesis 1 Prc1
1423775_s_at 0.95222 protein regulator of cytokinesis 1 Prc1
1423776_s_at 0.96052 "TBC1 domain family, member 22a" Tbc1d22a
1423777_at 0.71232 ubiquitin specific peptidase 20 Usp20
1423778_at 0.77434 ubiquitin specific peptidase 20 Usp20
1423779_at 0.96723 coiled-coil-helix-coiled-coil-helix domain containing 6 Chchd6
1423780_at 0.87287 3-hydroxyisobutyrate dehydrogenase Hibadh
1423781_at 1.09255 amyloid beta precursor protein binding protein 1 Appbp1
1423782_at 1.07816 "MOB1, Mps One Binder kinase activator-like 1B (yeast)" Mobk1b
1423783_at 1.01776 "torsin family 2, member A" Tor2a
1423784_at 1.10671 glycyl-tRNA synthetase Gars
1423785_at 1.01216 EGL nine homolog 1 (C. elegans) Egl1
1423786_at 0.98074 RIKEN cDNA 8430410A17 gene 8430410A17Rik
1423787_at 1.01021 nucleoporin 133 Nup133
1423788_at 0.92780 nucleoporin 133 Nup133
1423789_at 0.99398 cDNA sequence BC005624 BC005624
1423790_at 0.98866 death-associated protein Dap
1423791_at 0.95702 IK cytokine Ik
1423792_a_at 0.94168 CKLF-like MARVEL transmembrane domain containing 6 Cmtm6
1423793_at 0.92540 "DNA segment, Chr 2, ERATO Doi 391, expressed" D2Ertd391e
1423794_at 1.04406 "DNA segment, Chr 2, ERATO Doi 391, expressed" D2Ertd391e
1423795_at 1.03731 splicing factor proline/glutamine rich (polypyrimidine tract binding protein

associated) Sfpq
 1423796_at 1.10460 splicing factor proline/glutamine rich (polypyrimidine tract binding protein associated) Sfpq
 1423797_at 1.19866 acetoacetyl-CoA synthetase Aacs
 1423798_a_at 0.96571 eukaryotic translation initiation factor 1 Eif1
 1423799_at 1.02935 eukaryotic translation initiation factor 1 Eif1
 1423800_at 1.00841 aspartyl-tRNA synthetase Dars
 1423801_a_at 1.14016 adenine phosphoribosyl transferase Aprt
 1423802_at 0.98156 CaM kinase-like vesicle-associated Camkv
 1423803_s_at 1.03665 glioma tumor suppressor candidate region gene 2 Gltscr2
 1423804_a_at 1.45896 isopentenyl-diphosphate delta isomerase Idi1
 1423805_at 0.72971 disabled homolog 2 (Drosophila) Dab2
 1423806_at 1.03680 integrator complex subunit 4 Ints4
 1423807_a_at 0.95855 calmodulin 2 Calm2
 1423808_at 0.94734 WD repeat domain 79 Wdr79
 1423809_at 0.85444 transcription factor 19 Tcf19
 1423810_at 1.01186 protein phosphatase methylesterase 1 Ppme1
 1423811_at 1.09388 "splicing factor 3a, subunit 3" Sf3a3
 1423812_s_at 1.04481 expressed sequence AW146242 AW146242
 1423813_at 1.02935 kinesin family member 22 Kif22
 1423814_at 0.81907 DEAD (Asp-Glu-Ala-Asp) box polypeptide 41 Ddx41
 1423815_at 1.07388 DEAD (Asp-Glu-Ala-Asp) box polypeptide 56 Ddx56
 1423816_at 0.88557 CAAX box 1 homolog B (human) Cxx1b
 1423817_s_at 0.96367 RIKEN cDNA 2010315L10 gene 2010315L10Rik
 1423818_a_at 0.93829 ADP-ribosylation factor-like 6 interacting protein 1 Arl6ip1
 1423819_s_at 0.94993 ADP-ribosylation factor-like 6 interacting protein 1 Arl6ip1
 1423820_at 1.05378 "elongation factor 1 homolog (ELF1, S. cerevisiae)" Elof1
 1423821_at 0.76986 transmembrane protein 168 Tmem168
 1423822_a_at 0.83799 transmembrane protein 168 Tmem168
 1423823_at 1.20342 RIKEN cDNA 2610012O22 gene 2610012O22Rik
 1423824_at 0.85568 G protein-coupled receptor 177 Gpr177
 1423825_at 0.86937 G protein-coupled receptor 177 Gpr177
 1423826_at 0.97225 nucleolar complex associated 4 homolog (S. cerevisiae) Noc4l
 1423827_s_at 1.04479 nucleolar complex associated 4 homolog (S. cerevisiae) Noc4l
 1423828_at 1.19076 fatty acid synthase Fasn
 1423829_at 1.00993 RIKEN cDNA 0910001A06 gene 0910001A06Rik
 1423830_a_at 1.01731 "mesoderm induction early response 1, family member 2" Mier2
 1423831_at 0.81411 "protein kinase, AMP-activated, gamma 2 non-catalytic subunit" Prkag2
 1423832_at 0.82386 "protein kinase, AMP-activated, gamma 2 non-catalytic subunit" Prkag2
 1423833_a_at 0.89343 brain protein 44 Brp44
 1423834_s_at 0.97591 "golgi associated, gamma adaptin ear containing, ARF binding protein 1" Gga1
 1423835_at 0.90645 zinc finger protein 503 Zfp503
 1423836_at 0.90619 zinc finger protein 503 Zfp503

1423837_at 0.92730 RIKEN cDNA 2400003C14 gene 2400003C14Rik
1423838_s_at 0.94847 RIKEN cDNA 2400003C14 gene 2400003C14Rik
1423839_a_at 1.02014 basic transcription factor 3 Btf3
1423840_at 1.15908 coiled-coil domain containing 56 Ccdc56
1423841_at 0.91743 brix domain containing 2 Bxdc2
1423842_a_at 0.92265 ring finger protein 41 Rnf41
1423843_at 1.06249 leucine rich repeat containing 61 Lrrc61
1423844_s_at 0.57999 cystathionine beta-synthase Cbs
1423845_at 0.85657 "cold shock domain containing C2, RNA binding" Csdc2
1423846_x_at 0.93589 "tubulin, alpha 2" Tuba2
1423847_at 0.94525 RIKEN cDNA 2810406C15 gene 2810406C15Rik
1423848_at 1.06409 M phase phosphoprotein 6 Mphosph6
1423849_a_at 1.09606 CDC-like kinase 3 Clk3
1423850_at 1.20515 NOL1/NOP2/Sun domain family 2 Nsun2
1423851_a_at 0.94881 transmembrane protein 46 Tmem46
1423852_at 1.05911 transmembrane protein 46 Tmem46
1423853_at 0.92540 RIKEN cDNA 6330527O06 gene 6330527O06Rik
1423854_a_at 0.37482 "RAS-like, family 11, member B" Rasl11b
1423855_x_at 1.06708 ribosomal protein L17 Rpl17
1423856_at 1.12354 popeye domain containing 3 Popdc3
1423857_at 1.00505 mitochondrial ribosomal protein L30 Mrpl30
1423858_a_at 1.05247 3-hydroxy-3-methylglutaryl-Coenzyme A synthase 2 Hmgcs2
1423859_a_at 1.32571 prostaglandin D2 synthase (brain) Ptgds
1423860_at 1.44704 prostaglandin D2 synthase (brain) Ptgds
1423861_at 0.90389 "pleckstrin homology domain containing, family F (with FYVE domain)
member 2" Plekhf2
1423862_at 0.86253 "pleckstrin homology domain containing, family F (with FYVE domain)
member 2" Plekhf2
1423863_at 1.06704 "ATP-binding cassette, sub-family F (GCN20), member 2" Abcf2
1423864_at 0.93697 nuclear receptor binding protein Nrbp
1423865_at 0.72228 "solute carrier family 44, member 1" Slc44a1
1423866_at 1.05086 "serine (or cysteine) peptidase inhibitor, clade A, member 3K" Serpina3k
1423867_at 1.06633 "serine (or cysteine) peptidase inhibitor, clade A, member 3K" Serpina3k
1423868_at 1.24003 thioredoxin reductase 3 Txnrd3
1423869_s_at 0.91550 thioredoxin reductase 3 Txnrd3
1423870_at 1.01830 RIKEN cDNA 2610208M17 gene 2610208M17Rik
1423871_at 0.93437 transmembrane protein 63a Tmem63a
1423872_a_at 0.93687 dystroglycan 1 Dag1
1423873_at 0.99590 "LSM1 homolog, U6 small nuclear RNA associated (S. cerevisiae)" Lsm1
1423874_at 0.96240 WD repeat domain 33 Wdr33
1423875_at 1.22968 expressed sequence AI450540 AI450540
1423876_at 1.09672 expressed sequence AI450540 AI450540
1423877_at 1.02283 "chromatin assembly factor 1, subunit B (p60)" Chaf1b

1423878_at	1.07909	glycophorin C	Gypc	
1423879_at	1.02452	RIKEN cDNA D030056L22	gene	D030056L22Rik
1423880_at	1.07912	"DNA segment, Chr 10, Wayne State University 52, expressed"		D10Wsu52e
1423881_at	1.04777	"SAPS domain family, member 3"	Saps3	
1423882_at	1.12145	ring finger and WD repeat domain 3	Rfwd3	
1423883_at	1.04141	acyl-CoA synthetase long-chain family member 1	Acs11	
1423884_at	1.12170	"cirrhosis, autosomal recessive 1A (human)"	Cirh1a	
1423885_at	0.93179	"laminin, gamma 1"	Lamc1	
1423886_at	0.84338	"laminin, gamma 1"	Lamc1	
1423887_a_at	0.99647	RIKEN cDNA 1200003M09	gene	1200003M09Rik
1423888_at	1.05474	RIKEN cDNA 1200003M09	gene	1200003M09Rik
1423889_at	0.91236	similar to hypothetical protein FLJ20010		MGC5739
1423890_x_at	0.78492	"ATPase, Na ⁺ /K ⁺ transporting, beta 1 polypeptide"		Atp1b1
1423891_at	0.86307	"glutathione S-transferase, theta 3"	Gstt3	
1423892_at	0.86539	"amyloid beta (A4) precursor protein-binding, family B, member 1"		Apbb1
1423893_x_at	0.88165	"amyloid beta (A4) precursor protein-binding, family B, member 1"		Apbb1
1423894_a_at	0.86926	DALR anticodon binding domain containing 3	Dalrd3	
1423895_a_at	0.96551	"CUG triplet repeat, RNA binding protein 2"	Cugbp2	
1423896_a_at	0.92484	ring finger protein 187	Rnf187	
1423897_at	0.93990	ring finger protein 187	Rnf187	
1423898_a_at	1.06870	thyroid hormone receptor interactor 12	Trip12	
1423899_at	0.79017	thyroid hormone receptor interactor 12	Trip12	
1423900_at	1.05509	thyroid hormone receptor interactor 12	Trip12	
1423901_at	1.05715	thyroid hormone receptor interactor 12	Trip12	
1423902_s_at	0.82585	Rho guanine nucleotide exchange factor (GEF) 12	Arhgef12	
1423903_at	1.26269	poliovirus receptor	Pvr	
1423904_a_at	1.27019	poliovirus receptor	Pvr	
1423905_at	1.22497	poliovirus receptor	Pvr	
1423906_at	1.02346	heat shock factor binding protein 1	Hsbp1	
1423907_a_at	1.01300	NADH dehydrogenase (ubiquinone) Fe-S protein 8	Ndufs8	
1423908_at	1.02623	NADH dehydrogenase (ubiquinone) Fe-S protein 8	Ndufs8	
1423909_at	1.44486	RIKEN cDNA 0610011I04	gene	0610011I04Rik
1423910_at	1.03685	"centaurin, gamma 3"	Centg3	
1423911_at	0.80497	"protein phosphatase 2, regulatory subunit B (B56), alpha isoform"		Ppp2r5a
1423912_at	0.99184	"alveolar soft part sarcoma chromosome region, candidate 1 (human)"		Aspscr1
1423913_at	0.90236	HECT domain containing 3	Hectd3	
1423914_at	0.91525	RIKEN cDNA C630004H02	gene	C630004H02Rik
1423915_at	0.83774	olfactomedin-like 2B	Olfml2b	
1423916_s_at	0.94285	myeloid leukemia factor 2	Mlf2	
1423917_a_at	1.00337	cortactin	Ctnn	
1423918_at	0.87589	rhomboid domain containing 1	Rhbdd1	
1423919_at	0.96146	cDNA sequence BC023882		BC023882

1423920_at 0.97778 barren homolog (Drosophila) Brn1
1423921_at 1.01530 integrator complex subunit 3 Ints3
1423922_s_at 0.96509 integrator complex subunit 3 Ints3
1423923_a_at 0.99764 WD repeat domain 8 Wdr8
1423924_s_at 0.90638 tetraspanin 14 Tspan14
1423925_at 1.07182 DEAH (Asp-Glu-Ala-His) box polypeptide 16 Dhx16
1423926_at 0.90237 "solute carrier family 39 (metal ion transporter), member 13" Slc39a13
1423927_at 0.89691 "solute carrier family 35, member B2" Slc35b2
1423928_at 0.96890 phosphoglycerate dehydrogenase like 1 Phgdh1
1423929_at 0.94849 phosphoglycerate dehydrogenase like 1 Phgdh1
1423930_at 1.04298 anaphase promoting complex subunit 4 Anapc4
1423931_s_at 0.97931 anaphase promoting complex subunit 4 Anapc4
1423932_at 0.98778 "CUG triplet repeat, RNA binding protein 1" Cugbp1
1423933_a_at 0.87504 RIKEN cDNA 1600029D21 gene 1600029D21Rik
1423934_at 0.92854 --- ---
1423935_x_at 1.04710 keratin 14 Krt14
1423936_at 0.99596 potassium channel tetramerisation domain containing 5 Kctd5
1423937_at 1.05435 potassium channel tetramerisation domain containing 5 Kctd5
1423938_at 0.91490 lethal giant larvae homolog 2 (Drosophila) Llg2
1423939_a_at 0.91875 Yip1 interacting factor homolog A (S. cerevisiae) Yif1a
1423940_at 0.98879 Yip1 interacting factor homolog A (S. cerevisiae) Yif1a
1423941_at 1.07881 calcium/calmodulin-dependent protein kinase II gamma Camk2g
1423942_a_at 1.12861 calcium/calmodulin-dependent protein kinase II gamma Camk2g
1423943_at 1.02731 dihydrouridine synthase 1-like (S. cerevisiae) Dus11
1423944_at 0.86526 hemopexin Hpxn
1423945_a_at 0.82762 "protein kinase inhibitor, gamma" Pkig
1423946_at 1.10962 PDZ and LIM domain 2 Pdlim2
1423947_at 1.29195 RIKEN cDNA 1110008P14 gene 1110008P14Rik
1423948_at 1.14013 Bcl2-associated athanogene 2 Bag2
1423949_at 1.07107 ubiquitin-conjugating enzyme E2Z (putative) Ube2z
1423950_at 0.96874 "potassium voltage-gated channel, shaker-related subfamily, beta member 3"
Kcnab3
1423951_at 0.96300 TM2 domain containing 3 Tm2d3
1423952_a_at 1.10469 keratin 7 Krt7
1423953_at 0.86397 CDK5 regulatory subunit associated protein 1-like 1 Cdkal1
1423954_at 5.81380 complement component 3 C3
1423955_a_at 0.96805 longevity assurance homolog 2 (S. cerevisiae) Lass2
1423956_at 0.93089 stromal membrane-associated protein 1 Smap1
1423957_at 1.19762 interferon stimulated exonuclease gene 20-like 1 Isg20l1
1423958_a_at 0.93786 RIKEN cDNA 2900001O04 gene 2900001O04Rik
1423959_at 1.26256 ropporin 1-like Ropn11
1423960_at 0.97568 membrane bound O-acyltransferase domain containing 5 Mboat5
1423961_at 1.10835 WD repeat domain 26 Wdr26

1423962_at 0.92736 WD repeat domain 26 Wdr26
1423963_at 1.01901 WD repeat domain 26 Wdr26
1423964_at 1.02099 cleavage and polyadenylation specific factor 3-like Cpsf31
1423965_at 0.83126 Cd99 antigen-like 2 Cd99l2
1423966_at 0.82280 Cd99 antigen-like 2 Cd99l2
1423967_at 0.85861 paralemmin Palm
1423968_at 0.97305 "UDP glycosyltransferases 3 family, polypeptide A2" Ugt3a2
1423969_at 1.10240 nucleoporin 37 Nup37
1423970_at 1.01421 THO complex 3 Thoc3
1423971_at 1.09181 THO complex 3 Thoc3
1423972_at 1.01102 "electron transferring flavoprotein, alpha polypeptide" EtfA
1423973_a_at 1.06481 ADP-ribosylation factor 3 Arf3
1423974_at 0.90995 nuclear mitotic apparatus protein 1 Numa1
1423975_s_at 0.96965 nuclear mitotic apparatus protein 1 Numa1
1423976_at 0.94276 RIKEN cDNA 4930453N24 gene 4930453N24Rik
1423977_at 1.01822 RIKEN cDNA 4930453N24 gene /// similar to CG13876-PA /// similar to
CG13876-PA /// similar to CG13876-PA 4930453N24Rik /// LOC667795 /// LOC670686 ///
LOC676400
1423978_at 0.67183 SH3-binding kinase 1 Sbk1
1423979_a_at 0.89380 "solute carrier family 25 (mitochondrial carrier, palmitoylcarnitine transporter),
member 29" Slc25a29
1423980_at 0.84232 "solute carrier family 25 (mitochondrial carrier, palmitoylcarnitine transporter),
member 29" Slc25a29
1423981_x_at 0.98239 "solute carrier family 25 (mitochondrial carrier, palmitoylcarnitine
transporter), member 29" Slc25a29
1423982_at 1.11498 FUS interacting protein (serine-arginine rich) 1 Fusip1
1423983_at 0.82963 seizure related 6 homolog like 2 Sez6l2
1423984_a_at 0.99298 myeloid ecotropic viral integration site-related gene 2 /// similar to Homeobox
protein Meis3 (Meis1-related protein 2) Mrg2 /// LOC670770
1423985_at 0.90372 "guanine nucleotide binding protein (G protein), gamma 5 subunit" Gng5
1423986_a_at 0.93235 scotin gene Scotin
1423987_at 0.96027 cell division cycle 26 Cdc26
1423988_at 0.95307 adenylate kinase 1 Ak1
1423989_at 0.94849 RIKEN cDNA 2210010N04 gene 2210010N04Rik
1423990_at 0.92005 "RAB28, member RAS oncogene family" Rab28
1423991_at 1.05714 RIKEN cDNA 2610033H07 gene 2610033H07Rik
1423992_at 1.06958 GATA zinc finger domain containing 2A Gatad2a
1423993_at 1.13971 "ATPase, H+ transporting, lysosomal V1 subunit F" Atp6v1f
1423994_at 0.86460 kinesin family member 1B Kif1b
1423995_at 0.95601 kinesin family member 1B Kif1b
1423996_a_at 0.90404 "interleukin 4 receptor, alpha" Il4ra
1423997_at 1.12257 "cold shock domain containing E1, RNA binding" Csde1
1423998_at 0.99175 "general transcription factor IIIC, polypeptide 5" Gtf3c5

1423999_at 0.86805 v-abl Abelson murine leukemia oncogene 1 Abl1
1424000_a_at 1.04474 ribosomal protein S11 Rps11
1424001_at 1.02782 Mki67 (FHA domain) interacting nucleolar phosphoprotein Mki67ip
1424002_at 1.02821 phosducin-like 3 Pdc13
1424003_at 0.98245 RIKEN cDNA 4930444A02 gene 4930444A02Rik
1424004_x_at 0.81270 RIKEN cDNA 4930444A02 gene 4930444A02Rik
1424005_at 0.87663 RIKEN cDNA B230219D22 gene B230219D22Rik
1424006_at 0.97374 alanyl-tRNA synthetase domain containing 1 Aarsd1
1424007_at 0.88198 growth differentiation factor 10 Gdf10
1424008_a_at 1.07520 RNA binding protein with multiple splicing 2 Rbpms2
1424009_at 1.05072 regenerating islet-derived 3 delta Reg3d
1424010_at 0.80435 microfibrillar-associated protein 4 Mfap4
1424011_at 0.90158 aquaporin 9 Aqp9
1424012_at 0.77014 RIKEN cDNA 4930506L13 gene 4930506L13Rik
1424013_at 1.10025 eukaryotic translation termination factor 1 Etf1
1424014_at 1.09657 RIKEN cDNA 2900092E17 gene 2900092E17Rik
1424015_at 0.93522 Rab6 interacting protein 1 Rab6ip1
1424016_at 0.84497 RIKEN cDNA 2310007F21 gene 2310007F21Rik
1424017_a_at 0.98165 histidine triad nucleotide binding protein 1 Hint1
1424018_at 1.00802 histidine triad nucleotide binding protein 1 Hint1
1424019_at 1.08521 nucleolar protein 1 Noll
1424020_at 0.95356 ADP-ribosylation factor-like 6 interacting protein 6 /// similar to ADP-
ribosylation-like factor 6-interacting protein 6 Arl6ip6 /// LOC674449
1424021_at 0.90721 similar to ADP-ribosylation-like factor 6-interacting protein 6 LOC674449
1424022_at 1.02129 RIKEN cDNA 1700012B18 gene 1700012B18Rik
1424023_at 0.95275 ATP binding domain 3 Atpbd3
1424024_at 0.93794 multiple coagulation factor deficiency 2 Mcfd2
1424025_at 1.01487 cDNA sequence BC013529 BC013529
1424026_s_at 0.91084 cDNA sequence BC013529 BC013529
1424027_at 0.90465 paxillin Pxn
1424028_at 0.99202 RIKEN cDNA 5830457O10 gene 5830457O10Rik
1424029_at 0.75138 TSPY-like 4 Tspyl4
1424030_at 0.88219 grainyhead-like 1 (Drosophila) Grhl1
1424031_at 0.93957 sorting nexin 11 Snx11
1424032_at 1.09741 hydrogen voltage-gated channel 1 Hvcn1
1424033_at 0.74190 "splicing factor, arginine/serine-rich 7" Sfrs7
1424034_at 1.15373 RAR-related orphan receptor alpha Rora
1424035_at 1.14244 RAR-related orphan receptor alpha Rora
1424036_at 0.99962 PRP6 pre-mRNA splicing factor 6 homolog (yeast) Prpf6
1424037_at 0.88206 "inositol 1,4,5-trisphosphate 3-kinase A" Itpka
1424038_a_at 0.86242 RIKEN cDNA 2310044H10 gene 2310044H10Rik
1424039_at 0.84856 transmembrane protein 66 Tmem66
1424040_at 0.96212 arginine/proline rich coiled-coil 1 Rprc1

1424041_s_at	2.25994	"complement component 1, s subcomponent"	C1s
1424042_at	1.05586	transmembrane protein 5	Tmem5
1424043_at	1.20838	peptidylprolyl isomerase (cyclophilin)-like 4	Ppil4
1424044_at	1.06747	jumonji domain containing 2B	Jmjd2b
1424045_at	1.06072	RIKEN cDNA 5730437N04 gene	5730437N04Rik
1424046_at	1.00979	budding uninhibited by benzimidazoles 1 homolog (S. cerevisiae)	Bub1
1424047_at	1.04148	2-deoxyribose-5-phosphate aldolase homolog (C. elegans)	Dera
1424048_a_at	0.94564	cytochrome b5 reductase 1	Cyb5r1
1424049_at	0.89099	leucine rich repeat containing 42	Lrrc42
1424050_s_at	0.67295	fibroblast growth factor receptor 1	Fgfr1
1424051_at	0.88595	"procollagen, type IV, alpha 2"	Col4a2
1424052_at	1.01330	THAP domain containing 4	Thap4
1424053_a_at	0.96259	transcription factor 25 (basic helix-loop-helix)	Tcf25
1424054_at	0.93896	BTB (POZ) domain containing 2	Btbd2
1424055_at	0.87224	nuclear receptor coactivator 5	Ncoa5
1424056_at	1.00744	ubiquitin specific peptidase 48	Usp48
1424057_at	0.98123	ganglioside-induced differentiation-associated-protein 2	Gdap2
1424058_at	0.99455	RIKEN cDNA 1190002C06 gene	1190002C06Rik
1424059_at	0.97549	suppressor of variegation 4-20 homolog 2 (Drosophila)	Suv420h2
1424060_at	0.99430	nei like 3 (E. coli)	Neil3
1424061_at	0.95927	"mannosidase, beta A, lysosomal-like"	Manbal
1424062_at	0.91582	"ubiquitin-conjugating enzyme E2D 1, UBC4/5 homolog (yeast)"	Ube2d1
1424063_at	1.04213	androgen binding protein alpha	Abpa
1424064_at	0.92436	"RAB1B, member RAS oncogene family"	Rab1b
1424065_at	0.90184	"ER degradation enhancer, mannosidase alpha-like 1"	Edem1
1424066_at	0.95310	dihydrouridine synthase 3-like (S. cerevisiae)	Dus3l
1424067_at	1.29022	intercellular adhesion molecule	Icam1
1424068_at	0.87779	T-cell leukemia translocation altered gene	Tcta
1424069_at	0.99961	N-ethylmaleimide sensitive fusion protein attachment protein gamma	Napg
1424070_at	1.01956	N-ethylmaleimide sensitive fusion protein attachment protein gamma	Napg
1424071_s_at	0.84653	cDNA sequence BC018507	BC018507
1424072_at	0.96458	RIKEN cDNA 2010107G23 gene	2010107G23Rik
1424073_at	1.03595	RIKEN cDNA 5430437P03 gene	5430437P03Rik
1424074_at	0.87607	basic transcription factor 3-like 4	Btf3l4
1424075_at	1.02386	RIKEN cDNA 9430016H08 gene	9430016H08Rik
1424076_at	1.02017	glycerophosphodiester phosphodiesterase domain containing 1	Gdpd1
1424077_at	1.19009	glycerophosphodiester phosphodiesterase domain containing 1	Gdpd1
1424078_s_at	0.82498	peroxisomal biogenesis factor 6	Pex6
1424079_x_at	0.97104	RIKEN cDNA 2900073H19 gene	2900073H19Rik
1424080_at	0.97629	"decapping enzyme, scavenger"	Dcps
1424081_at	1.07985	polycomb group ring finger 6	Pcgf6
1424082_at	1.04674	"TBC1 domain family, member 13"	Tbc1d13
1424083_at	0.98829	ROD1 regulator of differentiation 1 (S. pombe)	Rod1

1424084_at 1.05755 ROD1 regulator of differentiation 1 (S. pombe) Rod1
1424085_at 0.93413 "NADH dehydrogenase (ubiquinone) 1 alpha subcomplex, 4" Ndufa4
1424086_at 0.96837 "DNA segment, Chr 9, University of California at Los Angeles 1" D9Ucla1
1424087_at 1.03271 RIKEN cDNA 1810042K04 gene 1810042K04Rik
1424088_at 0.96969 melanocyte proliferating gene 1 Myg1
1424089_a_at 0.90531 transcription factor 4 Tcf4
1424090_at 1.02497 syndecan binding protein (syntenin) 2 Sdcbp2
1424091_at 1.12354 expressed sequence AI449175 AI449175
1424092_at 1.07392 erythrocyte protein band 4.1 Epb4.1
1424093_x_at 0.89428 CD151 antigen Cd151
1424094_at 1.00892 NIMA (never in mitosis gene a)-related expressed kinase 9 Nek9
1424095_at 1.10331 RNA terminal phosphate cyclase domain 1 Rtdc1
1424096_at 1.00493 keratin 5 Krt5
1424097_at 0.89320 "ELOVL family member 7, elongation of long chain fatty acids (yeast)" Elovl7
1424098_at 0.85396 "ELOVL family member 7, elongation of long chain fatty acids (yeast)" Elovl7
1424099_at 0.99485 RIKEN cDNA 2310016C16 gene 2310016C16Rik
1424100_s_at 1.10051 cell cycle exit and neuronal differentiation 1 Cend1
1424101_at 0.98073 heterogeneous nuclear ribonucleoprotein L Hnrpl
1424102_at 1.03527 autophagy-related 4B (yeast) Atg4b
1424103_at 0.92961 autophagy-related 4B (yeast) Atg4b
1424104_at 1.01141 "SYF2 homolog, RNA splicing factor (S. cerevisiae)" Syf2
1424105_a_at 1.13182 pituitary tumor-transforming 1 Pttg1
1424106_at 0.85352 RIKEN cDNA 1200003C05 gene 1200003C05Rik
1424107_at 0.95136 kinesin family member 18A Kif18a
1424108_at 0.99031 glyoxalase 1 Glo1
1424109_a_at 0.97291 glyoxalase 1 Glo1
1424110_a_at 1.18210 "expressed in non-metastatic cells 1, protein" Nme1
1424111_at 0.90622 insulin-like growth factor 2 receptor Igf2r
1424112_at 0.88754 insulin-like growth factor 2 receptor Igf2r
1424113_at 0.96650 laminin B1 subunit 1 Lamb1-1
1424114_s_at 0.93240 laminin B1 subunit 1 Lamb1-1
1424115_at 0.91916 "protein phosphatase 5, catalytic subunit" Ppp5c
1424116_x_at 0.97594 "protein phosphatase 5, catalytic subunit" Ppp5c
1424117_at 0.94902 cDNA sequence BC056474 BC056474
1424118_a_at 0.93965 spindle pole body component 25 homolog (S. cerevisiae) Spbc25
1424119_at 0.96772 "protein kinase, AMP-activated, beta 1 non-catalytic subunit" Prkab1
1424120_at 0.97427 ring finger protein 8 Rnf8
1424121_at 0.97764 COMM domain containing 1 Commd1
1424122_s_at 1.02721 COMM domain containing 1 Commd1
1424123_at 0.99720 cDNA sequence BC011209 BC011209
1424124_at 1.08067 motile sperm domain containing 2 Mospd2
1424125_at 1.03314 "potassium channel, subfamily K, member 13" Kcnk13
1424126_at 1.08035 aminolevulinic acid synthase 1 Alas1

1424127_at 0.80959 eyes absent 2 homolog (Drosophila) Eya2
 1424128_x_at 0.98016 aurora kinase B Aurkb
 1424129_at 1.11475 major facilitator superfamily domain containing 1 Mfsd1
 1424130_a_at 1.06761 polymerase I and transcript release factor Ptrf
 1424131_at 0.86240 "procollagen, type VI, alpha 3 /// similar to alpha 3 type VI collagen isoform 1 precursor" Col6a3 /// LOC674521
 1424132_at 1.00650 Harvey rat sarcoma virus oncogene 1 Hras1
 1424133_at 0.89371 transmembrane protein 98 Tmem98
 1424134_at 0.98416 ring finger and SPRY domain containing 1 Rspry1
 1424135_at 0.88560 ring finger and SPRY domain containing 1 Rspry1
 1424136_a_at 1.07247 peptidyl prolyl isomerase H /// similar to peptidyl prolyl isomerase H /// hypothetical protein LOC624822 /// similar to peptidylprolyl isomerase H /// similar to peptidylprolyl isomerase H /// similar to peptidylprolyl isomerase H /// similar to peptidylprolyl isomerase H /// similar to peptidylprolyl isomerase H Ppih /// LOC433064 /// LOC624822 /// LOC665989 /// LOC667598 /// LOC669441 /// LOC670964 /// LOC676493
 1424137_at 1.08830 G protein-regulated inducer of neurite outgrowth 1 Gprin1
 1424138_at 1.12821 rhomboid family 1 (Drosophila) Rhbdf1
 1424139_at 1.00153 RAS-related protein-1a Rap1a
 1424140_at 1.23911 "galactose-4-epimerase, UDP" Gale
 1424141_at 0.99236 HECT domain containing 1 Hectd1
 1424142_at 0.88167 "inhibitor of kappa light polypeptide enhancer in B-cells, kinase complex-associated protein" Ikbkap
 1424143_a_at 0.98332 chromatin licensing and DNA replication factor 1 Cdt1
 1424144_at 0.88624 chromatin licensing and DNA replication factor 1 Cdt1
 1424145_at 0.96275 proline-rich polypeptide 3 Prr3
 1424146_at 0.94934 G protein-coupled receptor 37-like 1 Gpr3711
 1424147_at 0.98977 "AHA1, activator of heat shock protein ATPase homolog 1 (yeast)" Ahsa1
 1424148_a_at 1.00578 expressed sequence AW049765 AW049765
 1424149_at 1.10810 "non-SMC element 2 homolog (MMS21, S. cerevisiae)" Nsmce2
 1424150_at 1.39352 glycerophosphodiester phosphodiesterase domain containing 5 Gdpd5
 1424151_at 1.21742 JTV1 gene Jtv1
 1424152_at 1.08020 CDNA clone IMAGE:30031514 ---
 1424153_s_at 0.99432 sal-like 4 (Drosophila) Sall4
 1424154_a_at 1.04755 HESB like domain containing 1 Hbld1
 1424155_at 0.96048 "fatty acid binding protein 4, adipocyte" Fabp4
 1424156_at 1.00718 retinoblastoma-like 1 (p107) Rbl1
 1424157_at 0.95078 EH-domain containing 2 /// similar to EH-domain containing 2 Ehd2 /// LOC673251
 1424158_at 0.84602 similar to EH-domain containing 2 LOC673251
 1424159_at 0.92436 RIKEN cDNA 1300010M03 gene 1300010M03Rik
 1424160_at 0.97531 "asparagine-linked glycosylation 5 homolog (yeast, dolichyl-phosphate beta-glucosyltransferase)" Alg5
 1424161_at 1.24252 DEAD (Asp-Glu-Ala-Asp) box polypeptide 27 Ddx27

1424162_at	1.01341 tripartite motif protein 29	Trim29	
1424163_at	1.06920 required for meiotic nuclear division 5 homolog B (S. cerevisiae)		Rmnd5b
1424164_at	1.18059 mitochondrial ribosomal protein L50	Mrp150	
1424165_a_at	1.02242 SHANK-associated RH domain interacting protein	Sharpin	
1424166_at	0.90320 mutS homolog 3 (E. coli)	Msh3	
1424167_a_at	0.97580 phosphomannomutase 1	Pmm1	
1424168_a_at	0.95654 "capping protein (actin filament) muscle Z-line, beta"	Capzb	
1424169_at	0.82045 Tax1 (human T-cell leukemia virus type I) binding protein 3		Tax1bp3
1424170_at	0.95623 PHD finger protein 5A	Phf5a	
1424171_a_at	1.14107 hydroxyacyl glutathione hydrolase		Hagh
1424172_at	1.12980 hydroxyacyl glutathione hydrolase		Hagh
1424173_at	0.94317 transmembrane protein 48	Tmem48	
1424174_at	0.95985 Sh3kbp1 binding protein 1	Shkbp1	
1424175_at	0.89854 thyrotroph embryonic factor	Tef	
1424176_a_at	1.00052 annexin A4	Anxa4	
1424177_at	0.85986 transmembrane protein 38a	Tmem38a	
1424178_at	0.97531 transmembrane protein 38a	Tmem38a	
1424179_at	1.03592 "pleckstrin homology domain containing, family J member 1"		Plekhj1
1424180_a_at	1.06085 thyroid hormone receptor associated protein 4	Thrap4	
1424181_at	1.00276 septin 6	6-Sep	
1424182_at	0.91488 acetyl-Coenzyme A acetyltransferase 1	Acat1	
1424183_at	0.93028 acetyl-Coenzyme A acetyltransferase 1	Acat1	
1424184_at	0.84890 "acyl-Coenzyme A dehydrogenase, very long chain"		Acadvl
1424185_a_at	1.04550 mortality factor 4 like 1 /// similar to mortality factor 4 like 1 isoform b ///		
	RIKEN cDNA A730098P11 gene /// similar to mortality factor 4 like 1 /// similar to mortality factor 4		
	like 1 (predicted) Morf4l1 /// LOC433955 /// A730098P11Rik /// LOC627352 /// LOC635663		
1424186_at	0.96265 coiled-coil domain containing 80	Ccdc80	
1424187_at	0.94055 coiled-coil domain containing 80	Ccdc80	
1424188_at	1.07576 RAB GTPase activating protein 1	Rabgap1	
1424189_at	0.95938 "phosphatidylinositol glycan anchor biosynthesis, class C"		Pigc
1424190_at	1.00981 "phosphatidylinositol glycan anchor biosynthesis, class C"		Pigc
1424191_a_at	0.94577 transmembrane protein 41a	Tmem41a	
1424192_at	0.98794 RIKEN cDNA 1500011H22 gene	1500011H22Rik	
1424193_at	1.09594 "PWP2 (periodic tryptophan protein) homolog, yeast"		Pwp2h
1424194_at	1.02797 RCSD domain containing 1	Rcsd1	
1424195_a_at	0.84455 inositol polyphosphate-5-phosphatase D	Inpp5d	
1424196_at	0.96124 "Yip1 domain family, member 1"	Yipf1	
1424197_s_at	0.90486 "Fanconi anemia, complementation group E"		Fance
1424198_at	0.92616 "discs, large homolog 5 (Drosophila)"	Dlg5	
1424199_at	0.98340 SEH1-like (S. cerevisiae)	Seh1l	
1424200_s_at	1.04918 SEH1-like (S. cerevisiae)	Seh1l	
1424201_a_at	1.06598 SEH1-like (S. cerevisiae)	Seh1l	
1424202_at	0.98569 SEH1-like (S. cerevisiae)	Seh1l	

1424203_at 1.03538 nicalin homolog (zebrafish) Ncln
1424204_at 1.00573 mitochondrial ribosomal protein L13 Mrpl13
1424205_at 0.99721 "SWI/SNF related, matrix associated, actin dependent regulator of chromatin, subfamily a, member 5 /// RIKEN cDNA 9230115A19 gene /// similar to SWI/SNF related, matrix associated, actin dependent regulator of chromatin, subfamily a, member 5" Smarca5 /// 9230115A19Rik /// LOC675704
1424206_at 1.01670 "SWI/SNF related, matrix associated, actin dependent regulator of chromatin, subfamily a, member 5 /// RIKEN cDNA 9230115A19 gene /// similar to SWI/SNF related, matrix associated, actin dependent regulator of chromatin, subfamily a, member 5" Smarca5 /// 9230115A19Rik /// LOC675704
1424207_at 1.01403 "SWI/SNF related, matrix associated, actin dependent regulator of chromatin, subfamily a, member 5 /// RIKEN cDNA 9230115A19 gene /// similar to SWI/SNF related, matrix associated, actin dependent regulator of chromatin, subfamily a, member 5" Smarca5 /// 9230115A19Rik /// LOC675704
1424208_at 0.80049 prostaglandin E receptor 4 (subtype EP4) Ptger4
1424209_at 0.98737 arginyl-tRNA synthetase-like Rarsl
1424210_at 1.01303 "SPFH domain family, member 1" Spfh1
1424211_at 1.14379 RIKEN cDNA 5730438N18 gene 5730438N18Rik
1424212_at 0.95274 RIKEN cDNA 9430023L20 gene 9430023L20Rik
1424213_at 1.02987 UbiA prenyltransferase domain containing 1 Ubiad1
1424214_at 0.96265 RIKEN cDNA 9130213B05 gene 9130213B05Rik
1424215_at 0.84135 FUN14 domain containing 1 Fundc1
1424216_a_at 0.99354 poly (A) polymerase alpha Papola
1424217_at 1.05243 poly (A) polymerase alpha Papola
1424218_a_at 0.94184 cAMP responsive element binding protein 3-like 4 Creb3l4
1424219_at 1.03273 mitochondrial rRNA methyltransferase 1 homolog (S. cerevisiae) Mrm1
1424220_a_at 1.08102 porcupine homolog (Drosophila) Porcn
1424221_at 0.82058 sushi domain containing 4 Susd4
1424222_s_at 1.12591 RAD23b homolog (S. cerevisiae) Rad23b
1424223_at 0.88646 RIKEN cDNA 1700020C11 gene 1700020C11Rik
1424224_at 1.00472 ankyrin repeat and SOCS box-containing protein 8 Asb8
1424225_at 0.93522 ankyrin repeat and SOCS box-containing protein 8 Asb8
1424226_at 1.26427 RIKEN cDNA 9030617O03 gene 9030617O03Rik
1424227_at 1.02032 polymerase (RNA) III (DNA directed) polypeptide H Polr3h
1424228_at 0.94044 polymerase (RNA) III (DNA directed) polypeptide H Polr3h
1424229_at 0.88111 dual-specificity tyrosine-(Y)-phosphorylation regulated kinase 3 Dyrk3
1424230_at 1.04291 exocyst complex component 6 Exoc6
1424231_s_at 1.04951 exocyst complex component 6 Exoc6
1424232_a_at 0.88949 cDNA sequence BC025546 BC025546
1424233_at 1.02141 mesenchyme homeobox 2 Meox2
1424234_s_at 0.93310 mesenchyme homeobox 2 Meox2
1424235_at 1.00726 ORM1-like 2 (S. cerevisiae) Ormdl2
1424236_at 1.10719 "TBC1 domain family, member 10b" Tbc1d10b

1424237_at 1.00922 zinc finger protein 639 Zfp639
 1424238_at 0.99424 "sirtuin 7 (silent mating type information regulation 2, homolog) 7 (S. cerevisiae)" Sirt7
 1424239_at 0.84458 RIKEN cDNA 2310066E14 gene 2310066E14Rik
 1424240_at 0.98797 ADP-ribosylation factor interacting protein 2 Arfip2
 1424241_at 1.05153 "solute carrier family 30 (zinc transporter), member 6" Slc30a6
 1424242_at 0.85580 "biphenyl hydrolase-like (serine hydrolase, breast epithelial mucin-associated antigen)" Bphl
 1424243_at 0.95382 RWD domain containing 4A Rwdd4a
 1424244_at 0.98986 RWD domain containing 4A Rwdd4a
 1424245_at 0.95726 carboxylesterase 2 /// similar to carboxylesterase 2 Ces2 /// LOC667754
 1424246_a_at 1.00251 testis derived transcript Tes
 1424247_at 0.96123 ELKS/RAB6-interacting/CAST family member 1 Erc1
 1424248_at 0.94033 "cyclic AMP-regulated phosphoprotein, 21" Arpp21
 1424249_a_at 0.88306 Rho GTPase activating protein 9 Arhgap9
 1424250_a_at 1.07944 Rho guanine nucleotide exchange factor (GEF) 3 Arhgef3
 1424251_a_at 0.96894 heterogeneous nuclear ribonucleoprotein D-like Hnrpdl
 1424252_at 0.88314 heterogeneous nuclear ribonucleoprotein D-like Hnrpdl
 1424253_at 1.02072 RIKEN cDNA 1810073G14 gene 1810073G14Rik
 1424254_at 0.95812 interferon induced transmembrane protein 1 Ifitm1
 1424255_at 1.01564 suppressor of Ty 5 homolog (S. cerevisiae) Supt5h
 1424256_at 0.97760 retinol dehydrogenase 12 Rdh12
 1424257_at 0.99396 cyclin-dependent kinase 7 (homolog of Xenopus MO15 cdk-activating kinase) Cdk7
 1424258_at 1.16454 polymerase (RNA) II (DNA directed) polypeptide D Polr2d
 1424259_at 0.73132 RIKEN cDNA 2400010G15 gene 2400010G15Rik
 1424260_at 1.04392 "solute carrier family 12, member 1" Slc12a1
 1424261_at 0.82770 zinc finger protein 672 Zfp672
 1424262_at 0.90984 RIKEN cDNA 2810003C17 gene 2810003C17Rik
 1424263_at 1.04261 RIKEN cDNA 2810003C17 gene 2810003C17Rik
 1424264_at 1.00709 "mediator of RNA polymerase II transcription, subunit 6 homolog (yeast)" Med6
 1424265_at 0.91426 N-acetylneuraminatase pyruvate lyase Npl
 1424266_s_at 0.93973 expressed sequence AU018778 AU018778
 1424267_at 0.85981 RIKEN cDNA 1810043G02 gene 1810043G02Rik
 1424268_at 0.87142 spermine oxidase Smox
 1424269_a_at 0.92992 "myosin, light polypeptide 6, alkali, smooth muscle and non-muscle /// similar to myosin, light polypeptide 6, alkali, smooth muscle and non-muscle /// similar to Myosin light polypeptide 6 (Myosin light chain alkali 3) (Myosin light chain 3) (MLC-3) (LC17) /// similar to Myosin light polypeptide 6 (Myosin light chain alkali 3) (Myosin light chain 3) (MLC-3) (LC17)" Myl6 /// LOC433297 /// LOC667952 /// LOC671953
 1424270_at 1.12898 doublecortin and calcium/calmodulin-dependent protein kinase-like 1 Dcamk1l

1424271_at 1.18420 doublecortin and calcium/calmodulin-dependent protein kinase-like 1
 Dcamk1l
 1424272_at 1.02566 signal transducer and activator of transcription 3 Stat3
 1424273_at 0.96258 "cytochrome P450, family 2, subfamily c, polypeptide 70" Cyp2c70
 1424274_at 1.12517 vesicle docking protein Vdp
 1424275_s_at 0.96389 tripartite motif-containing 41 Trim41
 1424276_at 0.97683 sorting nexin 16 Snx16
 1424277_at 1.04884 RIKEN cDNA 1110020L19 gene 1110020L19Rik
 1424278_a_at 1.09373 baculoviral IAP repeat-containing 5 Birc5
 1424279_at 1.07865 "fibrinogen, alpha polypeptide" Fga
 1424280_at 0.79970 motile sperm domain containing 1 Mospd1
 1424281_at 0.96530 ubiquitin-associated protein 2 Ubap2
 1424282_at 1.01516 PET112-like (yeast) Pet112l
 1424283_at 0.98362 jumping translocation breakpoint Jtb
 1424284_at 0.85360 protein-O-mannosyltransferase 1 Pomt1
 1424285_s_at 1.03929 ADP-ribosylation factor-like 6 interacting protein 4 Arl6ip4
 1424286_at 0.95544 "protein kinase, X-linked" Prkx
 1424287_at 1.04183 "protein kinase, X-linked" Prkx
 1424288_at 0.91143 "bin3, bicoid-interacting 3, homolog (Drosophila)" Bcdin3
 1424289_at 0.88883 cDNA sequence BC010311 BC010311
 1424290_at 0.79232 cDNA sequence BC010311 BC010311
 1424291_at 1.06581 nucleoporin 93 Nup93
 1424292_at 0.87912 DEP domain containing 1a Depdc1a
 1424293_s_at 1.00250 transmembrane protein 55A Tmem55a
 1424294_at 0.99917 "protein phosphatase 4, regulatory subunit 1" Ppp4r1
 1424295_at 1.13337 developmental pluripotency-associated 3 Dppa3
 1424296_at 0.88143 "glutamate-cysteine ligase, catalytic subunit" Gclc
 1424297_at 0.90546 zinc finger protein 282 Zfp282
 1424298_at 1.03817 zinc finger protein 282 Zfp282
 1424299_at 1.01432 "OMA1 homolog, zinc metallopeptidase (S. cerevisiae)" Oma1
 1424300_at 1.15943 gem (nuclear organelle) associated protein 6 Gemin6
 1424301_at 0.97432 zinc finger protein 219 Zfp219
 1424302_at 0.97716 "leukocyte immunoglobulin-like receptor, subfamily B (with TM and ITIM
 domains), member 3 /// hypothetical LOC619608" Lilrb3 /// LOC619608
 1424303_at 0.97612 DEP domain containing 7 Depdc7
 1424304_at 1.12928 two pore segment channel 2 Tpcn2
 1424305_at 0.94079 immunoglobulin joining chain Igj
 1424306_at 0.98127 "elongation of very long chain fatty acids (FEN1/Elo2, SUR4/Elo3, yeast)-like
 4" Elovl4
 1424307_at 0.94220 Rho GTPase activating protein 1 Arhgap1
 1424308_at 0.80786 "solute carrier family 24 (sodium/potassium/calcium exchanger), member 3"
 Slc24a3
 1424309_a_at 1.05696 molybdenum cofactor synthesis 2 Mocs2

1424310_at 1.08245 --- ---
1424311_at 1.02172 RIKEN cDNA 0710008K08 gene 0710008K08Rik
1424312_at 0.99726 adiponectin receptor 1 Adipor1
1424313_a_at 0.95404 NADH dehydrogenase (ubiquinone) Fe-S protein 7 Ndufs7
1424314_at 0.99280 PRP3 pre-mRNA processing factor 3 homolog (yeast) Prpf3
1424315_at 1.18669 RIKEN cDNA 1110004E09 gene 1110004E09Rik
1424316_at 0.96146 "solute carrier family 25 (mitochondrial deoxynucleotide carrier), member 19"
Slc25a19
1424317_at 1.06593 "solute carrier family 25 (mitochondrial deoxynucleotide carrier), member 19"
Slc25a19
1424318_at 0.90980 RIKEN cDNA 1110067D22 gene 1110067D22Rik
1424319_at 0.98826 oral cancer overexpressed 1 Oraov1
1424320_a_at 1.02827 Tnf receptor-associated factor 7 Traf7
1424321_at 0.94483 replication factor C (activator 1) 4 Rfc4
1424322_at 0.92426 apurinic/aprimidinic endonuclease 2 Apex2
1424323_at 1.17140 nucleolar complex associated 2 homolog (S. cerevisiae) Noc21
1424324_at 1.02841 establishment of cohesion 1 homolog 1 (S. cerevisiae) Esco1
1424325_at 1.01093 establishment of cohesion 1 homolog 1 (S. cerevisiae) Esco1
1424326_at 1.00468 LEM domain containing 2 Lemd2
1424327_at 1.09726 RIKEN cDNA 3200002M19 gene 3200002M19Rik
1424328_s_at 1.04140 RIKEN cDNA 3200002M19 gene 3200002M19Rik
1424329_a_at 0.92452 proline-rich Gla (G-carboxyglutamic acid) polypeptide 2 Prrg2
1424330_at 0.99766 SUMO1/sentrin specific peptidase 1 Senp1
1424331_at 0.96968 "Rab40c, member RAS oncogene family" Rab40c
1424332_at 0.90428 "Rab40c, member RAS oncogene family" Rab40c
1424333_at 1.00633 RNA (guanine-9-) methyltransferase domain containing 1 Rg9mtd1
1424334_at 0.83636 tetraspanin 17 Tspan17
1424335_at 1.06137 phosphopantothenoylcysteine decarboxylase Ppcdc
1424336_at 1.14119 phosphopantothenoylcysteine decarboxylase Ppcdc
1424337_at 0.99755 sorting nexin 15 Snx15
1424338_at 1.07183 "solute carrier family 6 (neurotransmitter transporter, GABA), member 13"
Slc6a13
1424339_at 1.06343 2'-5' oligoadenylate synthetase-like 1 Oas11
1424340_at 0.73170 leucine rich repeat containing 48 Lrrc48
1424341_s_at 0.97154 "protocadherin alpha 4 /// protocadherin alpha 6 /// protocadherin alpha 7 ///
protocadherin alpha 5 /// protocadherin alpha 11 /// protocadherin alpha 10 /// protocadherin alpha 1 ///
protocadherin alpha 9 /// protocadherin alpha 3 /// protocadherin alpha 12 /// protocadherin alpha 2 ///
protocadherin alpha 8 /// protocadherin alpha subfamily C, 1 /// protocadherin alpha subfamily C, 2"
Pcdha4 /// Pcdha6 /// Pcdha7 /// Pcdha5 /// Pcdha11 /// Pcdha10 /// Pcdha1 /// Pcdha9 /// Pcdha3 ///
Pcdha12 /// Pcdha2 /// Pcdha8 /// Pcdhac1 /// Pcdhac2
1424342_at 0.94710 forty-two-three domain containing 1 Fyttl1
1424343_a_at 1.22124 eukaryotic translation initiation factor 1A Eif1a
1424344_s_at 1.14211 eukaryotic translation initiation factor 1A Eif1a

1424345_s_at	0.94239	"ubiquitin-conjugating enzyme E2M (UBC12 homolog, yeast)"	Ube2m
1424346_at	1.04248	"protein phosphatase 6, catalytic subunit"	Ppp6c
1424347_at	0.90111	"protein phosphatase 6, catalytic subunit"	Ppp6c
1424348_at	0.99534	RIKEN cDNA 1110007A13 gene	1110007A13Rik
1424349_a_at	0.85994	lysophosphatidylglycerol acyltransferase 1	Lpgat1
1424350_s_at	0.86211	lysophosphatidylglycerol acyltransferase 1	Lpgat1
1424351_at	1.17855	WAP four-disulfide core domain 2	Wfdc2
1424352_at	0.89880	"cytochrome P450, family 4, subfamily a, polypeptide 12"	Cyp4a12
1424353_at	1.15592	leucine-rich PPR-motif containing	Lrpprc
1424354_at	1.03365	RIKEN cDNA 1110007F12 gene	1110007F12Rik
1424355_a_at	1.13707	"transcriptional regulator, SIN3B (yeast)"	Sin3b
1424356_a_at	0.98472	"meteorin, glial cell differentiation regulator-like"	Metrn1
1424357_at	0.86205	transmembrane protein 45b	Tmem45b
1424358_at	0.92779	"ubiquitin-conjugating enzyme E2E 2 (UBC4/5 homolog, yeast)"	Ube2e2
1424359_at	0.81975	5-oxoprolinase (ATP-hydrolysing)	Oplah
1424360_at	0.96795	cDNA sequence BC019943	BC019943
1424361_at	0.99540	cDNA sequence BC019943	BC019943
1424362_at	0.92135	phosphatidic acid phosphatase type 2 domain containing 3	Ppapdc3
1424363_at	1.00124	Mycbp associated protein	Mycbpap
1424364_a_at	0.95270	RIKEN cDNA 1110020P15 gene	1110020P15Rik
1424365_at	0.98549	RIKEN cDNA 1810037I17 gene	1810037I17Rik
1424366_at	0.93455	transmembrane protein 15	Tmem15
1424367_a_at	0.99881	homer homolog 2 (Drosophila)	Homer2
1424368_s_at	0.99488	ubiquilin 1	Ubqln1
1424369_at	0.91523	"proteasome (prosome, macropain) inhibitor subunit 1"	Psmf1
1424370_s_at	0.99796	"proteasome (prosome, macropain) inhibitor subunit 1"	Psmf1
1424371_at	0.75547	"proteasome (prosome, macropain) inhibitor subunit 1"	Psmf1
1424372_at	1.06192	mitochondrial ribosomal protein L32	Mrpl32
1424373_at	1.02117	"armadillo repeat containing, X-linked 3"	Armxc3
1424374_at	0.92158	"GTPase, IMAP family member 4"	Gimap4
1424375_s_at	1.05814	"GTPase, IMAP family member 4"	Gimap4
1424376_at	0.89252	CDC42 effector protein (Rho GTPase binding) 1	Cdc42ep1
1424377_at	1.03282	similar to ribosomal protein L24-like	LOC670726
1424378_at	1.25436	low density lipoprotein receptor adaptor protein 1	Ldlrap1
1424379_at	0.89471	carbonic anhydrase 11	Car11
1424380_at	1.02973	vacuolar protein sorting 37B (yeast)	Vps37b
1424381_at	0.94495	splicing factor 4	Sf4
1424382_at	0.87897	"reticulocalbin 3, EF-hand calcium binding domain"	Rcn3
1424383_at	1.07647	transmembrane protein 51	Tmem51
1424384_a_at	1.14216	zinc and ring finger 1	Znrf1
1424385_at	1.02329	---	---
1424386_at	1.11860	receptor accessory protein 2	Reep2
1424387_at	1.11001	prolactin-like protein C 3	Prlpc3

1424388_at 0.83844 clusterin associated protein 1 Cluap1
1424389_at 1.00568 nucleoporin like 1 Nupl1
1424390_at 1.08160 Nucleoporin like 1 Nupl1
1424391_at 1.12002 "nardilysin, N-arginine dibasic convertase, NRD convertase 1" Nrd1
1424392_at 1.04659 "alcohol dehydrogenase, iron containing, 1" Adhfe1
1424393_s_at 0.93100 "alcohol dehydrogenase, iron containing, 1" Adhfe1
1424394_at 0.81194 selenoprotein M Selm
1424395_at 0.94635 asparaginase like 1 Asrg11
1424396_a_at 0.96108 asparaginase like 1 Asrg11
1424397_at 0.99956 DEAH (Asp-Glu-Ala-His) box polypeptide 36 Dhx36
1424398_at 1.12199 DEAH (Asp-Glu-Ala-His) box polypeptide 36 Dhx36
1424399_at 0.92644 uridine-cytidine kinase 1 Uck1
1424400_a_at 0.98513 "aldehyde dehydrogenase 1 family, member L1" Aldh111
1424401_at 0.97946 "aldehyde dehydrogenase 1 family, member L1" Aldh111
1424402_at 1.05179 RUN and FYVE domain containing 3 Rufy3
1424403_a_at 0.97742 RUN and FYVE domain containing 3 Rufy3
1424404_at 0.86933 RIKEN cDNA 0610040J01 gene 0610040J01Rik
1424405_at 1.02215 MAP3K12 binding inhibitory protein 1 Mbip
1424406_at 1.03442 BCL2-like 13 (apoptosis facilitator) Bcl2l13
1424407_s_at 0.94991 chromobox homolog 6 Cbx6
1424408_at 0.83583 LIM and senescent cell antigen like domains 2 Lims2
1424409_at 0.96794 claudin 23 Cldn23
1424410_at 0.83133 tetratricopeptide repeat domain 8 Ttc8
1424411_at 0.99354 expressed sequence AI840826 AI840826
1424412_at 0.85520 opioid growth factor receptor-like 1 Ogfr11
1424413_at 0.98773 opioid growth factor receptor-like 1 Ogfr11
1424414_at 0.92845 opioid growth factor receptor-like 1 /// similar to opioid growth factor receptor-like 1 Ogfr11 /// LOC673882
1424415_s_at 0.83608 "spondin 1, (f-spondin) extracellular matrix protein" Spon1
1424416_at 0.89389 NFkB inhibitor interacting Ras-like protein 2 Nkiras2
1424417_at 1.22008 --- ---
1424418_at 0.97160 cDNA sequence BC010801 BC010801
1424419_at 0.94678 leucine rich repeat containing 14 Lrrc14
1424420_at 0.68085 cell cycle progression 1 Ccpg1
1424421_at 0.99788 "RFad1, flavin adenine dinucleotide synthetase, homolog (yeast)" Flad1
1424422_s_at 1.00565 "lens epithelial protein /// RFad1, flavin adenine dinucleotide synthetase, homolog (yeast)" Lenep /// Flad1
1424423_at 0.92062 "lens epithelial protein /// RFad1, flavin adenine dinucleotide synthetase, homolog (yeast)" Lenep /// Flad1
1424424_at 1.05200 "solute carrier family 39 (zinc transporter), member 1" Slc39a1
1424425_a_at 0.87041 methylthioadenosine phosphorylase Mtap
1424426_at 0.91537 methylthioadenosine phosphorylase Mtap
1424427_at 0.89491 "transcriptional adaptor 1 (HF11 homolog, yeast) like" Tada11

1424428_at 1.06096 coiled-coil domain containing 95 Ccdc95
1424429_s_at 1.09267 coiled-coil domain containing 95 Ccdc95
1424430_at 1.13598 MTERF domain containing 2 Mterfd2
1424431_at 1.00786 chondroitin sulfate GalNAcT-2 Galnact2
1424432_at 1.03363 ubiquitin domain containing 1 Ubtd1
1424433_at 0.93998 methionine sulfoxide reductase B2 Msrb2
1424434_at 1.00700 cDNA sequence BC024814 BC024814
1424435_a_at 1.10908 phosphoribosylglycinamide formyltransferase Gart
1424436_at 0.98486 phosphoribosylglycinamide formyltransferase Gart
1424437_s_at 1.07127 "ATP-binding cassette, sub-family G (WHITE), member 4" Abcg4
1424438_a_at 0.85018 leptin receptor overlapping transcript Leprot
1424439_at 0.87545 RIKEN cDNA 1810065E05 gene 1810065E05Rik
1424440_at 1.08793 mitochondrial ribosomal protein S6 Mrps6
1424441_at 0.99322 "solute carrier family 27 (fatty acid transporter), member 4" Slc27a4
1424442_a_at 0.88986 "praja 2, RING-H2 motif containing" Pja2
1424443_at 0.86758 "hepatoma-derived growth factor, related protein 3 /// transmembrane 6 superfamily member 1" Hdgfrp3 /// Tm6sf1
1424444_a_at 0.97490 RIKEN cDNA 1600014C10 gene 1600014C10Rik
1424445_at 0.96370 transmembrane 4 superfamily member 5 Tm4sf5
1424446_at 0.94357 armadillo repeat containing 7 Armc7
1424447_at 0.95098 RIKEN cDNA 1700030K09 gene 1700030K09Rik
1424448_at 1.14508 tripartite motif protein 6 /// similar to Tripartite motif protein 6 Trim6 /// LOC633467
1424449_at 0.86953 fibroblast growth factor receptor substrate 3 Frs3
1424450_at 0.84344 "G protein-coupled receptor, family C, group 5, member C" Gprc5c
1424451_at 0.67622 acetyl-Coenzyme A acyltransferase 1B Acaa1b
1424452_at 1.18217 "SAFB-like, transcription modulator" Sltm
1424453_at 0.99864 "phosphate cytidylyltransferase 1, choline, alpha isoform" Pcyt1a
1424454_at 0.96567 transmembrane protein 87A Tmem87a
1424455_at 0.70583 G protein-coupled receptor associated sorting protein 1 Gprasp1
1424456_at 1.11574 poliovirus receptor-related 2 Pvr12
1424457_at 0.98692 "amyloid beta (A4) precursor protein-binding, family B, member 3" Apbb3
1424458_at 0.91097 jumonji domain containing 2C Jmjd2c
1424459_at 0.95440 acyltransferase like 2 Aytl2
1424460_s_at 0.93213 acyltransferase like 2 Aytl2
1424461_at 0.94036 dynactin 2 Dctn2
1424462_at 1.02020 ERGIC and golgi 2 Ergic2
1424463_at 0.79846 RIKEN cDNA 2210010L05 gene 2210010L05Rik
1424464_s_at 0.95045 RIKEN cDNA 2210010L05 gene 2210010L05Rik
1424465_at 1.25275 coiled-coil domain containing 58 Ccdc58
1424466_at 0.95267 importin 9 Ipo9
1424467_at 0.96441 "pleckstrin homology-like domain, family B, member 1" Phldb1
1424468_s_at 0.89317 "pleckstrin homology-like domain, family B, member 1" Phldb1

1424469_a_at 0.99815 cleavage and polyadenylation specific factor 4 Cpsf4
1424470_a_at 1.11368 Rap guanine nucleotide exchange factor (GEF) 3 Rapgef3
1424471_at 1.22536 Rap guanine nucleotide exchange factor (GEF) 3 Rapgef3
1424472_at 0.99919 nucleolar protein family 6 (RNA-associated) Nol6
1424473_at 1.01257 polymerase (RNA) II (DNA directed) polypeptide H Polr2h
1424474_a_at 1.08152 "calcium/calmodulin-dependent protein kinase kinase 2, beta" Camkk2
1424475_at 1.07546 "calcium/calmodulin-dependent protein kinase kinase 2, beta" Camkk2
1424476_at 1.09517 "calcium/calmodulin-dependent protein kinase kinase 2, beta" Camkk2
1424477_at 0.95059 cDNA sequence BC019731 BC019731
1424478_at 0.97341 Bardet-Biedl syndrome 2 homolog (human) Bbs2
1424479_at 0.99917 cystatin 8 (cystatin-related epididymal spermatogenic) Cst8
1424480_s_at 0.95564 thymoma viral proto-oncogene 2 Akt2
1424481_s_at 1.00503 "armadillo repeat containing, X-linked 5" Armcx5
1424482_at 1.01320 Rho guanine nucleotide exchange factor (GEF7) Arhgef7
1424483_at 1.08558 --- ---
1424484_at 0.99726 "MOB1, Mps One Binder kinase activator-like 1B (yeast)" Mobk1b
1424485_at 1.02737 angiopoietin-like 3 Angptl3
1424486_a_at 0.90321 thioredoxin reductase 1 Txnrd1
1424487_x_at 0.97962 thioredoxin reductase 1 Txnrd1
1424488_a_at 0.93581 pyrophosphatase (inorganic) 2 Ppa2
1424489_a_at 1.14589 tRNA isopentenyltransferase 1 Trit1
1424490_at 1.09817 zinc finger protein 428 Zfp428
1424491_at 0.90786 --- ---
1424492_at 0.88807 "transient receptor potential cation channel, subfamily C, member 2" Trpc2
1424493_s_at 0.99195 "UDP glycosyltransferases 3 family, polypeptide A1" Ugt3a1
1424494_s_at 0.83715 RIKEN cDNA 2810417J12 gene 2810417J12Rik
1424495_a_at 1.08663 chemokine-like factor Cklf
1424496_at 1.06970 RIKEN cDNA 5133401N09 gene 5133401N09Rik
1424497_at 0.97456 RIKEN cDNA 1110032O16 gene 1110032O16Rik
1424498_at 1.05568 RIKEN cDNA 5730596K20 gene 5730596K20Rik
1424499_s_at 1.06523 RIKEN cDNA 5730596K20 gene 5730596K20Rik
1424500_at 1.13882 "UTP6, small subunit (SSU) processome component, homolog (yeast)" Utp6
1424501_at 1.04374 "UTP6, small subunit (SSU) processome component, homolog (yeast)" Utp6
1424502_at 0.97001 oncoprotein induced transcript 1 Oit1
1424503_at 1.00312 "RAB22A, member RAS oncogene family" Rab22a
1424504_at 1.00208 "RAB22A, member RAS oncogene family" Rab22a
1424505_at 0.93065 RIKEN cDNA 0610042C05 gene 0610042C05Rik
1424506_at 0.91247 cDNA sequence BC026432 BC026432
1424507_at 1.16612 Ras and Rab interactor 1 Rin1
1424508_at 0.95770 tetratricopeptide repeat domain 5 Ttc5
1424509_at 1.09239 CD177 antigen Cd177
1424510_at 1.26594 nudix (nucleoside diphosphate linked moiety X)-type motif 6 Nudt6
1424511_at 1.02610 aurora kinase A Aurka

1424512_a_at 1.04741 thioredoxin domain containing 9 Txndc9
 1424513_at 1.05434 "ATPase, H⁺ transporting, lysosomal V0 subunit A2" Atp6v0a2
 1424514_at 1.10780 ring finger protein 126 Rnf126
 1424515_at 1.17407 RIKEN cDNA B230354K17 gene B230354K17Rik
 1424516_at 1.12569 RIKEN cDNA B230354K17 gene B230354K17Rik
 1424517_at 1.03209 coiled-coil domain containing 12 Ccdc12
 1424518_at 1.73497 RIKEN cDNA 2310016F22 gene /// cDNA sequence BC020489
 2310016F22Rik /// BC020489
 1424519_at 1.01124 mitochondrial GTPase 1 homolog (S. cerevisiae) Mtg1
 1424520_at 1.14661 RIKEN cDNA 2010305A19 gene 2010305A19Rik
 1424521_at 0.94913 "zinc finger, AN1 type domain 2B" Zfand2b
 1424522_at 1.24004 HEAT repeat containing 1 Heatr1
 1424523_at 0.97638 "engulfment and cell motility 1, ced-12 homolog (C. elegans)" Elmo1
 1424524_at 0.80223 RIKEN cDNA 1200002N14 gene 1200002N14Rik
 1424525_at 1.00548 gastrin releasing peptide Grp
 1424526_a_at 1.02566 "TDP-glucose 4,6-dehydratase" Tgds
 1424527_at 0.92791 "protein phosphatase 2, regulatory subunit B, delta isoform" Ppp2r2d
 1424528_at 0.86395 cell growth regulator with EF hand domain 1 Cgref1
 1424529_s_at 0.93367 cell growth regulator with EF hand domain 1 Cgref1
 1424530_at 0.88160 SEC14-like 2 (S. cerevisiae) Sec14l2
 1424531_a_at 0.71230 "transcription elongation factor A (SII), 3" Tcea3
 1424532_at 0.98919 YLP motif containing 1 Ylpm1
 1424533_a_at 0.97983 RIKEN cDNA 3110023B02 gene 3110023B02Rik
 1424534_at 1.13555 monocyte to macrophage differentiation-associated 2 Mmd2
 1424535_at 0.89333 RIKEN cDNA 4930521E07 gene 4930521E07Rik
 1424536_at 0.98301 2'-5' oligoadenylate synthetase 1E Oas1e
 1424537_at 1.04866 RIKEN cDNA 1700017G21 gene /// similar to Germ cell-less protein-like 2
 (mGclh) /// similar to Germ cell-less protein-like 2 (mGclh) /// similar to Germ cell-less protein-like 2
 (mGclh) /// similar to Germ cell-less protein-like 2 (mGclh) /// similar to Germ cell-less protein-like 2
 (mGclh) /// similar to Germ cell-less protein-like 2 (mGclh) /// similar to Germ cell-less protein-like 2
 (mGclh) /// similar to Germ cell-less protein-like 2 (mGclh) /// similar to Germ cell-less protein-like 2
 (mGclh) 1700017G21Rik /// LOC434724 /// LOC434725 /// LOC627264 /// LOC630022 ///
 LOC668958 /// LOC668960 /// LOC668963 /// LOC668964 /// LOC672864
 1424538_at 1.05886 ubiquitin-like 4 Ubl4
 1424539_at 0.93839 ubiquitin-like 4 Ubl4
 1424540_at 0.99955 homeodomain interacting protein kinase 1 Hipk1
 1424541_at 1.00077 transmembrane protein 70 Tmem70
 1424542_at 1.33595 S100 calcium binding protein A4 S100a4
 1424543_at 0.91692 non-catalytic region of tyrosine kinase adaptor protein 1 Nck1
 1424544_at 0.87220 nuclear receptor binding protein 2 Nrbp2
 1424545_at 1.04686 cDNA sequence BC003965 BC003965
 1424546_at 0.91595 cDNA sequence BC003965 BC003965
 1424547_at 1.03615 carbonic anhydrase 10 Car10

1424548_at 0.89139 "zinc finger, CCCH-type with G patch domain" Zgpat
 1424549_at 1.02413 "degenerative spermatocyte homolog 2 (Drosophila), lipid desaturase" Degs2
 1424550_at 0.93439 "zinc finger, FYVE domain containing 27" Zfyve27
 1424551_at 0.93893 "zinc finger, FYVE domain containing 27" Zfyve27
 1424552_at 0.96212 caspase 8 Casp8
 1424553_at 1.05165 "Gup1, glycerol uptake/transporter homolog (yeast)" Gup1
 1424554_at 0.96788 "protein phosphatase 1, regulatory (inhibitor) subunit 8" Ppp1r8
 1424555_at 0.86184 RIKEN cDNA 9430015G10 gene 9430015G10Rik
 1424556_at 0.94427 pyrroline-5-carboxylate reductase 1 Pycr1
 1424557_at 0.99253 "zinc finger, CCHC domain containing 9" Zcchc9
 1424558_a_at 1.01437 calcium-binding tyrosine-(Y)-phosphorylation regulated (fibrousheathin 2)
 Cabyr
 1424559_at 0.87104 expressed sequence AW060207 AW060207
 1424560_at 0.97401 proline-serine-threonine phosphatase-interacting protein 1 Pstpip1
 1424561_at 1.00877 endothelin converting enzyme 2 Ece2
 1424562_a_at 1.00545 "solute carrier family 25 (mitochondrial carrier, adenine nucleotide
 translocator), member 4" Slc25a4
 1424563_at 0.96979 "solute carrier family 25 (mitochondrial carrier, adenine nucleotide
 translocator), member 4" Slc25a4
 1424564_at 0.95686 RIKEN cDNA 2410001C21 gene 2410001C21Rik
 1424565_at 0.94872 polymerase (RNA) III (DNA directed) polypeptide D Polr3d
 1424566_s_at 0.95868 polymerase (RNA) III (DNA directed) polypeptide D Polr3d
 1424567_at 1.04386 tetraspanin 2 Tspan2
 1424568_at 0.98263 tetraspanin 2 Tspan2
 1424569_at 1.04533 DEAD (Asp-Glu-Ala-Asp) box polypeptide 46 Ddx46
 1424570_at 1.12640 DEAD (Asp-Glu-Ala-Asp) box polypeptide 46 Ddx46
 1424571_at 0.96142 DEAD (Asp-Glu-Ala-Asp) box polypeptide 46 Ddx46
 1424572_a_at 0.98845 "H2A histone family, member Y" H2afy
 1424573_at 1.03248 transmembrane emp24 protein transport domain containing 5 Tmed5
 1424574_at 1.09854 transmembrane emp24 protein transport domain containing 5 Tmed5
 1424575_at 0.93764 "RAB, member of RAS oncogene family-like 5" Rabl5
 1424576_s_at 1.06205 "cytochrome P450, family 2, subfamily c, polypeptide 44" Cyp2c44
 1424577_at 1.05995 misato homolog 1 (Drosophila) Msto1
 1424578_at 0.94808 arrestin domain containing 1 Arrdc1
 1424579_at 0.97043 "solute carrier family 35 (UDP-N-acetylglucosamine (UDP-GlcNAc)
 transporter), member 3" Slc35a3
 1424580_at 1.05103 "solute carrier family 35 (UDP-N-acetylglucosamine (UDP-GlcNAc)
 transporter), member 3" Slc35a3
 1424581_at 1.02106 SH3 and cysteine rich domain 2 Stac2
 1424582_at 1.07532 multiple substrate lipid kinase /// similar to multi-substrate lipid kinase ///
 similar to multi-substrate lipid kinase Mulk /// LOC667266 /// LOC672697
 1424583_at 1.01593 "FERM, RhoGEF and pleckstrin domain protein 2" Farp2
 1424584_a_at 0.99754 RAN binding protein 10 Ranbp10

1424585_at 1.03774 RAN binding protein 10 Ranbp10
1424586_at 0.85502 EH domain binding protein 1 Ehbp1
1424587_at 1.16009 cDNA sequence BC023488 BC023488
1424588_at 0.97448 SLIT-ROBO Rho GTPase activating protein 3 Srgap3
1424589_s_at 0.88132 "RNA-binding region (RNP1, RRM) containing 3" Rnpc3
1424590_at 0.90037 DEAD (Asp-Glu-Ala-Asp) box polypeptide 19b Ddx19b
1424591_at 0.85770 RIKEN cDNA 5830433M19 gene 5830433M19Rik
1424592_a_at 1.12099 deoxyribonuclease I Dnase1
1424593_at 0.96199 ecdysoneless homolog (Drosophila) Ecd
1424594_at 1.14609 "lectin, galactose binding, soluble 7" Lgals7
1424595_at 0.91170 F11 receptor F11r
1424596_s_at 0.74638 LIM and cysteine-rich domains 1 Lmcd1
1424597_at 1.00638 open reading frame 19 ORF19
1424598_at 0.95560 DEAD (Asp-Glu-Ala-Asp) box polypeptide 6 Ddx6
1424599_at 1.37209 fibrinogen-like protein 1 Fgl1
1424600_at 1.00249 "amiloride binding protein 1 (amine oxidase, copper-containing)" Abp1
1424601_at 0.92072 X-ray repair complementing defective repair in Chinese hamster cells 4 Xrcc4
1424602_s_at 0.88395 X-ray repair complementing defective repair in Chinese hamster cells 4 Xrcc4
1424603_at 1.02567 sulfatase modifying factor 1 Sumf1
1424604_s_at 1.00711 sulfatase modifying factor 1 Sumf1
1424605_at 1.05693 proprotein convertase subtilisin/kexin type 5 Pcsk5
1424606_at 1.07433 complexin 3 Cplx3
1424607_a_at 1.60719 cDNA sequence BC003993 /// hypothetical LOC544818 BC003993 ///
LOC544818
1424608_a_at 1.18935 basic leucine zipper and W2 domains 2 Bzw2
1424609_a_at 1.76789 cDNA sequence BC003993 /// hypothetical LOC544818 BC003993 ///
LOC544818
1424610_at 1.01209 TruB pseudouridine (psi) synthase homolog 2 (E. coli) Trub2
1424611_x_at 0.97109 TruB pseudouridine (psi) synthase homolog 2 (E. coli) Trub2
1424612_at 0.91322 NIPA-like domain containing 2 Npal2
1424613_at 0.96592 "G protein-coupled receptor, family C, group 5, member B" Gprc5b
1424614_at 0.95176 FGF receptor activating protein 1 Frag1
1424615_at 1.04115 FGF receptor activating protein 1 Frag1
1424616_s_at 0.90294 FGF receptor activating protein 1 Frag1
1424617_at 1.11692 interferon-induced protein 35 Ifi35
1424618_at 1.01250 4-hydroxyphenylpyruvic acid dioxygenase Hpd
1424619_at 0.94988 "splicing factor 3b, subunit 4 /// similar to splicing factor 3b, subunit 4
(predicted) /// similar to splicing factor 3b, subunit 4 (predicted)" Sf3b4 /// LOC634414 /// LOC666113
1424620_at 1.12793 "DNA segment, Chr 13, Wayne State University 177, expressed" D13Wsu177e
1424621_at 0.94509 EST AA792894 AA792894
1424622_at 0.98689 heat shock factor 1 Hsf1
1424623_at 1.07231 "serine (or cysteine) peptidase inhibitor, clade B, member 5" Serpinb5
1424624_at 0.98028 RIKEN cDNA 2900011O08 gene 2900011O08Rik

1424625_a_at 0.77555 DENN/MADD domain containing 1A Dennd1a
1424626_at 1.08815 RIKEN cDNA 2010003K11 gene 2010003K11Rik
1424627_at 1.11934 cystatin 12 Cst12
1424628_a_at 1.05725 RIKEN cDNA 1500032D16 gene 1500032D16Rik
1424629_at 0.89912 breast cancer 1 Brca1
1424630_a_at 0.95383 breast cancer 1 Brca1
1424631_a_at 1.00197 Immunoglobulin heavy chain (gamma polypeptide) Ighg
1424632_a_at 0.81230 "REV3-like, catalytic subunit of DNA polymerase zeta RAD54 like (S. cerevisiae)" Rev3l
1424633_at 0.87019 calcium/calmodulin-dependent protein kinase I gamma Camk1g
1424634_at 1.00151 transcription elongation factor A (SII)-like 1 Tceal1
1424635_at 0.99843 eukaryotic translation elongation factor 1 alpha 1 Eef1a1
1424636_at 0.95260 RIKEN cDNA 2610204L23 gene 2610204L23Rik
1424637_s_at 0.96792 RIKEN cDNA 2610204L23 gene 2610204L23Rik
1424638_at 1.16762 cyclin-dependent kinase inhibitor 1A (P21) Cdkn1a
1424639_a_at 1.00319 3-hydroxy-3-methylglutaryl-Coenzyme A lyase Hmgcl
1424640_at 1.05849 ADP-ribosylation factor-like 8A Arl8a
1424641_a_at 0.89571 THO complex 1 Thoc1
1424642_at 1.01318 THO complex 1 Thoc1
1424643_at 1.05508 "Treacher Collins Franceschetti syndrome 1, homolog" Tcof1
1424644_at 0.94849 tubulin-specific chaperone c Tbcc
1424645_at 0.99032 trinucleotide repeat containing 6C Tnrc6c
1424646_at 0.91842 uridine-cytidine kinase 1-like 1 Uck1l
1424647_at 1.02046 "gamma-aminobutyric acid (GABA-A) receptor, pi" Gabrp
1424648_at 1.08269 "RAB, member of RAS oncogene family-like 4" Rabl4
1424649_a_at 0.91144 tetraspanin 8 Tspan8
1424650_at 0.91591 protein disulfide isomerase associated 5 Pdia5
1424651_at 0.98015 cDNA sequence BC021611 BC021611
1424652_at 0.99722 transmembrane protein 166 Tmem166
1424653_at 1.04755 similar to Tetraspanin-15 (Tspan-15) (Transmembrane 4 superfamily member 15) (Tetraspan NET-7) LOC669875
1424654_at 0.96947 "acid phosphatase 2, lysosomal" Acp2
1424655_at 1.03227 "acid phosphatase 2, lysosomal" Acp2
1424656_s_at 0.97937 ubiquitin specific peptidase 19 Usp19
1424657_at 1.17614 TAO kinase 1 Taok1
1424658_at 1.07314 TAO kinase 1 Taok1
1424659_at 1.04373 slit homolog 2 (Drosophila) Slit2
1424660_s_at 0.95052 CREB regulated transcription coactivator 2 Crtc2
1424661_at 1.00439 peptidase (mitochondrial processing) alpha Pmpca
1424662_at 1.13156 peptidase (mitochondrial processing) alpha Pmpca
1424663_at 1.03235 cDNA sequence BC017647 BC017647
1424664_at 0.98509 cDNA sequence BC017647 BC017647
1424665_at 0.99828 RIKEN cDNA 5430405G24 gene 5430405G24Rik

1424666_at	1.04256	RIKEN cDNA 5430405G24 gene	5430405G24Rik
1424667_a_at	0.90940	cut-like 1 (Drosophila) Cut11	
1424668_a_at	0.88107	cut-like 1 (Drosophila) Cut11	
1424669_at	0.93145	"zinc finger, FYVE domain containing 21"	Zfyve21
1424670_s_at	0.97160	"zinc finger, FYVE domain containing 21"	Zfyve21
1424671_at	1.15093	"pleckstrin homology domain containing, family F (with FYVE domain) member 1"	Plekhf1
1424672_at	0.86275	Dmx-like 1	Dmx11
1424673_at	1.05803	"C-type lectin domain family 2, member h"	Clec2h
1424674_at	1.02565	"solute carrier family 39 (metal ion transporter), member 6"	Slc39a6
1424675_at	1.04099	"solute carrier family 39 (metal ion transporter), member 6"	Slc39a6
1424676_s_at	1.06480	SEC14-like 4 (<i>S. cerevisiae</i>)	Sec1414
1424677_at	0.90049	"cytochrome P450, family 2, subfamily j, polypeptide 9"	Cyp2j9
1424678_at	0.85272	actin-like 7b	Act17b
1424679_at	1.04776	mab-21-like 1 (<i>C. elegans</i>)	Mab2111
1424680_at	0.93096	expressed sequence BB146404	BB146404
1424681_a_at	1.11119	"proteasome (prosome, macropain) subunit, alpha type 5"	Psma5
1424682_at	1.08320	"ATP binding domain 1 family, member C"	Atpbd1c
1424683_at	1.02825	RIKEN cDNA 1810015C04 gene	1810015C04Rik
1424684_at	1.02091	"RAB5C, member RAS oncogene family"	Rab5c
1424685_at	1.06032	exosome component 4	Exosc4
1424686_at	1.08616	RIKEN cDNA 2700008B19 gene	2700008B19Rik
1424687_at	0.96481	RIKEN cDNA 2700008B19 gene	2700008B19Rik
1424688_at	1.02800	cAMP responsive element binding protein 3-like 3	Creb3l3
1424689_at	0.90702	"protease, serine, 32"	Prss32
1424690_at	0.99037	RIKEN cDNA 5930434B04 gene	5930434B04Rik
1424691_at	0.84466	RIKEN cDNA 5930434B04 gene	5930434B04Rik
1424692_at	0.99731	RIKEN cDNA 2810055F11 gene	2810055F11Rik
1424693_at	1.08083	RIKEN cDNA 4933407N01 gene	4933407N01Rik
1424694_at	0.53024	RIKEN cDNA 2010011I20 gene	2010011I20Rik
1424695_at	0.72375	RIKEN cDNA 2010011I20 gene	2010011I20Rik
1424696_at	0.97078	G protein-coupled receptor 89	Gpr89
1424697_at	1.30217	DTW domain containing 1	Dtwd1
1424698_s_at	0.84020	grancalcin	Gca
1424699_at	1.01501	RIKEN cDNA 4921511K06 gene	4921511K06Rik
1424700_at	0.92274	transmembrane protein 38B	Tmem38b
1424701_at	0.92424	protocadherin 20	Pcdh20
1424702_a_at	1.11149	RIKEN cDNA 2410024A21 gene	2410024A21Rik
1424703_at	1.02565	HemK methyltransferase family member 1	Hemk1
1424704_at	1.22375	runt related transcription factor 2	Runx2
1424705_at	0.97000	"RNA binding motif protein, X-linked 2"	RbmX2
1424706_at	0.88899	zinc finger protein 51	Zfp51
1424707_at	0.92638	transmembrane emp24-like trafficking protein 10 (yeast)	Tmed10

1424708_at 1.03205 transmembrane emp24-like trafficking protein 10 (yeast) Tmed10
 1424709_at 1.03986 "sterol-C5-desaturase (fungal ERG3, delta-5-desaturase) homolog (S. cerevisae)" Sc5d
 1424710_a_at 0.96771 golgi reassembly stacking protein 2 Gorasp2
 1424711_at 1.10157 transmembrane protein 2 Tmem2
 1424712_at 1.08105 AT hook containing transcription factor 1 Ahctf1
 1424713_at 0.60312 calmodulin-like 4 Calml4
 1424714_at 1.10251 "aldolase 3, C isoform" Aldoc
 1424715_at 0.80768 "retinol saturase (all trans retinol 13,14 reductase)" Retsat
 1424716_at 0.77045 "retinol saturase (all trans retinol 13,14 reductase)" Retsat
 1424717_at 1.16444 MIS12 homolog (yeast) Mis12
 1424718_at 0.89367 microtubule-associated protein tau Mapt
 1424719_a_at 0.98994 microtubule-associated protein tau Mapt
 1424720_at 0.93662 "mannoside acetylglucosaminyltransferase 4, isoenzyme B" Mgat4b
 1424721_at 1.03597 microfibrillar-associated protein 3 Mfap3
 1424722_at 1.15353 RIKEN cDNA 1300017J02 gene 1300017J02Rik
 1424723_s_at 1.04257 "cleavage stimulation factor, 3' pre-RNA, subunit 3" Cstf3
 1424724_a_at 1.18956 "DNA segment, Chr 16, ERATO Doi 472, expressed" D16Ertd472e
 1424725_at 0.92946 "DNA segment, Chr 16, ERATO Doi 472, expressed" D16Ertd472e
 1424726_at 1.11340 transmembrane protein 150 Tmem150
 1424727_at 1.02440 chemokine (C-C motif) receptor 5 Ccr5
 1424728_at 1.13788 cDNA sequence BC011248 BC011248
 1424729_at 1.02508 cDNA sequence BC054059 BC054059
 1424730_a_at 0.92932 "solute carrier family 15 (H⁺/peptide transporter), member 2" Slc15a2
 1424731_at 1.16206 notchless homolog 1 (Drosophila) Nle1
 1424732_s_at 1.19705 RIKEN cDNA 3110005G23 gene 3110005G23Rik
 1424733_at 0.97927 "purinergic receptor P2Y, G-protein coupled, 14" P2ry14
 1424734_at 1.08169 RAS protein-specific guanine nucleotide-releasing factor 1 Rasgrf1
 1424735_at 1.07174 "solute carrier family 25 (mitochondrial carrier, phosphate carrier), member 25" Slc25a25
 1424736_at 1.02203 eukaryotic translation elongation factor 2 Eef2
 1424737_at 0.93074 thyroid hormone responsive SPOT14 homolog (Rattus) Thrsp
 1424738_at 1.01141 RIKEN cDNA 4932432K03 gene 4932432K03Rik
 1424739_at 0.95291 RIKEN cDNA 4932432K03 gene 4932432K03Rik
 1424740_at 0.99938 cAMP responsive element binding protein 3 Creb3
 1424741_s_at 0.95706 cAMP responsive element binding protein 3 Creb3
 1424742_at 1.08028 --- ---
 1424743_at 1.03328 RIKEN cDNA 2610003J06 gene 2610003J06Rik
 1424744_at 0.85764 serine dehydratase Sds
 1424745_at 1.09083 RIKEN cDNA 2900006B13 gene 2900006B13Rik
 1424746_at 0.85071 --- ---
 1424747_at 0.80685 Kinesin family member 1C Kif1c
 1424748_at 0.82054 UDP-N-acetyl-alpha-D-galactosamine:polypeptide N-

acetylgalactosaminyltransferase 11 Galnt11
 1424749_at 0.94881 WD repeat and FYVE domain containing 1 Wdfy1
 1424750_at 0.99041 zinc finger and BTB domain containing 1 Zbtb1
 1424751_at 0.99635 activator of basal transcription Abt1
 1424752_x_at 1.05727 "zinc finger protein 71, related sequence 1" Zfp71-rs1
 1424753_at 0.92842 nudix (nucleoside diphosphate linked moiety X)-type motif 14 Nudt14
 1424754_at 0.84795 "membrane-spanning 4-domains, subfamily A, member 7" Ms4a7
 1424755_at 0.83427 huntingtin interacting protein 1 Hip1
 1424756_at 1.01910 huntingtin interacting protein 1 Hip1
 1424757_at 0.82820 cDNA sequence BC018242 BC018242
 1424758_s_at 1.09125 "serine (or cysteine) peptidase inhibitor, clade A (alpha-1 antiproteinase, antitrypsin), member 10" Serpina10
 1424759_at 0.95790 arrestin domain containing 4 Arrdc4
 1424760_a_at 1.09319 SET and MYND domain containing 2 Smyd2
 1424761_at 0.92581 cDNA sequence BC011487 BC011487
 1424762_at 0.87835 C1q and tumor necrosis factor related protein 5 /// membrane-type frizzled-related protein C1qtnf5 /// Mfrp
 1424763_at 0.88705 RIKEN cDNA 1700027N10 gene 1700027N10Rik
 1424764_at 0.99215 seizure related 6 homolog like Sez6l
 1424765_at 1.06610 epidermal growth factor receptor pathway substrate 15-like 1 Eps15l1
 1424766_at 0.97636 excision repair cross-complementing rodent repair deficiency complementation group 6 - like Ercc6l
 1424767_at 1.14163 cadherin 22 Cdh22
 1424768_at 1.06764 caldesmon 1 Cald1
 1424769_s_at 0.93774 caldesmon 1 Cald1
 1424770_at 1.04796 caldesmon 1 Cald1
 1424771_at 1.13535 "H2A histone family, member J /// similar to H2A histone family, member J isoform 2" H2afj /// LOC632401
 1424772_at 1.30113 "H2A histone family, member J /// similar to H2A histone family, member J isoform 2" H2afj /// LOC632401
 1424773_at 1.00099 RIKEN cDNA 1110012M11 gene 1110012M11Rik
 1424774_s_at 1.00987 RIKEN cDNA 3230401I01 gene 3230401I01Rik
 1424775_at 1.05372 2'-5' oligoadenylate synthetase 1A Oas1a
 1424776_a_at 0.98952 "solute carrier family 25, member 28" Slc25a28
 1424777_at 1.05609 WD repeat domain 21 Wdr21
 1424778_at 1.05778 "DNA segment, Chr 10, University of California at Los Angeles 1" D10Ucla1
 1424779_at 1.14898 "DNA segment, Chr 10, University of California at Los Angeles 1" D10Ucla1
 1424780_a_at 1.06301 "DNA segment, Chr 10, University of California at Los Angeles 1" D10Ucla1
 1424781_at 1.14304 "DNA segment, Chr 10, University of California at Los Angeles 1" D10Ucla1

1424782_at 0.94978 transmembrane protein 77 Tmem77
 1424783_a_at 0.88449 "UDP glucuronosyltransferase 1 family, polypeptide A2 /// UDP
 glucuronosyltransferase 1 family, polypeptide A6A /// UDP glycosyltransferase 1 family, polypeptide
 A10 /// UDP glucuronosyltransferase 1 family, polypeptide A7C /// UDP glucuronosyltransferase 1
 family, polypeptide A5 /// UDP glucuronosyltransferase 1 family, polypeptide A9 /// UDP
 glucuronosyltransferase 1 family, polypeptide A6B /// UDP glucuronosyltransferase 1 family,
 polypeptide A1" Ugt1a2 /// Ugt1a6a /// Ugt1a10 /// Ugt1a7c /// Ugt1a5 /// Ugt1a9 /// Ugt1a6b /// Ugt1a1
 1424784_at 1.08670 RIKEN cDNA 1700029I01 gene /// similar to zinc finger protein 665 /// similar
 to zinc finger protein 665 /// similar to zinc finger protein 665 1700029I01Rik /// LOC433791 ///
 LOC666532 /// LOC671566
 1424785_at 0.98958 angiopoietin-like 6 Angptl6
 1424786_s_at 0.90290 WD repeat domain 45 Wdr45
 1424787_a_at 1.02591 nuclear respiratory factor 1 Nrf1
 1424788_at 1.24612 cDNA sequence BC021438 BC021438
 1424789_at 0.80796 GLI-Kruppel family member HKR2 Hkr2
 1424790_at 0.99953 RIKEN cDNA 2900084M01 gene 2900084M01Rik
 1424791_a_at 0.84892 basal cell adhesion molecule Bcam
 1424792_at 1.11195 ribonuclease P 40 subunit (human) Rpp40
 1424793_a_at 1.08186 phosphatidylethanolamine binding protein 2 Pbp2
 1424794_at 1.07659 ring finger protein 186 Rnf186
 1424795_a_at 0.92410 RIKEN cDNA 1700001O22 gene 1700001O22Rik
 1424796_at 0.94179 RIKEN cDNA 1700054N08 gene 1700054N08Rik
 1424797_a_at 0.74461 paired-like homeodomain transcription factor 2 Pitx2
 1424798_a_at 0.93482 a disintegrin and metallopeptidase domain 5 Adam5
 1424799_a_at 1.00646 golgi coiled coil 1 Gcc1
 1424800_at 0.84937 enabled homolog (Drosophila) Enah
 1424801_at 0.86649 enabled homolog (Drosophila) Enah
 1424802_a_at 0.97219 RIKEN cDNA 3300001P08 gene 3300001P08Rik
 1424803_at 1.00521 cDNA sequence BC020002 BC020002
 1424804_at 0.95916 cDNA sequence BC020002 BC020002
 1424805_a_at 0.89697 RIKEN cDNA 1110039B18 gene 1110039B18Rik
 1424806_s_at 0.95274 RIKEN cDNA 1110039B18 gene 1110039B18Rik
 1424807_at 0.90175 "laminin, alpha 4" Lama4
 1424808_at 0.92657 "laminin, alpha 4" Lama4
 1424809_at 0.95378 crumbs homolog 3 (Drosophila) Crb3
 1424810_at 1.09394 "taspase, threonine aspartase 1" Tasp1
 1424811_at 0.98319 camello-like 5 Cml5
 1424812_at 0.90142 cDNA sequence BC017158 BC017158
 1424813_at 1.06896 expressed sequence AA536717 AA536717
 1424814_a_at 0.91482 Bcl2-like 14 (apoptosis facilitator) Bcl2l14
 1424815_at 0.99753 glycogen synthase 2 /// similar to glycogen synthase 2 Gys2 /// LOC633620
 1424816_at 0.93997 "cat eye syndrome chromosome region, candidate 5 homolog (human)"
 Cocr5

1424817_at	0.81466	sperm flagellar 1	Spef1	
1424818_at	0.95734	"asparagine-linked glycosylation 12 homolog (yeast, alpha-1,6-mannosyltransferase)"	Alg12	
1424819_a_at	1.08563	resistance to inhibitors of cholinesterase 8 homolog (C. elegans)		Ric8
1424820_a_at	0.79016	Nedd4 family interacting protein 1	Ndfip1	
1424821_at	0.83731	Nedd4 family interacting protein 1	Ndfip1	
1424822_at	1.01443	RIKEN cDNA 9630044O09 gene	9630044O09Rik	
1424823_s_at	1.04526	RIKEN cDNA 9630044O09 gene	9630044O09Rik	
1424824_at	0.97632	RIKEN cDNA 9630044O09 gene	9630044O09Rik	
1424825_a_at	1.08121	glycosylation dependent cell adhesion molecule 1		Glycam1
1424826_s_at	0.92496	metastasis suppressor 1	Mtss1	
1424827_a_at	0.90525	"casein kinase 1, alpha 1"	Csnk1a1	
1424828_a_at	1.05624	fumarate hydratase 1	Fh1	
1424829_at	1.07993	RIKEN cDNA A830007P12 gene	A830007P12Rik	
1424830_at	1.03673	cyclin K	Ccnk	
1424831_at	1.04265	copine II	Cpne2	
1424832_at	1.01598	RIKEN cDNA 4732429D16 gene	4732429D16Rik	
1424833_at	0.80385	"inositol 1,4,5-triphosphate receptor 2"	Itpr2	
1424834_s_at	0.90164	"inositol 1,4,5-triphosphate receptor 2"	Itpr2	
1424835_at	1.05674	"glutathione S-transferase, mu 4"	Gstm4	
1424836_a_at	1.01669	CLIP associating protein 2	Clasp2	
1424837_at	0.98981	ring finger protein 113A1	Rnf113a1	
1424838_at	1.03780	RIKEN cDNA A330049M08 gene	A330049M08Rik	
1424839_a_at	0.98145	"NOL1/NOP2/Sun domain family, member 4"	Nsun4	
1424840_at	0.92152	ribokinase	Rbks	
1424841_s_at	0.88650	ribokinase	Rbks	
1424842_a_at	0.63184	Rho GTPase activating protein 24		Arhgap24
1424843_a_at	1.26431	growth arrest specific 5	Gas5	
1424844_at	0.92381	angel homolog 1 (Drosophila)	Angel1	
1424845_a_at	1.11214	centrosomal protein 68	Cep68	
1424846_at	0.85507	centrosomal protein 68	Cep68	
1424847_at	1.06779	"neurofilament, heavy polypeptide"	Nefh	
1424848_at	0.92359	"potassium large conductance calcium-activated channel, subfamily M, alpha member 1"	Kcnma1	
1424849_at	1.14397	WD repeat domain 62	Wdr62	
1424850_at	0.89511	mitogen activated protein kinase kinase kinase 1 /// similar to Mitogen-activated protein kinase kinase kinase 1 (MAPK/ERK kinase kinase 1) (MEK kinase 1) (MEKK 1) /// similar to Mitogen-activated protein kinase kinase kinase 1 (MAPK/ERK kinase kinase 1) (MEK kinase 1) (MEKK 1)	Map3k1 /// LOC670493 /// LOC674573	
1424851_at	0.89585	coiled-coil-helix-coiled-coil-helix domain containing 5	Chchd5	
1424852_at	1.10565	myocyte enhancer factor 2C	Mef2c	
1424853_s_at	0.96012	"cytochrome P450, family 4, subfamily a, polypeptide 10 /// cDNA sequence BC013476"	Cyp4a10 /// BC013476	

1424854_at 1.16013 "histone 1, H4i" Hist1h4i
1424855_at 0.92877 oleoyl-ACP hydrolase Olah
1424856_at 1.06136 "ATPase, Na+/K+ transporting, alpha 3 polypeptide" Atp1a3
1424857_a_at 0.85848 tripartite motif protein 34 /// similar to Tripartite motif protein 34 Trim34 ///
LOC434218
1424858_at 1.17759 L-2-hydroxyglutarate dehydrogenase L2hgdh
1424859_at 1.01767 homer homolog 3 (Drosophila) Homer3
1424860_at 0.94400 RIKEN cDNA D930016D06 gene D930016D06Rik
1424861_at 1.03807 RIKEN cDNA D930016D06 gene D930016D06Rik
1424862_s_at 0.99503 mindbomb homolog 2 (Drosophila) Mib2
1424863_a_at 0.75102 homeodomain interacting protein kinase 2 Hipk2
1424864_at 0.84302 homeodomain interacting protein kinase 2 Hipk2
1424865_at 0.91784 peptide YY Pyy
1424866_at 0.96343 ubiquitin specific peptidase 43 Usp43
1424867_a_at 0.83708 glycine-N-acyltransferase Glyat
1424868_at 1.07071 glycine-N-acyltransferase Glyat
1424869_at 1.00150 dehydrogenase/reductase (SDR family) member 7B Dhhrs7b
1424870_at 1.05623 oxysterol binding protein-like 10 Osbpl10
1424871_s_at 0.98851 RIKEN cDNA 1500031H01 gene 1500031H01Rik
1424872_at 1.00499 RIKEN cDNA 2310001H12 gene /// similar to zinc finger protein 709 /// similar
to zinc finger protein 665 2310001H12Rik /// LOC636362 /// LOC666849
1424873_at 0.94431 ring finger protein 2 Rnf2
1424874_a_at 1.07374 polypyrimidine tract binding protein 1 Ptbp1
1424875_at 1.06064 "spastic paraplegia 20, spartin (Troyer syndrome) homolog (human)" Spg20
1424876_s_at 1.10428 "spastic paraplegia 20, spartin (Troyer syndrome) homolog (human)" Spg20
1424877_a_at 0.93691 "aminolevulinate, delta-, dehydratase" Alad
1424878_at 1.02135 leucine-rich repeats and calponin homology (CH) domain containing 4 Lrch4
1424879_at 0.90148 leucine-rich repeats and calponin homology (CH) domain containing 4 Lrch4
1424880_at 0.93672 tribbles homolog 1 (Drosophila) Trib1
1424881_at 1.06517 tribbles homolog 1 (Drosophila) Trib1
1424882_a_at 0.78623 5'-nucleotidase domain containing 2 Nt5dc2
1424883_s_at 1.07197 "splicing factor, arginine/serine-rich 7" Sfrs7
1424884_at 1.04771 F-box and WD-40 domain protein 2 Fbxw2
1424885_at 0.76188 kelch domain containing 8A Klhdc8a
1424886_at 0.71530 "protein tyrosine phosphatase, receptor type, D" Ptpkd
1424887_at 1.07327 kelch domain containing 4 Klhdc4
1424888_at 0.86052 membrane-associated ring finger (C3HC4) 2 2-Mar
1424889_at 0.98843 nucleoporin like 2 Nupl2
1424890_at 0.92659 basenuclin 1 Bnc1
1424891_a_at 1.03137 "ZW10 homolog (Drosophila), centromere/kinetochore protein" Zw10
1424892_at 0.88023 zinc finger protein 95 Zfp95
1424893_at 2.19071 nuclear distribution gene E-like homolog 1 (A. nidulans) Ndel1
1424894_at 0.98055 "RAB13, member RAS oncogene family" Rab13

1424895_at 1.02023 "G-protein signalling modulator 2 (AGS3-like, *C. elegans*)" Gpsm2
1424896_at 0.84096 G protein-coupled receptor 85 Gpr85
1424897_at 1.11727 G protein-coupled receptor 85 Gpr85
1424898_at 1.07516 "solute carrier family 10 (sodium/bile acid cotransporter family), member 1" Slc10a1
1424899_at 1.08488 nicotinamide nucleotide adenylyltransferase 3 Nmnat3
1424900_at 0.89798 "solute carrier family 29 (nucleoside transporters), member 4" Slc29a4
1424901_at 1.03916 "glucosaminyl (N-acetyl) transferase 3, mucin type" Gcnt3
1424902_at 0.81619 plexin domain containing 1 Plxdc1
1424903_at 0.95807 "jumonji, AT rich interactive domain 1D (Rbp2 like)" Jarid1d
1424904_at 0.91867 RIKEN cDNA 1300010F03 gene 1300010F03Rik
1424905_a_at 0.98540 "solute carrier family 39 (metal ion transporter), member 11" Slc39a11
1424906_at 1.02529 PQ loop repeat containing Pqlc3
1424907_a_at 1.04342 "phenylalanine-tRNA synthetase-like, alpha subunit" Farsla
1424908_at 1.00397 mitochondrial methionyl-tRNA formyltransferase Mtfmt
1424909_at 0.96893 leucine rich repeat containing 46 Lrrc46
1424910_at 1.11945 kinesin family member 12 Kif12
1424911_a_at 1.06616 lysozyme-like 4 Lyz14
1424912_at 0.98613 "solute carrier family 25 (mitochondrial carrier, peroxisomal membrane protein), member 17" Slc25a17
1424913_at 0.99960 RIKEN cDNA 2310044G17 gene 2310044G17Rik
1424914_at 0.97607 --- ---
1424915_s_at 1.03594 --- ---
1424916_x_at 1.04723 RIKEN cDNA 8030466O12 gene 8030466O12Rik
1424917_a_at 0.90842 "WD repeat domain, phosphoinositide interacting 1" Wipi1
1424918_at 0.81822 "TBC1 domain family, member 19" Tbc1d19
1424919_at 0.92069 "v-erb-b2 erythroblastic leukemia viral oncogene homolog 2, neuro/glioblastoma derived oncogene homolog (avian)" Erbb2
1424920_at 0.96336 "solute carrier family 37 (glycerol-3-phosphate transporter), member 3" Slc37a3
1424921_at 1.45613 bromodomain containing 4 /// bone marrow stromal cell antigen 2 Brd4 /// Bst2
1424922_a_at 1.24310 bromodomain containing 4 Brd4
1424923_at 2.95804 "serine (or cysteine) peptidase inhibitor, clade A, member 3G" Serpina3g
1424924_at 1.20635 SEC63-like (*S. cerevisiae*) Sec63
1424925_at 1.02342 SEC63-like (*S. cerevisiae*) Sec63
1424926_at 1.14756 SEC63-like (*S. cerevisiae*) Sec63
1424927_at 0.89142 GLI pathogenesis-related 1 (glioma) Glipr1
1424928_at 1.12488 RIKEN cDNA 2210018M11 gene 2210018M11Rik
1424929_a_at 1.01091 tripartite motif protein 26 Trim26
1424930_s_at 0.95160 expressed sequence AW544981 AW544981
1424931_s_at 1.08003 "immunoglobulin lambda chain, variable 1" Igl-V1
1424932_at 0.96316 epidermal growth factor receptor Egfr
1424933_at 1.03583 myosin VC Myo5c
1424934_at 0.90805 "UDP glucuronosyltransferase 2 family, polypeptide B1" Ugt2b1

1424935_at 1.02091 ganglioside-induced differentiation-associated protein 1-like 1 Gdap1l1
1424936_a_at 0.99915 "dynein, axonemal, heavy chain 8" Dnahc8
1424937_at 1.08024 RIKEN cDNA 2310076L09 gene 2310076L09Rik
1424938_at 1.23377 six transmembrane epithelial antigen of the prostate 1 Steap1
1424939_at 0.97640 "ankyrin repeat, SAM and basic leucine zipper domain containing 1" Asz1
1424940_s_at 1.00238 cDNA sequence BC022687 BC022687
1424941_at 1.04781 "cDNA sequence, BC016579" BC016579
1424942_a_at 1.17907 myelocytomatosis oncogene Myc
1424943_at 0.88785 cDNA sequence BC013476 BC013476
1424944_at 0.93988 Purkinje cell protein 2 (L7) Pcp2
1424945_at 1.07266 chordin-like 1 Chrdl1
1424946_a_at 1.02246 mitochondrial fission regulator 1 Mtfr1
1424947_at 1.08459 dynein cytoplasmic 1 light intermediate chain 1 Dync1li1
1424948_x_at 1.31213 "histocompatibility 2, D region locus 1 /// histocompatibility 2, K1, K region ///
MHC (A.CA/J(H-2K-f) class I antigen" H2-D1 /// H2-K1 /// LOC56628
1424949_at 0.95788 "HECT, UBA and WWE domain containing 1" Huwe1
1424950_at 0.97270 SRY-box containing gene 9 Sox9
1424951_at 0.75933 BAI1-associated protein 2-like 1 Baiap2l1
1424952_at 0.91194 OCIA domain containing 1 Ociad1
1424953_at 1.12593 cDNA sequence BC021614 BC021614
1424954_a_at 1.00259 --- ---
1424955_at 1.12520 coiled-coil domain containing 5 Ccdc5
1424956_at 0.99856 "AT hook, DNA binding motif, containing 1" Ahdc1
1424957_at 0.92966 "AT hook, DNA binding motif, containing 1" Ahdc1
1424958_at 1.06884 carbonic anhydrase 8 Car8
1424959_at 1.00215 annexin A13 Anxa13
1424960_at 0.89499 epsin 3 Epn3
1424961_at 1.08677 TAP binding protein-like Tapbpl
1424962_at 0.84273 transmembrane 4 superfamily member 4 Tm4sf4
1424963_at 1.00923 retinitis pigmentosa 1 homolog (human) Rp1h
1424964_at 0.91939 retinitis pigmentosa 1 homolog (human) Rp1h
1424965_at 1.00130 leupaxin Lpxn
1424966_at 1.12467 transmembrane protein 40 Tmem40
1424967_x_at 0.82839 "troponin T2, cardiac" Tnnt2
1424968_at 0.95667 RIKEN cDNA 2210023G05 gene 2210023G05Rik
1424969_s_at 0.97718 --- ---
1424970_at 0.74832 purine-rich element binding protein G Purg
1424971_at 0.96845 coiled-coil domain containing 99 Ccdc99
1424972_at 1.07636 --- ---
1424973_at 0.94542 "cytochrome P450, family 3, subfamily a, polypeptide 25 /// similar to
cytochrome P450, family 3, subfamily a, polypeptide 25" Cyp3a25 /// LOC622249
1424974_at 1.07369 zinc finger protein 418 Zfp418
1424975_at 0.98173 sialic acid binding Ig-like lectin F Siglecf

1424976_at 1.10706 "ras homolog gene family, member V" Rhov
1424977_at 1.01062 RIKEN cDNA 4930418G15 gene 4930418G15Rik
1424978_at 0.99937 outer dense fiber of sperm tails 4 Odf4
1424979_at 1.11204 anterior pharynx defective 1a homolog (C. elegans) Aph1a
1424980_s_at 0.94515 anterior pharynx defective 1a homolog (C. elegans) Aph1a
1424981_at 1.29216 neurolysin (metallopeptidase M3 family) Nln
1424982_a_at 1.04744 RIKEN cDNA 2700078E11 gene 2700078E11Rik
1424983_a_at 1.00870 RIKEN cDNA 2700078E11 gene 2700078E11Rik
1424984_at 0.99685 RIKEN cDNA 2700078E11 gene 2700078E11Rik
1424985_a_at 1.03952 SRY-box containing gene 10 Sox10
1424986_s_at 1.09959 "F-box and WD-40 domain protein 7, archipelago homolog (Drosophila)"
Fbxw7
1424987_at 1.01457 RIKEN cDNA 5430435G22 gene 5430435G22Rik
1424988_at 1.01727 myosin regulatory light chain interacting protein Mylip
1424989_at 1.01067 transmembrane protein 142A Tmem142a
1424990_at 1.07709 transmembrane protein 142A Tmem142a
1424991_s_at 1.01613 "thymidylate synthase /// thymidylate synthase, pseudogene" Tyms /// Tyms-
ps
1424992_at 1.22854 Tp53rk binding protein Tprkb
1424993_at 0.95589 RIKEN cDNA 8430410K20 gene 8430410K20Rik
1424994_at 1.06019 RIKEN cDNA 6230410P16 gene 6230410P16Rik
1424995_at 0.94176 RIKEN cDNA 6230410P16 gene 6230410P16Rik
1424996_at 1.04763 CASP8 and FADD-like apoptosis regulator Cflar
1424997_at 0.98614 "splicing factor, arginine/serine-rich 8" Sfrs8
1424998_at 0.93592 "EGF-like module containing, mucin-like, hormone receptor-like sequence
4" Emr4
1424999_at 1.17632 RIKEN cDNA 1700022C21 gene 1700022C21Rik
1425000_s_at 1.03917 RIKEN cDNA 5430407P10 gene 5430407P10Rik
1425001_at 0.72631 ring finger protein 146 Rnf146
1425002_at 0.99027 cDNA sequence BC010462 BC010462
1425003_at 0.93118 urocanase domain containing 1 Uroc1
1425004_s_at 1.00647 molybdenum cofactor synthesis 1 Mocs1
1425005_at 0.88941 "killer cell lectin-like receptor subfamily C, member 1 /// killer cell lectin-like
receptor subfamily C, member 2" Klrc1 /// Klrc2
1425006_a_at 1.08615 vaccinia related kinase 1 Vrk1
1425007_at 1.12512 zinc finger protein 566 Zfp566
1425008_a_at 1.04506 interferon activated gene 203 Ifi203
1425009_at 1.09893 t-complex-associated testis expressed 2 Tcte2
1425010_at 1.07314 zinc finger protein 119 Zfp119
1425011_x_at 0.96544 syntaxin 18 Stx18
1425012_at 0.94918 "guanine nucleotide binding protein (G protein), gamma 5 subunit ///
hypothetical protein LOC668891 /// similar to guanine nucleotide binding protein (G protein), gamma 5
subunit" Gng5 /// LOC668891 /// LOC671715

1425013_at 1.02294 "solute carrier family 25 (mitochondrial carrier, ornithine transporter) member 2" Slc25a2
 1425014_at 1.00905 "nuclear receptor subfamily 2, group C, member 2" Nr2c2
 1425015_at 0.78073 Eph receptor B2 Ephb2
 1425016_at 0.72821 Eph receptor B2 Ephb2
 1425017_at 0.97425 "protocadherin alpha subfamily C, 1" Pcdhac1
 1425018_at 1.15542 malignant T cell amplified sequence 1 Mcts1
 1425019_at 0.93984 UBX domain containing 4 Ubx4
 1425020_at 1.00234 UBX domain containing 4 /// similar to UBX domain containing 4 /// similar to UBX domain containing 4 Ubx4 /// LOC621629 /// LOC637151
 1425021_a_at 0.86701 peroxisome biogenesis factor 16 Pex16
 1425022_at 0.96722 ubiquitin specific peptidase 3 Usp3
 1425023_at 0.91116 ubiquitin specific peptidase 3 Usp3
 1425024_at 0.81466 RIKEN cDNA E430018J23 gene E430018J23Rik
 1425025_at 1.03160 transmembrane protein 106A Tmem106a
 1425026_at 0.97261 SFT2 domain containing 2 Sft2d2
 1425027_s_at 1.09768 SFT2 domain containing 2 Sft2d2
 1425028_a_at 0.81754 "tropomyosin 2, beta" Tpm2
 1425029_a_at 0.94286 membrane bound O-acyltransferase domain containing 2 Mboat2
 1425030_at 1.06069 zinc finger protein 622 Zfp622
 1425031_at 0.96030 Fukuyama type congenital muscular dystrophy homolog (human) Fcmd
 1425032_at 1.05596 androgen binding protein beta Abpb
 1425033_at 1.01291 "solute carrier family 17 (sodium phosphate), member 2" Slc17a2
 1425034_at 1.15649 "solute carrier family 17 (sodium phosphate), member 2" Slc17a2
 1425035_s_at 0.98765 DNA (cytosine-5-)-methyltransferase 3-like Dnmt3l
 1425036_a_at 0.93458 trinucleotide repeat containing 6a Tnrc6a
 1425037_at 1.06485 "FYVE, RhoGEF and PH domain containing 4" Fgd4
 1425038_at 0.98242 "solute carrier family 22 (organic anion transporter), member 19" Slc22a19
 1425039_at 0.99955 "integrin, beta-like 1" Itgb1
 1425040_at 1.09319 cytochrome b reductase 1 Cybrd1
 1425041_at 0.94521 LIM homeobox protein 3 Lhx3
 1425042_s_at 0.99715 "proline, glutamic acid and leucine rich protein 1" Pelp1
 1425043_s_at 1.03671 RIKEN cDNA 0610037D15 gene 0610037D15Rik
 1425044_at 0.93496 "potassium inwardly-rectifying channel, subfamily J, member 6" Kcnj6
 1425045_at 1.02074 "phospholipase A2, group IVB (cytosolic)" Pla2g4b
 1425046_at 1.05302 cDNA sequence BC018465 BC018465
 1425047_a_at 0.90092 RIKEN cDNA 4921517L17 gene 4921517L17Rik
 1425048_a_at 0.99035 high mobility group box 1 Hmgb1
 1425049_at 0.82819 2'-5' oligoadenylate synthetase 1H Oas1h
 1425050_at 0.90135 isochorismatase domain containing 1 Isoc1
 1425051_at 0.92208 isochorismatase domain containing 1 Isoc1
 1425052_at 0.84286 "CDNA, clone:Y2G0147N17, strand:unspecified" ---
 1425053_at 0.93030 isochorismatase domain containing 1 Isoc1

1425054_at 1.07997 RIKEN cDNA 2510006D16 gene 2510006D16Rik
 1425055_at 0.96602 "SAPS domain family, member 2" Saps2
 1425056_s_at 1.07387 "SAPS domain family, member 2" Saps2
 1425057_at 1.00319 RIKEN cDNA 0610038K03 gene 0610038K03Rik
 1425058_at 1.06498 zinc finger protein 472 Zfp472
 1425059_at 1.01736 protein arginine N-methyltransferase 6 Prmt6
 1425060_s_at 0.84731 restin (Reed-Steinberg cell-expressed intermediate filament-associated protein)
 Rsn
 1425061_at 0.90803 "WAS protein family, member 3" Wasf3
 1425062_at 1.20278 Fc receptor-like 1 Fcrl1
 1425063_at 1.06228 --- ---
 1425064_at 1.08480 aryl hydrocarbon receptor nuclear translocator Arnt
 1425065_at 0.92351 2'-5' oligoadenylate synthetase 2 Oas2
 1425066_a_at 0.97416 RIKEN cDNA 1110061O04 gene 1110061O04Rik
 1425067_at 1.05142 cadherin EGF LAG seven-pass G-type receptor 3 Celsr3
 1425068_a_at 0.99666 testis expressed gene 264 Tex264
 1425069_at 0.97180 cDNA sequence BC018285 BC018285
 1425070_at 0.97013 "neurotrophic tyrosine kinase, receptor, type 3" Ntrk3
 1425071_s_at 0.91554 "neurotrophic tyrosine kinase, receptor, type 3" Ntrk3
 1425072_at 1.66945 S-phase kinase-associated protein 2 (p45) Skp2
 1425073_at 0.99486 "pleckstrin homology domain containing, family G (with RhoGef domain)
 member 6" Plekhg6
 1425074_at 0.89992 Werner syndrome homolog (human) Wrn
 1425075_at 0.90644 GATA zinc finger domain containing 2B Gatad2b
 1425076_at 1.01303 "DnaJ (Hsp40) homolog, subfamily C, member 18" Dnajc18
 1425077_at 0.84726 "DnaJ (Hsp40) homolog, subfamily C, member 18" Dnajc18
 1425078_x_at 0.95607 similar to SP140 nuclear body protein (predicted) LOC620078
 1425079_at 1.08373 transmembrane 6 superfamily member 2 Tm6sf2
 1425080_at 0.95364 zinc finger protein 286 Zfp286
 1425081_at 1.12130 zinc finger protein 286 Zfp286
 1425082_s_at 1.17140 RIKEN cDNA 9030416H16 gene 9030416H16Rik
 1425083_at 1.00653 otoraplin Otor
 1425084_at 1.12905 "GTPase, IMAP family member 7" Gimap7
 1425085_at 1.02864 RIKEN cDNA 6330416L07 gene 6330416L07Rik
 1425086_a_at 1.00210 SLAM family member 6 Slamf6
 1425087_at 1.06543 RIKEN cDNA 2310003F16 gene 2310003F16Rik
 1425088_at 0.94108 "sodium channel, nonvoltage-gated, type I, alpha" Scnn1a
 1425089_at 1.09481 "potassium voltage gated channel, Shaw-related subfamily, member 4" Kcnc4
 1425090_s_at 0.91292 "potassium voltage gated channel, Shaw-related subfamily, member 4"
 Kcnc4
 1425091_at 0.98318 retinoic acid receptor responder (tazarotene induced) 2 Rarres2
 1425092_at 1.09979 cadherin 10 Cdh10
 1425093_at 1.01512 "purinergic receptor P2X, ligand-gated ion channel, 3" P2rx3

1425094_a_at 0.77368 LIM homeobox protein 6 Lhx6
1425095_at 1.08236 cDNA sequence BC002059 BC002059
1425096_a_at 0.87786 pentatricopeptide repeat domain 1 Ptc1
1425097_a_at 1.14778 zinc finger protein 106 Zfp106
1425098_at 0.95067 zinc finger protein 106 Zfp106
1425099_a_at 0.90532 aryl hydrocarbon receptor nuclear translocator-like Arntl
1425100_a_at 0.88687 "phosphodiesterase 6G, cGMP-specific, rod, gamma" Pde6g
1425101_a_at 1.00817 FK506 binding protein 6 Fkbp6
1425102_a_at 0.24823 angiotensin I converting enzyme (peptidyl-dipeptidase A) 2 Ace2
1425103_at 0.48144 angiotensin I converting enzyme (peptidyl-dipeptidase A) 2 Ace2
1425104_at 0.90644 potassium channel tetramerisation domain containing 1 Kctd1
1425105_at 0.96949 "retinol binding protein 3, interstitial" Rbp3
1425106_a_at 1.06703 tryptophanyl-tRNA synthetase Wars
1425107_a_at 0.86648 leukemia inhibitory factor receptor Lifr
1425108_a_at 0.88476 cDNA sequence BC004728 BC004728
1425109_at 0.86483 "solute carrier family 44, member 3" Slc44a3
1425110_at 0.95379 sortilin-related VPS10 domain containing receptor 3 Sorcs3
1425111_at 1.02357 sortilin-related VPS10 domain containing receptor 3 Sorcs3
1425112_at 1.07376 ankyrin repeat domain 27 (VPS9 domain) Ankrd27
1425113_x_at 1.03512 "PREDICTED: Mus musculus hypothetical protein LOC621686
(LOC621686), mRNA" ---
1425114_at 0.81203 retinoblastoma binding protein 6 Rbbp6
1425115_at 0.73288 retinoblastoma binding protein 6 Rbbp6
1425116_a_at 1.03625 spectrin beta 4 Spnb4
1425117_at 0.96516 RIKEN cDNA 0610012D14 gene 0610012D14Rik
1425118_at 1.04945 spire homolog 2 (Drosophila) Spire2
1425119_at 0.96167 --- ---
1425120_x_at 0.90013 RIKEN cDNA 1810023F06 gene 1810023F06Rik
1425121_a_at 0.92349 cDNA sequence BC006965 BC006965
1425122_at 0.93517 open reading frame 9 ORF9
1425123_at 0.91380 cDNA sequence BC025816 BC025816
1425124_at 1.02499 ring finger protein 183 Rnf183
1425125_at 1.30895 oncoprotein induced transcript 3 Oit3
1425126_at 0.89666 neural cell adhesion molecule 1 Ncam1
1425127_at 0.91534 "hydroxy-delta-5-steroid dehydrogenase, 3 beta- and steroid delta-isomerase 2"
Hsd3b2
1425128_at 0.84021 "UDP-GlcNAc:betaGal beta-1,3-N-acetylglucosaminyltransferase 8" B3gnt8
1425129_a_at 1.06735 transaldolase 1 Taldo1
1425130_a_at 0.98309 "protein tyrosine phosphatase, non-receptor type 5" Ptpn5
1425131_at 0.97973 "protein tyrosine phosphatase, non-receptor type 5" Ptpn5
1425132_at 0.92464 neuropilin (NRP) and tolloid (TLL)-like 1 Neto1
1425133_s_at 0.75904 RAB3A interacting protein (rabin3)-like 1 Rab3il1
1425134_a_at 0.96695 "phosphatidylinositol glycan anchor biosynthesis, class X" Pigx

1425135_a_at	0.89119	dynamamin 2	Dnm2	
1425136_x_at	0.96665	dynamamin 2	Dnm2	
1425137_a_at	1.01654	"histocompatibility 2, Q region locus 10"	H2-Q10	
1425138_at	1.07305	guanylate cyclase activator 1B	Guca1b	
1425139_at	1.00232	sestrin 2	Sesn2	
1425140_at	1.23245	"lactamase, beta 2"	Lactb2	
1425141_at	1.23539	"lactamase, beta 2"	Lactb2	
1425142_a_at	0.99707	heterogeneous nuclear ribonucleoprotein D	Hnrpd	
1425143_a_at	0.98722	NADH dehydrogenase (ubiquinone) Fe-S protein 1	Ndufs1	
1425144_at	1.02377	kallikrein 1-related peptidase b11	Klk1b11	
1425145_at	0.97007	interleukin 1 receptor-like 1	Il1rl1	
1425146_at	0.87830	RIKEN cDNA 2410075B13 gene	2410075B13Rik	
1425147_at	1.00334	RIKEN cDNA 2410075B13 gene	2410075B13Rik	
1425148_a_at	0.93937	sorting nexin 6	Snx6	
1425149_a_at	0.85610	phosducin-like	Pdcl	
1425150_at	1.00089	RIKEN cDNA C730036D15 gene	C730036D15Rik	
1425151_a_at	1.02360	NADPH oxidase organizer 1	Noxo1	
1425152_s_at	0.89897	"serine (or cysteine) peptidase inhibitor, clade I, member 2"	Serpini2	
1425153_at	1.01912	"myosin, heavy polypeptide 2, skeletal muscle, adult"	Myh2	
1425154_a_at	1.18408	colony stimulating factor 1 (macrophage)	Csf1	
1425155_x_at	1.32692	colony stimulating factor 1 (macrophage)	Csf1	
1425156_at	1.01325	guanylate binding protein 7	Gbp7	
1425157_x_at	1.03950	tetraspanin 33	Tspan33	
1425158_at	0.97960	T-box 20	Tbx20	
1425159_at	0.93112	golgi transport 1 homolog A (S. cerevisiae)	Golt1a	
1425160_at	1.00883	cDNA sequence AF067063	AF067063	
1425161_a_at	0.99193	TraB domain containing	Trabd	
1425162_at	0.85592	RAR-related orphan receptor beta	Rorb	
1425163_at	0.97659	expressed sequence AI661453	AI661453	
1425164_a_at	0.95267	phosphorylase kinase gamma 1	Phkg1	
1425165_at	1.01619	granzyme N	Gzmn	
1425166_at	1.03740	retinoblastoma-like 1 (p107)	Rbl1	
1425167_a_at	0.89628	"guanine nucleotide binding protein (G protein), gamma transducing activity polypeptide 1"	Gngt1	
1425168_at	0.98186	"guanine nucleotide binding protein (G protein), gamma transducing activity polypeptide 1"	Gngt1	
1425169_at	0.95266	testis serine protease 2	Tessp2	
1425170_a_at	0.65641	a disintegrin and metallopeptidase domain 15 (metargidin)	Adam15	
1425171_at	0.91103	rhodopsin	Rho	
1425172_at	1.05020	rhodopsin	Rho	
1425173_s_at	0.77565	golgi phosphoprotein 3-like	Golph3l	
1425174_at	0.92080	CDNA clone MGC:18942 IMAGE:3979577	---	
1425175_at	0.89091	C1q-like 3	C1ql3	

1425176_at 0.95341 C1q-like 3 C1ql3
1425177_at 1.04786 serine hydroxymethyl transferase 1 (soluble) Shmt1
1425178_s_at 0.98456 serine hydroxymethyl transferase 1 (soluble) Shmt1
1425179_at 1.09882 serine hydroxymethyl transferase 1 (soluble) Shmt1
1425180_at 1.01753 SH3-domain GRB2-like (endophilin) interacting protein 1 Sgip1
1425181_at 0.97132 SH3-domain GRB2-like (endophilin) interacting protein 1 Sgip1
1425182_x_at 0.91390 kallikrein 1-related peptidase b22 /// kallikrein 1-related peptidase b9
Klk1b22 /// Klk1b9
1425183_a_at 1.03849 ribosomal protein L4 Rpl4
1425184_at 0.98446 RIKEN cDNA 2810433K01 gene 2810433K01Rik
1425185_at 1.08765 RIKEN cDNA 5830417C01 gene 5830417C01Rik
1425186_at 1.04832 LMBR1 domain containing 1 Lmbrd1
1425187_at 0.94488 Sel1 (suppressor of lin-12) 1 homolog (C. elegans) Sel1h
1425188_s_at 1.02084 Sel1 (suppressor of lin-12) 1 homolog (C. elegans) Sel1h
1425189_a_at 1.03274 mitochondrial ribosomal protein L15 Mrpl15
1425190_a_at 1.01035 "phosphatase, orphan 2" Phospho2
1425191_at 0.87938 RIKEN cDNA 9430098E02 gene 9430098E02Rik
1425192_at 0.98762 kelch-like 25 (Drosophila) Klhl25
1425193_at 0.96524 RIKEN cDNA 2010106G01 gene 2010106G01Rik
1425194_a_at 1.06113 RIKEN cDNA 6330577E15 gene 6330577E15Rik
1425195_a_at 1.55069 acetyl-Coenzyme A acetyltransferase 2 /// acetyl-Coenzyme A
acetyltransferase 3 Acat2 /// Acat3
1425196_a_at 0.95949 histidine triad nucleotide binding protein 2 Hint2
1425197_at 0.84465 "protein tyrosine phosphatase, non-receptor type 2" Ptpn2
1425198_at 1.01510 "protein tyrosine phosphatase, non-receptor type 2" Ptpn2
1425199_a_at 1.00894 erythrocyte protein band 4.1-like 5 Epb4.115
1425200_at 1.07335 chloride channel CLIC-like 1 Clcc1
1425201_a_at 1.16545 hydroxypyruvate isomerase homolog (E. coli) Hyi
1425202_a_at 0.88506 "ankyrin 3, epithelial" Ank3
1425203_at 1.23061 DEAD (Asp-Glu-Ala-Asp) box polypeptide 19b Ddx19b
1425204_s_at 1.02082 DEAD (Asp-Glu-Ala-Asp) box polypeptide 19a /// DEAD (Asp-Glu-Ala-Asp)
box polypeptide 19b Ddx19a /// Ddx19b
1425205_at 0.83929 DEAD (Asp-Glu-Ala-Asp) box polypeptide 19b Ddx19b
1425206_a_at 0.81685 Ubiquitin protein ligase E3A Ube3a
1425207_at 0.97861 cDNA sequence BC026439 BC026439
1425208_at 0.85527 limb-bud and heart Lbh
1425209_at 0.96178 zinc finger protein 84 Zfp84
1425210_s_at 1.07681 zinc finger protein 84 Zfp84
1425211_at 1.04517 RIKEN cDNA 2010003O18 gene 2010003O18Rik
1425212_a_at 1.02476 "tumor necrosis factor receptor superfamily, member 19" Tnfrsf19
1425213_at 0.59292 RIKEN cDNA 6430514L14 gene 6430514L14Rik
1425214_at 1.29307 "pyrimidinergic receptor P2Y, G-protein coupled, 6" P2ry6
1425215_at 1.14718 free fatty acid receptor 2 Ffar2

1425216_at 1.03422 free fatty acid receptor 2 Ffar2
 1425217_a_at 0.92787 synaptojanin 2 Synj2
 1425218_a_at 1.04110 "secretoglobin, family 3A, member 2" Scgb3a2
 1425219_x_at 1.15927 "guanine nucleotide binding protein (G protein), gamma transducing activity polypeptide 1" Gngt1
 1425220_x_at 0.99438 "cDNA sequence AF067061 /// hypothetical protein LOC639910 /// similar to 2-cell-stage, variable group, member 1" AF067061 /// LOC639910 /// LOC675060
 1425221_at 1.05908 RIKEN cDNA E030025L21 gene E030025L21Rik
 1425222_x_at 1.02543 RIKEN cDNA D630002G06 gene /// RIKEN cDNA C730048C13 gene /// similar to integral membrane transport protein UST5r D630002G06Rik /// C730048C13Rik /// LOC673838
 1425223_at 0.92567 baculoviral IAP repeat-containing 3 Birc3
 1425224_at 0.94511 cDNA sequence BC019561 BC019561
 1425225_at 0.93383 "Fc fragment of IgG, low affinity IIIa, receptor" Fcgr3a
 1425226_x_at 1.03245 "T-cell receptor beta, variable 13 /// similar to T-cell receptor beta-2 chain C region /// similar to T-cell receptor beta-1 chain C region /// similar to T-cell receptor beta-2 chain C region" Tcrb-V13 /// LOC665506 /// LOC669847 /// LOC674048
 1425227_a_at 1.18678 "ATPase, H+ transporting, lysosomal V0 subunit A1" Atp6v0a1
 1425228_a_at 0.95632 deoxyguanosine kinase Dguok
 1425229_a_at 0.95982 "transcription factor 7-like 2, T-cell specific, HMG-box" Tcf7l2
 1425230_at 1.04767 N-acetylglutamate synthase Nags
 1425231_a_at 0.89635 zinc finger protein 46 Zfp46
 1425232_x_at 0.91675 "arrestin 3, retinal" Arr3
 1425233_at 0.97047 RIKEN cDNA 2210407C18 gene 2210407C18Rik
 1425234_at 0.93132 "collagen, type XX, alpha 1" Col20a1
 1425235_s_at 0.99338 "collagen, type XX, alpha 1" Col20a1
 1425236_at 1.01190 "collagen, type XX, alpha 1" Col20a1
 1425237_at 0.93565 keratin associated protein 16-8 Krtap16-8
 1425238_at 1.13569 40S ribosomal protein S20 pseudogene LOC625603
 1425239_at 0.91661 SET domain containing 4 Setd4
 1425240_at 1.05320 cDNA sequence BC011426 BC011426
 1425241_a_at 0.98651 WD repeat and SOCS box-containing 1 Wsb1
 1425242_at 1.00197 RIKEN cDNA 1810006K21 gene 1810006K21Rik
 1425243_at 0.86480 CD 207 antigen Cd207
 1425244_a_at 0.87171 testicular haploid expressed gene Theg
 1425245_a_at 0.89004 regulator of G-protein signaling 11 Rgs11
 1425246_at 0.93912 RIKEN cDNA 0610008F07 gene 0610008F07Rik
 1425247_a_at 0.96591 immunoglobulin heavy chain 4 (serum IgG1) Igh-4
 1425248_a_at 0.80113 TYRO3 protein tyrosine kinase 3 Tyro3
 1425249_a_at 1.07162 TYRO3 protein tyrosine kinase 3 Tyro3
 1425250_a_at 0.90942 "solute carrier family 14 (urea transporter), member 2" Slc14a2
 1425251_at 0.93792 prostaglandin E receptor 3 (subtype EP3) Ptger3
 1425252_a_at 0.97759 mitotic arrest deficient 1-like 1 Mad11

1425253_a_at 1.12045 mucosal vascular addressin cell adhesion molecule 1 Madcam1
1425254_at 1.03354 forkhead box N4 Foxn4
1425255_s_at 1.01577 heterogeneous nuclear ribonucleoprotein L-like Hnrpll
1425256_a_at 0.95540 DIX domain containing 1 Dixdc1
1425257_at 0.99317 acyl-CoA thioesterase 5 Acot5
1425258_at 0.94402 cystatin 11 Cst11
1425259_x_at 1.02407 "killer cell lectin-like receptor subfamily A, member 12" Klra12
1425260_at 1.03312 albumin 1 Alb1
1425261_at 1.07135 "CCAAT/enhancer binding protein (C/EBP), gamma" Cebpg
1425262_at 1.12011 "CCAAT/enhancer binding protein (C/EBP), gamma" Cebpg
1425263_a_at 1.17537 myelin basic protein Mbp
1425264_s_at 1.05486 myelin basic protein Mbp
1425265_a_at 0.93302 prion protein interacting protein 1 Prnpi1
1425266_a_at 1.00755 "RAP1, GTP-GDP dissociation stimulator 1" Rap1gds1
1425267_a_at 0.91334 RIKEN cDNA 3110045G13 gene 3110045G13Rik
1425268_a_at 0.96324 RIKEN cDNA 3110045G13 gene 3110045G13Rik
1425269_at 1.00184 "amyloid beta (A4) precursor protein-binding, family B, member 1 interacting protein" Apbb1p
1425270_at 1.20773 kinesin family member 1B Kif1b
1425271_at 1.00335 "proteasome (prosome, macropain) 26S subunit, ATPase 3, interacting protein" Psmc3ip
1425272_at 0.87591 epithelial membrane protein 2 Emp2
1425273_s_at 0.84407 epithelial membrane protein 2 Emp2
1425274_at 1.18503 aspartate-beta-hydroxylase Asph
1425275_at 1.12842 aspartate-beta-hydroxylase Asph
1425276_at 0.87229 fibrosin 1 Fbs1
1425277_at 1.04687 slit homolog 1 (Drosophila) Slit1
1425278_at 1.11573 "ubiquitination factor E4A, UFD2 homolog (S. cerevisiae)" Ube4a
1425279_at 0.85137 PDLIM1 interacting kinase 1 like Pdik11
1425280_at 0.93462 leukocyte receptor cluster (LRC) member 1 Leng1
1425281_a_at 0.73093 TSC22 domain family 3 Tsc22d3
1425282_at 0.89308 IBR domain containing 2 Ibrdc2
1425283_a_at 1.24271 "metallothionein-like 5, testis-specific (tesmin)" Mtl5
1425284_a_at 1.06958 "RAB27A, member RAS oncogene family" Rab27a
1425285_a_at 0.98410 "RAB27A, member RAS oncogene family" Rab27a
1425286_at 0.95094 "crystallin, gamma N" Crygn
1425287_at 0.91372 zinc finger protein 189 Zfp189
1425288_at 0.98054 sterile alpha motif domain containing 11 Samd11
1425289_a_at 0.94896 complement receptor 2 Cr2
1425290_at 1.06169 syntaxin 19 Stx19
1425291_at 0.78290 forkhead box J1 Foxj1
1425292_at 1.05971 dystrobrevin alpha Dtna
1425293_a_at 1.06236 "zinc finger, CCHC domain containing 18" Zcchc18

1425294_at 1.05746 SLAM family member 8 Slamf8
1425295_at 0.84421 "eosinophil-associated, ribonuclease A family, member 11" Ear11
1425296_a_at 0.95693 regulator of G-protein signaling 3 /// similar to Regulator of G-protein signaling 3 (RGS3) (C2PA) /// similar to Regulator of G-protein signaling 3 (RGS3) (C2PA) Rgs3 ///
LOC436235 /// LOC670092
1425297_at 0.96437 cDNA sequence BC012278 BC012278
1425298_a_at 0.84662 baculoviral IAP repeat-containing 1a Birc1a
1425299_s_at 1.06761 RIKEN cDNA 0610038D11 gene /// similar to UPF0315 protein /// similar to UPF0315 protein /// similar to UPF0315 protein 0610038D11Rik ///
LOC623390 /// LOC627985 /// LOC631562 /// LOC665860
1425300_at 1.11846 dihydroxyacetone kinase 2 homolog (yeast) Dak
1425301_at 1.12706 neural cell adhesion molecule 2 Ncam2
1425302_at 0.85832 cDNA sequence BC024997 BC024997
1425303_at 0.95991 glucokinase Gck
1425304_s_at 0.93283 proline rich membrane anchor 1 Prima1
1425305_at 1.08266 zinc finger protein 295 Zfp295
1425306_at 0.95615 cDNA sequence BC027072 BC027072
1425307_at 0.96591 "dynein, axonemal, heavy chain 1" Dnahc1
1425308_at 0.95565 tripartite motif-containing 31 Trim31
1425309_at 1.04535 "cation channel, sperm associated 2" Catsper2
1425310_a_at 0.91191 EMI domain containing 2 Emid2
1425311_at 1.10647 RIKEN cDNA 4930432F04 gene 4930432F04Rik
1425312_s_at 1.08766 "DNA segment, Chr 11, ERATO Doi 636, expressed" D11Erttd636e
1425313_at 1.03865 calcium response factor Carf
1425314_at 0.94962 G protein-coupled receptor 98 Gpr98
1425315_at 1.60249 dedicator of cytokinesis 7 Dock7
1425316_at 0.83045 "beta-1,3-glucuronyltransferase 1 (glucuronosyltransferase P)" B3gat1
1425317_x_at 0.92748 serine threonine kinase 31 Stk31
1425318_a_at 0.92245 transmembrane protein 116 Tmem116
1425319_s_at 1.06753 RIKEN cDNA 6530403A03 gene 6530403A03Rik
1425320_at 1.14641 RIKEN cDNA A830023I12 gene A830023I12Rik
1425321_a_at 1.00674 calmin Clmn
1425322_at 0.99418 interferon zeta /// similar to interferon zeta /// similar to interferon zeta /// similar to interferon zeta /// similar to interferon zeta /// similar to interferon zeta /// similar to interferon zeta /// similar to interferon zeta /// similar to interferon zeta /// similar to interferon zeta /// similar to interferon zeta /// similar to interferon zeta /// similar to interferon zeta /// similar to interferon zeta /// similar to interferon zeta /// similar to interferon zeta /// similar to interferon zeta /// similar to interferon zeta /// similar to interferon zeta
Ifnz /// LOC435791 /// LOC545645 /// LOC545646 /// LOC545647 ///
LOC545648 /// LOC545649 /// LOC545650 /// LOC545651 /// LOC545652 /// LOC545653 ///
LOC545654 /// LOC545655 /// LOC638085 /// LOC641172 /// LOC668147 /// LOC668208 ///
LOC671856 /// LOC673717
1425323_a_at 1.12137 cDNA sequence BC008155 BC008155
1425324_x_at 1.05663 immunoglobulin heavy chain 4 (serum IgG1) Igh-4

1425325_at 0.93450 "membrane-spanning 4-domains, subfamily A, member 4B" Ms4a4b
 1425326_at 1.36034 ATP citrate lyase Acly
 1425327_at 0.96667 cDNA sequence BC008163 BC008163
 1425328_at 0.97914 cDNA sequence BC008163 BC008163
 1425329_a_at 1.08855 cytochrome b5 reductase 3 Cyb5r3
 1425330_a_at 0.93378 "protein phosphatase 1B, magnesium dependent, beta isoform" Ppm1b
 1425331_at 1.08482 zinc finger protein 106 Zfp106
 1425332_at 1.04043 zinc finger protein 106 Zfp106
 1425333_at 0.90803 RIKEN cDNA 1810048P08 gene 1810048P08Rik
 1425334_at 0.99821 golgi reassembly stacking protein 1 Gorasp1
 1425335_at 0.99528 "CD8 antigen, alpha chain /// similar to T-cell surface glycoprotein CD8 alpha chain precursor (T-cell surface glycoprotein Lyt-2) /// similar to T-cell surface glycoprotein CD8 alpha chain precursor (T-cell surface glycoprotein Lyt-2) (CD8a antigen)" Cd8a /// LOC636147 /// LOC669166
 1425336_x_at 1.20515 "histocompatibility 2, K1, K region" H2-K1
 1425337_at 0.84854 "solute carrier family 12, member 5" Slc12a5
 1425338_at 0.92434 "phospholipase C, beta 4" Plcb4
 1425339_at 0.89414 "phospholipase C, beta 4" Plcb4
 1425340_a_at 0.98362 "protein tyrosine phosphatase, receptor type, A" Ptptra
 1425341_at 1.00784 "potassium channel, subfamily K, member 3" Kcnk3
 1425342_a_at 0.96756 "potassium channel, subfamily K, member 3" Kcnk3
 1425343_at 0.90714 haloacid dehalogenase-like hydrolase domain containing 3 Hdhd3
 1425344_at 1.02316 nuclear prelamin A recognition factor Narf
 1425345_at 0.88534 coiled-coil domain containing 28A Ccdc28a
 1425346_at 0.99259 zinc finger protein 318 Zfp318
 1425347_a_at 1.06152 zinc finger protein 318 Zfp318
 1425348_a_at 0.81260 "signal recognition particle receptor, B subunit" Srprb
 1425349_a_at 1.07518 "myelin basic protein expression factor 2, repressor" Myef2
 1425350_a_at 1.00808 "myelin basic protein expression factor 2, repressor" Myef2
 1425351_at 1.01303 sulfiredoxin 1 homolog (S. cerevisiae) Srxn1
 1425352_at 0.96749 REST corepressor 3 Rcor3
 1425353_at 0.98316 ectonucleotide pyrophosphatase/phosphodiesterase 3 Enpp3
 1425354_a_at 0.93295 angiogenic factor with G patch and FHA domains 1 Aggf1
 1425355_at 0.93860 cDNA sequence BC018371 /// hypothetical LOC433604 BC018371 /// LOC433604
 1425356_at 0.89746 zinc finger protein 142 Zfp142
 1425357_a_at 1.09458 gremlin 1 Grem1
 1425358_at 1.07743 RIO kinase 1 (yeast) Riok1
 1425359_at 1.02031 "mal, T-cell differentiation protein-like" Mall
 1425360_at 0.92420 similar to myeloid/lymphoid or mixed lineage-leukemia translocation to 6 homolog /// similar to myeloid/lymphoid or mixed lineage-leukemia translocation to 6 homolog LOC639490 /// LOC669933
 1425361_at 1.00315 zinc finger protein 397 Zfp397

1425362_at 2.96741 HIV-1 Rev binding protein-like Hrbl
 1425363_at 0.94237 "beta-1,4-N-acetyl-galactosaminyl transferase 1" B4galnt1
 1425364_a_at 0.90567 "solute carrier family 3 (activators of dibasic and neutral amino acid transport), member 2" Slc3a2
 1425365_a_at 0.97521 "cytochrome P450, family 2, subfamily d, polypeptide 13" Cyp2d13
 1425366_a_at 0.96091 Hus1 homolog (S. pombe) Hus1
 1425367_at 1.08562 integrin alpha L /// similar to Integrin alpha-L precursor (Leukocyte adhesion glycoprotein LFA-1 alpha chain) (LFA-1A) (Leukocyte function-associated molecule 1 alpha chain) (Lymphocyte antigen Ly-15) (CD11a antigen) Itgal /// LOC671163
 1425368_a_at 0.91836 numb gene homolog (Drosophila) Numb
 1425369_a_at 0.95314 SRY-box containing gene 10 Sox10
 1425370_a_at 0.93997 avian erythroblastosis virus E-26 (v-ets) oncogene related Erg
 1425371_at 1.42655 "polymerase (DNA directed), beta" Polb
 1425372_at 0.95711 "DnaJ (Hsp40) homolog, subfamily C, member 19" Dnajc19
 1425373_a_at 1.06306 "tumor necrosis factor superfamily, member 5-induced protein 1" Tnfsf5ip1
 1425374_at 1.11492 2'-5' oligoadenylate synthetase 3 Oas3
 1425375_at 1.09736 TD and POZ domain containing 1 Tdpoz1
 1425376_at 0.98643 arachidonate 8-lipoxygenase Alox8
 1425377_at 0.97844 wingless-related MMTV integration site 1 Wnt1
 1425378_at 1.02087 hect (homologous to the E6-AP (UBE3A) carboxyl terminus) domain and RCC1 (CHC1)-like domain (RLD) 1 Herc1
 1425379_at 0.98014 hepatocyte growth factor Hgf
 1425380_at 0.87362 RAS guanyl releasing protein 4 Rasgrp4
 1425381_a_at 0.94881 transferrin receptor 2 Trfr2
 1425382_a_at 0.98954 aquaporin 4 Aqp4
 1425383_a_at 0.75490 pre B-cell leukemia transcription factor 1 /// region containing RIKEN cDNA 2310056B04 gene; pre B-cell leukemia transcription factor 1 Pbx1 /// LOC676870
 1425384_a_at 1.09640 "ubiquitination factor E4A, UFD2 homolog (S. cerevisiae)" Ube4a
 1425385_a_at 0.96398 immunoglobulin heavy chain 1a (serum IgG2a) Igh-1a
 1425386_at 1.03978 RIKEN cDNA 4833422F24 gene 4833422F24Rik
 1425387_at 0.86965 "aldo-keto reductase family 1, member C20" Akr1c20
 1425388_a_at 0.84358 thiamine pyrophosphokinase Tpk1
 1425389_a_at 0.97632 runt related transcription factor 2 Runx2
 1425390_at 1.00590 "adenosine deaminase, tRNA-specific 1" Adat1
 1425391_a_at 0.94082 oxysterol binding protein-like 5 Osbp15
 1425392_a_at 0.94371 "nuclear receptor subfamily 1, group I, member 3" Nr1i3
 1425393_a_at 0.89975 mitogen activated protein kinase kinase 7 Map2k7
 1425394_at 0.87521 cDNA sequence BC023105 BC023105
 1425395_at 0.94026 a disintegrin and metallopeptidase domain 26A (testase 3) Adam26a
 1425396_a_at 0.92729 lymphocyte protein tyrosine kinase Lck
 1425397_at 0.99689 CDNA clone MGC:6909 IMAGE:2655941 ---
 1425398_at 0.94261 "histone 1, H2bc /// similar to Histone H2B F (H2B 291A) /// similar to Histone H2B F (H2B 291A) /// similar to Histone H2B F (H2B 291A)" Hist1h2bc /// LOC665596 ///

LOC665622 /// LOC671645

1425399_at 1.04635 RIKEN cDNA 9630055N22 gene 9630055N22Rik
 1425400_a_at 0.96056 "Cbp/p300-interacting transactivator, with Glu/Asp-rich carboxy-terminal domain, 4" Cited4
 1425401_at 1.04765 "DNA segment, Chr 11, ERATO Doi 636, expressed" D11Ertd636e
 1425402_at 1.10836 thimet oligopeptidase 1 Thop1
 1425403_at 1.01347 dynamin 3 Dnm3
 1425404_a_at 0.94438 transmembrane protein 110 Tmem110
 1425405_a_at 1.30733 "adenosine deaminase, RNA-specific" Adar
 1425406_at 1.09294 "C-type lectin domain family 4, member a2" Clec4a2
 1425407_s_at 1.02978 "C-type lectin domain family 4, member a2 /// C-type lectin domain family 4, member b1" Clec4a2 /// Clec4b1
 1425408_a_at 1.06349 RIKEN cDNA 2610034M16 gene 2610034M16Rik
 1425409_at 1.03887 "cholinergic receptor, nicotinic, alpha polypeptide 2 (neuronal)" Chrna2
 1425410_at 0.96348 Tp53rk binding protein Tprkb
 1425411_at 0.86894 ADP-ribosylation factor-like 4A Arl4a
 1425412_at 1.00692 cold autoinflammatory syndrome 1 homolog (human) Cias1
 1425413_at 0.85118 "regulatory factor X, 3 (influences HLA class II expression)" Rfx3
 1425414_at 0.95608 "protein phosphatase 1, regulatory (inhibitor) subunit 16B" Ppp1r16b
 1425415_a_at 0.88672 "solute carrier family 1 (neuronal/epithelial high affinity glutamate transporter, system Xag), member 1" Slc1a1
 1425416_s_at 0.97682 proline/serine-rich coiled-coil 1 Psrc1
 1425417_x_at 1.03122 "killer cell lectin-like receptor, subfamily A, member 8 /// killer cell lectin-like receptor subfamily A, member 21 /// similar to killer cell lectin-like receptor subfamily A, member 21 /// killer cell lectin-like receptor subfamily A member 29 /// similar to killer cell lectin-like receptor subfamily A, member 21 /// similar to Killer cell lectin-like receptor 8 (T-cell surface glycoprotein Ly-49H) (Ly49-H antigen) (Lymphocyte antigen 49H)" Klra8 /// Klra21 /// LOC640965 /// Klra29 ///
 LOC667769 /// LOC673223
 1425418_at 0.94603 WAP four-disulfide core domain 5 Wfdc5
 1425419_a_at 0.96392 v-raf-leukemia viral oncogene 1 Raf1
 1425420_s_at 1.20532 large tumor suppressor 2 Lats2
 1425421_at 1.23554 Retinoblastoma binding protein 6 Rbbp6
 1425422_a_at 0.97407 poly(A)-specific ribonuclease (deadenylation nuclease) Parn
 1425423_at 0.83561 GLIS family zinc finger 1 Glis1
 1425424_at 1.13825 hypothetical protein LOC620031 MGC7817
 1425425_a_at 1.00069 Wnt inhibitory factor 1 Wif1
 1425426_a_at 0.75378 myocyte enhancer factor 2A Mef2a
 1425427_at 0.89073 hypothetical protein LOC639910 LOC639910
 1425428_at 0.87822 "hypoxia inducible factor 3, alpha subunit" Hif3a
 1425429_s_at 0.92791 "hypoxia inducible factor 3, alpha subunit /// similar to hypoxia inducible factor 3, alpha subunit" Hif3a /// LOC641092
 1425430_at 0.90711 keratin associated protein 16-5 Krtap16-5
 1425431_at 1.01925 keratin associated protein 16-10 /// keratin associated protein 16-3 /// keratin

associated protein 16-6 Krtap16-10 /// Krtap16-3 /// Krtap16-6
 1425432_at 0.87237 "opioid receptor, mu 1" Oprm1
 1425433_a_at 0.94375 neuropilin 2 Nrp2
 1425434_a_at 0.91694 macrophage scavenger receptor 1 Msr1
 1425435_at 0.97368 macrophage scavenger receptor 1 Msr1
 1425436_x_at 1.03703 "killer cell lectin-like receptor, subfamily A, member 3" Klra3
 1425437_a_at 1.14611 "potassium channel, subfamily K, member 7" Kcnk7
 1425438_at 0.94615 --- ---
 1425439_a_at 0.81891 "solute carrier family 41, member 3" Slc41a3
 1425440_x_at 0.91689 "solute carrier family 41, member 3" Slc41a3
 1425441_at 0.93295 --- ---
 1425442_at 1.01858 osteoclast associated receptor Oscar
 1425443_at 1.08624 "transcription factor AP-2, delta" Tcfap2d
 1425444_a_at 1.00112 "transforming growth factor, beta receptor II" Tgfbr2
 1425445_a_at 1.02389 claudin 18 Cldn18
 1425446_at 0.90537 --- ---
 1425447_at 1.06648 dickkopf homolog 4 (Xenopus laevis) Dkk4
 1425448_x_at 1.00442 "ATPase, H+ transporting, lysosomal V0 subunit B" Atp6v0b
 1425449_at 0.90989 phosphatidic acid phosphatase 2a Ppap2a
 1425450_at 0.88902 chitinase 3-like 4 Chi3l4
 1425451_s_at 0.97096 chitinase 3-like 3 /// chitinase 3-like 4 Chi3l3 /// Chi3l4
 1425452_s_at 0.92275 "protein tyrosine phosphatase, receptor type, J /// expressed sequence
 AW125753" Ptpnj /// AW125753
 1425453_x_at 1.01977 "killer cell lectin-like receptor subfamily A, member 12" Klra12
 1425454_a_at 0.99882 interleukin 12a Il12a
 1425455_a_at 1.03731 churchill domain containing 1 Churc1
 1425456_a_at 1.05646 mitogen activated protein kinase kinase 3 Map2k3
 1425457_a_at 1.03315 growth factor receptor bound protein 10 Grb10
 1425458_a_at 0.86352 growth factor receptor bound protein 10 Grb10
 1425459_at 1.05614 myotubularin related protein 2 Mtmr2
 1425460_at 1.03564 myotubularin related protein 2 Mtmr2
 1425461_at 0.87754 F-box and WD-40 domain protein 11 Fbxw11
 1425462_at 0.96376 F-box and WD-40 domain protein 11 Fbxw11
 1425463_at 0.76842 GATA binding protein 6 Gata6
 1425464_at 0.77540 GATA binding protein 6 Gata6
 1425465_a_at 0.99764 SUMO/sentrin specific peptidase 2 Senp2
 1425466_at 1.04261 SUMO/sentrin specific peptidase 2 Senp2
 1425467_a_at 0.91063 proteolipid protein (myelin) 1 Plp1
 1425468_at 0.97990 proteolipid protein (myelin) 1 Plp1
 1425469_a_at 0.91943 RIKEN cDNA 9030208C03 gene 9030208C03Rik
 1425470_at 0.97190 --- ---
 1425471_x_at 0.82961 --- ---
 1425472_a_at 1.05883 lamin A Lmna

1425473_at	1.08974	"cofactor required for Sp1 transcriptional activation, subunit 6"	Crsp6
1425474_a_at	0.83002	vacuolar protein sorting 39 (yeast)	Vps39
1425475_at	0.72344	"procollagen, type IV, alpha 5"	Col4a5
1425476_at	0.75076	"procollagen, type IV, alpha 5"	Col4a5
1425477_x_at	0.90709	"histocompatibility 2, class II antigen A, beta 1"	H2-Ab1
1425478_x_at	0.90886	ubiquitin-conjugating enzyme E2I	Ube2i
1425479_at	1.05304	SET and MYND domain containing 5	Smyd5
1425480_at	1.01225	"CCR4-NOT transcription complex, subunit 6-like"	Cnot6l
1425481_at	0.93073	"CCR4-NOT transcription complex, subunit 6-like"	Cnot6l
1425482_s_at	0.91985	ankyrin repeat and MYND domain containing 2	Ankmy2
1425483_at	0.89312	thymocyte selection-associated HMG box gene	Tox
1425484_at	1.10343	thymocyte selection-associated HMG box gene	Tox
1425485_at	0.99147	myotubularin related protein 6	Mtmr6
1425486_s_at	1.00824	myotubularin related protein 6	Mtmr6
1425487_at	0.92019	"DNA segment, Chr 11, ERATO Doi 730, expressed"	D11Ertd730e
1425488_at	1.06917	"DNA segment, Chr 11, ERATO Doi 730, expressed"	D11Ertd730e
1425489_at	0.95108	"DNA segment, Chr 11, ERATO Doi 730, expressed"	D11Ertd730e
1425490_a_at	1.01216	WD repeat domain 13	Wdr13
1425491_at	0.94293	"bone morphogenetic protein receptor, type 1A"	Bmpr1a
1425492_at	1.02316	"bone morphogenetic protein receptor, type 1A"	Bmpr1a
1425493_at	0.87398	"bone morphogenetic protein receptor, type 1A"	Bmpr1a
1425494_s_at	0.93864	"bone morphogenetic protein receptor, type 1A"	Bmpr1a
1425495_at	0.88325	zinc finger protein 62	Zfp62
1425496_at	0.93302	"ATP-binding cassette, sub-family A (ABC1), member 3"	Abca3
1425497_a_at	0.98890	PRP4 pre-mRNA processing factor 4 homolog B (yeast)	Prpf4b
1425498_at	1.03625	PRP4 pre-mRNA processing factor 4 homolog B (yeast)	Prpf4b
1425499_at	1.00895	Hypothetical protein LOC625794	LOC625794
1425500_x_at	1.07324	Hypothetical protein LOC625794	LOC625794
1425501_at	0.88966	"Mammary gland RCB-0526 Jyg-MC(A) cDNA, RIKEN full-length enriched library, clone:G830022P11 product:unclassifiable, full insert sequence" ---	
1425502_x_at	0.92342	"Mammary gland RCB-0526 Jyg-MC(A) cDNA, RIKEN full-length enriched library, clone:G830022P11 product:unclassifiable, full insert sequence" ---	
1425503_at	1.13661	"glucosaminyl (N-acetyl) transferase 2, I-branching enzyme"	Gcnt2
1425504_at	0.92067	"myosin, light polypeptide kinase"	Mylk
1425505_at	0.93188	"myosin, light polypeptide kinase"	Mylk
1425506_at	1.11682	"myosin, light polypeptide kinase"	Mylk
1425507_at	1.12686	ADP-ribosylation factor related protein 1	Arfrp1
1425508_s_at	0.97844	ADP-ribosylation factor related protein 1	Arfrp1
1425509_at	0.99252	MAP/microtubule affinity-regulating kinase 1	Mark1
1425510_at	1.02297	MAP/microtubule affinity-regulating kinase 1	Mark1
1425511_at	0.86869	MAP/microtubule affinity-regulating kinase 1	Mark1
1425512_at	1.03786	mitogen activated protein kinase kinase 7	Map2k7
1425513_at	0.93882	mitogen activated protein kinase kinase 7	Map2k7

1425514_at 1.09358 "phosphatidylinositol 3-kinase, regulatory subunit, polypeptide 1 (p85 alpha)"
 Pik3r1
 1425515_at 0.92162 "phosphatidylinositol 3-kinase, regulatory subunit, polypeptide 1 (p85 alpha)"
 Pik3r1
 1425516_at 0.90877 O-linked N-acetylglucosamine (GlcNAc) transferase (UDP-N-acetylglucosamine:polypeptide-N-acetylglucosaminyl transferase) Ogt
 1425517_s_at 0.93836 O-linked N-acetylglucosamine (GlcNAc) transferase (UDP-N-acetylglucosamine:polypeptide-N-acetylglucosaminyl transferase) Ogt
 1425518_at 0.90701 Rap guanine nucleotide exchange factor (GEF) 4 Rapgef4
 1425519_a_at 1.41070 "CD74 antigen (invariant polypeptide of major histocompatibility complex, class II antigen-associated)" Cd74
 1425520_a_at 0.87053 dehydrodolichyl diphosphate synthase Dhdds
 1425521_at 1.04422 polyadenylate binding protein-interacting protein 1 Paip1
 1425522_at 1.56341 RIKEN cDNA 2600011C06 gene 2600011C06Rik
 1425523_at 1.08922 RIKEN cDNA 2600011C06 gene 2600011C06Rik
 1425524_at 1.84224 RIKEN cDNA 2600011C06 gene 2600011C06Rik
 1425525_a_at 1.12197 "purinergic receptor P2X, ligand-gated ion channel 4" P2rx4
 1425526_a_at 0.95720 paired related homeobox 1 Prrx1
 1425527_at 0.96113 paired related homeobox 1 Prrx1
 1425528_at 0.96804 paired related homeobox 1 Prrx1
 1425529_s_at 0.96357 "DNA segment, Chr 19, Wayne State University 162, expressed"
 D19Wsu162e
 1425530_a_at 0.98149 syntaxin 3 Stx3
 1425531_at 1.05204 "zinc finger, HIT domain containing 1" Znhit1
 1425532_a_at 0.95297 bridging integrator 1 Bin1
 1425533_a_at 0.90833 staufen (RNA binding protein) homolog 2 (Drosophila) Stau2
 1425534_at 0.78537 staufen (RNA binding protein) homolog 2 (Drosophila) Stau2
 1425535_at 0.83831 replication initiator 1 Repin1
 1425536_at 1.30666 syntaxin 3 Stx3
 1425537_at 1.10411 "protein phosphatase 1A, magnesium dependent, alpha isoform" Ppm1a
 1425538_x_at 1.06451 CEA-related cell adhesion molecule 1 Ceacam1
 1425539_a_at 0.97883 reticulon 3 Rtn3
 1425540_at 0.99790 ornithine transcarbamylase Otc
 1425541_at 1.01195 DEAD (Asp-Glu-Ala-Asp) box polypeptide 27 Ddx27
 1425542_a_at 1.09644 "protein phosphatase 2, regulatory subunit B (B56), gamma isoform"
 Ppp2r5c
 1425543_s_at 0.76084 "pleckstrin homology domain containing, family A member 5" Plekha5
 1425544_at 0.88618 "pleckstrin homology domain containing, family A member 5" Plekha5
 1425545_x_at 2.10424 "histocompatibility 2, D region locus 1" H2-D1
 1425546_a_at 0.82632 transferrin Trf
 1425547_a_at 0.89759 kinesin light chain 4 Klc4
 1425548_a_at 1.07125 leukocyte specific transcript 1 Lst1
 1425549_at 1.15704 presenilin 1 Psen1

1425550_a_at 0.98235 "protein kinase, cAMP dependent regulatory, type I, alpha" Prkar1a
 1425551_at 0.86226 huntingtin interacting protein 1 related Hip1r
 1425552_at 0.94219 huntingtin interacting protein 1 related Hip1r
 1425553_s_at 0.81963 huntingtin interacting protein 1 related Hip1r
 1425554_a_at 1.01308 CDC16 cell division cycle 16 homolog (*S. cerevisiae*) Cdc16
 1425555_at 0.92760 "Cdc2-related kinase, arginine/serine-rich" Crkrs
 1425556_at 1.11113 "Cdc2-related kinase, arginine/serine-rich" Crkrs
 1425557_x_at 0.90523 TSC22 domain family 3 Tsc22d3
 1425558_at 0.96304 kinesin light chain 3 Klc3
 1425559_a_at 0.86682 acyl-CoA synthetase medium-chain family member 3 Acsm3
 1425560_a_at 0.84986 S100 calcium binding protein A16 S100a16
 1425561_at 0.97144 "tRNA nucleotidyl transferase, CCA-adding, 1" Trnt1
 1425562_s_at 0.97134 "tRNA nucleotidyl transferase, CCA-adding, 1" Trnt1
 1425563_s_at 0.90199 protocadherin 10 Pcdh10
 1425564_at 0.81382 RE1-silencing transcription factor Rest
 1425565_at 0.96128 RE1-silencing transcription factor Rest
 1425566_at 0.94348 RE1-silencing transcription factor /// hypothetical protein LOC665152 ///
 hypothetical protein LOC669155 Rest /// LOC665152 /// LOC669155
 1425567_a_at 0.85866 annexin A5 Anxa5
 1425568_a_at 1.01417 transmembrane protein 33 Tmem33
 1425569_a_at 0.91276 signaling lymphocytic activation molecule family member 1 Slamf1
 1425570_at 0.98347 signaling lymphocytic activation molecule family member 1 Slamf1
 1425571_at 1.00137 signaling lymphocytic activation molecule family member 1 Slamf1
 1425572_a_at 1.01648 development and differentiation enhancing Ddef1
 1425573_a_at 1.02377 development and differentiation enhancing Ddef1
 1425574_at 1.00781 Eph receptor A3 Epha3
 1425575_at 0.92540 Eph receptor A3 Epha3
 1425576_at 0.99514 S-adenosylhomocysteine hydrolase-like 1 Ahcy11
 1425577_at 1.02016 RIKEN cDNA 9830124H08 gene 9830124H08Rik
 1425578_a_at 0.98898 glial cell line derived neurotrophic factor family receptor alpha 2 Gfra2
 1425579_at 0.99337 glial cell line derived neurotrophic factor family receptor alpha 2 Gfra2
 1425580_a_at 0.87612 "phosphoinositide-3-kinase, class 3" Pik3c3
 1425581_s_at 1.31811 UDP-N-acetyl-alpha-D-galactosamine: polypeptide N-
 acetylgalactosaminyltransferase 7 Galnt7
 1425582_a_at 0.97888 endomucin Emcn
 1425583_at 0.94782 "Mammary gland RCB-0526 Jyg-MC(A) cDNA, RIKEN full-length enriched
 library, clone:G830022P11 product:unclassifiable, full insert sequence" ---
 1425584_x_at 0.89838 "Mammary gland RCB-0526 Jyg-MC(A) cDNA, RIKEN full-length enriched
 library, clone:G830022P11 product:unclassifiable, full insert sequence" ---
 1425585_at 0.94934 "mediator of RNA polymerase II transcription, subunit 12 homolog (yeast)"
 Med12
 1425586_a_at 0.93364 melanophilin Mlph
 1425587_a_at 1.08224 "protein tyrosine phosphatase, receptor type, J" Ptprij

1425588_at 0.96190 "protein tyrosine phosphatase, receptor type, J" Ptpnj
1425589_at 0.83514 hydroxysteroid (17-beta) dehydrogenase 13 Hsd17b13
1425590_s_at 0.98185 aryl hydrocarbon receptor-interacting protein-like 1 Aipl1
1425591_a_at 1.08080 chromatin modifying protein 2A Chmp2a
1425592_at 1.06873 "transportin 2 (importin 3, karyopherin beta 2b)" Tnp2
1425593_at 1.08138 "transportin 2 (importin 3, karyopherin beta 2b)" Tnp2
1425594_at 0.84630 laminin gamma 3 Lamc3
1425595_at 0.96974 "gamma-aminobutyric acid (GABA-B) receptor, 1" Gabbr1
1425596_at 0.99934 expressed sequence AI317395 AI317395
1425597_a_at 1.06323 quaking Qk
1425598_a_at 1.07168 similar to Yamaguchi sarcoma viral (v-yes-1) oncogene homolog LOC676654
1425599_a_at 0.93180 GATA zinc finger domain containing 1 Gatad1
1425600_a_at 1.00757 "phospholipase C, beta 1" Plcb1
1425601_a_at 0.91255 rhotekin Rtkn
1425602_a_at 0.97633 "rabaptin, RAB GTPase binding effector protein 2" Rabep2
1425603_at 1.27999 RIKEN cDNA 0610011I04 gene 0610011I04Rik
1425604_at 1.09207 v-crk sarcoma virus CT10 oncogene homolog (avian)-like Crkl
1425605_a_at 0.93351 limb region 1 Lmbr1
1425606_at 0.99924 "solute carrier family 5 (iodide transporter), member 8" Slc5a8
1425607_at 1.01683 holocarboxylase synthetase (biotin- [propionyl-Coenzyme A-carboxylase (ATP-hydrolysing)] ligase) Hlcs
1425608_at 0.95699 dual specificity phosphatase 3 (vaccinia virus phosphatase VH1-related) Dusp3
1425609_at 0.95036 neutrophil cytosolic factor 1 Ncf1
1425610_s_at 0.78849 UDP-N-acetyl-alpha-D-galactosamine:polypeptide N-acetylgalactosaminyltransferase 2 Galnt2
1425611_a_at 0.76800 cut-like 1 (Drosophila) Cutl1
1425612_at 0.97397 cut-like 1 (Drosophila) Cutl1
1425613_at 1.00609 protocadherin alpha 11 Pcdha11
1425614_x_at 1.96110 "histocompatibility 2, D region locus 1" H2-D1
1425615_a_at 0.99363 phosphoenolpyruvate carboxykinase 2 (mitochondrial) Pck2
1425616_a_at 0.90184 coiled-coil domain containing 23 Ccdc23
1425617_at 1.18137 DEAH (Asp-Glu-Ala-His) box polypeptide 9 Dhx9
1425618_at 0.96818 DEAH (Asp-Glu-Ala-His) box polypeptide 9 Dhx9
1425619_s_at 0.83593 desmoglein 2 Dsg2
1425620_at 0.79859 "transforming growth factor, beta receptor III" Tgfbr3
1425621_at 0.83210 tripartite motif-containing 35 Trim35
1425622_at 1.06738 EGF-like repeats and discoidin I-like domains 3 Edil3
1425623_a_at 0.66224 cystathionine beta-synthase Cbs
1425624_at 0.86894 EPM2A (laforin) interacting protein 1 Epm2aip1
1425625_at 1.02285 "interleukin 13 receptor, alpha 1" Il13ra1
1425626_at 0.87020 "glutathione S-transferase, mu 1 /// similar to Glutathione S-transferase Mu 1 (GST class-mu 1) (Glutathione S-transferase GT8.7) (pmGT10) (GST 1-1) /// similar to Glutathione S-transferase Mu 1 (GST class-mu 1) (Glutathione S-transferase GT8.7) (pmGT10) (GST 1-1)" Gstm1 ///

LOC433943 /// LOC674086

1425627_x_at 0.82174 "glutathione S-transferase, mu 1 /// similar to Glutathione S-transferase Mu 1 (GST class-mu 1) (Glutathione S-transferase GT8.7) (pmGT10) (GST 1-1) /// similar to Glutathione S-transferase Mu 1 (GST class-mu 1) (Glutathione S-transferase GT8.7) (pmGT10) (GST 1-1)" Gstm1 /// LOC433943 /// LOC674086

1425628_a_at 0.86335 similar to General transcription factor II-I (GTFII-I) (TFII-I) (Bruton tyrosine kinase-associated protein 135) (BTK-associated protein 135) (BAP-135) LOC669007

1425629_a_at 1.04935 nucleolar protein family 6 (RNA-associated) Nol6

1425630_at 0.89617 "transcriptional regulator, SIN3B (yeast)" Sin3b

1425631_at 1.10489 "protein phosphatase 1, regulatory (inhibitor) subunit 3C" Ppp1r3c

1425632_a_at 0.98342 PQ loop repeat containing 2 Pqlc2

1425633_at 0.93082 cDNA sequence BC026782 BC026782

1425634_a_at 0.90450 "tyrosine kinase, non-receptor, 1" Tnk1

1425635_at 0.83957 "tyrosine kinase, non-receptor, 1" Tnk1

1425636_at 1.05948 hedgehog acyltransferase Hhat

1425637_at 0.91850 --- ---

1425638_at 0.97097 "centaurin, alpha 2" Centa2

1425639_at 0.91037 "centaurin, alpha 2" Centa2

1425640_at 0.96511 "AF4/FMR2 family, member 1" Aff1

1425641_at 1.02328 --- ---

1425642_at 0.86955 centrosomal protein 290 Cep290

1425643_at 0.99006 glycophorin A Gypa

1425644_at 0.91097 leptin receptor Lepr

1425645_s_at 0.97876 "cytochrome P450, family 2, subfamily b, polypeptide 10" Cyp2b10

1425646_at 1.09055 cDNA sequence BC016495 BC016495

1425647_at 1.02827 tripartite motif-containing 60 Trim60

1425648_at 0.89174 tripartite motif-containing 60 /// tripartite motif-containing 61 Trim60 /// Trim61

1425649_at 0.90493 "solute carrier family 39 (zinc transporter), member 14" Slc39a14

1425650_at 0.94403 "transducin-like enhancer of split 4, homolog of Drosophila E(spl)" Tle4

1425651_at 0.97062 nonhomologous end-joining factor 1 Nhej1

1425652_s_at 0.86967 RNA binding protein gene with multiple splicing Rbpms

1425653_at 1.00758 activating transcription factor 7 Atf7

1425654_a_at 0.93247 LIM domain and actin binding 1 Lima1

1425655_at 0.96087 keratin associated protein 16-1 Krtap16-1

1425656_a_at 0.93056 brain-specific angiogenesis inhibitor 1-associated protein 2 Baiap2

1425657_at 0.97700 RIKEN cDNA 1700016D06 gene 1700016D06Rik

1425658_at 0.90930 CD109 antigen Cd109

1425659_at 1.03113 target of myb1-like 2 (chicken) Tom112

1425660_at 0.99626 BTB (POZ) domain containing 3 Btbd3

1425661_at 0.93501 cytidine and dCMP deaminase domain containing 1 Cdadc1

1425662_at 0.99236 cytidine and dCMP deaminase domain containing 1 Cdadc1

1425663_at 0.91992 interleukin 1 receptor antagonist Il1rn

1425664_at 1.08492 kelch-like 20 (Drosophila) Klhl20
1425665_a_at 0.97685 signal recognition particle 54 /// similar to signal recognition particle 54
Srp54 /// LOC665155
1425666_at 1.02215 similar to zinc finger protein of the cerebellum 5 LOC624168
1425667_at 1.17298 acyl-CoA thioesterase 11 Acot11
1425668_a_at 0.91086 "ST3 beta-galactoside alpha-2,3-sialyltransferase 4" St3gal4
1425669_at 0.89634 "MOB1, Mps One Binder kinase activator-like 2B (yeast)" Mobk12b
1425670_at 0.88259 regulatory factor X-associated ankyrin-containing protein Rfxank
1425671_at 0.93635 homer homolog 1 (Drosophila) Homer1
1425672_a_at 0.96249 "transient receptor potential cation channel, subfamily C, member 2" Trpc2
1425673_at 1.72086 LIM domain containing preferred translocation partner in lipoma Lpp
1425674_a_at 1.08554 Ssu72 RNA polymerase II CTD phosphatase homolog (yeast) Ssu72
1425675_s_at 0.87601 CEA-related cell adhesion molecule 1 Ceacam1
1425676_a_at 1.13392 "elongation of very long chain fatty acids (FEN1/Elo2, SUR4/Elo3, yeast)-like
1" Elovl1
1425677_a_at 1.05786 "ankyrin 1, erythroid" Ank1
1425678_a_at 0.81377 SNF related kinase Snrk
1425679_a_at 0.77834 mitogen activated protein kinase 8 interacting protein 1 Mapk8ip1
1425680_a_at 1.03821 beta-transducin repeat containing protein Btrc
1425681_a_at 2.42038 prion protein dublet Prnd
1425682_a_at 1.29393 Tp53rk binding protein Tprkb
1425683_at 1.02567 --- ---
1425684_at 0.98648 RIKEN cDNA 2310005E10 gene 2310005E10Rik
1425685_at 1.07561 RIKEN cDNA 2310005E10 gene 2310005E10Rik
1425686_at 1.11040 CASP8 and FADD-like apoptosis regulator Cflar
1425687_at 1.18785 CASP8 and FADD-like apoptosis regulator Cflar
1425688_a_at 1.10847 dihydropyrimidinase Dpys
1425689_at 0.96481 dihydropyrimidinase Dpys
1425690_at 1.03704 "beta-1,3-glucuronyltransferase 1 (glucuronosyltransferase P)" B3gat1
1425691_at 1.01707 "beta-1,3-glucuronyltransferase 1 (glucuronosyltransferase P)" B3gat1
1425692_a_at 1.03004 "dynein, axonemal, heavy chain 8" Dnahc8
1425693_at 1.11620 Braf transforming gene Braf
1425694_at 1.17014 T-box 5 Tbx5
1425695_at 0.92103 T-box 5 Tbx5
1425696_at 1.04984 thioredoxin-like 6 Txnl6
1425697_at 0.98197 serologically defined colon cancer antigen 1 Sdccag1
1425698_a_at 0.84573 HCF-binding transcription factor Zhangfei Zf
1425699_a_at 0.88244 abhydrolase domain containing 14A Abhd14a
1425700_at 1.02957 "glutamate receptor, metabotropic 1" Grm1
1425701_a_at 0.94478 regulator of G-protein signaling 3 Rgs3
1425702_a_at 0.76988 ectonucleotide pyrophosphatase/phosphodiesterase 5 Enpp5
1425703_at 0.86538 peroxisome proliferator activator receptor delta Ppard
1425704_at 0.88154 cDNA sequence BC022224 BC022224

1425705_a_at 0.83124 ERO1-like beta (*S. cerevisiae*) Ero1lb
1425706_a_at 0.87776 damage specific DNA binding protein 2 Ddb2
1425707_a_at 1.01779 "potassium inwardly-rectifying channel, subfamily J, member 6" Kcnj6
1425708_at 1.01286 ring finger protein 17 Rnf17
1425709_at 0.96122 ring finger protein 17 Rnf17
1425710_a_at 0.90500 homer homolog 1 (*Drosophila*) Homer1
1425711_a_at 0.99263 thymoma viral proto-oncogene 1 Akt1
1425712_at 1.07584 cDNA sequence BC025446 BC025446
1425713_a_at 0.82942 ring finger protein 146 Rnf146
1425714_a_at 1.05169 Nfat activating molecule with ITAM motif 1 Nfam1
1425715_at 0.86092 "interleukin 1 family, member 8" Il1f8
1425716_s_at 1.00571 BCL2-antagonist/killer 1 Bak1
1425717_at 1.07212 LPS-responsive beige-like anchor Lrba
1425718_a_at 0.77956 influenza virus NS1A binding protein Ivns1abp
1425719_a_at 1.26129 N-myc (and STAT) interactor Nmi
1425720_at 0.92758 --- ---
1425721_at 0.85820 pleckstrin homology domain interacting protein Phip
1425722_at 1.03636 similar to interferon-inducible GTPase LOC435565
1425723_at 1.00791 "nuclear receptor subfamily 1, group I, member 2" Nr1i2
1425724_at 0.80627 "protein tyrosine phosphatase, receptor type, N polypeptide 2" Ptpn2
1425725_s_at 0.86584 "Protein phosphatase 2, regulatory subunit B (B56), gamma isoform"
Ppp2r5c
1425726_x_at 0.91973 "protein phosphatase 2, regulatory subunit B (B56), gamma isoform"
Ppp2r5c
1425727_at 0.99078 claudin 19 Cldn19
1425728_at 0.99320 hypothetical gene Rp23-14f5.7 RP23-14F5.7
1425729_at 0.99508 vitelliform macular dystrophy 2-like protein 1 Vmd211
1425730_at 0.99946 "calcium channel, voltage-dependent, gamma subunit 6" Cacng6
1425731_at 0.97740 ankyrin repeat domain 24 Ankrd24
1425732_a_at 0.86091 Max interacting protein 1 Mxi1
1425733_a_at 1.33040 epidermal growth factor receptor pathway substrate 8 Eps8
1425734_a_at 1.10138 coiled-coil domain containing 77 Ccdc77
1425735_at 1.10919 MAP-kinase activating death domain Madd
1425736_at 0.98994 CD37 antigen Cd37
1425737_at 0.93610 --- ---
1425738_at 0.96629 immunoglobulin kappa chain variable 21 (V21)-12 /// similar to Ig kappa chain
V-III region PC 6684 /// similar to Ig kappa chain V-III region PC 6684 Iggk-V21-12 /// LOC669090 ///
LOC676193
1425739_at 0.83244 phospholipase D1 Pld1
1425740_at 1.05550 --- ---
1425741_at 0.90214 SLIT-ROBO Rho GTPase activating protein 3 Srgap3
1425742_a_at 0.85033 "TSC22 domain family, member 1" Tsc22d1
1425743_at 0.99000 tripartite motif protein 7 Trim7

1425744_a_at 1.06318 mannoside acetylglucosaminyltransferase 5 Mgat5
 1425745_a_at 0.86391 "transforming, acidic coiled-coil containing protein 2" Tacc2
 1425746_at 1.18980 RIKEN cDNA D730039F16 gene D730039F16Rik
 1425747_at 1.01127 RIKEN cDNA 1110060D06 gene /// dedicator of cytokinesis 5
 1110060D06Rik /// Dock5
 1425748_at 0.91522 "DIRAS family, GTP-binding RAS-like 1" Diras1
 1425749_at 0.90182 syntaxin binding protein 6 (amisyn) Stxbp6
 1425750_a_at 0.97313 Janus kinase 3 Jak3
 1425751_at 1.01522 cDNA sequence BC014805 BC014805
 1425752_at 0.93338 cDNA sequence BC014805 BC014805
 1425753_a_at 0.90600 uracil DNA glycosylase Ung
 1425754_a_at 0.97919 "butyrophilin, subfamily 1, member A1" Btn1a1
 1425755_at 1.11308 mature T-cell proliferation 1 Mtcp1
 1425756_at 0.91139 "Rab40b, member RAS oncogene family" Rab40b
 1425757_a_at 1.09090 interphotoreceptor matrix proteoglycan 1 Impg1
 1425758_a_at 1.24992 interphotoreceptor matrix proteoglycan 1 Impg1
 1425759_at 0.84722 NOBOX oogenesis homeobox Nobox
 1425760_a_at 1.00337 phosphatidylinositol membrane-associated 1 Pitpm1
 1425761_a_at 1.00099 "nuclear factor of activated T-cells, cytoplasmic, calcineurin-dependent 1"
 Nfatc1
 1425762_a_at 0.90188 retinoid X receptor alpha Rxra
 1425763_x_at 1.01899 immunoglobulin heavy chain (J558 family) /// similar to immunoglobulin
 heavy chain variable region /// similar to immunoglobulin mu-chain /// similar to anti-poly(dC)
 monoclonal antibody heavy chain /// similar to Ig heavy chain V region 23 precursor /// similar to Ig
 heavy chain V region 102 precursor /// similar to Ig heavy chain V region VH558 A1/A4 precursor Igh-
 VJ558 /// LOC238447 /// LOC544903 /// LOC544907 /// LOC633774 /// LOC676478 /// LOC677563
 1425764_a_at 1.00759 "branched chain aminotransferase 2, mitochondrial" Bcat2
 1425765_at 1.09333 "LOC622757 gene /// similar to transcription elongation factor B (SIII),
 polypeptide 2" LOC622757 /// LOC676501
 1425766_x_at 1.10003 "LOC622757 gene /// similar to transcription elongation factor B (SIII),
 polypeptide 2" LOC622757 /// LOC676501
 1425767_a_at 1.20028 sine oculis-related homeobox 4 homolog (Drosophila) Six4
 1425768_at 1.03318 diablo homolog (Drosophila) Diablo
 1425769_x_at 1.09931 chemokine-like factor Cklf
 1425770_at 0.98877 RIKEN cDNA A730028G07 gene A730028G07Rik
 1425771_at 1.09963 "aldo-keto reductase family 1, member D1" Akr1d1
 1425772_at 0.52842 "procollagen, type IV, alpha 4" Col4a4
 1425773_s_at 1.01496 nicotinamide nucleotide adenylyltransferase 1 Nmnat1
 1425774_at 0.79961 RIKEN cDNA 1500001A10 gene 1500001A10Rik
 1425775_at 1.08546 hypothetical gene MGC29393 MGC29393
 1425776_a_at 0.87860 expressed sequence C87436 C87436
 1425777_at 1.02915 "calcium channel, voltage-dependent, beta 1 subunit" Cacnb1
 1425778_at 0.86815 RIKEN cDNA C230043N17 gene C230043N17Rik

1425779_a_at 1.02220 T-box 1 Tbx1
1425780_a_at 1.02186 transmembrane protein 167 Tmem167
1425781_a_at 1.03969 "phospholipase C, beta 1" Plcb1
1425782_at 0.93012 "phospholipase C, beta 1" Plcb1
1425783_at 1.02282 membrane targeting (tandem) C2 domain containing 1 Mtac2d1
1425784_a_at 0.95115 olfactomedin 1 Olfm1
1425785_a_at 0.91805 TXK tyrosine kinase Txk
1425786_a_at 1.05263 heat shock transcription factor 4 Hsf4
1425787_a_at 1.06568 synaptotagmin-like 3 Sytl3
1425788_a_at 0.88662 enoyl Coenzyme A hydratase domain containing 2 Echdc2
1425789_s_at 1.15201 annexin A8 Anxa8
1425790_a_at 0.98935 "glutamate receptor, ionotropic, kainate 2 (beta 2)" Grik2
1425791_at 1.03727 paraoxonase 2 Pon2
1425792_a_at 0.68771 RAR-related orphan receptor gamma Rorc
1425793_a_at 0.89542 RAR-related orphan receptor gamma Rorc
1425794_at 1.21755 "polymerase (DNA directed), alpha 2" Pola2
1425795_a_at 0.99633 mitogen activated protein kinase kinase kinase 7 Map3k7
1425796_a_at 0.73735 fibroblast growth factor receptor 3 Fgfr3
1425797_a_at 1.02483 spleen tyrosine kinase Syk
1425798_a_at 0.99655 RecQ protein-like Recql
1425799_at 1.04280 flavin containing monooxygenase 4 Fmo4
1425800_at 0.96612 RAD9 homolog B (S. cerevisiae) Rad9b
1425801_x_at 0.89727 coactosin-like 1 (Dictyostelium) Cotl1
1425802_a_at 0.93909 Fc receptor-like A Fcrla
1425803_a_at 0.97041 methyl-CpG binding domain protein 2 Mbd2
1425804_at 0.98575 H6 homeo box 2 Hmx2
1425805_a_at 0.92747 ubiquitin specific peptidase 12 Usp12
1425806_a_at 1.03226 SRB7 (suppressor of RNA polymerase B) homolog (S. cerevisiae) Surb7
1425807_at 0.92819 cDNA sequence BC021891 BC021891
1425808_a_at 0.99087 myocardin Myocd
1425809_at 0.94710 "fatty acid binding protein 4, adipocyte" Fabp4
1425810_a_at 0.80631 cysteine and glycine-rich protein 1 Csrp1
1425811_a_at 0.90912 cysteine and glycine-rich protein 1 Csrp1
1425812_a_at 0.97598 "calcium channel, voltage-dependent, N type, alpha 1B subunit" Cacna1b
1425813_at 0.94992 "phosphatidylinositol glycan anchor biosynthesis, class N" Pign
1425814_a_at 0.97056 calcitonin receptor-like Calcrl
1425815_a_at 0.94460 hyaluronan mediated motility receptor (RHAMM) Hmnr
1425816_at 0.87937 zinc finger protein 287 Zfp287
1425817_a_at 1.03225 "solute carrier family 8 (sodium/calcium exchanger), member 1" Slc8a1
1425818_at 0.97556 --- ---
1425819_at 0.96274 zinc finger and BTB domain containing 7C Zbtb7c
1425820_x_at 1.25619 G patch domain containing 4 Gpatc4
1425821_at 1.08227 chloride channel 7 Clcn7

1425822_a_at 1.06607 deltex 1 homolog (Drosophila) Dtx1
1425823_at 1.05915 cDNA sequence BC026782 /// similar to complement factor H-related protein B BC026782 /// LOC640602
1425824_a_at 1.06939 proprotein convertase subtilisin/kexin type 4 Pcsk4
1425825_at 0.93379 RIKEN cDNA C230094A16 gene C230094A16Rik
1425826_a_at 0.70146 sorbin and SH3 domain containing 1 Sorbs1
1425827_at 0.94143 "NK2 transcription factor related, locus 3 (Drosophila)" Nkx2-3
1425828_at 1.17334 "NK6 transcription factor related, locus 1 (Drosophila)" Nkx6-1
1425829_a_at 1.07973 STEAP family member 4 Steap4
1425830_a_at 0.98466 RIKEN cDNA 2810452K22 gene /// similar to cyclin-dependent kinase 2-interacting protein 2810452K22Rik /// LOC640972
1425831_at 1.18480 zinc finger protein 101 Zfp101
1425832_a_at 1.09413 chemokine (C-X-C motif) receptor 6 Cxcr6
1425833_a_at 0.90070 hippocalcin Hpca
1425834_a_at 0.95721 "glycerol-3-phosphate acyltransferase, mitochondrial" Gpam
1425835_a_at 0.99581 bobby sox homolog (Drosophila) Bbx
1425836_a_at 1.03482 "LIM-domain containing, protein kinase" Limk1
1425837_a_at 1.21401 CCR4 carbon catabolite repression 4-like (S. cerevisiae) Ccrn4l
1425838_at 1.23019 "ATPase, class II, type 9A" Atp9a
1425839_at 0.96183 FK506 binding protein 11 Fkbp11
1425840_a_at 0.79977 "sema domain, immunoglobulin domain (Ig), short basic domain, secreted, (semaphorin) 3 F" Sema3f
1425841_at 0.89698 "solute carrier family 26, member 7" Slc26a7
1425842_at 0.95557 EGF-like repeats and discoidin I-like domains 3 Edil3
1425843_at 0.97866 mitochondrial ribosomal protein L33 Mrpl33
1425844_a_at 0.91447 RNA guanylyltransferase and 5'-phosphatase Rngtt
1425845_a_at 0.91877 soc-2 (suppressor of clear) homolog (C. elegans) Shoc2
1425846_a_at 0.97437 calneuron 1 Caln1
1425847_a_at 0.81456 RIKEN cDNA B230120H23 gene B230120H23Rik
1425848_a_at 1.00971 dual specificity phosphatase 26 (putative) Dusp26
1425849_at 1.00087 "cholinergic receptor, nicotinic, beta polypeptide 4" Chrn4
1425850_a_at 1.09628 NIMA (never in mitosis gene a)-related expressed kinase 6 /// similar to NIMA (never in mitosis gene a)-related expressed kinase 6 Nek6 /// LOC674247
1425851_a_at 1.07203 adhesion molecule with Ig like domain 1 Amigo1
1425852_at 0.91353 RIKEN cDNA A230107C01 gene A230107C01Rik
1425853_s_at 1.00900 prolactin receptor Prlr
1425854_x_at 1.00429 "T-cell receptor beta, variable 13 /// similar to T-cell receptor beta-2 chain C region /// similar to T-cell receptor beta-1 chain C region /// similar to T-cell receptor beta-2 chain C region" Tcrb-V13 /// LOC665506 /// LOC669847 /// LOC674048
1425855_a_at 0.98715 v-crk sarcoma virus CT10 oncogene homolog (avian) Crk
1425856_at 0.95398 Calcium and integrin binding 1 (calmyrin) Cib1
1425857_at 1.11982 F-box and WD-40 domain protein 9 Fbxw9
1425858_at 0.96879 "ubiquitin-conjugating enzyme E2M (UBC12 homolog, yeast)" Ube2m

1425859_a_at 1.11525 "proteasome (prosome, macropain) 26S subunit, non-ATPase, 4" Psm4
 1425860_x_at 0.99462 chemokine-like factor Cklf
 1425861_x_at 1.09076 "calcium channel, voltage-dependent, alpha2/delta subunit 1" Cacna2d1
 1425862_a_at 0.92087 "phosphatidylinositol 3-kinase, C2 domain containing, alpha polypeptide"
 Pik3c2a
 1425863_a_at 0.90464 "protein tyrosine phosphatase, receptor type, O" Ptpro
 1425864_a_at 1.01108 VPS10 domain receptor protein SORCS 1 Sorcs1
 1425865_a_at 0.96990 "ligase III, DNA, ATP-dependent" Lig3
 1425866_a_at 0.93690 "pleckstrin homology domain containing, family A (phosphoinositide binding
 specific) member 4" Plekha4
 1425867_at 0.86579 "pleckstrin homology domain containing, family A (phosphoinositide binding
 specific) member 4" Plekha4
 1425868_at 0.99504 "Histone 2, H2bb" Hist2h2bb
 1425869_a_at 0.92496 presenilin 2 Psen2
 1425870_a_at 1.09993 Kv channel-interacting protein 2 Kcnip2
 1425871_a_at 1.23266 Similar to immunoglobulin light chain variable region LOC384413
 1425872_at 1.06341 keratin associated protein 6-1 Krtap6-1
 1425873_a_at 1.13365 leptin receptor Lepr
 1425874_at 1.02922 homeo box C13 Hoxc13
 1425875_a_at 1.06735 leptin receptor Lepr
 1425876_a_at 1.05394 glucuronyl C5-epimerase Glce
 1425877_at 0.97565 hyaluronoglucosaminidase 3 Hyal3
 1425878_at 1.23704 calcium binding protein 4 Cabp4
 1425879_at 0.93352 zinc finger protein 352 Zfp352
 1425880_x_at 1.02096 zinc finger protein 352 Zfp352
 1425881_at 1.37765 pregnancy-specific glycoprotein 28 Psg28
 1425882_at 1.03815 growth differentiation factor 2 Gdf2
 1425883_at 1.26300 "Smg-6 homolog, nonsense mediated mRNA decay factor (C. elegans)"
 Smg6
 1425884_at 0.82937 brix domain containing 1 Bxdc1
 1425885_a_at 0.94990 "potassium voltage-gated channel, shaker-related subfamily, beta member
 2" Kcnab2
 1425886_at 0.90363 FEV (ETS oncogene family) Fev
 1425887_at 0.95459 RIKEN cDNA 4930511J11 gene 4930511J11Rik
 1425888_at 0.99937 "killer cell lectin-like receptor, subfamily A, member 17" Klra17
 1425889_at 0.97892 wingless-type MMTV integration site 9A Wnt9a
 1425890_at 1.05671 "lymphocyte antigen 6 complex, locus I" Ly6i
 1425891_a_at 0.88365 GH regulated TBC protein 1 Grtp1
 1425892_a_at 0.96040 prepronociceptin Pnoc
 1425893_a_at 1.03030 fragile histidine triad gene Fhit
 1425894_at 0.95974 "MAS-related GPR, member F" Mrgprf
 1425895_a_at 0.59093 inhibitor of DNA binding 1 Id1
 1425896_a_at 0.94804 fibrillin 1 Fbn1

1425897_at 1.03648 baculoviral IAP repeat-containing 1c Birc1c
1425898_x_at 0.98291 olfactomedin 3 Olfm3
1425899_a_at 1.07390 intersectin 1 (SH3 domain protein 1A) Itsn1
1425900_at 0.90124 hexokinase domain containing 1 Hkdc1
1425901_at 0.89173 "nuclear factor of activated T-cells, cytoplasmic, calcineurin-dependent 2"
Nfatc2
1425902_a_at 1.19293 "nuclear factor of kappa light polypeptide gene enhancer in B-cells 2, p49/
p100" Nfkb2
1425903_at 0.83509 "sema domain, transmembrane domain (TM), and cytoplasmic domain,
(semaphorin) 6A" Sema6a
1425904_at 0.89872 special AT-rich sequence binding protein 2 Satb2
1425905_at 1.05908 --- ---
1425906_a_at 0.97435 "sema domain, immunoglobulin domain (Ig), short basic domain, secreted,
(semaphorin) 3E" Sema3e
1425907_s_at 0.93894 angiominin Amot
1425908_at 1.06860 "guanine nucleotide binding protein, beta 1" Gnb1
1425909_at 0.91369 CDNA clone MGC:11898 IMAGE:3598523 ---
1425910_at 0.99382 "DnaJ (Hsp40) homolog, subfamily C, member 2" Dnajc2
1425911_a_at 0.83846 fibroblast growth factor receptor 1 Fgfr1
1425912_at 0.97873 centrosomal protein 164 Cep164
1425913_a_at 0.94711 RIKEN cDNA 2810022L02 gene 2810022L02Rik
1425914_a_at 1.07891 "armadillo repeat containing, X-linked 1" Armcx1
1425915_at 1.02039 "solute carrier family 26, member 8" Slc26a8
1425916_at 0.87571 calpain 8 Capn8
1425917_at 1.00421 histocompatibility 28 H28
1425918_at 0.99729 EGL nine homolog 3 (C. elegans) Egl3
1425919_at 1.34488 "NADH dehydrogenase (ubiquinone) 1 alpha subcomplex, 12" Ndufa12
1425920_at 1.18437 CUE domain containing 1 Cuedc1
1425921_a_at 0.97825 RIKEN cDNA 1810055G02 gene 1810055G02Rik
1425922_a_at 0.76901 "v-myc myelocytomatosis viral related oncogene, neuroblastoma derived
(avian)" Mycn
1425923_at 0.98960 "v-myc myelocytomatosis viral related oncogene, neuroblastoma derived
(avian)" Mycn
1425924_at 0.91106 MyoD family inhibitor Mdf1
1425925_at 0.97977 "Fc receptor, IgA, IgM, high affinity" Fcgr2
1425926_a_at 1.10620 orthodenticle homolog 2 (Drosophila) Otx2
1425927_a_at 1.02506 activating transcription factor 5 Atf5
1425928_at 0.96559 X Kell blood group precursor related family member 6 homolog Xkr6
1425929_a_at 1.05548 ring finger protein 14 Rnf14
1425930_a_at 1.01636 MAX-like protein X Mlx
1425931_a_at 0.97370 aryl hydrocarbon receptor nuclear translocator-like 2 Arntl2
1425932_a_at 1.15759 "CUG triplet repeat, RNA binding protein 1" Cugbp1
1425933_a_at 0.93241 "5'-nucleotidase, cytosolic II" Nt5c2

1425934_a_at 1.10252 "UDP-Gal:betaGlcNAc beta 1,4-galactosyltransferase, polypeptide 4"
 B4galt4
 1425935_at 1.04269 RIKEN cDNA 2900042B11 gene 2900042B11Rik
 1425936_a_at 1.17396 ankyrin repeat and MYND domain containing 2 Ankmy2
 1425937_a_at 1.02441 hexamethylene bis-acetamide inducible 1 Hexim1
 1425938_a_at 1.00593 FK506 binding protein 11 Fkbp11
 1425939_at 0.99341 RAD50 homolog (*S. cerevisiae*) Rad50
 1425940_a_at 0.73683 single-stranded DNA binding protein 3 Ssbp3
 1425941_a_at 1.03225 cDNA sequence BC025462 BC025462
 1425942_a_at 0.94264 glycoprotein m6b Gpm6b
 1425943_at 0.89085 neuromedin U receptor 2 Nmur2
 1425944_a_at 0.91759 RAD51-like 3 (*S. cerevisiae*) Rad51l3
 1425945_at 1.04178 RIKEN cDNA 4933426I21 gene 4933426I21Rik
 1425946_at 1.02876 "glutathione S-transferase, mu 7" Gstm7
 1425947_at 0.92885 interferon gamma Ifng
 1425948_a_at 0.92660 "solute carrier family 25, member 30" Slc25a30
 1425949_at 0.98442 "solute carrier family 25, member 30" Slc25a30
 1425950_at 0.99556 cDNA sequence BC019537 BC019537
 1425951_a_at 0.98575 "C-type lectin domain family 4, member n" Clec4n
 1425952_a_at 1.08466 glucagon Gcg
 1425953_at 0.97000 Dynein cytoplasmic 1 light intermediate chain 1 Dync1li1
 1425954_a_at 0.96549 apurinic/aprimidinic endonuclease 2 Apex2
 1425955_at 1.03859 caveolin 2 Cav2
 1425956_a_at 0.91398 cytidine and dCMP deaminase domain containing 1 Cdadc1
 1425957_x_at 0.91624 RIKEN cDNA 1700021K02 gene 1700021K02Rik
 1425958_at 0.89861 "interleukin 1 family, member 9" Il1f9
 1425959_x_at 1.02522 "killer cell lectin-like receptor, subfamily A, member 1 /// killer cell lectin-like
 receptor subfamily A, member 12 /// killer cell lectin-like receptor subfamily A, member 13 /// killer cell
 lectin-like receptor, subfamily A, member 4 /// killer cell lectin-like receptor, subfamily A, member
 15 /// killer cell lectin-like receptor, subfamily A, member 16 /// killer cell lectin-like receptor subfamily
 A, member 23 /// killer cell lectin-like receptor subfamily A, member 20 /// killer cell lectin-like receptor
 subfamily A, member 22 /// killer cell lectin-like receptor, subfamily A, member 18 /// similar to Killer
 cell lectin-like receptor 4 (T-cell surface glycoprotein Ly-49D) (Ly49-D antigen) (Lymphocyte antigen
 49D) /// Ly49p/d /// similar to killer cell lectin-like receptor, subfamily A, member 16 /// killer cell lectin-
 like receptor subfamily A member 33 /// similar to killer cell lectin-like receptor, subfamily A, member
 18 /// similar to killer cell lectin-like receptor subfamily A, member 13" Klra1 /// Klra12 /// Klra13 ///
 Klra4 /// Klra15 /// Klra16 /// Klra23 /// Klra20 /// Klra22 /// Klra18 /// LOC634585 /// LOC635513 ///
 LOC640058 /// Klra33 /// LOC667764 /// LOC673407
 1425960_s_at 0.97339 paired box gene 6 Pax6
 1425961_at 0.99493 cDNA sequence BC016548 BC016548
 1425962_at 0.96310 killer cell lectin-like receptor subfamily B member 1F Klrb1f
 1425963_at 0.86621 calcium binding protein 7 Cabp7
 1425964_x_at 0.66481 heat shock protein 1 Hspb1

1425965_at 0.79839 ubiquitin C Ubc
1425966_x_at 0.84548 ubiquitin C Ubc
1425967_a_at 0.88526 mast cell protease 4 Mcpt4
1425968_s_at 0.86393 SPEG complex locus Speg
1425969_a_at 1.07246 Huntington disease gene homolog Hdh
1425970_a_at 1.12079 Ros1 proto-oncogene Ros1
1425971_at 1.03740 (ORF 1) and reverse transcriptase (ORF 2) genes ---
1425972_a_at 1.04398 zinc finger protein X-linked Zfx
1425973_at 0.90382 lysosomal trafficking regulator Lyst
1425974_a_at 1.05397 tripartite motif protein 25 Trim25
1425975_a_at 0.90548 mitogen-activated protein kinase 8 interacting protein 3 Mapk8ip3
1425976_x_at 1.04831 zinc finger protein 353 Zfp353
1425977_a_at 1.02545 STE20-like kinase (yeast) Slk
1425978_at 0.90876 myocardin Myocd
1425979_a_at 0.88116 Fas (TNFRSF6) binding factor 1 Fbf1
1425980_at 0.98617 WD repeat domain 54 Wdr54
1425981_a_at 0.96789 retinoblastoma-like 2 /// similar to retinoblastoma-like 2 Rbl2 ///
LOC635075
1425982_a_at 0.88776 Werner syndrome homolog (human) Wrn
1425983_x_at 0.76888 homeodomain interacting protein kinase 2 Hipk2
1425984_at 1.01171 "procollagen, type V, alpha 3" Col5a3
1425985_s_at 0.62027 mannan-binding lectin serine peptidase 1 Masp1
1425986_a_at 0.89480 "DCUN1D1 DCN1, defective in cullin neddylation 1, domain containing 1 (S. cerevisiae)" Dcun1d1
1425987_a_at 0.93409 "potassium large conductance calcium-activated channel, subfamily M, alpha member 1" Kcnma1
1425988_a_at 0.88695 homeodomain interacting protein kinase 1 /// similar to homeodomain-interacting protein kinase 1 Hipk1 /// LOC634033
1425989_a_at 1.05407 eyes absent 3 homolog (Drosophila) Eya3
1425990_a_at 0.98867 "nuclear factor of activated T-cells, cytoplasmic, calcineurin-dependent 2" Nfatc2
1425991_a_at 0.95752 ankyrin repeat domain 25 Ankrd25
1425992_at 0.88162 "solute carrier family 6 (neurotransmitter transporter, glycine), member 5" Slc6a5
1425993_a_at 0.94915 heat shock protein 110 Hsp110
1425994_a_at 0.99179 N-acylsphingosine amidohydrolase 2 Asah2
1425995_s_at 1.06524 Wilms tumor homolog Wt1
1425996_a_at 0.77872 "SWI/SNF related, matrix associated, actin dependent regulator of chromatin, subfamily a, member 3" Smarca3
1425997_a_at 0.93286 "phosphatidylinositol glycan anchor biosynthesis, class N" Pign
1425998_at 0.96678 synaptotagmin-like 4 Sytl4
1425999_at 0.91980 complement factor H-related protein LOC214403
1426000_at 0.91712 oxytocin receptor Oxt

1426001_at	1.00497	eomesodermin homolog (<i>Xenopus laevis</i>)	Eomes
1426002_a_at	1.11076	cell division cycle 7 (<i>S. cerevisiae</i>)	Cdc7
1426003_at	0.96405	"neurotrophic tyrosine kinase, receptor, type 3"	Ntrk3
1426004_a_at	0.89870	"transglutaminase 2, C polypeptide"	Tgm2
1426005_at	0.91215	dentin matrix protein 1	Dmp1
1426006_at	0.94409	"potassium voltage-gated channel, subfamily Q, member 2"	Kcnq2
1426007_a_at	0.97615	UBX domain containing 3	Ubx3
1426008_a_at	0.90804	"solute carrier family 7 (cationic amino acid transporter, y+ system), member 2"	Slc7a2
1426009_a_at	1.15056	"phosphatidylinositol-4-phosphate 5-kinase, type 1 beta"	Pip5k1b
1426010_a_at	0.92459	erythrocyte protein band 4.1-like 3	Epb4.113
1426011_a_at	1.08917	gametogenetin binding protein 2	Ggnbp2
1426012_a_at	0.98994	RIKEN cDNA 2610301G19 gene	2610301G19Rik
1426013_s_at	1.00864	"pleckstrin homology domain containing, family A (phosphoinositide binding specific) member 4"	Plekha4
1426014_a_at	1.09084	mucin and cadherin like	Mucdhl
1426015_s_at	0.86115	aspartate-beta-hydroxylase	Asph
1426016_a_at	1.12696	trophinin	Tro
1426017_a_at	1.05971	RIKEN cDNA 0610011L14 gene	0610011L14Rik
1426018_a_at	1.01780	SRY-box containing gene 6	Sox6
1426019_at	1.04757	"phospholipase A2, activating protein"	Plaa
1426020_at	1.06205	thymopoietin	Tmpo
1426021_a_at	0.98423	cell division cycle 7 (<i>S. cerevisiae</i>)	Cdc7
1426022_a_at	0.97996	villin-like	Vill
1426023_a_at	1.01873	"rabaptin, RAB GTPase binding effector protein 1"	Rabep1
1426024_a_at	0.80027	drebrin 1	Dbn1
1426025_s_at	0.97043	lysosomal-associated protein transmembrane 5	Laptn5
1426026_at	0.97183	PRP6 pre-mRNA splicing factor 6 homolog (yeast)	Prpf6
1426027_a_at	0.83080	Rho GTPase activating protein 10	Arhgap10
1426028_a_at	0.77895	citron	Cit
1426029_a_at	0.81925	nuclear factor of activated T-cells 5	Nfat5
1426030_a_at	0.86622	acylpeptide hydrolase	Apeh
1426031_a_at	0.99396	"nuclear factor of activated T-cells, cytoplasmic, calcineurin-dependent 2"	Nfatc2
1426032_at	0.91166	"nuclear factor of activated T-cells, cytoplasmic, calcineurin-dependent 2"	Nfatc2
1426033_at	0.95767	regulator of G-protein signaling 9	Rgs9
1426034_a_at	1.05744	runt related transcription factor 2	Runx2
1426035_at	0.97732	hypothetical LOC574418	LOC574418
1426036_a_at	0.95547	phosphodiesterase 1C	Pde1c
1426037_a_at	1.03357	regulator of G-protein signaling 16	Rgs16
1426038_at	1.04894	---	---
1426039_a_at	0.88381	"arachidonate lipoxygenase, epidermal"	Alox12e

1426040_a_at 1.16643 outer dense fiber of sperm tails 2 Odf2
 1426041_a_at 0.96794 "FYVE, RhoGEF and PH domain containing 4" Fgd4
 1426042_at 0.88391 "FYVE, RhoGEF and PH domain containing 4" Fgd4
 1426043_a_at 0.75751 calpain 3 Capn3
 1426044_a_at 0.85746 "protein kinase C, theta" Prkcq
 1426045_at 0.93461 kininogen 1 Kng1
 1426046_a_at 1.19172 "Rab geranylgeranyl transferase, a subunit" Rabggta
 1426047_a_at 1.00580 "protein tyrosine phosphatase, receptor type, R" Ptprr
 1426048_s_at 0.97574 "histidine rich calcium binding protein /// transcription factor AP-2, alpha"
 Hrc /// Tcfap2a
 1426049_a_at 0.95820 --- ---
 1426050_at 0.89451 Bcl2-like 1 Bcl2l1
 1426051_a_at 1.16510 centromere protein B Cenpb
 1426052_at 0.83128 mutL homolog 3 (E coli) Mlh3
 1426053_a_at 0.92419 xenotropic and polytropic retrovirus receptor 1 Xpr1
 1426054_at 0.96368 neuropeptide Y receptor Y1 Npy1r
 1426055_a_at 0.96083 "phosphatidylinositol glycan anchor biosynthesis, class Q" Pigq
 1426056_at 1.01594 "phosphatidylinositol glycan anchor biosynthesis, class Q" Pigq
 1426057_a_at 0.92362 Eph receptor A3 Epha3
 1426058_a_at 0.98480 "potassium channel, subfamily K, member 3" Kcnk3
 1426059_at 0.94052 glucokinase regulatory protein Gckr
 1426060_at 1.26939 --- ---
 1426061_x_at 1.13292 --- ---
 1426062_a_at 0.91115 caspase 7 Casp7
 1426063_a_at 0.89126 GTP binding protein (gene overexpressed in skeletal muscle) Gem
 1426064_at 1.01033 "cytochrome P450, family 3, subfamily a, polypeptide 44" Cyp3a44
 1426065_a_at 1.27316 tribbles homolog 3 (Drosophila) Trib3
 1426066_a_at 1.01429 dystrobrevin alpha Dtna
 1426067_x_at 1.00094 chaperonin subunit 3 (gamma) Cct3
 1426068_at 0.91185 "solute carrier family 7 (cationic amino acid transporter, y+ system), member
 4" Slc7a4
 1426069_s_at 0.88588 "solute carrier family 7 (cationic amino acid transporter, y+ system), member
 4" Slc7a4
 1426070_a_at 0.82684 "potassium voltage-gated channel, Shal-related family, member 3" Kcnd3
 1426071_at 1.05082 TGF-beta1-induced anti-apoptotic factor 2 Tiaf2
 1426072_at 0.99982 chemokine-like receptor 1 Cmk1r1
 1426073_at 1.07896 twisted gastrulation homolog 1 (Drosophila) Twsg1
 1426074_at 0.93327 LOC434536 LOC434536
 1426075_at 1.03199 klotho beta Klb
 1426076_at 0.87250 cell division cycle 27 homolog (S. cerevisiae) Cdc27
 1426077_at 0.89733 "monooxygenase, DBH-like 2" Moxd2
 1426078_a_at 0.91812 G protein-coupled receptor 108 Gpr108
 1426079_at 1.01922 CD247 antigen Cd247

1426080_a_at 0.90787 "potassium voltage-gated channel, subfamily Q, member 2" Kcnq2
 1426081_a_at 1.07821 "deiodinase, iodothyronine, type II" Dio2
 1426082_a_at 0.85967 "solute carrier family 16 (monocarboxylic acid transporters), member 4"
 Slc16a4
 1426083_a_at 0.92854 "B-cell translocation gene 1, anti-proliferative" Btg1
 1426084_a_at 0.97467 torsin A interacting protein 1 Tor1aip1
 1426085_a_at 0.90406 paxillin Pxn
 1426086_a_at 0.92063 fragile X mental retardation syndrome 1 homolog Fmr1
 1426087_at 1.11769 icos ligand Icosl
 1426088_at 1.05654 --- ---
 1426089_a_at 0.98042 cDNA sequence BC003331 BC003331
 1426090_a_at 1.09303 "fer (fms/fps related) protein kinase, testis specific 2" Fert2
 1426091_a_at 1.04075 wingless related MMTV integration site 10b Wnt10b
 1426092_a_at 0.91797 tripartite motif protein 34 Trim34
 1426093_at 0.93829 tripartite motif protein 34 Trim34
 1426094_at 1.02636 "rhomboid, veinlet-like 1 (Drosophila)" Rhbdl1
 1426095_a_at 1.21120 "tumor necrosis factor receptor superfamily, member 22" Tnfrsf22
 1426096_at 0.90431 protein arginine N-methyltransferase 7 Prmt7
 1426097_a_at 1.00673 coiled-coil domain containing 106 Ccdc106
 1426098_a_at 1.08072 calpastatin Cast
 1426099_at 1.01880 histamine H4 receptor Hrh4
 1426100_a_at 0.91994 "thymidine kinase 2, mitochondrial" Tk2
 1426101_at 0.99525 ORFs (putative primordial protein) ---
 1426102_at 0.95679 "cytochrome P450, family 2, subfamily j, polypeptide 13" Cyp2j13
 1426103_a_at 0.98455 estrogen receptor 2 (beta) Esr2
 1426104_at 0.98759 mitogen activated protein kinase 14 Mapk14
 1426105_a_at 0.88600 UBX domain containing 5 Ubx5
 1426106_a_at 0.98337 synaptotagmin VI Syt6
 1426107_at 0.93580 VPR domain containing 9 Prdm9
 1426108_s_at 0.95606 "calcium channel, voltage-dependent, beta 1 subunit" Cacnb1
 1426109_a_at 1.03233 "solute carrier family 14 (urea transporter), member 2" Slc14a2
 1426110_a_at 0.93991 "endothelial differentiation, lysophosphatidic acid G-protein-coupled receptor,
 2" Edg2
 1426111_x_at 0.90772 interferon regulatory factor 3 Irf3
 1426112_a_at 0.90187 CD72 antigen Cd72
 1426113_x_at 0.89445 T-cell receptor alpha chain /// RIKEN cDNA A430107P09 gene Tcra ///
 A430107P09Rik
 1426114_at 0.93748 heterogeneous nuclear ribonucleoprotein A/B Hnripab
 1426115_a_at 0.83965 "potassium inwardly-rectifying channel, subfamily J, member 9" Kcnj9
 1426116_at 0.91350 ankyrin repeat and SOCS box-containing 18 Asb18
 1426117_a_at 1.18921 "solute carrier family 19 (thiamine transporter), member 2" Slc19a2
 1426118_a_at 1.07194 translocase of outer mitochondrial membrane 40 homolog (yeast) Tomm40
 1426119_at 1.00590 "thymidylate synthase, pseudogene" Tyms-ps

1426120_a_at 1.04545 CD244 natural killer cell receptor 2B4 /// similar to Natural killer cell receptor 2B4 precursor (NKR2B4) (NK cell type I receptor protein 2B4) (CD244 antigen) (Non MHC restricted killing associated) Cd244 /// LOC677008
 1426121_at 0.86115 "MAS-related GPR, member A3" Mrgpra3
 1426122_a_at 0.77711 "coronin, actin binding protein 6" Coro6
 1426123_a_at 1.32850 ribosome binding protein 1 Rrbp1
 1426124_a_at 1.17107 CDC-like kinase 1 Clk1
 1426125_a_at 0.88017 caspase 9 Casp9
 1426126_a_at 0.90701 hypothetical protein MGC30332 MGC30332
 1426127_x_at 0.91012 "killer cell lectin-like receptor, subfamily A, member 18" Klra18
 1426128_a_at 1.00473 "potassium voltage-gated channel, subfamily Q, member 2" Kcnq2
 1426129_at 1.04544 breast cancer metastasis-suppressor 1 Brms1
 1426130_at 0.94021 thioredoxin-like 6 Txnl6
 1426131_at 1.03033 RIKEN cDNA 1700036D21 gene 1700036D21Rik
 1426132_at 0.98008 --- ---
 1426133_a_at 1.20169 "MIT, microtubule interacting and transport, domain containing 1" Mitd1
 1426134_at 1.01093 triadin Trdn
 1426135_a_at 0.97505 parkin Park2
 1426136_x_at 0.94057 "killer cell lectin-like receptor subfamily A, member 21" Klra21
 1426137_at 1.04703 reduced expression 2 Rex2
 1426138_a_at 1.04319 "ubiquitin-conjugating enzyme E2, J2 homolog (yeast)" Ube2j2
 1426139_a_at 1.02523 chemokine (C-C motif) receptor-like 1 Ccr1
 1426140_x_at 1.00907 "killer cell lectin-like receptor, subfamily A, member 19" Klra19
 1426141_at 0.89558 "MAS-related GPR, member A1" Mrgpra1
 1426142_a_at 1.07823 triadin Trdn
 1426143_at 1.03493 triadin Trdn
 1426144_x_at 0.94525 triadin Trdn
 1426145_at 0.91442 --- ---
 1426146_a_at 0.96269 choline phosphotransferase 1 Chpt1
 1426147_s_at 0.96864 claudin 10 Cldn10
 1426148_at 0.81040 "globoside alpha-1,3-N-acetylgalactosaminyltransferase 1" Gbgt1
 1426149_at 0.90328 serine protease inhibitor 15 Spi15
 1426150_at 0.91665 "GIPC PDZ domain containing family, member 3" Gipc3
 1426151_a_at 1.04886 syntaxin 3 Stx3
 1426152_a_at 0.99376 kit ligand Kitl
 1426153_a_at 0.99219 desmoglein 2 Dsg2
 1426154_s_at 0.99349 major urinary protein 1 /// major urinary protein 3 Mup1 /// Mup3
 1426155_a_at 1.01329 odd-skipped related 2 (Drosophila) Osr2
 1426156_at 0.95039 large tumor suppressor 2 Lats2
 1426157_a_at 0.91244 CD209b antigen Cd209b
 1426158_at 0.99925 "T-cell receptor beta, variable 13" Tcrb-V13
 1426159_x_at 1.01651 "T-cell receptor beta, variable 13" Tcrb-V13
 1426160_a_at 0.96510 serine/threonine kinase 16 Stk16

1426161_at 1.03299 --- ---
1426162_a_at 1.02169 ribosomal protein L7 Rpl7
1426163_x_at 1.04635 ribosomal protein L7 Rpl7
1426164_a_at 0.87659 upstream transcription factor 1 Usf1
1426165_a_at 1.00470 caspase 3 Casp3
1426166_at 0.80472 major urinary protein 5 Mup5
1426167_a_at 0.91533 calcium/calmodulin-dependent protein kinase IV Camk4
1426168_a_at 0.85941 T-cell receptor alpha chain /// RIKEN cDNA A130082M07 gene /// RIKEN
cDNA A430107P09 gene Tcra /// A130082M07Rik /// A430107P09Rik
1426169_a_at 1.02388 "linker for activation of T cells family, member 2" Lat2
1426170_a_at 0.93650 "CD8 antigen, beta chain 1" Cd8b1
1426171_x_at 0.93778 "killer cell lectin-like receptor, subfamily A, member 7" Klra7
1426172_a_at 1.00460 CD209a antigen Cd209a
1426173_at 1.07672 "Cytochrome P450, family 51" Cyp51
1426174_s_at 1.07094 Immunoglobulin heavy chain (gamma polypeptide) Ighg
1426175_a_at 1.01153 tryptase alpha/beta 1 Tpsab1
1426176_a_at 0.93874 prokineticin 2 Prok2
1426177_a_at 1.00504 "NK1 transcription factor related, locus 2 (Drosophila)" Nkx1-2
1426178_at 1.07317 Immunoglobulin kappa chain complex ---
1426179_a_at 0.81129 twisted gastrulation homolog 1 (Drosophila) Twsg1
1426180_a_at 0.92167 submaxillary gland androgen regulated protein 2 Smr2
1426181_a_at 1.08015 interleukin 24 Il24
1426182_a_at 1.03229 "killer cell lectin-like receptor subfamily C, member 1" Klrc1
1426183_a_at 0.86372 CD209d antigen Cd209d
1426184_a_at 1.00736 programmed cell death 6 interacting protein Pdc6ip
1426185_at 1.03535 "calcium channel, voltage-dependent, alpha 2/delta subunit 2" Cacna2d2
1426186_a_at 0.94442 fibroblast growth factor 5 Fgf5
1426187_a_at 1.11026 HCLS1 associated X-1 Hax1
1426188_s_at 0.92688 cDNA sequence BC005685 BC005685
1426189_at 1.08267 dual specificity phosphatase-like 15 Dusp15
1426190_at 1.08436 activator of yeast meiotic promoters 1 Aym1
1426191_a_at 0.87845 Bcl2-like 1 Bcl2l1
1426192_at 0.81738 "SWI/SNF related, matrix associated, actin dependent regulator of chromatin,
subfamily d, member 2" Smardc2
1426193_at 1.07084 otospiralin Otos
1426194_x_at 0.94966 "REV3-like, catalytic subunit of DNA polymerase zeta RAD54 like (S.
cerevisiae)" Rev3l
1426195_a_at 0.87681 cystatin C Cst3
1426196_at 0.89371 Immunoglobulin heavy chain 1a (serum IgG2a) Igh-1a
1426197_at 1.06906 Anti-human CD20 antibody 1F5 gamma heavy chain variable region ---
1426198_at 1.16510 Anti-human CD37 antibody WR17 gamma heavy chain variable region ---
1426199_x_at 0.86434 Anti-human CD37 antibody WR17 gamma heavy chain variable region ---
1426200_at 0.97730 Immunoglobulin kappa chain complex ---

1426201_at 0.83167 "Gene model 1499, (NCBI)" Gm1499
1426202_at 0.99967 "Histocompatibility 2, class II antigen E alpha /// MHC class II E-sub-alpha (E-sub-alpha) mRNA, 5' flanking region and partial cds" H2-Ea
1426203_at 1.00215 keratin associated protein 16-4 Krtap16-4
1426204_a_at 1.03235 opioid receptor-like 1 Oprl1
1426205_at 0.93802 "protein phosphatase 1, catalytic subunit, beta isoform" Ppp1cb
1426206_at 1.03780 roundabout homolog 4 (Drosophila) Robo4
1426207_at 0.96696 inhibitor of kappaB kinase beta Ikbkb
1426208_x_at 1.11247 pleiomorphic adenoma gene-like 1 Plagl1
1426209_at 0.95446 "striatin, calmodulin binding protein 4" Strn4
1426210_x_at 1.12749 "poly (ADP-ribose) polymerase family, member 3" Parp3
1426211_at 0.96530 transmembrane protein 161A Tmem161a
1426212_s_at 0.94058 transmembrane protein 161A Tmem161a
1426213_at 0.98378 "IMP4, U3 small nucleolar ribonucleoprotein, homolog (yeast)" Imp4
1426214_at 0.99435 "IMP4, U3 small nucleolar ribonucleoprotein, homolog (yeast)" Imp4
1426215_at 0.66404 dopa decarboxylase Ddc
1426216_at 0.97097 component of oligomeric golgi complex 6 Cog6
1426217_at 0.94211 RIKEN cDNA 2810441K11 gene 2810441K11Rik
1426218_at 0.89092 glucocorticoid induced transcript 1 Glcci1
1426219_at 0.97599 "sterol carrier protein 2, liver" Scp2
1426220_at 0.98137 RIKEN cDNA 4930471M23 gene 4930471M23Rik
1426221_at 0.85343 "loss of heterozygosity, 11, chromosomal region 2, gene A homolog (human)" Loh11cr2a
1426222_s_at 0.97735 "loss of heterozygosity, 11, chromosomal region 2, gene A homolog (human)" Loh11cr2a
1426223_at 0.83999 RIKEN cDNA 2810439F02 gene 2810439F02Rik
1426224_x_at 1.03324 CKLF-like MARVEL transmembrane domain containing 2A Cmtm2a
1426225_at 0.95888 "retinol binding protein 4, plasma" Rbp4
1426226_at 0.90766 dual-specificity tyrosine-(Y)-phosphorylation regulated kinase 1a Dyrk1a
1426227_s_at 0.96178 vacuolar protein sorting 37C (yeast) Vps37c
1426228_at 0.91182 v-Ki-ras2 Kirsten rat sarcoma viral oncogene homolog Kras
1426229_s_at 0.83413 v-Ki-ras2 Kirsten rat sarcoma viral oncogene homolog Kras
1426230_at 1.06724 sphingosine kinase 2 Sphk2
1426231_at 0.98318 vitrin Vit
1426232_at 1.05844 cDNA sequence BC024479 BC024479
1426233_at 0.89560 mitogen activated protein kinase kinase 4 Map2k4
1426234_s_at 1.08862 cDNA sequence BC002199 BC002199
1426235_a_at 0.82473 glutamate-ammonia ligase (glutamine synthetase) Glul
1426236_a_at 0.79571 glutamate-ammonia ligase (glutamine synthetase) Glul
1426237_at 0.91376 Sp2 transcription factor Sp2
1426238_at 0.92478 bone morphogenetic protein 1 Bmp1
1426239_s_at 1.01894 "arrestin, beta 2" Arrb2
1426240_at 1.01537 chromatin modifying protein 4B Chmp4b

1426241_a_at 0.87201 sex comb on midleg homolog 1 Scmh1
 1426242_at 0.98769 polymerase (RNA) II (DNA directed) polypeptide A Polr2a
 1426243_at 0.98429 cystathionase (cystathionine gamma-lyase) Cth
 1426244_at 1.00018 "microtubule-associated protein, RP/EB family, member 2" Mapre2
 1426245_s_at 0.89802 "microtubule-associated protein, RP/EB family, member 2" Mapre2
 1426246_at 0.84940 protein S (alpha) Pros1
 1426247_at 0.97275 "serine/threonine kinase 24 (STE20 homolog, yeast)" Stk24
 1426248_at 1.02049 "serine/threonine kinase 24 (STE20 homolog, yeast)" Stk24
 1426249_at 1.02586 "adrenergic receptor kinase, beta 1" Adrbk1
 1426250_s_at 1.09769 RIKEN cDNA 9130404D08 gene 9130404D08Rik
 1426251_at 1.01002 carboxypeptidase Z Cpz
 1426252_a_at 0.92840 uromodulin Umod
 1426253_at 1.03846 RIKEN cDNA 4933428G09 gene 4933428G09Rik
 1426254_at 0.88445 TM2 domain containing 1 Tm2d1
 1426255_at 0.96840 "neurofilament, light polypeptide" Nefl
 1426256_at 1.15073 translocase of inner mitochondrial membrane 17a Timm17a
 1426257_a_at 1.05364 seryl-aminoacyl-tRNA synthetase Sars
 1426258_at 0.66613 "sortilin-related receptor, LDLR class A repeats-containing" Sorl1
 1426259_at 1.03513 pantothenate kinase 3 Pank3
 1426260_a_at 0.88308 "UDP glucuronosyltransferase 1 family, polypeptide A2 /// UDP
 glucuronosyltransferase 1 family, polypeptide A6A /// UDP glycosyltransferase 1 family, polypeptide
 A10 /// UDP glucuronosyltransferase 1 family, polypeptide A7C /// UDP glucuronosyltransferase 1
 family, polypeptide A5 /// UDP glucuronosyltransferase 1 family, polypeptide A9 /// UDP
 glucuronosyltransferase 1 family, polypeptide A6B /// UDP glucuronosyltransferase 1 family,
 polypeptide A1" Ugt1a2 /// Ugt1a6a /// Ugt1a10 /// Ugt1a7c /// Ugt1a5 /// Ugt1a9 /// Ugt1a6b /// Ugt1a1
 1426261_s_at 0.84020 "UDP glucuronosyltransferase 1 family, polypeptide A2 /// UDP
 glucuronosyltransferase 1 family, polypeptide A6A /// UDP glycosyltransferase 1 family, polypeptide
 A10 /// UDP glucuronosyltransferase 1 family, polypeptide A7C /// UDP glucuronosyltransferase 1
 family, polypeptide A5 /// UDP glucuronosyltransferase 1 family, polypeptide A9 /// UDP
 glucuronosyltransferase 1 family, polypeptide A6B /// UDP glucuronosyltransferase 1 family,
 polypeptide A1" Ugt1a2 /// Ugt1a6a /// Ugt1a10 /// Ugt1a7c /// Ugt1a5 /// Ugt1a9 /// Ugt1a6b /// Ugt1a1
 1426262_at 0.89352 zinc finger protein 508 Zfp508
 1426263_at 0.82574 "immunoglobulin superfamily, member 4C" Igsf4c
 1426264_at 0.89223 dihydrolipoamide S-acetyltransferase (E2 component of pyruvate
 dehydrogenase complex) Dlat
 1426265_x_at 0.90323 dihydrolipoamide S-acetyltransferase (E2 component of pyruvate
 dehydrogenase complex) Dlat
 1426266_s_at 1.21370 zinc finger and BTB domain containing 8 opposite strand Zbtb8os
 1426267_at 0.99131 zinc finger and BTB domain containing 8 opposite strand Zbtb8os
 1426268_at 0.87957 RIKEN cDNA C130090K23 gene C130090K23Rik
 1426269_at 0.89851 synaptobrevin like 1 Sybl1
 1426270_at 1.18656 structural maintenance of chromosomes 5 Smc5
 1426271_at 1.14754 structural maintenance of chromosomes 5 Smc5

1426272_at	0.98773	limb region 1	Lmbr1	
1426273_at	0.89026	limb region 1	Lmbr1	
1426274_at	1.04412	"solute carrier family 9 (sodium/hydrogen exchanger), member 8"	Slc9a8	
1426275_a_at	1.04675	UDP-glucuronate decarboxylase 1	Uxs1	
1426276_at	0.79714	interferon induced with helicase C domain 1	Ifih1	
1426277_at	1.02391	RIKEN cDNA C730025P13 gene	C730025P13Rik	
1426278_at	1.03642	"interferon, alpha-inducible protein 27"	Ifi27	
1426279_at	1.07104	RIKEN cDNA 5830415L20 gene	5830415L20Rik	
1426280_at	0.93945	expressed sequence AV249152	AV249152	
1426281_at	1.12953	cation channel of sperm 1	Catsper1	
1426282_at	1.02347	neurotrimin	Hnt	
1426283_at	1.04135	neurotrimin	Hnt	
1426284_at	0.98378	keratin 20	Krt20	
1426285_at	0.89333	"laminin, alpha 2"	Lama2	
1426286_at	1.18855	nucleolar complex associated 3 homolog (S. cerevisiae)	Noc3l	
1426287_at	1.00385	ataxin 7	Atxn7	
1426288_at	1.04142	low density lipoprotein receptor-related protein 4	Lrp4	
1426289_at	0.94642	RIKEN cDNA 2610028H07 gene	2610028H07Rik	
1426290_at	1.04239	DIM1 dimethyladenosine transferase 1-like (S. cerevisiae)	Dimt1	
1426291_at	1.03436	DIM1 dimethyladenosine transferase 1-like (S. cerevisiae)	Dimt1	
1426292_at	0.88777	RIKEN cDNA 6330581L23 gene	6330581L23Rik	
1426293_at	0.82467	RIKEN cDNA 6330581L23 gene	6330581L23Rik	
1426294_at	0.95627	hyaluronan and proteoglycan link protein 1	Hapln1	
1426295_at	0.93425	hyaluronan and proteoglycan link protein 1	Hapln1	
1426296_at	1.03713	RAD52 homolog (S. cerevisiae)	Rad52	
1426297_at	0.95061	transcription factor E2a	Tcfe2a	
1426298_at	0.91807	Iroquois related homeobox 2 (Drosophila)	Irx2	
1426299_at	0.93432	RIKEN cDNA 9130017C17 gene	9130017C17Rik	
1426300_at	0.90433	activated leukocyte cell adhesion molecule	Alcam	
1426301_at	0.85033	activated leukocyte cell adhesion molecule	Alcam	
1426302_at	0.90566	"transmembrane protease, serine 4"	Tmprss4	
1426303_at	1.10723	"xylosylprotein beta1,4-galactosyltransferase, polypeptide 7 (galactosyltransferase I)"	B4galt7	
1426304_x_at	1.00154	"xylosylprotein beta1,4-galactosyltransferase, polypeptide 7 (galactosyltransferase I)"	B4galt7	
1426305_at	0.87503	uroplakin 1A	Upk1a	
1426306_a_at	0.92986	"melanoma antigen, family D, 2"	Maged2	
1426307_at	1.00278	cytochrome b5 reductase 4	Cyb5r4	
1426308_at	0.98843	methyl-CpG binding domain protein 3-like 1	Mbd3l1	
1426309_at	0.89750	ankyrin repeat and SOCS box-containing protein 9	Asb9	
1426310_at	0.96594	"zinc finger, DHHC domain containing 5"	Zdhhc5	
1426311_s_at	0.94984	"zinc finger, DHHC domain containing 5"	Zdhhc5	
1426312_at	1.02681	brain and reproductive organ-expressed protein	Bre	

1426313_at 0.99386 brain and reproductive organ-expressed protein Bre
1426314_at 1.10638 endothelin receptor type B Ednrb
1426315_a_at 1.03203 RIKEN cDNA 6330416G13 gene 6330416G13Rik
1426316_at 1.05659 RIKEN cDNA 6330416G13 gene 6330416G13Rik
1426317_at 0.95486 StAR-related lipid transfer (START) domain containing 6 Stard6
1426318_at 1.26169 "serine (or cysteine) peptidase inhibitor, clade B, member 1b" Serpinb1b
1426319_at 1.01145 "platelet-derived growth factor, D polypeptide" Pdgfd
1426320_at 1.04278 RIKEN cDNA 4933437K13 gene 4933437K13Rik
1426321_at 0.95458 T-box 4 Tbx4
1426322_a_at 0.98504 "potassium large conductance calcium-activated channel, subfamily M, beta member 2" Kcnmb2
1426323_x_at 0.99326 Cd27 binding protein (Hindu God of destruction) Siva
1426324_at 1.39686 "histocompatibility 2, D region locus 1" H2-D1
1426325_at 0.86701 kinesin family member 1C Kif1c
1426326_at 0.96108 zinc finger protein 91 Zfp91
1426327_s_at 1.01866 ciliary neurotrophic factor /// zinc finger protein 91 Cntf /// Zfp91
1426328_a_at 0.84737 "sodium channel, voltage-gated, type III, beta" Scn3b
1426329_s_at 0.99011 "brain and acute leukemia, cytoplasmic" Baalc
1426330_at 0.85686 "calcium channel, voltage-dependent, gamma subunit 5" Cacng5
1426331_a_at 0.84348 FERM domain containing 4B Frmd4b
1426332_a_at 0.86597 claudin 3 Cldn3
1426333_a_at 0.89972 inhibitor of kappaB kinase beta Ikbkb
1426334_a_at 0.72393 BCL2-like 11 (apoptosis facilitator) Bcl2l11
1426335_at 0.90358 "potassium voltage-gated channel, subfamily Q, member 2" Kcnq2
1426336_at 0.89897 "calcium channel, voltage-dependent, gamma subunit 7" Cacng7
1426337_a_at 1.09594 TEA domain family member 4 Tead4
1426338_a_at 0.91164 netrin G1 Ntng1
1426339_at 0.96254 adenylate kinase 5 Ak5
1426340_at 0.96574 "solute carrier family 1 (glial high affinity glutamate transporter), member 3" Slc1a3
1426341_at 1.05389 "solute carrier family 1 (glial high affinity glutamate transporter), member 3" Slc1a3
1426342_at 1.02403 "STT3, subunit of the oligosaccharyltransferase complex, homolog B (S. cerevisiae)" Stt3b
1426343_at 1.01173 "STT3, subunit of the oligosaccharyltransferase complex, homolog B (S. cerevisiae)" Stt3b
1426344_at 0.87855 GLE1 RNA export mediator-like (yeast Gle1
1426345_at 0.89905 prolyl endopeptidase-like Prepl
1426346_at 0.90032 prolyl endopeptidase-like Prepl
1426347_at 0.93600 RIKEN cDNA 2010321M09 gene 2010321M09Rik
1426348_at 0.87914 "procollagen, type IV, alpha 1" Col4a1
1426349_s_at 0.90720 thymopoietin Tmpo
1426350_at 0.99706 mannoside acetylglucosaminyltransferase 2 Mgat2

1426351_at 1.17220 heat shock protein 1 (chaperonin) Hspd1
1426352_s_at 0.88980 Tial1 cytotoxic granule-associated RNA binding protein-like 1 Tial1
1426353_at 1.10176 signal transducer and activator of transcription 6 Stat6
1426354_at 0.91824 Brca1 associated protein 1 Bap1
1426355_a_at 0.95842 RIKEN cDNA 6330578E17 gene 6330578E17Rik
1426356_at 0.98312 RIKEN cDNA 6330578E17 gene 6330578E17Rik
1426357_at 0.94401 TAO kinase 1 Taok1
1426358_at 1.02119 TAO kinase 1 Taok1
1426359_at 1.04798 zinc finger CCCH type containing 11A Zc3h11a
1426360_at 1.01451 zinc finger CCCH type containing 11A Zc3h11a
1426361_at 1.00745 zinc finger CCCH type containing 11A Zc3h11a
1426362_at 0.90391 transmembrane protein 144 Tmem144
1426363_x_at 1.08141 "H2A histone family, member Y2" H2afy2
1426364_at 0.97408 mitochondrial ribosome recycling factor Mrrf
1426365_at 1.02420 RIKEN cDNA 2810403A07 gene 2810403A07Rik
1426366_at 1.10366 "eukaryotic translation initiation factor 2C, 2" Eif2c2
1426367_at 0.95542 calcium binding protein 39-like Cab39l
1426368_at 0.76025 Ras and Rab interactor 2 Rin2
1426369_at 0.98827 male sterility domain containing 2 Mlstd2
1426370_at 0.95353 male sterility domain containing 2 Mlstd2
1426371_at 1.00299 male sterility domain containing 2 Mlstd2
1426372_a_at 0.96015 blocked early in transport 1 homolog (*S. cerevisiae*)-like Bet11
1426373_at 0.87782 Sloan-Kettering viral oncogene homolog Ski
1426374_at 0.98370 RIKEN cDNA 2410166I05 gene 2410166I05Rik
1426375_s_at 0.99399 oxidoreductase NAD-binding domain containing 1 Oxnad1
1426376_at 0.90236 receptor accessory protein 5 Reep5
1426377_at 1.00921 zinc finger protein 281 Zfp281
1426378_at 0.99316 eukaryotic translation initiation factor 4B Eif4b
1426379_at 0.96080 eukaryotic translation initiation factor 4B Eif4b
1426380_at 0.97078 eukaryotic translation initiation factor 4B Eif4b
1426381_at 1.20627 "peroxisome proliferative activated receptor, gamma, coactivator-related 1"
Pprc1
1426382_at 1.01038 "protein phosphatase 1B, magnesium dependent, beta isoform" Ppm1b
1426383_at 0.94042 cryptochrome 2 (photolyase-like) Cry2
1426384_a_at 1.10382 "tyrosine 3-monooxygenase/tryptophan 5-monooxygenase activation protein,
epsilon polypeptide" Ywhae
1426385_x_at 1.06145 "tyrosine 3-monooxygenase/tryptophan 5-monooxygenase activation protein,
epsilon polypeptide" Ywhae
1426386_at 1.13505 ribosomal protein L7-like 1 Rpl7l1
1426387_x_at 0.85210 RIKEN cDNA 4933439C20 gene /// phosphatidylserine decarboxylase
4933439C20Rik /// Pisd
1426388_s_at 0.95342 receptor-like tyrosine kinase Ryk
1426389_at 1.02290 calcium/calmodulin-dependent protein kinase ID Camk1d

1426390_a_at 0.97405 ADP-ribosylation factor 1 Arf1
 1426391_at 0.99854 ADP-ribosylation factor 1 /// similar to ADP-ribosylation factor 1 Arf1 ///
 LOC545267
 1426392_a_at 0.93007 ARP3 actin-related protein 3 homolog (yeast) Actr3
 1426393_a_at 0.96126 stromal cell derived factor 4 Sdf4
 1426394_at 1.02718 "eukaryotic translation initiation factor 3, subunit 1 alpha" Eif3s1
 1426395_s_at 1.14176 "eukaryotic translation initiation factor 3, subunit 1 alpha" Eif3s1
 1426396_at 0.91770 CD247 antigen Cd247
 1426397_at 1.03905 "transforming growth factor, beta receptor II" Tgfbr2
 1426398_at 0.91735 ubiquitin-conjugating enzyme E2W (putative) Ube2w
 1426399_at 0.65137 von Willebrand factor A domain containing 1 Vwa1
 1426400_a_at 1.01778 "calpain, small subunit 1" Capns1
 1426401_at 0.95169 "protein phosphatase 3, catalytic subunit, alpha isoform" Ppp3ca
 1426402_at 1.06843 "synaptotagmin binding, cytoplasmic RNA interacting protein" Syncrip
 1426403_at 1.00513 ARP1 actin-related protein 1 homolog B (yeast) Actr1b
 1426404_a_at 0.86389 ring finger protein 11 Rnf11
 1426405_at 0.97257 ring finger protein 11 Rnf11
 1426406_at 1.00214 SET domain containing (lysine methyltransferase) 8 Setd8
 1426407_at 1.00590 "CUG triplet repeat, RNA binding protein 1" Cugbp1
 1426408_at 1.04495 "CUG triplet repeat, RNA binding protein 1" Cugbp1
 1426409_at 0.87085 "leucine zipper, putative tumor suppressor 2" Lzts2
 1426410_at 1.10840 "pyruvate dehydrogenase kinase, isoenzyme 3" Pdk3
 1426411_a_at 0.89926 spermatid perinuclear RNA binding protein /// RIKEN cDNA C230082I21
 gene Strbp /// C230082I21Rik
 1426412_at 1.10085 neurogenic differentiation 1 Neurod1
 1426413_at 1.05982 neurogenic differentiation 1 Neurod1
 1426414_a_at 1.08569 ring finger protein 7 Rnf7
 1426415_a_at 1.16339 tripartite motif protein 25 Trim25
 1426416_a_at 0.86822 "Yip1 domain family, member 4" Yipf4
 1426417_at 1.23885 "Yip1 domain family, member 4" Yipf4
 1426418_at 0.70982 atonal homolog 8 (Drosophila) Atoh8
 1426419_at 1.03275 RNA binding motif protein 26 Rbm26
 1426420_at 0.94294 RNA binding motif protein 26 Rbm26
 1426421_s_at 1.01550 RNA binding motif protein 26 Rbm26
 1426422_at 0.96786 coiled-coil and C2 domain containing 1A Cc2d1a
 1426423_at 1.07704 serine hydroxymethyl transferase 2 (mitochondrial) Shmt2
 1426424_at 0.95837 "SGT1, suppressor of G2 allele of SKP1 (S. cerevisiae)" Sugt1
 1426425_at 1.12311 "SGT1, suppressor of G2 allele of SKP1 (S. cerevisiae)" Sugt1
 1426426_at 1.03651 RNA binding motif protein 13 Rbm13
 1426427_at 0.99190 tubulin tyrosine ligase-like 1 Ttl1
 1426428_at 1.00088 THAP domain containing 7 Thap7
 1426429_at 0.89270 THAP domain containing 7 Thap7
 1426430_at 0.94507 jagged 2 Jag2

1426431_at	0.93249 jagged 2	Jag2	
1426432_a_at	0.99595 "solute carrier family 4 (anion exchanger), member 4"	Slc4a4	
1426433_at	1.03110 myc target 1	Myct1	
1426434_at	0.92333 transmembrane protein 43	Tmem43	
1426435_at	0.86406 transmembrane protein 135	Tmem135	
1426436_at	0.93973 transmembrane protein 159	Tmem159	
1426437_s_at	0.99593 histone deacetylase 3	Hdac3	
1426438_at	0.97569 "DEAD (Asp-Glu-Ala-Asp) box polypeptide 3, Y-linked"	Ddx3y	
1426439_at	1.09898 "DEAD (Asp-Glu-Ala-Asp) box polypeptide 3, Y-linked"	Ddx3y	
1426440_at	0.90476 dehydrogenase/reductase (SDR family) member 7	Dhrs7	
1426441_at	1.50484 "solute carrier family 11 (proton-coupled divalent metal ion transporters), member 2"	Slc11a2	
1426442_at	1.00989 glycoprotein m6a	Gpm6a	
1426443_at	0.86815 rhomboid domain containing 2	Rhbdd2	
1426444_at	0.98106 rhomboid domain containing 2	Rhbdd2	
1426445_at	0.97165 meningioma expressed antigen 6 (coiled-coil proline-rich)	Mgea6	
1426446_at	1.12723 RIKEN cDNA 6430548M08 gene	6430548M08Rik	
1426447_at	1.09914 nucleoporin 35	Nup35	
1426448_at	0.86252 "praja1, RING-H2 motif containing"	Pja1	
1426449_a_at	0.87427 "praja1, RING-H2 motif containing"	Pja1	
1426450_at	0.91889 phospholipase C-like 2	Plcl2	
1426451_at	0.94038 RIKEN cDNA 6030465E24 gene	6030465E24Rik	
1426452_a_at	1.08867 "RAB30, member RAS oncogene family"	Rab30	
1426453_at	0.96893 pitrilysin metalloproteinase 1	Pitrm1	
1426454_at	1.82764 "Rho, GDP dissociation inhibitor (GDI) beta"	Arhgdib	
1426455_at	1.07429 serologically defined colon cancer antigen 10	Sdccag10	
1426456_a_at	0.95546 protein inhibitor of activated STAT 2	Pias2	
1426457_at	1.19193 sarcolemma associated protein	Slmap	
1426458_at	1.10886 sarcolemma associated protein	Slmap	
1426459_s_at	0.90296 expressed sequence AW549877	AW549877	
1426460_a_at	1.00091 UDP-glucose pyrophosphorylase 2	Ugp2	
1426461_at	0.93342 UDP-glucose pyrophosphorylase 2	Ugp2	
1426462_at	1.18767 gephyrin	Gphn	
1426463_at	1.11541 gephyrin	Gphn	
1426464_at	0.81429 "nuclear receptor subfamily 1, group D, member 1"	Nr1d1	
1426465_at	1.01509 "discs, large homolog-associated protein 4 (Drosophila)"	Dlgap4	
1426466_s_at	0.91917 ribosomal protein S6 kinase-like 1	Rps6kl1	
1426467_s_at	1.00536 RIKEN cDNA 0610037L13 gene	0610037L13Rik	
1426468_at	0.98530 RIKEN cDNA 0610037L13 gene	0610037L13Rik	
1426469_a_at	0.94170 TATA box binding protein	Tbp	
1426470_at	1.00144 --- ---		
1426471_at	0.96132 zinc finger protein 52	Zfp52	
1426472_at	1.10417 zinc finger protein 52	Zfp52	

1426473_at 0.90497 "DnaJ (Hsp40) homolog, subfamily C, member 9" Dnajc9
1426474_at 1.05834 ATP synthase mitochondrial F1 complex assembly factor 2 Atpaf2
1426475_at 1.07901 hydroxymethylbilane synthase Hmbs
1426476_at 0.89968 RAS p21 protein activator 1 Rasa1
1426477_at 1.00512 RAS p21 protein activator 1 Rasa1
1426478_at 0.73405 RAS p21 protein activator 1 Rasa1
1426479_a_at 0.88715 trinucleotide repeat containing 5 Tnrc5
1426480_at 0.96817 Shwachman-Bodian-Diamond syndrome homolog (human) Sbds
1426481_at 0.94608 kelch-like 22 (Drosophila) Klhl22
1426482_at 1.13424 "protein-kinase, interferon-inducible double stranded RNA dependent inhibitor, repressor of (P58 repressor)" Prkrir
1426483_at 1.01249 "protein-kinase, interferon-inducible double stranded RNA dependent inhibitor, repressor of (P58 repressor)" Prkrir
1426484_at 0.95296 UBX domain containing 2 Ubxd2
1426485_at 0.99625 UBX domain containing 2 Ubxd2
1426486_at 0.98220 UBX domain containing 2 Ubxd2
1426487_a_at 1.14781 retinoblastoma binding protein 6 Rbbp6
1426488_at 0.96902 bifunctional apoptosis regulator Bfar
1426489_s_at 0.96075 bifunctional apoptosis regulator Bfar
1426490_at 0.91189 bifunctional apoptosis regulator Bfar
1426491_at 0.97567 hect (homologous to the E6-AP (UBE3A) carboxyl terminus) domain and RCC1 (CHC1)-like domain (RLD) 2 Herc2
1426492_at 1.01542 tyrosyl-DNA phosphodiesterase 1 Tdp1
1426493_a_at 0.90782 kinesin family member C2 Kifc2
1426494_at 0.92030 RNA (guanine-9-) methyltransferase domain containing 3 Rg9mtd3
1426495_at 1.06795 RIKEN cDNA 2410042D21 gene 2410042D21Rik
1426496_at 1.09747 WD repeat domain 55 Wdr55
1426497_at 0.95827 "jumonji, AT rich interactive domain 1C (Rbp2 like)" Jarid1c
1426498_at 1.19943 "jumonji, AT rich interactive domain 1C (Rbp2 like)" Jarid1c
1426499_at 1.01739 SH3-domain GRB2-like endophilin B2 Sh3glb2
1426500_at 1.01154 isoprenylcysteine carboxyl methyltransferase Icmt
1426501_a_at 1.04995 Traf2 binding protein T2bp
1426502_s_at 1.12336 "glutamic pyruvic transaminase 1, soluble" Gpt1
1426503_a_at 1.01752 ring finger protein 121 Rnf121
1426504_a_at 0.95684 ring finger protein 121 Rnf121
1426505_at 0.96229 ecotropic viral integration site 2b Evi2b
1426506_at 0.95682 ecotropic viral integration site 2a /// ecotropic viral integration site 2b Evi2a /// Evi2b
1426507_at 0.99563 "interleukin 1 family, member 5 (delta)" Il1f5
1426508_at 0.94840 glial fibrillary acidic protein Gfap
1426509_s_at 1.12608 glial fibrillary acidic protein Gfap
1426510_at 1.01899 saccharopine dehydrogenase (putative) Sccpdh
1426511_at 0.94337 sushi domain containing 2 Susd2

1426512_at 0.98794 olfactomedin 3 Olfm3
1426513_at 1.15909 RNA binding motif protein 28 Rbm28
1426514_at 0.91548 RIKEN cDNA 4631426J05 gene 4631426J05Rik
1426515_a_at 0.99098 "torsin family 1, member A (torsin A)" Tor1a
1426516_a_at 1.32974 lipin 1 Lpin1
1426517_at 0.98672 "guanine nucleotide binding protein, alpha z subunit" Gnaz
1426518_at 0.89535 "tubulin, gamma complex associated protein 5" Tubgcp5
1426519_at 0.81754 "procollagen-proline, 2-oxoglutarate 4-dioxygenase (proline 4-hydroxylase), alpha 1 polypeptide" P4ha1
1426520_at 0.96762 B-cell translocation gene 4 Btg4
1426521_at 0.91907 RIKEN cDNA D230025D16 gene D230025D16Rik
1426522_at 0.86813 "hydroxyacyl-Coenzyme A dehydrogenase/3-ketoacyl-Coenzyme A thiolase/enoyl-Coenzyme A hydratase (trifunctional protein), beta subunit /// similar to hydroxyacyl-Coenzyme A dehydrogenase/3-ketoacyl-Coenzyme A thiolase/enoyl-Coenzyme A hydratase (trifunctional protein), beta subunit" Hadhb /// LOC623031
1426523_a_at 1.01083 glucosamine-6-phosphate deaminase 2 Gnpda2
1426524_at 1.08149 glucosamine-6-phosphate deaminase 2 Gnpda2
1426525_at 0.93345 AT rich interactive domain 2 (Arid-rfx like) Arid2
1426526_s_at 1.05289 oviductal glycoprotein 1 Ovgp1
1426527_at 1.00521 "target of EGR1, member 1 (nuclear)" Toe1
1426528_at 1.30951 neuropilin 2 Nrp2
1426529_a_at 1.00700 transgelin 2 Tagln2
1426530_a_at 0.86815 kelch-like 5 (Drosophila) Klh15
1426531_at 0.90843 "zinc finger, MYND domain containing 11" Zmynd11
1426532_at 0.95300 "zinc finger, MYND domain containing 11" Zmynd11
1426533_at 1.13851 nucleolar protein 5A Nol5a
1426534_a_at 0.97725 ADP-ribosylation factor GTPase activating protein 3 Arfgap3
1426535_at 0.97208 RIKEN cDNA 9630046K23 gene 9630046K23Rik
1426536_at 0.96248 NMDA receptor-regulated gene 2 Narg2
1426537_at 1.04967 NMDA receptor-regulated gene 2 Narg2
1426538_a_at 1.02281 transformation related protein 53 Trp53
1426539_at 0.89298 ubiquitin specific peptidase 11 Usp11
1426540_at 1.01269 endonuclease domain containing 1 Endod1
1426541_a_at 0.97816 endonuclease domain containing 1 Endod1
1426542_at 0.94670 endonuclease domain containing 1 Endod1
1426543_x_at 1.05646 endonuclease domain containing 1 Endod1
1426544_a_at 1.02777 tetratricopeptide repeat domain 14 Ttc14
1426545_at 0.93383 trinucleotide repeat containing 6b Tnrc6b
1426546_at 1.16304 testis-specific kinase 2 Tesk2
1426547_at 0.75594 group specific component Gc
1426548_a_at 0.92071 ATP binding domain 4 Atpbd4
1426549_at 1.04335 RIKEN cDNA 2410018C20 gene 2410018C20Rik
1426550_at 0.97910 "SID1 transmembrane family, member 1" Sidt1

1426551_at 0.93324 "SID1 transmembrane family, member 1" Sidt1
 1426552_a_at 0.98892 B-cell CLL/lymphoma 11A (zinc finger protein) Bcl11a
 1426553_at 0.91852 "DnaJ (Hsp40) homolog, subfamily C, member 14" Dnajc14
 1426554_a_at 1.10608 phosphoglycerate mutase 1 /// similar to Phosphoglycerate mutase 1 (Phosphoglycerate mutase isozyme B) (PGAM-B) (BPG-dependent PGAM 1) /// similar to Phosphoglycerate mutase 1 (Phosphoglycerate mutase isozyme B) (PGAM-B) (BPG-dependent PGAM 1) /// similar to Phosphoglycerate mutase 1 (Phosphoglycerate mutase isozyme B) (PGAM-B) (BPG-dependent PGAM 1) /// similar to Phosphoglycerate mutase 1 (Phosphoglycerate mutase isozyme B) (PGAM-B) (BPG-dependent PGAM 1) Pgam1 /// LOC668045 /// LOC672147 /// LOC675179 /// LOC677000
 1426555_at 0.95641 serine carboxypeptidase 1 Scpep1
 1426556_at 0.98365 suppressor of hairy wing homolog 4 (Drosophila) Suhw4
 1426557_at 0.97818 mesoderm posterior 1 Mesp1
 1426558_x_at 0.91125 RIKEN cDNA 3100002L24 gene /// similar to zinc finger protein 14 /// RIKEN cDNA 0610010B08 gene /// similar to zinc finger protein 91 /// similar to zinc finger protein 709 /// similar to gonadotropin inducible ovarian transcription factor 1 /// similar to zinc finger protein 97 /// similar to gonadotropin inducible ovarian transcription factor 1 3100002L24Rik /// LOC627901 /// 0610010B08Rik /// LOC628084 /// LOC628147 /// LOC668030 /// LOC668039 /// LOC668066
 1426559_at 1.11630 "sno, strawberry notch homolog 1 (Drosophila)" Sbno1
 1426560_a_at 0.76723 nephronectin Npnt
 1426561_a_at 0.76668 nephronectin Npnt
 1426562_a_at 0.92125 olfactomedin 1 Olm1
 1426563_at 0.86768 zinc finger protein 553 Zfp553
 1426564_at 1.08640 zinc finger protein 553 Zfp553
 1426565_at 1.00922 insulin-like growth factor I receptor Igflr
 1426566_s_at 0.78928 interleukin 17 receptor E Il17re
 1426567_a_at 0.90388 PQ loop repeat containing 1 Pqlc1
 1426568_at 1.10421 "solute carrier family 2 (facilitated glucose transporter), member 9" Slc2a9
 1426569_a_at 1.24620 fyn-related kinase Frk
 1426570_a_at 1.02677 fyn-related kinase Frk
 1426571_at 0.74502 transmembrane protein 16A Tmem16a
 1426572_at 1.19200 "malic enzyme 2, NAD(+)-dependent, mitochondrial" Me2
 1426573_at 1.00778 "malic enzyme 2, NAD(+)-dependent, mitochondrial" Me2
 1426574_a_at 0.88418 adducin 3 (gamma) Add3
 1426575_at 0.96738 transmembrane protein 23 Tmem23
 1426576_at 0.99890 transmembrane protein 23 Tmem23
 1426577_a_at 1.03666 RIKEN cDNA 1810054G18 gene 1810054G18Rik
 1426578_s_at 0.96860 SNAP-associated protein Snapap
 1426579_at 0.98255 guanine nucleotide binding protein-like 2 (nucleolar) Gnl2
 1426580_at 1.01080 polo-like kinase 4 (Drosophila) Plk4
 1426581_at 1.02428 "protein tyrosine phosphatase, mitochondrial 1" Ptpmt1
 1426582_at 1.14295 activating transcription factor 2 Atf2
 1426583_at 1.05259 activating transcription factor 2 Atf2

1426584_a_at 0.80633 sorbitol dehydrogenase Sord
1426585_s_at 1.06394 mitogen activated protein kinase 1 Mapk1
1426586_at 0.92733 "solute carrier family 25 (mitochondrial carrier oxoglutarate carrier), member 11" Slc25a11
1426587_a_at 1.02680 signal transducer and activator of transcription 3 Stat3
1426588_at 0.95712 tetraspanin 10 Tspan10
1426589_at 1.07291 growth factor receptor bound protein 2-associated protein 3 Gab3
1426590_at 0.90490 "G elongation factor, mitochondrial 2" Gfm2
1426591_at 1.05488 "G elongation factor, mitochondrial 2" Gfm2
1426592_a_at 1.04754 F-box only protein 22 Fbxo22
1426593_a_at 1.03584 F-box only protein 22 Fbxo22
1426594_at 0.85890 FERM domain containing 4B Frmd4b
1426595_at 1.02097 "solute carrier family 18 (vesicular monoamine), member 1" Slc18a1
1426596_a_at 1.16162 survival motor neuron 1 Smn1
1426597_s_at 1.08809 expressed sequence C79267 C79267
1426598_at 1.08088 "ubiquitously transcribed tetratricopeptide repeat gene, Y chromosome" Uty
1426599_a_at 1.40866 "solute carrier family 2 (facilitated glucose transporter), member 1" Slc2a1
1426600_at 1.25073 "solute carrier family 2 (facilitated glucose transporter), member 1" Slc2a1
1426601_at 0.94922 "solute carrier family 37 (glycerol-3-phosphate transporter), member 1" Slc37a1
1426602_at 1.04647 v-raf murine sarcoma 3611 viral oncogene homolog Araf
1426603_at 0.87759 "ribonuclease L (2', 5'-oligoadenylate synthetase-dependent)" Rnasel
1426604_at 0.86229 "ribonuclease L (2', 5'-oligoadenylate synthetase-dependent)" Rnasel
1426605_at 0.81559 "BRCA1/BRCA2-containing complex, subunit 3" Brcc3
1426606_at 0.97483 cartilage acidic protein 1 Crtac1
1426607_at 0.77850 similar to CG10866-PA LOC633640
1426608_at 1.02836 similar to CG10866-PA LOC633640
1426609_at 1.12161 RIKEN cDNA 2810028N01 gene 2810028N01Rik
1426610_a_at 0.99068 transcription termination factor 1 Ttf1
1426611_at 1.05097 "proteasome (prosome, macropain) 26S subunit, ATPase 2" Psmc2
1426612_at 1.02128 timeless interacting protein Tipin
1426613_a_at 1.04456 U2 small nuclear ribonucleoprotein B Snrpb2
1426614_at 0.93714 protein kinase C binding protein 1 Prkcbp1
1426615_s_at 0.74869 N-myc downstream regulated gene 4 Ndr4
1426616_at 1.00512 TLC domain containing 1 Tlcd1
1426617_a_at 1.09196 tweety homolog 1 (Drosophila) Ttyh1
1426618_a_at 0.91268 "protein O-linked mannose beta1,2-N-acetylglucosaminyltransferase"
Pomgnt1
1426619_at 0.87971 "gene model 1313, (NCBI)" Gm1313
1426620_at 0.91179 carbohydrate sulfotransferase 10 Chst10
1426621_a_at 0.73016 "protein phosphatase 2 (formerly 2A), regulatory subunit B (PR 52), beta isoform" Ppp2r2b
1426622_a_at 0.95705 glutaminyl-peptide cyclotransferase (glutaminyl cyclase) /// similar to
Glutaminyl-peptide cyclotransferase precursor (QC) (Glutaminyl-tRNA cyclotransferase) (Glutaminyl

cyclase) Qpct /// LOC639942
1426623_a_at 0.88411 Rho GTPase activating protein 17 Arhgap17
1426624_a_at 0.68708 yippee-like 3 (Drosophila) Ypel3
1426625_at 0.97410 zinc finger protein 623 Zfp623
1426626_at 1.18511 "general transcription factor IIF, polypeptide 2" Gtf2f2
1426627_at 0.99794 mitogen activated protein kinase kinase kinase 7 Map3k7
1426628_at 0.87677 transmembrane protein 34 Tmem34
1426629_at 1.04212 DEAH (Asp-Glu-Ala-His) box polypeptide 8 Dhx8
1426630_at 1.12333 exosome component 2 Exosc2
1426631_at 1.07599 pseudouridylate synthase 7 homolog (S. cerevisiae) Pus7
1426632_at 0.97797 potassium channel tetramerisation domain containing 14 Kctd14
1426633_s_at 0.88974 potassium channel tetramerisation domain containing 14 Kctd14
1426634_at 1.01762 "solute carrier family 5 (sodium/glucose cotransporter), member 9" Slc5a9
1426635_at 0.99001 acyl-Coenzyme A binding domain containing 3 Acbd3
1426636_a_at 0.96600 baculoviral IAP repeat-containing 4 Birc4
1426637_a_at 0.92495 sine oculis-related homeobox 3 homolog (Drosophila) Six3
1426638_at 1.10780 sine oculis-related homeobox 3 homolog (Drosophila) Six3
1426639_a_at 0.91436 "transcription factor 7-like 2, T-cell specific, HMG-box" Tcf712
1426640_s_at 0.87507 tribbles homolog 2 (Drosophila) Trib2
1426641_at 1.07334 tribbles homolog 2 (Drosophila) Trib2
1426642_at 0.75945 fibronectin 1 Fn1
1426643_at 0.98626 elongation protein 3 homolog (S. cerevisiae) Elp3
1426644_at 1.10018 "TBC1 domain family, member 20" Tbc1d20
1426645_at 1.05798 "heat shock protein 90kDa alpha (cytosolic), class A member 1" Hsp90aa1
1426646_at 0.88903 RIKEN cDNA 9130011J15 gene 9130011J15Rik
1426647_at 0.88296 RIKEN cDNA 9130011J15 gene 9130011J15Rik
1426648_at 1.00439 MAP kinase-activated protein kinase 2 Mapkapk2
1426649_at 0.83755 transmembrane protein with EGF-like and two follistatin-like domains 1 Tmeff1
1426650_at 1.01568 "myosin, heavy polypeptide 8, skeletal muscle, perinatal" Myh8
1426651_at 1.15870 mitochondrial ribosomal protein L44 Mrpl44
1426652_at 1.05920 minichromosome maintenance deficient 3 (S. cerevisiae) /// similar to DNA replication licensing factor MCM3 (DNA polymerase alpha holoenzyme-associated protein P1) (P1-MCM3) Mcm3 /// LOC671904
1426653_at 1.04203 minichromosome maintenance deficient 3 (S. cerevisiae) Mcm3
1426654_at 1.02115 "zinc finger, C3HC type 1" Zc3hc1
1426655_a_at 1.04298 RIKEN cDNA 4930504E06 gene 4930504E06Rik
1426656_at 0.91177 RIKEN cDNA 4930504E06 gene 4930504E06Rik
1426657_s_at 1.12307 3-phosphoglycerate dehydrogenase /// similar to 3-phosphoglycerate dehydrogenase /// similar to 3-phosphoglycerate dehydrogenase /// similar to 3-phosphoglycerate dehydrogenase /// similar to 3-phosphoglycerate dehydrogenase Phgdh /// LOC666422 /// LOC666875 /// LOC669985 /// LOC671102 /// LOC673015 /// LOC675010
1426658_x_at 1.07357 3-phosphoglycerate dehydrogenase /// similar to 3-phosphoglycerate

dehydrogenase Phgdh /// LOC673015
 1426659_a_at 1.00978 ribosomal protein L23a Rpl23a
 1426660_x_at 1.00598 ribosomal protein L23a Rpl23a
 1426661_at 1.03080 ribosomal protein L27a /// similar to ribosomal protein L27a Rpl27a ///
 LOC665189
 1426662_at 0.91510 cytidine monophospho-N-acetylneuraminic acid synthetase Cmas
 1426663_s_at 1.09059 "solute carrier family 45, member 3" Slc45a3
 1426664_x_at 1.04734 "solute carrier family 45, member 3" Slc45a3
 1426665_at 1.02474 katanin p80 (WD40-containing) subunit B 1 Katnb1
 1426666_a_at 0.93520 unc-84 homolog A (C. elegans) Unc84a
 1426667_a_at 0.91596 unc-84 homolog A (C. elegans) Unc84a
 1426668_at 0.90413 "solute carrier family 30 (zinc transporter), member 9" Slc30a9
 1426669_at 0.83997 RIKEN cDNA C530044N13 gene C530044N13Rik
 1426670_at 0.92439 agrin Agrin
 1426671_a_at 0.88587 RNA binding motif protein 39 Rbm39
 1426672_at 0.91972 transmembrane protein 16K Tmem16k
 1426673_at 0.98069 cadherin 3 Cdh3
 1426674_at 1.02509 "eukaryotic translation initiation factor 3, subunit 9 (eta)" Eif3s9
 1426675_at 1.02472 translocase of outer mitochondrial membrane 70 homolog A (yeast)
 Tomm70a
 1426676_s_at 1.02275 translocase of outer mitochondrial membrane 70 homolog A (yeast)
 Tomm70a
 1426677_at 0.95786 "filamin, alpha" Flna
 1426678_at 0.94932 zinc finger protein 706 Zfp706
 1426679_at 0.98402 zinc finger protein 706 Zfp706
 1426680_at 0.77143 "selenoprotein N, 1" Sepn1
 1426681_at 0.91939 zinc finger CCCH type containing 5 Zc3h5
 1426682_at 0.86524 "CCR4-NOT transcription complex, subunit 6" Cnot6
 1426683_at 0.98549 "CCR4-NOT transcription complex, subunit 6" Cnot6
 1426684_at 0.91602 "CCR4-NOT transcription complex, subunit 6" Cnot6
 1426685_a_at 0.98667 "CCR4-NOT transcription complex, subunit 6" Cnot6
 1426686_s_at 0.93685 mitogen activated protein kinase kinase kinase 3 Map3k3
 1426687_at 1.03293 mitogen activated protein kinase kinase kinase 3 Map3k3
 1426688_at 0.95454 "succinate dehydrogenase complex, subunit A, flavoprotein (Fp)" Sdha
 1426689_s_at 0.94856 "succinate dehydrogenase complex, subunit A, flavoprotein (Fp)" Sdha
 1426690_a_at 0.90132 sterol regulatory element binding factor 1 Srebf1
 1426691_at 1.05738 tight junction associated protein 1 Tjap1
 1426692_at 1.00743 coiled-coil domain containing 97 Ccdc97
 1426693_x_at 0.99147 "COX15 homolog, cytochrome c oxidase assembly protein (yeast)" Cox15
 1426694_at 0.96553 RIKEN cDNA 9030624J02 gene 9030624J02Rik
 1426695_at 1.06070 RIKEN cDNA 9030624J02 gene 9030624J02Rik
 1426696_at 0.78810 low density lipoprotein receptor-related protein associated protein 1 Lrpap1
 1426697_a_at 0.75301 low density lipoprotein receptor-related protein associated protein 1 Lrpap1

1426698_a_at 1.07643 heterogeneous nuclear ribonucleoprotein M Hnrpm
1426699_at 0.91357 expressed sequence AU040320 AU040320
1426700_a_at 1.05837 ubiquitin specific peptidase 52 Usp52
1426701_at 0.89717 RIKEN cDNA 4632419K20 gene 4632419K20Rik
1426702_at 0.89348 RIKEN cDNA 4632419K20 gene 4632419K20Rik
1426703_at 1.04815 cyclin G associated kinase Gak
1426704_at 0.88196 cyclin G associated kinase Gak
1426705_s_at 1.20688 isoleucine-tRNA synthetase Iars
1426706_s_at 1.07715 xylulokinase homolog (H. influenzae) Xylb
1426707_at 0.87843 "tubulin, gamma complex associated protein 3" Tubgcp3
1426708_at 0.90330 anthrax toxin receptor 2 Antxr2
1426709_a_at 0.98522 ubiquitin specific peptidase 33 Usp33
1426710_at 0.99069 calmodulin 3 Calm3
1426711_at 0.75189 transmembrane and coiled-coil domains 3 Tmco3
1426712_at 1.02046 "solute carrier family 6 (neurotransmitter transporter), member 15" Slc6a15
1426713_s_at 1.12103 glutamyl-prolyl-tRNA synthetase /// similar to Bifunctional aminoacyl-tRNA
synthetase Eprs /// LOC633677
1426714_at 1.04648 "DNA segment, Chr 11, ERATO Doi 18, expressed" D11Ert18e
1426715_s_at 1.09137 "DNA segment, Chr 11, ERATO Doi 18, expressed" D11Ert18e
1426716_at 1.11205 tudor domain containing 7 Tdrd7
1426717_at 0.99457 non imprinted in Prader-Willi/Angelman syndrome 2 homolog (human)
Nipa2
1426718_at 1.11174 superkiller viralicidic activity 2-like 2 (S. cerevisiae) Skiv2l2
1426719_at 0.94600 "amyloid beta (A4) precursor protein-binding, family B, member 2" Apbb2
1426720_at 0.81488 "amyloid beta (A4) precursor protein-binding, family B, member 2" Apbb2
1426721_s_at 1.13862 TCDD-inducible poly(ADP-ribose) polymerase Tiparp
1426722_at 0.86304 "solute carrier family 38, member 2" Slc38a2
1426723_at 0.96961 WD repeat domain 48 Wdr48
1426724_at 0.90931 "calponin 3, acidic" Cnn3
1426725_s_at 1.01116 "E26 avian leukemia oncogene 1, 5' domain" Ets1
1426726_at 1.00201 "protein phosphatase 1, regulatory subunit 10" Ppp1r10
1426727_s_at 1.00725 "protein phosphatase 1, regulatory subunit 10" Ppp1r10
1426728_x_at 0.94301 phosphatidylserine synthase 2 Ptdss2
1426729_at 0.97105 RIKEN cDNA 2900046G09 gene 2900046G09Rik
1426730_a_at 0.98891 prolactin-like protein K Prlplk
1426731_at 0.83521 desmin Des
1426732_at 1.04705 desmin Des
1426733_at 1.03611 "inositol 1,3,4-triphosphate 5/6 kinase" Itpk1
1426734_at 0.92887 cDNA sequence BC022623 BC022623
1426735_at 1.00149 "isoleucine-tRNA synthetase 2, mitochondrial" Iars2
1426736_at 0.98070 G1 to S phase transition 1 Gspt1
1426737_at 1.12141 G1 to S phase transition 1 Gspt1
1426738_at 0.87049 diacylglycerol kinase zeta Dgkz

1426739_at 0.96770 downstream neighbor of SON Donson
1426740_s_at 1.07190 armadillo repeat containing 5 Armc5
1426741_a_at 0.92520 FAST kinase domains 2 Fastkd2
1426742_at 0.95200 "ATP synthase, H+ transporting, mitochondrial F0 complex, subunit b, isoform 1" Atp5f1
1426743_at 0.89537 Dip3 beta Dip3b
1426744_at 1.17206 sterol regulatory element binding factor 2 Srebf2
1426745_at 0.96748 "DNA segment, Chr 11, Brigham & Women's Genetics 0434 expressed" D11Bwg0434e
1426746_at 0.91399 RIKEN cDNA 1810026J23 gene 1810026J23Rik
1426747_at 1.08845 "ATP-binding cassette, sub-family F (GCN20), member 3" Abcf3
1426748_s_at 0.82382 "ATP-binding cassette, sub-family F (GCN20), member 3" Abcf3
1426749_at 1.09922 protein arginine N-methyltransferase 3 Prmt3
1426750_at 1.12219 "filamin, beta" Flnb
1426751_s_at 1.05725 nucleoporin 107 Nup107
1426752_at 0.83286 PHD finger protein 17 Phf17
1426753_at 0.94202 PHD finger protein 17 Phf17
1426754_x_at 0.89295 cytoskeleton-associated protein 4 Ckap4
1426755_at 0.83080 cytoskeleton-associated protein 4 Ckap4
1426756_at 0.75695 UDP-N-acetyl-alpha-D-galactosamine:polypeptide N-acetylgalactosaminyltransferase 2 Galnt2
1426757_at 1.04094 adenosine monophosphate deaminase 2 (isoform L) Ampd2
1426758_s_at 1.05696 "GTL2, imprinted maternally expressed untranslated mRNA" Gtl2
1426759_at 0.86046 mitogen-activated protein kinase kinase kinase 3 /// similar to mitogen-activated protein kinase kinase kinase 3 Map4k3 /// LOC675560
1426760_at 0.95880 importin 8 Ipo8
1426761_at 0.96232 amine oxidase (flavin containing) domain 2 Aof2
1426762_s_at 0.91048 amine oxidase (flavin containing) domain 2 Aof2
1426763_at 0.88323 ornithine decarboxylase antizyme 2 Oaz2
1426764_at 0.93192 ornithine decarboxylase antizyme 2 Oaz2
1426765_at 1.08880 COMM domain containing 7 /// similar to COMM domain containing protein 7 /// similar to COMM domain-containing protein 7 Commd7 /// LOC631742 /// LOC674161
1426766_at 1.06405 RIKEN cDNA 6330403K07 gene 6330403K07Rik
1426767_at 0.93594 RIKEN cDNA 3230401M21 gene 3230401M21Rik
1426768_at 0.94872 RIKEN cDNA 1700060H10 gene 1700060H10Rik
1426769_s_at 0.93624 mastermind like 1 (Drosophila) Maml1
1426770_at 0.89898 peroxisome biogenesis factor 5 Pex5
1426771_at 0.88459 taurine upregulated gene 1 Tug1
1426772_x_at 0.97245 "T-cell receptor beta, joining region /// T-cell receptor beta, variable 13 /// similar to T-cell receptor beta-1 chain C region /// similar to T-cell receptor beta-2 chain C region /// similar to T-cell receptor beta-1 chain C region /// similar to T-cell receptor beta-1 chain C region /// similar to T-cell receptor beta-2 chain C region /// similar to T-cell receptor beta-1 chain C region" Tcrb-J /// Tcrb-V13 /// LOC665500 ///

LOC665506 /// LOC669844 /// LOC669847 /// LOC674047 /// LOC674048 /// LOC676782
 1426773_at 0.90905 mitofusin 1 Mfn1
 1426774_at 0.99782 "poly (ADP-ribose) polymerase family, member 12" Parp12
 1426775_s_at 0.91507 secretory carrier membrane protein 1 Scamp1
 1426776_at 0.98718 Wiskott-Aldrich syndrome-like (human) Wasl
 1426777_a_at 1.02150 Wiskott-Aldrich syndrome-like (human) Wasl
 1426778_at 0.89111 dystroglycan 1 Dag1
 1426779_x_at 0.90647 dystroglycan 1 Dag1
 1426780_at 0.95640 "DNA segment, Chr 14, ERATO Doi 436, expressed" D14Ertd436e
 1426781_at 0.84295 tRNA-yW synthesizing protein 1 homolog (S. cerevisiae) Tyw1
 1426782_at 0.85490 G protein-coupled receptor 125 Gpr125
 1426783_at 0.96181 GCN5 general control of amino acid synthesis-like 2 (yeast) Gcn512
 1426784_at 0.90057 tripartite motif protein 47 Trim47
 1426785_s_at 1.24084 monoglyceride lipase Mgl1
 1426786_s_at 1.01521 DEAH (Asp-Glu-Ala-His) box polypeptide 38 Dhx38
 1426787_at 0.96821 "Sfi1 homolog, spindle assembly associated (yeast) /// similar to spindle
 assembly associated Sfi1 homolog isoform b /// similar to spindle assembly associated Sfi1 homolog
 isoform b /// similar to Sfi1 homolog, spindle assembly associated (yeast)" Sfi1 /// LOC673404 ///
 LOC673420 /// LOC673566
 1426788_a_at 1.07009 structure specific recognition protein 1 Ssrp1
 1426789_s_at 1.01749 structure specific recognition protein 1 Ssrp1
 1426790_at 1.00055 structure specific recognition protein 1 Ssrp1
 1426791_at 0.93345 RUN and SH3 domain containing 2 Rusc2
 1426792_s_at 0.99430 RUN and SH3 domain containing 2 Rusc2
 1426793_a_at 1.21569 ribosomal protein L14 Rpl14
 1426794_at 0.94038 "protein tyrosine phosphatase, receptor type, S" Ptprs
 1426795_at 0.89400 "protein tyrosine phosphatase, receptor type, S" Ptprs
 1426796_at 0.97007 karyopherin (importin) alpha 6 Kpna6
 1426797_at 1.01291 RIKEN cDNA 2700094F01 gene 2700094F01Rik
 1426798_a_at 1.02959 --- ---
 1426799_at 1.09600 "RAB8B, member RAS oncogene family" Rab8b
 1426800_at 0.91272 "RAB8B, member RAS oncogene family /// similar to RAB8B, member RAS
 oncogene family" Rab8b /// LOC670268
 1426801_at 1.03795 septin 8 8-Sep
 1426802_at 1.13832 septin 8 8-Sep
 1426803_at 1.10904 RNA binding motif protein 26 Rbm26
 1426804_at 0.99693 "SWI/SNF related, matrix associated, actin dependent regulator of chromatin,
 subfamily a, member 4" Smarca4
 1426805_at 1.06020 "SWI/SNF related, matrix associated, actin dependent regulator of chromatin,
 subfamily a, member 4" Smarca4
 1426806_at 1.27003 RIKEN cDNA 5830411E10 gene 5830411E10Rik
 1426807_at 0.91354 Leukotriene A4 hydrolase Lta4h
 1426808_at 1.21593 "lectin, galactose binding, soluble 3" Lgals3

1426809_at 1.03209 RIKEN cDNA C430004E15 gene C430004E15Rik
1426810_at 1.15244 jumonji domain containing 1A Jmjd1a
1426811_at 1.00750 "protein phosphatase 2, regulatory subunit B (B56), beta isoform" Ppp2r5b
1426812_a_at 1.02985 RIKEN cDNA 9130404D14 gene 9130404D14Rik
1426813_at 1.05160 LTV1 homolog (S. cerevisiae) Ltv1
1426814_at 0.92964 expressed sequence AU024582 AU024582
1426815_s_at 0.92083 expressed sequence AU024582 AU024582
1426816_at 0.90105 coiled-coil domain containing 64 Ccdc64
1426817_at 1.05301 antigen identified by monoclonal antibody Ki 67 /// similar to Antigen KI-67
Mki67 /// LOC638774
1426818_at 0.76042 arrestin domain containing 4 Arrdc4
1426819_at 0.94875 homeodomain interacting protein kinase 3 Hipk3
1426820_at 0.97323 RIKEN cDNA 2610507B11 gene 2610507B11Rik
1426821_at 0.87765 component of oligomeric golgi complex 8 Cog8
1426822_at 0.91856 "ras homolog gene family, member T2" Rhot2
1426823_s_at 1.09056 "proteasome (prosome, macropain) activator subunit 4" Psme4
1426824_at 1.02661 "proteasome (prosome, macropain) activator subunit 4" Psme4
1426825_at 1.07969 formin-like 3 Fmnl3
1426826_at 1.03913 RNA binding motif protein 16 Rbm16
1426827_at 0.88852 RIKEN cDNA A730098D12 gene A730098D12Rik
1426828_at 0.90634 RIKEN cDNA 1300018I17 gene 1300018I17Rik
1426829_at 1.10842 retinoid X receptor interacting protein 110 Rxrip110
1426830_a_at 0.98658 S-adenosylhomocysteine hydrolase-like 1 Ahcyl1
1426831_at 0.95516 S-adenosylhomocysteine hydrolase-like 1 Ahcyl1
1426832_at 0.78383 RIKEN cDNA 6330505F04 gene 6330505F04Rik
1426833_at 0.97349 "eukaryotic translation initiation factor 4 gamma, 3" Eif4g3
1426834_s_at 0.93292 RIKEN cDNA D930015E06 gene D930015E06Rik
1426835_at 1.07719 methionyl aminopeptidase 1 Metap1
1426836_s_at 1.12332 methionyl aminopeptidase 1 Metap1
1426837_at 1.14196 methionyl aminopeptidase 1 Metap1
1426838_at 1.00041 "polymerase (DNA-directed), delta 3, accessory subunit" Pold3
1426839_at 1.00122 "polymerase (DNA-directed), delta 3, accessory subunit" Pold3
1426840_at 0.97426 YTH domain family 3 Ythdf3
1426841_at 1.03202 YTH domain family 3 Ythdf3
1426842_at 0.94450 YTH domain family 3 Ythdf3
1426843_at 1.05324 RIKEN cDNA 1810013C15 gene 1810013C15Rik
1426844_a_at 1.01115 programmed cell death 2-like Pcd2l
1426845_at 0.95548 programmed cell death 2-like Pcd2l
1426846_at 0.91980 centromere protein T Cenpt
1426847_at 0.92698 sirtuin 4 (silent mating type information regulation 2 homolog) 4 (S.
cerevisiae) Sirt4
1426848_at 1.01745 "SEC24 related gene family, member B (S. cerevisiae)" Sec24b
1426849_at 1.04115 "SEC24 related gene family, member B (S. cerevisiae)" Sec24b

1426850_a_at 0.87502 mitogen activated protein kinase kinase 6 Map2k6
 1426851_a_at 1.30404 nephroblastoma overexpressed gene Nov
 1426852_x_at 1.46551 nephroblastoma overexpressed gene Nov
 1426853_at 1.26757 SET translocation Set
 1426854_a_at 1.07353 SET translocation Set
 1426855_at 0.91747 "DNA segment, Chr 10, ERATO Doi 610, expressed" D10Erttd610e
 1426856_at 0.87369 hydroxysteroid dehydrogenase like 2 Hsd12
 1426857_a_at 0.83636 hydroxysteroid dehydrogenase like 2 Hsd12
 1426858_at 0.51837 inhibin beta-B Inhbb
 1426859_at 0.79621 inhibin beta-B Inhbb
 1426860_at 1.08971 E1A binding protein p400 Ep400
 1426861_at 0.97096 aftiphilin Aftph
 1426862_at 1.00992 aftiphilin Aftph
 1426863_at 0.83131 "RNA binding motif protein, X chromosome" Rbmx
 1426864_a_at 0.67627 neural cell adhesion molecule 1 Ncam1
 1426865_a_at 0.72284 neural cell adhesion molecule 1 Ncam1
 1426866_at 0.91301 dermatan 4 sulfotransferase 1 D4st1
 1426867_at 1.07262 doublesex and mab-3 related transcription factor 2 Dmrt2
 1426868_x_at 1.11707 lamin A Lmna
 1426869_at 1.10357 biregional cell adhesion molecule-related/down-regulated by oncogenes (Cdon)
 binding protein Boc
 1426870_at 0.89451 F-box only protein 33 Fbxo33
 1426871_at 0.86147 F-box only protein 33 Fbxo33
 1426872_at 1.08451 Fc fragment of IgG binding protein Fcgbp
 1426873_s_at 0.85630 junction plakoglobin Jup
 1426874_at 0.95617 enhancer of mRNA decapping 4 Edc4
 1426875_s_at 1.10957 sulfiredoxin 1 homolog (S. cerevisiae) Srxn1
 1426876_at 0.87397 RIKEN cDNA 4732466D17 gene 4732466D17Rik
 1426877_a_at 0.96430 polybromo 1 Pb1
 1426878_at 0.94796 polybromo 1 Pb1
 1426879_at 1.03265 RIKEN cDNA 1190005F20 gene 1190005F20Rik
 1426880_at 0.88277 cDNA sequence BC026657 BC026657
 1426881_at 0.99582 ubiquitin protein ligase E3C Ube3c
 1426882_at 0.97562 ubiquitin protein ligase E3C Ube3c
 1426883_at 0.78617 expressed sequence AW491445 AW491445
 1426884_at 0.92311 required for meiotic nuclear division 5 homolog A (S. cerevisiae) Rmnd5a
 1426885_a_at 0.90676 CDK2 (cyclin-dependent kinase 2)-associated protein 1 Cdk2ap1
 1426886_at 1.03330 "ceroid-lipofuscinosis, neuronal 5" Cln5
 1426887_at 0.89430 nudix (nucleoside diphosphate linked moiety X)-type motif 11 Nudt11
 1426888_at 0.95952 euchromatic histone lysine N-methyltransferase 2 Ehmt2
 1426889_at 0.87523 RIKEN cDNA 4930566A11 gene 4930566A11Rik
 1426890_a_at 1.09384 RNA polymerase II associated protein 1 Rpap1
 1426891_at 0.94840 RNA polymerase II associated protein 1 Rpap1

1426892_at 0.85869 utrophin Utn
1426893_at 0.88783 RIKEN cDNA C230093N12 gene C230093N12Rik
1426894_s_at 0.78102 RIKEN cDNA C230093N12 gene C230093N12Rik
1426895_at 0.87464 zinc finger protein 191 Zfp191
1426896_at 1.02784 zinc finger protein 191 Zfp191
1426897_at 0.97146 regulator of chromosome condensation 2 Rcc2
1426898_at 0.91088 mitogen-activated protein kinase kinase kinase 7 interacting protein 1
Map3k7ip1
1426899_at 0.94601 "TBC1 domain family, member 23" Tbc1d23
1426900_at 1.04163 jumonji domain containing 1C Jmjd1c
1426901_s_at 1.15782 calmodulin binding transcription activator 2 Camta2
1426902_at 0.98207 coenzyme Q6 homolog (yeast) Coq6
1426903_at 1.06114 fibronectin type III domain containing 3a Fndc3a
1426904_s_at 0.93261 "DnaJ (Hsp40) homolog, subfamily C, member 10" Dnajc10
1426905_a_at 0.93309 "DnaJ (Hsp40) homolog, subfamily C, member 10" Dnajc10
1426906_at 1.34391 interferon activated gene 203 Ifi203
1426907_s_at 0.94805 DEAH (Asp-Glu-Ala-Asp/His) box polypeptide 57 Dhx57
1426908_at 1.16746 UDP-N-acetyl-alpha-D-galactosamine: polypeptide N-
acetylgalactosaminyltransferase 7 Galnt7
1426909_at 1.07764 uridine-cytidine kinase 2 Uck2
1426910_at 0.95977 --- ---
1426911_at 0.93643 desmocollin 2 Dsc2
1426912_at 1.02279 ring finger and WD repeat domain 2 /// similar to constitutive
photomorphogenic protein 1 Rfwd2 /// LOC621205
1426913_at 2.03842 lanosterol synthase Lss
1426914_at 1.04522 MARVEL (membrane-associating) domain containing 2 Marveld2
1426915_at 1.00630 death associated protein kinase 1 Dapk1
1426916_at 0.89543 secernin 3 Scrn3
1426917_s_at 0.90275 secernin 3 Scrn3
1426918_at 0.94170 integrin beta 1 (fibronectin receptor beta) Itgb1
1426919_at 1.10555 --- ---
1426920_x_at 0.87320 integrin beta 1 (fibronectin receptor beta) Itgb1
1426921_at 0.99772 "ATP-binding cassette, sub-family F (GCN20), member 1" Abcf1
1426922_s_at 1.02655 HIV-1 Rev binding protein Hrb
1426923_at 1.06052 HIV-1 Rev binding protein Hrb
1426924_at 1.11850 membrane associated DNA binding protein MnaB
1426925_at 0.94754 membrane associated DNA binding protein MnaB
1426926_at 0.83863 "phospholipase C, gamma 2" Plcg2
1426927_at 0.91958 "adaptor-related protein complex 3, beta 2 subunit" Ap3b2
1426928_at 1.04787 coiled-coil domain containing 93 Ccdc93
1426929_at 1.04129 "bruno-like 4, RNA binding protein (Drosophila)" Brunol4
1426930_at 1.04266 "bruno-like 4, RNA binding protein (Drosophila)" Brunol4
1426931_s_at 1.08777 "DNA segment, Chr 19, Brigham & Women's Genetics 1357 expressed /// SWI/

SNF related, matrix associated, actin dependent regulator of chromatin, subfamily a, member 2"

D19Bwg1357e /// Smarca2

1426932_at 1.01968 "SWI/SNF related, matrix associated, actin dependent regulator of chromatin, subfamily a, member 2" Smarca2

1426933_at 1.11189 oxidative-stress responsive 1 Oxsrl

1426934_at 0.82757 NHS-like 1 Nhs1

1426935_at 1.00057 RIKEN cDNA 6330590E21 gene 6330590E21Rik

1426936_at 1.22192 RIKEN cDNA 2610305J24 gene /// cDNA sequence BC005512 /// hypothetical protein LOC215866 /// hypothetical protein LOC629242 /// similar to melanoma antigen /// similar to melanoma antigen /// similar to melanoma antigen /// similar to melanoma antigen /// hypothetical ENV polyprotein /// similar to melanoma antigen /// hypothetical protein LOC669019 /// similar to melanoma antigen /// hypothetical protein LOC669436 /// hypothetical protein LOC670270 /// similar to melanoma antigen /// similar to melanoma antigen /// similar to melanoma antigen /// similar to melanoma antigen /// similar to melanoma antigen /// similar to melanoma antigen /// similar to melanoma antigen /// similar to melanoma antigen /// similar to melanoma antigen /// similar to melanoma antigen /// similar to melanoma antigen /// similar to melanoma antigen 2610305J24Rik /// BC005512 /// LOC215866 /// LOC629242 /// LOC630153 /// LOC630525 /// LOC631386 /// LOC637482 /// LOC641366 /// LOC668904 /// LOC669019 /// LOC669231 /// LOC669436 /// LOC670270 /// LOC670719 /// LOC672061 /// LOC672196 /// LOC672939 /// LOC673876 /// LOC674912 /// LOC674927 /// LOC675168 /// LOC675410 /// LOC675706 /// LOC676247 /// LOC676718 /// LOC677506

1426937_at 0.92073 RIKEN cDNA 6330406I15 gene 6330406I15Rik

1426938_at 0.71507 neuro-oncological ventral antigen 1 /// similar to neuro-oncological ventral antigen 1 isoform 2 Nova1 /// LOC664883

1426939_at 0.94521 "TSR2, 20S rRNA accumulation, homolog (*S. cerevisiae*)" Tsr2

1426940_at 0.84905 "SID1 transmembrane family, member 2" Sidt2

1426941_at 1.01450 --- ---

1426942_at 0.98947 absent in melanoma 1 Aim1

1426943_at 0.93574 integrator complex subunit 1 Ints1

1426944_at 0.85419 F-box and WD-40 domain protein 8 Fbxw8

1426945_at 1.14581 RAN binding protein 5 Ranbp5

1426946_at 1.15267 RAN binding protein 5 Ranbp5

1426947_x_at 0.85493 "procollagen, type VI, alpha 2" Col6a2

1426948_at 1.05219 translocated promoter region Tpr

1426949_s_at 1.07174 translocated promoter region Tpr

1426950_at 0.91218 "poly (ADP-ribose) polymerase family, member 16" Parp16

1426951_at 1.24860 cysteine rich transmembrane BMP regulator 1 (chordin like) Crim1

1426952_at 1.03837 Rho GTPase activating protein 18 Arhgap18

1426953_at 0.98116 high mobility group box 2-like 1 Hmgb2l1

1426954_at 1.00191 tripartite motif protein 33 /// RIKEN cDNA 8030451N04 gene Trim33 /// 8030451N04Rik

1426955_at 1.03000 "procollagen, type XVIII, alpha 1" Col18a1

1426956_a_at 0.95945 transformation related protein 53 binding protein 1 Trp53bp1

1426957_at 0.97536 transformation related protein 53 binding protein 1 Trp53bp1
 1426958_at 1.27748 ribosomal protein S9 /// similar to ribosomal protein S9 Rps9 /// LOC272691
 1426959_at 0.96381 "3-hydroxybutyrate dehydrogenase, type 1" Bdh1
 1426960_a_at 0.99079 fatty acid 2-hydroxylase Fa2h
 1426961_at 0.98243 PHD finger protein 20 Phf20
 1426962_at 1.03919 PHD finger protein 20 Phf20
 1426963_at 1.09402 phosphofurin acidic cluster sorting protein 2 Pacs2
 1426964_at 1.00648 RIKEN cDNA 3110003A17 gene 3110003A17Rik
 1426965_at 0.96446 RAS related protein 2a Rap2a
 1426966_at 1.02729 axin 1 Axin1
 1426967_at 1.02172 axin 1 Axin1
 1426968_a_at 0.86595 retinol dehydrogenase 10 (all-trans) Rdh10
 1426969_at 1.05921 tripartite motif protein 23 Trim23
 1426970_a_at 1.13137 ubiquitin-activating enzyme E1-like /// RIKEN cDNA D330022A01 gene
 Ube11 /// D330022A01Rik
 1426971_at 1.32364 ubiquitin-activating enzyme E1-like Ube11
 1426972_at 0.89795 "SEC24 related gene family, member D (S. cerevisiae)" Sec24d
 1426973_at 0.92286 G protein-coupled receptor 153 Gpr153
 1426974_at 0.85887 RIKEN cDNA 4632413K17 gene 4632413K17Rik
 1426975_at 0.87827 RIKEN cDNA 4632413K17 gene 4632413K17Rik
 1426976_at 0.93425 ubiquitin specific peptidase 47 Usp47
 1426977_at 0.94977 ubiquitin specific peptidase 47 Usp47
 1426978_at 0.90583 "kelch-like 2, Mayven (Drosophila)" Klh12
 1426979_at 0.94473 MLX interacting protein Mlxip
 1426980_s_at 0.97108 RIKEN cDNA E130012A19 gene E130012A19Rik
 1426981_at 0.89110 proprotein convertase subtilisin/kexin type 6 Pcsk6
 1426982_at 0.92055 FLYWCH-type zinc finger 1 Flywch1
 1426983_at 1.14792 formin binding protein 1 Fnbp1
 1426984_at 0.91877 RIKEN cDNA 2310067B10 gene 2310067B10Rik
 1426985_s_at 1.03136 RIKEN cDNA 2810485I05 gene 2810485I05Rik
 1426986_at 0.97774 RIKEN cDNA 2810485I05 gene 2810485I05Rik
 1426987_at 0.95171 RIKEN cDNA 5430417L22 gene 5430417L22Rik
 1426988_at 0.97393 kelch domain containing 5 Klhdc5
 1426989_at 0.88328 calsynenin 3 Clstn3
 1426990_at 0.95273 cubilin (intrinsic factor-cobalamin receptor) Cubn
 1426991_at 0.78229 RIKEN cDNA 1810048J11 gene 1810048J11Rik
 1426992_at 0.83656 xenotropic and polytropic retrovirus receptor 1 Xpr1
 1426993_at 0.79931 xenotropic and polytropic retrovirus receptor 1 Xpr1
 1426994_at 0.85925 PH domain and leucine rich repeat protein phosphatase Phlpp
 1426995_a_at 1.10139 "growth factor, erv1 (S. cerevisiae)-like (augmenter of liver regeneration)"
 Gfer
 1426996_at 0.98720 expressed sequence AA409316 AA409316
 1426997_at 0.92898 thyroid hormone receptor alpha Thra

1426998_at 0.98605 "zinc finger, AN1-type domain 3" Zfand3
 1426999_at 1.02036 zinc finger CCCH type containing 14 Zc3h14
 1427000_at 0.90302 "hepatic nuclear factor 4, alpha" Hnf4a
 1427001_s_at 0.81690 "hepatic nuclear factor 4, alpha" Hnf4a
 1427002_s_at 1.06164 arylsulfatase G Arsg
 1427003_at 1.04907 "protein phosphatase 2, regulatory subunit B (B56), gamma isoform" Ppp2r5c
 1427004_at 0.98484 F-box only protein 2 Fbxo2
 1427005_at 1.18058 polo-like kinase 2 (Drosophila) Plk2
 1427006_at 1.00020 Rap guanine nucleotide exchange factor (GEF) 1 Rapgef1
 1427007_at 0.91719 RIKEN cDNA 1200013B08 gene 1200013B08Rik
 1427008_at 1.11021 ring finger protein 43 Rnf43
 1427009_at 0.96389 "laminin, alpha 5 /// similar to Laminin alpha-5 chain precursor" Lama5 ///
 LOC671909
 1427010_s_at 0.99368 "laminin, alpha 5 /// similar to Laminin alpha-5 chain precursor" Lama5 ///
 LOC671909
 1427011_a_at 0.83396 LanC (bacterial lantibiotic synthetase component C)-like 1 Lancl1
 1427012_at 1.05567 LanC (bacterial lantibiotic synthetase component C)-like 1 Lancl1
 1427013_at 0.97895 carbonic anhydrase 9 Car9
 1427014_at 0.96011 DENN/MADD domain containing 4B Dennd4b
 1427015_at 1.02605 Riken cDNA C230021P08 gene C230021P08Rik
 1427016_at 0.80548 RIKEN cDNA 4932438A13 gene /// similar to CG4841-PA
 4932438A13Rik /// LOC665732
 1427017_at 1.15694 special AT-rich sequence binding protein 2 Satb2
 1427018_at 1.03804 translin-associated factor X (Tsnax) interacting protein 1 Tsnaxip1
 1427019_at 1.02116 "protein tyrosine phosphatase, receptor type Z, polypeptide 1" Ptporz1
 1427020_at 1.04337 "scavenger receptor class A, member 3" Scara3
 1427021_s_at 1.03629 ferritin heavy chain 1 Fth1
 1427022_at 1.05764 DEAD (Asp-Glu-Ala-Asp) box polypeptide 42 Ddx42
 1427023_at 0.90565 phytanoyl-CoA hydroxylase interacting protein-like Phyhip1
 1427024_at 1.19916 RIKEN cDNA 5033405K12 gene 5033405K12Rik
 1427025_at 0.86898 myotubularin related protein 7 Mtmr7
 1427026_at 0.88949 "myosin, heavy polypeptide 4, skeletal muscle" Myh4
 1427027_a_at 0.94982 general transcription factor III A Gtf3a
 1427028_at 0.88978 leucine-rich repeat-containing G protein-coupled receptor 6 Lgr6
 1427029_at 0.86042 HtrA serine peptidase 3 Htra3
 1427030_at 0.81277 coiled-coil domain containing 52 Ccdc52
 1427031_s_at 0.90401 coiled-coil domain containing 52 Ccdc52
 1427032_at 1.05219 hect domain and RLD 4 Herc4
 1427033_at 0.92588 dynamin binding protein Dnmbp
 1427034_at 1.04856 angiotensin I converting enzyme (peptidyl-dipeptidase A) 1 Ace
 1427035_at 1.11224 "solute carrier family 39 (zinc transporter), member 14" Slc39a14
 1427036_a_at 1.10591 "eukaryotic translation initiation factor 4, gamma 1" Eif4g1
 1427037_at 0.95309 "eukaryotic translation initiation factor 4, gamma 1" Eif4g1

1427038_at	0.96420	preproenkephalin 1	Penk1
1427039_at	0.95060	epsin 1	Epn1
1427040_at	0.98009	MyoD family inhibitor domain containing	Mdfic
1427041_at	0.99709	cDNA sequence BC013712	BC013712
1427042_at	0.84150	"mal, T-cell differentiation protein 2"	Mal2
1427043_s_at	0.85051	cytosolic ovarian carcinoma antigen 1	Cova1
1427044_a_at	0.95113	amphiphysin	Amph
1427045_at	0.95720	synaptopodin	Synpo
1427046_at	0.91773	grainyhead-like 2 (Drosophila)	Grhl2
1427047_at	1.03133	nucleoporin 188	Nup188
1427048_at	0.88507	smoothened homolog (Drosophila)	Smo
1427049_s_at	0.91252	smoothened homolog (Drosophila)	Smo
1427050_at	0.79573	RIKEN cDNA 5730420B22 gene	5730420B22Rik
1427051_at	0.98644	tankyrase 1 binding protein 1	Tnks1bp1
1427052_at	0.92054	acetyl-Coenzyme A carboxylase beta	Acacb
1427053_at	1.21470	"ABI gene family, member 3 (NESH) binding protein"	Abi3bp
1427054_s_at	0.91727	"ABI gene family, member 3 (NESH) binding protein"	Abi3bp
1427055_at	0.82923	thioesterase superfamily member 4	Them4
1427056_at	0.94593	"a disintegrin-like and metallopeptidase (reprolysin type) with thrombospondin type 1 motif, 15"	Adamts15
1427057_at	1.11589	RIKEN cDNA C630002B14 gene	C630002B14Rik
1427058_at	1.13181	eukaryotic translation initiation factor 4A1	Eif4a1
1427059_at	0.98645	RIKEN cDNA 4732495E13 gene	4732495E13Rik
1427060_at	0.97068	mitogen activated protein kinase 3	Mapk3
1427061_at	0.91091	retinoblastoma binding protein 8	Rbbp8
1427062_at	0.92821	retinoblastoma binding protein 8	Rbbp8
1427063_at	0.92503	RIKEN cDNA 5330417C22 gene	5330417C22Rik
1427064_a_at	1.05031	scribbled homolog (Drosophila)	Scrib
1427065_at	1.02565	RIKEN cDNA 4933439F18 gene	4933439F18Rik
1427066_a_at	0.90227	RIKEN cDNA 4933439F18 gene	4933439F18Rik
1427067_at	0.92330	RIKEN cDNA 4933439F18 gene	4933439F18Rik
1427068_x_at	1.06905	RIKEN cDNA 4933439F18 gene	4933439F18Rik
1427069_at	0.79328	F-box protein 28	Fbxo28
1427070_at	0.94118	RIKEN cDNA 5730407K14 gene	5730407K14Rik
1427071_at	1.14754	F-box protein 42	Fbxo42
1427072_at	0.61436	START domain containing 8	Stard8
1427073_at	0.84727	lactation elevated 1	Lace1
1427074_at	0.91059	protein-L-isoaspartate (D-aspartate) O-methyltransferase domain containing 2	Pcmdt2
1427075_s_at	0.92043	protein-L-isoaspartate (D-aspartate) O-methyltransferase domain containing 2	Pcmdt2
1427076_at	0.99818	macrophage expressed gene 1 /// similar to macrophage expressed gene 1	Mpeg1 /// LOC671359

1427077_a_at 1.06020 "adaptor-related protein complex 2, beta 1 subunit" Ap2b1
 1427078_at 1.00150 sorting nexin 19 Snx19
 1427079_at 0.93105 "microtubule-associated protein, RP/EB family, member 3" Mapre3
 1427080_at 0.93037 RIKEN cDNA 2610036D13 gene 2610036D13Rik
 1427081_at 1.00747 RIKEN cDNA A630072M18 gene A630072M18Rik
 1427082_at 0.87137 RIKEN cDNA 4632417N05 gene 4632417N05Rik
 1427083_a_at 0.98212 mitogen-activated protein kinase kinase kinase kinase 5 Map4k5
 1427084_a_at 1.01030 mitogen-activated protein kinase kinase kinase kinase 5 Map4k5
 1427085_at 1.03167 RIKEN cDNA 2810432D09 gene 2810432D09Rik
 1427086_at 0.87737 slit homolog 3 (Drosophila) Slit3
 1427087_at 1.16542 LUC7-like 2 (S. cerevisiae) Luc7l2
 1427088_at 0.95358 cyclin T2 Ccnt2
 1427089_at 0.97068 cyclin T2 Ccnt2
 1427090_at 0.95623 "zinc finger, BED domain containing 4" Zbed4
 1427091_at 0.96915 expressed sequence AI481105 AI481105
 1427092_at 1.06846 --- ---
 1427093_at 0.91772 zinc finger protein 707 Zfp707
 1427094_at 0.95247 "polymerase (DNA directed), epsilon 2 (p59 subunit)" Pole2
 1427095_at 1.08519 CUB domain containing protein 1 Cdcpl
 1427096_s_at 0.93317 "signal sequence receptor, delta" Ssr4
 1427097_at 1.01972 WW domain containing E3 ubiquitin protein ligase 1 Wwp1
 1427098_at 0.86029 WW domain containing E3 ubiquitin protein ligase 1 Wwp1
 1427099_at 0.92855 MYC-associated zinc finger protein (purine-binding transcription factor)
 Maz
 1427100_at 1.00691 "meteorin, glial cell differentiation regulator" Metrn
 1427101_at 0.98250 "meteorin, glial cell differentiation regulator" Metrn
 1427102_at 1.00850 schlafen 4 Slfn4
 1427103_at 0.96242 "pleckstrin homology domain containing, family Q member 1" Plekhq1
 1427104_at 0.81106 zinc finger protein 612 Zfp612
 1427105_at 0.93343 RIKEN cDNA 2610510J17 gene 2610510J17Rik
 1427106_at 0.84850 zinc finger and BTB domain containing 3 Zbtb3
 1427107_at 0.86666 "solute carrier family 16 (monocarboxylic acid transporters), member 11"
 Slc16a11
 1427108_at 0.97199 RIKEN cDNA 9530068E07 gene 9530068E07Rik
 1427109_at 0.98964 open reading frame 61 ORF61
 1427110_at 0.99794 "ribonucleoprotein, PTB-binding 1" Raver1
 1427111_s_at 0.98149 "ribonucleoprotein, PTB-binding 1" Raver1
 1427112_at 0.95016 RIKEN cDNA 2700049H19 gene 2700049H19Rik
 1427113_s_at 0.90802 RIKEN cDNA 2700049H19 gene 2700049H19Rik
 1427114_at 0.95943 tetratricopeptide repeat domain 19 Ttc19
 1427115_at 0.95082 "myosin, heavy polypeptide 3, skeletal muscle, embryonic" Myh3
 1427116_at 1.12307 SET domain containing 1A Setd1a
 1427117_at 1.03686 myotubularin related protein 3 Mtmr3

1427118_at	0.95748	keratin 86	krt86	
1427119_at	1.09928	"serine peptidase inhibitor, Kazal type 4"	Spink4	
1427120_at	1.01191	zinc finger protein 26	Zfp26	
1427121_at	1.03067	F-box only protein 4	Fbxo4	
1427122_at	1.02920	"coatamer protein complex, subunit gamma 2, antisense 2"		Copg2as2
1427123_s_at	0.99155	"coatamer protein complex, subunit gamma 2, antisense 2"		Copg2as2
1427124_at	1.06094	RIKEN cDNA 4932416H05 gene	4932416H05Rik	
1427125_s_at	0.92283	leucine rich repeat containing 41	Lrrc41	
1427126_at	0.95029	heat shock protein 1B	Hspa1b	
1427127_x_at	0.85299	heat shock protein 1B	Hspa1b	
1427128_at	0.98986	"protein tyrosine phosphatase, non-receptor type 23"	Ptpn23	
1427129_a_at	0.99099	heterogeneous nuclear ribonucleoprotein R	Hnrpr	
1427130_x_at	0.97875	RIKEN cDNA 1700021K02 gene	1700021K02Rik	
1427131_s_at	0.95421	leucine rich repeat containing 58	Lrrc58	
1427132_at	0.98994	SET binding factor 2	Sbf2	
1427133_s_at	0.96212	low density lipoprotein receptor-related protein 2	Lrp2	
1427134_at	1.43169	"splicing factor, arginine/serine-rich 12"	Sfrs12	
1427135_at	1.46769	"splicing factor, arginine/serine-rich 12"	Sfrs12	
1427136_s_at	1.41384	"splicing factor, arginine/serine-rich 12"	Sfrs12	
1427137_at	0.97224	carboxylesterase 5	Ces5	
1427138_at	0.83240	RIKEN cDNA 0610010D24 gene	0610010D24Rik	
1427139_at	0.81197	"a disintegrin-like and metallopeptidase (reprolysin type) with thrombospondin type 1 motif, 10"	Adamts10	
1427140_at	1.12587	plasmacytoma variant translocation 1	Pvt1	
1427141_at	1.01644	RIKEN cDNA 2700099C18 gene	2700099C18Rik	
1427142_s_at	0.73572	"jumonji, AT rich interactive domain 1B (Rbp2 like)"	Jarid1b	
1427143_at	0.87792	"jumonji, AT rich interactive domain 1B (Rbp2 like)"	Jarid1b	
1427144_at	0.99976	heterogeneous nuclear ribonucleoprotein L-like	Hnrpll	
1427145_at	0.88397	IQ motif and Sec7 domain 1	Iqsec1	
1427146_at	0.83009	expressed sequence AI790298	AI790298	
1427147_at	1.01408	RIKEN cDNA F730047E07 gene	F730047E07Rik	
1427148_at	0.97393	"praja 2, RING-H2 motif containing"	Pja2	
1427149_at	0.82744	"pleckstrin homology domain containing, family A member 6"	Plekha6	
1427150_at	0.87214	myeloid/lymphoid or mixed-lineage leukemia 3	Mll3	
1427151_at	0.82625	glutamine and serine rich 1	Qser1	
1427152_at	0.98953	glutamine and serine rich 1	Qser1	
1427153_at	1.03096	"branched chain ketoacid dehydrogenase E1, beta polypeptide"	Bckdhb	
1427154_at	0.96187	keratin 2	Krt2	
1427155_at	0.92351	FCH and double SH3 domains 1	Fchsd1	
1427156_s_at	0.93445	activating signal cointegrator 1 complex subunit 2	Ascc2	
1427157_at	1.05095	coiled-coil domain containing 85A	Ccdc85a	
1427158_at	0.97802	mitochondrial ribosomal protein S30	Mrps30	
1427159_at	0.99009	cleavage and polyadenylation factor subunit homolog (S. cerevisiae)	Pcf11	

1427160_at 1.00641 cleavage and polyadenylation factor subunit homolog (*S. cerevisiae*) Pcf11
 1427161_at 1.01225 centromere protein F Cenpf
 1427162_a_at 1.03898 "ELK4, member of ETS oncogene family" Elk4
 1427163_at 0.93063 ubiquitin protein ligase E3 component n-recogin 2 Ubr2
 1427164_at 1.07738 "interleukin 13 receptor, alpha 1" Il13ra1
 1427165_at 1.10982 "interleukin 13 receptor, alpha 1" Il13ra1
 1427166_a_at 0.93854 receptor-like tyrosine kinase /// spastic paraplegia 7 homolog (human) Ryk /// Spg7
 1427167_at 0.96202 expressed sequence AI448196 AI448196
 1427168_a_at 0.73570 "procollagen, type XIV, alpha 1" Col14a1
 1427169_at 0.77393 "amyotrophic lateral sclerosis 2 (juvenile) chromosome region, candidate 19 (human)" Als2cr19
 1427170_at 1.03295 "proteasome (prosome, macropain) subunit, alpha type, 8" Psma8
 1427171_at 1.15522 rearranged L-myc fusion sequence Rlf
 1427172_at 0.99295 oral-facial-digital syndrome 1 gene homolog (human) Ofd1
 1427173_a_at 1.00241 mitochondrial ribosomal protein S33 Mrps33
 1427174_at 1.07779 RIKEN cDNA 3100002L24 gene /// RIKEN cDNA 9230108I15 gene /// hypothetical LOC329575 /// similar to gonadotropin inducible ovarian transcription factor 1 /// similar to zinc finger protein 709 /// similar to zinc finger protein 709 /// similar to zinc finger protein 709 /// similar to zinc finger protein 709 /// similar to gonadotropin inducible ovarian transcription factor 1 /// similar to zinc finger protein 97 /// similar to gonadotropin inducible ovarian transcription factor 1 3100002L24Rik /// 9230108I15Rik /// LOC329575 /// LOC626832 /// LOC627914 /// LOC628147 /// LOC665043 /// LOC665078 /// LOC665211 /// LOC668030 /// LOC668039 /// LOC668066
 1427175_at 0.97104 expressed sequence AI428936 AI428936
 1427176_s_at 0.91510 expressed sequence AI428936 AI428936
 1427177_at 0.99398 FYVE and coiled-coil domain containing 1 Fyco1
 1427178_at 1.07945 transmembrane channel-like gene family 4 Tmc4
 1427179_at 0.92616 keratin 33B Krt33b
 1427180_at 0.85197 "solute carrier family 27 (fatty acid transporter), member 3" Slc27a3
 1427181_at 1.08761 "DNA segment, Chr 18, ERATO Doi 653, expressed" D18Ertd653e
 1427182_s_at 0.96888 "DNA segment, Chr 18, ERATO Doi 653, expressed" D18Ertd653e
 1427183_at 1.01062 epidermal growth factor-containing fibulin-like extracellular matrix protein 1 Efemp1
 1427184_at 1.05917 "T-cell receptor beta, variable 13" Tcrb-V13
 1427185_at 0.78190 myocyte enhancer factor 2A Mef2a
 1427186_a_at 0.87223 myocyte enhancer factor 2A Mef2a
 1427187_at 0.85136 RIKEN cDNA 6030413G23 gene 6030413G23Rik
 1427188_at 0.94740 ariadne ubiquitin-conjugating enzyme E2 binding protein homolog 1 (Drosophila) Arih1
 1427189_at 1.02866 ariadne ubiquitin-conjugating enzyme E2 binding protein homolog 1 (Drosophila) Arih1
 1427190_at 0.88787 RIKEN cDNA 2900057D21 gene 2900057D21Rik

1427191_at 0.61311 natriuretic peptide receptor 2 Npr2
1427192_a_at 1.06324 bromodomain containing 8 Brd8
1427193_at 1.31707 bromodomain containing 8 Brd8
1427194_a_at 1.06873 "excision repair cross-complementing rodent repair deficiency, complementation group 8" Ercc8
1427195_at 1.06058 "Mus musculus, clone IMAGE:3983419, mRNA" ---
1427196_at 0.96489 WNK lysine deficient protein kinase 4 Wnk4
1427197_at 1.17536 ataxia telangiectasia and Rad3 related Atr
1427198_at 1.09375 cDNA sequence BC022960 BC022960
1427199_at 0.96146 furry homolog-like (Drosophila) Fryl
1427200_at 0.97361 "zinc finger, RAN-binding domain containing 1" Zranb1
1427201_at 0.90595 "musculoskeletal, embryonic nuclear protein 1" Mustn1
1427202_at 1.00788 RIKEN cDNA 4833442J19 gene 4833442J19Rik
1427203_at 1.10317 myosin XVB Myo15b
1427204_at 0.97889 RIKEN cDNA 3322402L07 gene 3322402L07Rik
1427205_x_at 0.95109 coiled-coil domain containing 46 Ccdc46
1427206_at 1.01355 AFG3(ATPase family gene 3)-like 2 (yeast) Afg3l2
1427207_s_at 1.00276 AFG3(ATPase family gene 3)-like 2 (yeast) Afg3l2
1427208_at 0.95937 zinc finger protein 451 Zfp451
1427209_at 0.94844 "bromodomain adjacent to zinc finger domain, 2A" Baz2a
1427210_at 1.01069 "bromodomain adjacent to zinc finger domain, 2A" Baz2a
1427211_at 1.23789 keratin associated protein 8-1 Krtap8-1
1427212_at 1.03494 mitogen-activated protein kinase associated protein 1 Mapkap1
1427213_at 1.05598 "6-phosphofructo-2-kinase/fructose-2,6-biphosphatase 1" Pfkfb1
1427214_at 0.83246 agmatine ureohydrolase (agmatinase) Agmat
1427215_at 0.96847 acyl-CoA synthetase medium-chain family member 2 Acsm2
1427216_at 0.87410 interferon zeta Ifnz
1427217_at 0.88139 zinc finger protein 455 Zfp455
1427218_at 1.05693 kelch-like 11 (Drosophila) Klhl11
1427219_at 0.96805 --- ---
1427220_a_at 0.97161 seminal vesicle secretion 5 Svs5
1427221_at 1.06014 X transporter protein 3 similar 1 gene Xtrp3s1
1427222_a_at 1.02678 seminal vesicle protein 2 Svp2
1427223_a_at 1.19251 acyl-CoA synthetase medium-chain family member 2 Acsm2
1427224_at 0.97385 Acyl-CoA synthetase medium-chain family member 2 Acsm2
1427225_at 1.23535 epsin 2 Epn2
1427226_at 1.12623 epsin 2 Epn2
1427227_at 1.01440 "gamma-aminobutyric acid (GABA-A) receptor, subunit gamma 1" Gabrg1
1427228_at 0.88653 "palladin, cytoskeletal associated protein" Palld
1427229_at 1.48997 3-hydroxy-3-methylglutaryl-Coenzyme A reductase Hmgcr
1427230_at 0.95068 RIKEN cDNA B930041F14 gene B930041F14Rik
1427231_at 0.76273 roundabout homolog 1 (Drosophila) Robo1
1427232_at 0.77395 teashirt zinc finger family member 1 Tshz1

1427233_at 0.83827 teashirt zinc finger family member 1 Tshz1
 1427234_at 0.92077 "vang-like 1 (van gogh, Drosophila)" Vangl1
 1427235_at 0.95278 "ubiquitously transcribed tetratricopeptide repeat gene, X chromosome" Utx
 1427236_a_at 0.93366 myeloid/lymphoid or mixed-lineage leukemia 5 Mll5
 1427237_at 0.96330 RIKEN cDNA 2410015M20 gene 2410015M20Rik
 1427238_at 0.96229 F-box protein 15 Fbxo15
 1427239_at 0.77406 intraflagellar transport 122 homolog (Chlamydomonas) Ift122
 1427240_at 1.04106 dedicator of cytokinesis 6 /// similar to Dedicator of cytokinesis protein 6
 Dock6 /// LOC670024
 1427241_at 1.00122 poly(A) polymerase gamma Papolg
 1427242_at 0.99548 DEAD (Asp-Glu-Ala-Asp) box polypeptide 4 Ddx4
 1427243_at 0.84133 expressed sequence AA536743 AA536743
 1427244_at 0.88903 tetratricopeptide repeat domain 15 Ttc15
 1427245_at 0.91927 ADP-ribosylation factor GTPase activating protein 1 Arfgap1
 1427246_at 0.85979 "membrane associated guanylate kinase, WW and PDZ domain containing 1"
 Magi1
 1427247_at 1.05045 "DNA segment, Chr 3, Brigham & Women's Genetics 0562 expressed"
 D3Bwg0562e
 1427248_at 0.98736 Wolf-Hirschhorn syndrome candidate 2 (human) Whsc2
 1427249_x_at 1.04564 major urinary protein 3 Mup3
 1427250_at 0.92468 "ATPase, Ca⁺⁺ transporting, cardiac muscle, slow twitch 2" Atp2a2
 1427251_at 0.84145 "ATPase, Ca⁺⁺ transporting, cardiac muscle, slow twitch 2" Atp2a2
 1427252_at 0.99645 "DMRT-like family B with proline-rich C-terminal, 1" Dmrtb1
 1427253_s_at 1.07706 suppressor of zeste 12 homolog (Drosophila) Suz12
 1427254_at 0.98862 zinc finger protein 445 Zfp445
 1427255_s_at 0.92319 zinc finger protein 445 Zfp445
 1427256_at 0.84703 chondroitin sulfate proteoglycan 2 Cspg2
 1427257_at 0.95810 chondroitin sulfate proteoglycan 2 Cspg2
 1427258_at 0.95673 tripartite motif protein 24 Trim24
 1427259_at 0.93642 tripartite motif protein 24 Trim24
 1427260_a_at 0.98019 "tropomyosin 3, gamma" Tpm3
 1427261_at 1.06849 "WW, C2 and coiled-coil domain containing 1" Wwc1
 1427262_at 1.25650 inactive X specific transcripts Xist
 1427263_at 1.13864 inactive X specific transcripts Xist
 1427264_at 0.90998 "crystallin, gamma B" Crygb
 1427265_at 0.94073 breakpoint cluster region homolog Bcr
 1427266_at 1.02190 polybromo 1 Pb1
 1427267_at 1.05911 zinc finger protein 469 Zfp469
 1427268_at 0.97269 filaggrin /// similar to Filaggrin /// similar to Filaggrin Flg /// LOC620355 ///
 LOC675741
 1427269_at 0.93065 "splicing factor, arginine/serine-rich 11" Sfrs11
 1427270_a_at 0.93109 BSD domain containing 1 Bsdc1
 1427271_at 1.11836 BTB (POZ) domain containing 15 Btbd15

1427272_at	1.01497 BTB (POZ) domain containing 15	Btbd15	
1427273_at	1.00454 RIKEN cDNA D130054N24 gene	D130054N24Rik	
1427274_at	0.91620 RIKEN cDNA D130054N24 gene	D130054N24Rik	
1427275_at	0.91640 structural maintenance of chromosomes 4	Smc4	
1427276_at	0.89924 structural maintenance of chromosomes 4	Smc4	
1427277_at	0.98574 sine oculis-related homeobox 1 homolog (Drosophila)	Six1	
1427278_at	1.04875 restin-like 2	Rsnl2	
1427279_at	0.83142 ---	---	
1427280_at	1.07572 "sodium channel, voltage-gated, type II, alpha 1"	Scn2a1	
1427281_at	1.08314 "sodium channel, voltage-gated, type II, alpha 1"	Scn2a1	
1427282_a_at	1.10718 frataxin	Fxn	
1427283_at	1.05755 myeloid/lymphoid or mixed-lineage leukemia 1	Mll1	
1427284_a_at	0.95028 tocopherol (alpha) transfer protein	Ttpa	
1427285_s_at	1.21274 metastasis associated lung adenocarcinoma transcript 1 (non-coding RNA)	Malat1	
1427286_at	0.88590 "DNA segment, Chr 11, Brigham & Women's Genetics 0517 expressed"	D11Bwg0517e	
1427287_s_at	0.73887 "inositol 1,4,5-triphosphate receptor 2"	Itpr2	
1427288_at	1.06928 "amyloid beta (A4) precursor protein-binding, family A, member 2"	Apba2	
1427289_at	0.97563 DDHD domain containing 1	Ddhd1	
1427290_at	1.08612 keratin 81	Krt81	
1427291_at	1.05290 synaptonemal complex protein 1	Sycp1	
1427292_at	0.90246 "immunoglobulin lambda chain, variable 1"	Igl-V1	
1427293_a_at	1.08025 autism susceptibility candidate 2	Auts2	
1427294_a_at	0.91600 RIKEN cDNA 1810073N04 gene	1810073N04Rik	
1427295_at	0.93771 RIKEN cDNA 1810073N04 gene	1810073N04Rik	
1427296_at	1.00799 cDNA sequence BC010304	BC010304	
1427297_at	0.97941 mitochondrial ribosomal protein L9	Mrpl9	
1427298_at	0.95408 "dynamin 3, opposite strand"	Dnm3os	
1427299_at	1.02886 ribosomal protein S6 kinase polypeptide 3	Rps6ka3	
1427300_at	0.95846 LIM homeobox protein 8	Lhx8	
1427301_at	1.04857 CD48 antigen	Cd48	
1427302_at	0.96475 ectonucleotide pyrophosphatase/phosphodiesterase 3	Enpp3	
1427303_at	1.05418 ectonucleotide pyrophosphatase/phosphodiesterase 3	Enpp3	
1427304_at	0.99450 vacuolar protein sorting 18 (yeast)	Vps18	
1427305_at	1.23459 "phosphatidylinositol glycan anchor biosynthesis, class A"	Piga	
1427306_at	1.05870 "ryanodine receptor 1, skeletal muscle"	Ryr1	
1427307_a_at	1.08468 disabled homolog 1 (Drosophila)	Dab1	
1427308_at	1.03045 disabled homolog 1 (Drosophila)	Dab1	
1427309_at	0.97574 prolyl-tRNA synthetase (mitochondrial)(putative)	Pars2	
1427310_at	1.17648 fetal Alzheimer antigen Falz		
1427311_at	0.97663 fetal Alzheimer antigen Falz		
1427312_at	0.90695 cardiomyopathy associated 5	Cmya5	

1427313_at 1.00246 prostaglandin I receptor (IP) Ptgir
1427314_at 1.00075 transmembrane emp24 protein transport domain containing 7 Tmed7
1427315_at 0.99476 transmembrane emp24 protein transport domain containing 7 Tmed7
1427316_s_at 0.76647 EST AA881470 AA881470
1427317_at 0.94882 antigenic determinant of rec-A protein Kin
1427318_s_at 1.10815 "dysferlin /// fer-1-like 3, myoferlin (C. elegans)" Dysf /// Fer113
1427319_at 1.18169 RIKEN cDNA A230046K03 gene A230046K03Rik
1427320_at 1.06813 "coatomer protein complex, subunit gamma 2, antisense 2" Copg2as2
1427321_s_at 1.03875 coxsackievirus and adenovirus receptor Cxadr
1427322_at 0.90267 bromodomain and WD repeat domain containing 1 Brwd1
1427323_s_at 0.96359 "WD repeat domain, phosphoinositide interacting 1" Wipi1
1427324_at 0.81930 cDNA sequence BC022593 BC022593
1427325_s_at 1.00554 expressed sequence AI597013 AI597013
1427326_at 1.08752 RIKEN cDNA 4732471D19 gene 4732471D19Rik
1427327_at 1.07174 paired immunoglobulin-like type 2 receptor alpha Pilra
1427328_a_at 0.86904 CLIP associating protein 2 Clasp2
1427329_a_at 1.00858 immunoglobulin heavy chain 6 (heavy chain of IgM) Igh-6
1427330_at 0.92703 cysteinyl-tRNA synthetase Cars
1427331_at 0.92613 adenosine A1 receptor Adora1
1427332_at 0.97177 "mediator of RNA polymerase II transcription, subunit 19 homolog (yeast)"
Med19
1427333_s_at 1.04563 "splicing factor, arginine/serine-rich 15 /// similar to splicing factor, arginine/
serine-rich 15" Sfrs15 /// LOC670368
1427334_s_at 1.06201 RIKEN cDNA 2810474O19 gene 2810474O19Rik
1427335_at 0.94892 RIKEN cDNA 6720456H20 gene 6720456H20Rik
1427336_at 0.95536 "aldehyde dehydrogenase 8 family, member A1" Aldh8a1
1427337_at 0.99032 "aldehyde dehydrogenase 8 family, member A1" Aldh8a1
1427338_at 0.90368 "ciliary rootlet coiled-coil, rootletin" Crocc
1427339_at 0.93425 "solute carrier family 30 (zinc transporter), member 2" Slc30a2
1427340_at 0.89317 regulator of G-protein signaling 6 Rgs6
1427341_at 0.87674 RIKEN cDNA E130103I17 gene E130103I17Rik
1427342_at 1.01206 FAST kinase domains 1 Fastkd1
1427343_at 0.97835 "RASD family, member 2" Rasd2
1427344_s_at 1.10642 "RASD family, member 2" Rasd2
1427345_a_at 0.76534 "sulfotransferase family 1A, phenol-preferring, member 1" Sult1a1
1427346_at 0.99118 ovary testis transcribed /// similar to ovary testis transcribed /// similar to ovary
testis transcribed /// similar to ovary testis transcribed /// similar to ovary testis transcribed /// similar to
ovary testis transcribed Ott /// LOC434863 /// LOC434864 /// LOC434865 /// LOC666184 ///
LOC666260
1427347_s_at 0.89628 "tubulin, beta 2a /// tubulin, beta 2b /// similar to tubulin, beta 3 /// similar to
tubulin, beta 3" Tubb2a /// Tubb2b /// LOC665524 /// LOC671656
1427348_at 1.55759 zinc finger CCCH type containing 12A Zc3h12a
1427349_x_at 0.98171 RIKEN cDNA 2810021G02 gene 2810021G02Rik

1427350_a_at 0.98231 GTP binding protein 6 (putative) Gtpbp6
1427351_s_at 0.98935 immunoglobulin heavy chain 6 (heavy chain of IgM) Igh-6
1427352_at 0.82275 cDNA sequence BC031593 BC031593
1427353_at 1.50699 CLIP associating protein 1 Clasp1
1427354_at 0.92604 homeo box A4 Hoxa4
1427355_at 1.07849 "calcitonin/calcitonin-related polypeptide, alpha" Calca
1427356_at 0.82387 RIKEN cDNA 2310031A18 gene 2310031A18Rik
1427357_at 1.10336 cytidine deaminase Cda
1427358_a_at 1.09128 death associated protein kinase 1 Dapk1
1427359_at 1.04666 RIKEN cDNA A630082K20 gene A630082K20Rik
1427360_at 0.89851 RIKEN cDNA 4930507D05 gene 4930507D05Rik
1427361_at 0.92098 homeo box C6 Hoxc6
1427362_x_at 0.91702 homeo box C6 Hoxc6
1427363_at 2.21629 Myosin X Myo10
1427364_a_at 1.08193 "ornithine decarboxylase, structural 1 /// similar to Ornithine decarboxylase (ODC) /// similar to Ornithine decarboxylase (ODC) /// similar to Ornithine decarboxylase (ODC) /// similar to Ornithine decarboxylase (ODC) /// hypothetical protein LOC671973 /// similar to Ornithine decarboxylase (ODC)" Odc1 /// LOC639787 /// LOC666231 /// LOC668343 /// LOC670279 /// LOC671154 /// LOC671973 /// LOC673302
1427365_at 0.90383 keratin 86 krt86
1427366_at 0.99444 keratin associated protein 3-1 Krtap3-1
1427367_at 0.89779 EF-hand calcium binding domain 4A Efcab4a
1427368_x_at 0.88673 feline sarcoma oncogene Fes
1427369_at 1.01184 "NACHT, leucine rich repeat and PYD containing 6" Nalp6
1427370_at 0.86771 amidohydrolase domain containing 1 Amdhd1
1427371_at 0.87948 "ATP-binding cassette, sub-family A (ABC1), member 8a" Abca8a
1427372_at 0.96152 "cytochrome P450, family 27, subfamily b, polypeptide 1" Cyp27b1
1427373_at 1.05027 adhesion molecule with Ig like domain 1 Amigo1
1427374_at 0.99967 "mucin 3, intestinal /// similar to mucin 17 /// similar to mucin 17" Muc3 /// LOC666339 /// LOC677034
1427375_at 1.11905 RNA (guanine-9-) methyltransferase domain containing 2 Rg9mtd2
1427376_a_at 1.01935 mitogen-activated protein kinase kinase kinase 5 Map4k5
1427377_x_at 0.93296 "hydroxy-delta-5-steroid dehydrogenase, 3 beta- and steroid delta-isomerase 3" Hsd3b3
1427378_at 0.85705 keratin 75 Krt75
1427379_at 1.20362 Neuropathy target esterase Nte
1427380_at 1.02187 kallikrein 1-related peptidase b3 Klk1b3
1427381_at 0.94331 immunoresponsive gene 1 Irg1
1427382_a_at 1.04530 suppressor of variegation 3-9 homolog 1 (Drosophila) Suv39h1
1427383_at 0.94258 Iroquois related homeobox 6 (Drosophila) Irx6
1427384_at 0.85728 chromodomain helicase DNA binding protein 6 Chd6
1427385_s_at 1.02366 "actinin, alpha 1" Actn1
1427386_at 0.99668 Rho guanine nucleotide exchange factor (GEF) 16 Arhgef16

1427387_a_at 0.75938 integrin beta 4 Itgb4
1427388_at 0.98564 leucine rich repeat containing 2 Lrrc2
1427389_at 1.17753 cDNA sequence BC021785 BC021785
1427390_at 1.00072 "biogenesis of lysosome-related organelles complex-1, subunit 3" Bloc1s3
1427391_a_at 0.84560 "procollagen, type XII, alpha 1" Col12a1
1427392_at 0.99629 Down syndrome cell adhesion molecule-like 1 Dscaml1
1427393_at 1.00203 coagulation factor IX F9
1427394_at 1.01260 "insulin-like growth factor 2, antisense" Igf2as
1427395_a_at 0.84762 "aldehyde dehydrogenase family 1, subfamily A3" Aldh1a3
1427396_a_at 0.97013 "cold shock domain containing E1, RNA binding" Csde1
1427397_at 1.13597 RIKEN cDNA 2810046L04 gene 2810046L04Rik
1427398_at 0.96703 mucin 4 Muc4
1427399_a_at 0.88528 nuclear RNA export factor 7 Nxf7
1427400_at 0.95191 ladybird homeobox homolog 1 (Drosophila) Lbx1
1427401_at 0.97220 "cholinergic receptor, nicotinic, alpha polypeptide 5" Chrna5
1427402_at 0.96047 polymerase (RNA) II (DNA directed) polypeptide F Polr2f
1427403_at 0.92775 "solute carrier organic anion transporter family, member 1a5" Slco1a5
1427404_x_at 1.04680 "similar to enolase 1, alpha non-neuron" LOC433182
1427405_s_at 0.93581 RAB11 family interacting protein 5 (class I) Rab11fip5
1427406_at 1.09055 thyroid hormone receptor interactor 11 Trip11
1427407_s_at 0.98767 thyroid hormone receptor interactor 11 Trip11
1427408_a_at 0.96973 thyroid hormone receptor associated protein 3 Thrap3
1427409_at 0.92267 membrane-associated ring finger (C3HC4) 9 9-Mar
1427410_at 0.54320 "deleted in lymphocytic leukemia, 2" Dleu2
1427411_s_at 0.49405 "deleted in lymphocytic leukemia, 2" Dleu2
1427412_s_at 0.96650 Rap guanine nucleotide exchange factor (GEF) 6 Rapgef6
1427413_a_at 1.17879 "CUG triplet repeat, RNA binding protein 1" Cugbp1
1427414_at 1.00148 "protein kinase, cAMP dependent regulatory, type II alpha" Prkar2a
1427415_at 1.04510 F-box protein 43 Fbxo43
1427416_x_at 1.03900 similar to dual specificity phosphatase 7 LOC674944
1427417_at 1.13222 sex comb on midleg-like 4 (Drosophila) Scml4
1427418_a_at 1.17056 "hypoxia inducible factor 1, alpha subunit" Hif1a
1427419_x_at 1.03888 chemokine (C-C motif) receptor 9 Ccr9
1427420_at 0.98956 "NK6 transcription factor related, locus 2 (Drosophila)" Nkx6-2
1427421_at 1.03584 t-complex protein 10b Tcp10b
1427422_at 0.93612 similar to hepatocellular carcinoma-associated gene TD26 LOC624219
1427423_at 0.99185 RIKEN cDNA 2810410C14 gene 2810410C14Rik
1427424_at 1.07788 UDP-N-acetyl-alpha-D-galactosamine:polypeptide N-acetylgalactosaminyltransferase 6 Galnt6
1427425_at 1.22119 RIKEN cDNA 9130208E07 gene 9130208E07Rik
1427426_at 1.00339 "potassium voltage-gated channel, subfamily Q, member 5" Kcnq5
1427427_at 0.97669 ryanodine receptor 3 Ryr3
1427428_at 1.02378 "C-type lectin domain family 4, member g" Clec4g

1427429_at 1.36065 colony stimulating factor 2 (granulocyte-macrophage) Csf2
1427430_at 0.86648 expressed sequence AI848100 AI848100
1427431_at 0.89145 --- ---
1427432_a_at 1.07096 "splicing factor, arginine/serine-rich 10 (transformer 2 homolog, Drosophila)" Sfrs10
1427433_s_at 0.94992 homeo box A3 Hoxa3
1427434_at 0.91360 --- ---
1427435_at 1.03738 --- ---
1427436_at 0.88935 sine oculis-related homeobox 2 homolog (Drosophila) Six2
1427437_at 0.80356 RIKEN cDNA 2610203C20 gene 2610203C20Rik
1427438_at 1.08972 "Phosphatidylinositol-4-phosphate 5-kinase, type II, beta" Pip5k2b
1427439_s_at 1.11195 protein arginine N-methyltransferase 5 Prmt5
1427440_a_at 0.64371 afamin Afm
1427441_a_at 0.95987 "succinate-Coenzyme A ligase, GDP-forming, beta subunit" Suclg2
1427442_a_at 0.86711 amyloid beta (A4) precursor protein App
1427443_at 1.11731 Hypothetical protein LOC625794 LOC625794
1427444_at 0.86067 "ATP-binding cassette, sub-family F (GCN20), member 1" Abcf1
1427445_a_at 1.16473 titin Ttn
1427446_s_at 1.05038 titin Ttn
1427447_a_at 0.91297 TRIO and F-actin binding protein Triobp
1427448_at 0.97197 "rabaptin, RAB GTPase binding effector protein 1" Rabep1
1427449_a_at 0.92444 ADP-ribosylhydrolase like 2 Adprhl2
1427450_x_at 1.01997 myosin IB Myo1b
1427451_a_at 0.89082 cDNA sequence BC018473 BC018473
1427452_at 1.01207 ribosomal protein L7 Rpl7
1427453_at 0.96311 cDNA sequence BC018473 BC018473
1427454_at 1.01830 homeo box C6 Hoxc6
1427455_x_at 1.01837 "immunoglobulin kappa chain, constant region /// immunoglobulin kappa chain variable 28 (V28) /// immunoglobulin kappa chain variable 8 (V8)-16" Igk-C /// Igk-V28 /// Igk-V8-16
1427456_at 0.85142 WD repeat and FYVE domain containing 3 Wdfy3
1427457_a_at 0.91023 bone morphogenetic protein 1 Bmp1
1427458_at 0.94252 bone morphogenetic protein 1 Bmp1
1427459_at 0.80399 "carboxypeptidase N, polypeptide 2" Cpn2
1427460_at 1.02664 "TAF4A RNA polymerase II, TATA box binding protein (TBP)-associated factor" Taf4a
1427461_at 1.02167 cDNA sequence BC005561 BC005561
1427462_at 1.21825 E2F transcription factor 3 E2f3
1427463_at 0.87106 CAS1 domain containing 1 Casd1
1427464_s_at 0.94173 heat shock 70kD protein 5 (glucose-regulated protein) Hspa5
1427465_at 0.83207 "ATPase, Na+/K+ transporting, alpha 2 polypeptide" Atp1a2
1427466_at 1.01254 CDC91 cell division cycle 91-like 1 (S. cerevisiae) Cdc9111
1427467_a_at 0.93425 retinitis pigmentosa GTPase regulator Rpgr

1427468_at 0.94621 "protein phosphatase 3, catalytic subunit, beta isoform" Ppp3cb
1427469_at 0.86874 helicase with zinc finger domain Helz
1427470_s_at 1.14205 N-ethylmaleimide sensitive fusion protein attachment protein beta Napb
1427471_at 0.83286 F-box and leucine-rich repeat protein 3 Fbxl3
1427472_a_at 0.92840 "complement component 8, beta subunit" C8b
1427473_at 0.98389 "glutathione S-transferase, mu 3" Gstm3
1427474_s_at 0.94345 "glutathione S-transferase, mu 3" Gstm3
1427475_a_at 0.89486 similar to PDZ and LIM domain protein 5 (Enigma homolog) (Enigma-like
PDZ and LIM domains protein) LOC669660
1427476_a_at 0.96250 tripartite motif protein 32 Trim32
1427477_at 1.05184 "transmembrane protease, serine 13" Tmprss13
1427478_at 0.88857 ubiquitin specific peptidase 12 Usp12
1427479_at 1.09634 eukaryotic translation initiation factor 1A /// similar to Eukaryotic translation
initiation factor 1A (eIF-1A) (eIF-4C) /// similar to GLE1 RNA export mediator-like (yeast /// similar to
Eukaryotic translation initiation factor 1A (eIF-1A) (eIF-4C) /// similar to Eukaryotic translation
initiation factor 1A (eIF-1A) (eIF-4C) /// similar to X-linked eukaryotic translation initiation factor
1A /// similar to X-linked eukaryotic translation initiation factor 1A /// similar to X-linked eukaryotic
translation initiation factor 1A /// similar to X-linked eukaryotic translation initiation factor 1A ///
similar to X-linked eukaryotic translation initiation factor 1A /// similar to X-linked eukaryotic
translation initiation factor 1A /// similar to X-linked eukaryotic translation initiation factor 1A ///
similar to X-linked eukaryotic translation initiation factor 1A /// similar to X-linked eukaryotic
translation initiation factor 1A /// similar to X-linked eukaryotic translation initiation factor 1A
Eif1a /// MGC107533 /// LOC432715 /// LOC435337 /// LOC544881 /// LOC544883 /// LOC546233 ///
LOC623167 /// LOC627878 /// LOC627881 /// LOC630607 /// LOC640794 /// LOC641136 ///
LOC666806 /// LOC666862
1427480_at 0.99967 liver-expressed antimicrobial peptide 2 Leap2
1427481_a_at 1.02657 "ATPase, Na+/K+ transporting, alpha 3 polypeptide" Atp1a3
1427482_a_at 1.06432 carbonic anhydrase 8 /// similar to Carbonic anhydrase-related protein (CARP)
(CA-VIII) Car8 /// LOC676792
1427483_at 1.06357 "solute carrier family 25 (mitochondrial carrier, phosphate carrier), member
24" Slc25a24
1427484_at 1.16592 echinoderm microtubule associated protein like 5 Eml5
1427485_at 0.93639 leiomodulin 1 (smooth muscle) Lmod1
1427486_at 1.26490 "protein tyrosine phosphatase, receptor type, B" Ptpnb2
1427487_at 1.04107 --- ---
1427488_a_at 0.99181 baculoviral IAP repeat-containing 6 Birc6
1427489_at 0.95521 integrin alpha 8 Itga8
1427490_at 0.92982 "ATP-binding cassette, sub-family B (MDR/TAP), member 7" Abcb7
1427491_at 1.22705 NOL1/NOP2/Sun domain family 6 Nsun6
1427492_at 0.99974 premature ovarian failure 1B Pof1b
1427493_at 0.95886 sodium channel and clathrin linker 1 Sclt1
1427494_at 0.95249 homeo box B7 /// homeo box B8 Hoxb7 /// Hoxb8
1427495_at 0.89405 "sodium channel, voltage-gated, type VII, alpha" Scn7a

1427496_at 1.00069 centrosomal protein 152 Cep152
1427497_at 0.92739 RIKEN cDNA 2610015P09 gene 2610015P09Rik
1427498_a_at 0.98639 sperm associated antigen 5 Spag5
1427499_at 0.86380 zinc finger protein 81 Zfp81
1427500_at 1.01292 signal peptidase complex subunit 1 homolog (*S. cerevisiae*) Spcs1
1427501_at 0.97816 cDNA sequence BC007180 BC007180
1427502_at 1.06245 a disintegrin and metallopeptidase domain 4 Adam4
1427503_at 0.92163 expressed sequence AI324046 AI324046
1427504_s_at 0.99464 "splicing factor, arginine/serine-rich 2 (SC-35)" Sfrs2
1427505_a_at 0.82794 CASP2 and RIPK1 domain containing adaptor with death domain Cradd
1427506_at 1.06579 peptidylprolyl isomerase (cyclophilin) like 5 Ppil5
1427507_at 0.91527 tripartite motif protein 17 Trim17
1427508_at 0.97757 arylsulfatase i Arsi
1427509_at 0.98604 BAI1-associated protein 3 Baiap3
1427510_at 0.49545 "guanine nucleotide binding protein, alpha inhibiting 1" Gnai1
1427511_at 1.56679 Beta-2 microglobulin B2m
1427512_a_at 1.02173 "laminin, alpha 3 /// similar to Laminin alpha-3 chain precursor (Nicein alpha subunit)" Lama3 /// LOC669814
1427513_at 1.12925 cDNA sequence BC024137 BC024137
1427514_at 1.06121 hypothetical protein LOC624295 LOC624295
1427515_at 1.07314 RIKEN cDNA A530088I07 gene A530088I07Rik
1427516_a_at 0.98062 biregional cell adhesion molecule-related/down-regulated by oncogenes (Cdon) binding protein Boc
1427517_at 0.91299 biregional cell adhesion molecule-related/down-regulated by oncogenes (Cdon) binding protein Boc
1427518_at 0.96108 cDNA sequence D10627 D10627
1427519_at 0.86525 adenosine A2a receptor Adora2a
1427520_a_at 0.90326 "myosin, heavy polypeptide 1, skeletal muscle, adult" Myh1
1427521_a_at 1.03797 RIKEN cDNA 9930023K05 gene 9930023K05Rik
1427522_at 0.88297 Rho GTPase activating protein 20 Arhgap20
1427523_at 0.93012 sine oculis-related homeobox 3 homolog (*Drosophila*) /// RIKEN cDNA E130112M24 gene Six3 /// E130112M24Rik
1427524_a_at 0.98000 RIKEN cDNA 4930548G07 gene 4930548G07Rik
1427525_at 1.00663 CD151 antigen Cd151
1427526_at 1.07978 FGFR1 oncogene partner 2 Fgfr1op2
1427527_a_at 1.03643 parathyroid hormone-like peptide Pthlh
1427528_a_at 0.66881 Eph receptor A7 EphA7
1427529_at 0.81426 frizzled homolog 9 (*Drosophila*) Fzd9
1427530_at 1.00702 bicaudal D homolog 1 (*Drosophila*) Bicd1
1427531_a_at 0.95176 "solute carrier family 22 (organic cation transporter), member 18" Slc22a18
1427532_at 0.95984 T cell receptor associated transmembrane adaptor 1 Trat1
1427533_at 1.44959 ALS2 C-terminal like Als2cl
1427534_at 0.95117 RIKEN cDNA 4930535I16 gene 4930535I16Rik

1427535_s_at 0.86454 expressed sequence AW822216 AW822216
1427536_at 0.95893 "Adult male olfactory brain cDNA, RIKEN full-length enriched library, clone:6430562O12 product:hypothetical protein, full insert sequence" ---
1427537_at 0.90665 epiplakin 1 Eppk1
1427538_at 0.91051 zinc finger protein 369 Zfp369
1427539_a_at 1.19052 ZW10 interactor Zwint
1427540_at 1.37087 ZW10 interactor Zwint
1427541_x_at 0.98191 hyaluronan mediated motility receptor (RHAMM) Hmnr
1427542_at 0.87089 RIKEN cDNA 5330439J01 gene 5330439J01Rik
1427543_s_at 0.97960 "ubiquitin-activating enzyme E1, Chr Y 1" Ube1y1
1427544_a_at 1.34219 poly (A) polymerase alpha Papola
1427545_at 1.01405 sex comb on midleg-like 4 (Drosophila) Scml4
1427546_at 1.05959 "ATP-binding cassette, sub-family A (ABC1), member 8b" Abca8b
1427547_a_at 1.03401 "solute carrier family 26, member 3" Slc26a3
1427548_a_at 1.09747 "chloride channel, nucleotide-sensitive, 1A" Clns1a
1427549_s_at 0.95543 keratin associated protein 16-10 /// keratin associated protein 16-3 /// keratin associated protein 16-6 Krtap16-10 /// Krtap16-3 /// Krtap16-6
1427550_at 0.98827 paternally expressed 10 Peg10
1427551_at 0.97620 ubiquitin specific peptidase 29 Usp29
1427552_a_at 0.95573 glutathione transferase zeta 1 (maleylacetoacetate isomerase) Gstz1
1427553_at 1.08223 hypothetical protein LOC628926 LOC628926
1427554_at 1.00277 "helicase, mus308-like (Drosophila)" Hel308
1427555_at 0.89991 cDNA sequence BC032281 BC032281
1427556_at 0.98497 "myosin, light polypeptide kinase 2, skeletal muscle /// similar to Myosin light chain kinase 2, skeletal/cardiac muscle (MLCK2)" Mylk2 /// LOC669275
1427557_at 0.99175 "asparagine-linked glycosylation 12 homolog (yeast, alpha-1,6-mannosyltransferase)" Alg12
1427558_s_at 0.95565 "asparagine-linked glycosylation 12 homolog (yeast, alpha-1,6-mannosyltransferase)" Alg12
1427559_a_at 1.08652 activating transcription factor 2 Atf2
1427560_at 0.99687 sine oculis-related homeobox 5 homolog (Drosophila) Six5
1427561_a_at 0.70387 afamin Afm
1427562_a_at 1.09682 "protein kinase C, alpha" Prkca
1427563_at 0.91821 "distal-less homeobox 6, antisense" Dlx6as
1427564_at 1.27548 diaphanous homolog 2 (Drosophila) Diap2
1427565_a_at 1.02686 "ATP-binding cassette, sub-family C (CFTR/MRP), member 5" Abcc5
1427566_at 1.06788 "WAP, FS, Ig, KU, and NTR-containing protein 1" Wfikn1
1427567_a_at 1.02053 "tropomyosin 3, gamma" Tpm3
1427568_a_at 0.81472 intraflagellar transport 80 homolog (Chlamydomonas) Ift80
1427569_a_at 1.12048 utrophin Utrn
1427570_at 1.04266 signal recognition particle receptor ('docking protein') Srpr
1427571_at 1.08973 sonic hedgehog Shh
1427572_at 1.12720 Leucine rich repeat containing 51 Lrrc51

1427573_at 1.03243 cysteine-rich hydrophobic domain 1 Chic1
 1427574_s_at 1.16237 SH3 domain protein D19 Sh3d19
 1427575_at 0.96472 F-box and WD-40 domain protein 14 Fbxw14
 1427576_at 0.98005 immunoglobulin kappa chain variable 28 (V28) /// immunoglobulin kappa chain variable 8 (V8) /// immunoglobulin kappa chain /// immunoglobulin kappa chain variable 21 (V21) Igk-V28 /// Igk-V8 /// LOC545854 /// Igk-V21
 1427577_x_at 0.79810 immunoglobulin kappa chain variable 28 (V28) /// immunoglobulin kappa chain variable 8 (V8) /// immunoglobulin kappa chain /// immunoglobulin kappa chain variable 6-23 /// immunoglobulin kappa chain variable 19 (V19)-17 /// immunoglobulin kappa chain variable 19 (V19)-14 /// immunoglobulin kappa chain variable 19 (V19)-13 /// similar to Ig kappa chain V-V region MPC11 precursor /// similar to Ig kappa chain V-V region MPC11 precursor /// similar to Ig kappa chain V-V region MPC11 precursor /// similar to Ig kappa chain V-V region MPC11 precursor /// immunoglobulin kappa chain variable 21 (V21) Igk-V28 /// Igk-V8 /// LOC545854 /// Igkv6-23 /// Igk-V19-17 /// Igk-V19-14 /// Igk-V19-13 /// LOC669070 /// LOC672450 /// LOC673123 /// LOC676136 /// LOC676175 /// Igk-V21
 1427578_a_at 1.14985 integrin beta 4 binding protein Itgb4bp
 1427579_at 0.85712 "rhomoid, veinlet-like 3 (Drosophila)" Rhbdl3
 1427580_a_at 1.06130 RNA imprinted and accumulated in nucleus Rian
 1427581_at 0.96078 hypermethylated in cancer 2 Hic2
 1427582_at 0.89416 fibroblast growth factor 6 Fgf6
 1427583_at 0.89820 RIKEN cDNA 4921505C17 gene 4921505C17Rik
 1427584_at 0.86956 angiominin Amot
 1427585_at 1.02524 --- ---
 1427586_at 1.04861 "sema domain, immunoglobulin domain (Ig), transmembrane domain (TM) and short cytoplasmic domain, (semaphorin) 4B" Sema4b
 1427587_at 1.08051 zinc finger protein 28 Zfp28
 1427588_a_at 0.96019 dystrobrevin alpha Dtna
 1427589_at 0.93731 RIKEN cDNA 8430419L09 gene 8430419L09Rik
 1427590_at 0.92281 zinc finger protein 39 Zfp39
 1427591_at 0.95456 chloride channel 1 Clcn1
 1427592_at 0.97883 protocadherin 7 Pcdh7
 1427593_at 0.87797 tripartite motif protein 8 Trim8
 1427594_at 1.05636 zinc finger protein 697 Zfp697
 1427595_at 1.15444 acetyl-Coenzyme A carboxylase alpha /// similar to acetyl-coenzyme A carboxylase alpha Acaca /// LOC672245
 1427596_at 0.93575 "X (inactive)-specific transcript, antisense" Tsix
 1427597_at 1.08973 "pleckstrin homology domain containing, family H (with MyTH4 domain) member 2" Plekhh2
 1427598_at 1.03897 Prader-Willi chromosome region 1 homolog (human) Pwcr1
 1427599_at 0.97251 "bol, boule-like (Drosophila)" Boll
 1427600_at 1.01086 "Tumor necrosis factor receptor superfamily, member 19" Tnfrsf19
 1427601_at 1.17694 RIKEN cDNA 9430022F06 gene 9430022F06Rik
 1427602_at 1.16764 diabetic embryopathy 1 Dep1

1427603_at 1.02226 activating transcription factor 7 interacting protein 2 Atf7ip2
 1427604_a_at 0.97990 "ATPase, class II, type 9A" Atp9a
 1427605_at 0.97885 homeo box B3 Hoxb3
 1427606_at 1.01268 "ubiquitin specific peptidase 27, X chromosome" Usp27x
 1427607_at 0.94985 "calcium channel, voltage-dependent, T type, alpha 1H subunit" Cacna1h
 1427608_a_at 0.95129 "T-cell receptor gamma, constant region" Tcr-g-C
 1427609_at 0.91741 SRY-box containing gene 14 Sox14
 1427610_at 0.64304 desmoplakin Dsp
 1427611_at 1.03225 growth hormone releasing hormone receptor Ghrhr
 1427612_at 1.06995 defensin beta 9 Defb9
 1427613_at 0.90286 cDNA sequence AY074887 AY074887
 1427614_at 1.00879 prolactin induced protein Pip
 1427615_at 1.06849 integrin alpha 4 Itga4
 1427616_at 0.97917 "Mus musculus, clone IMAGE:5362978" ---
 1427617_at 0.94173 fucosyltransferase 10 Fut10
 1427618_at 0.96679 cadherin 9 Cdh9
 1427619_a_at 0.95352 SH3 domain and tetratricopeptide repeats 1 Sh3tc1
 1427620_at 1.00659 p53-associated parkin-like cytoplasmic protein Parc
 1427621_at 0.99703 Low density lipoprotein receptor-related protein 2 Lrp2
 1427622_at 0.97970 maternal embryonic message 1 Mem1
 1427623_at 0.96610 hect (homologous to the E6-AP (UBE3A) carboxyl terminus) domain and
 RCC1 (CHC1)-like domain (RLD) 2 Herc2
 1427624_s_at 1.01050 interleukin 22 /// interleukin 10-related T cell-derived inducible factor beta
 Il22 /// Iltifb
 1427625_a_at 0.83161 hect (homologous to the E6-AP (UBE3A) carboxyl terminus) domain and
 RCC1 (CHC1)-like domain (RLD) 2 Herc2
 1427626_at 0.92828 "mucin 5, subtype B, tracheobronchial /// similar to mucin 5, subtype B,
 tracheobronchial" Muc5b /// LOC640195
 1427627_at 1.06744 "Solute carrier family 34 (sodium phosphate), member 1" Slc34a1
 1427628_at 0.95925 "T-cell receptor beta, variable 8.2" Tcrb-V8.2
 1427629_at 1.17839 "protein tyrosine phosphatase, receptor type, J" Ptpnj
 1427630_x_at 0.93246 CEA-related cell adhesion molecule 1 Ceacam1
 1427631_x_at 0.91513 major urinary protein 3 Mup3
 1427632_x_at 1.04137 decay accelerating factor 2 Daf2
 1427633_a_at 1.02431 pregnancy-associated plasma protein A Pappa
 1427634_at 1.01431 "potassium voltage-gated channel, subfamily H (eag-related), member 7" Kcnh7
 1427635_at 1.01081 kinesin family member 5C Kif5c
 1427636_at 0.95418 "testis specific protein-Y encoded, pseudogene" Tspy-ps
 1427637_a_at 0.83789 desmocollin 3 Dsc3
 1427638_at 0.98946 zinc finger and BTB domain containing 16 Zbtb16
 1427639_a_at 1.06231 NIMA (never in mitosis gene a)-related expressed kinase 4 Nek4
 1427640_a_at 1.09091 "runt-related transcription factor 1; translocated to, 1 (cyclin D-related)"
 Runx1t1

1427641_at 1.02822 Distal-less homeobox 4 Dlx4
1427642_at 1.03539 "transient receptor potential cation channel, subfamily M, member 6" Trpm6
1427643_at 1.03310 RIKEN cDNA 1200009O22 gene 1200009O22Rik
1427644_at 1.05271 "DNA segment, KIST 4" D0Kist4
1427645_a_at 0.94445 interleukin 1 receptor accessory protein-like 2 Il1rapl2
1427646_a_at 0.91471 rho/rac guanine nucleotide exchange factor (GEF) 2 Arhgef2
1427647_at 0.97515 cDNA sequence BC022150 BC022150
1427648_at 0.83557 Regulator of G-protein signaling 3 Rgs3
1427649_at 1.07487 WD repeat domain 22 Wdr22
1427650_a_at 1.16802 runt related transcription factor 1 Runx1
1427651_x_at 2.08011 "histocompatibility 2, D region locus 1" H2-D1
1427652_x_at 0.97569 synaptojanin 2 Synj2
1427653_at 0.89376 "T-cell receptor alpha chain /// RIKEN cDNA A430107P09 gene /// gene model
1220, (NCBI) /// similar to T-cell receptor alpha chain V region CTL-F3 precursor /// similar to T-cell
receptor alpha chain V region CTL-F3 precursor /// T cell receptor alpha variable 6D-6" Tcra ///
A430107P09Rik /// Gm1220 /// LOC620297 /// LOC634046 /// Trav6d-6
1427654_a_at 1.08306 5 hydroxytryptamine (serotonin) receptor 4 Htr4
1427655_a_at 0.87001 RIKEN cDNA A630038E17 gene A630038E17Rik
1427656_at 0.98698 "T-cell receptor beta, variable 13" Tcrb-V13
1427657_at 0.99260 RIKEN cDNA 2310042L06 gene 2310042L06Rik
1427658_at 1.05188 "chitobiase, di-N-acetyl-" Ctbs
1427659_at 0.89934 "Mus musculus, clone IMAGE:4006890" ---
1427660_x_at 0.87026 "immunoglobulin kappa chain, constant region /// immunoglobulin kappa
chain variable 28 (V28) /// immunoglobulin kappa chain variable 8 (V8)-16" Igk-C /// Igk-V28 /// Igk-
V8-16
1427661_a_at 0.98174 tumor-suppressing subchromosomal transferable fragment 4 Tssc4
1427662_at 1.01120 Supervillin Svll
1427663_a_at 1.06961 CDC like kinase 4 Clk4
1427664_at 0.95358 RIKEN cDNA 4930453N24 gene 4930453N24Rik
1427665_a_at 0.88576 nuclear factor I/C Nfic
1427666_a_at 0.95897 "T-cell receptor beta, joining region" Tcrb-J
1427667_s_at 0.96616 "T-cell receptor beta, joining region" Tcrb-J
1427668_at 0.81131 "sialic acid binding Ig-like lectin 1, sialoadhesin" Siglec1
1427669_a_at 1.01317 citron Cit
1427670_a_at 0.98187 transcription factor 12 Tcf12
1427671_a_at 0.89758 formin 1 Fmn1
1427672_a_at 1.00925 "ubiquitously transcribed tetratricopeptide repeat gene, X chromosome" Utx
1427673_a_at 1.00155 "sema domain, immunoglobulin domain (Ig), short basic domain, secreted,
(semaphorin) 3E" Sema3e
1427674_a_at 0.95619 seizure related gene 6 Sez6
1427675_at 1.08266 "vomeronasal 1 receptor, A2" V1ra2
1427676_a_at 0.94318 "glutamate receptor, ionotropic, kainate 1" Grik1
1427677_a_at 0.83906 SRY-box containing gene 6 Sox6

1427678_at 0.98656 "zinc finger, imprinted 3" Zim3
 1427679_at 1.21211 large tumor suppressor Lats1
 1427680_a_at 1.02473 nuclear factor I/B Nfib
 1427681_s_at 0.86231 "vomeronasal 2, receptor, 13" V2r13
 1427682_a_at 0.69791 early growth response 2 Egr2
 1427683_at 0.38888 early growth response 2 Egr2
 1427684_at 1.11202 zinc finger protein 264 Zfp264
 1427685_a_at 1.26587 synaptojanin 2 Synj2
 1427686_at 0.97890 "vomeronasal 2, receptor, 13 /// similar to vomeronasal 2, receptor, 8" V2r13 ///
 LOC632660
 1427687_at 1.06918 protocadherin alpha 10 Pcdha10
 1427688_a_at 0.72675 "protein tyrosine phosphatase, receptor type, S" Ptpns
 1427689_a_at 1.18680 TNFAIP3 interacting protein 1 Tnip1
 1427690_a_at 0.87364 transmembrane phosphatase with tensin homology Tpte
 1427691_a_at 1.23862 interferon (alpha and beta) receptor 2 Ifnar2
 1427692_a_at 0.95680 calcium/calmodulin-dependent serine protein kinase (MAGUK family)
 Cask
 1427693_at 0.92558 "inositol 1,4,5-triphosphate receptor 2" Itpr2
 1427694_at 0.89483 gonadotropin releasing hormone receptor Gnhr
 1427695_a_at 1.07216 "POU domain, class 2, transcription factor 1" Pou2f1
 1427696_at 1.02927 ferrochelatase Fech
 1427697_a_at 0.85999 transformation related protein 73 Trp73
 1427698_at 1.07693 breast cancer 1 Brca1
 1427699_a_at 1.16063 "protein tyrosine phosphatase, non-receptor type 11" Ptpn11
 1427700_x_at 0.86590 keratin 6A Krt6a
 1427701_a_at 1.19544 sine oculis-related homeobox 4 homolog (Drosophila) Six4
 1427702_at 1.11551 zinc finger protein 1 Zfp1
 1427703_at 1.14308 "platelet-activating factor acetylhydrolase, isoform 1b, beta1 subunit" Pafah1b1
 1427704_a_at 1.01736 galanin receptor 1 Galr1
 1427705_a_at 1.04553 "nuclear factor of kappa light chain gene enhancer in B-cells 1, p105" Nfkb1
 1427706_a_at 1.05651 "sialic acid binding Ig-like lectin 1, sialoadhesin" Siglec1
 1427707_a_at 1.10034 --- ---
 1427708_a_at 0.97289 neurofibromatosis 2 Nf2
 1427709_at 1.08661 "glutamate receptor, ionotropic, kainate 3" Grik3
 1427710_at 1.07966 cDNA sequence BC013491 BC013491
 1427711_a_at 1.04969 CEA-related cell adhesion molecule 1 Ceacam1
 1427712_at 0.96371 CEA-related cell adhesion molecule 1 Ceacam1
 1427713_x_at 0.90500 "POU domain, class 2, transcription factor 2" Pou2f2
 1427714_at 0.89800 SET and MYND domain containing 1 Smyd1
 1427715_a_at 1.09962 "5'-nucleotidase, cytosolic IB" Nt5c1b
 1427716_at 1.00531 RIKEN cDNA 1700036D21 gene 1700036D21Rik
 1427717_at 1.11007 CD80 antigen Cd80
 1427718_a_at 1.09400 transformed mouse 3T3 cell double minute 2 Mdm2

1427719_s_at 0.96483 "keratin complex 2, basic, pseudogene 1 /// keratin 82" Krt2-ps1 /// Krt82
1427720_a_at 0.99822 novel nuclear protein 1 Nnp1
1427721_at 1.00953 --- ---
1427722_at 0.94546 RIKEN cDNA 9830102E05 gene 9830102E05Rik
1427723_at 0.87149 growth differentiation factor 11 Gdf11
1427724_at 1.08808 topoisomerase (DNA) II alpha Top2a
1427725_a_at 0.91020 "POU domain, class 2, transcription factor 2" Pou2f2
1427726_at 0.99553 "similar to histocompatibility 2, T region locus 3" LOC633417
1427727_x_at 0.96109 pregnancy specific glycoprotein 19 Psg19
1427728_at 0.95159 "cholinergic receptor, nicotinic, gamma polypeptide" Chrng
1427729_at 1.17533 EH-domain containing 2 Ehd2
1427730_a_at 0.85948 zinc finger protein 148 Zfp148
1427731_at 1.13179 gonadotropin releasing hormone receptor Gnhr
1427732_s_at 0.89432 "ATP-binding cassette, sub-family G (WHITE), member 4" Abcg4
1427733_a_at 0.96003 nuclear factor I/A Nfia
1427734_a_at 1.19876 Down syndrome cell adhesion molecule-like 1 Dscaml1
1427735_a_at 0.95128 "actin, alpha 1, skeletal muscle" Acta1
1427736_a_at 0.99970 chemokine (C-C motif) receptor-like 2 Ccr12
1427737_a_at 0.97307 RIKEN cDNA 1200011M11 gene 1200011M11Rik
1427738_at 1.14195 "DNA segment, KIST 2" D0Kist2
1427739_a_at 1.08750 transformation related protein 53 Trp53
1427740_a_at 1.10688 Strain C3H/HeN mRNA preferentially expressed in LPS-normoresponsive
macrophages ---
1427741_x_at 0.96749 "NK2 transcription factor related, locus 3 (Drosophila)" Nkx2-3
1427742_a_at 1.05889 Kruppel-like factor 6 Klf6
1427743_at 1.14061 "Gene model 672, (NCBI)" Gm672
1427744_at 0.97640 cyclin B3 Ccnb3
1427745_x_at 0.82340 proline rich protein MP4 /// similar to proline rich protein MP4 /// cDNA
sequence BC061494 /// proline-rich protein BstNI subfamily 1 /// similar to proline rich protein MP4 ///
similar to proline rich protein MP4 MP4 /// MGC78233 /// BC061494 /// Prb1 /// LOC667929 ///
LOC674323
1427746_x_at 1.28425 "histocompatibility 2, K1, K region" H2-K1
1427747_a_at 37.64685 lipocalin 2 Lcn2
1427748_at 1.19405 Immunoglobulin kappa chain variable 1 (V1) Igk-V1
1427749_at 1.15964 hypothetical protein LOC620893 LOC620893
1427750_at 1.03726 Immunoglobulin kappa chain variable 1 (V1) Igk-V1
1427751_a_at 0.91775 keratin 36 Krt36
1427752_a_at 0.99115 "T-cell receptor beta, variable 13 /// similar to T-cell receptor beta chain V
region PHDS203 precursor /// similar to T-cell receptor beta chain V region PHDS203 precursor ///
similar to T-cell receptor beta chain V region PHDS203 precursor /// similar to T-cell receptor beta
chain V region PHDS203 precursor" Tcrb-V13 /// LOC665446 /// LOC669809 /// LOC674045 ///
LOC676754
1427753_at 0.97326 Immunoglobulin light chain variable region Igv1-135

1427754_a_at 0.93075 dynamin 1 Dnm1
 1427755_at 1.03498 "mast cell protease, pseudogene 1" Mcpt-ps1
 1427756_x_at 0.97926 immunoglobulin heavy chain 4 (serum IgG1) Igh-4
 1427757_at 0.88129 immunoglobulin kappa chain variable 21 (V21) Igh-V21
 1427758_x_at 0.87486 immunoglobulin heavy chain 4 (serum IgG1) /// immunoglobulin heavy chain 6 (heavy chain of IgM) /// immunoglobulin heavy chain (V7183 family) /// immunoglobulin heavy chain (J558 family) /// immunoglobulin heavy chain (S107 family) /// immunoglobulin heavy chain 1a (serum IgG2a) /// Immunoglobulin heavy chain (gamma polypeptide) /// expressed sequence AI324046 /// similar to immunoglobulin mu-chain /// similar to Ig heavy chain V region M315 precursor /// similar to Ig heavy chain V region M315 precursor Igh-4 /// Igh-6 /// Igh-V7183 /// Igh-VJ558 /// Igh-VS107 /// Igh-1a /// Ighg /// AI324046 /// LOC544903 /// LOC633674 /// LOC641025 /// LOC676462
 1427759_a_at 0.99248 ADP-ribosyltransferase 5 Art5
 1427760_s_at 1.15699 "proliferin /// proliferin 2 /// mitogen regulated protein, proliferin 3" Plf /// Plf2 /// Mrpplf3
 1427761_at 0.97210 prion protein interacting protein 1 Prnpi1
 1427762_x_at 1.04034 "histone 1, H2bp" Hist1h2bp
 1427763_a_at 1.16165 "calcium/calmodulin-dependent protein kinase II, delta" Camk2d
 1427764_a_at 0.94560 transcription factor E2a Tcf2a
 1427765_a_at 0.95263 "T-cell receptor beta, variable 13 /// similar to T-cell receptor beta chain V region CTL-L17 precursor /// similar to T-cell receptor beta chain V region CTL-L17 precursor" Tcrb-V13 /// LOC384373 /// LOC669782
 1427766_at 1.07715 "T-cell receptor alpha, variable 8" Tcra-V8
 1427767_a_at 0.93832 cystic fibrosis transmembrane conductance regulator homolog Cftr
 1427768_s_at 0.94783 "myosin, light polypeptide 3" Myl3
 1427769_x_at 0.89198 "myosin, light polypeptide 3" Myl3
 1427770_a_at 0.99658 "solute carrier family 2 (facilitated glucose transporter), member 3" Slc2a3
 1427771_x_at 1.05773 integrin beta 1 (fibronectin receptor beta) Itgb1
 1427772_at 0.97886 defensin beta 15 Defb15
 1427773_a_at 0.93163 Rab acceptor 1 (prenylated) Rabac1
 1427774_at 1.00547 defensin beta 7 Defb7
 1427775_at 0.96185 defensin beta 10 Defb10
 1427776_a_at 0.97807 fibroblast growth factor receptor 4 Fgfr4
 1427777_x_at 1.00142 fibroblast growth factor receptor 4 Fgfr4
 1427778_at 0.89932 defensin beta 8 Defb8
 1427779_a_at 1.07621 CD4 antigen Cd4
 1427780_at 0.97821 defensin beta 35 Defb35
 1427781_at 1.00852 upregulated during skeletal muscle growth 2 Usmg2
 1427782_a_at 0.93678 corticotropin releasing hormone receptor 1 Crhr1
 1427783_at 0.94669 v-erb-a erythroblastic leukemia viral oncogene homolog 4 (avian) Erbb4
 1427784_at 0.90364 defensin beta 13 Defb13
 1427785_x_at 0.98118 Small optic lobes homolog (Drosophila) Solh
 1427786_at 0.89289 trans-acting transcription factor 6 Sp6

1427787_at 0.99410 trans-acting transcription factor 6 Sp6
 1427788_at 1.14019 neuroendocrine secretory protein antisense Nespas
 1427789_s_at 0.95469 "GNAS (guanine nucleotide binding protein, alpha stimulating) complex locus" Gnas
 1427790_at 0.83081 MAP kinase-activated protein kinase 5 /// a disintegrin and metallopeptidase domain 1a Mapkapk5 /// Adam1a
 1427791_a_at 1.00197 a disintegrin and metallopeptidase domain 1a Adam1a
 1427792_at 0.90935 Similar to immunoglobulin light chain variable region LOC384413
 1427793_at 0.89973 "hypothetical protein, P4(21)n gene" AJ242955
 1427794_at 0.87934 "hypothetical protein, P4(21)n gene" AJ242955
 1427795_s_at 1.02491 "Mus musculus, clone IMAGE:5068294" ---
 1427796_at 1.11332 zinc finger protein 133 Zfp133
 1427797_s_at 0.99898 cathepsin E /// similar to melanoma antigen /// similar to melanoma antigen /// similar to melanoma antigen /// similar to melanoma antigen /// similar to melanoma antigen /// similar to melanoma antigen /// similar to melanoma antigen /// similar to melanoma antigen /// similar to melanoma antigen /// similar to melanoma antigen /// similar to melanoma antigen /// similar to melanoma antigen /// similar to melanoma antigen /// similar to melanoma antigen /// similar to melanoma antigen /// hypothetical protein LOC670510 /// similar to melanoma antigen Ctse /// LOC622147 /// LOC665955 /// LOC666402 /// LOC667328 /// LOC667538 /// LOC668269 /// LOC668639 /// LOC668769 /// LOC668842 /// LOC669176 /// LOC669407 /// LOC669658 /// LOC669821 /// LOC670510 /// LOC671134
 1427798_x_at 1.04496 "Envelope protein, mRNA sequence" ---
 1427799_x_at 0.99404 immunoglobulin kappa chain variable 28 (V28) /// immunoglobulin kappa chain variable 8 (V8) /// immunoglobulin kappa chain /// immunoglobulin kappa chain variable 19 (V19)-14 /// immunoglobulin kappa chain variable 21 (V21) Igc-V28 /// Igc-V8 /// LOC545854 /// Igc-V19-14 /// Igc-V21
 1427800_at 0.94325 keratin associated protein 16-2 Krtap16-2
 1427801_at 0.98516 keratin associated protein 16-2 Krtap16-2
 1427802_a_at 1.00669 T-cell receptor gamma chain Tcrg
 1427803_at 1.00524 --- ---
 1427804_at 0.90527 --- ---
 1427805_at 0.98987 --- ---
 1427806_at 1.07590 --- ---
 1427807_at 0.87797 hypothetical protein LOC676582 LOC676582
 1427808_at 0.83351 nuclear transcription factor-Y alpha Nfya
 1427809_at 0.98172 Latrophilin 3 Lphn3
 1427810_at 1.00662 thymidylate synthase Tyms
 1427811_at 0.98002 thymidylate synthase Tyms
 1427812_at 0.95547 iduronate 2-sulfatase Ids
 1427813_at 0.98753 iduronate 2-sulfatase Ids
 1427814_at 1.02336 oncofetal antigen Ofa
 1427815_at 1.01922 --- ---
 1427816_at 0.96078 "splicing factor, arginine/serine-rich 2 (SC-35)" Sfrs2
 1427817_at 1.08949 hypothetical protein LOC641050 LOC641050

1427818_at	1.21536	B-cell leukemia/lymphoma 2	Bcl2
1427819_at	0.93698	---	---
1427820_at	1.03561	---	---
1427821_at	1.05296	"coatomer protein complex, subunit gamma 2, antisense 2"	Copg2as2
1427822_a_at	0.90453	"coatomer protein complex, subunit gamma 2, antisense 2"	Copg2as2
1427823_at	0.95191	---	---
1427824_at	1.01948	---	---
1427825_at	0.94083	---	---
1427826_a_at	1.10557	"solute carrier organic anion transporter family, member 1b2"	Slco1b2
1427827_at	0.95093	Murine retrovirus readthrough RNA sequence	---
1427828_at	0.83177	Murine retrovirus readthrough RNA sequence	---
1427829_at	0.92503	"ATP-binding cassette, sub-family D (ALD), member 4"	Abcd4
1427830_at	0.80431	zinc finger protein 260	Zfp260
1427831_s_at	0.78219	zinc finger protein 260 /// similar to Zinc finger protein OZF (Zinc finger protein 260) (Zfp-260)	Zfp260 /// LOC635007
1427832_at	0.93809	"tubulin alpha, related sequence 1"	Tuba-rs1
1427833_at	0.96744	serine protease inhibitor 16	Spi16
1427834_at	0.90691	serine protease inhibitor 16	Spi16
1427835_at	0.96768	"POU domain, class 2, transcription factor 1"	Pou2f1
1427836_at	0.93542	stretch responsive protein 278	Sr278
1427837_at	0.98934	Immunoglobulin kappa chain variable 32 (V32)	Igk-V32
1427838_at	0.93681	"tubulin, beta 2a"	Tubb2a
1427839_at	0.96432	Immunoglobulin heavy chain (J558 family)	Igh-VJ558
1427840_at	0.99831	---	---
1427841_at	0.95389	keratin 14 /// keratin 17	Krt14 /// Krt17
1427842_at	0.88335	keratin associated protein 6-3	Krtap6-3
1427843_at	0.95799	expressed sequence AI840826	AI840826
1427844_a_at	0.97233	"CCAAT/enhancer binding protein (C/EBP), beta"	Cebpb
1427845_at	0.95790	fibroblast growth factor receptor 4	Fgfr4
1427846_x_at	0.96858	fibroblast growth factor receptor 4	Fgfr4
1427847_at	0.94258	runt related transcription factor 1	Runx1
1427848_at	0.99672	"T-cell receptor beta, variable 13"	Tcrb-V13
1427849_a_at	0.87706	"T-cell receptor beta, variable 13"	Tcrb-V13
1427850_x_at	0.96421	immunoglobulin heavy chain (J558 family)	Igh-VJ558
1427851_x_at	0.94438	immunoglobulin heavy chain (J558 family)	Igh-VJ558
1427852_x_at	0.84321	immunoglobulin heavy chain (J558 family)	Igh-VJ558
1427853_a_at	0.82062	heat shock protein 1	Hspb1
1427854_x_at	0.72501	heat shock protein 1	Hspb1
1427855_at	0.83636	---	---
1427856_a_at	0.93291	Immunoglobulin kappa chain complex	---
1427857_x_at	1.10637	"Ig kappa chain V-region mRNA, partial cds, clone 2F2"	---
1427858_at	1.04620	Immunoglobulin kappa chain variable 1 (V1)	Igk-V1
1427859_at	1.00059	Immunoglobulin kappa chain complex	---

1427860_at 1.03805 immunoglobulin kappa chain variable 6-23 /// immunoglobulin kappa chain
 variable 19 (V19)-14 /// immunoglobulin kappa chain variable 19 (V19)-13 /// similar to Ig kappa chain
 V-V region MPC11 precursor /// similar to Ig kappa chain V-V region MPC11 precursor /// similar to Ig
 kappa chain V-V region MPC11 precursor /// similar to Ig kappa chain V-V region MPC11 precursor ///
 similar to Ig kappa chain V-V region MPC11 precursor Igkv6-23 /// Igk-V19-14 /// Igk-V19-13 ///
 LOC669053 /// LOC669070 /// LOC676136 /// LOC676162 /// LOC676175
 1427861_at 1.06725 ADP-ribosylation factor-like 6 interacting protein 3 Arl6ip3
 1427862_at 1.08001 ADP-ribosylation factor-like 6 interacting protein 3 Arl6ip3
 1427863_at 0.98447 "histone1, H3d" Hist1h3d
 1427864_at 1.09334 "histone 2, H3c1 /// histone1, H3d /// histone 1, H3b /// histone 1, H3e /// histone
 2, H3b" Hist2h3c1 /// Hist1h3d /// Hist1h3b /// Hist1h3e /// Hist2h3b
 1427865_at 1.00264 --- ---
 1427866_x_at 0.89434 "hemoglobin, beta adult major chain" Hbb-b1
 1427867_at 0.96078 "Myosin, heavy polypeptide 1, skeletal muscle, adult" Myh1
 1427868_x_at 0.99871 "myosin, heavy polypeptide 1, skeletal muscle, adult" Myh1
 1427869_at 0.96502 immunoglobulin heavy chain 4 (serum IgG1) Igh-4
 1427870_x_at 0.99742 immunoglobulin heavy chain 4 (serum IgG1) Igh-4
 1427871_at 0.89330 platelet-activating factor receptor Ptafr
 1427872_at 0.88983 platelet-activating factor receptor Ptafr
 1427873_at 0.94728 defensin related cryptdin 15 Defcr15
 1427874_at 0.89039 zinc finger protein 313 Zfp313
 1427875_a_at 1.07898 RIKEN cDNA 1100001I22 gene 1100001I22Rik
 1427876_at 1.07583 RIKEN cDNA 2610312B22 gene 2610312B22Rik
 1427877_at 0.97671 RIKEN cDNA 2610312B22 gene 2610312B22Rik
 1427878_at 0.99388 RIKEN cDNA 0610010O12 gene 0610010O12Rik
 1427879_at 0.93774 RIKEN cDNA 1810031K17 gene 1810031K17Rik
 1427880_at 1.02851 "ubiquinol-cytochrome c reductase, complex III subunit VII" Uqcrq
 1427881_at 1.06217 "deoxynucleotidyltransferase, terminal, interacting protein 2" Dnttip2
 1427882_at 1.07078 "deoxynucleotidyltransferase, terminal, interacting protein 2" Dnttip2
 1427883_a_at 0.84851 "procollagen, type III, alpha 1" Col3a1
 1427884_at 1.06664 "procollagen, type III, alpha 1" Col3a1
 1427885_at 0.93994 "polymerase (DNA-directed), delta 4" Pold4
 1427886_at 0.96162 nuclear pore membrane protein 121 Pom121
 1427887_at 1.03988 RIKEN cDNA 2610304G08 gene 2610304G08Rik
 1427888_a_at 0.92627 "spectrin alpha 2 /// similar to Spectrin alpha chain, brain (Spectrin, non-
 erythroid alpha chain) (Alpha-II spectrin) (Fodrin alpha chain)" Spna2 /// LOC381358
 1427889_at 0.93659 "spectrin alpha 2 /// similar to Spectrin alpha chain, brain (Spectrin, non-
 erythroid alpha chain) (Alpha-II spectrin) (Fodrin alpha chain)" Spna2 /// LOC381358
 1427890_a_at 0.89987 leucine rich repeat containing 56 Lrrc56
 1427891_at 0.98616 "GTPase, IMAP family member 6" Gimap6
 1427892_at 1.13768 myosin IG Myo1g
 1427893_a_at 1.41999 phosphomevalonate kinase Pmvk
 1427894_at 0.79656 vasorin Vasn

1427895_at	0.96084	RIKEN cDNA 2310004N24	gene	2310004N24Rik
1427896_at	0.90018	RIKEN cDNA 2400003N08	gene	2400003N08Rik
1427897_s_at	0.84126	RIKEN cDNA 2400003N08	gene	2400003N08Rik
1427898_at	1.05450	ring finger protein (C3H2C3 type) 6		Rnf6
1427899_at	0.97426	ring finger protein (C3H2C3 type) 6		Rnf6
1427900_at	1.15251	phosphatidylinositol-4-phosphate 5-kinase-like 1		Pip5k1l
1427901_at	1.09679	mitochondrial ribosomal protein S18C		Mrps18c
1427902_at	1.32502	serine/arginine repetitive matrix 2		Srrm2
1427903_at	1.07067	phosphohistidine phosphatase 1		Phpt1
1427904_s_at	1.01787	RIKEN cDNA 2410091C18	gene	2410091C18Rik
1427905_at	1.11495	RIKEN cDNA 1810063B07	gene	1810063B07Rik
1427906_at	1.01277	RIKEN cDNA 1110037F02	gene	1110037F02Rik
1427907_at	1.03626	RIKEN cDNA 1110037F02	gene	1110037F02Rik
1427908_at	1.12341	"BCL2/adenovirus E1B interacting protein 1, NIP1"		Bnip1
1427909_at	1.02959	RIKEN cDNA 2410015M20	gene	2410015M20Rik
1427910_at	1.07365	cystatin E/M		Cst6
1427911_at	1.04865	RIKEN cDNA 2610307O08	gene	2610307O08Rik
1427912_at	1.13759	carbonyl reductase 3		Cbr3
1427913_at	1.04063	RWD domain containing 1		Rwdd1
1427914_a_at	1.04378	"transcription elongation factor B (SIII), polypeptide 1"		Tceb1
1427915_s_at	0.91656	"transcription elongation factor B (SIII), polypeptide 1"		Tceb1
1427916_at	0.78519	suppression of tumorigenicity 7-like		St7l
1427917_s_at	0.89785	single-stranded DNA binding protein 3		Ssbp3
1427918_a_at	0.94442	"ras homolog gene family, member Q"		Rhoq
1427919_at	1.01646	"sushi-repeat-containing protein, X-linked 2"		SrpX2
1427920_at	1.00924	PHD finger protein 19		Phf19
1427921_s_at	0.95488	RIKEN cDNA 2310061C15	gene	2310061C15Rik
1427922_at	0.88298	RIKEN cDNA 2310061C15	gene	2310061C15Rik
1427923_at	0.91989	"zinc metalloproteinase, STE24 homolog (S. cerevisiae)"		Zmpste24
1427924_at	0.94876	syntaxin 17		Stx17
1427925_at	1.11764	syntaxin 17		Stx17
1427926_at	1.09803	mastermind like 2 (Drosophila) /// similar to mastermind-like 2; mastermind (Drosophila)-like 2		Maml2 /// LOC434369
1427927_at	0.97750	expressed sequence AW049829		AW049829
1427928_s_at	0.90893	CDC91 cell division cycle 91-like 1 (S. cerevisiae)		Cdc91l1
1427929_a_at	0.90409	"pyridoxal (pyridoxine, vitamin B6) kinase"		Pdxk
1427930_at	1.05920	"pyridoxal (pyridoxine, vitamin B6) kinase"		Pdxk
1427931_s_at	1.09876	"pyridoxal (pyridoxine, vitamin B6) kinase"		Pdxk
1427932_s_at	1.34453	RIKEN cDNA 1200003I10 gene /// RIKEN cDNA 1200015M12 gene /// RIKEN cDNA 1200016E24 gene /// RIKEN cDNA A130040M12 gene /// RIKEN cDNA E430024C06 gene 1200003I10Rik /// 1200015M12Rik /// 1200016E24Rik /// A130040M12Rik /// E430024C06Rik		
1427933_at	1.02004	vacuolar protein sorting 33B (yeast)		Vps33b
1427934_at	0.97370	RIKEN cDNA 2610208E05	gene	2610208E05Rik

1427935_at 1.03944 RIKEN cDNA 2610208E05 gene 2610208E05Rik
1427936_at 0.91253 threonine synthase-like 1 (bacterial) Thns11
1427937_at 1.01553 RIKEN cDNA 2610030H06 gene 2610030H06Rik
1427938_at 0.83273 c-myc binding protein Mycbp
1427939_s_at 0.84245 c-myc binding protein Mycbp
1427940_s_at 0.89069 c-myc binding protein Mycbp
1427941_at 1.08184 "Dicer1, Dcr-1 homolog (Drosophila)" Dicer1
1427942_at 0.78703 peptidylglycine alpha-amidating monooxygenase COOH-terminal interactor
Pamci
1427943_at 0.98724 "acylphosphatase 2, muscle type" Acyp2
1427944_at 0.90906 C1q domain containing 1 C1qdc1
1427945_at 1.01151 dihydropyrimidine dehydrogenase Dpyd
1427946_s_at 0.99757 dihydropyrimidine dehydrogenase Dpyd
1427947_at 1.09559 cDNA sequence BC028440 BC028440
1427948_a_at 1.00895 zinc finger protein 99 Zfp99
1427949_at 1.04790 zinc finger protein 294 Zfp294
1427950_at 1.02044 zinc finger protein 294 Zfp294
1427951_s_at 0.83722 coiled-coil domain containing 28A Ccdc28a
1427952_at 0.94590 aryl hydrocarbon receptor-interacting protein-like 1 Aipl1
1427953_at 0.95229 cDNA sequence BC025462 BC025462
1427954_at 1.04483 cDNA sequence BC048403 BC048403
1427955_a_at 0.85929 differentially expressed in B16F10 1 Deb1
1427956_at 1.11259 polycomb group ring finger 1 Pcgf1
1427957_at 1.03613 RIKEN cDNA 9530008L14 gene 9530008L14Rik
1427958_at 0.83300 abhydrolase domain containing 10 Abhd10
1427959_at 0.99326 abhydrolase domain containing 10 Abhd10
1427960_at 0.72341 "UDP glucuronosyltransferase 2 family, polypeptide B34" Ugt2b34
1427961_s_at 0.68199 "UDP glucuronosyltransferase 2 family, polypeptide B34" Ugt2b34
1427962_at 0.89988 coiled-coil domain containing 102A /// similar to CG31638-PA Ccdc102a ///
LOC672218
1427963_s_at 1.02495 retinol dehydrogenase 9 Rdh9
1427964_at 0.79395 CKLF-like MARVEL transmembrane domain containing 8 Cmtm8
1427965_at 1.06773 single-stranded DNA binding protein 1 Ssbp1
1427966_at 0.92110 cDNA sequence BC087945 BC087945
1427967_at 0.88526 SLIT-ROBO Rho GTPase activating protein 2 Srgap2
1427968_at 0.92524 TGIF homeobox 1 Tgifx1
1427969_s_at 1.08518 zinc finger protein 654 Zfp654
1427970_at 0.94959 zinc finger protein 689 Zfp689
1427971_at 0.99325 "Vcell division cycle 73, Paf1/RNA polymerase II complex component,
homolog (S. cerevisiae)" Cdc73
1427972_at 0.90869 "Vcell division cycle 73, Paf1/RNA polymerase II complex component,
homolog (S. cerevisiae)" Cdc73
1427973_s_at 1.01914 "Vcell division cycle 73, Paf1/RNA polymerase II complex component,

homolog (*S. cerevisiae*)" Cdc73

1427974_s_at 0.97406 "calcium channel, voltage-dependent, L type, alpha 1D subunit" Cacna1d

1427975_at 0.73926 RIKEN cDNA 2210403B10 gene 2210403B10Rik

1427976_at 0.89686 oogenesisin 1 Oog1

1427977_x_at 1.07387 oogenesisin 1 Oog1

1427978_at 1.08563 RIKEN cDNA 4732418C07 gene 4732418C07Rik

1427979_at 0.99902 RIKEN cDNA 4732418C07 gene 4732418C07Rik

1427980_at 0.90311 RIKEN cDNA 4933407C03 gene 4933407C03Rik

1427981_a_at 0.86712 cysteine sulfinic acid decarboxylase Csad

1427982_s_at 1.01907 synaptic nuclear envelope 2 Syne2

1427983_at 1.33185 suppressor of hairy wing homolog 3 (*Drosophila*) Suhw3

1427984_at 1.02356 SUMO/sentrin specific peptidase 6 Senp6

1427985_at 0.83650 RIKEN cDNA 9630042H07 gene 9630042H07Rik

1427986_a_at 0.88029 "procollagen, type XVI, alpha 1" Col16a1

1427987_at 1.15235 scaffold attachment factor B2 Safb2

1427988_s_at 1.13170 scaffold attachment factor B2 Safb2

1427989_at 0.93064 anti-Mullerian hormone type 2 receptor Amhr2

1427990_at 0.98901 ubiquitin specific peptidase 45 Usp45

1427991_s_at 0.92007 ubiquitin specific peptidase 45 Usp45

1427992_a_at 1.00957 "RAB12, member RAS oncogene family" Rab12

1427993_at 0.84520 RUN and FYVE domain-containing 2 Rufy2

1427994_at 0.83236 CD300 antigen like family member F Cd300lf

1427995_at 0.98577 carboxypeptidase A5 Cpa5

1427996_at 0.97224 cDNA sequence BC028528 BC028528

1427997_at 1.09162 RIKEN cDNA 1110007M04 gene 1110007M04Rik

1427998_at 0.96775 RIKEN cDNA 2600001B17 gene 2600001B17Rik

1427999_at 1.05007 "small nuclear RNA activating complex, polypeptide 5" Snapc5

1428000_at 0.99653 transmembrane protein 60 Tmem60

1428001_at 1.05228 thyroid stimulating hormone receptor Tshr

1428002_at 0.97787 serum amyloid A-like 1 Saal1

1428003_s_at 0.93372 "thiamine triphosphatase /// zinc finger homeobox 2, antisense" Thtpa /// Zfhx2as

1428004_at 1.07655 RIKEN cDNA 3300001G02 gene 3300001G02Rik

1428005_at 0.86220 MOCO sulphurase C-terminal domain containing 1 Moscl1

1428006_at 0.94854 sec1 family domain containing 1 Scfd1

1428007_at 0.98643 keratin associated protein 13-1 Krtap13-1

1428008_at 1.03092 dynamin 1-like Dnm1l

1428009_a_at 1.01892 "ATP-binding cassette, sub-family C (CFTR/MRP), member 10" Abcc10

1428010_at 1.08178 translocase of inner mitochondrial membrane 9 homolog (yeast) Timm9

1428011_a_at 0.91292 Erbb2 interacting protein Erbb2ip

1428012_at 0.98629 "complement component 8, alpha polypeptide" C8a

1428013_at 1.28507 RIKEN cDNA 6030458C11 gene 6030458C11Rik

1428014_at 0.95902 "histone 1, H4h" Hist1h4h

1428015_at 1.18195 "DNA segment, Chr 12, Wayne State University 118, expressed" D12Wsu118e
1428016_a_at 0.96706 Ras interacting protein 1 Rasip1
1428017_at 1.08330 Pbx/knotted 1 homeobox 2 Pknox2
1428018_a_at 0.96091 Cd300D antigen Cd300d
1428019_at 0.89848 RIKEN cDNA 4932425I24 gene 4932425I24Rik
1428020_at 0.88576 brix domain containing 5 Bxdc5
1428021_at 0.98973 methylcrotonoyl-Coenzyme A carboxylase 2 (beta) Mccc2
1428022_at 0.97635 lipocalin 13 Lcn13
1428023_at 1.03727 RIKEN cDNA 3110009E18 gene 3110009E18Rik
1428024_at 1.08045 Microtubule-associated protein 1 light chain 3 beta Map1lc3b
1428025_s_at 1.02897 "phosphatidylinositol transfer protein, cytoplasmic 1" Pitpnc1
1428026_at 0.87335 teashirt zinc finger family member 2 Tshz2
1428027_at 1.18187 "BIC noncoding mRNA, complete sequence" ---
1428028_at 0.99836 gene trap insertion site 4-1 Gt4-1
1428029_a_at 0.96101 "H2A histone family, member V" H2afv
1428030_at 0.98752 a disintegrin and metallopeptidase domain 34 Adam34
1428031_at 0.97660 melanin-concentrating hormone receptor 1 Mchr1
1428032_at 0.95654 "activin A receptor, type IC" Acvr1c
1428033_at 0.89826 hypothetical protein LOC245174 LOC245174
1428034_a_at 1.53846 "tumor necrosis factor receptor superfamily, member 9" Tnfrsf9
1428035_at 1.04789 fibrous sheath-interacting protein 1 Fsip1
1428036_at 0.93192 "MRNA similar to phosphatidylinositol glycan, class S (cDNA clone MGC:41618 IMAGE:1265093)" ---
1428037_at 0.98026 otoancorin Otoa
1428038_at 1.08288 "gene model 568, (NCBI)" Gm568
1428039_at 0.95988 RIKEN cDNA 2310040A13 gene 2310040A13Rik
1428040_at 0.93971 "one cut domain, family member 3" Onecut3
1428041_at 0.99788 RIKEN cDNA 1600002H07 gene 1600002H07Rik
1428042_at 0.99908 Similar to RIKEN cDNA B430319H21 gene MGC41410
1428043_a_at 1.04759 cytidine monophospho-N-acetylneuraminic acid hydroxylase Cmah
1428044_at 1.09046 "adaptor-related protein complex 3, sigma 2 subunit" Ap3s2
1428045_a_at 0.87426 E74-like factor 2 Elf2
1428046_a_at 0.80730 zinc finger protein X-linked Zfx
1428047_s_at 0.87270 "zinc finger protein, autosomal /// zinc finger protein X-linked" Zfa /// Zfx
1428048_at 0.99979 cDNA sequence BC027582 BC027582
1428049_a_at 0.98761 nudix (nucleoside diphosphate linked moiety X)-type motif 16-like 1 Nudt16l1
1428050_a_at 0.96057 transmembrane BAX inhibitor motif containing 4 Tmbim4
1428051_a_at 1.11206 "calcium channel, voltage-dependent, L type, alpha 1D subunit" Cacna1d
1428052_a_at 1.00254 "zinc finger, MYM domain containing 1" Zmym1
1428053_at 1.04085 "G protein-coupled receptor, family C, group 6, member A" Gprc6a
1428054_at 0.95562 "solute carrier family 8 (sodium/calcium exchanger), member 2" Slc8a2
1428055_at 0.93869 RNA imprinted and accumulated in nucleus Rian

1428056_at	0.97310	"Killer cell lectin-like receptor, subfamily A, member 4"	Klra4
1428057_a_at	0.91702	AHNAK nucleoprotein (desmoyokin)	Ahnak
1428058_at	0.92311	AHNAK nucleoprotein (desmoyokin)	Ahnak
1428059_at	1.00731	CD247 antigen	Cd247
1428060_at	0.89296	CD247 antigen	Cd247
1428061_at	1.06475	histone aminotransferase 1	Hat1
1428062_at	1.12624	carboxypeptidase A1	Cpa1
1428063_at	0.88123	ankyrin repeat domain 46	Ankrd46
1428064_at	1.04319	"centaurin, delta 2"	Centd2
1428065_at	0.86973	"solute carrier family 44, member 2"	Slc44a2
1428066_at	1.07686	coiled-coil domain containing 120	Ccdc120
1428067_at	1.02403	"RAS-like, family 12"	Ras12
1428068_at	1.02427	sorting and assembly machinery component 50 homolog (S. cerevisiae)	Samm50
1428069_at	1.02622	cell division cycle associated 7	Cdca7
1428070_at	0.92431	"synovial apoptosis inhibitor 1, synoviolin"	Syvn1
1428071_at	0.94809	RIKEN cDNA 1110038D17 gene	1110038D17Rik
1428072_a_at	0.99714	dolichyl pyrophosphate phosphatase 1	Dolpp1
1428073_a_at	1.01708	nucleoporin 88	Nup88
1428074_at	1.23537	transmembrane protein 158	Tmem158
1428075_at	0.92319	NADH dehydrogenase (ubiquinone) 1 beta subcomplex 4	Ndufb4
1428076_s_at	0.99370	NADH dehydrogenase (ubiquinone) 1 beta subcomplex 4	Ndufb4
1428077_at	0.87357	transmembrane protein 163	Tmem163
1428078_at	0.85457	RIKEN cDNA 0610013E23 gene	0610013E23Rik
1428079_at	0.99049	"fibrinogen, B beta polypeptide"	Fgb
1428080_at	0.95838	phosphoglycerate mutase family member 5	Pgam5
1428081_at	1.06218	kelch-like 21 (Drosophila)	Klhl21
1428082_at	1.02764	acyl-CoA synthetase long-chain family member 5	Acsl5
1428083_at	1.89701	RIKEN cDNA 2310043N10 gene	2310043N10Rik
1428084_at	0.99243	"KRR1, small subunit (SSU) processome component, homolog (yeast)"	Krr1
1428085_at	1.03359	RIKEN cDNA 1110057K04 gene	1110057K04Rik
1428086_at	0.97779	dynamamin 1-like	Dnm11
1428087_at	0.95044	dynamamin 1-like	Dnm11
1428088_at	0.89616	RIKEN cDNA 2410002I01 gene	2410002I01Rik
1428089_at	1.01416	"SLIT and NTRK-like family, member 1"	Slitrk1
1428090_at	1.03361	Pentatricopeptide repeat domain 3	Ptcd3
1428091_at	0.93459	kelch-like 7 (Drosophila)	Klhl7
1428092_at	1.01656	cell division cycle 5-like (S. pombe)	Cdc51
1428093_at	0.91225	TraB domain containing	Trabd
1428094_at	0.86681	lysosomal membrane glycoprotein 2	Lamp2
1428095_a_at	0.96406	transmembrane protein 24	Tmem24
1428096_at	1.10436	importin 11	Ipo11
1428097_at	1.16865	RIKEN cDNA 2510009E07 gene	2510009E07Rik

1428098_a_at 0.91338 transmembrane 7 superfamily member 3 Tm7sf3
1428099_a_at 1.00616 "splicing factor, arginine/serine-rich 1 (ASF/SF2)" Sfrs1
1428100_at 1.07007 "splicing factor, arginine/serine-rich 1 (ASF/SF2)" Sfrs1
1428101_at 0.91172 ring finger protein 38 Rnf38
1428102_at 0.94624 carboxypeptidase B1 (tissue) Cpb1
1428103_at 0.82392 --- ---
1428104_at 1.07201 "TPX2, microtubule-associated protein homolog (Xenopus laevis)" Tpx2
1428105_at 0.97392 "TPX2, microtubule-associated protein homolog (Xenopus laevis)" Tpx2
1428106_at 1.03604 RIKEN cDNA 1300001I01 gene 1300001I01Rik
1428107_at 0.84738 SH3-binding domain glutamic acid-rich protein like Sh3bgrl
1428108_x_at 1.03449 transmembrane and coiled-coil domains 2 Tmcc2
1428109_at 1.03429 vacuolar protein sorting 11 (yeast) Vps11
1428110_x_at 1.09748 vacuolar protein sorting 11 (yeast) Vps11
1428111_at 0.91378 "solute carrier family 38, member 4" Slc38a4
1428112_at 1.02728 "arginine-rich, mutated in early stage tumors" Armet
1428113_at 0.82000 transmembrane and tetratricopeptide repeat containing 4 Tmtc4
1428114_at 0.97225 "solute carrier family 14 (urea transporter), member 1" Slc14a1
1428115_a_at 0.85476 "RAB2B, member RAS oncogene family" Rab2b
1428116_a_at 0.92016 dynein light chain Tctex-type 1 Dynlt1
1428117_x_at 0.79196 G protein-coupled receptor 178 Gpr178
1428118_at 0.93059 leucine rich repeat neuronal 6A Lrrn6a
1428119_a_at 1.03201 "RAP2C, member of RAS oncogene family" Rap2c
1428120_at 0.95198 F-box and WD-40 domain protein 9 Fbxw9
1428121_at 0.96244 RIKEN cDNA 2610528K11 gene 2610528K11Rik
1428122_s_at 0.78206 RIKEN cDNA 2610528K11 gene 2610528K11Rik
1428123_at 0.93152 RIKEN cDNA 2610528K11 gene 2610528K11Rik
1428124_at 0.97431 "general transcription factor II E, polypeptide 1 (alpha subunit)" Gtf2e1
1428125_at 0.82519 RIKEN cDNA 4921506J03 gene 4921506J03Rik
1428126_a_at 0.94431 RIKEN cDNA 4921506J03 gene 4921506J03Rik
1428127_at 0.89383 RIKEN cDNA 4921506J03 gene 4921506J03Rik
1428128_at 0.91951 RIKEN cDNA 4921506J03 gene 4921506J03Rik
1428129_at 0.87781 "lectin, mannose-binding, 1" Lman1
1428130_at 0.99462 "lectin, mannose-binding, 1" Lman1
1428131_a_at 0.92024 CDC42 small effector 1 Cdc42se1
1428132_at 0.87832 CDC42 small effector 1 Cdc42se1
1428133_at 0.99810 Smad nuclear interacting protein 1 Snip1
1428134_at 0.79585 coenzyme Q9 homolog (yeast) Coq9
1428135_a_at 1.01282 eukaryotic translation elongation factor 1 delta (guanine nucleotide exchange protein) Eef1d
1428136_at 0.96769 secreted frizzled-related sequence protein 1 Sfrp1
1428137_at 0.98893 ADP-ribosylation factor-like 8B Arl8b
1428138_s_at 0.98208 ADP-ribosylation factor-like 8B Arl8b
1428139_at 1.06669 RIKEN cDNA 4930538D17 gene 4930538D17Rik

1428140_at 0.93445 3-oxoacid CoA transferase 1 Oxct1
1428141_at 0.98709 "golgi associated, gamma adaptin ear containing, ARF binding protein 2" Gga2
1428142_at 1.35082 ets variant gene 5 Etv5
1428143_a_at 0.93066 patatin-like phospholipase domain containing 2 Pnpla2
1428144_at 0.94206 RIKEN cDNA 1200011O22 gene 1200011O22Rik
1428145_at 0.90114 acetyl-Coenzyme A acyltransferase 2 (mitochondrial 3-oxoacyl-Coenzyme A thiolase) Acaa2
1428146_s_at 0.94981 acetyl-Coenzyme A acyltransferase 2 (mitochondrial 3-oxoacyl-Coenzyme A thiolase) Acaa2
1428147_at 1.03080 coronin 7 Coro7
1428148_s_at 0.95984 coronin 7 Coro7
1428149_at 0.88218 coronin 7 Coro7
1428150_at 0.93743 coronin 7 Coro7
1428151_x_at 0.89890 cysteine conjugate-beta lyase 1 Ccb11
1428152_a_at 1.00421 Ribosomal protein L18A Rpl18a
1428153_at 1.15488 mitochondrial ribosomal protein S10 Mrps10
1428154_s_at 0.89303 phosphatidic acid phosphatase type 2 domain containing 1 Ppapdc1
1428155_at 0.99208 COMM domain containing 9 Commd9
1428156_at 0.99384 "guanine nucleotide binding protein (G protein), gamma 2 subunit" Gng2
1428157_at 0.98217 "guanine nucleotide binding protein (G protein), gamma 2 subunit" Gng2
1428158_at 0.95457 AKT1 substrate 1 (proline-rich) Akt1s1
1428159_s_at 1.01115 "NADH dehydrogenase (ubiquinone) 1, alpha/beta subcomplex, 1" Ndufab1
1428160_at 0.75467 "NADH dehydrogenase (ubiquinone) 1, alpha/beta subcomplex, 1" Ndufab1
1428161_a_at 0.96510 coiled-coil-helix-coiled-coil-helix domain containing 2 Chchd2
1428162_at 0.93474 RIKEN cDNA 4933421E11 gene 4933421E11Rik
1428163_at 1.01335 SAR1 gene homolog B (S. cerevisiae) Sar1b
1428164_at 0.96143 nudix (nucleoside diphosphate linked moiety X)-type motif 9 Nudt9
1428165_at 0.94542 vacuolar protein sorting 24 (yeast) Vps24
1428166_at 1.04004 "congenital dyserythropoietic anemia, type I (human)" Cdan1
1428167_a_at 0.72030 myelin protein zero-like 1 Mpzl1
1428168_at 0.95238 myelin protein zero-like 1 Mpzl1
1428169_at 0.92448 autophagy-related 16-like 1 (yeast) Atg16l1
1428170_at 1.02159 zinc finger protein 180 Zfp180
1428171_at 1.11370 PRP39 pre-mRNA processing factor 39 homolog (yeast) Prpf39
1428172_at 1.08633 PRP39 pre-mRNA processing factor 39 homolog (yeast) Prpf39
1428173_at 1.04177 echinoderm microtubule associated protein like 2 Eml2
1428174_x_at 1.00207 KH-type splicing regulatory protein Khgrp
1428175_at 0.90821 transmembrane protein 161B Tmem161b
1428176_at 1.06024 "endothelial differentiation, sphingolipid G-protein-coupled receptor, 5" Edg5
1428177_at 1.02009 trafficking protein particle complex 6B /// similar to Trafficking protein particle complex subunit 6B Trappc6b /// LOC433458
1428178_s_at 0.95928 trafficking protein particle complex 6B Trappc6b

1428179_at 1.02813 NADH dehydrogenase (ubiquinone) flavoprotein 2 Ndufv2
 1428180_at 1.01258 RIKEN cDNA 2810422J05 gene 2810422J05Rik
 1428181_at 1.02348 "electron transferring flavoprotein, beta polypeptide" Etfb
 1428182_at 1.02723 phosphoribosyl pyrophosphate synthetase-associated protein 1 Prpsap1
 1428183_at 0.92680 ankyrin repeat domain 23 Ankrd23
 1428184_at 1.02286 RIKEN cDNA 3110035E14 gene 3110035E14Rik
 1428185_at 0.93288 potassium channel tetramerisation domain containing 18 Kctd18
 1428186_at 0.87784 potassium channel tetramerisation domain containing 6 Kctd6
 1428187_at 1.21323 "CD47 antigen (Rh-related antigen, integrin-associated signal transducer)"
 Cd47
 1428188_at 1.02738 RIKEN cDNA 2610019A05 gene 2610019A05Rik
 1428189_at 0.94099 RIKEN cDNA 5730494M16 gene 5730494M16Rik
 1428190_at 1.21422 "solute carrier family 25 (mitochondrial carrier, citrate transporter), member
 1" Slc25a1
 1428191_s_at 0.99731 methyltransferase 11 domain containing 1 Mett11d1
 1428192_at 0.93633 kelch repeat and BTB (POZ) domain containing 7 Kbtbd7
 1428193_at 1.06004 "ubiquitin specific peptidase 9, X chromosome" Usp9x
 1428194_at 0.98259 "ubiquitin specific peptidase 9, X chromosome" Usp9x
 1428195_at 0.92388 RIKEN cDNA 4631427C17 gene 4631427C17Rik
 1428196_a_at 0.99189 RIKEN cDNA 1200015F23 gene 1200015F23Rik
 1428197_at 0.89288 tetraspanin 9 Tspan9
 1428198_at 0.93746 adenosine deaminase-like Adal
 1428199_at 0.81879 adenosine deaminase-like Adal
 1428200_a_at 0.84655 RIKEN cDNA 2210010A19 gene 2210010A19Rik
 1428201_at 1.02666 RIKEN cDNA 2310036O22 gene 2310036O22Rik
 1428202_at 0.86944 RIKEN cDNA 1810037C20 gene 1810037C20Rik
 1428203_at 1.24304 RIKEN cDNA C030002O17 gene C030002O17Rik
 1428204_at 0.98824 RIKEN cDNA C030002O17 gene C030002O17Rik
 1428205_x_at 1.13384 RIKEN cDNA C030002O17 gene C030002O17Rik
 1428206_at 0.97583 coiled-coil domain containing 69 Ccdc69
 1428207_at 0.70015 B-cell CLL/lymphoma 7A Bcl7a
 1428208_at 1.20619 B-cell CLL/lymphoma 7A Bcl7a
 1428209_at 0.87795 brain expressed gene 4 Bex4
 1428210_s_at 1.01344 conserved helix-loop-helix ubiquitous kinase Chuk
 1428211_at 0.87715 RIKEN cDNA 4933406E20 gene 4933406E20Rik
 1428212_x_at 1.04930 ribosomal protein L31 /// similar to ribosomal protein L31 /// similar to
 ribosomal protein L31 /// similar to ribosomal protein L31 /// similar to ribosomal protein L31 /// similar
 to ribosomal protein L31 /// similar to ribosomal protein L31 /// similar to ribosomal protein L31 ///
 similar to ribosomal protein L31 /// similar to ribosomal protein L31 Rpl31 /// LOC544868 ///
 LOC632618 /// LOC638399 /// LOC665562 /// LOC667682 /// LOC670381 /// LOC672957 ///
 LOC673582 /// LOC676592
 1428213_at 0.95989 RIKEN cDNA 2410003A14 gene 2410003A14Rik
 1428214_at 0.97690 translocase of outer mitochondrial membrane 7 homolog (yeast) Tomm7

1428215_x_at 0.92910 translocase of outer mitochondrial membrane 7 homolog (yeast) Tomm7
1428216_s_at 0.96505 translocase of outer mitochondrial membrane 7 homolog (yeast) Tomm7
1428217_at 0.89550 RIKEN cDNA 1600012H06 gene 1600012H06Rik
1428218_a_at 0.99481 RIKEN cDNA 1600012H06 gene 1600012H06Rik
1428219_at 0.86075 RING1 and YY1 binding protein /// similar to RING1 and YY1 binding protein Rybp /// LOC628746
1428220_at 1.03745 RIKEN cDNA 5730419I09 gene 5730419I09Rik
1428221_at 0.70273 kelch domain containing 8B Klhdc8b
1428222_at 0.93786 doublecortin and CaM kinase-like 2 Dcamk2
1428223_at 0.93331 major facilitator superfamily domain containing 2 Mfsd2
1428224_at 0.77297 heterogeneous nuclear ribonucleoprotein D-like Hnrpd1
1428225_s_at 1.04029 heterogeneous nuclear ribonucleoprotein D-like Hnrpd1
1428226_at 1.06382 "neuroguidin, EIF4E binding protein" Ngdn
1428227_at 1.00254 RE1-silencing transcription factor Rest
1428228_at 0.95902 phosphoglucomutase 3 Pgm3
1428229_at 0.95345 "protein kinase C, nu" Prkc
1428230_at 0.94129 "protein kinase C, nu" Prkc
1428231_at 1.00826 cleavage and polyadenylation specific factor 6 Cpsf6
1428232_at 1.06317 cleavage and polyadenylation specific factor 6 Cpsf6
1428233_at 0.97845 cleavage and polyadenylation specific factor 6 Cpsf6
1428234_at 1.08717 cleavage and polyadenylation specific factor 6 Cpsf6
1428235_at 1.04766 "succinate dehydrogenase complex, subunit D, integral membrane protein" Sdhd
1428236_at 1.03735 acyl-Coenzyme A binding domain containing 5 Acbd5
1428237_at 0.97934 RIKEN cDNA 2700059D21 gene 2700059D21Rik
1428238_at 0.92108 RIKEN cDNA 2700059D21 gene 2700059D21Rik
1428239_at 1.15036 ankyrin repeat domain 16 Ankrd16
1428240_at 0.96960 neurexin I Nrxa1
1428241_at 0.83705 RIKEN cDNA 2310035K24 gene 2310035K24Rik
1428242_at 0.95807 histocompatibility (minor) HA-1 Hmha1
1428243_at 1.02783 RIKEN cDNA 1700021K19 gene 1700021K19Rik
1428244_at 1.16107 "La ribonucleoprotein domain family, member 1 /// similar to la related protein isoform 1" Larpl /// LOC631268
1428245_at 1.00124 "glucose 6 phosphatase, catalytic, 3" G6pc3
1428246_at 1.03416 vacuolar protein sorting 26 homolog B (yeast) Vps26b
1428247_at 0.97168 vacuolar protein sorting 26 homolog B (yeast) Vps26b
1428248_at 1.07870 "nuclear transcription factor, X-box binding 1" Nfx1
1428249_at 0.98410 pygopus 2 Pygo2
1428250_at 0.76202 G protein-coupled receptor 30 Gpr30
1428251_at 0.99307 SMC hinge domain containing 1 Smchd1
1428252_at 1.03077 chromatin modifying protein 2B Chmp2b
1428253_at 1.01785 chromatin modifying protein 2B Chmp2b
1428254_at 1.06767 Purine rich element binding protein B Purb
1428255_at 0.89160 Luc7 homolog (S. cerevisiae)-like Luc7l

1428256_at 0.92826 RIKEN cDNA 2310047H23 gene 2310047H23Rik
1428257_s_at 1.13296 dynein light chain roadblock-type 1 Dynlrb1
1428258_at 0.95781 RIKEN cDNA 2010107E04 gene 2010107E04Rik
1428259_at 0.95822 peroxidasin homolog (Drosophila) Pxdn
1428260_at 0.82771 spastic paraplegia 3A homolog (human) Spg3a
1428261_at 0.99265 RIKEN cDNA 2310042L06 gene 2310042L06Rik
1428262_s_at 1.11911 heterogeneous nuclear ribonucleoprotein A3 /// similar to heterogeneous nuclear ribonucleoprotein A3 /// similar to heterogeneous nuclear ribonucleoprotein A3 /// similar to heterogeneous nuclear ribonucleoprotein A3 /// similar to Hnrpa3 /// LOC545592 /// LOC629025 /// LOC630401 /// LOC668559
1428263_a_at 1.07442 "transcription elongation factor B (SIII), polypeptide 2" Tceb2
1428264_at 1.07261 WD repeat domain 57 (U5 snRNP specific) Wdr57
1428265_at 1.26192 "protein phosphatase 2 (formerly 2A), regulatory subunit A (PR 65), beta isoform" Ppp2r1b
1428266_at 0.94012 "myosin, light polypeptide 3" Myl3
1428267_at 0.97500 DEAH (Asp-Glu-Ala-His) box polypeptide 40 Dhx40
1428268_at 0.87864 pleckstrin and Sec7 domain containing 2 Psd2
1428269_a_at 0.85001 glycosyltransferase 8 domain containing 1 Glt8d1
1428270_at 0.91439 glycosyltransferase 8 domain containing 1 Glt8d1
1428271_at 1.00480 acyl-Coenzyme A binding domain containing 4 Acbd4
1428272_at 0.97359 eukaryotic translation initiation factor 1B Eif1b
1428273_at 0.93099 abhydrolase domain containing 13 Abhd13
1428274_s_at 0.88582 abhydrolase domain containing 13 Abhd13
1428275_at 0.87835 abhydrolase domain containing 13 Abhd13
1428276_at 0.95861 abhydrolase domain containing 13 Abhd13
1428277_at 1.00209 OTU domain containing 6B Otud6b
1428278_at 1.01091 "zinc finger, DHHC domain containing 24" Zdhhc24
1428279_a_at 1.17062 ataxin 7-like 4 Atxn7l4
1428280_at 0.86660 FIP1 like 1 (S. cerevisiae) Fip111
1428281_at 1.02699 TruB pseudouridine (psi) synthase homolog 1 (E. coli) Trub1
1428282_at 1.06429 tubulin-specific chaperone e Tbce
1428283_at 1.29980 "cytochrome P450, family 2, subfamily s, polypeptide 1" Cyp2s1
1428284_at 0.94744 RIKEN cDNA 8430427H17 gene 8430427H17Rik
1428285_at 0.92905 RIKEN cDNA 8430427H17 gene 8430427H17Rik
1428286_at 1.03164 --- ---
1428287_at 1.04842 cullin 5 Cul5
1428288_at 0.82676 Kruppel-like factor 9 Klf9
1428289_at 0.79912 Kruppel-like factor 9 Klf9
1428290_at 0.97173 mitochondrial intermediate peptidase Mipep
1428291_at 1.06132 exosome component 8 Exosc8
1428292_at 0.99405 NADPH dependent diflavin oxidoreductase 1 Ndor1
1428293_at 0.95308 RIKEN cDNA 2310022M17 gene 2310022M17Rik
1428294_at 0.96532 zinc finger protein 259 Zfp259

1428295_at 1.01413 synaptopodin 2-like Synpo2l
1428296_at 0.91706 polymerase (RNA) II (DNA directed) polypeptide L Polr2l
1428297_at 1.01880 mitogen activated protein kinase kinase kinase 2 Map4k2
1428298_at 0.85420 RIKEN cDNA 1700029G01 gene 1700029G01Rik
1428299_at 0.96382 dual-specificity tyrosine-(Y)-phosphorylation regulated kinase 1a Dyrk1a
1428300_at 0.91439 RIKEN cDNA 4932439K10 gene 4932439K10Rik
1428301_at 0.96423 hypothetical protein LOC544988 /// RIKEN cDNA B930046C15 gene ///
hypothetical protein LOC545007 /// RIKEN cDNA 2610042L04 gene /// similar to Spetex-2H protein ///
similar to Spetex-2H protein /// hypothetical protein LOC664825 /// similar to Spetex-2F protein ///
similar to Spetex-2H protein /// similar to Spetex-2H protein /// similar to Spetex-2H protein ///
hypothetical protein LOC666740 /// hypothetical protein LOC666772 /// hypothetical protein
LOC666890 /// hypothetical protein LOC671957 LOC544988 /// B930046C15Rik /// LOC545007 ///
2610042L04Rik /// LOC622931 /// LOC638853 /// LOC664825 /// LOC666193 /// LOC666329 ///
LOC666442 /// LOC666684 /// LOC666740 /// LOC666772 /// LOC666890 /// LOC671957
1428302_at 1.11540 mitochondrial ribosomal protein L48 Mrpl48
1428303_at 1.00120 RIKEN cDNA 1500005I02 gene 1500005I02Rik
1428304_at 0.94247 establishment of cohesion 1 homolog 2 (S. cerevisiae) Esco2
1428305_at 0.93725 proprotein convertase subtilisin/kexin type 2 Pcsk2
1428306_at 1.56003 DNA-damage-inducible transcript 4 Ddit4
1428307_at 0.99102 "zinc finger, DHHC domain containing 13" Zdhhc13
1428308_at 1.00685 p53 and DNA damage regulated 1 Pdrg1
1428309_s_at 1.04176 p53 and DNA damage regulated 1 Pdrg1
1428310_at 0.96222 "La ribonucleoprotein domain family, member 7" Larp7
1428311_at 1.05244 coiled-coil domain containing 6 Ccdc6
1428312_at 1.02690 leucine rich repeat containing 57 Lrrc57
1428313_at 1.09133 "zinc finger, SWIM domain containing 3" Zswim3
1428314_at 0.97713 PEST proteolytic signal containing nuclear protein Pcnp
1428315_at 1.18887 EBNA1 binding protein 2 Ebna1bp2
1428316_a_at 1.04571 FUN14 domain containing 2 Fundc2
1428317_at 0.94620 --- ---
1428318_at 1.00085 submandibular gland protein C Smgc
1428319_at 0.78780 PDZ and LIM domain 7 Pdlim7
1428320_at 0.97365 jumonji domain containing 1B Jmjd1b
1428321_at 0.85631 echinoderm microtubule associated protein like 1 /// similar to echinoderm
microtubule associated protein like 1 Eml1 /// LOC634102
1428322_a_at 0.95802 "NADH dehydrogenase (ubiquinone) 1 beta subcomplex, 10" Ndufb10
1428323_at 1.13121 "glycerol phosphate dehydrogenase 2, mitochondrial" Gpd2
1428324_at 1.19684 --- ---
1428325_at 0.94103 RIKEN cDNA 2610019P18 gene 2610019P18Rik
1428326_s_at 1.04334 heat-responsive protein 12 Hrsp12
1428327_at 0.93472 "trafficking protein, kinesin binding 1" Trak1
1428328_at 0.94481 nucleoporin 50 Nup50
1428329_a_at 0.79665 intraflagellar transport 80 homolog (Chlamydomonas) Ift80

1428330_at	1.02471	dopey family member 2	Dopey2	
1428331_at	0.97672	RIKEN cDNA 2210016F16	gene	2210016F16Rik
1428332_at	0.70462	RIKEN cDNA 1500004A08	gene	1500004A08Rik
1428333_at	0.99268	RIKEN cDNA 2900062L11	gene	2900062L11Rik
1428334_at	1.06128	osteopetrosis associated transmembrane protein 1		Ostm1
1428335_a_at	0.96294	sec1 family domain containing 1	Scfd1	
1428336_at	0.99648	"1-acylglycerol-3-phosphate O-acyltransferase 1 (lysophosphatidic acid acyltransferase, delta)"	Agpat4	
1428337_at	0.98864	RIKEN cDNA 1810034K20	gene	1810034K20Rik
1428338_at	0.90290	RIKEN cDNA 2610039E05	gene	2610039E05Rik
1428339_at	0.94042	nudix (nucleoside diphosphate linked moiety X)-type motif 21	Nudt21	
1428340_s_at	0.84006	ATPase type 13A2	Atp13a2	
1428341_at	0.98422	zinc finger protein 655	Zfp655	
1428342_at	0.85986	REST corepressor 3	Rcor3	
1428343_at	0.94100	REST corepressor 3	Rcor3	
1428344_at	0.90107	phosphatidic acid phosphatase type 2 domain containing 2		Ppapdc2
1428345_at	0.89123	phosphatidic acid phosphatase type 2 domain containing 2		Ppapdc2
1428346_at	0.86297	TRAF type zinc finger domain containing 1	Trafd1	
1428347_at	0.92180	cytoplasmic FMR1 interacting protein 2	Cyfp2	
1428348_at	0.95840	zinc finger protein 336	Zfp336	
1428349_s_at	0.99585	early B-cell factor 3	Ebf3	
1428350_at	1.07155	RIKEN cDNA 2310061F22	gene	2310061F22Rik
1428351_at	0.97001	protein phosphatase 1M	Ppm1m	
1428352_at	1.38048	arrestin domain containing 2	Arrdc2	
1428353_at	1.12701	forkhead box K2	Foxk2	
1428354_at	1.19533	forkhead box K2	Foxk2	
1428355_at	1.08437	oxysterol binding protein 2	Osbp2	
1428356_at	0.92472	oxysterol binding protein 2	Osbp2	
1428357_at	0.93286	RIKEN cDNA 2610019F03	gene	2610019F03Rik
1428358_at	0.90500	RIKEN cDNA 1810010M01	gene	1810010M01Rik
1428359_s_at	0.92811	RIKEN cDNA 1810010M01	gene	1810010M01Rik
1428360_x_at	0.99029	"NADH dehydrogenase (ubiquinone) 1 alpha subcomplex, 7 (B14.5a)"		Ndufa7
1428361_x_at	0.87159	"hemoglobin alpha, adult chain 1"	Hba-a1	
1428362_at	0.98851	"eukaryotic translation initiation factor 4, gamma 2"	Eif4g2	
1428363_at	0.96253	"eukaryotic translation initiation factor 4, gamma 2"	Eif4g2	
1428364_at	1.10120	sodium channel modifier 1	Scnm1	
1428365_a_at	1.27400	"protease, serine, 15"	Prss15	
1428366_at	1.05730	RIKEN cDNA 1600027N09	gene	1600027N09Rik
1428367_at	0.94486	N-deacetylase/N-sulfotransferase (heparan glucosaminyl) 1		Ndst1
1428368_at	0.86218	Rho GTPase activating protein 21	Arhgap21	
1428369_s_at	0.88771	Rho GTPase activating protein 21	Arhgap21	
1428370_at	1.00004	RIKEN cDNA 1500011B03	gene	1500011B03Rik

1428371_at 0.94385 RIKEN cDNA 2610507N02 gene 2610507N02Rik
1428372_at 1.09422 suppression of tumorigenicity 5 St5
1428373_at 1.10464 inositol hexaphosphate kinase 2 Ihpk2
1428374_at 1.04120 glucuronyl C5-epimerase Glce
1428375_at 0.74256 RIKEN cDNA 4932415G12 gene 4932415G12Rik
1428376_at 0.90315 RIKEN cDNA 4932415G12 gene 4932415G12Rik
1428377_at 0.78816 BTB (POZ) domain containing 11 Btbd11
1428378_at 1.04783 "zinc finger CCCH type, antiviral 1" Zc3hav1
1428379_at 1.11915 "solute carrier family 17 (sodium-dependent inorganic phosphate cotransporter), member 6" Slc17a6
1428380_at 1.01721 RIKEN cDNA 0610007C21 gene 0610007C21Rik
1428381_a_at 0.93331 RIKEN cDNA 2700038C09 gene 2700038C09Rik
1428382_at 0.94332 "SWI/SNF related, matrix associated, actin dependent regulator of chromatin, subfamily c, member 2" Smarcc2
1428383_a_at 1.08930 RIKEN cDNA 2310021P13 gene 2310021P13Rik
1428384_at 0.98358 "DNA segment, Chr 4, Brigham & Women's Genetics 0951 expressed" D4Bwg0951e
1428385_at 0.92830 membrane-associated ring finger (C3HC4) 8 8-Mar
1428386_at 1.17532 acyl-CoA synthetase long-chain family member 3 Acsl3
1428387_at 1.13362 acyl-CoA synthetase long-chain family member 3 Acsl3
1428388_at 0.97155 "tankyrase, TRF1-interacting ankyrin-related ADP-ribose polymerase 2" Tnks2
1428389_s_at 1.10944 WD repeat domain 43 Wdr43
1428390_at 1.21596 WD repeat domain 43 Wdr43
1428391_at 0.75456 RAB3A interacting protein (rabin3)-like 1 Rab3il1
1428392_at 0.98002 Ras association (RalGDS/AF-6) domain family 2 Rassf2
1428393_at 0.88795 neuritin 1 Nrn1
1428394_at 0.49605 phytanoyl-CoA dioxygenase domain containing 1 /// leucine rich repeat containing 8A Phyhd1 /// Lrrc8a
1428395_at 1.04828 SMAD specific E3 ubiquitin protein ligase 1 /// similar to Smad ubiquitination regulatory factor 1 (Ubiquitin--protein ligase SMURF1) (Smad-specific E3 ubiquitin ligase 1) (hSMURF1) Smurf1 /// LOC640390
1428396_at 0.96993 SMAD specific E3 ubiquitin protein ligase 1 /// similar to Smad ubiquitination regulatory factor 1 (Ubiquitin--protein ligase SMURF1) (Smad-specific E3 ubiquitin ligase 1) (hSMURF1) Smurf1 /// LOC640390
1428397_at 0.95661 "UDP-Gal:betaGlcNAc beta 1,3-galactosyltransferase, polypeptide 5" B3galt5
1428398_at 0.98971 "UDP-Gal:betaGlcNAc beta 1,3-galactosyltransferase, polypeptide 5" B3galt5
1428399_a_at 1.00835 armadillo repeat containing 9 Armc9
1428400_at 0.95427 RIKEN cDNA 2200002K05 gene 2200002K05Rik
1428401_at 0.87741 "zinc finger, CCHC domain containing 3" Zcchc3
1428402_at 0.90824 "zinc finger, CCHC domain containing 3" Zcchc3
1428403_at 0.98242 RIKEN cDNA 2410025L10 gene 2410025L10Rik

1428404_at	0.85407	RIKEN cDNA 2410025L10 gene	2410025L10Rik	
1428405_at	0.93289	host cell factor C1 regulator 1 (XPO1-dependent)	Hcfc1r1	
1428406_s_at	0.92998	host cell factor C1 regulator 1 (XPO1-dependent)	Hcfc1r1	
1428407_at	0.96222	heterogeneous nuclear ribonucleoprotein A0	Hnrpa0	
1428408_a_at	1.01712	"DNA segment, Chr 12, ERATO Doi 551, expressed"	D12Ert551e	
1428409_at	1.17514	N-acetyltransferase 13	Nat13	
1428410_at	1.00160	N-acetyltransferase 13	Nat13	
1428411_at	0.94892	RIKEN cDNA 1700020I14 gene	1700020I14Rik	
1428412_at	0.96742	transmembrane 9 superfamily member 3	Tm9sf3	
1428413_at	0.80796	RIKEN cDNA 5730405I09 gene	5730405I09Rik	
1428414_at	1.09158	RIKEN cDNA 5730405I09 gene	5730405I09Rik	
1428415_at	0.88830	ring finger protein 113A2	Rnf113a2	
1428416_at	0.84673	RIKEN cDNA 3110050N22 gene	3110050N22Rik	
1428417_at	0.82833	RIKEN cDNA 3110050N22 gene	3110050N22Rik	
1428418_s_at	1.01180	RIKEN cDNA 3110050N22 gene	3110050N22Rik	
1428419_at	1.04103	RIKEN cDNA 5430411K18 gene	5430411K18Rik	
1428420_a_at	1.06787	RIKEN cDNA 1200009I06 gene	1200009I06Rik	
1428421_a_at	1.00618	RIKEN cDNA 2700085E05 gene	2700085E05Rik	
1428422_at	1.07328	UBX domain containing 8	Ubx8	
1428423_at	0.93029	polycomb group ring finger 3	Pcgf3	
1428424_at	1.01869	polycomb group ring finger 3	Pcgf3	
1428425_at	1.00632	"transforming growth factor, beta receptor associated protein 1"	Tgfbrap1	
1428426_s_at	0.96577	"transforming growth factor, beta receptor associated protein 1"	Tgfbrap1	
1428427_at	1.03022	F-box and leucine-rich repeat protein 2	Fbxl2	
1428428_at	1.17206	abhydrolase domain containing 11	Abhd11	
1428429_at	1.03919	"RGM domain family, member B"	Rgmb	
1428430_at	0.91452	"RGM domain family, member B"	Rgmb	
1428431_at	0.77125	RIKEN cDNA 2310047A01 gene	2310047A01Rik	
1428432_at	0.78783	RIKEN cDNA 2310047A01 gene	2310047A01Rik	
1428433_at	0.69561	homeodomain interacting protein kinase 2	Hipk2	
1428434_at	1.00864	"zinc finger, CCHC domain containing 12"	Zcchc12	
1428435_at	0.93764	mucin 2	Muc2	
1428436_at	0.98426	"LSM14 homolog A (SCD6, S. cerevisiae)"	Lsm14a	
1428437_at	0.84814	"LSM14 homolog A (SCD6, S. cerevisiae)"	Lsm14a	
1428438_s_at	0.93610	"LSM14 homolog A (SCD6, S. cerevisiae)"	Lsm14a	
1428439_at	1.02809	negative regulator of ubiquitin-like proteins 1	Nub1	
1428440_at	0.92304	"solute carrier family 25 (mitochondrial carrier, Aralar), member 12"	Slc25a12	
1428441_at	1.06870	RIKEN cDNA 1500009M05 gene	1500009M05Rik	
1428442_at	1.05457	cDNA sequence BC029722	BC029722	
1428443_a_at	0.74635	Rap1 GTPase-activating protein	Rap1gap	
1428444_at	0.99892	ankyrin repeat and SOCS box-containing protein 2	Asb2	
1428445_at	0.88534	Yip1 interacting factor homolog B (S. cerevisiae)	Yif1b	
1428446_at	0.82509	dynein cytoplasmic 2 light intermediate chain 1	Dync2li1	

1428447_at	0.99635	transmembrane protein 14A	Tmem14a	
1428448_a_at	0.99208	"general transcription factor IIIC, polypeptide 2, beta"		Gtf3c2
1428449_at	0.86906	"general transcription factor IIIC, polypeptide 2, beta"		Gtf3c2
1428450_at	0.90536	RIKEN cDNA 2610034B18 gene	2610034B18Rik	
1428451_at	1.01069	leucine rich repeat containing 47	Lrrc47	
1428452_at	1.06400	RIKEN cDNA 2810025M15 gene	2810025M15Rik	
1428453_at	0.92450	N-acetyltransferase 12	Nat12	
1428454_at	0.90178	breast carcinoma amplified sequence 3	Bcas3	
1428455_at	0.93215	"procollagen, type XIV, alpha 1"	Col14a1	
1428456_at	1.05631	bolA-like 1 (E. coli)	Bol1	
1428457_at	1.03534	RIKEN cDNA 5830472M02 gene	5830472M02Rik	
1428458_at	1.28853	"processing of precursor 1, ribonuclease P/MRP family, (S. cerevisiae)"	Pop1	
1428459_at	1.09917	PRAME family member 12	Pramef12	
1428460_at	0.97832	synapsin II	Syn2	
1428461_at	1.09680	"protein phosphatase 2, regulatory subunit B (B56), epsilon isoform"		Ppp2r5e
1428462_at	0.86322	"protein phosphatase 2, regulatory subunit B (B56), epsilon isoform"		Ppp2r5e
1428463_a_at	0.89631	"protein phosphatase 2, regulatory subunit B (B56), epsilon isoform"		Ppp2r5e
1428464_at	0.92571	"NADH dehydrogenase (ubiquinone) 1 alpha subcomplex, 3"	Ndufa3	
1428465_at	1.01435	transmembrane protein 147	Tmem147	
1428466_at	0.96602	chromodomain helicase DNA binding protein 3	Chd3	
1428467_at	0.84268	TAR DNA binding protein	Tardbp	
1428468_at	0.98693	RIKEN cDNA 3110043O21 gene	3110043O21Rik	
1428469_a_at	1.03447	DAZ interacting protein 1	Dzip1	
1428470_at	0.98952	exocyst complex component 2	Exoc2	
1428471_at	0.67202	sorbin and SH3 domain containing 1	Sorbs1	
1428472_at	0.75912	splA/ryanodine receptor domain and SOCS box containing 1	Spsb1	
1428473_at	0.98008	"protein phosphatase 3, catalytic subunit, beta isoform"		Ppp3cb
1428474_at	0.87363	"protein phosphatase 3, catalytic subunit, beta isoform"		Ppp3cb
1428475_at	1.21477	RIKEN cDNA 1190005P17 gene	1190005P17Rik	
1428476_a_at	1.12133	elaC homolog 2 (E. coli)	Elac2	
1428477_at	1.02232	elaC homolog 2 (E. coli)	Elac2	
1428478_at	1.03616	receptor accessory protein 4	Reep4	
1428479_at	1.07668	"nuclear factor of activated T-cells, cytoplasmic, calcineurin-dependent 1"		Nfatc1
1428480_at	1.08796	cell division cycle associated 8	Cdca8	
1428481_s_at	0.99490	cell division cycle associated 8	Cdca8	
1428482_at	1.09132	A kinase (PRKA) anchor protein 10	Akap10	
1428483_a_at	0.97547	RIKEN cDNA 2610039C10 gene	2610039C10Rik	
1428484_at	1.11702	oxysterol binding protein-like 3	Osbpl3	
1428485_at	0.88463	carbonic anhydrase 12	Car12	
1428486_at	0.98259	RIKEN cDNA 1700123L14 gene	1700123L14Rik	
1428487_s_at	0.87762	coenzyme Q10 homolog B (S. cerevisiae)	Coq10b	
1428488_at	0.95950	"phosphatidylinositol glycan anchor biosynthesis, class K"		Pigk

1428489_at 0.99659 zinc finger CCHC-type and RNA binding motif 1 Zcrb1
 1428490_at 0.99145 "core 1 UDP-galactose:N-acetylgalactosamine-alpha-R beta 1,3-galactosyltransferase" C1galt1
 1428491_at 1.06587 COMM domain containing 10 Commd10
 1428492_at 0.99066 GLI pathogenesis-related 2 Glipr2
 1428493_at 1.07078 RIKEN cDNA 2610511M17 gene /// similar to signal-induced proliferation-associated 1 like 3 2610511M17Rik /// LOC672426
 1428494_a_at 0.94567 polymerase (RNA) II (DNA directed) polypeptide I Polr2i
 1428495_at 1.07386 RIKEN cDNA 2410003K15 gene 2410003K15Rik
 1428496_at 1.08075 SECIS binding protein 2 Secisbp2
 1428497_at 1.08064 SECIS binding protein 2 Secisbp2
 1428498_at 0.90922 RIKEN cDNA 2610206B13 gene 2610206B13Rik
 1428499_at 0.93260 RIKEN cDNA 2810454L23 gene 2810454L23Rik
 1428500_at 0.93064 RIKEN cDNA 2210419D22 gene 2210419D22Rik
 1428501_at 0.89853 RIKEN cDNA 1110051M20 gene 1110051M20Rik
 1428502_at 0.97365 ARP6 actin-related protein 6 homolog (yeast) Actr6
 1428503_a_at 1.10822 NFkB inhibitor interacting Ras-like protein 1 Nkiras1
 1428504_at 1.01906 X-ray repair complementing defective repair in Chinese hamster cells 6 Xrcc6
 1428505_at 0.94600 RIKEN cDNA 2310015N07 gene 2310015N07Rik
 1428506_at 1.13061 5-aminoimidazole-4-carboxamide ribonucleotide formyltransferase/IMP cyclohydrolase Atic
 1428507_at 0.99514 haloacid dehalogenase-like hydrolase domain containing 2 Hdhd2
 1428508_at 0.88581 "TBC1 domain family, member 2B" Tbc1d2b
 1428509_at 0.90884 myosin IE Myo1e
 1428510_at 0.69305 latrophilin 1 Lphn1
 1428511_at 1.10724 "phosphorylase kinase, gamma 2 (testis)" Phkg2
 1428512_at 0.98973 "basic helix-loop-helix domain containing, class B9" Bhlhb9
 1428513_at 0.89054 calcium binding and coiled coil domain 1 Calcoco1
 1428514_at 1.14957 copine III Cpne3
 1428515_at 1.16142 RIKEN cDNA 2410012H22 gene 2410012H22Rik
 1428516_a_at 1.06384 "alkB, alkylation repair homolog 7 (E. coli)" Alkbh7
 1428517_at 0.89555 WD repeat and FYVE domain containing 3 Wdfy3
 1428518_at 1.06757 myeloid leukemia factor 1 interacting protein Mlf1ip
 1428519_at 1.08485 RIKEN cDNA 2610528E23 gene 2610528E23Rik
 1428520_at 1.00490 RIKEN cDNA 1110032A13 gene 1110032A13Rik
 1428521_at 1.01325 "THAP domain containing, apoptosis associated protein 3" Thap3
 1428522_at 0.96179 "transcription termination factor, RNA polymerase II" Ttf2
 1428523_at 0.90084 latrophilin 3 Lphn3
 1428524_at 0.81141 RIKEN cDNA 2310051N18 gene 2310051N18Rik
 1428525_at 0.97166 RIKEN cDNA 4930488B01 gene 4930488B01Rik
 1428526_at 1.08195 LSM domain containing 1 Lsmd1
 1428527_at 0.99053 sorting nexin 7 Snx7
 1428528_at 0.96155 RIKEN cDNA 1110007L15 gene 1110007L15Rik

1428529_at 1.05554 RIKEN cDNA 2810026P18 gene 2810026P18Rik
 1428530_x_at 1.07182 ribosomal protein S24 /// similar to ribosomal protein S24 /// similar to
 ribosomal protein S24 /// similar to ribosomal protein S24 Rps24 /// LOC629116 /// LOC636056 ///
 LOC677113
 1428531_at 1.03659 integrator complex subunit 7 Ints7
 1428532_at 0.99754 integrator complex subunit 7 Ints7
 1428533_at 0.97691 "DNA segment, Chr 1, Brigham & Women's Genetics 0212 expressed"
 D1Bwg0212e
 1428534_at 1.09462 RIKEN cDNA 2310073E15 gene 2310073E15Rik
 1428535_at 1.01782 RIKEN cDNA 9430020K01 gene 9430020K01Rik
 1428536_at 0.95411 "potassium voltage-gated channel, subfamily G, member 4" Kcng4
 1428537_at 1.02502 "casein kinase 1, alpha 1" Csnk1a1
 1428538_s_at 1.04338 retinoic acid receptor responder (tazarotene induced) 2 Rarres2
 1428539_at 1.05191 RIKEN cDNA 2610207I05 gene 2610207I05Rik
 1428540_at 0.83057 RIKEN cDNA 3321401G04 gene 3321401G04Rik
 1428541_at 0.98753 RIKEN cDNA 3321401G04 gene 3321401G04Rik
 1428542_at 1.10757 RIKEN cDNA 2010003J03 gene 2010003J03Rik
 1428543_at 1.05607 phosphoribosyl pyrophosphate amidotransferase Ppat
 1428544_at 0.99955 RIKEN cDNA 0610007L01 gene 0610007L01Rik
 1428545_at 1.02428 RIKEN cDNA 0610007L01 gene 0610007L01Rik
 1428546_at 1.01694 "synaptotagmin binding, cytoplasmic RNA interacting protein" Syncrip
 1428547_at 0.44609 "5' nucleotidase, ecto" Nt5e
 1428548_at 0.94049 p21 (CDKN1A)-activated kinase 4 Pak4
 1428549_at 0.87903 coiled-coil domain containing 3 Ccdc3
 1428550_at 0.87345 RIKEN cDNA 1810015A11 gene 1810015A11Rik
 1428551_at 1.11452 RIKEN cDNA 2410075D05 gene 2410075D05Rik
 1428552_at 1.04036 RIKEN cDNA 2610001J05 gene 2610001J05Rik
 1428553_at 1.10739 glutaredoxin 5 homolog (*S. cerevisiae*) Glrx5
 1428554_a_at 1.00885 RIKEN cDNA 1810035L17 gene 1810035L17Rik
 1428555_at 1.04048 SET domain containing 2 Setd2
 1428556_at 1.02123 "phosphatidylinositol glycan anchor biosynthesis, class Y" Pigy
 1428557_a_at 1.06774 O-sialoglycoprotein endopeptidase-like 1 Osgepl1
 1428558_at 1.01494 O-sialoglycoprotein endopeptidase-like 1 Osgepl1
 1428559_at 0.97295 follicular lymphoma variant translocation 1 Fvt1
 1428560_at 1.16690 exportin 5 Xpo5
 1428561_at 1.02377 RIKEN cDNA 2610002J23 gene 2610002J23Rik
 1428562_at 1.26344 RIKEN cDNA 2210403K04 gene 2210403K04Rik
 1428563_at 1.13286 DEAD (Asp-Glu-Ala-Asp) box polypeptide 10 Ddx10
 1428564_at 0.93933 zinc finger protein 579 Zfp579
 1428565_at 1.08783 unkempt-like (*Drosophila*) Unkl
 1428566_at 1.04849 Hspb associated protein 1 Hspbap1
 1428567_at 1.08060 Hspb associated protein 1 Hspbap1
 1428568_at 1.12262 RIKEN cDNA B230217C12 gene B230217C12Rik

1428569_at 1.04644 RIKEN cDNA B230217C12 gene B230217C12Rik
1428570_at 1.05151 cyclin C Ccnc
1428571_at 0.87028 "procollagen, type IX, alpha 1" Col9a1
1428572_at 1.07585 "brain abundant, membrane attached signal protein 1" Basp1
1428573_at 0.78386 chimerin (chimaerin) 2 Chn2
1428574_a_at 0.95333 chimerin (chimaerin) 2 Chn2
1428575_at 0.87761 FCH domain only 1 Fcho1
1428576_at 1.01317 "hypoxia-inducible factor 1, alpha subunit inhibitor" Hif1an
1428577_at 0.95715 "protein tyrosine phosphatase, receptor type, f polypeptide (PTPRF), interacting protein (liprin), alpha 4" Ppfia4
1428578_s_at 1.00694 "protein tyrosine phosphatase, receptor type, f polypeptide (PTPRF), interacting protein (liprin), alpha 4" Ppfia4
1428579_at 1.12779 formin-like 2 Fmnl2
1428580_at 1.09809 biliverdin reductase A Blvra
1428581_at 0.95586 RIKEN cDNA 1700024G10 gene 1700024G10Rik
1428582_at 1.13351 RIKEN cDNA 2010208K18 gene 2010208K18Rik
1428583_at 0.98746 nuclear fragile X mental retardation protein interacting protein 2 Nufip2
1428584_a_at 1.14604 hydroxyacylglutathione hydrolase-like Haghl
1428585_at 1.02059 "actinin, alpha 1" Actn1
1428586_at 1.02494 transmembrane protein 41B Tmem41b
1428587_at 1.08426 transmembrane protein 41B Tmem41b
1428588_a_at 0.99529 mitochondrial ribosomal protein L41 Mrpl41
1428589_at 1.10078 mitochondrial ribosomal protein L41 Mrpl41
1428590_at 0.87063 mitochondrial ribosomal protein L41 Mrpl41
1428591_at 0.86853 patatin-like phospholipase domain containing 2 Pnpla2
1428592_s_at 0.81432 ubiquitin specific peptidase 38 Usp38
1428593_at 1.10584 RIKEN cDNA 1700029F09 gene 1700029F09Rik
1428594_at 0.92258 GTPase activating RANGAP domain-like 1 Garnl1
1428595_at 0.84713 "solute carrier family 6 (neurotransmitter transporter), member 19" Slc6a19
1428596_at 0.81722 "TBC1 domain family, member 9B" Tbc1d9b
1428597_at 0.90236 "TBC1 domain family, member 9B" Tbc1d9b
1428598_at 1.06553 "TBC1 domain family, member 7" Tbc1d7
1428599_at 1.25170 kinase non-catalytic C-lobe domain (KIND) containing 1 Kndc1
1428600_at 1.07639 RIKEN cDNA 3110068G20 gene 3110068G20Rik
1428601_at 0.91454 RIKEN cDNA 1700003E16 gene 1700003E16Rik
1428602_at 1.17356 "potassium inwardly-rectifying channel, subfamily J, member 9" Kcnj9
1428603_at 0.93303 glucocorticoid induced transcript 1 Glcci1
1428604_at 0.99193 RIKEN cDNA 2610305D13 gene 2610305D13Rik
1428605_at 1.05610 similar to Y73E7A.1 LOC669879
1428606_at 0.83078 coiled-coil domain containing 124 Ccdc124
1428607_at 0.90136 v-raf murine sarcoma 3611 viral oncogene homolog Araf
1428608_at 0.93988 "myosin light chain, regulatory B" Mylc2b
1428609_at 0.88235 "myosin light chain, regulatory B /// similar to myosin regulatory light chain-

like /// similar to Myosin regulatory light chain 2-B, smooth muscle isoform (Myosin RLC-B)"
Mylc2b /// LOC630701 /// LOC640689

1428610_at	0.89331	autophagy-related 7 (yeast)	Atg7	
1428611_at	1.02410	autophagy-related 7 (yeast)	Atg7	
1428612_at	0.87656	autophagy-related 7 (yeast)	Atg7	
1428613_at	0.96650	lactate dehydrogenase D	Ldhd	
1428614_at	1.04302	lactate dehydrogenase D	Ldhd	
1428615_at	0.98106	"purinergic receptor P2Y, G-protein coupled, 5"	P2ry5	
1428616_at	1.00214	zinc finger protein 131	Zfp131	
1428617_at	1.03735	host cell factor C2	Hcfc2	
1428618_at	1.13143	host cell factor C2	Hcfc2	
1428619_at	0.81660	RIKEN cDNA 2310005N03 gene	2310005N03Rik	
1428620_at	0.97151	endosulfine alpha	Ensa	
1428621_a_at	1.06980	vacuolar protein sorting 25 (yeast)	Vps25	
1428622_at	0.82827	DEP domain containing 6	Depdc6	
1428623_at	0.94232	plexin A1	Plxna1	
1428624_at	1.03348	RIKEN cDNA 2810482I07 gene	2810482I07Rik	
1428625_a_at	0.99543	"RMI1, RecQ mediated genome instability 1, homolog (S. cerevisiae)"	Rmi1	
1428626_at	0.99710	"LysM, putative peptidoglycan-binding, domain containing 2"	Lysmd2	
1428627_at	0.97980	zinc finger protein 509	Zfp509	
1428628_at	1.00864	RIKEN cDNA 1190002A17 gene	1190002A17Rik	
1428629_at	0.95503	zinc finger protein 518	Zfp518	
1428630_x_at	1.16076	hydroxyacylglutathione hydrolase-like	Haghl	
1428631_a_at	0.99822	ubiquinol cytochrome c reductase core protein 2	Uqcrc2	
1428632_at	0.95930	RIKEN cDNA 1810006K21 gene	1810006K21Rik	
1428633_at	1.14382	TWIST neighbor /// similar to TWIST neighbor /// similar to TWIST neighbor	Twistnb /// LOC627278 /// LOC667501	
1428634_at	1.00582	TWIST neighbor	Twistnb	
1428635_at	1.09829	catechol-O-methyltransferase domain containing 1	Comtd1	
1428636_at	1.12143	six transmembrane epithelial antigen of prostate 2	Steap2	
1428637_at	1.05374	dual-specificity tyrosine-(Y)-phosphorylation regulated kinase 2	Dyrk2	
1428638_at	1.00809	EF-hand domain (C-terminal) containing 2	Efhc2	
1428639_at	1.00066	lin-9 homolog (C. elegans)	Lin9	
1428640_at	1.03158	heat shock transcription factor 2 binding protein	Hsf2bp	
1428641_at	0.87880	heat shock transcription factor 2 binding protein	Hsf2bp	
1428642_at	1.06409	"solute carrier family 35, member D3"	Slc35d3	
1428643_at	0.79354	mannoside acetylglucosaminyltransferase 5	Mgat5	
1428644_at	1.02347	mannoside acetylglucosaminyltransferase 5	Mgat5	
1428645_at	0.96471	"guanine nucleotide binding protein, alpha inhibiting 3"	Gnai3	
1428646_at	0.82543	region containing RIKEN cDNA 2310056B04 gene; pre B-cell leukemia transcription factor 1	LOC676870	
1428647_at	0.80128	region containing RIKEN cDNA 2310056B04 gene; pre B-cell leukemia transcription factor 1	LOC676870	

1428648_at 1.07468 cullin associated and neddylation disassociated 1 Cand1
1428649_at 1.01751 cullin associated and neddylation disassociated 1 Cand1
1428650_at 0.67816 tensin 1 Tns1
1428651_at 0.81792 kelch-like 24 (Drosophila) Klh24
1428652_at 0.95365 RIKEN cDNA 0610010F05 gene 0610010F05Rik
1428653_x_at 0.91344 "ELAV (embryonic lethal, abnormal vision, Drosophila)-like 1 (Hu antigen R)" Elavl1
1428654_at 0.88774 RIKEN cDNA 1200016B10 gene 1200016B10Rik
1428655_at 1.03039 RIKEN cDNA 1110018J12 gene 1110018J12Rik
1428656_at 1.03005 "ribonuclease III, nuclear" Rnasen
1428657_at 0.87457 ras responsive element binding protein 1 Rreb1
1428658_at 0.98894 "protein (peptidyl-prolyl cis/trans isomerase) NIMA-interacting, 4 (parvulin) /// similar to Peptidyl-prolyl cis-trans isomerase NIMA-interacting 4 (Rotamase Pin4) (PPIase Pin4) /// similar to Peptidyl-prolyl cis-trans isomerase NIMA-interacting 4 (Rotamase Pin4) (PPIase Pin4)" Pin4 /// LOC628161 /// LOC635999
1428659_at 1.05658 PHD finger protein 7 Phf7
1428660_s_at 1.32230 "torsin family 3, member A" Tor3a
1428661_at 0.91235 nuclear factor of kappa light polypeptide gene enhancer in B-cells inhibitor-like 2 Nfkbil2
1428662_a_at 0.88460 homeobox only domain Hod
1428663_at 0.98343 RIKEN cDNA 5133401H06 gene 5133401H06Rik
1428664_at 1.00903 vasoactive intestinal polypeptide Vip
1428665_at 1.01976 --- ---
1428666_at 1.09358 asparaginyl-tRNA synthetase Nars
1428667_at 0.85003 monoamine oxidase A Maoa
1428668_at 0.90808 acyl-Coenzyme A binding domain containing 3 Acbd3
1428669_at 1.00671 brain expressed myelocytomatosis oncogene Bmyc
1428670_at 1.18196 RIKEN cDNA 2610305J24 gene /// similar to melanoma antigen /// hypothetical protein LOC669436 /// hypothetical protein LOC670270 /// hypothetical protein LOC672052 /// similar to melanoma antigen /// similar to melanoma antigen 2610305J24Rik /// LOC668904 /// LOC669436 /// LOC670270 /// LOC672052 /// LOC672939 /// LOC677506
1428671_at 1.18346 RIKEN cDNA 2200002D01 gene 2200002D01Rik
1428672_at 1.11122 small nuclear ribonucleoprotein polypeptide F Snrpf
1428673_at 1.05387 symplekin Sympk
1428674_at 1.05247 PRP38 pre-mRNA processing factor 38 (yeast) domain containing B Prpf38b
1428675_at 0.96092 RIKEN cDNA 1110049F12 gene 1110049F12Rik
1428676_at 1.05786 transmembrane serine protease 6 Tmprss6
1428677_at 1.00773 WD repeat domain 73 Wdr73
1428678_s_at 1.06122 WD repeat domain 73 Wdr73
1428679_s_at 1.08386 RIKEN cDNA 0610010K14 gene 0610010K14Rik
1428680_at 1.07387 CDP-diacylglycerol synthase 1 Cds1
1428681_at 0.97889 "gene model 608, (NCBI)" Gm608
1428682_at 0.72193 zinc finger CCCH type containing 6 Zc3h6

1428683_at 1.02006 RIKEN cDNA A930016P21 gene A930016P21Rik
1428684_at 1.08790 RIKEN cDNA 1500001M20 gene 1500001M20Rik
1428685_at 1.03270 synaptonemal complex central element protein 1 Syce1
1428686_at 1.04009 RIKEN cDNA 2810029C07 gene 2810029C07Rik
1428687_at 0.97321 zinc finger protein 687 Zfp687
1428688_at 1.16371 programmed cell death protein 11 Pcd11
1428689_at 1.01824 trypsin domain containing 1 Tysnd1
1428690_at 1.06391 trypsin domain containing 1 Tysnd1
1428691_at 1.01413 RIKEN cDNA 5630401D06 gene 5630401D06Rik
1428692_at 0.81908 HD domain containing 3 Hddc3
1428693_at 0.85209 RIKEN cDNA 2610044O15 gene 2610044O15Rik
1428694_at 1.16331 RIKEN cDNA 5033413D16 gene 5033413D16Rik
1428695_at 1.01648 RIKEN cDNA 2310014G06 gene 2310014G06Rik
1428696_at 1.04530 RIKEN cDNA 2310015N21 gene 2310015N21Rik
1428697_at 0.89986 dipeptidylpeptidase 8 Dpp8
1428698_at 0.88098 RIKEN cDNA 2310004I03 gene 2310004I03Rik
1428699_at 1.17251 spermine synthase Sms
1428700_at 0.95299 "purinergic receptor P2Y, G-protein coupled 13" P2ry13
1428701_at 1.00168 poliovirus receptor-related 4 Pvr14
1428702_at 1.07215 DEAD (Asp-Glu-Ala-Asp) box polypeptide 28 Ddx28
1428703_at 0.95494 RIKEN cDNA 1700012B07 gene 1700012B07Rik
1428704_at 1.08384 zinc finger protein 661 Zfp661
1428705_at 1.05195 RIKEN cDNA 1700007K13 gene 1700007K13Rik
1428706_at 0.97035 proline-rich polypeptide 6 Prr6
1428707_at 0.85889 parathymosin Ptms
1428708_x_at 0.87699 parathymosin Ptms
1428709_a_at 1.08377 mitochondrial ribosomal protein L24 Mrpl24
1428710_at 0.85109 Ras-like without CAAX 1 Rit1
1428711_at 1.07931 MON1 homolog b (yeast) Mon1b
1428712_at 1.04565 MON1 homolog b (yeast) Mon1b
1428713_s_at 0.99936 RIKEN cDNA 4833427B12 gene 4833427B12Rik
1428714_at 0.96718 progesterone receptor membrane component 2 Pgrmc2
1428715_at 1.00424 RIKEN cDNA 2810423A18 gene 2810423A18Rik
1428716_at 0.96862 peroxisome biogenesis factor 1 Pex1
1428717_at 1.17938 secernin 1 Scrn1
1428718_at 1.13826 secernin 1 Scrn1
1428719_at 0.92198 RIKEN cDNA 2010309G21 gene /// hypothetical protein LOC207685
2010309G21Rik /// LOC207685
1428720_s_at 0.96657 "immunoglobulin lambda chain, variable 1 /// RIKEN cDNA 2010309G21
gene /// hypothetical protein LOC207685 /// similar to Ig lambda-3 chain C region - mouse" Igl-V1 ///
2010309G21Rik /// LOC207685 /// LOC386520
1428721_at 1.01964 BTB (POZ) domain containing 5 Btbd5
1428722_at 0.96187 "creatine kinase, mitochondrial 2" Ckmt2

1428723_at 1.10196 RIKEN cDNA 2310047M10 gene 2310047M10Rik
1428724_at 0.78555 cleavage and polyadenylation factor subunit homolog (*S. cerevisiae*) Pcf11
1428725_at 0.93516 protein inhibitor of activated STAT 2 Pias2
1428726_at 1.01473 THUMP domain containing 2 Thumpd2
1428727_at 1.10972 RIKEN cDNA 4631422C13 gene 4631422C13Rik
1428728_at 0.97566 DEAD (Asp-Glu-Ala-Asp) box polypeptide 51 Ddx51
1428729_at 1.10953 "KRIT1, ankyrin repeat containing" Krit1
1428730_at 1.10741 "KRIT1, ankyrin repeat containing" Krit1
1428731_at 0.98124 ubiquitin specific peptidase 54 Usp54
1428732_at 0.99656 RIKEN cDNA 1700008J07 gene 1700008J07Rik
1428733_at 1.02261 "guanine nucleotide binding protein (G protein), gamma transducing activity polypeptide 2" Gngt2
1428734_at 0.80122 RIKEN cDNA 3200002M19 gene 3200002M19Rik
1428735_at 0.99804 CD69 antigen Cd69
1428736_at 0.86050 GRAM domain containing 3 Gramd3
1428737_s_at 0.81832 GRAM domain containing 3 Gramd3
1428738_a_at 0.97348 "DNA segment, Chr 14, ERATO Doi 449, expressed" D14Ertd449e
1428739_at 0.96457 RIKEN cDNA 2310040A07 gene 2310040A07Rik
1428740_a_at 0.75868 "phosphatidylinositol glycan anchor biosynthesis, class T" Pigt
1428741_at 0.97384 "ELAV (embryonic lethal, abnormal vision, *Drosophila*)-like 4 (Hu antigen D)" Elavl4
1428742_at 1.02767 F-box protein 45 Fbxo45
1428743_at 0.92732 Bri3 binding protein Bri3bp
1428744_s_at 0.97912 Bri3 binding protein Bri3bp
1428745_a_at 0.92192 RIKEN cDNA 2310003L22 gene 2310003L22Rik
1428746_a_at 1.06453 tRNA selenocysteine associated protein 1 Trspap1
1428747_at 1.00315 tRNA selenocysteine associated protein 1 Trspap1
1428748_at 1.45079 RIKEN cDNA 5830428H23 gene 5830428H23Rik
1428749_at 1.09155 Dmx-like 2 Dmxl2
1428750_at 1.11431 CDC42 effector protein (Rho GTPase binding) 2 Cdc42ep2
1428751_at 0.95701 Park2 co-regulated Pacrg
1428752_at 0.98746 "solute carrier family 5 (sodium/glucose cotransporter), member 11" Slc5a11
1428753_a_at 1.01317 DiGeorge syndrome critical region gene 6 Dgcr6
1428754_at 1.12452 RIKEN cDNA 3300001M20 gene 3300001M20Rik
1428755_at 1.03452 cAMP responsive element binding protein 1 Creb1
1428756_at 0.96229 aminoadipate-semialdehyde dehydrogenase-phosphopantetheinyl transferase Aasdhppt
1428757_at 1.01508 aminoadipate-semialdehyde dehydrogenase-phosphopantetheinyl transferase Aasdhppt
1428758_at 0.69435 transmembrane protein 86A Tmem86a
1428759_s_at 0.81182 coiled-coil domain containing 49 Ccdc49
1428760_at 1.05378 "small nuclear RNA activating complex, polypeptide 3" Snapc3
1428761_a_at 1.25152 "small nuclear RNA activating complex, polypeptide 3" Snapc3

1428762_at	0.86223	mitogen-activated protein kinase kinase kinase 7 interacting protein 3	
Map3k7ip3			
1428763_at	1.06738	single-strand selective monofunctional uracil DNA glycosylase	Smug1
1428764_at	1.06798	"GTL2, imprinted maternally expressed untranslated mRNA"	Gtl2
1428765_at	1.04266	"GTL2, imprinted maternally expressed untranslated mRNA"	Gtl2
1428766_at	0.99298	RNA methyltransferase like 1	Rnmtl1
1428767_at	1.20188	gasdermin domain containing 1	Gsdmdc1
1428768_at	0.97855	PAN3 polyA specific ribonuclease subunit homolog (S. cerevisiae)	Pan3
1428769_at	1.01010	TatD DNase domain containing 3	Tatdn3
1428770_at	0.98993	DIP2 disco-interacting protein 2 homolog A (Drosophila)	Dip2a
1428771_at	0.87863	RIKEN cDNA 2410127E18 gene	2410127E18Rik
1428772_at	1.25137	"exportin, tRNA (nuclear export receptor for tRNAs)"	Xpot
1428773_s_at	0.79492	Bcl6 interacting corepressor	Bcor
1428774_at	0.71097	glypican 6	Gpc6
1428775_at	1.01956	RIKEN cDNA 1110008L16 gene	1110008L16Rik
1428776_at	0.95346	"solute carrier family 10 (sodium/bile acid cotransporter family), member 6"	
Slc10a6			
1428777_at	1.09295	"sprouty protein with EVH-1 domain 1, related sequence"	Spred1
1428778_at	1.11421	"Sfi1 homolog, spindle assembly associated (yeast)"	Sfi1
1428779_at	0.79844	zinc finger and BTB domain containing 41 homolog	Zbtb41
1428780_at	0.98981	threonine aldolase 1	Tha1
1428781_at	0.97241	dermokine	Dmkn
1428782_a_at	0.98269	ubiquinol-cytochrome c reductase core protein 1	Uqcrc1
1428783_at	0.95531	"protein kinase, cAMP dependent regulatory, type II alpha"	Prkar2a
1428784_at	1.06156	Gem-interacting protein	Gmip
1428785_at	1.06983	angiomin-like 1	Amotl1
1428786_at	0.93677	NCK associated protein 1 like	Nckap11
1428787_at	1.03314	NCK associated protein 1 like	Nckap11
1428788_at	1.39226	RIKEN cDNA 1700012G19 gene	1700012G19Rik
1428789_at	1.05814	Ral GEF with PH domain and SH3 binding motif 2	Ralgps2
1428790_at	0.95135	Nfat activating molecule with ITAM motif 1	Nfam1
1428791_at	0.82891	ubiquitin-conjugating enzyme E2H	Ube2h
1428792_at	1.19973	breast carcinoma amplified sequence 1	Bcas1
1428793_at	1.05234	"solute carrier family 36 (proton/amino acid symporter), member 1"	Slc36a1
1428794_at	1.07697	spectrin domain with coiled-coils 1	Specc1
1428795_at	0.76483	RIKEN cDNA 1110021L09 gene	1110021L09Rik
1428796_at	0.95685	RIKEN cDNA 5530401J07 gene	5530401J07Rik
1428797_at	1.01621	RIKEN cDNA 0610039J04 gene	0610039J04Rik
1428798_s_at	1.06572	RIKEN cDNA 0610039J04 gene	0610039J04Rik
1428799_at	1.06038	RIKEN cDNA 4930431B11 gene	4930431B11Rik
1428800_a_at	0.93186	pseudouridylate synthase 7 homolog (S. cerevisiae)-like	Pus7l
1428801_at	1.04633	mannoside acetylglucosaminyltransferase 3	Mgat3
1428802_at	0.90132	mannoside acetylglucosaminyltransferase 3	Mgat3

1428803_at 1.01418 RIKEN cDNA 4632408A20 gene 4632408A20Rik
1428804_at 1.14003 microfibrillar-associated protein 3-like Mfap3l
1428805_at 1.02555 "solute carrier family 35, member E3" Slc35e3
1428806_at 1.03691 "casein kinase 1, gamma 1" Csnk1g1
1428807_at 1.07465 RIKEN cDNA 4932702K14 gene 4932702K14Rik
1428808_at 0.99715 prickle-like 2 (Drosophila) Prickle2
1428809_at 0.97045 RIKEN cDNA 1810010H24 gene /// hypothetical protein LOC671668 ///
hypothetical protein LOC674875 1810010H24Rik /// LOC671668 /// LOC674875
1428810_at 1.02299 RIKEN cDNA 2700097O09 gene 2700097O09Rik
1428811_at 1.13924 "WAP, follistatin/kazal, immunoglobulin, kunitz and netrin domain containing
2" Wfikkn2
1428812_at 0.93733 RIKEN cDNA 1700040L02 gene 1700040L02Rik
1428813_a_at 0.92768 dopamine receptor D1 interacting protein Drd1ip
1428814_at 1.10496 --- ---
1428815_at 1.09223 RIKEN cDNA 4930583C14 gene 4930583C14Rik
1428816_a_at 0.92775 GATA binding protein 2 Gata2
1428817_at 0.89780 MON2 homolog (yeast) Mon2
1428818_at 0.86584 MON2 homolog (yeast) Mon2
1428819_at 1.20750 "microtubule-associated protein, RP/EB family, member 1" Mapre1
1428820_at 1.39067 "microtubule-associated protein, RP/EB family, member 1" Mapre1
1428821_at 0.95688 "1-acylglycerol-3-phosphate O-acyltransferase 2 (lysophosphatidic acid
acyltransferase, beta)" Agpat2
1428822_a_at 0.81242 sorting nexin 24 Snx24
1428823_at 1.26503 HD domain containing 2 Hddc2
1428824_at 1.01173 RIKEN cDNA 2310003C23 gene 2310003C23Rik
1428825_at 1.05460 "nuclear receptor subfamily 6, group A, member 1" Nr6a1
1428826_at 1.19379 "nuclear receptor subfamily 6, group A, member 1" Nr6a1
1428827_at 1.01352 Wolf-Hirschhorn syndrome candidate 1 (human) Whsc1
1428828_at 1.04850 RPA interacting protein Rpain
1428829_at 1.00240 RIKEN cDNA 6820401H01 gene 6820401H01Rik
1428830_at 1.28233 ataxia telangiectasia mutated homolog (human) Atm
1428831_at 1.03151 RIKEN cDNA 6230429P13 gene 6230429P13Rik
1428832_at 0.98202 RIKEN cDNA 1600002H07 gene 1600002H07Rik
1428833_at 0.95266 RIKEN cDNA 4930406D14 gene 4930406D14Rik
1428834_at 1.44978 dual specificity phosphatase 4 Dusp4
1428835_at 0.84320 "myosin, heavy polypeptide 14 /// similar to nonmuscle myosin heavy chain"
Myh14 /// LOC672865
1428836_at 0.92051 RIKEN cDNA 2300009A05 gene 2300009A05Rik
1428837_at 1.00437 kelch-like 14 (Drosophila) Kllh14
1428838_a_at 0.94842 deoxycytidine kinase Dck
1428839_at 1.08279 WD repeat domain 53 Wdr53
1428840_s_at 1.01048 WD repeat domain 53 Wdr53
1428841_at 0.85706 vitelliform macular dystrophy 2 homolog (human) Vmd2

1428842_a_at 0.99996 nerve growth factor receptor (TNFRSF16) associated protein 1 Ngfrap1
 1428843_at 0.96488 membrane-associated ring finger (C3HC4) 5 5-Mar
 1428844_a_at 1.04582 BCL2-associated transcription factor 1 Bclaf1
 1428845_at 1.01476 BCL2-associated transcription factor 1 Bclaf1
 1428846_at 0.96932 tetratricopeptide repeat domain 14 Ttc14
 1428847_a_at 0.94563 microtubule-actin crosslinking factor 1 Macf1
 1428848_a_at 1.04376 microtubule-actin crosslinking factor 1 Macf1
 1428849_at 0.95483 "ribosomal protein S6 kinase, polypeptide 1" Rps6kb1
 1428850_x_at 0.94047 CD99 antigen Cd99
 1428851_at 1.02285 RIKEN cDNA 1300014I06 gene 1300014I06Rik
 1428852_at 0.86390 dedicator of cyto-kinesis 3 Dock3
 1428853_at 0.88381 patched homolog 1 Ptch1
 1428854_at 0.91608 transmembrane emp24 domain containing 8 Tmed8
 1428855_at 1.09333 histocompatibility 13 H13
 1428856_at 1.00596 Histocompatibility 13 H13
 1428857_at 0.90638 "Sin3A associated protein, 130kDa" Sap130
 1428858_at 0.95061 WD repeat domain 70 Wdr70
 1428859_at 0.98922 polyamine oxidase (exo-N4-amino) Paox
 1428860_at 0.90646 RIKEN cDNA 4930572J05 gene 4930572J05Rik
 1428861_at 0.88450 RIKEN cDNA 4631422O05 gene /// similar to downregulated in ovarian cancer
 1 isoform 2 /// similar to downregulated in ovarian cancer 1 isoform 2 4631422O05Rik ///
 LOC667770 /// LOC676704
 1428862_at 0.93873 tetratricopeptide repeat domain 17 Ttc17
 1428863_at 0.92255 ankyrin repeat domain 39 Ankrd39
 1428864_at 0.98689 RIKEN cDNA 5530400B01 gene 5530400B01Rik
 1428865_at 0.99106 BCL2-like 12 (proline rich) Bcl2l12
 1428866_at 0.97783 RIKEN cDNA 2810037O22 gene 2810037O22Rik
 1428867_at 0.88386 RIKEN cDNA 4933417E01 gene 4933417E01Rik
 1428868_a_at 1.08680 ornithine decarboxylase antizyme Oaz1
 1428869_at 1.33253 nucleolar and coiled-body phosphoprotein 1 Nolc1
 1428870_at 1.12587 nucleolar and coiled-body phosphoprotein 1 Nolc1
 1428871_at 1.07591 RIKEN cDNA 4121402D02 gene 4121402D02Rik
 1428872_at 1.01966 RIKEN cDNA 4121402D02 gene 4121402D02Rik
 1428873_a_at 1.07086 RIKEN cDNA 4121402D02 gene 4121402D02Rik
 1428874_at 1.01695 RIKEN cDNA 1110019N10 gene 1110019N10Rik
 1428875_at 0.93434 golgi phosphoprotein 4 Golph4
 1428876_at 0.99465 signal recognition particle 72 Srp72
 1428877_at 0.97906 signal recognition particle 72 Srp72
 1428878_a_at 0.90081 "phosphatidylinositol transfer protein, cytoplasmic 1" Pitpnc1
 1428879_at 0.88459 "phosphatidylinositol transfer protein, cytoplasmic 1" Pitpnc1
 1428880_at 1.04140 RIKEN cDNA 0610038F07 gene 0610038F07Rik
 1428881_at 1.09869 kinesin 2 Kns2
 1428882_at 1.09139 "WD repeat domain, phosphoinositide interacting 2" Wipi2

1428883_at	0.84966	transmembrane protein 57	Tmem57
1428884_at	0.99652	transmembrane protein 57	Tmem57
1428885_at	1.12600	arylformamidase	Afmid
1428886_at	1.05705	retinoblastoma binding protein 5	Rbbp5
1428887_at	0.87831	RIKEN cDNA 2610020C11 gene	2610020C11Rik
1428888_at	0.96170	transmembrane protein 33	Tmem33
1428889_at	1.00768	"alkB, alkylation repair homolog 3 (E. coli)"	Alkbh3
1428890_at	0.93278	fem-1 homolog c (C.elegans)	Fem1c
1428891_at	0.84003	---	---
1428892_at	1.04396	peptidylprolyl isomerase (cyclophilin)-like 1	Ppil1
1428893_at	1.04611	oxysterol binding protein-like 7	Osbp17
1428894_at	1.00357	RIKEN cDNA 1300018J18 gene	1300018J18Rik
1428895_at	1.03602	RIKEN cDNA 3222401M22 gene	3222401M22Rik
1428896_at	0.95677	platelet-derived growth factor receptor-like	Pdgfrl
1428897_at	1.11477	RIKEN cDNA 2610029I01 gene	2610029I01Rik
1428898_at	0.82971	MON1 homolog A (yeast)	Mon1a
1428899_at	1.04926	RIKEN cDNA 2310079P10 gene	2310079P10Rik
1428900_s_at	0.74197	methyltransferase 5 domain containing 1	Mett5d1
1428901_at	1.06758	DTW domain containing 2	Dtwd2
1428902_at	1.00196	carbohydrate sulfotransferase 11	Chst11
1428903_at	1.03395	RIKEN cDNA 3110037I16 gene	3110037I16Rik
1428904_at	1.16784	RIKEN cDNA E230022H04 gene	E230022H04Rik
1428905_at	1.04374	Ras-related GTP binding A	Rraga
1428906_at	1.02622	"general transcription factor IIH, polypeptide 5"	Gtf2h5
1428907_at	1.12092	RIKEN cDNA 2600011C06 gene	2600011C06Rik
1428908_at	1.20614	RIKEN cDNA 2600011C06 gene	2600011C06Rik
1428909_at	1.95939	RIKEN cDNA A130040M12 gene	A130040M12Rik
1428910_at	0.98378	RIKEN cDNA 2310022B05 gene	2310022B05Rik
1428911_at	1.02382	"tubulin tyrosine ligase-like family, member 4"	Ttll4
1428912_at	0.96842	"tubulin tyrosine ligase-like family, member 4"	Ttll4
1428913_at	1.04564	"DNA segment, Chr 2, Brigham & Women's Genetics 1335 expressed"	D2Bwg1335e
1428914_at	0.81796	RIKEN cDNA 2310014D11 gene	2310014D11Rik
1428915_at	0.87845	sirtuin 5 (silent mating type information regulation 2 homolog) 5 (S. cerevisiae)	Sirt5
1428916_s_at	0.87657	sirtuin 5 (silent mating type information regulation 2 homolog) 5 (S. cerevisiae)	Sirt5
1428917_at	0.83939	syntaxin 17	Stx17
1428918_at	1.04826	SCY1-like 3 (S. cerevisiae)	Scyl3
1428919_at	1.39676	Fgfr1 oncogene partner	Fgfr1op
1428920_at	1.37494	---	---
1428921_at	1.15070	RIKEN cDNA 2810021B07 gene	2810021B07Rik
1428922_at	1.30324	RIKEN cDNA 1200009O22 gene	1200009O22Rik

1428923_at	0.97446	"protein phosphatase 1, regulatory (inhibitor) subunit 3G"	Ppp1r3g
1428924_at	0.89816	molybdenum cofactor synthesis 3	Mocs3
1428925_at	0.91759	SUMO1/sentrin specific peptidase 1	Senp1
1428926_at	0.83753	RIKEN cDNA 1110003O08 gene	1110003O08Rik
1428927_at	1.06331	SHQ1 homolog (<i>S. cerevisiae</i>)	Shq1
1428928_at	0.83207	"calcium channel, voltage-dependent, beta 4 subunit"	Cacnb4
1428929_s_at	0.87453	"solute carrier family 25 (mitochondrial carrier, phosphate carrier), member 26"	Slc25a26
1428930_at	0.89958	transmembrane protein 29	Tmem29
1428931_a_at	0.92973	"poly (ADP-ribose) polymerase family, member 6"	Parp6
1428932_at	1.07160	tetratricopeptide repeat domain 21A	ttc21a
1428933_at	1.03341	histone deacetylase 8	Hdac8
1428934_at	0.94190	CKLF-like MARVEL transmembrane domain containing 2B	Cmtm2b
1428935_at	0.88808	calnexin	Canx
1428936_at	1.20390	"ATPase, Ca ⁺⁺ transporting, plasma membrane 1"	Atp2b1
1428937_at	1.07929	"ATPase, Ca ⁺⁺ transporting, plasma membrane 1"	Atp2b1
1428938_at	0.98224	"guanine nucleotide binding protein, alpha q polypeptide"	Gnaq
1428939_s_at	0.95041	"guanine nucleotide binding protein, alpha q polypeptide"	Gnaq
1428940_at	1.03678	"guanine nucleotide binding protein, alpha q polypeptide"	Gnaq
1428941_at	0.73037	"zinc finger, MYM-type 2"	Zmym2
1428942_at	0.90220	metallothionein 2	Mt2
1428943_at	1.12441	nudix (nucleoside diphosphate linked moiety X)-type motif 13	Nudt13
1428944_at	1.03719	ubiquitin-activating enzyme E1-like 2	Ube1l2
1428945_at	1.06608	ubiquitin-activating enzyme E1-like 2	Ube1l2
1428946_at	1.02692	ubiquitin-activating enzyme E1-like 2	Ube1l2
1428947_at	1.07471	RIKEN cDNA 2010001M09 gene	2010001M09Rik
1428948_at	0.93078	RIKEN cDNA 5730414M22 gene	5730414M22Rik
1428949_at	1.12572	"exportin, tRNA (nuclear export receptor for tRNAs)"	Xpot
1428950_s_at	1.18105	nucleolar protein 8	Nol8
1428951_at	1.20096	nucleolar protein 8	Nol8
1428952_at	1.04919	protein disulfide isomerase associated 2	Pdia2
1428953_at	0.96357	OTU domain containing 7B	Otud7b
1428954_at	0.95845	"solute carrier family 9 (sodium/hydrogen exchanger), isoform 3 regulator 2"	Slc9a3r2
1428955_x_at	0.91705	"solute carrier family 9 (sodium/hydrogen exchanger), isoform 3 regulator 2"	Slc9a3r2
1428956_at	0.89368	RIKEN cDNA 2300008B03 gene	2300008B03Rik
1428957_at	1.06419	RIKEN cDNA 2300008B03 gene	2300008B03Rik
1428958_at	0.80629	progesterin and adipoQ receptor family member VIII	Paqr8
1428959_at	1.00654	RIKEN cDNA 5830405N20 gene	5830405N20Rik
1428960_at	0.82082	RIKEN cDNA 4933434I06 gene	4933434I06Rik
1428961_a_at	0.89665	"splicing factor, arginine/serine-rich 16 (suppressor-of-white-apricot homolog, <i>Drosophila</i>)"	Sfrs16

1428962_at	0.91913	RIKEN cDNA 1700013F07 gene	1700013F07Rik
1428963_at	0.73194	RWD domain containing 2	Rwdd2
1428964_at	0.95073	"solute carrier family 25 (mitochondrial carrier), member 18"	Slc25a18
1428965_at	0.99905	coiled-coil domain containing 54	Ccdc54
1428966_at	0.99695	RIKEN cDNA 2610204K14 gene	2610204K14Rik
1428967_at	0.95123	insulin-like growth factor I receptor	Igflr
1428968_at	1.02204	centrosomal protein 57	Cep57
1428969_at	0.97311	RIKEN cDNA D030074E01 gene	D030074E01Rik
1428970_at	1.02493	N-acetyltransferase 13	Nat13
1428971_at	0.98054	RIKEN cDNA 5730405I09 gene	5730405I09Rik
1428972_at	1.13631	RIKEN cDNA 0610012D17 gene	0610012D17Rik
1428973_s_at	1.18917	RIKEN cDNA 0610012D17 gene	0610012D17Rik
1428974_s_at	0.85293	leucine zipper transcription factor-like 1	Lztfl1
1428975_at	1.02739	sushi domain containing 3	Susd3
1428976_at	0.91063	thymopoietin	Tmpo
1428977_at	0.56866	carbohydrate (N-acetylgalactosamine 4-0) sulfotransferase 8	Chst8
1428978_at	0.90384	RIKEN cDNA 2900009J20 gene	2900009J20Rik
1428979_at	0.98480	metal response element binding transcription factor 1	Mtf1
1428980_at	0.99270	RIKEN cDNA 1110001M24 gene	1110001M24Rik
1428981_at	1.09388	RIKEN cDNA 2810007J24 gene	2810007J24Rik
1428982_at	1.02548	RIKEN cDNA D530031C13 gene	D530031C13Rik
1428983_at	0.85450	scleraxis	Scx
1428984_a_at	0.91461	RIKEN cDNA 1700012B09 gene	1700012B09Rik
1428985_at	0.75797	integrator complex subunit 12	Ints12
1428986_at	0.80226	"solute carrier family 17 (sodium-dependent inorganic phosphate cotransporter), member 7"	Slc17a7
1428987_at	0.83829	dynein light chain roadblock-type 2	Dynlrb2
1428988_at	1.14673	"ATP-binding cassette, sub-family C (CFTR/MRP), member 3"	Abcc3
1428989_at	0.90636	RIKEN cDNA 0710001D07 gene	0710001D07Rik
1428990_at	0.80122	RIKEN cDNA 2310047K21 gene	2310047K21Rik
1428991_at	1.16194	HRAS-like suppressor	Hrasls
1428992_at	0.94236	unc-13 homolog D (C. elegans)	Unc13d
1428993_at	0.98116	RIKEN cDNA 1110017I16 gene	1110017I16Rik
1428994_s_at	0.92996	RIKEN cDNA 1110017I16 gene	1110017I16Rik
1428995_at	0.88430	5-oxoprolinase (ATP-hydrolysing)	Oplah
1428996_at	1.19649	RIKEN cDNA 4833426J09 gene	4833426J09Rik
1428997_at	1.03415	PHD finger protein 23	Phf23
1428998_at	0.99445	PHD finger protein 3	Phf3
1428999_at	1.01385	PHD finger protein 3	Phf3
1429000_at	0.98114	PHD finger protein 3	Phf3
1429001_at	1.11974	pirin	Pir
1429002_at	0.90993	SNW domain containing 1	Snw1
1429003_at	0.85510	SNW domain containing 1	Snw1

1429004_at	0.97286	pleckstrin homology domain interacting protein	Phip
1429005_at	1.01813	malignant fibrous histiocytoma amplified sequence 1	Mfhas1
1429006_s_at	1.04519	RIKEN cDNA 2610110G12 gene	2610110G12Rik
1429007_at	1.10293	"solute carrier family 35, member B2"	Slc35b2
1429008_at	1.15016	RIKEN cDNA 1810074P20 gene	1810074P20Rik
1429009_at	1.23258	U1 small nuclear ribonucleoprotein polypeptide A	Snrp70
1429010_at	0.97060	U1 small nuclear ribonucleoprotein polypeptide A	Snrp70
1429011_x_at	0.93595	U1 small nuclear ribonucleoprotein polypeptide A	Snrp70
1429012_at	0.99758	Rac/Cdc42 guanine nucleotide exchange factor (GEF) 6	Arhgef6
1429013_at	0.92710	RIKEN cDNA 2900002G04 gene	2900002G04Rik
1429014_at	0.81603	"DNA segment, Chr 1, ERATO Doi 161, expressed"	D1Ert161e
1429015_at	0.95259	"DNA segment, Chr 1, ERATO Doi 161, expressed"	D1Ert161e
1429016_at	0.99935	"DNA segment, Chr 1, ERATO Doi 161, expressed /// galactosidase, beta 1-like"	D1Ert161e /// Glb1l
1429017_at	1.04498	"Smith-Magenis syndrome chromosome region, candidate 8 homolog (human)"	Smcr8
1429018_at	1.09312	RIKEN cDNA 3830408D24 gene	3830408D24Rik
1429019_s_at	0.95747	paraoxonase 2	Pon2
1429020_at	0.99326	paraoxonase 2	Pon2
1429021_at	0.87320	Eph receptor A4	Epha4
1429022_at	0.99751	adenylate cyclase activating polypeptide 1 receptor 1	Adcyap1r1
1429023_at	1.07858	RIKEN cDNA 2900042E01 gene	2900042E01Rik
1429024_at	0.86696	RIKEN cDNA 1110018J23 gene	1110018J23Rik
1429025_a_at	0.75702	hexamethylene bis-acetamide inducible 2	Hexim2
1429026_at	0.85680	hexamethylene bis-acetamide inducible 2	Hexim2
1429027_at	0.89374	RIKEN cDNA 0610007N19 gene	0610007N19Rik
1429028_at	1.27003	dedicator of cytokinesis 11	Dock11
1429029_at	1.03670	RIKEN cDNA 4933405A16 gene	4933405A16Rik
1429030_at	0.91115	C1q and tumor necrosis factor related protein 7	C1qtnf7
1429031_at	0.87245	RIKEN cDNA 4833408D11 gene	4833408D11Rik
1429032_at	0.87469	MICAL C-terminal like	Micalcl
1429033_at	0.75837	golgi coiled coil 1	Gcc1
1429034_at	1.13538	essential meiotic endonuclease 1 homolog 2 (S. pombe)	Eme2
1429035_at	1.01273	dipeptidase 3	Dpep3
1429036_at	0.99006	otopetrin 3	Otop3
1429037_at	0.97377	RIKEN cDNA 1700019A02 gene	1700019A02Rik
1429038_at	1.04057	RIKEN cDNA 1500034J01 gene	1500034J01Rik
1429039_s_at	0.88619	RIKEN cDNA 1500034J01 gene	1500034J01Rik
1429040_at	0.93838	RIKEN cDNA 2610005L07 gene	2610005L07Rik
1429041_at	1.30826	RIKEN cDNA 2610005L07 gene	2610005L07Rik
1429042_at	0.89924	RIKEN cDNA 2010200O16 gene	2010200O16Rik
1429043_at	0.92955	survival motor neuron domain containing 1	Smndc1
1429044_at	0.96953	calmodulin regulated spectrin-associated protein 1-like 1	Camsap111

1429045_at 1.08591 SMAD specific E3 ubiquitin protein ligase 2 Smurf2
1429046_at 0.94159 SMAD specific E3 ubiquitin protein ligase 2 Smurf2
1429047_at 1.02000 "Rtf1, Paf1/RNA polymerase II complex component, homolog (S. cerevisiae)" Rtf1
1429048_at 1.08613 "biogenesis of lysosome-related organelles complex-1, subunit 2" Bloc1s2
1429049_at 0.89660 "NUAK family, SNF1-like kinase, 2" Nuak2
1429050_at 0.82006 cysteine-rich hydrophobic domain 2 Chic2
1429051_s_at 0.86170 SRY-box containing gene 11 Sox11
1429052_at 0.58090 "protein tyrosine phosphatase, receptor type, D" Ptpnd
1429053_at 0.97146 RIKEN cDNA 1110012J17 gene 1110012J17Rik
1429054_at 1.09966 mitochondrial ribosomal protein L47 Mrpl47
1429055_at 0.94459 RIKEN cDNA 4930506M07 gene 4930506M07Rik
1429056_at 1.11950 NMDA receptor regulated 1-like Narg11
1429057_at 1.26906 NMDA receptor regulated 1-like Narg11
1429058_at 0.86789 RIKEN cDNA 1110004B13 gene 1110004B13Rik
1429059_s_at 0.88317 RIKEN cDNA 1110004B13 gene 1110004B13Rik
1429060_at 1.45758 --- ---
1429061_at 1.11910 RIKEN cDNA 1810063B05 gene 1810063B05Rik
1429062_at 0.81636 kinesin family member 16B Kif16b
1429063_s_at 0.86111 kinesin family member 16B Kif16b
1429064_at 0.91559 DIP2 disco-interacting protein 2 homolog C (Drosophila) Dip2c
1429065_at 0.77990 RIKEN cDNA 1200009F10 gene 1200009F10Rik
1429066_at 0.97748 RIKEN cDNA 4930565B19 gene 4930565B19Rik
1429067_at 1.00870 "calpain, small subunit 2" Capns2
1429068_at 0.95122 RIKEN cDNA 2810488G03 gene 2810488G03Rik
1429069_at 0.96610 STIP1 homology and U-Box containing protein 1 Stub1
1429070_at 0.91923 RIKEN cDNA 4933440H19 gene 4933440H19Rik
1429071_at 0.94243 "malic enzyme 3, NADP(+)-dependent, mitochondrial" Me3
1429072_at 0.91969 RIKEN cDNA 1110001D15 gene 1110001D15Rik
1429073_at 0.93222 RIKEN cDNA 2210015D19 gene 2210015D19Rik
1429074_at 0.95372 RIKEN cDNA 1700026D08 gene 1700026D08Rik
1429075_a_at 0.97299 RIKEN cDNA 1700018B08 gene 1700018B08Rik
1429076_a_at 0.96217 glycerophosphodiester phosphodiesterase domain containing 2 Gdpd2
1429077_x_at 1.04409 ribosomal protein L21 /// similar to 60S ribosomal protein L21 /// similar to ribosomal protein L21 /// similar to ribosomal protein L21 /// similar to ribosomal protein L21 /// similar to ribosomal protein L21 /// similar to ribosomal protein L21 /// similar to ribosomal protein L21 /// similar to ribosomal protein L21 /// similar to ribosomal protein L21 /// similar to ribosomal protein L21 /// similar to ribosomal protein L21 Rpl21 /// LOC433125 /// LOC434067 /// LOC545423 /// LOC635258 /// LOC636855 /// LOC639424 /// LOC674827 /// LOC675278 /// LOC676253 /// LOC676590 /// LOC676825 /// LOC676903
1429078_a_at 1.20490 "COP9 (constitutive photomorphogenic) homolog, subunit 7a (Arabidopsis thaliana)" Cops7a
1429079_a_at 1.09632 RIKEN cDNA 2310045N01 gene 2310045N01Rik

1429080_at 1.08296 M-phase phosphoprotein 10 (U3 small nucleolar ribonucleoprotein) Mphosph10

1429081_at 0.91050 GRIP and coiled-coil domain containing 2 Gcc2

1429082_at 1.09093 "DNA segment, Chr 5, ERATO Doi 585, expressed" D5Ertd585e

1429083_at 1.02343 "amylo-1,6-glucosidase, 4-alpha-glucanotransferase" Agl

1429084_at 0.90791 vascular endothelial zinc finger 1 Vezf1

1429085_at 1.00945 vascular endothelial zinc finger 1 Vezf1

1429086_at 0.99039 grainyhead-like 2 (Drosophila) Grhl2

1429087_at 0.85626 RIKEN cDNA 1110054O05 gene 1110054O05Rik

1429088_at 0.68425 limb-bud and heart Lbh

1429089_s_at 0.90056 RIKEN cDNA 2900026A02 gene 2900026A02Rik

1429090_at 1.07875 RIKEN cDNA A030007L17 gene A030007L17Rik

1429091_at 0.99217 RIKEN cDNA 1600002K03 gene 1600002K03Rik

1429092_at 1.01278 "vitamin K epoxide reductase complex, subunit 1-like 1" Vkorc111

1429093_at 1.04039 DNA-damage inducible protein 2 Ddi2

1429094_at 1.05585 DNA-damage inducible protein 2 Ddi2

1429095_at 0.97945 centromere protein P Cenpp

1429096_at 0.90969 RIKEN cDNA 2810455D13 gene 2810455D13Rik

1429097_at 0.94566 RIKEN cDNA C030044C12 gene C030044C12Rik

1429098_s_at 0.89340 nonhomologous end-joining factor 1 Nhej1

1429099_at 1.04421 RIKEN cDNA 1110051B16 gene 1110051B16Rik

1429100_at 0.82327 RIKEN cDNA 2010109K11 gene 2010109K11Rik

1429101_at 1.04170 gap junction membrane channel protein alpha 5 Gja5

1429102_at 1.00333 arrestin domain containing 5 Arrdc5

1429103_at 1.02190 translocase of outer mitochondrial membrane 22 homolog (yeast) Tomm22

1429104_at 0.95863 LIM domain containing 2 Limd2

1429105_at 1.03346 "discs, large (Drosophila) homolog-associated protein 1" Dlgap1

1429106_at 1.23925 RIKEN cDNA 4921509J17 gene 4921509J17Rik

1429107_at 1.00362 RIKEN cDNA 1110059H15 gene 1110059H15Rik

1429108_at 0.87869 male-specific lethal 2-like 1 (Drosophila) Msl211

1429109_at 0.96957 male-specific lethal 2-like 1 (Drosophila) Msl211

1429110_a_at 1.06910 "NOL1/NOP2/Sun domain family, member 4" Nsun4

1429111_at 0.95888 talin 2 /// similar to talin 2 Tln2 /// LOC639214

1429112_at 1.00440 talin 2 /// similar to talin 2 Tln2 /// LOC639214

1429113_at 0.79202 RIKEN cDNA 2900092E17 gene /// RIKEN cDNA 1500031I19 gene 2900092E17Rik /// 1500031I19Rik

1429114_at 1.00473 SEC14 and spectrin domains 1 Sestd1

1429115_at 0.95636 RIKEN cDNA 2010003O02 gene 2010003O02Rik

1429116_at 0.90699 "solute carrier family 17 (anion/sugar transporter), member 5" Slc17a5

1429117_at 0.87000 TNFRSF1A-associated via death domain Tradd

1429118_a_at 1.02203 RIKEN cDNA 4930429M06Rik 4930429M06Rik

1429119_at 1.01707 RIKEN cDNA 4833421E05 gene 4833421E05Rik

1429120_at 0.87179 limb and neural patterns Lnp

1429121_at	1.02643	RIKEN cDNA 4921517N04 gene	4921517N04Rik
1429122_a_at	0.95515	RIKEN cDNA 1700040I03 gene	1700040I03Rik
1429123_at	0.89423	"RAB27A, member RAS oncogene family"	Rab27a
1429124_s_at	1.04551	"REX1, RNA exonuclease 1 homolog (S. cerevisiae)"	Rexo1
1429125_at	0.96106	zinc finger and BTB domain containing 9	Zbtb9
1429126_at	0.95414	RIKEN cDNA 2600001M11 gene	2600001M11Rik
1429127_at	0.92296	neurotrophin receptor associated death domain	Nradd
1429128_x_at	1.14779	"nuclear factor of kappa light polypeptide gene enhancer in B-cells 2, p49/p100"	Nfkb2
1429129_at	0.73710	leucine rich repeat and coiled-coil domain containing 1	Lrrcc1
1429130_at	0.98709	tetratricopeptide repeat domain 25	Ttc25
1429131_at	0.89897	ubiquitin-conjugating enzyme E2 variant 2	Ube2v2
1429132_at	0.85974	ubiquitin-conjugating enzyme E2 variant 2	Ube2v2
1429133_at	0.95232	RIKEN cDNA 4930519N16 gene	4930519N16Rik
1429134_at	1.05249	human immunodeficiency virus type I enhancer binding protein 3	Hivep3
1429135_at	0.89527	RIKEN cDNA 1110059M19 gene	1110059M19Rik
1429136_at	0.90220	RIKEN cDNA 1700052K11 gene	1700052K11Rik
1429137_at	0.97519	RIKEN cDNA 2810422O20 gene	2810422O20Rik
1429138_at	0.78434	neuronal PAS domain protein 3	Npas3
1429139_at	0.82850	OTU domain containing 7B	Otud7b
1429140_at	1.07126	RIKEN cDNA 9830002I17 gene	9830002I17Rik
1429141_at	1.05314	neuralized-like 2 (Drosophila)	Neurl2
1429142_at	0.96732	RIKEN cDNA 1700012A03 gene	1700012A03Rik
1429143_at	1.00473	tektin 3	Tekt3
1429144_at	0.87424	preimplantation protein 4	Prei4
1429145_at	1.03156	NHL repeat containing 2	Nhlrc2
1429146_at	1.06034	RIKEN cDNA 6620401M08 gene	6620401M08Rik
1429147_at	0.87630	growth arrest specific 2	Gas2
1429148_at	1.03485	nuclear factor I/C	Nfic
1429149_at	0.99564	GINS complex subunit 3 (Psf3 homolog)	Gins3
1429150_at	1.01497	coiled-coil domain containing 77	Ccdc77
1429151_at	0.94544	WD repeat domain 68	Wdr68
1429152_at	0.91688	zinc finger with KRAB and SCAN domains 1	Zkscan1
1429153_at	1.15128	RIKEN cDNA 6530406A20 gene	6530406A20Rik
1429154_at	0.98319	"solute carrier family 35, member F2"	Slc35f2
1429155_at	0.92246	RIKEN cDNA 4933411K20 gene	4933411K20Rik
1429156_at	0.93264	RIKEN cDNA 2610036L11 gene	2610036L11Rik
1429157_at	0.96098	RIKEN cDNA 4930507C10 gene	4930507C10Rik
1429158_at	1.04642	F-box protein 28	Fbxo28
1429159_at	0.83536	RIKEN cDNA 4631408O11 gene	4631408O11Rik
1429160_at	0.96993	RIKEN cDNA 2810012L14 gene	2810012L14Rik
1429161_at	1.03301	ARV1 homolog (yeast)	Arv1
1429162_at	0.97396	RIKEN cDNA 1500015A07 gene	1500015A07Rik

1429163_at 1.04264 dachsous 1 (Drosophila) Dchs1
1429164_at 0.99718 "protease, serine, 36" Prss36
1429165_at 1.10487 RIKEN cDNA 3110001I22 gene 3110001I22Rik
1429166_s_at 0.58016 calmin Clmn
1429167_at 0.99991 coiled-coil domain containing 112 Ccdc112
1429168_at 0.98864 BTB (POZ) domain containing 4 Btbd4
1429169_at 1.09392 RNA binding motif protein 3 Rbm3
1429170_a_at 1.09795 metal response element binding transcription factor 1 Mtf1
1429171_a_at 1.05434 RIKEN cDNA 5730507H05 gene 5730507H05Rik
1429172_a_at 1.01667 RIKEN cDNA 5730507H05 gene 5730507H05Rik
1429173_at 1.12015 deoxyribonuclease 1-like 1 Dnase111
1429174_at 0.92508 WD repeat domain 34 Wdr34
1429175_at 0.93400 RIKEN cDNA 2810417M05 gene 2810417M05Rik
1429176_at 0.96667 leucine rich repeat and sterile alpha motif containing 1 Lrsam1
1429177_x_at 1.02164 SRY-box containing gene 17 Sox17
1429178_at 0.87474 odd Oz/ten-m homolog 3 (Drosophila) Odz3
1429179_at 1.11171 collectin sub-family member 11 Colec11
1429180_at 1.20721 guanosine monophosphate reductase 2 Gmpr2
1429181_at 0.93431 RIKEN cDNA 1700009P17 gene 1700009P17Rik
1429182_at 1.02672 RIKEN cDNA 5430400D12 gene 5430400D12Rik
1429183_at 1.07159 plakophilin 2 Pkp2
1429184_at 0.90506 "GTPase, very large interferon inducible 1" Gvin1
1429185_at 0.89707 RIKEN cDNA 4631416L12 gene 4631416L12Rik
1429186_a_at 0.84066 cytidine and dCMP deaminase domain containing 1 Cdadc1
1429187_at 0.91706 transmembrane emp24 protein transport domain containing 7 Tmed7
1429188_at 0.96892 "COX11 homolog, cytochrome c oxidase assembly protein (yeast)" Cox11
1429189_at 0.85919 RIKEN cDNA 1110007C02 gene 1110007C02Rik
1429190_at 0.99199 arylsulfatase B Arsb
1429191_at 1.03239 DEAH (Asp-Glu-Ala-His) box polypeptide 33 Dhx33
1429192_at 0.90232 Sloan-Kettering viral oncogene homolog Ski
1429193_at 0.99669 ankyrin repeat and IBR domain containing 1 Ankib1
1429194_at 0.94931 tigger transposable element derived 2 Tigd2
1429195_at 1.01291 --- ---
1429196_at 0.70052 RAB GTPase activating protein 1-like Rabgap11
1429197_s_at 0.90976 RAB GTPase activating protein 1-like Rabgap11
1429198_at 0.95229 RIKEN cDNA 1810030O07 gene 1810030O07Rik
1429199_s_at 1.02099 serum amyloid A-like 1 Saal1
1429200_at 0.97340 serum amyloid A-like 1 Saal1
1429201_at 0.93343 cylindromatosis (turban tumor syndrome) Cyld
1429202_at 1.00364 RIKEN cDNA 2610019N06 gene 2610019N06Rik
1429203_at 1.12733 RIKEN cDNA 2410076I21 gene 2410076I21Rik
1429204_at 0.83708 calcium/calmodulin-dependent protein kinase II inhibitor 2 Camk2n2
1429205_at 0.84732 myeloid/lymphoid or mixed lineage-leukemia translocation to 3 homolog

(Drosophila)	Mllt3		
1429206_at	0.97985 Rho-related BTB domain containing 1	Rhobtb1	
1429207_at	1.24791 RIKEN cDNA 5730408K05 gene	5730408K05Rik	
1429208_at	0.97740 RIKEN cDNA 5730408K05 gene	5730408K05Rik	
1429209_at	0.98270 "procollagen, type XXIII, alpha 1"	Col23a1	
1429210_at	0.98636 "procollagen, type XXIII, alpha 1"	Col23a1	
1429211_at	0.89970 RIKEN cDNA 2900078E11 gene	2900078E11Rik	
1429212_a_at	0.87072 leucine rich repeat containing 51	Lrrc51	
1429213_at	1.09284 RIKEN cDNA 2310030N02 gene	2310030N02Rik	
1429214_at	1.00182 ADAMTS-like 2	Adamtsl2	
1429215_at	1.07869 RIKEN cDNA 2310058N22 gene	2310058N22Rik	
1429216_at	1.11682 progesterin and adipoQ receptor family member III	Paqr3	
1429217_at	0.98046 zinc finger protein 655	Zfp655	
1429218_at	0.89219 de-etiolated homolog 1 (Arabidopsis)	Det1	
1429219_at	0.91230 RIKEN cDNA 1200009F10 gene	1200009F10Rik	
1429220_at	0.83777 WD40 repeat domain 85	Wdr85	
1429221_at	1.10704 WD40 repeat domain 85	Wdr85	
1429222_at	1.20476 --- ---		
1429223_a_at	1.00529 hemochromatosis type 2 (juvenile) (human homolog)	Hfe2	
1429224_at	1.03136 paraneoplastic antigen MA1	Pnma1	
1429225_at	0.97122 RIKEN cDNA 6330417K15 gene	6330417K15Rik	
1429226_at	0.96421 RIKEN cDNA 2310014F06 gene	2310014F06Rik	
1429227_x_at	1.06093 nucleosome assembly protein 1-like 1	Nap111	
1429228_at	0.75352 RIKEN cDNA 4930534B04 gene	4930534B04Rik	
1429229_s_at	0.86821 RIKEN cDNA 4930534B04 gene	4930534B04Rik	
1429230_at	1.02862 kallikrein 5	Klk5	
1429231_at	1.14808 --- ---		
1429232_at	0.88894 RIKEN cDNA 2610528B01 gene	2610528B01Rik	
1429233_at	1.02787 septin 11	11-Sep	
1429234_s_at	0.99755 septin 11	11-Sep	
1429235_at	1.02524 UDP-N-acetyl-alpha-D-galactosamine:polypeptide N-acetylgalactosaminyltransferase-like 2	Galntl2	
1429236_at	1.02398 UDP-N-acetyl-alpha-D-galactosamine:polypeptide N-acetylgalactosaminyltransferase-like 2	Galntl2	
1429237_at	1.10416 syntaxin 16	Stx16	
1429238_a_at	0.95934 2-oxoglutarate and iron-dependent oxygenase domain containing 2	Ogfod2	
1429239_a_at	1.34894 StAR-related lipid transfer (START) domain containing 4	Stard4	
1429240_at	1.32412 StAR-related lipid transfer (START) domain containing 4	Stard4	
1429241_at	1.00129 NIPA-like domain containing 3	Npal3	
1429242_at	0.93955 RIKEN cDNA 1110054O05 gene	1110054O05Rik	
1429243_at	1.06776 RIKEN cDNA 1110054O05 gene	1110054O05Rik	
1429244_at	1.02006 RIKEN cDNA 2610524H06 gene	2610524H06Rik	
1429245_at	0.98608 RIKEN cDNA 2510022D24 gene	2510022D24Rik	

1429246_a_at	0.82816	annexin A6	Anxa6	
1429247_at	1.01747	annexin A6	Anxa6	
1429248_at	1.05676	RIKEN cDNA 2810002N01	gene	2810002N01Rik
1429249_at	1.04293	RIKEN cDNA 4833424O15	gene	4833424O15Rik
1429250_at	0.83403	dynein cytoplasmic 2 heavy chain 1	Dync2h1	
1429251_at	0.83307	"PR domain containing 2, with ZNF domain"	Prdm2	
1429252_at	1.14562	RIKEN cDNA 0610010K14	gene	0610010K14Rik
1429253_at	0.96084	"zinc finger, MYM-type 4"	Zmym4	
1429254_at	0.96539	aquaporin 11	Aqp11	
1429255_at	0.95791	RIKEN cDNA 2010007H12	gene	2010007H12Rik
1429256_at	0.95831	"GTL2, imprinted maternally expressed untranslated mRNA"		Gtl2
1429257_at	0.87389	"GTL2, imprinted maternally expressed untranslated mRNA"		Gtl2
1429258_at	1.06983	RIKEN cDNA 1110025D03	gene	1110025D03Rik
1429259_a_at	1.02831	RIKEN cDNA 1810014B01	gene	1810014B01Rik
1429260_at	1.03353	RIKEN cDNA 1810014B01	gene	1810014B01Rik
1429261_at	1.07642	hypothetical protein LOC664968 /// hypothetical protein LOC669621 ///		
		hypothetical protein LOC675341 LOC664968 /// LOC669621 /// LOC675341		
1429262_at	1.06165	Ras association (RalGDS/AF-6) domain family 6	Rassf6	
1429263_at	0.93871	RIKEN cDNA 4933425D22	gene	4933425D22Rik
1429264_at	0.90727	RIKEN cDNA C030044B11	gene	C030044B11Rik
1429265_a_at	0.85832	ring finger protein 130	Rnf130	
1429266_at	1.04255	"tetraspan transmembrane protein, hair cell stereocilia"		Tmhs
1429267_at	0.95624	acyl-CoA thioesterase 11	Acot11	
1429268_at	1.02251	RIKEN cDNA 2610318N02	gene	2610318N02Rik
1429269_at	1.03090	cDNA sequence BC068157	BC068157	
1429270_a_at	1.07499	synaptonemal complex central element protein 2	Syce2	
1429271_at	1.05193	RIKEN cDNA 9130011E15	gene	9130011E15Rik
1429272_a_at	0.90178	"RIKEN cDNA 9130022K13 gene /// RIKEN cDNA 2210421G13 gene ///		
		hypothetical protein LOC193676 /// similar to apolipoprotein L, 3 (predicted) /// similar to		
		apolipoprotein L, 3 (predicted)"	9130022K13Rik /// 2210421G13Rik /// LOC193676 ///	
		LOC666661 /// LOC672691		
1429273_at	1.29194	BMP-binding endothelial regulator	Bmper	
1429274_at	0.96949	RIKEN cDNA 2310010M24	gene	2310010M24Rik
1429275_at	0.72047	RIKEN cDNA 9530059J11	gene	9530059J11Rik
1429276_at	0.99975	lupus brain antigen 1	Lba1	
1429277_at	1.10871	---	---	
1429278_at	1.00732	nucleotide binding protein-like	Nubpl	
1429279_at	1.11641	tryptophanyl tRNA synthetase 2 (mitochondrial)	Wars2	
1429280_at	0.89130	"collagen, type XXII, alpha 1"	Col22a1	
1429281_at	0.86293	RIKEN cDNA 2610008E11	gene	2610008E11Rik
1429282_at	1.00027	"THAP domain containing, apoptosis associated protein 1"		Thap1
1429283_at	1.18319	RIKEN cDNA 1500009M05	gene	1500009M05Rik
1429284_at	0.84640	RIKEN cDNA 8430436F23	gene	8430436F23Rik

1429285_at 0.74595 "serine (or cysteine) peptidase inhibitor, clade A (alpha-1 antitrypsin), member 9" Serpina9

1429286_at 1.00960 RIKEN cDNA 1190003M12 gene 1190003M12Rik

1429287_a_at 0.92735 prolactin Prl

1429288_x_at 1.00110 syntaxin 18 Stx18

1429289_at 1.07305 LETM1 domain containing 1 Letmd1

1429290_at 0.87766 chromobox homolog 6 Cbx6

1429291_at 1.12008 "proteasome (prosome, macropain) 26S subunit, non-ATPase, 1" Psm1

1429292_a_at 0.95969 RIKEN cDNA 2310046K01 gene 2310046K01Rik

1429293_at 1.04856 glypican 2 (cerebroglycan) Gpc2

1429294_at 1.00271 thyroid hormone receptor interactor 13 Trip13

1429295_s_at 1.05717 thyroid hormone receptor interactor 13 Trip13

1429296_at 0.90055 "RAB10, member RAS oncogene family" Rab10

1429297_at 1.05354 "serine (or cysteine) peptidase inhibitor, clade B (ovalbumin), member 12" Serpinb12

1429298_at 1.07479 dimethylarginine dimethylaminohydrolase 1 Ddah1

1429299_at 1.01263 dimethylarginine dimethylaminohydrolase 1 Ddah1

1429300_at 0.96399 ankyrin repeat domain 9 Ankrd9

1429301_at 0.95204 "casein kinase 2, alpha prime polypeptide" Csnk2a2

1429302_at 0.99510 "casein kinase 2, alpha prime polypeptide" Csnk2a2

1429303_at 0.91695 Kruppel-like factor 17 Klf17

1429304_at 1.03851 ankyrin repeat domain 10 Ankrd10

1429305_at 0.93974 ankyrin repeat domain 10 Ankrd10

1429306_at 1.12544 leucine zipper and CTNNBIP1 domain containing Lzic

1429307_s_at 0.96514 leucine zipper and CTNNBIP1 domain containing Lzic

1429308_at 1.06117 PR domain containing 16 Prdm16

1429309_at 1.03017 PR domain containing 16 Prdm16

1429310_at 0.79068 fibronectin leucine rich transmembrane protein 3 Flrt3

1429311_at 1.05817 ubiquitin-conjugating enzyme E2Q (putative) Ube2q

1429312_s_at 0.94799 receptor tyrosine kinase-like orphan receptor 1 Ror1

1429313_at 0.97090 receptor tyrosine kinase-like orphan receptor 1 Ror1

1429314_at 0.86430 synaptotagmin XI Syt11

1429315_at 0.86364 synaptotagmin XI Syt11

1429316_at 0.96508 "RasGEF domain family, member 1A" Rasgef1a

1429317_at 0.98437 glutaminyl-tRNA synthase (glutamine-hydrolyzing)-like 1 Qrs11

1429318_a_at 0.92681 quaking Qk

1429319_at 0.95054 "ras homolog gene family, member H" Rhoh

1429320_at 0.89233 RIKEN cDNA 4921511I16 gene 4921511I16Rik

1429321_at 0.91011 ring finger protein 149 Rnf149

1429322_at 1.00601 Exosome component 6 Exosc6

1429323_at 0.90598 RIKEN cDNA 1700001E04 gene 1700001E04Rik

1429324_at 0.96253 RIKEN cDNA 1700012A16 gene 1700012A16Rik

1429325_at 1.16049 WD repeat domain 51B Wdr51b

1429326_at 0.96044 centromere protein L Cenpl
 1429327_at 1.18123 serologically defined colon cancer antigen 1 Sdccag1
 1429328_at 1.17208 NSFL1 (p97) cofactor (p47) Nsfl1c
 1429329_at 1.10445 "COX10 homolog, cytochrome c oxidase assembly protein, heme A: farnesyltransferase (yeast)" Cox10
 1429330_at 0.98898 "gamma-aminobutyric acid (GABA-A) receptor, subunit alpha 4" Gabra4
 1429331_at 1.23991 RIKEN cDNA 4632427E13 gene 4632427E13Rik
 1429332_at 1.20611 RIKEN cDNA 4632427E13 gene 4632427E13Rik
 1429333_at 1.00437 kinesin family member 2B Kif2b
 1429334_at 1.08805 RIKEN cDNA 2300002O18 gene 2300002O18Rik
 1429335_at 1.02091 "small nuclear RNA activating complex, polypeptide 1" Snapc1
 1429336_at 0.87680 RIKEN cDNA 2610301K12 gene 2610301K12Rik
 1429337_at 1.14595 RIKEN cDNA 2610301K12 gene 2610301K12Rik
 1429338_a_at 1.11304 nucleolar protein 9 Nol9
 1429339_a_at 0.93262 "acyl-Coenzyme A dehydrogenase family, member 10" Acad10
 1429340_at 1.03855 "HRAS-like suppressor family, member 5" Hrasls5
 1429341_at 0.85743 ornithine decarboxylase antizyme 3 Oaz3
 1429342_s_at 0.88317 RIKEN cDNA 2310021H06 gene 2310021H06Rik
 1429343_at 0.96031 RIKEN cDNA 1700012H05 gene 1700012H05Rik
 1429344_at 0.99608 cortactin binding protein 2 Ctnbp2
 1429345_at 1.19746 "DNA segment, Chr 2, ERATO Doi 435, expressed" D2Ertd435e
 1429346_a_at 0.79119 RIKEN cDNA 4933440J22 gene 4933440J22Rik
 1429347_at 1.02424 Bcl2-like 14 (apoptosis facilitator) Bcl2l14
 1429348_at 0.89689 "sema domain, immunoglobulin domain (Ig), short basic domain, secreted, (semaphorin) 3C" Sema3c
 1429349_at 1.14006 high-mobility group box 4 Hmgb4
 1429350_at 0.90627 RIKEN cDNA 1700027M21 gene 1700027M21Rik
 1429351_at 0.86636 kelch-like 24 (Drosophila) Klh24
 1429352_at 0.93941 molybdenum cofactor sulfurase Mocos
 1429353_a_at 0.93249 RIKEN cDNA 4930511M11 gene 4930511M11Rik
 1429354_at 1.03953 thioesterase superfamily member 5 Them5
 1429355_at 1.09099 RIKEN cDNA 3300001K11 gene 3300001K11Rik
 1429356_s_at 0.98134 geranylgeranyl diphosphate synthase 1 Ggps1
 1429357_at 1.12757 RIKEN cDNA 4921533L14 gene 4921533L14Rik
 1429358_at 0.97082 RIKEN cDNA 4921533L14 gene 4921533L14Rik
 1429359_s_at 0.91285 RNA binding protein gene with multiple splicing Rbpms
 1429360_at 0.93943 Kruppel-like factor 3 (basic) Klf3
 1429361_at 0.95344 pro-melanin-concentrating hormone Pmch
 1429362_a_at 0.94353 "splicing factor 3b, subunit 2" Sf3b2
 1429363_at 1.12667 vacuolar protein sorting 37A (yeast) Vps37a
 1429364_at 1.03686 RIKEN cDNA 4930579G24 gene 4930579G24Rik
 1429365_at 0.89828 RIKEN cDNA 1700016M24 gene 1700016M24Rik
 1429366_at 0.89583 leucine rich repeat containing 34 Lrrc34

1429367_at 0.91777 "WD repeat domain, phosphoinositide interacting 2" Wipi2
1429368_at 1.02175 leucine-rich repeats and immunoglobulin-like domains 3 Lrig3
1429369_at 1.05203 transportin 3 Tnp3
1429370_a_at 1.17516 "proteasome (prosome, macropain) 26S subunit, non-ATPase, 11" Psmd11
1429371_at 0.72434 RIKEN cDNA 2810426N06 gene 2810426N06Rik
1429372_at 0.93942 SRY-box containing gene 11 Sox11
1429373_x_at 0.94208 CREB regulated transcription coactivator 2 Crtc2
1429374_at 0.95818 cysteine-rich perinuclear theca 12 Cypt12
1429375_at 1.04221 anaphase promoting complex subunit 10 Anapc10
1429376_s_at 1.03923 anaphase promoting complex subunit 10 Anapc10
1429377_at 1.06781 RIKEN cDNA 2410004A20 gene 2410004A20Rik
1429378_x_at 0.95496 cDNA sequence AY761184 AY761184
1429379_at 0.95720 extra cellular link domain-containing 1 Xlkd1
1429380_at 0.97341 regulator of G-protein signaling 12 Rgs12
1429381_x_at 1.09473 immunoglobulin heavy chain (J558 family) /// similar to immunoglobulin heavy chain variable region /// similar to immunoglobulin mu-chain /// similar to anti-poly(dC) monoclonal antibody heavy chain /// similar to Ig heavy chain V region 23 precursor /// similar to Ig heavy chain V region 102 precursor /// similar to Ig heavy chain V region VH558 A1/A4 precursor Igh-VJ558 /// LOC238447 /// LOC544903 /// LOC544907 /// LOC633774 /// LOC676478 /// LOC677563
1429382_at 0.93147 translocase of outer mitochondrial membrane 40 homolog-like (yeast) Tomm40l
1429383_at 0.85708 "casein kinase 1, gamma 3" Csnk1g3
1429384_at 0.97486 "casein kinase 1, gamma 3" Csnk1g3
1429385_at 0.96003 WD repeat domain 68 Wdr68
1429386_at 0.90200 WD repeat domain 68 Wdr68
1429387_at 1.00223 GRB2-related adaptor protein Grap
1429388_at 0.90407 Nanog homeobox Nanog
1429389_at 0.90070 SET domain and mariner transposase fusion gene Setmar
1429390_at 0.66487 acid phosphatase-like 2 Acpl2
1429391_at 1.07305 acid phosphatase-like 2 Acpl2
1429392_at 1.06812 WD repeat domain 40A Wdr40a
1429393_at 0.93069 WD repeat domain 40A Wdr40a
1429394_at 1.11625 RIKEN cDNA A130010J15 gene A130010J15Rik
1429395_at 0.90327 "glutathione S-transferase, C-terminal domain containing" Gstcd
1429396_at 0.72767 autophagy related 16 like 2 (S. cerevisiae) Atg16l2
1429397_a_at 0.90786 RIKEN cDNA 4921509E07 gene 4921509E07Rik
1429398_at 1.10027 RIKEN cDNA 5730446D14 gene 5730446D14Rik
1429399_at 0.83565 ring finger protein 125 Rnf125
1429400_at 1.08697 chloride channel 5 Clcn5
1429401_at 0.88900 storkhead box 2 Stox2
1429402_at 1.03061 glycosyltransferase 8 domain containing 2 Glt8d2
1429403_x_at 1.12574 glycosyltransferase 8 domain containing 2 Glt8d2
1429404_at 1.13257 RIKEN cDNA 2010317E24 gene 2010317E24Rik

1429405_at	0.99620	RIKEN cDNA 2010317E24 gene	2010317E24Rik
1429406_at	0.99414	"solute carrier organic anion transporter family, member 6c1"	Slco6c1
1429407_at	0.96423	peroxisomal biogenesis factor 11c	Pex11c
1429408_at	0.96291	"TBC1 domain family, member 21"	Tbc1d21
1429409_at	1.05931	"NACHT, leucine rich repeat and PYD containing 14"	Nalp14
1429410_at	1.03368	enhancer of yellow 2 homolog (Drosophila)	Eny2
1429411_a_at	0.99597	enhancer of yellow 2 homolog (Drosophila)	Eny2
1429412_at	1.14402	enhancer of yellow 2 homolog (Drosophila)	Eny2
1429413_at	0.89923	carboxypeptidase M	Cpm
1429414_at	0.91712	RIKEN cDNA 4933400E14 gene	4933400E14Rik
1429415_at	0.93734	protein kinase C binding protein 1	Prkcbp1
1429416_at	1.05512	tetratricopeptide repeat domain 9B	Ttc9b
1429417_at	0.90882	RIKEN cDNA 4833446K15 gene	4833446K15Rik
1429418_at	1.12664	CDC14 cell division cycle 14 homolog B (S. cerevisiae)	Cdc14b
1429419_at	1.02036	RIKEN cDNA 2310007A19 gene	2310007A19Rik
1429420_at	0.87081	RIKEN cDNA 4931414P19 gene	4931414P19Rik
1429421_at	0.90380	RIKEN cDNA 2610203E10 gene	2610203E10Rik
1429422_at	0.94297	RIKEN cDNA 4933412E12 gene	4933412E12Rik
1429423_at	1.23237	RIKEN cDNA 4930518I15 gene	4930518I15Rik
1429424_at	0.80657	cystatin 13	Cst13
1429425_at	0.88933	ring finger protein 139	Rnf139
1429426_at	0.94613	ring finger protein 139	Rnf139
1429427_s_at	0.73368	"transcription factor 7-like 2, T-cell specific, HMG-box"	Tcf7l2
1429428_at	0.82665	"transcription factor 7-like 2, T-cell specific, HMG-box"	Tcf7l2
1429429_s_at	0.98206	protein-L-isoaspartate (D-aspartate) O-methyltransferase domain containing 1	Pcmd1
1429430_at	0.79991	---	---
1429431_at	0.97932	"zinc finger protein, subfamily 1A, 5"	Zfpn1a5
1429432_at	0.92998	BAT2 domain containing 1	Bat2d
1429433_at	1.20691	BAT2 domain containing 1	Bat2d
1429434_at	0.94155	"phosphatidylinositol 3-kinase, catalytic, alpha polypeptide"	Pik3ca
1429435_x_at	0.99407	"phosphatidylinositol 3-kinase, catalytic, alpha polypeptide"	Pik3ca
1429436_at	0.98700	PRP40 pre-mRNA processing factor 40 homolog A (yeast)	Prpf40a
1429437_at	0.99059	PRP40 pre-mRNA processing factor 40 homolog A (yeast)	Prpf40a
1429438_at	0.81071	Bcl6 interacting corepressor	Bcor
1429439_at	0.99434	"excision repair cross-complementing rodent repair deficiency, complementation group 8"	Ercc8
1429440_at	1.11436	RIKEN cDNA 1810041L15 gene	1810041L15Rik
1429441_at	1.03002	F-box protein 30	Fbxo30
1429442_at	0.97353	RNA pseudouridylate synthase domain containing 2	Rpsud2
1429443_at	0.86537	copine IV	Cpne4
1429444_at	0.67076	"RAS-like, family 11, member A"	Rasl11a
1429445_at	0.89472	RIKEN cDNA 4921510J17 gene	4921510J17Rik

1429446_at 1.03401 serologically defined colon cancer antigen 1 Sdccag1
1429447_at 0.98312 Ellis van Creveld syndrome 2 homolog (human) Evc2
1429448_s_at 0.89663 CXXC finger 6 Cxxc6
1429449_at 1.21272 sterile alpha motif domain containing 4 Samd4
1429450_at 1.00348 RIKEN cDNA 1700026J04 gene 1700026J04Rik
1429451_at 0.93699 RIKEN cDNA 2610301B20 gene 2610301B20Rik
1429452_x_at 0.91655 RIKEN cDNA 4933439C20 gene 4933439C20Rik
1429453_a_at 1.14475 mitochondrial ribosomal protein L55 Mrpl55
1429454_at 1.04305 GTPase activating protein and VPS9 domains 1 Gapvd1
1429455_at 1.07988 GTPase activating protein and VPS9 domains 1 Gapvd1
1429456_a_at 1.01361 polymerase (RNA) III (DNA directed) polypeptide E Polr3e
1429457_at 0.95278 RIKEN cDNA 2310020A21 gene 2310020A21Rik
1429458_at 1.01391 "RIKEN cDNA 2410127L17 gene /// similar to RIKEN cDNA 2410127L17 ///
similar to CG11596-PA, isoform A /// similar to CG11596-PA, isoform A /// similar to RIKEN cDNA
2410127L17 gene" 2410127L17Rik /// LOC626107 /// LOC666456 /// LOC677519 /// LOC677553
1429459_at 1.10995 "sema domain, immunoglobulin domain (Ig), short basic domain, secreted,
(semaphorin) 3D" Sema3d
1429460_at 1.03147 G protein-coupled receptor 115 Gpr115
1429461_at 1.04684 integrator complex subunit 2 Ints2
1429462_at 1.08313 "solute carrier family 25, member 32" Slc25a32
1429463_at 0.88136 "protein kinase, AMP-activated, alpha 2 catalytic subunit" Prkaa2
1429464_at 0.89741 "protein kinase, AMP-activated, alpha 2 catalytic subunit" Prkaa2
1429465_at 1.11540 RIKEN cDNA 1700010M22 gene 1700010M22Rik
1429466_s_at 0.87056 anterior pharynx defective 1c homolog (C. elegans) /// anterior pharynx
defective 1b homolog (C. elegans) Aph1c /// Aph1b
1429467_s_at 0.97897 "solute carrier family 26, member 3" Slc26a3
1429468_at 1.02169 RIKEN cDNA 1110018F16 gene 1110018F16Rik
1429469_at 1.27317 RIKEN cDNA A930013F10 gene A930013F10Rik
1429470_at 0.94452 speedy homolog A (Drosophila) Spdya
1429471_at 0.90787 RIKEN cDNA 1110017D15 gene 1110017D15Rik
1429472_at 1.04807 RIKEN cDNA 4921525O09 gene 4921525O09Rik
1429473_at 0.91857 RIKEN cDNA 1200003I07 gene 1200003I07Rik
1429474_at 0.85173 "zinc binding alcohol dehydrogenase, domain containing 1" Zadh1
1429475_at 0.99134 RIKEN cDNA 2810457I06 gene 2810457I06Rik
1429476_s_at 0.98918 "DnaJ (Hsp40) homolog, subfamily A, member 2" Dnaja2
1429477_at 1.55783 "DNA segment, Chr 15, ERATO Doi 785, expressed" D15Ertd785e
1429478_at 1.60827 RIKEN cDNA 6720463M24 gene 6720463M24Rik
1429479_at 0.95052 RIKEN cDNA 6430590I03 gene 6430590I03Rik
1429480_at 1.04091 RIKEN cDNA 1700011N24 gene 1700011N24Rik
1429481_at 0.97997 RIKEN cDNA 4833426I10 gene 4833426I10Rik
1429482_at 0.95830 RIKEN cDNA 1700003H04 gene 1700003H04Rik
1429483_at 1.04845 nuclear domain 10 protein 52 Ndp52
1429484_at 1.03338 RIKEN cDNA 1110002L01 gene 1110002L01Rik

1429485_a_at 1.14810 "UTP11-like, U3 small nucleolar ribonucleoprotein, (yeast)" Utp11
1429486_at 0.69089 "6-phosphofructo-2-kinase/fructose-2,6-biphosphatase 2" Pfkfb2
1429487_at 1.01828 "protein phosphatase 1, regulatory (inhibitor) subunit 12A" Ppp1r12a
1429488_at 1.24900 "zinc finger, DHHC domain containing 21" Zdhhc21
1429489_at 0.95405 "REX1, RNA exonuclease 1 homolog (S. cerevisiae)" Rexo1
1429490_at 1.02746 similar to Telomere-associated protein RIF1 (Rap1-interacting factor 1 homolog) (mRif1) LOC671598
1429491_s_at 1.04563 Rap1 interacting factor 1 homolog (yeast) /// similar to Telomere-associated protein RIF1 (Rap1-interacting factor 1 homolog) (mRif1) Rif1 /// LOC671598
1429492_x_at 0.96040 phosphatidylserine synthase 2 Ptdss2
1429493_at 1.04512 methyltransferase like 4 Mettl4
1429494_at 1.60966 tripartite motif-containing 35 Trim35
1429495_at 1.01124 RIKEN cDNA 1110004B13 gene 1110004B13Rik
1429496_x_at 1.01237 RIKEN cDNA 2300002M23 gene 2300002M23Rik
1429497_s_at 0.97797 sorting nexin 6 Snx6
1429498_at 0.86613 SFT2 domain containing 3 Sft2d3
1429499_at 1.02314 F-box only protein 5 Fbxo5
1429500_at 0.94765 "protein phosphatase 1A, magnesium dependent, alpha isoform" Ppm1a
1429501_s_at 0.89378 "protein phosphatase 1A, magnesium dependent, alpha isoform" Ppm1a
1429502_at 0.99025 "stress 70 protein chaperone, microsomal-associated, human homolog" Stch
1429503_at 0.62694 RIKEN cDNA 2900024C23 gene 2900024C23Rik
1429504_at 0.94547 "RNA-binding region (RNP1, RRM) containing 3" Rnpc3
1429505_at 0.94947 RIKEN cDNA 2310076G13 gene 2310076G13Rik
1429506_at 1.13900 naked cuticle 1 homolog (Drosophila) /// similar to naked cuticle 1 homolog /// similar to naked cuticle 1 homolog Nkd1 /// LOC634379 /// LOC671838
1429507_at 0.98800 similar to naked cuticle 1 homolog LOC671838
1429508_at 0.99616 RIKEN cDNA 2310057M21 gene 2310057M21Rik
1429509_at 0.96058 RIKEN cDNA 2600001B17 gene 2600001B17Rik
1429510_at 0.93904 RIKEN cDNA 2810410L24 gene 2810410L24Rik
1429511_at 1.05454 RIKEN cDNA 4933402E13 gene 4933402E13Rik
1429512_at 0.96020 zona pellucida binding protein 2 Zpbp2
1429513_at 0.84378 RIKEN cDNA 1700019M22 gene 1700019M22Rik
1429514_at 0.80864 phosphatidic acid phosphatase type 2B Ppap2b
1429515_at 1.00089 ubiquitin protein ligase E3 component n-recogin 2 Ubr2
1429516_at 0.92878 ubiquitin protein ligase E3 component n-recogin 2 Ubr2
1429517_at 0.91225 "zinc finger, FYVE domain containing 20" Zfyve20
1429518_at 1.04289 Fas apoptotic inhibitory molecule 2 Faim2
1429519_at 0.82709 fucose-1-phosphate guanylyltransferase Fpgt
1429520_a_at 0.97427 "phytoceramidase, alkaline" Phca
1429521_at 1.12645 "alkB, alkylation repair homolog 8 (E. coli)" Alkbh8
1429522_at 0.93195 ankyrin repeat domain 42 Ankrd42
1429523_a_at 0.99500 "solute carrier family 39 (metal ion transporter), member 5" Slc39a5
1429524_at 1.08688 myosin IF Myo1f

1429525_s_at 0.91523 myosin IF Myo1f
1429526_at 1.55721 bromodomain containing 8 Brd8
1429527_a_at 1.06945 phospholipid scramblase 1 /// hypothetical LOC433328 /// hypothetical protein
LOC677340 Plscr1 /// LOC433328 /// LOC677340
1429528_at 1.04906 RAE1 RNA export 1 homolog (S. pombe) Rae1
1429529_at 0.98394 "tRNA splicing endonuclease 54 homolog (SEN54, S. cerevisiae)" Tsen54
1429530_a_at 1.01944 sphingomyelin phosphodiesterase 4 Smpd4
1429531_at 0.88546 sphingomyelin phosphodiesterase 4 Smpd4
1429532_at 1.01820 microrchidia 2A Morc2a
1429533_at 0.98944 "inner membrane protein, mitochondrial" Immt
1429534_a_at 1.05323 "inner membrane protein, mitochondrial" Immt
1429535_at 1.27946 armadillo repeat containing 8 Armc8
1429536_at 1.07814 RIKEN cDNA 2310040C09 gene 2310040C09Rik
1429537_at 1.01354 RIKEN cDNA 5730406M06 gene 5730406M06Rik
1429538_a_at 1.04766 RIKEN cDNA 5730406M06 gene 5730406M06Rik
1429539_at 0.88414 BCL2-like 13 (apoptosis facilitator) Bcl2l13
1429540_at 1.06804 cornifelin Cnfn
1429541_at 1.00268 RIKEN cDNA 2500002G23 gene 2500002G23Rik
1429542_at 1.08376 RIKEN cDNA 1700007B14 gene 1700007B14Rik
1429543_at 0.80979 RIKEN cDNA 6230424C14 gene 6230424C14Rik
1429544_at 1.13935 RIKEN cDNA 1700029J11 gene 1700029J11Rik
1429545_at 1.15494 "ubiquitin-conjugating enzyme E2I /// RIKEN cDNA F830028O17 gene ///
similar to Chain A, Human Ubiquitin-Conjugating Enzyme Ubc9 /// similar to Ubiquitin-conjugating
enzyme E2 I (Ubiquitin-protein ligase I) (Ubiquitin carrier protein I) (SUMO-1-protein ligase) (SUMO-
1-conjugating enzyme) (Ubiquitin-conjugating enzyme UbcE2A)" Ube2i /// F830028O17Rik ///
LOC546265 /// LOC669417
1429546_at 0.79731 RIKEN cDNA 4930423K06 gene 4930423K06Rik
1429547_at 0.98825 RIKEN cDNA 4930578I06 gene 4930578I06Rik
1429548_at 0.96919 testis expressed gene 22 Tex22
1429549_at 0.93735 "procollagen, type XXVII, alpha 1" Col27a1
1429550_at 0.90488 ectonucleoside triphosphate diphosphohydrolase 8 Entpd8
1429551_at 0.85405 RIKEN cDNA 4930579G22 gene 4930579G22Rik
1429552_at 0.96443 WD repeat domain 16 Wdr16
1429553_at 0.99782 cartilage intermediate layer protein 2 Cilp2
1429554_at 1.01757 serologically defined colon cancer antigen 3 Sdccag3
1429555_at 0.89575 claudin domain containing 1 Cldnd1
1429556_at 1.16086 RIKEN cDNA 2610024B07 gene 2610024B07Rik
1429557_at 1.08971 minichromosome maintenance deficient 8 (S. cerevisiae) Mcm8
1429558_a_at 0.94751 "La ribonucleoprotein domain family, member 7" Larp7
1429559_at 0.97040 "guanine nucleotide binding protein, alpha q polypeptide" Gnaq
1429560_at 0.98384 "zinc finger protein 422, related sequence 1" Zfp422-rs1
1429561_at 1.00950 "BRF2, subunit of RNA polymerase III transcription initiation factor, BRF1-
like" Brf2

1429562_at	1.03244	RIKEN cDNA 5031415C07 gene	5031415C07Rik
1429563_x_at	0.99029	RIKEN cDNA 5031415C07 gene	5031415C07Rik
1429564_at	1.07807	polycomb group ring finger 5	Pcgf5
1429565_s_at	0.73306	late cornified envelope 5A	Lce5a
1429566_a_at	0.94159	homeodomain interacting protein kinase 2	Hipk2
1429567_at	0.87089	RIKEN cDNA 4632411J06 gene	4632411J06Rik
1429568_x_at	1.03695	ubiquitin-conjugating enzyme E2F (putative)	Ube2f
1429569_a_at	0.88243	coiled-coil domain containing 46	Ccdc46
1429570_at	1.29155	mixed lineage kinase domain-like	Mlkl
1429571_a_at	0.99728	sperm acrosome associated 1	Spaca1
1429572_at	1.01171	"NOL1/NOP2/Sun domain family, member 7"	Nsun7
1429573_at	0.96822	DMRT-like family C1a	Dmrta1a
1429574_at	0.93640	chloride intracellular channel 3	Clic3
1429575_at	0.90213	---	---
1429576_at	1.11183	RIKEN cDNA 1700060E18 gene	1700060E18Rik
1429577_at	0.86197	IQ motif containing F4	Iqcf4
1429578_at	0.94571	RIKEN cDNA 1700049K14 gene	1700049K14Rik
1429579_at	0.86934	RIKEN cDNA 6330407I18 gene	6330407I18Rik
1429580_x_at	1.12494	RIKEN cDNA 5133400G04 gene	5133400G04Rik
1429581_at	1.12030	"acyl-Coenzyme A dehydrogenase family, member 9"	Acad9
1429582_at	0.86675	BTB (POZ) domain containing 14A	Btbd14a
1429583_at	0.98639	RIKEN cDNA 2900053A13 gene	2900053A13Rik
1429584_at	0.86490	myoneurin	Mynn
1429585_s_at	0.79914	myoneurin	Mynn
1429586_at	1.11533	RIKEN cDNA 4930558N01 gene	4930558N01Rik
1429587_at	1.06642	RNA binding motif protein 34	Rbm34
1429588_at	0.98433	RIKEN cDNA 2810474O19 gene	2810474O19Rik
1429589_at	1.01130	RIKEN cDNA 6330404F12 gene	6330404F12Rik
1429590_at	0.77239	"transforming, acidic coiled-coil containing protein 1"	Tacc1
1429591_at	0.79057	"transforming, acidic coiled-coil containing protein 1"	Tacc1
1429592_at	0.94396	lipoma HMGIC fusion partner-like 3	Lhfp13
1429593_at	1.02210	"solute carrier family 38, member 2"	Slc38a2
1429594_at	0.84381	"solute carrier family 38, member 2"	Slc38a2
1429595_at	0.96537	RIKEN cDNA 2700049A03 gene	2700049A03Rik
1429596_at	1.06511	"solute carrier family 7, member 6 opposite strand"	Slc7a6os
1429597_at	1.00971	developmental pluripotency associated 4	Dppa4
1429598_at	0.88428	RIKEN cDNA 2310042D19 gene	2310042D19Rik
1429599_a_at	1.03598	RIKEN cDNA C630010D07 gene	/// similar to methylenetetrahydrofolate dehydrogenase (NADP+ dependent) 2-like
		/// region containing RIKEN cDNA C630010D07 gene;	
		RIKEN cDNA 1110019K23 gene	C630010D07Rik
		/// LOC665563	/// LOC674367
1429600_at	1.05807	RIKEN cDNA C630010D07 gene	/// similar to methylenetetrahydrofolate dehydrogenase (NADP+ dependent) 2-like
		/// region containing RIKEN cDNA C630010D07 gene;	
		RIKEN cDNA 1110019K23 gene	C630010D07Rik
		/// LOC665563	/// LOC674367

1429601_x_at 1.05608 RIKEN cDNA C630010D07 gene /// similar to methylenetetrahydrofolate dehydrogenase (NADP+ dependent) 2-like /// region containing RIKEN cDNA C630010D07 gene; RIKEN cDNA 1110019K23 gene C630010D07Rik /// LOC665563 /// LOC674367

1429602_at 0.89247 D164 sialomucin-like 2 Cd164l2

1429603_at 1.19809 RIKEN cDNA 2610110G12 gene 2610110G12Rik

1429604_at 1.03144 RIKEN cDNA 1810011H11 gene 1810011H11Rik

1429605_at 0.92043 RIKEN cDNA 4930532L20 gene 4930532L20Rik

1429606_at 0.93618 RIKEN cDNA 4930527B16 gene 4930527B16Rik

1429607_at 0.92052 "trafficking protein, kinesin binding 2" Trak2

1429608_at 1.00528 alcohol dehydrogenase 6A (class V) Adh6a

1429609_at 1.07103 RIKEN cDNA 1700001D09 gene 1700001D09Rik

1429610_a_at 1.04696 zinc finger protein 511 Zfp511

1429611_at 0.99891 RIKEN cDNA 1700034E13 gene 1700034E13Rik

1429612_at 1.05417 echinoderm microtubule associated protein like 4 Eml4

1429613_at 0.94677 RIKEN cDNA 0710001B24 gene 0710001B24Rik

1429614_at 1.08629 PRP18 pre-mRNA processing factor 18 homolog (yeast) Prpf18

1429615_at 0.96702 zinc finger protein 91 Zfp91

1429616_at 1.08805 zinc finger protein 91 Zfp91

1429617_at 1.10968 cylindromatosis (turban tumor syndrome) Cyld

1429618_at 1.22584 cylindromatosis (turban tumor syndrome) Cyld

1429619_a_at 1.02566 RIKEN cDNA 8430406I07 gene 8430406I07Rik

1429620_at 0.97818 RIKEN cDNA 8430406I07 gene 8430406I07Rik

1429621_at 0.69383 cullin-associated and neddylation-dissociated 2 (putative) Cand2

1429622_at 0.88066 cullin-associated and neddylation-dissociated 2 (putative) Cand2

1429623_at 0.91904 zinc finger protein 644 Zfp644

1429624_at 1.22684 "SAFB-like, transcription modulator" Sltm

1429625_at 0.77265 RIKEN cDNA 2900054C01 gene 2900054C01Rik

1429626_at 1.05083 surfactant associated protein A1 Sftpa1

1429627_at 1.03952 peptidylprolyl isomerase (cyclophilin)-like 3 Ppil3

1429628_at 1.12438 RIKEN cDNA 6330407J23 gene 6330407J23Rik

1429629_at 0.90653 enhancer of zeste homolog 1 (Drosophila) Ezh1

1429630_at 1.06578 enhancer of zeste homolog 1 (Drosophila) Ezh1

1429631_at 0.94387 "sirtuin 6 (silent mating type information regulation 2, homolog) 6 (S. cerevisiae)" Sirt6

1429632_at 1.11087 trafficking protein particle complex 4 Trappc4

1429633_at 1.22685 ligand dependent nuclear receptor corepressor Lcor

1429634_at 0.82259 zinc finger protein 580 Zfp580

1429635_at 1.01622 coiled-coil domain containing 89 Ccdc89

1429636_at 0.87902 RIKEN cDNA 1700010D01 gene /// similar to hypothetical protein LOC76386 1700010D01Rik /// MGC118250

1429637_at 1.03916 RIKEN cDNA 2210419I08 gene 2210419I08Rik

1429638_at 0.96916 Werner helicase interacting protein 1 Wrnip1

1429639_at 1.19179 preimplantation protein 4 Prei4

1429640_at 1.02595 RIKEN cDNA 1700025F22 gene 1700025F22Rik
1429641_x_at 1.00329 RIKEN cDNA 2310007F04 gene 2310007F04Rik
1429642_at 0.75939 "AN1, ubiquitin-like, homolog (Xenopus laevis)" Anubl1
1429643_a_at 0.97527 phosphodiesterase 1C Pde1c
1429644_at 0.96003 RIKEN cDNA 1700011I03 gene /// hypothetical gene supported by AK005868;
BC048576 1700011I03Rik /// LOC433188
1429645_at 0.98251 RIKEN cDNA 1700017G19 gene 1700017G19Rik
1429646_at 0.92849 RIKEN cDNA 1700112C13 gene 1700112C13Rik
1429647_at 0.93010 RIKEN cDNA 1700027L20 gene 1700027L20Rik
1429648_at 1.05449 "solute carrier family 35 (UDP-N-acetylglucosamine (UDP-GlcNAc)
transporter), member 3" Slc35a3
1429649_at 0.89877 "solute carrier family 35 (UDP-N-acetylglucosamine (UDP-GlcNAc)
transporter), member 3" Slc35a3
1429650_at 0.94455 serine/threonine kinase 40 Stk40
1429651_at 1.10021 phosphatase and actin regulator 3 Phactr3
1429652_at 0.79164 RIKEN cDNA 1190002C06 gene 1190002C06Rik
1429653_at 0.85621 genetic suppressor element 1 Gse1
1429654_at 0.95685 developmental pluripotency associated 2 Dppa2
1429655_at 1.10398 NudC domain containing 1 Nudcd1
1429656_at 1.10474 Rho-related BTB domain containing 1 Rhobtb1
1429657_at 1.31079 "zinc finger, AN1-type domain 5" Zfand5
1429658_a_at 1.10946 structural maintenance of chromosomes 2 Smc2
1429659_at 0.82716 structural maintenance of chromosomes 2 Smc2
1429660_s_at 0.98570 structural maintenance of chromosomes 2 Smc2
1429661_at 1.05393 Rho-related BTB domain containing 3 Rhobtb3
1429662_at 0.95110 spermatogenesis associated 1 Spata1
1429663_at 0.71455 katanin p60 subunit A-like 2 Katnal2
1429664_at 0.94191 cyclin-dependent kinase-like 1 (CDC2-related kinase) Cdkl1
1429665_at 0.93165 RIKEN cDNA 6230416J20 gene 6230416J20Rik
1429666_at 0.99227 potassium channel tetramerisation domain containing 16 Kctd16
1429667_at 0.90686 "POU domain, class 4, transcription factor 1" Pou4f1
1429668_at 1.02371 "POU domain, class 4, transcription factor 1" Pou4f1
1429669_at 0.86790 RIKEN cDNA 4930488L21 gene 4930488L21Rik
1429670_a_at 0.94000 leucine-rich repeats and IQ motif containing 2 Lrriq2
1429671_at 0.74019 RIKEN cDNA 2410018M08 gene 2410018M08Rik
1429672_at 1.01873 RIKEN cDNA 5830407E08 gene 5830407E08Rik
1429673_at 1.05244 RIKEN cDNA 5830407E08 gene 5830407E08Rik
1429674_at 0.98412 RIKEN cDNA 1700113H08 gene 1700113H08Rik
1429675_at 1.13081 RIKEN cDNA 1700023A16 gene 1700023A16Rik
1429676_at 0.96484 "Adult male stomach cDNA, RIKEN full-length enriched library,
clone:2210408O09 product:unclassifiable, full insert sequence" ---
1429677_at 1.05892 RIKEN cDNA 1700056E22 gene 1700056E22Rik
1429678_at 1.07032 RIKEN cDNA 5730508B09 gene 5730508B09Rik

1429679_at 1.00533 leucine rich repeat containing 17 Lrrc17
1429680_at 1.23699 transformer 2 alpha homolog (Drosophila) Tra2a
1429681_a_at 1.12465 "glycoprotein, synaptic 2" Gpsn2
1429682_at 0.76820 RIKEN cDNA 4930431B09 gene 4930431B09Rik
1429683_at 0.95021 RIKEN cDNA 5830472M02 gene 5830472M02Rik
1429684_at 1.05145 RIKEN cDNA 5830472M02 gene 5830472M02Rik
1429685_at 0.99493 RIKEN cDNA C030002O17 gene C030002O17Rik
1429686_at 1.15397 polymerase (RNA) III (DNA directed) polypeptide F Polr3f
1429687_at 0.96685 RIKEN cDNA 4930503B16 gene 4930503B16Rik
1429688_at 0.99534 aryl hydrocarbon receptor nuclear translocator-like 2 Arntl2
1429689_at 1.00142 RIKEN cDNA 4932433N03 gene 4932433N03Rik
1429690_at 0.88583 RIKEN cDNA 1300003B13 gene 1300003B13Rik
1429691_at 0.87009 RIKEN cDNA 5430405N12 gene 5430405N12Rik
1429692_s_at 1.01057 GTP cyclohydrolase 1 Gch1
1429693_at 1.23305 disabled homolog 2 (Drosophila) Dab2
1429694_at 1.13320 RIKEN cDNA 4930402H24 gene 4930402H24Rik
1429695_at 0.98297 izumo sperm-egg fusion 1 Izumo1
1429696_at 0.92243 G protein-coupled receptor 123 Gpr123
1429697_at 0.87981 RIKEN cDNA 1700023D19 gene 1700023D19Rik
1429698_at 1.10452 mitochondrial transcription termination factor Mterf
1429699_at 1.01939 "3-oxoacyl-ACP synthase, mitochondrial" Oxsm
1429700_at 0.94040 RIKEN cDNA 3110040M04 gene /// similar to U2-associated SR140 protein ///
hypothetical protein LOC668055 /// similar to U2-associated SR140 protein /// similar to U2-associated
SR140 protein /// similar to U2-associated SR140 protein /// similar to U2-associated SR140 protein ///
similar to U2-associated SR140 protein /// similar to U2-associated SR140 protein
3110040M04Rik /// LOC668050 /// LOC668055 /// LOC671025 /// LOC671035 /// LOC671043 ///
LOC671049 /// LOC676755 /// LOC676758
1429701_at 1.01752 RIKEN cDNA 2410003J06 gene 2410003J06Rik
1429702_at 2.18015 RIKEN cDNA 2900072G11 gene 2900072G11Rik
1429703_at 2.09242 RIKEN cDNA 2900072G11 gene 2900072G11Rik
1429704_at 1.00860 Lrp2 binding protein Lrp2bp
1429705_at 0.99587 sperm acrosome associated 4 Spaca4
1429706_at 0.96311 IQ motif containing F5 Iqcf5
1429707_at 1.16386 "phospholipase A2, activating protein" Plaa
1429708_at 0.94797 NADH dehydrogenase (ubiquinone) 1 alpha subcomplex 11 Ndufa11
1429709_at 0.99732 peptidase (mitochondrial processing) beta Pmpcb
1429710_at 1.08540 phosphoserine/threonine/tyrosine interaction protein Styx
1429711_at 1.03855 phosphoserine/threonine/tyrosine interaction protein Styx
1429712_at 1.10707 ethanol induced 1 /// RIKEN cDNA 6230416C02 gene /// similar to zinc finger
protein 97 /// similar to zinc finger protein 97 Etoh1 /// 6230416C02Rik /// LOC667962 ///
LOC668009
1429713_at 0.87333 sulfatase modifying factor 2 Sumf2
1429714_at 0.94993 sulfatase modifying factor 2 Sumf2

1429715_at 1.01688 "protein phosphatase 2 (formerly 2A), regulatory subunit B (PR 52), alpha isoform" Ppp2r2a

1429716_at 1.07503 "zinc finger, CCHC domain containing 4" Zcchc4

1429717_at 1.02169 importin 11 Ipo11

1429718_at 0.93558 "SLIT and NTRK-like family, member 5" Slitrk5

1429719_at 0.86730 forkhead box P4 Foxp4

1429720_at 0.93203 "MAK10 homolog, amino-acid N-acetyltransferase subunit, (*S. cerevisiae*)" Mak10

1429721_s_at 0.91263 RIKEN cDNA 4930579C15 gene 4930579C15Rik

1429722_at 0.88860 zinc finger and BTB domain containing 4 Zbtb4

1429723_at 1.10829 RIKEN cDNA 6330409N04 gene 6330409N04Rik

1429724_at 1.09351 PRP4 pre-mRNA processing factor 4 homolog (yeast) Prpf4

1429725_at 1.20906 AT motif binding factor 1 Atbf1

1429726_at 0.96630 "solute carrier family 16 (monocarboxylic acid transporters), member 9" Slc16a9

1429727_at 1.01765 "solute carrier family 16 (monocarboxylic acid transporters), member 9" Slc16a9

1429728_at 1.10415 RIKEN cDNA 4930429M06Rik 4930429M06Rik

1429729_at 0.86864 synaptotagmin XI Syt11

1429730_at 1.01592 RIKEN cDNA 1810009N02 gene 1810009N02Rik

1429731_at 1.08857 RIKEN cDNA 6530403G13 gene 6530403G13Rik

1429732_at 1.06101 "protein phosphatase 1, regulatory (inhibitor) subunit 1C" Ppp1r1c

1429733_at 0.94842 coiled-coil domain containing 94 Ccdc94

1429734_at 1.08364 RIKEN cDNA 4632434I11 gene 4632434I11Rik

1429735_at 1.14176 RIKEN cDNA 1110003F05 gene 1110003F05Rik

1429736_at 0.90973 RIKEN cDNA 1110003F05 gene 1110003F05Rik

1429737_a_at 1.11856 cardiolipin synthase 1 Crls1

1429738_at 1.02426 myelin transcription factor 1-like Myt11

1429739_a_at 0.81779 zinc finger protein 278 Zfp278

1429740_at 0.91381 tetratricopeptide repeat domain 7 Ttc7

1429741_at 1.05872 "potassium channel, subfamily V, member 1" Kcnv1

1429742_at 1.09395 regulator of chromosome condensation (RCC1) and BTB (POZ) domain containing protein 2 Rcbtb2

1429743_at 1.06221 RIKEN cDNA 6720468P15 gene 6720468P15Rik

1429744_at 0.88577 RIKEN cDNA 4931407G18 gene 4931407G18Rik

1429745_at 0.87141 similar to melanoma ubiquitous mutated protein /// similar to UBE-1C2 protein /// similar to UBE-1C2 protein /// hypothetical protein LOC574404 /// hypothetical protein LOC574405 /// similar to melanoma associated antigen (mutated) 1-like 1 LOC434795 /// LOC434797 /// LOC546325 /// MGC106740 /// LOC574405 /// LOC674060

1429746_at 1.10728 RIKEN cDNA 1700007I06 gene 1700007I06Rik

1429747_at 0.88018 RIKEN cDNA 1700009N14 gene 1700009N14Rik

1429748_at 1.05733 --- ---

1429749_at 0.93939 RIKEN cDNA 9330180L21 gene 9330180L21Rik

1429750_at 1.19041 multiple inositol polyphosphate histidine phosphatase 1 Minpp1

1429751_at 1.00903 "zinc finger, SWIM domain containing 2" Zswim2

1429752_x_at 0.97976 restin-like 2 Rsnl2
1429753_at 0.97964 neurexophilin 4 Nxph4
1429754_a_at 0.92260 RIKEN cDNA 4933406F09 gene 4933406F09Rik
1429755_a_at 1.07527 RIKEN cDNA 4933406F09 gene 4933406F09Rik
1429756_at 1.01533 RIKEN cDNA 4931428F04 gene 4931428F04Rik
1429757_at 0.94895 RIKEN cDNA 1700094C09 gene 1700094C09Rik
1429758_at 1.16284 RIKEN cDNA 1700017B05 gene /// hypothetical protein LOC640563
1700017B05Rik /// LOC640563
1429759_at 0.88950 ribosomal protein S6 kinase polypeptide 6 Rps6ka6
1429760_at 1.07644 ribosomal protein S6 kinase polypeptide 6 Rps6ka6
1429761_at 1.15875 reticulon 1 Rtn1
1429762_a_at 1.03796 Bardet-Biedl syndrome 5 (human) Bbs5
1429763_at 1.12011 cornichon homolog 4 (Drosophila) Cnih4
1429764_at 0.96440 RIKEN cDNA 1500005K14 gene 1500005K14Rik
1429765_at 0.98268 RIKEN cDNA 1500005K14 gene 1500005K14Rik
1429766_at 0.87177 RIKEN cDNA 2810013P06 gene 2810013P06Rik
1429767_at 0.96225 RIKEN cDNA 2810013P06 gene 2810013P06Rik
1429768_at 0.91500 dystrobrevin alpha Dtna
1429769_at 0.89090 "protein geranylgeranyltransferase type I, beta subunit" Pgg1b
1429770_at 1.06988 "protein geranylgeranyltransferase type I, beta subunit" Pgg1b
1429771_at 1.22991 RIKEN cDNA 3110073H01 gene 3110073H01Rik
1429772_at 0.93484 plexin A2 Plxna2
1429773_at 1.04678 RIKEN cDNA 2810451A06 gene 2810451A06Rik
1429774_a_at 0.99994 coiled-coil domain containing 98 Ccdc98
1429775_a_at 1.08229 G protein-coupled receptor 137B /// similar to transmembrane 7 superfamily
member 1 /// similar to transmembrane 7 superfamily member 1 Gpr137b /// LOC664862 ///
LOC673335
1429776_a_at 0.91425 "DnaJ (Hsp40) homolog, subfamily B, member 6" Dnajb6
1429777_at 0.99693 "DnaJ (Hsp40) homolog, subfamily B, member 6" Dnajb6
1429778_at 1.01057 optineurin Optn
1429779_at 1.07320 "eukaryotic translation initiation factor 2C, 4" Eif2c4
1429780_at 1.00985 coiled-coil domain containing 39 Ccdc39
1429781_s_at 1.00695 coiled-coil domain containing 39 Ccdc39
1429782_at 1.11576 "gene rich cluster, C10 gene" Grcc10
1429783_at 1.01381 PDZ and LIM domain 5 Pdlim5
1429784_at 1.99801 RIKEN cDNA C130032J12 gene C130032J12Rik
1429785_at 0.90973 RIKEN cDNA 5830458C19 gene 5830458C19Rik
1429786_a_at 1.02272 ZW10 interactor Zwint
1429787_x_at 1.08461 ZW10 interactor Zwint
1429788_at 1.04512 actin related protein M1 Arpm1
1429789_at 0.94578 actin related protein M1 Arpm1
1429790_at 1.06198 "bruno-like 6, RNA binding protein (Drosophila) /// similar to bruno-like 6,
RNA binding protein" Brunol6 /// LOC669789

1429791_at	0.86717	RIKEN cDNA A930004D18	gene	A930004D18Rik
1429792_at	1.06681	RIKEN cDNA 9530048O09	gene	9530048O09Rik
1429793_at	0.96490	RIKEN cDNA 1600014K23	gene	1600014K23Rik
1429794_a_at	1.01994	"purinergic receptor P2X, ligand-gated ion channel, 1"		P2rx1
1429795_at	0.98146	RIKEN cDNA 1700001L05	gene	1700001L05Rik
1429796_at	0.99176	"kalirin, RhoGEF kinase"		Kalrn
1429797_at	1.32468	RIKEN cDNA 5730596K20	gene	5730596K20Rik
1429798_s_at	1.01212	CKLF-like MARVEL transmembrane domain containing 5		Cmtm5
1429799_at	0.91143	zinc finger protein 606		Zfp606
1429800_at	0.99257	RIKEN cDNA 9130221H12	gene	9130221H12Rik
1429801_at	1.12072	RIKEN cDNA 2310028H24	gene	2310028H24Rik
1429802_at	0.94289	dehydrogenase/reductase (SDR family) member 10		Dhrs10
1429803_at	1.13234	RIKEN cDNA 1700021F07	gene	1700021F07Rik
1429804_at	1.00137	"solute carrier family 22 (organic cation transporter), member 16"		Slc22a16
1429805_at	1.02219	myosin IA /// similar to Myosin Ia (Brush border myosin I) (BBM-I) (BBMI) (Myosin I heavy chain) (MIHC)		Myo1a /// LOC674795
1429806_at	1.26972	Era (G-protein)-like 1 (E. coli)		Eral1
1429807_at	0.95094	RIKEN cDNA 1700011H22	gene	1700011H22Rik
1429808_at	0.89540	RIKEN cDNA 1110020C03	gene	1110020C03Rik
1429809_at	0.87819	transmembrane and tetratricopeptide repeat containing 2		Tmtc2
1429810_at	1.48811	RIKEN cDNA 4921505C17	gene	4921505C17Rik
1429811_at	1.02516	RIKEN cDNA 4933424B01	gene	4933424B01Rik
1429812_at	1.16452	RIKEN cDNA 2610002D18	gene	2610002D18Rik
1429813_at	1.00703	pantothenate kinase 1		Pank1
1429814_at	1.01140	pantothenate kinase 1		Pank1
1429815_at	1.09775	armadillo repeat containing 3		Armc3
1429816_at	0.95217	armadillo repeat containing 3		Armc3
1429817_at	1.07772	spermatogenesis and oogenesis specific basic helix-loop-helix 2		Sohlh2
1429818_at	0.91529	RIKEN cDNA 2410012M07	gene	2410012M07Rik
1429819_at	1.01051	nicotinamide nucleotide adenylyltransferase 1		Nmnat1
1429820_at	1.02814	RUN and TBC1 domain containing 2		Rutbc2
1429821_at	1.15911	RIKEN cDNA 2810046L04	gene	2810046L04Rik
1429822_at	1.75026	RIKEN cDNA 4633401B06	gene	4633401B06Rik
1429823_at	0.90728	RIKEN cDNA 5430420E18	gene	5430420E18Rik
1429824_at	1.01531	RIKEN cDNA 4930550C14	gene	4930550C14Rik
1429825_at	0.88784	RIKEN cDNA 1700030E15	gene	1700030E15Rik
1429826_at	0.95004	prolactin-like protein N		Prlpn
1429827_at	0.97806	RIKEN cDNA 1700010B08	gene	1700010B08Rik
1429828_at	1.05280	RIKEN cDNA 1700024P04	gene	1700024P04Rik
1429829_at	0.94317	RIKEN cDNA 9530003J23	gene	9530003J23Rik
1429830_a_at	0.90688	CD59a antigen		Cd59a
1429831_at	1.10210	phosphoinositide-3-kinase adaptor protein 1		Pik3ap1
1429832_at	1.21608	peptidyl prolyl isomerase H /// similar to peptidyl prolyl isomerase H ///		

hypothetical protein LOC624822 /// similar to Peptidyl-prolyl cis-trans isomerase H (PPIase H) (Rotamase H) /// similar to peptidylprolyl isomerase H /// similar to Peptidyl-prolyl cis-trans isomerase H (PPIase H) (Rotamase H) /// similar to peptidylprolyl isomerase H /// similar to peptidylprolyl isomerase H /// similar to peptidylprolyl isomerase H /// similar to Peptidyl-prolyl cis-trans isomerase H (PPIase H) (Rotamase H) /// similar to peptidylprolyl isomerase H Ppih /// LOC433064 /// LOC624822 /// LOC629952 /// LOC665989 /// LOC666411 /// LOC667598 /// LOC669441 /// LOC670964 /// LOC671009 /// LOC676493

1429833_at 1.12194 "lymphocyte antigen 6 complex, locus G6E" Ly6g6e
1429834_a_at 0.97935 RIKEN cDNA 1110014N23 gene 1110014N23Rik
1429835_at 0.92615 RIKEN cDNA 2310033E01 gene 2310033E01Rik
1429836_at 0.99403 UDP-glucose ceramide glucosyltransferase-like 2 Ugcgl2
1429837_at 0.92464 RIKEN cDNA 1700016D18 gene 1700016D18Rik
1429838_at 1.02488 sperm associated antigen 16 Spag16
1429839_a_at 0.90100 YY1 associated factor 2 Yaf2
1429840_at 1.00935 RIKEN cDNA 4933439G12 gene 4933439G12Rik
1429841_at 0.99024 multiple EGF-like-domains 10 Megf10
1429842_at 0.92063 "malate dehydrogenase 1B, NAD (soluble)" Mdh1b
1429843_at 1.00830 glycine/arginine rich protein 1 Grp1
1429844_at 1.10758 RIKEN cDNA 2310043J07 gene 2310043J07Rik
1429845_at 0.99961 RIKEN cDNA 1700019L03 gene 1700019L03Rik
1429846_at 1.24726 RIKEN cDNA 9030411K21 gene 9030411K21Rik
1429847_a_at 1.04131 RIKEN cDNA 4833418A01 gene 4833418A01Rik
1429848_at 1.03194 poliovirus receptor Pvr
1429849_at 1.34171 RIKEN cDNA 4632411B12 gene 4632411B12Rik
1429850_x_at 0.89703 "alkB, alkylation repair homolog 4 (E. coli)" Alkbh4
1429851_at 1.03577 RIKEN cDNA 1700003M02 gene 1700003M02Rik
1429852_at 0.94454 coiled-coil domain containing 57 Ccdc57
1429853_at 0.89463 RIKEN cDNA 1700018C11 gene 1700018C11Rik
1429854_at 0.90051 RIKEN cDNA 2900092C05 gene 2900092C05Rik
1429855_at 1.03338 RIKEN cDNA 1700023I07 gene 1700023I07Rik
1429856_at 0.89035 tetraspanin 18 Tspan18
1429857_at 1.19030 RIKEN cDNA 2900092D14 gene 2900092D14Rik
1429858_at 1.13476 RIKEN cDNA 1700011E24 gene 1700011E24Rik
1429859_a_at 0.99225 ADP-ribosylation factor-like 2 binding protein Arl2bp
1429860_at 1.03427 similar to RIKEN cDNA 5730590G19-like LOC677447
1429861_at 0.82119 protocadherin 9 /// similar to protocadherin 9 isoform 1 precursor Pcdh9 ///
LOC638275
1429862_at 0.98894 "phospholipase A2, group IVE" Pla2g4e
1429863_at 1.10139 LON peptidase N-terminal domain and ring finger 3 Lonrf3
1429864_at 1.01957 spermatogenesis and centriole associated 1 Spatc1
1429865_at 0.99081 RIKEN cDNA 4931417G12 gene 4931417G12Rik
1429866_at 1.12476 AHNAK nucleoprotein (desmoyokin) Ahnak
1429867_at 0.95669 RIKEN cDNA 4933424C08 gene 4933424C08Rik

1429868_at	0.99246 --- ---	
1429869_at	1.02242	RIKEN cDNA 1110020C03 gene 1110020C03Rik
1429870_at	0.92230	RIKEN cDNA C630040K21 gene C630040K21Rik
1429871_at	1.33672	hyaluronan mediated motility receptor (RHAMM) Hmnr
1429872_at	0.94536	RIKEN cDNA 1700007N14 gene 1700007N14Rik
1429873_at	0.89687	RIKEN cDNA 1700125H20 gene 1700125H20Rik
1429874_at	0.91239	spermatogenesis associated 17 Spata17
1429875_at	0.95185	RIKEN cDNA 1700034J05 gene /// hypothetical protein LOC635035 1700034J05Rik /// LOC635035
1429876_at	0.92951	suppressor of Ty 7 (<i>S. cerevisiae</i>)-like Supt7l
1429877_at	1.04379	leucine rich repeat containing 44 Lrrc44
1429878_a_at	0.97463	nucleoporin like 1 Nupl1
1429879_at	1.12479	RIKEN cDNA 0610037L13 gene 0610037L13Rik
1429880_at	1.03002	RIKEN cDNA 4921531C22 gene 4921531C22Rik
1429881_at	0.99290	Rho GTPase activating protein 15 Arhgap15
1429882_at	1.80946	RIKEN cDNA 2610005L07 gene 2610005L07Rik
1429883_at	1.28497	actin-like 6A Actl6a
1429884_at	0.93579	SLIT-ROBO Rho GTPase activating protein 2 Srgap2
1429885_at	1.06102	RIKEN cDNA 4930431P19 gene 4930431P19Rik
1429886_at	0.88583	zinc finger protein 294 Zfp294
1429887_at	0.98173	RIKEN cDNA 2310005C01 gene 2310005C01Rik
1429888_a_at	0.88601	heat shock protein 2 Hspb2
1429889_at	0.89239	Fas apoptotic inhibitory molecule 3 Faim3
1429890_at	1.00512	RIKEN cDNA A930007D18 gene A930007D18Rik
1429891_at	0.98302	calcyphosine-like Capsl
1429892_at	1.05136	PLAC8-like 1 Plac8l1
1429893_at	0.98423	RIKEN cDNA 2810004A10 gene 2810004A10Rik
1429894_a_at	0.98929	microtubule-associated protein 7 Mtap7
1429895_at	1.01351	RIKEN cDNA 2310010G23 gene 2310010G23Rik
1429896_at	1.15316	RIKEN cDNA 5830408B19 gene 5830408B19Rik
1429897_a_at	1.07072	"DNA segment, Chr 16, ERATO Doi 472, expressed" D16Ert472e
1429898_at	1.22014	RIKEN cDNA 2310008M10 gene 2310008M10Rik
1429899_at	0.89526	RIKEN cDNA 5730414N17 gene 5730414N17Rik
1429900_at	2.87693	RIKEN cDNA 5330406M23 gene 5330406M23Rik
1429901_at	1.07555	T-cell lymphoma breakpoint associated target 1 Tcba1
1429902_at	1.63161	RIKEN cDNA 5830443J22 gene 5830443J22Rik
1429903_at	1.07990	"gap junction membrane channel protein alpha 10 /// D4, zinc and double PHD fingers family 2" Gja10 /// Dpf2
1429904_at	0.97108	RIKEN cDNA 4930507A01 gene 4930507A01Rik
1429905_at	1.05400	RIKEN cDNA 3110009O07 gene 3110009O07Rik
1429906_at	0.86405	RIKEN cDNA A930035E12 gene A930035E12Rik
1429907_at	1.09049	RIKEN cDNA 1700094D03 gene 1700094D03Rik
1429908_at	0.93668	RIKEN cDNA 6530403A03 gene 6530403A03Rik

1429909_at	0.79306	RIKEN cDNA 4833411O04	gene	4833411O04Rik
1429910_at	0.94323	RIKEN cDNA 5730499H23	gene	5730499H23Rik
1429911_at	1.14550	"microcephaly, primary autosomal recessive 1"		Mcph1
1429912_at	1.04744	sperm associated antigen 4-like		Spag4l
1429913_at	0.98835	"potassium channel, subfamily K, member 16"		Kcnk16
1429914_at	0.71766	enhancer of polycomb homolog 1 (Drosophila)		Epc1
1429915_at	0.99074	RIKEN cDNA 4930426L09	gene	4930426L09Rik
1429916_at	0.94979	RIKEN cDNA 9530077C05	gene	9530077C05Rik
1429917_at	0.82253	RIKEN cDNA 9030625N01	gene	9030625N01Rik
1429918_at	0.92375	Rho GTPase activating protein 20		Arhgap20
1429919_at	1.01281	ankyrin repeat domain 48		Ankrd48
1429920_at	1.11986	RIKEN cDNA 4933403M22	gene	4933403M22Rik
1429921_at	1.25580	RIKEN cDNA 9530068E07	gene	9530068E07Rik
1429922_at	1.06745	"Sodium channel, voltage-gated, type IX, alpha"		Scn9a
1429923_x_at	0.97643	spermatogenesis associated 3		Spata3
1429924_at	1.07520	RIKEN cDNA 1700019O17	gene	1700019O17Rik
1429925_at	0.94125	RIKEN cDNA 4933425B16	gene	4933425B16Rik
1429926_at	1.26950	RIKEN cDNA 6720473G16	gene	6720473G16Rik
1429927_at	0.93945	RIKEN cDNA 5830409B07	gene	5830409B07Rik
1429928_at	1.20706	RIKEN cDNA 1700030B21	gene	1700030B21Rik
1429929_at	0.99044	meiosis defective 1		Mei1
1429930_at	0.91635	coiled-coil domain containing 19		Ccdc19
1429931_at	0.91476	RIKEN cDNA 4930553P18	gene	4930553P18Rik
1429932_at	0.90563	RIKEN cDNA 4930566F21	gene	4930566F21Rik
1429933_at	0.90598	ATP/GTP binding protein 1		Agtpbp1
1429934_at	0.96386	RIKEN cDNA 4930502E18	gene	4930502E18Rik
1429935_at	0.94334	lipocalin 12		Lcn12
1429936_at	0.95029	RIKEN cDNA 2700059L22	gene	2700059L22Rik
1429937_at	1.20059	RIKEN cDNA D530033C11	gene	D530033C11Rik
1429938_at	0.99700	RIKEN cDNA A930009E05	gene	A930009E05Rik
1429939_at	0.95807	"SAPS domain family, member 3"		Saps3
1429940_at	0.86154	"Vcell division cycle 73, Paf1/RNA polymerase II complex component, homolog (S. cerevisiae)"		Cdc73
1429941_at	0.89966	RIKEN cDNA 2410193C02	gene	2410193C02Rik
1429942_at	0.93815	glucocorticoid modulatory element binding protein 1		Gmeb1
1429943_at	0.88568	"chitinase, di-N-acetyl-"		Ctbs
1429944_at	1.00308	trichohyalin-like 1		Tchhl1
1429945_at	1.09714	RIKEN cDNA 2810406K13	gene	2810406K13Rik
1429946_at	1.05201	RIKEN cDNA 2610301F02	gene	2610301F02Rik
1429947_a_at	1.07234	Z-DNA binding protein 1		Zbp1
1429948_x_at	0.91735	"crystallin, gamma F"		Crygf
1429949_at	0.75128	RIKEN cDNA 6530415H11	gene	6530415H11Rik
1429950_at	0.95006	unc-5 homolog C (C. elegans)-like		Unc5cl

1429951_at	1.60359	single-stranded DNA binding protein 2	Ssbp2	
1429952_at	1.01914	motile sperm domain containing 4	Mospd4	
1429953_at	0.84199	RIKEN cDNA 2210011C24	gene	2210011C24Rik
1429954_at	1.01707	"C-type lectin domain family 4, member a3"	Clec4a3	
1429955_at	0.91893	RIKEN cDNA 5031434O11	gene	5031434O11Rik
1429956_at	1.12594	RIKEN cDNA 1700021K02	gene	1700021K02Rik
1429957_at	0.81350	RIKEN cDNA 2310002B14	gene	2310002B14Rik
1429958_x_at	1.24140	hydroxyacylglutathione hydrolase-like	Haghl	
1429959_at	0.96954	---	---	
1429960_at	0.87641	RIKEN cDNA 4930438A08	gene	4930438A08Rik
1429961_at	0.94151	RIKEN cDNA 1700021C14	gene	1700021C14Rik
1429962_at	1.15035	CCAAT/enhancer binding protein zeta	Cebpz	
1429963_at	1.37613	mitogen-activated protein kinase 6	Mapk6	
1429964_at	1.04048	RIKEN cDNA 5031420N21	gene	5031420N21Rik
1429965_at	0.96120	LON peptidase N-terminal domain and ring finger 2		Lonrf2
1429966_at	1.00841	RIKEN cDNA 3110057O12	gene	3110057O12Rik
1429967_at	1.02209	RIKEN cDNA 2600002B07	gene	2600002B07Rik
1429968_at	1.28914	RIKEN cDNA 2310047L11	gene	2310047L11Rik
1429969_at	0.95092	RIKEN cDNA 4833403J16	gene	4833403J16Rik
1429970_at	1.04771	sperm acrosome associated 3	Spaca3	
1429971_at	0.97025	thioredoxin reductase 2	Txnrd2	
1429972_s_at	0.90613	thioredoxin reductase 2	Txnrd2	
1429973_at	0.94560	RIKEN cDNA 4921509B22	gene	4921509B22Rik
1429974_at	0.68642	T-box18	Tbx18	
1429975_at	0.96936	RIKEN cDNA 4931432M23	gene /// similar to plasma kallikrein-like protein 4	4931432M23Rik /// LOC636530
1429976_at	0.88365	CLIP associating protein 2	Clasp2	
1429977_at	1.00419	RIKEN cDNA 9030425L15	gene	9030425L15Rik
1429978_at	0.95986	RIKEN cDNA 5830467E07	gene	5830467E07Rik
1429979_a_at	0.99462	RIKEN cDNA 1810073N04	gene	1810073N04Rik
1429980_x_at	0.96014	RIKEN cDNA 1810073N04	gene	1810073N04Rik
1429981_a_at	0.98625	RIKEN cDNA 4933426K21	gene	4933426K21Rik
1429982_at	1.03893	RIKEN cDNA 4933426K21	gene	4933426K21Rik
1429983_at	0.82395	RIKEN cDNA 2010002M09	gene	2010002M09Rik
1429984_at	1.07535	RIKEN cDNA 5730455O13	gene	5730455O13Rik
1429985_at	0.95520	RIKEN cDNA 5430439G13	gene	5430439G13Rik
1429986_at	0.94971	G-protein coupled receptor 173	Gpr173	
1429987_at	0.93245	RIKEN cDNA 9930013L23	gene	9930013L23Rik
1429988_at	0.89691	zinc finger protein 235	Zfp235	
1429989_at	0.85886	RIKEN cDNA 1700008A04	gene	1700008A04Rik
1429990_at	1.07391	hyaluronoglucosaminidase 4	Hyal4	
1429991_at	1.08450	Fez family zinc finger 1	Fezf1	
1429992_at	1.04134	spermatogenesis associated glutamate (E)-rich protein 4b		Speer4b

1429993_s_at	0.97482	spermatogenesis associated glutamate (E)-rich protein 4b	Speer4b
1429994_s_at	1.05370	"cytochrome P450, family 2, subfamily c, polypeptide 65"	Cyp2c65
1429995_at	1.05415	RIKEN cDNA 1700036D21 gene	1700036D21Rik
1429996_at	3.25059	RIKEN cDNA 6330417A16 gene	6330417A16Rik
1429997_at	1.01714	RIKEN cDNA 4832441B07 gene	4832441B07Rik
1429998_at	1.02429	tripartite motif protein 23	Trim23
1429999_at	1.00659	RIKEN cDNA 4933403M19 gene	4933403M19Rik
1430000_at	1.23420	RIKEN cDNA B230117O15 gene	B230117O15Rik
1430001_at	1.01222	interleukin 31	Il31
1430002_at	0.91457	IQ motif containing F1	Iqcf1
1430003_at	0.96802	RIKEN cDNA 6330412A17 gene	6330412A17Rik
1430004_s_at	0.96431	WD repeat domain 20	Wdr20
1430005_a_at	0.89171	"basic leucine zipper transcription factor, ATF-like 2"	Batf2
1430006_x_at	1.06509	RIKEN cDNA 1700024B05 gene	1700024B05Rik
1430007_a_at	0.85397	calcium-binding tyrosine-(Y)-phosphorylation regulated (fibrousheathin 2)	Cabyr
1430008_x_at	1.00034	"spermatogenesis associated glutamate (E)-rich protein 5, pseudogene 1"	Speer5-ps1
1430009_at	0.93507	"heat shock transcription factor, Y linked 2"	Hsfy2
1430010_at	1.18575	RIKEN cDNA 2810465G24 gene	2810465G24Rik
1430011_at	1.05612	RIKEN cDNA 2610317O13 gene	2610317O13Rik
1430012_at	0.92385	RIKEN cDNA 1110050K14 gene	1110050K14Rik
1430013_at	0.92269	RIKEN cDNA 2210008F06 gene	2210008F06Rik
1430014_at	1.09891	RIKEN cDNA 5430419D17 gene	5430419D17Rik
1430015_at	1.07120	"peroxisomal, testis specific 1"	Pxt1
1430016_at	0.99391	RIKEN cDNA 4930584F24 gene	4930584F24Rik
1430017_at	1.10014	RIKEN cDNA 1600016N20 gene	1600016N20Rik
1430018_at	0.89441	CDC42 effector protein (Rho GTPase binding) 5	Cdc42ep5
1430019_a_at	1.07442	heterogeneous nuclear ribonucleoprotein A1 /// heterogeneous nuclear ribonucleoprotein A1 pseudogene /// heterogeneous nuclear ribonucleoprotein A1 pseudogene /// heterogeneous nuclear ribonucleoprotein A1 pseudogene /// similar to Heterogeneous nuclear ribonucleoprotein A1 (Helix-destabilizing protein) (Single-strand binding protein) (hnRNP core protein A1) (HDP-1) (Topoisomerase-inhibitor suppressed)	Hnrpa1 /// LOC225307 /// LOC434858 /// LOC654467 /// LOC665646
1430020_x_at	1.10213	heterogeneous nuclear ribonucleoprotein A1	Hnrpa1
1430021_a_at	1.08087	ubiquitin-like 1 (sentrin) activating enzyme E1A	Uble1a
1430022_at	0.94328	ubiquitin-like 1 (sentrin) activating enzyme E1A	Uble1a
1430023_at	1.04291	RIKEN cDNA 5133400G04 gene	5133400G04Rik
1430024_at	1.07354	"Yip1 domain family, member 6"	Yipf6
1430025_at	0.94283	"protein phosphatase 3, catalytic subunit, gamma isoform"	Ppp3cc
1430026_at	0.99500	"stress 70 protein chaperone, microsome-associated, human homolog"	Stch
1430027_at	1.04608	RIKEN cDNA 1810028F09 gene	1810028F09Rik
1430028_at	1.01217	RIKEN cDNA 2210018M11 gene	2210018M11Rik

1430029_a_at	0.93874	tetraspanin 31	Tspan31	
1430030_at	1.28130	RIKEN cDNA 5330426P16	gene	5330426P16Rik
1430031_at	0.87709	inositol polyphosphate multikinase	Ipmk	
1430032_at	0.81723	RNA binding motif protein 4B	Rbm4b	
1430033_at	1.46525	RIKEN cDNA 5330431K02	gene	5330431K02Rik
1430034_at	1.04447	chaperonin subunit 4 (delta)	Cct4	
1430035_at	1.05646	RIKEN cDNA 1700034M03	gene	1700034M03Rik
1430036_at	1.01429	RIKEN cDNA 2310015B20	gene	2310015B20Rik
1430037_at	1.06914	sorting nexin family member 27	Snx27	
1430038_at	0.98335	gephyrin	Gphn	
1430039_at	0.93565	CDK5 regulatory subunit associated protein 1-like 1	Cdkal1	
1430040_at	0.86464	heat shock protein 12A	Hspa12a	
1430041_at	1.12700	RIKEN cDNA 6330548G22	gene	6330548G22Rik
1430042_at	1.24439	RIKEN cDNA A930012M21	gene	A930012M21Rik
1430043_at	1.70833	tetratricopeptide repeat domain 19	Ttc19	
1430044_at	0.96491	RIKEN cDNA 4930415F15	gene	4930415F15Rik
1430045_at	1.08124	translin-associated factor X	Tsnax	
1430046_at	1.08436	OTU domain containing 3	Otud3	
1430047_at	1.00624	BRCT domain containing 1	Brctd1	
1430048_at	1.04080	RIKEN cDNA 7630402I04	gene	7630402I04Rik
1430049_at	1.04843	RIKEN cDNA 4932701A20	gene	4932701A20Rik
1430050_at	1.06097	DEAD (Asp-Glu-Ala-Asp) box polypeptide 23	Ddx23	
1430051_at	1.07542	RIKEN cDNA 4930486L24	gene	4930486L24Rik
1430052_at	1.05666	RIKEN cDNA C030048H21	gene	C030048H21Rik
1430053_a_at	1.17240	RIKEN cDNA 2810409H07	gene	2810409H07Rik
1430054_at	0.98185	RIKEN cDNA 1700030F18	gene	1700030F18Rik
1430055_at	0.93305	RIKEN cDNA 1700080G18	gene	1700080G18Rik
1430056_at	1.00670	RIKEN cDNA D130059P03	gene	D130059P03Rik
1430057_s_at	0.87004	leucine rich repeat containing 57	Lrrc57	
1430058_at	1.86700	stem-loop binding protein	Slbp	
1430059_at	1.15877	WD repeat domain 40A	Wdr40a	
1430060_at	1.00317	RIKEN cDNA 1700034G24	gene	1700034G24Rik
1430061_at	0.99094	RIKEN cDNA 1700105P06	gene	1700105P06Rik
1430062_at	1.04826	RIKEN cDNA 1600002O04	gene	1600002O04Rik
1430063_at	0.98107	RIKEN cDNA 4930571K23	gene	4930571K23Rik
1430064_at	1.13938	RIKEN cDNA 4933424A20	gene	4933424A20Rik
1430065_at	1.00969	RIKEN cDNA A930005N03	gene	A930005N03Rik
1430066_at	0.93618	RIKEN cDNA 4930589M24	gene	4930589M24Rik
1430067_at	1.00858	RIKEN cDNA 5430401F13	gene	5430401F13Rik
		5430401F13Rik	///	LOC625758
1430068_at	1.00892	RIKEN cDNA C030011L09	gene	C030011L09Rik
1430069_at	1.00130	RIKEN cDNA 4921517L17	gene	4921517L17Rik
1430070_at	1.01498	RIKEN cDNA 1500035N22	gene	1500035N22Rik

1430071_at	1.03469	brain protein 44	Brp44	
1430072_at	0.97376	RIKEN cDNA 1700057K13	gene	1700057K13Rik
1430073_at	1.02355	RIKEN cDNA 2900016B01	gene	2900016B01Rik
1430074_x_at	1.10887	RIKEN cDNA 8430426H19	gene	8430426H19Rik
1430075_at	1.56313	"splicing factor 3b, subunit 3"	Sf3b3	
1430076_at	0.99078	coiled-coil domain containing 116	Ccdc116	
1430077_at	1.27437	"splicing factor, arginine/serine-rich 11"	Sfrs11	
1430078_a_at	0.94337	8-oxoguanine DNA-glycosylase 1	Ogg1	
1430079_at	1.01615	RIKEN cDNA 5033406O09	gene	5033406O09Rik
1430080_at	1.06059	ficolin B	Fcnb	
1430081_at	0.94559	PHD finger protein 15	Phf15	
1430082_at	0.84237	RIKEN cDNA 4930511H01	gene	4930511H01Rik
1430083_at	0.77669	RIKEN cDNA 2610307P16	gene	2610307P16Rik
1430084_at	1.03879	TLC domain containing 1	Tlcd1	
1430085_at	1.07244	RIKEN cDNA 4933412A08	gene	4933412A08Rik
1430086_at	1.02357	"cholinergic receptor, nicotinic, alpha polypeptide 9"	Chrna9	
1430087_at	0.92359	"NTPase, KAP family P-loop domain containing 1"	Nkpd1	
1430088_at	0.96255	RIKEN cDNA 3000002G13	gene	3000002G13Rik
1430089_at	2.42141	RIKEN cDNA 5830469G19	gene	5830469G19Rik
1430090_at	0.98402	coiled coil domain containing 28B	Ccdc28b	
1430091_at	1.00126	RIKEN cDNA 1700001P01	gene	1700001P01Rik
1430092_at	1.13481	serine active site containing 1	Serac1	
1430093_at	0.98571	RIKEN cDNA 4933406P04	gene	4933406P04Rik
1430094_at	1.15032	RIKEN cDNA 4931440L10	gene	4931440L10Rik
1430095_at	1.02534	RIKEN cDNA 1500017I17	gene	1500017I17Rik
1430096_at	0.96431	RIKEN cDNA 2900017F05	gene	2900017F05Rik
1430097_at	1.24961	RIKEN cDNA 8430436C05	gene	8430436C05Rik
1430098_at	0.97102	RIKEN cDNA 6330409D20	gene	6330409D20Rik
1430099_at	0.96874	RIKEN cDNA 4930509E16	gene	4930509E16Rik
1430100_at	1.06160	mitochondrial ribosomal protein S15	Mrps15	
1430101_at	0.93042	RIKEN cDNA 6330406O05	gene	6330406O05Rik
1430102_at	0.91845	ring finger protein 148	Rnf148	
1430103_at	1.00430	RIKEN cDNA 9030607L20	gene	9030607L20Rik
1430104_at	0.96277	R-spondin 2 homolog (<i>Xenopus laevis</i>)	Rspo2	
1430105_at	0.88453	RIKEN cDNA 1700042O10	gene	1700042O10Rik
1430106_at	1.01808	RIKEN cDNA 1700126L10	gene	1700126L10Rik
1430107_at	1.12589	RIKEN cDNA 9230116B18	gene	9230116B18Rik
1430108_at	1.04609	RIKEN cDNA 9030622M22	gene	9030622M22Rik
1430109_at	1.61961	RIKEN cDNA 4833412E19	gene	4833412E19Rik
1430110_at	1.07560	RIKEN cDNA D230044M03	gene	D230044M03Rik
1430111_a_at	1.01863	"branched chain aminotransferase 1, cytosolic"	Bcat1	
1430112_at	0.93389	WD repeat domain 66	Wdr66	
1430113_at	0.91319	RIKEN cDNA 4933424C13	gene	4933424C13Rik

1430114_at	0.95080	RIKEN cDNA 5430420C16	gene	5430420C16Rik
1430115_at	1.19144	RIKEN cDNA 4833413E03	gene	4833413E03Rik
1430116_at	1.01944	RIKEN cDNA 6030422M01	gene	6030422M01Rik
1430117_a_at	0.89512	zinc finger protein 64		Zfp64
1430118_at	0.88716	RIKEN cDNA 2700046A07	gene	2700046A07Rik
1430119_at	0.90720	fibronectin type III domain containing 1		Fndc1
1430120_at	0.92898	RIKEN cDNA 4930519G04	gene	4930519G04Rik
1430121_at	0.97520	RIKEN cDNA 1700006E09	gene	1700006E09Rik
1430122_at	1.11445	RIKEN cDNA 9430002A10	gene	9430002A10Rik
1430123_a_at	0.95645	"aldo-keto reductase family 1, member A4 (aldehyde reductase)"		Akr1a4
1430124_x_at	0.93893	"aldo-keto reductase family 1, member A4 (aldehyde reductase)"		Akr1a4
1430125_s_at	0.72378	PQ loop repeat containing 1		Pqlc1
1430126_at	0.88624	RIKEN cDNA 2310035C23	gene	2310035C23Rik
1430127_a_at	1.05984	cyclin D2		Ccnd2
1430128_a_at	0.89874	receptor accessory protein 6		Reep6
1430129_a_at	1.01274	COMM domain containing 8		Commd8
1430130_at	1.00223	von Willebrand factor C and EGF domains		Vwce
1430131_at	0.98245	complement receptor related protein		Crry
1430132_at	1.00093	keratin 28		Krt28
1430133_at	1.06509	"TBC1 domain family, member 8B"		Tbc1d8b
1430134_a_at	1.03772	tyrosyl-tRNA synthetase 2 (mitochondrial)		Yars2
1430135_at	0.95130	deoxyribonuclease II alpha		Dnase2a
1430136_at	1.00950	"glutamate receptor, metabotropic 3"		Grm3
1430137_at	0.98279	mitogen-activated protein kinase kinase kinase 13		Map3k13
1430138_at	1.19688	"CD3E antigen, epsilon polypeptide associated protein"		Cd3eap
1430139_at	1.05706	"helicase, lymphoid specific"		Hells
1430140_at	0.94340	RIKEN cDNA 1700017N19	gene	1700017N19Rik
1430141_at	1.10059	RIKEN cDNA 1200015N20	gene	1200015N20Rik
1430142_at	0.98631	transglutaminase 5		Tgm5
1430143_at	1.03434	RIKEN cDNA 4930426D05	gene	4930426D05Rik
1430144_at	1.11194	RIKEN cDNA 5830411G16	gene	5830411G16Rik
1430145_at	1.06233	leucine rich repeat containing 28		Lrrc28
1430146_at	1.05173	RIKEN cDNA 4933402N03	gene	4933402N03Rik
1430147_a_at	1.11450	Josephin domain containing 3		Josd3
1430148_at	1.05479	"RAB19, member RAS oncogene family"		Rab19
1430149_at	1.08342	"NADH dehydrogenase (ubiquinone) 1 alpha subcomplex, 3"		Ndufa3
1430150_at	1.06543	DCP2 decapping enzyme homolog (S. cerevisiae)		Dcp2
1430151_at	1.85976	nischarin		Nisch
1430152_at	1.43956	epidermal growth factor receptor pathway substrate 15		Eps15
1430153_at	0.92820	tropomodulin 2		Tmod2
1430154_at	1.02337	RIKEN cDNA 4930543C13	gene	4930543C13Rik
1430155_at	1.04495	RIKEN cDNA 4933427E11	gene	4933427E11Rik
1430156_at	1.02217	RIKEN cDNA 4930520O04	gene	4930520O04Rik

1430157_at 1.08350 RIKEN cDNA 1700095J03 gene 1700095J03Rik
1430158_at 1.18194 RIKEN cDNA 3110021A11 gene 3110021A11Rik
1430159_at 1.40498 RIKEN cDNA 5830408C22 gene 5830408C22Rik
1430160_at 1.00015 ankyrin repeat and IBR domain containing 1 Ankib1
1430161_at 1.10791 dihydrolipoamide S-succinyltransferase (E2 component of 2-oxo-glutarate complex) Dlst
1430162_at 0.84731 RIKEN cDNA 3830417A13 gene 3830417A13Rik
1430163_at 1.03963 RIKEN cDNA 1810048P08 gene 1810048P08Rik
1430164_a_at 1.11084 growth factor receptor bound protein 10 Grb10
1430165_at 1.09840 serine/threonine kinase 17b (apoptosis-inducing) Stk17b
1430166_at 0.98314 RIKEN cDNA 1700042D18 gene 1700042D18Rik
1430167_a_at 0.95026 RWD domain containing 3 Rwdd3
1430168_at 1.07129 "CSA-conditional, T cell activation-dependent protein" Cstad
1430169_at 0.92068 RIKEN cDNA 1700007K09 gene 1700007K09Rik
1430170_at 0.83822 Bardet-Biedl syndrome 10 (human) Bbs10
1430171_at 1.29306 fibroblast growth factor (acidic) intracellular binding protein Fibp
1430172_a_at 0.83974 "cytochrome P450, family 4, subfamily f, polypeptide 16 /// similar to cytochrome P450, family 4, subfamily f, polypeptide 16 /// similar to cytochrome P450, family 4, subfamily f, polypeptide 16" Cyp4f16 /// LOC623748 /// LOC677156
1430173_x_at 1.09153 "cytochrome P450, family 4, subfamily f, polypeptide 16" Cyp4f16
1430174_at 0.96614 RIKEN cDNA A930010G16 gene A930010G16Rik
1430175_at 1.03418 RIKEN cDNA 4930588G05 gene 4930588G05Rik
1430176_at 0.94831 RIKEN cDNA 5430433E21 gene 5430433E21Rik
1430177_at 1.12680 "ubiquitin-conjugating enzyme E2B, RAD6 homology (S. cerevisiae)" Ube2b
1430178_at 1.01959 RIKEN cDNA 2810408A11 gene 2810408A11Rik
1430179_at 1.14692 RIKEN cDNA 4921507G05 gene 4921507G05Rik
1430180_at 1.43003 mitogen activated protein kinase kinase 5 Map2k5
1430181_at 1.03824 "H1 histone family, member N, testis-specific" H1fnt
1430182_a_at 0.96975 RIKEN cDNA 1700006J14 gene 1700006J14Rik
1430183_at 2.48506 Pre B-cell leukemia transcription factor 1 Pbx1
1430184_at 1.06873 zinc finger protein 597 Zfp597
1430185_at 1.11426 RIKEN cDNA 5830460E08 gene 5830460E08Rik
1430186_at 1.01868 "odontogenic, ameloblast associated" Odam
1430187_at 0.93239 zinc finger protein 60 Zfp60
1430188_at 1.00204 RIKEN cDNA 1700037C18 gene 1700037C18Rik
1430189_at 1.03347 nucleolar protein 4 Nol4
1430190_at 0.89710 RIKEN cDNA 1700041C02 gene 1700041C02Rik
1430191_at 4.18034 RIKEN cDNA 9130004J05 gene 9130004J05Rik
1430192_at 1.07514 caspase 3 Casp3
1430193_at 0.90640 cancer susceptibility candidate 5 /// similar to cancer susceptibility candidate 5 isoform 2 Casc5 /// LOC677208
1430194_at 0.91074 RIKEN cDNA 1110018H23 gene 1110018H23Rik

1430195_at	2.61970	RIKEN cDNA 2810043O03	gene	2810043O03Rik
1430196_at	1.11850	RIKEN cDNA 8430408J09	gene	8430408J09Rik
1430197_a_at	0.90746	"phosphatidylinositol transfer protein, membrane-associated 2"		Pitpnm2
1430198_at	1.24668	RIKEN cDNA 3110009F21	gene	3110009F21Rik
1430199_at	1.01250	RIKEN cDNA 1700007E05	gene	1700007E05Rik
1430200_at	1.00671	RIKEN cDNA 4930579D07	gene	4930579D07Rik
1430201_at	1.24814	RIKEN cDNA 9130017K11	gene	9130017K11Rik
1430202_at	1.02647	RIKEN cDNA 1700108F19	gene	1700108F19Rik
1430203_at	0.95172	ubiquitin specific peptidase 16		Usp16
1430204_at	0.99238	RIKEN cDNA 5033425G24	gene	5033425G24Rik
1430205_a_at	1.00615	cell division cycle 37 homolog (S. cerevisiae)-like 1		Cdc3711
1430206_at	0.86392	RIKEN cDNA 4933407L21	gene	4933407L21Rik
1430207_at	1.09159	RIKEN cDNA 4933437N03	gene	4933437N03Rik
1430208_at	0.90144	keratin 42		Krt42
1430209_at	0.98289	RIKEN cDNA 4930404H21	gene	4930404H21Rik
1430210_at	1.19512	RIKEN cDNA C330024D12	gene	C330024D12Rik
1430211_at	1.82800	RIKEN cDNA 4930415O20	gene	4930415O20Rik
1430212_at	0.84711	zinc finger protein 712		Zfp712
1430213_at	0.84398	regulator of chromosome condensation (RCC1) and BTB (POZ) domain containing protein 1		Rcbtb1
1430214_a_at	1.07681	RIKEN cDNA 2610020H08	gene	2610020H08Rik
1430215_at	0.81228	RIKEN cDNA 2610020H08	gene	2610020H08Rik
1430216_at	0.90196	zinc finger protein 292		Zfp292
1430217_at	0.93392	RIKEN cDNA 4921528H16	gene	4921528H16Rik
1430218_at	1.03780	RIKEN cDNA 4933424M12	gene	4933424M12Rik
1430219_at	0.91433	fused toes		Fts
1430220_at	1.11491	RIKEN cDNA 4833420G17	gene	4833420G17Rik
1430221_at	0.95199	RIKEN cDNA 9130008F23	gene	9130008F23Rik
1430222_at	1.01621	RIKEN cDNA 9130007G19	gene	9130007G19Rik
1430223_at	0.81479	RIKEN cDNA 4930442P07	gene	4930442P07Rik
1430224_at	0.87572	WAP four-disulfide core domain 3		Wfdc3
1430225_at	1.02453	RIKEN cDNA 5730442G03	gene	5730442G03Rik
1430226_at	1.05139	X-linked lymphocyte-regulated 4C		Xlr4c
1430227_at	1.01562	RIKEN cDNA 1700007I08	gene	1700007I08Rik
1430228_at	2.05113	denticleless homolog (Drosophila)		Dtl
1430229_at	0.94319	RIKEN cDNA 6330416L11	gene	6330416L11Rik
1430230_at	1.01150	RCSD domain containing 1		Rcsd1
1430231_a_at	0.74894	centrosomal protein 70		Cep70
1430232_at	0.94847	RIKEN cDNA 4933407E14	gene	4933407E14Rik
1430233_a_at	0.93093	RIKEN cDNA 4933425K02	gene	4933425K02Rik
1430234_at	1.00580	ADP-ribosylation factor-like 14		Arl14
1430235_at	0.93917	RIKEN cDNA 4933424G06	gene	4933424G06Rik
1430236_s_at	1.08440	gasdermin 2		Gsdm2

1430237_at	1.02103	claudin 22	Cldn22	
1430238_at	1.07043	glutamic-oxaloacetic transaminase 1-like 1	Got111	
1430239_at	0.98134	RIKEN cDNA 5430421F17 gene	5430421F17Rik	
1430240_a_at	1.03462	calmegin	Clgn	
1430241_at	0.88262	F-box and leucine-rich repeat protein 17	Fbxl17	
1430242_at	0.94771	RWD domain containing 1	Rwdd1	
1430243_at	0.84155	RIKEN cDNA 1700058C01 gene	1700058C01Rik	
1430244_at	1.22800	RIKEN cDNA 4921509J17 gene	4921509J17Rik	
1430245_at	1.00237	---	---	
1430246_at	1.03064	RIKEN cDNA 4933406C10 gene	4933406C10Rik	
1430247_at	0.82844	dishevelled associated activator of morphogenesis 2	Daam2	
1430248_at	1.05331	RIKEN cDNA 4930415O11 gene	4930415O11Rik	
1430249_at	0.99580	RIKEN cDNA 4930473A02 gene	4930473A02Rik	
1430250_at	1.00687	"SEC24 related gene family, member C (<i>S. cerevisiae</i>)"	Sec24c	
1430251_at	1.19258	RIKEN cDNA D330022H12 gene	D330022H12Rik	
1430252_at	0.87671	RIKEN cDNA 3110027N22 gene	3110027N22Rik	
1430253_at	1.05495	RIKEN cDNA 2900006B11 gene	2900006B11Rik	
1430254_at	1.01006	RIKEN cDNA 4921517D22 gene	4921517D22Rik	
1430255_at	1.00516	Kruppel-like factor 5	Klf5	
1430256_at	1.01328	RIKEN cDNA 2310033K02 gene	2310033K02Rik	
1430257_at	0.83823	"caspase recruitment domain family, member 11"	Card11	
1430258_at	1.14701	RIKEN cDNA 2810422J05 gene	2810422J05Rik	
1430259_at	0.94811	"tumor necrosis factor receptor superfamily, member 11a"	Tnfrsf11a	
1430260_at	1.03376	---	---	
1430261_at	0.93449	---	---	
1430262_at	0.92961	RIKEN cDNA 5330411J11 gene	5330411J11Rik	
1430263_at	1.07069	RIKEN cDNA 1700109F18 gene	1700109F18Rik	
1430264_at	1.12957	RIKEN cDNA 2610030P05 gene	2610030P05Rik	
1430265_at	1.11156	amyloid beta precursor protein (cytoplasmic tail) binding protein 2	Appbp2	
1430266_at	0.91297	RIKEN cDNA 1700003D09 gene	1700003D09Rik	
1430267_at	0.81887	RIKEN cDNA 4833422M21 gene	4833422M21Rik	
1430268_at	1.06948	RIKEN cDNA 9630005C17 gene	9630005C17Rik	
1430269_at	1.02427	myosin binding protein H-like	Mybphl	
1430270_at	1.09073	RIKEN cDNA 4933411K05 gene	4933411K05Rik	
1430271_x_at	1.03205	Josephin domain containing 3	Josd3	
1430272_at	1.24682	GTP binding protein 5	Gtpbp5	
1430273_at	1.04974	RIKEN cDNA 2410087M07 gene	2410087M07Rik	
1430274_a_at	0.94125	STARD3 N-terminal like	Stard3nl	
1430275_a_at	1.08861	aquarius	Aqr	
1430276_at	1.08835	RIKEN cDNA 1700003I16 gene	1700003I16Rik	
1430277_at	1.30837	RIKEN cDNA 9430034F23 gene	9430034F23Rik	
1430278_a_at	1.04783	DEAQ RNA-dependent ATPase	Dqx1	
1430279_at	1.02721	RIKEN cDNA 3110038A09 gene	3110038A09Rik	

1430280_at	1.14422	RIKEN cDNA 1810062G17 gene	1810062G17Rik
1430281_at	0.96493	GLI pathogenesis-related 1 like 1	Glipr111
1430282_at	0.86722	RIKEN cDNA 5730437N04 gene	5730437N04Rik
1430283_s_at	1.13450	RIKEN cDNA 4930418G15 gene	4930418G15Rik
1430284_at	0.96218	RIKEN cDNA 1700008P02 gene	1700008P02Rik
1430285_at	1.00509	"protein phosphatase 1, regulatory (inhibitor) subunit 14c"	Ppp1r14c
1430286_s_at	0.97923	"protein phosphatase 1, regulatory (inhibitor) subunit 14c"	Ppp1r14c
1430287_s_at	0.99655	HemK methyltransferase family member 1	Hemk1
1430288_x_at	0.96503	ribosomal protein S21	Rps21
1430289_a_at	1.11541	WD repeat domain 77	Wdr77
1430290_at	1.18094	ribosomal protein S15a /// similar to ribosomal protein S15a /// similar to ribosomal protein S15a /// similar to ribosomal protein S15a	Rps15a /// LOC434460 /// LOC664903 /// LOC675729
1430291_at	0.99833	RIKEN cDNA 1110060D06 gene /// dedicator of cytokinesis 5	1110060D06Rik /// Dock5
1430292_a_at	0.94166	RIKEN cDNA 1810030N24 gene	1810030N24Rik
1430293_a_at	1.07245	RIKEN cDNA B230118G17 gene	B230118G17Rik
1430294_at	1.10270	single-stranded DNA binding protein 1	Ssbp1
1430295_at	1.02902	"guanine nucleotide binding protein, alpha 13"	Gna13
1430296_at	0.86819	RIKEN cDNA 4733401H21 gene	4733401H21Rik
1430297_a_at	1.05440	RIKEN cDNA 2010301N04 gene	2010301N04Rik
1430298_at	1.09603	RIKEN cDNA 4930473A06 gene	4930473A06Rik
1430299_at	0.99733	TSC22 domain family 4	Tsc22d4
1430300_at	0.93953	SCO cytochrome oxidase deficient homolog 1 (yeast)	Sco1
1430301_at	0.95880	syntaxin binding protein 5 (tomosyn)	Stxbp5
1430302_at	0.90704	RIKEN cDNA 1500041B16 gene	1500041B16Rik
1430303_at	1.17186	RIKEN cDNA 4921537D05 gene	4921537D05Rik
1430304_at	0.99870	WD repeat domain 76	Wdr76
1430305_at	1.07195	RIKEN cDNA 2500004C02 gene	2500004C02Rik
1430306_a_at	0.99284	"ATPase, H+ transporting, lysosomal V1 subunit C2"	Atp6v1c2
1430307_a_at	1.11376	"malic enzyme, supernatant /// similar to NADP-dependent malic enzyme (NADP-ME) (Malic enzyme 1) /// similar to NADP-dependent malic enzyme (NADP-ME) (Malic enzyme 1)"	Mod1 /// LOC624892 /// LOC677317
1430308_at	1.08907	RIKEN cDNA 5730409N16 gene	5730409N16Rik
1430309_at	1.11092	Nipped-B homolog (Drosophila)	Nipbl
1430310_at	0.88679	tetraspanin 11	Tspan11
1430311_at	1.25525	Myristoylated alanine rich protein kinase C substrate	Marcks
1430312_at	0.96966	tryptophan rich basic protein	Wrb
1430313_at	0.94100	ADAMTS-like 1	Adamts11
1430314_at	0.86478	RIKEN cDNA 4933437F05 gene	4933437F05Rik
1430315_at	0.91723	RIKEN cDNA 1700010N08 gene	1700010N08Rik
1430316_at	0.96581	RIKEN cDNA 4930579E17 gene	4930579E17Rik
1430317_at	1.29946	"ubiquitin-conjugating enzyme E2, J2 homolog (yeast)"	Ube2j2

1430318_at 0.94959 spermidine/spermine N1-acetyl transferase 2 Sat2
1430319_at 1.07594 RIKEN cDNA 4833411C07 gene 4833411C07Rik
1430320_at 0.90104 "dystrophin, muscular dystrophy" Dmd
1430321_at 0.92045 RIKEN cDNA A930040G15 gene A930040G15Rik
1430322_at 0.96206 ADP-ribosylation factor-like 13B Arl13b
1430323_at 0.97171 RIKEN cDNA 1700049E17 gene /// hypothetical protein LOC624112 /// similar
to Spetex-2C protein /// similar to Spetex-2C protein 1700049E17Rik /// LOC624112 ///
LOC638262 /// LOC666214
1430324_x_at 1.02725 RIKEN cDNA 1700049E17 gene /// similar to Spetex-2C protein
1700049E17Rik /// LOC666214
1430325_at 0.96472 --- ---
1430326_s_at 1.02884 "ubiquinol-cytochrome c reductase, complex III subunit VII" Uqcrcq
1430327_at 0.98028 secretory carrier membrane protein 4 Scamp4
1430328_at 1.33472 polymerase (RNA) III (DNA directed) polypeptide F Polr3f
1430329_at 1.20710 cingulin Cgn
1430330_at 1.02911 widely-interspaced zinc finger motifs Wiz
1430331_at 0.94396 RIKEN cDNA 1110005A03 gene 1110005A03Rik
1430332_a_at 0.84813 "glucuronidase, beta" Gusb
1430333_at 1.05398 elongation factor Tu GTP binding domain containing 1 Eftud1
1430334_at 0.91232 zinc finger protein 192 Zfp192
1430335_a_at 1.02119 paired box gene 3 Pax3
1430336_at 0.97430 guanylate cyclase 2g Gucy2g
1430337_at 0.91148 RIKEN cDNA 6530401N04 gene 6530401N04Rik
1430338_at 0.98074 RIKEN cDNA 4122401K19 gene 4122401K19Rik
1430339_at 0.90760 RIKEN cDNA 2610021K21 gene 2610021K21Rik
1430340_at 0.84936 RIKEN cDNA 4930431B11 gene 4930431B11Rik
1430341_at 1.30082 nudix (nucleoside diphosphate linked moiety X)-type motif 5 Nudt5
1430342_at 0.93604 RIKEN cDNA 4930518J20 gene 4930518J20Rik
1430343_at 0.90286 nucleoporin 205 Nup205
1430344_at 1.05412 RIKEN cDNA 2900064P18 gene 2900064P18Rik
1430345_at 1.07977 RIKEN cDNA 5530402H23 gene 5530402H23Rik
1430346_at 0.93741 RIKEN cDNA 5730507A09 gene 5730507A09Rik
1430347_at 1.00136 RIKEN cDNA 4930524J08 gene 4930524J08Rik
1430348_at 1.13217 RIKEN cDNA 2900019E01 gene 2900019E01Rik
1430349_at 0.88627 "zinc finger, DHHC domain containing 11" Zdhhc11
1430350_at 0.99772 RIKEN cDNA 2610035F20 gene 2610035F20Rik
1430351_at 1.04325 spermatogenesis associated 18 Spata18
1430352_at 1.76198 RIKEN cDNA 8430417A20 gene 8430417A20Rik
1430353_at 0.83702 GLIS family zinc finger 3 Glis3
1430354_x_at 0.88423 SCY1-like 1 (S. cerevisiae) Scyl1
1430355_a_at 0.98816 STEAP family member 3 Steap3
1430356_at 1.12324 RIKEN cDNA 2210402A03 gene 2210402A03Rik
1430357_at 2.71064 "H3 histone, family 3B" H3f3b

1430358_at 1.06853 "beclin 1 (coiled-coil, myosin-like BCL2-interacting protein)" Becn1
1430359_a_at 0.94870 NAD synthetase 1 Nadsyn1
1430360_at 1.18186 RIKEN cDNA 4833412C15 gene 4833412C15Rik
1430361_at 0.89605 RIKEN cDNA 2610005M20 gene 2610005M20Rik
1430362_at 0.98697 RIKEN cDNA 5730409N24 gene 5730409N24Rik
1430363_at 0.81366 RIKEN cDNA 4933407O12 gene 4933407O12Rik
1430364_at 1.08431 activating transcription factor 7 interacting protein 2 Atf7ip2
1430365_at 1.02893 "nuclear transcription factor, X-box binding-like 1" Nfxl1
1430366_at 0.95977 RIKEN cDNA 5430405H02 gene 5430405H02Rik
1430367_at 1.25235 Stam binding protein like 1 Stambpl1
1430368_s_at 0.99859 RIKEN cDNA 1700019D03 gene 1700019D03Rik
1430369_at 1.13710 erythrocyte protein band 4.1 Epb4.1
1430370_at 1.13508 glutamyl-prolyl-tRNA synthetase Eprs
1430371_x_at 1.06205 eukaryotic translation initiation factor 2 alpha kinase 3 Eif2ak3
1430372_at 0.86534 RIKEN cDNA D730045B01 gene D730045B01Rik
1430373_at 0.98216 RIKEN cDNA 5430427O19 gene 5430427O19Rik
1430374_at 1.00266 "solute carrier family 25, member 37" Slc25a37
1430375_a_at 0.98247 chemokine (C-C motif) ligand 27 Ccl27
1430376_at 0.98089 leucine rich repeat containing 9 Lrrc9
1430377_at 1.02933 RIKEN cDNA 1300015D01 gene 1300015D01Rik
1430378_at 0.85780 RIKEN cDNA 2900011G08 gene 2900011G08Rik
1430379_at 1.16904 RIKEN cDNA 5830411K21 gene 5830411K21Rik
1430380_at 1.13458 HemK methyltransferase family member 2 Hemk2
1430381_at 0.98865 Ftsj homolog 1 (E. coli) Ftsj1
1430382_at 0.87761 RIKEN cDNA 4833413G10 gene 4833413G10Rik
1430383_at 1.03856 CEA-related cell adhesion molecule 20 Ceacam20
1430384_at 1.47389 "transducin-like enhancer of split 4, homolog of Drosophila E(spl)" Tle4
1430385_a_at 0.91133 "galactosidase, beta 1-like" Glb1l
1430386_at 1.91338 RIKEN cDNA E030024N20 gene E030024N20Rik
1430387_at 1.03050 RIKEN cDNA 1810073O08 gene 1810073O08Rik
1430388_a_at 0.72866 sulfatase 2 Sulf2
1430389_at 0.94447 RIKEN cDNA 4930509H03 gene 4930509H03Rik
1430390_x_at 0.92613 RIKEN cDNA 4930444G20 gene /// similar to sentrin 18 /// similar to Sumo1/
sentrin/Smt3 specific protease 17 /// similar to Sumo1/sentrin/Smt3 specific protease 17
4930444G20Rik /// LOC625660 /// LOC639644 /// LOC672736
1430391_a_at 0.94767 "ST8 alpha-N-acetyl-neuraminide alpha-2,8-sialyltransferase 4" St8sia4
1430392_at 1.73779 RIKEN cDNA 9530086O07 gene 9530086O07Rik
1430393_at 1.72508 RIKEN cDNA 2510017J16 gene 2510017J16Rik
1430394_a_at 0.96405 "ATP-binding cassette, sub-family B (MDR/TAP), member 9" Abcb9
1430395_at 0.95781 ankyrin repeat domain 45 Ankrd45
1430396_at 0.93526 RIKEN cDNA 5730403I07 gene 5730403I07Rik
1430397_at 1.11601 "glutamate oxaloacetate transaminase 2, mitochondrial /// similar to Aspartate
aminotransferase, mitochondrial precursor (Transaminase A) (Glutamate oxaloacetate transaminase

2)" Got2 /// LOC640847

1430398_at	0.95203	RIKEN cDNA 4921517O11	gene	4921517O11Rik	
1430399_at	1.13566	RIKEN cDNA 2810458H16	gene	2810458H16Rik	
1430400_at	1.09158	RIKEN cDNA 4930521A18	gene	4930521A18Rik	
1430401_at	1.20621	RIKEN cDNA 3110045C21	gene	3110045C21Rik	
1430402_at	0.92816	"succinate-Coenzyme A ligase, ADP-forming, beta subunit"			Sucla2
1430403_at	1.31674	ubiquitin specific peptidase 32	Usp32		
1430404_at	2.01209	RIKEN cDNA 4833416J08	gene	4833416J08Rik	
1430405_at	0.89832	---	---		
1430406_at	0.91922	RIKEN cDNA 4930467M19	gene	4930467M19Rik	
1430407_at	0.76105	RIKEN cDNA 3110035C09	gene	3110035C09Rik	
1430408_at	0.93881	"calcium channel, voltage-dependent, P/Q type, alpha 1A subunit"			Cacna1a
1430409_at	0.98225	N-ethylmaleimide sensitive fusion protein	Nsf		
1430410_at	0.89166	RIKEN cDNA 1810019N24	gene	1810019N24Rik	
1430411_at	0.95732	RIKEN cDNA 6330525I24	gene	6330525I24Rik	
1430412_at	1.28442	RIKEN cDNA 2310068C19	gene	2310068C19Rik	
1430413_at	1.04221	transmembrane protein 29	Tmem29		
1430414_at	1.10576	RIKEN cDNA 5730406G12	gene	5730406G12Rik	
1430415_at	1.69190	PHD finger protein 6	Phf6		
1430416_at	1.01478	RIKEN cDNA 4931431B13	gene	4931431B13Rik	
1430417_s_at	0.99620	RIKEN cDNA 0610025P10	gene	0610025P10Rik	
1430418_at	1.49489	transmembrane protein 57	Tmem57		
1430419_at	1.02191	RIKEN cDNA 2310031A07	gene	2310031A07Rik	
1430420_at	0.97334	"asparagine-linked glycosylation 9 homolog (yeast, alpha 1,2 mannosyltransferase)"	Alg9		
1430421_a_at	1.05825	cDNA sequence BC010787	BC010787		
1430422_at	1.30126	RIKEN cDNA 1700020A23	gene	1700020A23Rik	
1430423_s_at	1.01761	RIKEN cDNA 2610528E23	gene /// FGF receptor activating protein 1		
			2610528E23Rik /// Frag1		
1430424_at	1.01334	RIKEN cDNA 4933433C11	gene	4933433C11Rik	
1430425_at	0.96482	sidekick homolog 2 (chicken)	Sdk2		
1430426_at	1.01156	RIKEN cDNA 4930540M05	gene	4930540M05Rik	
1430427_a_at	0.78777	protocadherin 18	Pcdh18		
1430428_at	1.22512	RIKEN cDNA 2310066F23	gene	2310066F23Rik	
1430429_at	0.95090	phosphatidylglycerophosphate synthase 1	Pgs1		
1430430_at	1.08223	RIKEN cDNA 6130401L20	gene	6130401L20Rik	
1430431_at	1.06458	RIKEN cDNA 4933407H18	gene	4933407H18Rik	
1430432_at	0.83115	tripartite motif-containing 42	Trim42		
1430433_at	0.94095	RIKEN cDNA 4933406J08	gene /// hypothetical protein LOC640572		
			4933406J08Rik /// LOC640572		
1430434_at	1.08333	transketolase-like 2	Tktl2		
1430435_at	0.84590	"AF4/FMR2 family, member 3"	Aff3		
1430436_at	0.94112	RIKEN cDNA 3321401G04	gene	3321401G04Rik	

1430437_a_at	0.93342	RIKEN cDNA 1300018I17 gene	1300018I17Rik	
1430438_at	1.14522	RIKEN cDNA 8430439J12 gene	8430439J12Rik	
1430439_at	0.94648	"multiple C2 domains, transmembrane 1"	Mctp1	
1430440_at	0.92519	RIKEN cDNA 4930442J19 gene	4930442J19Rik	
1430441_at	1.05115	RIKEN cDNA 4933438K21 gene	4933438K21Rik	
1430442_at	1.01177	nitric oxide synthase 1 (neuronal) adaptor protein	Nos1ap	
1430443_at	1.04670	annexin A10	Anxa10	
1430444_at	0.83518	RIKEN cDNA 0610006L08 gene	0610006L08Rik	
1430445_at	0.96674	A kinase (PRKA) anchor protein 13	Akap13	
1430446_at	1.01822	RIKEN cDNA 2310057B04 gene	2310057B04Rik	
1430447_a_at	0.96152	leukocyte-associated Ig-like receptor 1	Lair1	
1430448_at	1.01842	RIKEN cDNA 6720418B01 gene	6720418B01Rik	
1430449_at	0.99291	RIKEN cDNA C330002I19 gene	C330002I19Rik	
1430450_at	1.01699	RIKEN cDNA 2310004L02 gene	2310004L02Rik	
1430451_at	1.00250	RIKEN cDNA 4930481B07 gene	4930481B07Rik	
1430452_at	1.03709	"cytochrome P450, family 20, subfamily A, polypeptide 1"	Cyp20a1	
1430453_a_at	1.00813	Bcl2-like 2	Bcl2l2	
1430454_x_at	0.97510	Bcl2-like 2	Bcl2l2	
1430455_at	0.99633	olfactory receptor 701	Olf701	
1430456_at	0.91775	RIKEN cDNA 5830406C21 gene	5830406C21Rik	
1430457_at	0.93325	decay accelerating factor 2	Daf2	
1430458_at	0.97802	suppressor of Ty 6 homolog (S. cerevisiae)	Supt6h	
1430459_at	0.99024	RIKEN cDNA 4632428C04 gene	4632428C04Rik	
1430460_at	1.22593	RIKEN cDNA 5830410F13 gene	5830410F13Rik	
1430461_at	0.94414	RIKEN cDNA 1700120E14 gene	1700120E14Rik	
1430462_at	0.99439	RIKEN cDNA 2310002L09 gene	2310002L09Rik	
1430463_a_at	0.94648	"synuclein, alpha interacting protein (synphilin)"	Sncaip	
1430464_at	0.98606	RIKEN cDNA 9430021M05 gene	9430021M05Rik	
1430465_at	1.16627	---	---	
1430466_at	0.94361	RIKEN cDNA 2310005G13 gene	2310005G13Rik	
1430467_at	1.03058	RIKEN cDNA 4921511H03 gene	4921511H03Rik	
1430468_at	0.99141	RIKEN cDNA 2810449G22 gene	2810449G22Rik	
1430469_at	0.84766	tubulin tyrosine ligase like family 5	Tll5	
1430470_at	0.93519	RIKEN cDNA 5730411F24 gene	5730411F24Rik	
1430471_at	1.03558	RIKEN cDNA 9230110K08 gene	9230110K08Rik	
1430472_at	1.04465	armadillo repeat containing 1	Armcl	
1430473_at	0.94796	RIKEN cDNA 9530034A14 gene	9530034A14Rik	
1430474_a_at	1.00843	mitochondrial carrier homolog 2 (C. elegans)	Mtch2	
1430475_at	0.93514	---	---	
1430476_at	1.05303	signal recognition particle 54	Srp54	
1430477_s_at	0.90404	---	---	
1430478_at	0.96440	RIKEN cDNA 3110043A19 gene	3110043A19Rik	
1430479_at	0.97181	RIKEN cDNA 2010007H06 gene	2010007H06Rik	

1430480_at	0.83800	negative regulator of ubiquitin-like proteins 1	Nub1	
1430481_at	1.05682	RIKEN cDNA 4930545L23	gene	4930545L23Rik
1430482_at	0.94281	---	---	
1430483_a_at	0.99563	transmembrane protein 79	Tmem79	
1430484_at	0.83830	RIKEN cDNA 8430428J23	gene	8430428J23Rik
1430485_at	2.09424	RIKEN cDNA 3010009O07	gene	3010009O07Rik
1430486_at	0.97501	RAD51-like 1 (<i>S. cerevisiae</i>)	Rad5111	
1430487_at	1.03094	AU RNA binding protein/enoyl-coenzyme A hydratase		Auh
1430488_at	0.99461	RIKEN cDNA 1700066B19	gene	1700066B19Rik
1430489_at	0.94580	RIKEN cDNA 5430416G10	gene	5430416G10Rik
1430490_at	0.90493	---	---	
1430491_at	1.46176	block of proliferation 1	Bop1	
1430492_at	0.94863	phosphorylase kinase alpha 2	Phka2	
1430493_at	0.95088	RIKEN cDNA 2310061G22	gene	2310061G22Rik
1430494_at	1.15615	RIKEN cDNA 2010005J08	gene	2010005J08Rik
1430495_at	0.95347	inositol (myo)-1(or 4)-monophosphatase 1	Imp1	
1430496_at	1.15461	RIKEN cDNA 6720408I04	gene	6720408I04Rik
1430497_at	1.10718	retinoid X receptor alpha	Rxra	
1430498_at	1.17912	RIKEN cDNA 9130009I01	gene	9130009I01Rik
1430499_at	0.91813	RIKEN cDNA 1700058M13	gene	1700058M13Rik
1430500_s_at	0.96716	metaxin 2	Mtx2	
1430501_at	0.98574	RIKEN cDNA C030047K22	gene	C030047K22Rik
1430502_at	1.19340	RIKEN cDNA 9130206N08	gene	9130206N08Rik
1430503_at	1.05953	RIKEN cDNA 6330522J23	gene	6330522J23Rik
1430504_at	0.86047	RIKEN cDNA 1700108E19	gene	1700108E19Rik
1430505_at	1.07361	RIKEN cDNA 1110032O16	gene	1110032O16Rik
1430506_at	1.02824	RIKEN cDNA 8430406P12	gene	8430406P12Rik
1430507_at	0.95568	RIKEN cDNA 1700027J07	gene	1700027J07Rik
1430508_at	0.98718	RIKEN cDNA 1700084F23	gene	1700084F23Rik
1430509_at	0.79478	---	---	
1430510_at	0.88237	RIKEN cDNA B130050I23	gene	B130050I23Rik
1430511_at	0.95569	RIKEN cDNA 1500037O19	gene	1500037O19Rik
1430512_a_at	0.95886	immunoglobulin (CD79A) binding protein 1b	Igbp1b	
1430513_at	0.92602	RIKEN cDNA 1700055N04	gene	1700055N04Rik
1430514_a_at	0.93209	CD99 antigen	Cd99	
1430515_s_at	1.00344	aminoadipate-semialdehyde dehydrogenase-phosphopantetheinyl transferase		
Aasdhppt				
1430516_at	0.93722	RIKEN cDNA 4930428B01	gene	4930428B01Rik
1430517_at	1.00583	RIKEN cDNA 4930544O15	gene	4930544O15Rik
1430518_at	0.96383	RIKEN cDNA 5430402E10	gene	5430402E10Rik
1430519_a_at	0.99065	"CCR4-NOT transcription complex, subunit 7"	Cnot7	
1430520_at	0.93506	copine VIII	Cpne8	
1430521_s_at	0.92322	copine VIII	Cpne8	

1430522_a_at 1.09716 vesicle-associated membrane protein 5 Vamp5
1430523_s_at 0.94866 "immunoglobulin lambda chain, variable 1" Igl-V1
1430524_at 0.86563 RIKEN cDNA 1700020N15 gene /// hypothetical LOC627927
1700020N15Rik /// LOC627927
1430525_at 0.86614 RIKEN cDNA 2310001K24 gene 2310001K24Rik
1430526_a_at 0.94536 "SWI/SNF related, matrix associated, actin dependent regulator of chromatin, subfamily a, member 2" Smarca2
1430527_a_at 0.79827 ring finger protein 167 Rnf167
1430528_at 1.08528 syndecan binding protein Sdcbp
1430529_at 1.46459 "casein kinase 1, alpha 1" Csnk1a1
1430530_s_at 0.94292 NmrA-like family domain containing 1 Nmral1
1430531_at 1.03806 "biphenyl hydrolase-like (serine hydrolase, breast epithelial mucin-associated antigen)" Bphl
1430532_at 0.97519 dymeclin Dym
1430533_a_at 1.00822 "catenin (cadherin associated protein), beta 1" Ctnnb1
1430534_at 0.93453 "ribonuclease, RNase A family, 6" Rnase6
1430535_at 0.86933 TSC22 domain family 2 Tsc22d2
1430536_a_at 1.16628 enhancer of rudimentary homolog (Drosophila) Erh
1430537_at 0.95887 tRNA methyltransferase 12 homolog (S. cerevisiae) Trmt12
1430538_at 0.86458 RIKEN cDNA 2210013O21 gene 2210013O21Rik
1430539_at 1.01582 RIKEN cDNA 1810057P16 gene 1810057P16Rik
1430540_at 0.92685 coiled-coil-helix-coiled-coil-helix domain containing 8 Chchd8
1430541_at 0.96815 : F-box protein 44 Fbxo44
1430542_a_at 1.15428 "solute carrier family 25 (mitochondrial carrier, adenine nucleotide translocator), member 5 /// similar to ADP/ATP translocase 2 (Adenine nucleotide translocator 2) (ANT 2) (ADP,ATP carrier protein 2) (Solute carrier family 25 member 5) /// similar to solute carrier family 25, member 5" Slc25a5 /// LOC433923 /// LOC670371
1430543_at 1.05515 RIKEN cDNA 1500005P14 gene 1500005P14Rik
1430544_at 0.91124 RIKEN cDNA 5830404H04 gene 5830404H04Rik
1430545_at 1.10507 DBF4 homolog (S. cerevisiae) Dbf4
1430546_at 0.95998 "crystallin, zeta (quinone reductase)-like 1" Cryz11
1430547_s_at 1.06343 "crystallin, zeta (quinone reductase)-like 1" Cryz11
1430548_at 1.02758 --- ---
1430549_at 1.04001 blocked early in transport 1 homolog (S. cerevisiae)-like Bet11
1430550_at 0.85805 "lipase-like, ab-hydrolase domain containing 3" Lip13
1430551_s_at 1.05059 "lipase-like, ab-hydrolase domain containing 3" Lip13
1430552_a_at 0.94862 SET binding factor 1 Sbf1
1430553_at 0.98834 V-set and immunoglobulin domain containing 1 Vsig1
1430554_at 0.96949 leucine-rich repeats and immunoglobulin-like domains 3 Lrig3
1430555_s_at 0.82116 leucine-rich repeats and immunoglobulin-like domains 3 Lrig3
1430556_at 1.20474 sperm associated antigen 9 Spag9
1430557_at 0.91566 RIKEN cDNA 4930434J08 gene 4930434J08Rik
1430558_at 0.98247 zinc finger protein 318 Zfp318

1430559_at	1.01939	"5'-nucleotidase, cytosolic III-like"	Nt5c3l
1430560_at	1.09510	"protein phosphatase 1, regulatory subunit 10"	Ppp1r10
1430561_at	0.84579	"DnaJ (Hsp40) homolog, subfamily B, member 14"	Dnajb14
1430562_at	1.00497	zinc finger protein 95	Zfp95
1430563_at	0.98993	minichromosome maintenance deficient domain containing 1	Mcmdc1
1430564_at	1.00938	"MOB1, Mps One Binder kinase activator-like 1A (yeast)"	Mobk11a
1430565_at	0.84156	RIKEN cDNA 1700030J22 gene	1700030J22Rik
1430566_at	1.44306	RIKEN cDNA 4733401A01 gene	4733401A01Rik
1430567_at	1.04803	"serine peptidase inhibitor, Kazal type 5"	Spink5
1430568_at	0.99521	zinc finger CCCH type containing 13	Zc3h13
1430569_at	0.99782	tetratricopeptide repeat domain 9C	Ttc9c
1430570_at	0.91639	kynureninase (L-kynurenine hydrolase)	Kynu
1430571_s_at	1.01645	armadillo repeat containing 6	Armc6
1430572_at	1.15121	RIKEN cDNA 1810042K04 gene	1810042K04Rik
1430573_s_at	0.90016	ethanol decreased 2	Etohd2
1430574_at	1.01736	cyclin-dependent kinase inhibitor 3	Cdkn3
1430575_a_at	1.41028	tripeptidyl peptidase II	Tpp2
1430576_at	1.29776	tripeptidyl peptidase II	Tpp2
1430577_at	0.77249	centrosomal protein 27	Cep27
1430578_at	1.04658	"eukaryotic translation initiation factor 4 gamma, 3"	Eif4g3
1430579_at	0.63607	TRAF2 and NCK interacting kinase /// similar to [Segment 1 of 2] Traf2 and NCK-interacting kinase /// region containing TRAF2 and NCK interacting kinase; RIKEN cDNA C630040K21 gene Tnik /// LOC665113 /// LOC674765	
1430580_at	0.95158	RIKEN cDNA 2810030E01 gene	2810030E01Rik
1430581_at	0.99161	---	---
1430582_at	0.90202	SNF2 histone linker PHD RING helicase	Shprh
1430583_at	0.92159	RIKEN cDNA C130076O07 gene	C130076O07Rik
1430584_s_at	0.99614	carbonic anhydrase 3	Car3
1430585_at	1.95957	RIKEN cDNA 5930436O19 gene	5930436O19Rik
1430586_at	1.05170	RIKEN cDNA 2700007P21 gene	2700007P21Rik
1430587_at	1.02266	RIKEN cDNA 4933425L03 gene	4933425L03Rik
1430588_at	0.89348	Maestro	Mro
1430589_at	0.98852	"phosphatidylinositol glycan anchor biosynthesis, class H"	Pigh
1430590_at	0.93450	"DNA segment, Chr 3, ERATO Doi 751, expressed"	D3Ertd751e
1430591_at	0.97332	dopa decarboxylase	Ddc
1430592_at	1.13626	transmembrane protein 106B	Tmem106b
1430593_at	0.99357	RIKEN cDNA 1700041C02 gene	1700041C02Rik
1430594_at	0.97083	RAB11 family interacting protein 1 (class I)	Rab11fip1
1430595_at	0.91711	growth arrest-specific 2 like 1	Gas2l1
1430596_s_at	0.88205	RIKEN cDNA 1700110N18 gene	1700110N18Rik
1430597_at	0.99859	janus kinase and microtubule interacting protein 1	Jakmip1
1430598_at	1.01119	cAMP responsive element modulator	Crem
1430599_at	1.07750	myelin transcription factor 1-like	Myt1l

1430600_at	0.79921 CKLF-like MARVEL transmembrane domain containing 5	Cmtm5
1430601_at	0.92395 TBCC domain containing 1	Tbccd1
1430602_at	0.98915 RIKEN cDNA 4930520K10 gene	4930520K10Rik
1430603_at	0.96014 RIKEN cDNA 4930579K19 gene	4930579K19Rik
1430604_a_at	0.92270 disabled homolog 2 (Drosophila)	Dab2
1430605_at	1.02985 coiled-coil domain containing 101	Ccdc101
1430606_at	1.00743 RIKEN cDNA 2310020F24 gene	2310020F24Rik
1430607_at	0.99206 RIKEN cDNA 4930505D03 gene	4930505D03Rik
1430608_at	1.10661 RIKEN cDNA 4930535I16 gene	4930535I16Rik
1430609_at	1.10040 MORN repeat containing 1	Morn1
1430610_at	1.05580 mitochondrial ribosomal protein L38	Mrpl38
1430611_at	0.97815 "zinc finger, DHHC domain containing 25"	Zdhhc25
1430612_at	1.06423 RIKEN cDNA 1810033B17 gene	1810033B17Rik
1430613_at	1.03652 RIKEN cDNA 1520402A15 gene	1520402A15Rik
1430614_at	0.90415 RIKEN cDNA 4632415K11 gene	4632415K11Rik
1430615_at	1.05662 RIKEN cDNA 4921517B04 gene	4921517B04Rik
1430616_at	0.88147 RIKEN cDNA 4930528A17 gene	4930528A17Rik
1430617_at	0.96827 Opa interacting protein 5	Oip5
1430618_at	1.20517 RIKEN cDNA 2610020C07 gene	2610020C07Rik
1430619_a_at	1.10810 mevalonate kinase /// similar to mevalonate kinase	Mvk /// LOC637711
1430620_at	0.94786 RIKEN cDNA 4930588N13 gene	4930588N13Rik
1430621_at	1.01313 multimerin 1	Mmrn1
1430622_at	1.84174 RIKEN cDNA 4833423F13 gene	4833423F13Rik
1430623_s_at	1.18202 RIKEN cDNA 5830411E10 gene	5830411E10Rik
1430624_at	0.93695 RIKEN cDNA 4921511M17 gene /// hypothetical LOC434874 /// hypothetical LOC434879 /// hypothetical LOC546397 /// hypothetical LOC625650 /// hypothetical LOC625679 /// hypothetical LOC625713 /// hypothetical protein LOC635502 /// hypothetical protein LOC635523 /// hypothetical protein LOC635580 /// hypothetical protein LOC635709 /// similar to RIKEN cDNA 4921511M17 gene /// hypothetical protein LOC667285 /// hypothetical protein LOC670713 /// hypothetical protein LOC673177 4921511M17Rik /// LOC434874 /// LOC434879 /// LOC546397 /// LOC625650 /// LOC625679 /// LOC625713 /// LOC635502 /// LOC635523 /// LOC635580 /// LOC635709 /// LOC667256 /// LOC667285 /// LOC670713 /// LOC673177	
1430625_at	0.98796 keratin associated protein 3-2	Krtap3-2
1430626_at	1.11748 zinc finger protein 398	Zfp398
1430627_at	1.06288 RIKEN cDNA 5330420P17 gene	5330420P17Rik
1430628_at	0.96881 jumonji domain containing 2C	Jmjd2c
1430629_at	1.05254 "solute carrier family 16 (monocarboxylic acid transporters), member 14"	Slc16a14
1430630_at	1.14236 --- ---	
1430631_at	1.05798 protein phosphatase 1F (PP2C domain containing)	Ppm1f
1430632_at	1.01986 ankyrin repeat and SOCS box-containing protein 1	Asb1
1430633_s_at	1.38239 RIKEN cDNA C430045I18 gene	C430045I18Rik
1430634_a_at	0.94389 "phosphofructokinase, platelet"	Pfkp

1430635_at	1.37470 melan-A Mlana	
1430636_at	1.22879 RIKEN cDNA C030010B13 gene	C030010B13Rik
1430637_at	1.01449 RIKEN cDNA 2210016H18 gene	2210016H18Rik
1430638_at	1.01077 RIKEN cDNA 1700041C02 gene	1700041C02Rik
1430639_at	1.04134 RIKEN cDNA 4930524L23 gene	4930524L23Rik
1430640_a_at	0.93414 "protein kinase, cAMP dependent regulatory, type II beta"	Prkar2b
1430641_at	0.89046 RIKEN cDNA 9030605I04 gene /// expressed sequence AI987692	
	9030605I04Rik /// AI987692	
1430642_at	0.96998 RIKEN cDNA 2900001G08 gene	2900001G08Rik
1430643_at	0.91361 mutS homolog 3 (E. coli)	Msh3
1430644_at	0.87216 Williams Beuren syndrome chromosome region 25 (human)	Wbscr25
1430645_at	0.96797 CD200 receptor 3	Cd200r3
1430646_at	1.19247 RIKEN cDNA 5830407P18 gene	5830407P18Rik
1430647_at	1.09601 cytoskeleton associated protein 5	Ckap5
1430648_at	1.18251 "sodium channel, voltage-gated, type II, beta"	Scn2b
1430649_at	0.98930 "brain and acute leukemia, cytoplasmic"	Baalc
1430650_at	0.94458 zinc finger protein 191	Zfp191
1430651_s_at	0.92715 zinc finger protein 191	Zfp191
1430652_at	0.99873 RIKEN cDNA 9030612E09 gene	9030612E09Rik
1430653_at	1.43322 RIKEN cDNA A930026B05 gene	A930026B05Rik
1430654_at	1.12381 NOL1/NOP2/Sun domain family 3	Nsun3
1430655_at	0.86330 RIKEN cDNA 4631405K08 gene	4631405K08Rik
1430656_a_at	1.00357 asparagine synthetase domain containing 1	Asnsd1
1430657_at	1.05365 RIKEN cDNA 1810034E14 gene	1810034E14Rik
1430658_a_at	1.07278 gasdermin 2	Gsdm2
1430659_at	0.95617 RIKEN cDNA 4930548H24 gene	4930548H24Rik
1430660_at	0.99838 RIKEN cDNA 5830417I10 gene	5830417I10Rik
1430661_at	1.11015 "myosin, light polypeptide 6B"	Myl6b
1430662_at	0.90167 RIKEN cDNA 9430091E24 gene	9430091E24Rik
1430663_at	1.32433 RIKEN cDNA 9430037O13 gene	9430037O13Rik
1430664_at	0.95815 zinc finger protein 169	Zfp169
1430665_at	1.10764 RIKEN cDNA 5730480H06 gene	5730480H06Rik
1430666_at	0.93069 similar to phospholipase B	LOC665270
1430667_at	0.94896 protocadherin 10	Pcdh10
1430668_a_at	0.82295 "ankyrin repeat, family A (RFXANK-like), 2"	Ankra2
1430669_at	1.01920 keratin associated protein 4-7	Krtap4-7
1430670_at	1.03417 coiled-coil domain containing 91	Ccdc91
1430671_a_at	0.89987 RIKEN cDNA 0610010K06 gene	0610010K06Rik
1430672_at	1.84795 RIKEN cDNA 5033418A18 gene	5033418A18Rik
1430673_a_at	0.97466 RIKEN cDNA 2210009G21 gene	2210009G21Rik
1430674_at	1.01270 RIKEN cDNA 1700016C15 gene	1700016C15Rik
1430675_at	0.96433 RIKEN cDNA 2900055J20 gene	2900055J20Rik
1430676_at	0.93296 "procollagen, type XIX, alpha 1"	Col19a1

1430677_at 0.96162 RIKEN cDNA 4930442L01 gene 4930442L01Rik
1430678_at 0.99118 RIKEN cDNA 5730457N03 gene 5730457N03Rik
1430679_at 1.02687 RIKEN cDNA 4933404O12 gene 4933404O12Rik
1430680_a_at 1.11573 RIKEN cDNA 5830417I10 gene /// similar to Dingo protein isoform 2 ///
similar to Dingo protein isoform 2 5830417I10Rik /// LOC673610 /// LOC677582
1430681_at 0.93944 "crystallin, lamda 1" Cryl1
1430682_at 0.80420 RIKEN cDNA 4930555I21 gene 4930555I21Rik
1430683_at 0.95851 dedicator of cytokinesis 6 /// similar to Dedicator of cytokinesis protein 6
Dock6 /// LOC670024
1430684_s_at 0.77527 Lrp2 binding protein Lrp2bp
1430685_at 0.92199 RIKEN cDNA 6330503C03 gene 6330503C03Rik
1430686_at 0.89386 RIKEN cDNA 4833418N02 gene 4833418N02Rik
1430687_at 1.00122 RIKEN cDNA 6330545A04 gene 6330545A04Rik
1430688_s_at 0.77672 RIKEN cDNA D730039F16 gene D730039F16Rik
1430689_at 0.95636 RIKEN cDNA 2810019C22 gene 2810019C22Rik
1430690_at 1.21553 RIKEN cDNA 5430411C19 gene 5430411C19Rik
1430691_at 0.88102 RIKEN cDNA 4632411P08 gene 4632411P08Rik
1430692_a_at 0.95895 Sell1 (suppressor of lin-12) 1 homolog (C. elegans) Sellh
1430693_at 0.99892 patatin-like phospholipase domain containing 5 Pnpla5
1430694_at 0.91626 --- ---
1430695_at 0.91892 toll-like receptor 4 Tlr4
1430696_at 1.05752 eukaryotic translation initiation factor 2 alpha kinase 1 Eif2ak1
1430697_at 1.16859 "Alport syndrome, mental retardation, midface hypoplasia and elliptocytosis
chromosomal region gene 1 homolog (human)" Ammecr1
1430698_a_at 0.92485 "DNA segment, Chr 14, ERATO Doi 581, expressed" D14Ert581e
1430699_at 1.18646 G protein-coupled receptor 165 Gpr165
1430700_a_at 0.99148 "phospholipase A2, group VII (platelet-activating factor acetylhydrolase,
plasma)" Pla2g7
1430701_a_at 1.03205 RIKEN cDNA 5730528L13 gene 5730528L13Rik
1430702_at 2.14875 RIKEN cDNA 5830427D03 gene 5830427D03Rik
1430703_at 0.96109 --- ---
1430704_at 0.90816 interleukin-1 receptor-associated kinase 3 Irak3
1430705_at 1.47765 RIKEN cDNA 2410104I19 gene 2410104I19Rik
1430706_at 1.00520 RIKEN cDNA 1700081D17 gene 1700081D17Rik
1430707_s_at 0.98754 RIKEN cDNA 1700081D17 gene 1700081D17Rik
1430708_a_at 0.93702 ubiquitin specific petidase 45 Usp45
1430709_at 0.99132 RIKEN cDNA 4833405L11 gene 4833405L11Rik
1430710_at 1.06097 --- ---
1430711_at 1.01238 RIKEN cDNA 4933416E03 gene 4933416E03Rik
1430712_at 0.77846 Rho GTPase activating protein 24 Arhgap24
1430713_s_at 0.87787 "NADH dehydrogenase (ubiquinone) 1 alpha subcomplex, 13" Ndufa13
1430714_at 1.04077 RIKEN cDNA 4930451C15 gene 4930451C15Rik
1430715_at 0.89578 RIKEN cDNA 2310014L17 gene 2310014L17Rik

1430716_at	0.98268	RIKEN cDNA 5330421F07	gene	5330421F07Rik
1430717_at	0.93102	RIKEN cDNA 4930403C10	gene	4930403C10Rik
1430718_s_at	1.11092	"zinc finger, CSL-type containing 3"		Zcs13
1430719_at	0.91997	RIKEN cDNA 4833447P13	gene	4833447P13Rik
1430720_at	1.24374	Y box protein 1		Ybx1
1430721_at	0.91014	RIKEN cDNA 4931431C16	gene	4931431C16Rik
1430722_at	0.91777	RIKEN cDNA 4921515J06	gene	4921515J06Rik
1430723_at	0.98061	"prenyl (solanesyl) diphosphate synthase, subunit 2"		Pdss2
1430724_at	0.90458	abhydrolase domain containing 9		Abhd9
1430725_at	0.92953	RIKEN cDNA 4933432I09	gene	4933432I09Rik
1430726_at	1.06027	Ras association (RalGDS/AF-6) domain family 8		Rassf8
1430727_at	0.99844	RIKEN cDNA 4930432O21	gene	4930432O21Rik
1430728_at	0.87830	keratin associated protein 5-5		Krtap5-5
1430729_at	0.91020	melanocyte proliferating gene 1		Myg1
1430730_at	1.11547	RIKEN cDNA 9430024F10	gene	9430024F10Rik
1430731_at	0.95874	keratin associated protein 2-4		Krtap2-4
1430732_at	0.84770	RIKEN cDNA 4921525D07	gene	4921525D07Rik
1430733_at	1.10226	RIKEN cDNA 4932441P12	gene	4932441P12Rik
1430734_at	1.08281	cytochrome b5 reductase 3		Cyb5r3
1430735_at	0.99315	RIKEN cDNA 4930424G05	gene	4930424G05Rik
1430736_at	1.01186	"RIKEN cDNA 9030411M15	gene	/// similar to CG8486-PA, isoform A
		similar to CG8486-PA, isoform A"		9030411M15Rik
				/// LOC629416
				/// LOC667742
1430737_at	1.01366	RIKEN cDNA 2810409K11	gene	2810409K11Rik
1430738_at	0.92820	RIKEN cDNA 4833419K08	gene	4833419K08Rik
1430739_at	0.98787	RIKEN cDNA 4930471M09	gene	4930471M09Rik
1430740_at	0.91247	---		---
1430741_at	0.98981	RIKEN cDNA 4933433G08	gene	4933433G08Rik
1430742_a_at	0.81148	spermatogenesis associated 16		Spata16
1430743_at	1.03083	RIKEN cDNA 4930412F12	gene	4930412F12Rik
1430744_at	0.98765	napsin A aspartic peptidase		Napsa
1430745_at	0.93588	RIKEN cDNA 5930409G06	gene	5930409G06Rik
1430746_at	1.05063	RIKEN cDNA 1110050P16	gene	1110050P16Rik
1430747_at	1.03870	PQ loop repeat containing 1		Pqlc1
1430748_at	0.93460	RIKEN cDNA 1810063B07	gene	1810063B07Rik
1430749_at	1.08359	RIKEN cDNA 2810040C05	gene	2810040C05Rik
1430750_at	1.01071	dihydrofolate reductase		Dhfr
1430751_at	0.89243	---		---
1430752_at	1.12014	RIKEN cDNA C330006D17	gene	C330006D17Rik
1430753_at	0.86546	RIKEN cDNA 8030491N06	gene	8030491N06Rik
1430754_at	1.10553	RIKEN cDNA 4930511H11	gene	4930511H11Rik
1430755_at	0.98864	RIKEN cDNA 4930452G13	gene	4930452G13Rik
1430756_at	1.07511	RIKEN cDNA 5430427G11	gene	5430427G11Rik
1430757_at	1.07023	RIKEN cDNA A330102H22	gene	A330102H22Rik

1430758_x_at 0.94847 breast carcinoma amplified sequence 1 Bcas1
1430759_at 1.00113 RIKEN cDNA 4930523C07 gene 4930523C07Rik
1430760_a_at 0.92262 ubiquitin specific peptidase 50 Usp50
1430761_at 1.01029 RIKEN cDNA 0610039K10 gene 0610039K10Rik
1430762_at 1.06062 RIKEN cDNA 4833427G06 gene 4833427G06Rik
1430763_at 0.95868 RIKEN cDNA 4930563E22 gene 4930563E22Rik
1430764_at 0.91421 RIKEN cDNA 1700023F06 gene 1700023F06Rik
1430765_at 1.15482 RIKEN cDNA 5730521K06 gene 5730521K06Rik
1430766_at 0.92697 RIKEN cDNA 5033403F01 gene 5033403F01Rik
1430767_a_at 1.02101 RIKEN cDNA 4930543E12 gene 4930543E12Rik
1430768_at 0.88694 RIKEN cDNA 9530018H14 gene 9530018H14Rik
1430769_s_at 0.94551 RIKEN cDNA 2900009I07 gene 2900009I07Rik
1430770_at 0.95809 RIKEN cDNA 3110080E11 gene 3110080E11Rik
1430771_a_at 1.06276 mutS homolog 5 (E. coli) Msh5
1430772_at 0.80254 --- ---
1430773_a_at 1.09351 t-complex-associated testis expressed 2 Tcte2
1430774_at 1.66431 RIKEN cDNA A430106A12 gene A430106A12Rik
1430775_at 0.99540 WD repeat domain 44 Wdr44
1430776_s_at 0.92969 ankyrin repeat domain 24 Ankrd24
1430777_a_at 0.93827 golgi phosphoprotein 3 Golph3
1430778_a_at 1.19278 nucleotide binding protein 1 Nubp1
1430779_at 0.83520 netrin G1 Ntng1
1430780_a_at 0.90265 phosphomannomutase 1 Pmm1
1430781_at 0.92019 adenylate kinase 7 Ak7
1430782_at 1.18633 RIKEN cDNA B130055D15 gene B130055D15Rik
1430783_at 0.96204 RIKEN cDNA 4932416J16 gene 4932416J16Rik
1430784_a_at 1.06329 RIKEN cDNA 4932417H02 gene 4932417H02Rik
1430785_at 1.01960 orphan short chain dehydrogenase/reductase Sdro
1430786_at 1.07684 RIKEN cDNA 1110002E22 gene 1110002E22Rik
1430787_at 1.06200 RIKEN cDNA 2310050B05 gene 2310050B05Rik
1430788_at 1.02699 --- ---
1430789_at 1.04188 RIKEN cDNA 4930452B06 gene 4930452B06Rik
1430790_at 1.03175 RIKEN cDNA 4930485G23 gene 4930485G23Rik
1430791_at 0.94193 RIKEN cDNA 6330415B21 gene 6330415B21Rik
1430792_at 1.43133 --- ---
1430793_at 0.93571 RIKEN cDNA 4931429I11 gene 4931429I11Rik
1430794_at 1.06650 RIKEN cDNA 4933426K07 gene 4933426K07Rik
1430795_at 1.04332 RIKEN cDNA 5830407F19 gene 5830407F19Rik
1430796_at 1.13651 tripartite motif-containing 14 Trim14
1430797_at 1.10694 exportin 4 Xpo4
1430798_x_at 1.13059 mitochondrial ribosomal protein L15 Mrpl15
1430799_at 0.99156 RIKEN cDNA 5830432E09 gene 5830432E09Rik
1430800_at 0.97058 RIKEN cDNA 4930518F22 gene 4930518F22Rik

1430801_at 0.97311 RIKEN cDNA 2310042E16 gene /// vacuolar protein sorting 13B (yeast) /// similar to vacuolar protein sorting 13B isoform 4 /// region containing RIKEN cDNA C330002D13 gene; RIKEN cDNA 2310042E16 gene; vacuolar protein sorting 13B (yeast) 2310042E16Rik /// Vps13b /// LOC666173 /// LOC669978
 1430802_at 1.04606 "histocompatibility 2, Q region locus 8" H2-Q8
 1430803_at 0.97664 lin-28 homolog B (C. elegans) Lin28b
 1430804_at 0.98983 "solute carrier family 13 (sodium/sulphate symporters), member 1" Slc13a1
 1430805_s_at 1.00129 "RMI1, RecQ mediated genome instability 1, homolog (S. cerevisiae)" Rmi1
 1430806_at 0.91132 RIKEN cDNA 2810047F03 gene 2810047F03Rik
 1430807_at 0.95530 kelch-like 22 (Drosophila) Klh22
 1430808_at 1.28199 "TBC1 domain family, member 5" Tbc1d5
 1430809_at 1.02021 Frizzled homolog 5 (Drosophila) Fzd5
 1430810_at 0.93863 RIKEN cDNA 5730409E15 gene 5730409E15Rik
 1430811_a_at 0.98069 cell division cycle associated 1 Cdca1
 1430812_at 0.92963 RIKEN cDNA 4930511A02 gene 4930511A02Rik
 1430813_at 0.96236 RIKEN cDNA 1810057I13 gene 1810057I13Rik
 1430814_at 1.12880 "RIKEN cDNA 1300013D18 gene /// similar to cytochrome P450, family 2, subfamily d, polypeptide 13" 1300013D18Rik /// LOC271300
 1430815_at 0.79323 RIKEN cDNA 4930534B04 gene 4930534B04Rik
 1430816_at 0.98448 RIKEN cDNA 3110003A22 gene 3110003A22Rik
 1430817_at 0.99514 sterile alpha motif domain containing 7 Samd7
 1430818_at 1.09363 transmembrane channel-like gene family 1 Tmc1
 1430819_at 1.01024 N-acylsphingosine amidohydrolase (acid ceramidase)-like Asah1
 1430820_a_at 1.06943 bobby sox homolog (Drosophila) Bbx
 1430821_at 1.08034 RIKEN cDNA 1700024C24 gene 1700024C24Rik
 1430822_at 1.07437 "SWI/SNF related, matrix associated, actin dependent regulator of chromatin, subfamily e, member 1" Smarce1
 1430823_at 0.97591 RIKEN cDNA 2700029L08 gene 2700029L08Rik
 1430824_at 1.03758 Rho GTPase activating protein 23 Arhgap23
 1430825_at 1.17170 RIKEN cDNA E130102C15 gene E130102C15Rik
 1430826_s_at 1.39918 "glucosaminyl (N-acetyl) transferase 2, I-branching enzyme" Gcnt2
 1430827_a_at 1.06165 PTK2 protein tyrosine kinase 2 Ptk2
 1430828_at 1.07475 tRNA 5-methylaminomethyl-2-thiouridylate methyltransferase Trmu
 1430829_s_at 0.94185 fatso Fto
 1430830_at 1.09546 RIKEN cDNA A430105J06 gene A430105J06Rik
 1430831_at 0.96695 RIKEN cDNA 4930404J24 gene 4930404J24Rik
 1430832_at 0.92317 RIKEN cDNA 4930426I24 gene 4930426I24Rik
 1430833_at 0.90999 IL2-inducible T-cell kinase Itk
 1430834_at 1.04210 --- ---
 1430835_at 1.07465 RIKEN cDNA 5830436D01 gene 5830436D01Rik
 1430836_at 0.95650 RIKEN cDNA 4933437G19 gene 4933437G19Rik
 1430837_a_at 0.90020 methyl-CpG binding domain protein 1 Mbd1
 1430838_x_at 0.89594 methyl-CpG binding domain protein 1 Mbd1

1430839_at	0.96535	RIKEN cDNA 9430076G02	gene	9430076G02Rik
1430840_at	0.98303	RIKEN cDNA 4930449C09	gene	4930449C09Rik
1430841_at	1.10884	RIKEN cDNA 3300002P13	gene	3300002P13Rik
1430842_at	0.96457	RIKEN cDNA 1500002F19	gene	1500002F19Rik
1430843_at	1.07132	glutamate decarboxylase-like 1	Gad1l	
1430844_at	0.88887	RIKEN cDNA 1700018M17	gene	1700018M17Rik
1430845_at	0.93510	RIKEN cDNA 2410018L13	gene	2410018L13Rik
1430846_at	0.88004	RIKEN cDNA 1700061G19	gene	1700061G19Rik
1430847_a_at	1.05793	cAMP responsive element modulator	Crem	
1430848_a_at	0.94585	RIKEN cDNA 2410017P07	gene	2410017P07Rik
1430849_a_at	0.89700	cell growth regulator with ring finger domain 1	Cgrrf1	
1430850_x_at	1.06772	cell growth regulator with ring finger domain 1	Cgrrf1	
1430851_at	1.08461	zinc finger protein 444	Zfp444	
1430852_at	1.09815	RIKEN cDNA 1600029I14	gene	1600029I14Rik
1430853_a_at	1.07037	RIKEN cDNA 3110045G13	gene	3110045G13Rik
1430854_at	0.90607	"asparagine-linked glycosylation 5 homolog (yeast, dolichyl-phosphate beta-glucosyltransferase)"	Alg5	
1430855_at	0.86176	"collagen, type XX, alpha 1"	Col20a1	
1430856_at	0.87873	peroxisomal biogenesis factor 11c	Pex11c	
1430857_s_at	1.03879	peroxisomal biogenesis factor 11c	Pex11c	
1430858_at	1.00716	RIKEN cDNA 4632433K11	gene	4632433K11Rik
1430859_at	0.95080	RIKEN cDNA 5330427O13	gene	5330427O13Rik
1430860_at	1.10878	RIKEN cDNA 4833422M21	gene	4833422M21Rik
1430861_at	1.03471	RIKEN cDNA 2010111I01	gene	2010111I01Rik
1430862_at	1.11349	RIKEN cDNA 2310057M21	gene	2310057M21Rik
1430863_at	0.98788	RIKEN cDNA 4933411B09	gene	4933411B09Rik
1430864_at	0.85176	tubulin tyrosine ligase like family 9	Ttl19	
1430865_s_at	1.12210	tubulin tyrosine ligase like family 9	Ttl19	
1430866_at	0.91090	RIKEN cDNA 4921537D05	gene	4921537D05Rik
1430867_at	0.97989	RIKEN cDNA 4933431M02	gene	4933431M02Rik
1430868_at	0.95001	RIKEN cDNA 5830403M04	gene	5830403M04Rik
1430869_a_at	0.93765	homeo box C13 /// hyaluronic acid binding protein 4 /// similar to hyaluronic acid binding protein 4	Hoxc13 /// Habp4 /// LOC671769	
1430870_at	0.87177	RIKEN cDNA 4930442P19	gene	4930442P19Rik
1430871_at	1.04903	"phospholipase C, beta 4"	Plcb4	
1430872_at	1.07507	RIKEN cDNA 4930412O13	gene	4930412O13Rik
1430873_at	1.16592	---	---	
1430874_at	1.01239	RIKEN cDNA 9430085M18	gene	9430085M18Rik
1430875_a_at	1.07918	PAK1 interacting protein 1	Pak1ip1	
1430876_at	1.01454	RIKEN cDNA 2610312I10	gene	2610312I10Rik
1430877_at	1.04964	RIKEN cDNA 8030425K09	gene	8030425K09Rik
1430878_at	1.01893	RIKEN cDNA 2210406H18	gene	2210406H18Rik
1430879_at	1.03454	RIKEN cDNA C030008P14	gene	C030008P14Rik

1430880_at	0.88930	pleckstrin and Sec7 domain containing 3	Psd3
1430881_at	0.96800	RIKEN cDNA 7630403G23	gene 7630403G23Rik
1430882_at	0.86819	---	---
1430883_at	1.01461	RIKEN cDNA 4933402C05	gene 4933402C05Rik
1430884_at	0.98311	RIKEN cDNA 4930568G15	gene 4930568G15Rik
1430885_at	0.91758	RIKEN cDNA 1110012N22	gene 1110012N22Rik
1430886_at	0.90059	RIKEN cDNA 1700112E06	gene 1700112E06Rik
1430887_s_at	1.04342	RIKEN cDNA 1700112E06	gene 1700112E06Rik
1430888_at	0.99614	RIKEN cDNA 4930535E21	gene 4930535E21Rik
1430889_a_at	0.97252	thiopurine methyltransferase	Tpmt
1430890_at	1.05850	RIKEN cDNA 2210010C17	gene 2210010C17Rik
1430891_at	0.88274	dual specificity phosphatase 12	Dusp12
1430892_at	0.99099	RIKEN cDNA 1700011F03	gene 1700011F03Rik
1430893_at	1.11498	major urinary protein 1	Mup1
1430894_at	0.98230	RIKEN cDNA 5530401N12	gene 5530401N12Rik
1430895_at	0.89494	RIKEN cDNA 2010109A12	gene 2010109A12Rik
1430896_s_at	1.07791	nudix (nucleoside diphosphate linked moiety X)-type motif 7	Nudt7
1430897_at	0.95199	RIKEN cDNA 4931428L18	gene 4931428L18Rik
1430898_s_at	1.02584	RIKEN cDNA 4931428L18	gene 4931428L18Rik
1430899_at	0.97916	"mucin 5, subtypes A and C, tracheobronchial/gastric"	Muc5ac
1430900_at	0.95148	RIKEN cDNA 4930430O22	gene 4930430O22Rik
1430901_at	1.02354	RIKEN cDNA 1700003F17	gene 1700003F17Rik
1430902_at	1.06535	ankyrin repeat domain 24	Ankrd24
1430903_at	0.89049	RIKEN cDNA 4933416O17	gene 4933416O17Rik
1430904_at	1.08887	ADP-ribosylation factor GTPase activating protein 3	Arfgap3
1430905_at	1.04228	RIKEN cDNA 4921504E06	gene 4921504E06Rik
1430906_at	1.00018	RIKEN cDNA 2510016G02	gene 2510016G02Rik
1430907_at	1.05981	RIKEN cDNA 4930455J16	gene 4930455J16Rik
1430908_at	0.94016	RIKEN cDNA 4933429O19	gene 4933429O19Rik
1430909_at	0.88563	nucleoporin 210-like	Nup210l
1430910_at	1.01328	RIKEN cDNA 4930544L04	gene 4930544L04Rik
1430911_at	1.06543	"DNA segment, Chr 19, ERATO Doi 652, expressed"	D19Ertd652e
1430912_a_at	1.04426	tectorin beta	Tectb
1430913_at	0.85293	RIKEN cDNA 1810033M07	gene 1810033M07Rik
1430914_at	1.02970	RIKEN cDNA 5330426L24	gene 5330426L24Rik
1430915_at	0.93798	fibronectin type III domain containing 8	Fndc8
1430916_at	1.32397	"Fanconi anemia, complementation group D2"	Fancd2
1430917_at	0.84658	RIKEN cDNA 9130401L11	gene 9130401L11Rik
1430918_at	0.98232	membrane-associated ring finger (C3HC4) 3	3-Mar
1430919_at	0.90334	RIKEN cDNA 4930525F21	gene 4930525F21Rik
1430920_at	1.18157	RIKEN cDNA 2410075D05	gene 2410075D05Rik
1430921_at	0.89747	RIKEN cDNA 9130015G15	gene 9130015G15Rik
1430922_at	1.02653	RIKEN cDNA 5530401A14	gene 5530401A14Rik

1430923_at	1.09819	RIKEN cDNA 2610300M13	gene	2610300M13Rik
1430924_at	0.98058	RIKEN cDNA 4933440M02	gene	4933440M02Rik
1430925_at	1.02782	RIKEN cDNA 4930556L07	gene	4930556L07Rik
1430926_at	1.04189	tetratricopeptide repeat domain 18		Ttc18
1430927_at	1.02156	CEA-related cell adhesion molecule 1		Ceacam18
1430928_at	0.99159	RIKEN cDNA 2900002J02	gene	2900002J02Rik
1430929_at	1.14629	RIKEN cDNA 9430082L08	gene	9430082L08Rik
1430930_at	1.01708	RIKEN cDNA 0610038B21	gene	0610038B21Rik
1430931_at	0.90162	RIKEN cDNA 4933403O03	gene	4933403O03Rik
1430932_at	1.29647	"solute carrier family 9 (sodium/hydrogen exchanger), member 8"		Slc9a8
1430933_at	0.85976	RIKEN cDNA 5730433K22	gene	5730433K22Rik
1430934_at	0.83782	RIKEN cDNA 2810043G22	gene	2810043G22Rik
1430935_at	1.10825	RIKEN cDNA 4933413G19	gene	4933413G19Rik
1430936_at	1.10436	"CHMP family, member 7"		Chmp7
1430937_at	1.14060	RIKEN cDNA A930016O22	gene	A930016O22Rik
1430938_at	1.00191	zinc finger protein 64		Zfp64
1430939_at	0.95748	RIKEN cDNA 4930528D03	gene	4930528D03Rik
1430940_at	1.68694	RIKEN cDNA 3110045A19	gene	3110045A19Rik
1430941_at	1.12212	Bardet-Biedl syndrome 7		Bbs7
1430942_at	1.35624	RIKEN cDNA 8430437O03	gene	8430437O03Rik
1430943_at	0.95786	RIKEN cDNA 6030498E09	gene	6030498E09Rik
1430944_at	1.09932	RIKEN cDNA A930030B08	gene	A930030B08Rik
1430945_at	0.92311	RIKEN cDNA A930028C08	gene	A930028C08Rik
1430946_at	1.04694	RIKEN cDNA 2600014E21	gene	2600014E21Rik
1430947_at	0.90184	RIKEN cDNA 4930549C15	gene	4930549C15Rik
1430948_at	0.95978	RIKEN cDNA 4921525B02	gene	4921525B02Rik
1430949_at	0.93049	RIKEN cDNA 9130231A04	gene	9130231A04Rik
1430950_at	0.99554	odd-skipped related 2 (Drosophila)		Osr2
1430951_at	0.99925	RIKEN cDNA 2810011L19	gene	2810011L19Rik
1430952_at	1.03472	RIKEN cDNA 3110039C02	gene	3110039C02Rik
1430953_at	1.05257	RIKEN cDNA 3200001G23	gene	3200001G23Rik
1430954_at	1.00934	peptidylprolyl isomerase (cyclophilin)-like 3		Ppil3
1430955_at	1.05871	RIKEN cDNA 2810403A07	gene	2810403A07Rik
1430956_at	0.95504	"Protease, serine, 23"		Prss23
1430957_at	0.99721	Rho GTPase activating protein 26		Arhgap26
1430958_at	1.07430	RIKEN cDNA 4933430N04	gene	4933430N04Rik
1430959_at	0.90617	RIKEN cDNA 4930546H06	gene	4930546H06Rik
1430960_at	1.00774	RIKEN cDNA 4933440J02	gene	4933440J02Rik
1430961_at	0.99662	zinc finger protein 292		Zfp292
1430962_at	1.12190	ubiquitin-conjugating enzyme E2S		Ube2s
1430963_at	0.92256	"glucosaminyl (N-acetyl) transferase 3, mucin type"		Gcnt3
1430964_at	1.01004	RIKEN cDNA 2310034O05	gene	2310034O05Rik
1430965_at	1.02411	RIKEN cDNA 9430064K01	gene	9430064K01Rik

1430966_at 0.90761 camello-like 3 Cml3
1430967_at 1.07667 RIKEN cDNA 3110080O07 gene 3110080O07Rik
1430968_at 1.08705 RIKEN cDNA 4930517O19 gene 4930517O19Rik
1430969_at 1.34870 mitochondrial carrier homolog 2 (C. elegans) Mtch2
1430970_a_at 0.97486 RIKEN cDNA 4930438O03 gene 4930438O03Rik
1430971_a_at 1.01961 aquarius Aqr
1430972_x_at 1.14348 aquarius Aqr
1430973_at 1.02656 zinc finger protein 558 Zfp558
1430974_a_at 1.05125 SUMO1/sentrin specific peptidase 7 Senp7
1430975_at 1.05604 RIKEN cDNA 8430416G17 gene 8430416G17Rik
1430976_a_at 0.90211 mitochondrial ribosomal protein L9 Mrpl9
1430977_at 0.87645 RIKEN cDNA A930021C24 gene A930021C24Rik
1430978_at 1.24899 ribosomal protein S25 Rps25
1430979_a_at 0.98940 peroxiredoxin 2 Prdx2
1430980_a_at 1.14838 eukaryotic translation initiation factor 4A1 Eif4a1
1430981_s_at 0.92689 GC-rich promoter binding protein 1 Gbp1
1430982_at 1.03133 "splicing factor, arginine/serine-rich 1 (ASF/SF2)" Sfrs1
1430983_at 1.06431 protein disulfide isomerase associated 6 Pdia6
1430984_at 0.92522 antizyme inhibitor 1 Azin1
1430985_at 1.05388 RIKEN cDNA 1810027O10 gene 1810027O10Rik
1430986_at 1.05164 "phenylalanine-tRNA synthetase-like, beta subunit" Farslb
1430987_s_at 0.94549 WW domain binding protein 11 Wbp11
1430988_at 0.92977 RIKEN cDNA 2810407C02 gene 2810407C02Rik
1430989_a_at 0.98606 RIKEN cDNA 1700020I14 gene 1700020I14Rik
1430990_s_at 0.98121 mitochondrial ribosomal protein L44 Mrpl44
1430991_at 1.02975 RIKEN cDNA 1810014B01 gene 1810014B01Rik
1430992_s_at 1.10468 RIKEN cDNA 1500009M05 gene 1500009M05Rik
1430993_at 0.98256 ubiquitin associated protein 2-like Ubap2l
1430994_at 1.06142 membrane-associated ring finger (C3HC4) 5 5-Mar
1430995_at 1.10028 --- ---
1430996_at 0.92995 ethanolamine kinase 1 Etnk1
1430997_at 1.04402 "CD47 antigen (Rh-related antigen, integrin-associated signal transducer)"
Cd47
1430998_at 0.87742 sulfide quinone reductase-like (yeast) Sqrdl
1430999_a_at 0.91567 short coiled-coil protein Scoc
1431000_at 0.88120 RIKEN cDNA 2310002B06 gene 2310002B06Rik
1431001_at 0.93057 RIKEN cDNA 9030624J02 gene 9030624J02Rik
1431002_x_at 0.93567 fumarylacetoacetate hydrolase domain containing 2A Fahd2a
1431003_a_at 1.01948 RIKEN cDNA 2610002J02 gene 2610002J02Rik
1431004_at 1.19403 lysyl oxidase-like 2 Loxl2
1431005_at 0.91226 "DNA segment, Chr 10, Wayne State University 102, expressed" D10Wsu102e
1431006_at 1.06126 --- ---
1431007_at 1.08651 RIKEN cDNA 9030025P20 gene 9030025P20Rik

1431008_at 1.31628 RIKEN cDNA 0610037M15 gene 0610037M15Rik
1431009_at 0.88208 RIKEN cDNA B230219D22 gene B230219D22Rik
1431010_a_at 1.01190 retinol dehydrogenase 12 Rdh12
1431011_at 1.03380 dihydrolipoamide S-succinyltransferase (E2 component of 2-oxo-glutarate complex) Dlst
1431012_a_at 0.89242 "peroxisomal delta3, delta2-enoyl-Coenzyme A isomerase" Peci
1431013_at 0.94603 "proteasome (prosome, macropain) 26S subunit, non-ATPase, 11 /// similar to 26S proteasome non-ATPase regulatory subunit 11 (26S proteasome regulatory subunit S9) (26S proteasome regulatory subunit p44.5) /// similar to 26S proteasome non-ATPase regulatory subunit 11 (26S proteasome regulatory subunit S9) (26S proteasome regulatory subunit p44.5)" Psm11 /// LOC627316 /// LOC671671
1431014_at 0.99838 "cat eye syndrome chromosome region, candidate 2 homolog (human)" Cacr2
1431015_at 0.91293 transmembrane protein 129 Tmem129
1431016_at 1.09974 "CWF19-like 1, cell cycle control (S. pombe)" Cwf19l1
1431017_at 1.06408 regulator of chromosome condensation (RCC1) and BTB (POZ) domain containing protein 2 Rcbtb2
1431018_at 1.09470 RIKEN cDNA 1810013L24 gene 1810013L24Rik
1431019_at 1.03516 leucine rich repeat containing 28 Lrrc28
1431020_a_at 1.10966 FGFR1 oncogene partner 2 Fgfr1op2
1431021_at 0.93779 cytochrome b-561 domain containing 1 Cyb561d1
1431022_at 1.12582 "synuclein, alpha" Snca
1431023_at 0.96406 RIKEN cDNA C030046E11 gene C030046E11Rik
1431024_a_at 0.96068 AT rich interactive domain 4B (Rbp1 like) Arid4b
1431025_at 1.01600 fuzzy homolog (Drosophila) Fuz
1431026_at 1.02458 N-acetyltransferase 12 Nat12
1431027_at 1.03279 RIKEN cDNA 2810455O05 gene 2810455O05Rik
1431028_a_at 1.04900 pantothenate kinase 1 Pank1
1431029_at 1.04605 tubulin tyrosine ligase like family 9 Ttl9
1431030_a_at 0.94149 ring finger protein 14 Rnf14
1431031_at 0.87107 AT rich interactive domain 4B (Rbp1 like) Arid4b
1431032_at 1.14681 "amylo-1,6-glucosidase, 4-alpha-glucanotransferase" Agl
1431033_x_at 0.99553 "amylo-1,6-glucosidase, 4-alpha-glucanotransferase" Agl
1431034_at 0.99323 G protein-coupled receptor 125 Gpr125
1431035_at 1.04942 dishevelled associated activator of morphogenesis 1 Daam1
1431036_a_at 0.93328 transmembrane emp24 domain trafficking protein 2 Tmed2
1431037_a_at 0.94021 "ELAV (embryonic lethal, abnormal vision, Drosophila)-like 1 (Hu antigen R)" Elavl1
1431038_at 1.14734 Ras association (RalGDS/AF-6) domain family 4 Rassf4
1431039_at 1.03744 OTU domain containing 4 Otud4
1431040_at 0.99525 RIKEN cDNA 6530401N04 gene /// hypothetical protein LOC638977 6530401N04Rik /// LOC638977
1431041_at 0.97643 GTP-binding protein 8 (putative) Gtpbp8

1431042_at	0.90156	progesterin and adipoQ receptor family member VIII	Paqr8
1431043_at	0.90062	kelch repeat and BTB (POZ) domain containing 5	Kbtbd5
1431044_at	1.82315	THO complex 1	Thoc1
1431045_at	0.98942	"DNA segment, Chr 12, ERATO Doi 553, expressed"	D12ErtD553e
1431046_at	0.95905	RIKEN cDNA 2410127E16 gene	2410127E16Rik
1431047_at	0.88048	chromodomain helicase DNA binding protein 6	Chd6
1431048_at	1.21806	phosphoserine/threonine/tyrosine interaction protein	Styx
1431049_at	1.04864	islet cell autoantigen 1-like	Ica11
1431050_at	0.87066	"ribosomal protein S6 kinase, polypeptide 5"	Rps6ka5
1431051_at	0.92928	RIKEN cDNA 4930577N17 gene	4930577N17Rik
1431052_at	0.72763	Rho GTPase activating protein 12	Arhgap12
1431053_at	1.05754	M-phase phosphoprotein 9	Mphosph9
1431054_at	1.05350	"LSM6 homolog, U6 small nuclear RNA associated (S. cerevisiae)"	Lsm6
1431055_a_at	1.07510	sorting nexin 10	Snx10
1431056_a_at	0.61094	lipoprotein lipase /// similar to Lipoprotein lipase precursor (LPL)	Lpl ///
LOC669888			
1431057_a_at	1.04885	"protease, serine, 23"	Prss23
1431058_at	1.00895	HIV TAT specific factor 1	Htatsf1
1431059_x_at	0.82415	HIV TAT specific factor 1	Htatsf1
1431060_at	1.65614	pellino 1	Peli1
1431061_s_at	0.91987	pellino 1	Peli1
1431062_a_at	1.04181	exocyst complex component 4	Exoc4
1431063_at	0.74580	---	---
1431064_at	0.90845	dipeptidylpeptidase 8	Dpp8
1431065_at	1.09405	ELL associated factor 1	Eaf1
1431066_at	0.95246	fucosyltransferase 11	Fut11
1431067_at	0.89040	RIKEN cDNA 6330404A07 gene	6330404A07Rik
1431068_at	0.87985	required for meiotic nuclear division 5 homolog A (S. cerevisiae)	Rmnd5a
1431069_at	0.92363	RIKEN cDNA 4930544G21 gene	4930544G21Rik
1431070_a_at	0.93100	RIKEN cDNA 9130404D08 gene	9130404D08Rik
1431071_at	0.85534	metaxin 3	Mtx3
1431072_a_at	1.03283	coiled-coil domain containing 50	Ccdc50
1431073_at	1.10947	protein prenyltransferase alpha subunit repeat containing 1	Ptar1
1431074_a_at	0.86005	"phosphatidylinositol transfer protein, cytoplasmic 1"	Pitpnc1
1431075_a_at	0.97641	eukaryotic translation initiation factor 4E nuclear import factor 1	Eif4enif1
1431076_at	0.94900	adducin 2 (beta)	Add2
1431077_at	1.10430	"alkaline phosphatase, intestinal"	Alpi
1431078_at	0.97475	F-box only protein 3	Fbxo3
1431079_at	1.02745	C1q and tumor necrosis factor related protein 2	C1qtnf2
1431080_at	0.96507	RIKEN cDNA 1700021K10 gene	1700021K10Rik
1431081_a_at	0.90125	phospholipid scramblase 3	Plscr3
1431082_a_at	0.96324	"RAB, member of RAS oncogene family-like 3"	Rab13
1431083_a_at	0.89407	RIKEN cDNA 1810014B01 gene	1810014B01Rik

1431084_x_at 0.95556 RIKEN cDNA 1810014B01 gene 1810014B01Rik
1431085_a_at 1.04525 protein-L-isoaspartate (D-aspartate) O-methyltransferase 1 Pcmt1
1431086_s_at 1.08779 protein-L-isoaspartate (D-aspartate) O-methyltransferase 1 Pcmt1
1431087_at 1.03993 spindle pole body component 24 homolog (S. cerevisiae) Spbc24
1431088_at 0.89674 forkhead box P4 Foxp4
1431089_at 0.95483 cleavage and polyadenylation specific factor 2 Cpsf2
1431090_at 0.87480 Ssu72 RNA polymerase II CTD phosphatase homolog (yeast) Ssu72
1431091_at 1.02331 pygopus 1 Pygo1
1431092_at 0.89595 "protein phosphatase 1, regulatory (inhibitor) subunit 12C" Ppp1r12c
1431093_at 1.02700 nephronophthisis 3 (adolescent) Nphp3
1431094_at 1.06904 RIKEN cDNA 1110006E14 gene 1110006E14Rik
1431095_a_at 0.86864 hect domain and RLD 5 Herc5
1431096_at 1.03266 integrator complex subunit 8 Ints8
1431097_at 0.85296 GTPase activating RANGAP domain-like 1 Garnl1
1431098_at 0.93135 restin (Reed-Steinberg cell-expressed intermediate filament-associated protein)
Rsn
1431099_at 0.99052 homeo box D8 Hoxd8
1431100_at 1.01882 RIKEN cDNA 9930116N10 gene 9930116N10Rik
1431101_a_at 0.93712 steroid 5 alpha-reductase 1 Srd5a1
1431102_at 0.96820 centrosomal protein 350 Cep350
1431103_at 0.94009 RIKEN cDNA 1700003P14 gene 1700003P14Rik
1431104_at 0.98401 RIKEN cDNA 4930522H14 gene 4930522H14Rik
1431105_a_at 1.02098 transmembrane protein 33 Tmem33
1431106_a_at 0.92756 RIKEN cDNA 9530053H05 gene 9530053H05Rik
1431107_at 0.95239 serine/threonine kinase 35 Stk35
1431108_at 0.83348 RIKEN cDNA 2410131K14 gene 2410131K14Rik
1431109_at 0.91403 RIKEN cDNA 5430406M13 gene 5430406M13Rik
1431110_at 1.03944 RIKEN cDNA 5430431D22 gene 5430431D22Rik
1431111_at 1.17164 "enolase 1, alpha non-neuron /// similar to enolase 1, alpha non-neuron ///
similar to enolase 1, alpha non-neuron /// similar to T21B10.2b /// similar to enolase 1, alpha non-
neuron" Eno1 /// LOC238943 /// LOC433182 /// LOC671462 /// LOC674877
1431112_at 1.02239 unc-51-like kinase 3 (C. elegans) Ulk3
1431113_at 1.48553 RNA binding motif protein 4B Rbm4b
1431114_at 1.06514 dedicator of cytokinesis 4 Dock4
1431115_at 1.13345 TGFB-induced factor 2 Tgif2
1431116_at 1.01653 RIKEN cDNA C030003D03 gene C030003D03Rik
1431117_x_at 1.06389 RIKEN cDNA 1810029B16 gene /// hypothetical LOC383495 /// hypothetical
protein LOC675103 1810029B16Rik /// LOC383495 /// LOC675103
1431118_at 1.59625 RIKEN cDNA 6720427H10 gene 6720427H10Rik
1431119_at 1.15969 "DnaJ (Hsp40) related, subfamily B, member 13" Dnajb13
1431120_a_at 1.24860 "golgi autoantigen, golgin subfamily a, 1" Golga1
1431121_at 1.00642 angiomotin-like 1 Amotl1
1431122_at 0.86167 armadillo repeat containing 9 Armc9

1431123_s_at 1.09923 "DnaJ (Hsp40) homolog, subfamily C, member 8 /// similar to DnaJ (Hsp40) homolog, subfamily C, member 8 (predicted)" Dnajc8 /// LOC669266
 1431124_at 1.01544 AU RNA binding protein/enoyl-coenzyme A hydratase Auh
 1431125_a_at 0.93772 threonyl-tRNA synthetase-like 1 Tarsl1
 1431126_a_at 1.07047 RIKEN cDNA 0610011F06 gene 0610011F06Rik
 1431127_at 0.93411 zinc finger protein 297B Zfp297b
 1431128_at 0.96866 "DNA segment, Chr 8, Brigham & Women's Genetics 1414 expressed" D8Bwg1414e
 1431129_at 0.94811 ubiquitin specific peptidase 31 Usp31
 1431130_at 1.01869 RIKEN cDNA 2010110P09 gene 2010110P09Rik
 1431131_s_at 1.00043 RIKEN cDNA A630007B06 gene /// similar to oocyte-testis gene 1 A630007B06Rik /// LOC671029
 1431132_x_at 1.09501 RIKEN cDNA 0610007P22 gene 0610007P22Rik
 1431133_at 1.05353 Rho GTPase activating protein 18 Arhgap18
 1431134_at 1.17268 "inhibitor of growth family, member 5" Ing5
 1431135_at 0.96763 --- ---
 1431136_at 0.90241 "RAB36, member RAS oncogene family" Rab36
 1431137_at 1.06541 RUN and SH3 domain containing 1 Ruscl
 1431138_at 0.88630 armadillo repeat containing 9 Armc9
 1431139_at 1.02395 "chromodomain protein, Y chromosome-like 2" Cdy12
 1431140_at 1.01711 thioesterase superfamily member 4 Them4
 1431141_at 0.97795 RIKEN cDNA 2610507I01 gene 2610507I01Rik
 1431142_s_at 0.98044 apoptosis-inducing factor (AIF)-like mitochondrion-associated inducer of death Amid
 1431143_x_at 0.97009 apoptosis-inducing factor (AIF)-like mitochondrion-associated inducer of death Amid
 1431144_at 1.10707 RIKEN cDNA 2810002G02 gene /// region containing RIKEN cDNA 2810002G02 gene; tuftelin interacting protein 11 2810002G02Rik /// LOC671202
 1431145_a_at 0.99187 CUE domain containing 2 Cuedc2
 1431146_a_at 0.83409 copine VIII Cpne8
 1431147_at 1.14911 RAD50 interactor 1 Rint1
 1431148_at 0.81112 --- ---
 1431149_at 0.86933 RIKEN cDNA B430105A11 gene B430105A11Rik
 1431150_at 0.85821 RIKEN cDNA 1700110M21 gene 1700110M21Rik
 1431151_at 0.92491 "similar to Gastric triacylglycerol lipase precursor (Gastric lipase) (GL) /// similar to Gastric triacylglycerol lipase precursor (Gastric lipase) (GL) /// similar to Gastric triacylglycerol lipase precursor (Gastric lipase) (GL) /// similar to lipase, gastric /// similar to Gastric triacylglycerol lipase precursor (Gastric lipase) (GL)" LOC628236 /// LOC668099 /// LOC668114 /// LOC673023 /// LOC676650
 1431152_at 1.09590 hyaluronan and proteoglycan link protein 3 Hapln3
 1431153_at 0.87434 Neurexin III Nrxn3
 1431154_at 1.00121 zinc and ring finger 2 Znrf2
 1431155_at 1.01755 calmodulin regulated spectrin-associated protein 1-like 1 Camsap111

1431156_at 1.05922 RIKEN cDNA 9430040K09 gene 9430040K09Rik
1431157_at 1.02492 RIKEN cDNA 1700025I17 gene 1700025I17Rik
1431158_at 2.07253 RIKEN cDNA 8430406H22 gene 8430406H22Rik
1431159_at 1.07046 RIKEN cDNA 2900006F19 gene 2900006F19Rik
1431160_x_at 0.95651 similar to gonadotropin inducible ovarian transcription factor 1 /// similar to Hippocalcin-like protein 1 (Visinin-like protein 3) (VILIP-3) (Neural visinin-like protein 3) (NVL-3) (NVP-3) LOC628850 /// LOC628882
1431161_at 1.02928 Rho-guanine nucleotide exchange factor Rgnef
1431162_a_at 1.11060 enabled homolog (Drosophila) Enah
1431163_at 1.01749 RIKEN cDNA 2700046G09 gene 2700046G09Rik
1431164_at 0.94548 Ras-related GTP binding D Rragd
1431165_at 1.05877 RIKEN cDNA A030009A09 gene A030009A09Rik
1431166_at 1.02073 --- ---
1431167_at 1.01134 "diacylglycerol kinase, gamma" Dgkg
1431168_at 0.96418 Six3 opposite strand transcript 1 Six3os1
1431169_at 1.00685 RIKEN cDNA D230012E17 gene D230012E17Rik
1431170_at 1.04623 ephrin A3 Efna3
1431171_at 1.02168 RIKEN cDNA D730001G18 gene D730001G18Rik
1431172_at 0.92029 "origin recognition complex, subunit 4-like (S. cerevisiae)" Orc4l
1431173_at 1.01340 RIKEN cDNA A930008G19 gene A930008G19Rik
1431174_at 1.02386 RIKEN cDNA A930036K24 gene A930036K24Rik
1431175_at 0.91655 RIKEN cDNA 1810019D21 gene 1810019D21Rik
1431176_at 0.91938 RIKEN cDNA 2310010I16 gene 2310010I16Rik
1431177_a_at 1.06828 ribosomal protein L10A Rpl10a
1431178_at 0.74335 similar to murine ULBP-like transcript 1 LOC640488
1431179_at 1.00267 ectonucleoside triphosphate diphosphohydrolase 7 Entpd7
1431180_at 0.97604 RIKEN cDNA 1700123A16 gene /// similar to zinc finger protein 709 1700123A16Rik /// LOC666461
1431181_a_at 1.00091 Luc7 homolog (S. cerevisiae)-like Luc7l
1431182_at 1.11360 heat shock protein 8 /// similar to heat shock protein 8 /// similar to heat shock protein 8 /// similar to heat shock protein 8 /// heat shock protein 8 pseudogene /// heat shock protein 8 pseudogene /// similar to heat shock protein 8 /// similar to heat shock protein 8 /// Hspa8 /// LOC621284 /// LOC624853 /// LOC632214 /// LOC641192 /// LOC666508 /// LOC666904 /// LOC669493 /// LOC671270 /// LOC675417
1431183_at 0.96370 RIKEN cDNA 1700066M21 gene 1700066M21Rik
1431184_a_at 1.02782 RIKEN cDNA 4930503B20 gene 4930503B20Rik
1431185_at 0.96572 RIKEN cDNA 4930503B20 gene 4930503B20Rik
1431186_at 1.03382 "discs, large homolog 5 (Drosophila)" Dlg5
1431187_s_at 1.18849 "discs, large homolog 5 (Drosophila)" Dlg5
1431188_a_at 1.01392 target of myb1 homolog (chicken) /// similar to TOM1 Tom1 /// LOC545878
1431189_a_at 0.85651 fumarylacetoacetate hydrolase domain containing 2A Fahd2a
1431190_x_at 0.94787 fumarylacetoacetate hydrolase domain containing 2A Fahd2a

1431191_a_at 0.99839 synaptotagmin I Syt1
1431192_at 1.06167 RIKEN cDNA 2010203O03 gene 2010203O03Rik
1431193_at 0.94056 "TAF4B RNA polymerase II, TATA box binding protein (TBP)-associated factor" Taf4b
1431194_at 1.00292 RIKEN cDNA 6030468B19 gene 6030468B19Rik
1431195_at 1.07078 ring finger protein 170 Rnf170
1431196_at 0.97663 "ATPase, Ca⁺⁺-sequestering" Atp2c1
1431197_at 0.89534 ADP-ribosylation factor-like 6 interacting protein 2 Arl6ip2
1431198_x_at 1.29130 RIKEN cDNA 9430038I01 gene 9430038I01Rik
1431199_at 0.87969 RIKEN cDNA 0610031G08 gene 0610031G08Rik
1431200_a_at 0.96235 "zinc finger, CSL domain containing 2" Zcsl2
1431201_at 0.95524 "solute carrier family 25, member 27" Slc25a27
1431202_at 1.02265 hect domain and RLD 3 Herc3
1431203_at 0.96624 serologically defined colon cancer antigen 8 Sdccag8
1431204_at 0.78367 RIKEN cDNA 4930578N16 gene 4930578N16Rik
1431205_at 0.94693 "Solute carrier family 9 (sodium/hydrogen exchanger), member 8" Slc9a8
1431206_at 0.81043 RIKEN cDNA 5730601F06 gene 5730601F06Rik
1431207_at 1.04137 RIKEN cDNA 2900024O10 gene 2900024O10Rik
1431208_a_at 0.86089 "solute carrier family 9 (sodium/hydrogen exchanger), isoform 3 regulator 2" Slc9a3r2
1431209_s_at 1.02721 transmembrane protein 42 Tmem42
1431210_at 1.03007 ATP/GTP binding protein-like 3 Agbl3
1431211_s_at 0.97356 thioesterase superfamily member 5 Them5
1431212_a_at 1.04753 RIKEN cDNA 3300001M20 gene 3300001M20Rik
1431213_a_at 1.31174 RIKEN cDNA 1300007C21 gene /// similar to RIKEN cDNA 1300007C21 gene /// similar to NADH-ubiquinone oxidoreductase MWFE subunit (Complex I-MWFE) (CI-MWFE) /// hypothetical protein LOC670993 /// hypothetical protein LOC673100 1300007C21Rik /// LOC668462 /// LOC668581 /// LOC670993 /// LOC673100
1431214_at 0.99084 RIKEN cDNA 1300007C21 gene 1300007C21Rik
1431215_at 0.97025 "DnaJ (Hsp40) homolog, subfamily C, member 6" Dnajc6
1431216_s_at 1.04651 "DnaJ (Hsp40) homolog, subfamily C, member 6" Dnajc6
1431217_at 1.22789 leucine rich repeat containing 27 Lrrc27
1431218_at 1.35006 "zinc finger, DHHC domain containing 20" Zdhhc20
1431219_at 1.29789 "beta-1,3-glucuronyltransferase 3 (glucuronosyltransferase I)" B3gat3
1431220_at 0.70539 RIKEN cDNA 2810416G20 gene 2810416G20Rik
1431221_at 1.07004 RIKEN cDNA 5830405M20 gene 5830405M20Rik
1431222_at 0.98826 T-cell lymphoma invasion and metastasis 2 Tiam2
1431223_at 0.95282 RIKEN cDNA 2700054A10 gene 2700054A10Rik
1431224_at 0.92874 ubiquitin protein ligase E3A Ube3a
1431225_at 0.71521 SRY-box containing gene 11 Sox11
1431226_a_at 0.85481 fibronectin type III domain containing 4 Fndc4
1431227_at 1.05733 RIKEN cDNA 2310040A13 gene 2310040A13Rik
1431228_s_at 1.06466 RIKEN cDNA 4930526I15 gene 4930526I15Rik

1431229_at 0.95728 imprinted gene in the Prader-Willi syndrome region Ipw
1431230_a_at 0.98345 BTB (POZ) domain containing 9 Btbd9
1431231_at 1.22423 "histone 2, H3c1 /// histone 1, H3f /// histone1, H3d /// histone 1, H3b /// histone 1, H3e /// histone 2, H3b" Hist2h3c1 /// Hist1h3f /// Hist1h3d /// Hist1h3b /// Hist1h3e /// Hist2h3b
1431232_a_at 1.23365 MAX gene associated Mga
1431233_at 0.72642 cyclin M4 Cnm4
1431234_at 1.00503 RIKEN cDNA 1700041B20 gene 1700041B20Rik
1431235_at 1.88541 --- ---
1431236_at 0.87220 serine/threonine kinase 32B Stk32b
1431237_at 0.97260 "nuclear receptor subfamily 2, group F, member 2" Nr2f2
1431238_at 0.92174 PQ loop repeat containing 1 Pqlc1
1431239_at 1.51523 "non-POU-domain-containing, octamer binding protein" Nono
1431240_at 1.03397 "C-type lectin domain family 2, member h" Clec2h
1431241_at 1.14786 coiled-coil-helix-coiled-coil-helix domain containing 3 Chchd3
1431242_at 0.94582 RIKEN cDNA 6330575P09 gene 6330575P09Rik
1431243_at 0.97704 RIKEN cDNA 1700120J03 gene 1700120J03Rik
1431244_s_at 0.84890 dynamin binding protein Dnmbp
1431245_at 0.80474 RIKEN cDNA 1700063H04 gene 1700063H04Rik
1431246_at 0.96434 IQ motif containing E Iqce
1431247_at 1.02132 RIKEN cDNA 1300014J16 gene 1300014J16Rik
1431248_at 2.34493 RIKEN cDNA 5031426D15 gene 5031426D15Rik
1431249_at 1.02508 --- ---
1431250_at 0.95178 RIKEN cDNA 9030624G23 gene /// similar to development- and differentiation-
enhancing factor 2 /// similar to Hippocalcin-like protein 1 (Visinin-like protein 3) (VILIP-3) (Neural
visinin-like protein 3) (NVL-3) (NVP-3) /// similar to development- and differentiation-enhancing factor
2 /// similar to development- and differentiation-enhancing factor 2 /// similar to gonadotropin inducible
ovarian transcription factor 1 /// similar to development- and differentiation-enhancing factor 2 ///
similar to development- and differentiation-enhancing factor 2 /// similar to development- and
differentiation-enhancing factor 2 /// hypothetical protein LOC673340 9030624G23Rik ///
LOC272713 /// LOC544848 /// LOC623886 /// LOC668643 /// LOC668669 /// LOC671762 ///
LOC673231 /// LOC673232 /// LOC673340
1431251_at 0.99365 RIKEN cDNA 1300011L04 gene 1300011L04Rik
1431252_a_at 0.87671 zinc finger protein 655 Zfp655
1431253_s_at 0.82254 "TBC1 domain family, member 9" Tbc1d9
1431254_at 0.96170 kelch repeat and BTB (POZ) domain containing 11 /// similar to Protein
KIAA0711 Kbtbd11 /// LOC632344
1431255_at 1.13878 calreticulin 3 Calr3
1431256_at 1.33634 RIKEN cDNA 5033423O07 gene 5033423O07Rik
1431257_at 0.91475 RIKEN cDNA 6230409E13 gene 6230409E13Rik
1431258_at 1.07743 death inducer-obliterator 1 Dido1
1431259_at 0.81546 --- ---
1431260_at 1.02270 RIKEN cDNA 4833417C18 gene 4833417C18Rik
1431261_at 0.91002 chloride intracellular channel 5 Clic5

1431262_at 1.00949 hypothetical protein LOC544988 /// similar to Spetex-2H protein /// similar to Spetex-2H protein /// hypothetical protein LOC666772 LOC544988 /// LOC622931 /// LOC666442 /// LOC666772

1431263_at 0.97388 "protein kinase, AMP-activated, gamma 2 non-catalytic subunit" Prkag2

1431264_at 1.03431 --- ---

1431265_at 0.97884 zinc finger protein 28 Zfp28

1431266_at 1.04014 RIKEN cDNA 5830433G22 gene 5830433G22Rik

1431267_at 1.01069 RIKEN cDNA 5430425J12 gene 5430425J12Rik

1431268_at 1.09947 RIKEN cDNA 1700047I17 gene 1700047I17Rik

1431269_at 1.17710 peroxiredoxin 4 Prdx4

1431270_a_at 0.90748 asteroid homolog 1 (Drosophila) Aste1

1431271_at 0.98489 RIKEN cDNA 2810428J06 gene 2810428J06Rik

1431272_at 0.96160 RIKEN cDNA 5430405H02 gene 5430405H02Rik

1431273_at 1.10387 RIKEN cDNA 2810417H13 gene 2810417H13Rik

1431274_a_at 1.03976 --- ---

1431275_at 0.93334 "chromatin assembly factor 1, subunit B (p60)" Chaf1b

1431276_at 0.91103 RIKEN cDNA 9130019P16 gene 9130019P16Rik

1431277_at 0.93299 --- ---

1431278_s_at 0.91123 "phospholipase A2, group VI" Pla2g6

1431279_s_at 1.07332 tubulin tyrosine ligase like family 5 Ttl15

1431280_at 1.18463 expressed sequence AI597468 AI597468

1431281_at 1.00798 dysferlin interacting protein 1 Dysfip1

1431282_at 0.99592 thyroid hormone receptor associated protein 4 Thrap4

1431283_at 0.86220 --- ---

1431284_a_at 1.05104 RIKEN cDNA 2210013K02 gene 2210013K02Rik

1431285_at 0.89416 "mahogunin, ring finger 1" Mgrn1

1431286_at 1.02009 --- ---

1431287_at 1.13847 pericentriolar material 1 Pcm1

1431288_at 1.02755 carbonic anhydrase 10 Car10

1431289_at 0.97033 tetraspanin 32 Tspan32

1431290_at 1.03160 RIKEN cDNA 4930588J15 gene 4930588J15Rik

1431291_at 1.07861 RIKEN cDNA 2700071J12 gene 2700071J12Rik

1431292_a_at 1.00240 protein tyrosine kinase 9-like (A6-related protein) Ptk9l

1431293_a_at 0.90326 claudin domain containing 1 Cldnd1

1431294_at 0.95714 "spermatogenesis associated glutamate (E)-rich protein 8, pseudogene 1 /// spermatogenesis associated glutamate (E)-rich protein 7, pseudogene 1" Speer8-ps1 /// Speer7-ps1

1431295_a_at 0.96088 syntaxin 18 Stx18

1431296_at 0.96704 G protein-coupled receptor 15 Gpr15

1431297_a_at 1.03200 RIKEN cDNA 4933436C20 gene 4933436C20Rik

1431298_at 1.18895 RIKEN cDNA 5930416I19 gene 5930416I19Rik

1431299_a_at 0.94268 RIKEN cDNA 2310014H01 gene 2310014H01Rik

1431300_at 0.94545 SH3-domain GRB2-like (endophilin) interacting protein 1 Sgip1

1431301_at 1.07959 RIKEN cDNA E130201N16 gene E130201N16Rik

1431302_a_at 1.05800 nudix (nucleoside diphosphate linked moiety X)-type motif 7 Nudt7
1431303_at 1.12408 serine/threonine kinase 38 Stk38
1431304_a_at 0.97491 RIKEN cDNA 1300007B12 gene 1300007B12Rik
1431305_at 0.99892 mucolipin 1 Mcoln1
1431306_at 0.99442 "5'-nucleotidase, cytosolic III" Nt5c3
1431307_at 1.08234 RIKEN cDNA 4930589M24 gene 4930589M24Rik
1431308_at 0.95601 RIKEN cDNA 5730557B15 gene 5730557B15Rik
1431309_at 0.99873 "tubulin tyrosine ligase-like family, member 10" Ttl10
1431310_s_at 0.95714 "tubulin tyrosine ligase-like family, member 10" Ttl10
1431311_at 1.02124 RIKEN cDNA 4430401P03 gene 4430401P03Rik
1431312_at 0.87810 RIKEN cDNA 5730577I03 gene 5730577I03Rik
1431313_at 0.92016 RIKEN cDNA 5730442P18 gene 5730442P18Rik
1431314_a_at 1.03729 RIKEN cDNA 5830417I10 gene 5830417I10Rik
1431315_at 1.02980 hydrolethalus syndrome 1 Hyls1
1431316_at 1.41922 itchy Itch
1431317_at 1.13660 RIKEN cDNA 2410018L13 gene 2410018L13Rik
1431318_at 0.97529 RIKEN cDNA 4833420G17 gene 4833420G17Rik
1431319_at 0.86420 unc-51-like kinase 4 (C. elegans) Ulk4
1431320_a_at 1.12677 myosin Va Myo5a
1431321_at 0.99348 RIKEN cDNA 4930513E20 gene 4930513E20Rik
1431322_at 0.91000 "immunoglobulin superfamily, member 3" Igsf3
1431323_at 0.98542 leucine zipper transcription factor-like 1 Lztf1
1431324_at 0.97775 "extracellular matrix protein 2, female organ and adipocyte specific" Ecm2
1431325_at 0.95319 camello-like 3 Cml3
1431326_a_at 1.01307 tropomodulin 2 Tmod2
1431327_at 1.18128 vacuolar protein sorting 13A (yeast) Vps13a
1431328_at 0.87135 "protein phosphatase 1, catalytic subunit, beta isoform" Ppp1cb
1431329_at 0.96581 nephronophthisis 4 (juvenile) homolog (human) Nphp4
1431330_at 0.93763 RIKEN cDNA 3110002H16 gene 3110002H16Rik
1431331_at 0.91102 monoglyceride lipase Mgl
1431332_a_at 1.09468 telomeric repeat binding factor 1 Terf1
1431333_at 0.98670 "histone 1, H4h" Hist1h4h
1431334_a_at 1.03737 RIKEN cDNA 4933433P14 gene 4933433P14Rik
1431335_a_at 0.59213 WAP four-disulfide core domain 1 Wfdc1
1431336_at 0.99317 "scavenger receptor class A, member 5 (putative)" Scara5
1431337_a_at 0.95773 RIKEN cDNA 1810055E12 gene 1810055E12Rik
1431338_at 0.97290 CASK interacting protein 1 Caskin1
1431339_a_at 1.04014 EF hand domain containing 2 Efhd2
1431340_a_at 1.12744 RIKEN cDNA 2310002J21 gene 2310002J21Rik
1431341_at 0.88065 "Protein phosphatase 2 (formerly 2A), catalytic subunit, beta isoform" Ppp2cb
1431342_at 0.91429 --- ---
1431343_at 1.04628 --- ---
1431344_at 0.93311 "G protein-coupled receptor, family C, group 1, member B" Grm2

1431345_a_at	0.90781	"TATA box binding protein (Tbp)-associated factor, RNA polymerase I, B"	
Taf1b			
1431346_at	0.85471	RIKEN cDNA 6330405D24 gene	6330405D24Rik
1431347_at	1.09523	RIKEN cDNA 5730407M17 gene	5730407M17Rik
1431348_at	0.99783	"peptidylprolyl isomerase F, opposite strand transcription unit"	Ppifos
1431349_at	1.06605	heterogeneous nuclear ribonucleoprotein A/B	Hnrpab
1431350_at	0.93980	RIKEN cDNA C130036L24 gene	C130036L24Rik
1431351_at	1.13279	plasmacytoma variant translocation 1	Pvt1
1431352_s_at	1.30902	plasmacytoma variant translocation 1	Pvt1
1431353_at	1.12882	RIKEN cDNA C330050A14 gene	C330050A14Rik
1431354_a_at	0.95912	phenylalanine-tRNA synthetase 2 (mitochondrial)	Fars2
1431355_s_at	2.23628	"transient receptor potential cation channel, subfamily M, member 7"	Trpm7
1431356_at	1.07972	RIKEN cDNA 6430710C18 gene	6430710C18Rik
1431357_a_at	0.93565	retinitis pigmentosa GTPase regulator interacting protein 1	Rpgrip1
1431358_at	1.00816	RIKEN cDNA 4930547N16 gene	4930547N16Rik
1431359_a_at	1.10057	RIKEN cDNA 1110007C09 gene	1110007C09Rik
1431360_s_at	1.05634	DMRT-like family C1a	Dmrta1a
1431361_at	0.97038	prolylcarboxypeptidase (angiotensinase C)	Prpc
1431362_a_at	1.02599	SPARC related modular calcium binding 2	Smoc2
1431363_at	0.90595	RIKEN cDNA 4933425K02 gene	4933425K02Rik
1431364_a_at	1.09090	RIKEN cDNA 2810047C21 gene	2810047C21Rik
1431365_at	0.94198	ring finger protein 32	Rnf32
1431366_at	0.95865	RIKEN cDNA 4930519G04 gene	4930519G04Rik
1431367_at	0.92905	membrane-associated ring finger (C3HC4) 1	1-Mar
1431368_at	1.03906	RIKEN cDNA 4930404N11 gene	4930404N11Rik
1431369_at	1.03242	development and differentiation enhancing factor 2	Ddef2
1431370_at	1.04436	RIKEN cDNA 4930512B01 gene	4930512B01Rik
1431371_at	1.00654	RIKEN cDNA 9030411M13 gene	9030411M13Rik
1431372_at	1.31418	serine/arginine-rich protein specific kinase 2	Srpk2
1431373_at	0.97381	"processing of precursor 5, ribonuclease P/MRP family (S. cerevisiae)"	Pop5
1431374_at	1.22722	RIKEN cDNA 6330407A03 gene	6330407A03Rik
1431375_s_at	0.98430	"parvin, alpha"	Parva
1431376_at	1.19334	WD repeat domain 62	Wdr62
1431377_at	1.31239	ataxin 10	Atxn10
1431378_at	0.95332	---	---
1431379_a_at	1.00979	"solute carrier family 13 (sodium/sulphate symporters), member 1"	Slc13a1
1431380_at	1.06279	RIKEN cDNA 5730409L17 gene	5730409L17Rik
1431381_at	0.95335	RIKEN cDNA 3110005L24 gene	3110005L24Rik
1431382_a_at	0.98072	restin-like 2	Rsnl2
1431383_at	0.97047	RIKEN cDNA 1110028F11 gene	1110028F11Rik
1431384_at	1.03093	RIKEN cDNA 1700095G12 gene	1700095G12Rik
1431385_a_at	0.79905	"membrane-bound transcription factor peptidase, site 1"	Mbtps1
1431386_s_at	0.99713	"membrane-bound transcription factor peptidase, site 1"	Mbtps1

1431387_at 0.96161 Annexin A11 Anxa11
1431388_at 1.11740 M-phase phosphoprotein 10 (U3 small nucleolar ribonucleoprotein)
Mphosph10
1431389_at 0.92348 TBK1 binding protein 1 Tbkbp1
1431390_a_at 0.83109 "glutamate receptor, ionotropic, N-methyl D-aspartate-like 1A" Grin1a
1431391_at 1.41934 Ral GEF with PH domain and SH3 binding motif 1 Ralgps1
1431392_at 1.00118 homeo box A3 /// RIKEN cDNA 5730596B20 gene Hoxa3 ///
5730596B20Rik
1431393_at 1.08665 Six6 opposite strand transcript 1 Six6os1
1431394_a_at 0.83164 leucine-rich repeat kinase 2 Lrrk2
1431395_a_at 1.09563 IQ motif containing GTPase activating protein 1 Iqgap1
1431396_at 1.13366 IQ motif containing GTPase activating protein 1 Iqgap1
1431397_at 0.98771 --- ---
1431398_at 0.89754 RIKEN cDNA 5730446D14 gene 5730446D14Rik
1431399_at 1.19663 "a disintegrin-like and metallopeptidase (reprolysin type) with thrombospondin
type 1 motif, 9" Adamts9
1431400_a_at 0.92850 growth arrest specific 7 Gas7
1431401_at 1.17314 RIKEN cDNA 4930539J05 gene 4930539J05Rik
1431402_at 1.14723 kin of IRRE like 3 (Drosophila) Kirrel3
1431403_a_at 0.93607 RIKEN cDNA 2900002G04 gene 2900002G04Rik
1431404_at 0.90845 transmembrane protein 56 Tmem56
1431405_a_at 1.41338 RIKEN cDNA 4930413O22 gene 4930413O22Rik
1431406_at 0.94288 alanine-glyoxylate aminotransferase 2-like 1 Agxt2l1
1431407_at 1.00639 RIKEN cDNA 2310024H09 gene 2310024H09Rik
1431408_at 0.98651 RIKEN cDNA 4930412B13 gene 4930412B13Rik
1431409_at 0.89686 RIKEN cDNA C030015A19 gene C030015A19Rik
1431410_at 0.96609 "DNA segment, Chr 16, ERATO Doi 472, expressed" D16Ertd472e
1431411_a_at 0.99754 retinoic acid induced 12 Rai12
1431412_at 1.14409 RIKEN cDNA 2810455B08 gene 2810455B08Rik
1431413_at 1.05526 RIKEN cDNA 9130218E19 gene 9130218E19Rik
1431414_at 1.03451 RIKEN cDNA 2610312K03 gene 2610312K03Rik
1431415_a_at 1.08907 TATA box binding protein-like 1 Tbp1l
1431416_a_at 0.97366 junction adhesion molecule 2 Jam2
1431417_at 1.07211 junction adhesion molecule 2 Jam2
1431418_at 0.93853 RIKEN cDNA C030026M15 gene C030026M15Rik
1431419_at 1.01166 RIKEN cDNA 2610524G07 gene 2610524G07Rik
1431420_s_at 1.09525 RIKEN cDNA 2610524G07 gene 2610524G07Rik
1431421_x_at 1.05096 RIKEN cDNA 2610524G07 gene 2610524G07Rik
1431422_a_at 1.16724 dual specificity phosphatase 14 Dusp14
1431423_a_at 1.10311 "mediator of RNA polymerase II transcription, subunit 8 homolog (yeast)"
Med8
1431424_at 0.89854 RIKEN cDNA 2810055G20 gene 2810055G20Rik
1431425_a_at 0.94665 RIKEN cDNA 4930535B03 gene 4930535B03Rik

1431426_at	0.96595	RIKEN cDNA 4930535B03 gene	4930535B03Rik
1431427_at	1.01963	RIKEN cDNA 5430427M07 gene	5430427M07Rik
1431428_a_at	1.02943	nitric oxide synthase interacting protein	Nosip
1431429_a_at	0.85464	ADP-ribosylation factor-like 4A	Arl4a
1431430_s_at	0.83152	tripartite motif-containing 59 /// similar to mouse RING finger 1	Trim59 ///
LOC630539			
1431431_a_at	0.96136	nitrogen fixation gene 1 (<i>S. cerevisiae</i>)	Nfs1
1431432_at	1.21558	"cofilin 2, muscle"	Cfl2
1431433_at	1.05328	SET binding factor 2	Sbf2
1431434_at	0.94434	"serine/threonine kinase 39, STE20/SPS1 homolog (yeast)"	Stk39
1431435_at	0.95541	---	---
1431436_a_at	0.72334	katanin p60 subunit A-like 2	Katnal2
1431437_at	1.02390	forkhead box P4	Foxp4
1431438_at	1.07915	RIKEN cDNA C530001K22 gene	C530001K22Rik
1431439_at	1.12078	RIKEN cDNA 5830487J09 gene	5830487J09Rik
1431440_at	1.09608	"polymerase (DNA directed), epsilon 2 (p59 subunit)"	Pole2
1431441_at	0.81701	RIKEN cDNA 1700106J16 gene	1700106J16Rik
1431442_at	0.91838	RIKEN cDNA 2310047B19 gene	2310047B19Rik
1431443_at	0.99448	RIKEN cDNA 1810008B01 gene	1810008B01Rik
1431444_at	0.97893	"serine protease inhibitor, Kunitz type 4"	Spint4
1431445_at	0.95308	RIKEN cDNA 4921509A18 gene	4921509A18Rik
1431446_at	1.04896	RIKEN cDNA 0610009B14 gene	0610009B14Rik
1431447_at	0.97258	RIKEN cDNA 4930432K09 gene	4930432K09Rik
1431448_at	0.97974	RIKEN cDNA 4930474N09 gene	4930474N09Rik
1431449_at	1.03129	"solute carrier organic anion transporter family, member 6d1"	Slco6d1
1431450_at	1.06105	RIKEN cDNA 4930444P10 gene /// hypothetical protein LOC638108	
4930444P10Rik		/// LOC638108	
1431451_at	0.82825	glycosyltransferase 6 domain containing 1	Glt6d1
1431452_at	0.98721	N-deacetylase/N-sulfotransferase (heparin glucosaminyl) 4	Ndst4
1431453_at	1.00205	RIKEN cDNA 4933406K04 gene	4933406K04Rik
1431454_at	1.03560	RIKEN cDNA 4930523O13 gene	4930523O13Rik
1431455_at	1.02444	RIKEN cDNA 4933401B01 gene	4933401B01Rik
1431456_at	0.96645	RIKEN cDNA 4931439C15 gene	4931439C15Rik
1431457_at	0.97955	RIKEN cDNA 7420700N18 gene	7420700N18Rik
1431458_at	0.93645	RIKEN cDNA 4921507L20 gene	4921507L20Rik
1431459_at	1.00165	RIKEN cDNA 9430013L14 gene	9430013L14Rik
1431460_at	0.90354	RIKEN cDNA 4930519D14 gene	4930519D14Rik
1431461_at	1.05748	RIKEN cDNA 4933417E11 gene	4933417E11Rik
1431462_at	0.94765	"protein phosphatase 1, regulatory (inhibitor) subunit 12B"	Ppp1r12b
1431463_at	1.00347	RIKEN cDNA 1700041C02 gene	1700041C02Rik
1431464_a_at	1.03094	phosphomannomutase 2	Pmm2
1431465_s_at	1.02717	forty-two-three domain containing 1	Fytd1
1431466_at	1.06664	RIKEN cDNA 4930553D19 gene	4930553D19Rik

1431467_at	0.96287	RIKEN cDNA 4931402G19 gene	4931402G19Rik
1431468_at	0.95464	cysteine-rich secretory protein 4	Crisp4
1431469_a_at	0.60163	CXXC finger 5 Cxxc5	
1431470_a_at	1.11302	RIKEN cDNA 4933414I15 gene	4933414I15Rik
1431471_at	1.07594	RIKEN cDNA 4833428M15 gene	4833428M15Rik
1431472_at	1.05085	RIKEN cDNA 2310043D23 gene	2310043D23Rik
1431473_at	1.13103	RIKEN cDNA 5330423I11 gene	5330423I11Rik
1431474_at	0.96587	RIKEN cDNA 4833414L08 gene	4833414L08Rik
1431475_a_at	0.91140	homeo box A10 Hoxa10	
1431476_at	0.95165	RIKEN cDNA 4933407I05 gene	4933407I05Rik
1431477_at	0.99865	RIKEN cDNA 4921515L22 gene	4921515L22Rik
1431478_at	1.01140	RIKEN cDNA 3110035G12 gene	3110035G12Rik
1431479_at	0.97085	RIKEN cDNA 4921524L21 gene	4921524L21Rik
1431480_at	1.02914	RIKEN cDNA 4833419E13 gene	4833419E13Rik
1431481_at	0.98983	RIKEN cDNA 4930443O20 gene	4930443O20Rik
1431482_at	1.07206	RIKEN cDNA 2900012M01 gene	2900012M01Rik
1431483_at	1.00207	RIKEN cDNA 4930544M13 gene	4930544M13Rik
1431484_at	1.03311	RIKEN cDNA 2010007L08 gene	2010007L08Rik
1431485_at	1.12904	RIKEN cDNA 4833447I15 gene	4833447I15Rik
1431486_at	0.92892	RIKEN cDNA 4930503E24 gene	4930503E24Rik
1431487_at	0.99441	kinesin light chain 2 Klc2	
1431488_at	0.96456	RIKEN cDNA 4833428L15 gene	4833428L15Rik
1431489_at	0.99270	RIKEN cDNA 4933432K03 gene	4933432K03Rik
1431490_at	0.92132	RIKEN cDNA A930033C23 gene	A930033C23Rik
1431491_at	1.01854	RIKEN cDNA 9430087N24 gene	9430087N24Rik
1431492_at	1.02236	RIKEN cDNA 2210418G03 gene	2210418G03Rik
1431493_at	1.01259	RIKEN cDNA 9530046B11 gene	9530046B11Rik
1431494_at	1.15373	RIKEN cDNA A930008B05 gene	A930008B05Rik
1431495_at	0.99004	RIKEN cDNA 6030440G07 gene	6030440G07Rik
1431496_at	1.08784	RIKEN cDNA B230104F01 gene	B230104F01Rik
1431497_at	0.88007	RIKEN cDNA C330022C24 gene	C330022C24Rik
1431498_at	1.03223	RIKEN cDNA 9530097N15 gene	9530097N15Rik
1431499_at	0.95843	RIKEN cDNA 4933436F18 gene	4933436F18Rik
1431500_at	0.93558	RIKEN cDNA 4933406J10 gene	4933406J10Rik
1431501_at	0.97589	RIKEN cDNA 1700037C18 gene	1700037C18Rik
1431502_a_at	1.21255	exosome component 3 Exosc3	
1431503_at	1.03874	RIKEN cDNA 4930513O06 gene	4930513O06Rik
1431504_at	1.00385	RIKEN cDNA 2310061N02 gene	2310061N02Rik
1431505_at	1.05219	similar to Peptidyl-prolyl cis-trans isomerase H (PPIase H) (Rotamase H) /// similar to Peptidyl-prolyl cis-trans isomerase H (PPIase H) (Rotamase H) /// similar to Peptidyl-prolyl cis-trans isomerase H (PPIase H) (Rotamase H) LOC629952 /// LOC666411 /// LOC671009	
1431506_s_at	1.06499	peptidyl prolyl isomerase H /// similar to peptidyl prolyl isomerase H /// similar to Peptidyl-prolyl cis-trans isomerase H (PPIase H) (Rotamase H) /// similar to peptidylprolyl isomerase	

H /// similar to Peptidyl-prolyl cis-trans isomerase H (PPIase H) (Rotamase H) /// similar to peptidylprolyl isomerase H /// similar to Peptidyl-prolyl cis-trans isomerase H (PPIase H) (Rotamase H) /// similar to peptidylprolyl isomerase H Ppih /// LOC433064 /// LOC629952 /// LOC665989 /// LOC666411 /// LOC669441 /// LOC671009 /// LOC676493

1431507_a_at 1.01097 synaptojanin 2 binding protein /// RIKEN cDNA 1810020G14 gene Synj2bp /// 1810020G14Rik

1431508_at 0.93134 RIKEN cDNA 4930423M02 gene 4930423M02Rik

1431509_at 0.92683 RIKEN cDNA 4933404G15 gene 4933404G15Rik

1431510_s_at 0.99689 RIKEN cDNA 2010110K16 gene 2010110K16Rik

1431511_at 0.93035 RIKEN cDNA 9230110F15 gene 9230110F15Rik

1431512_at 0.99498 RIKEN cDNA 4933437F24 gene 4933437F24Rik

1431513_at 0.92495 RIKEN cDNA 4921537P18 gene 4921537P18Rik

1431514_at 0.65876 RIKEN cDNA 5730453C05 gene 5730453C05Rik

1431515_at 0.95043 RIKEN cDNA 4932414N04 gene 4932414N04Rik

1431516_at 0.77235 RIKEN cDNA 4930565N06 gene 4930565N06Rik

1431517_at 0.90558 RIKEN cDNA 4933417C20 gene 4933417C20Rik

1431518_at 0.94908 patched domain containing 3 Ptchd3

1431519_at 0.98007 RIKEN cDNA 4933406L23 gene 4933406L23Rik

1431520_at 1.02687 RIKEN cDNA 4933406J09 gene 4933406J09Rik

1431521_at 1.04205 RIKEN cDNA 5033428B15 gene 5033428B15Rik

1431522_at 0.97773 RIKEN cDNA A930001A20 gene A930001A20Rik

1431523_at 0.91078 RIKEN cDNA 4930564K09 gene 4930564K09Rik

1431524_at 0.94850 RIKEN cDNA 4933427I22 gene 4933427I22Rik

1431525_at 1.10602 RIKEN cDNA 9130002K18 gene 9130002K18Rik

1431526_at 0.92169 hypothetical protein LOC672430 LOC672430

1431527_at 0.96510 CD164 antigen Cd164

1431528_at 1.58326 RIKEN cDNA 5830427D02 gene 5830427D02Rik

1431529_at 1.09869 RIKEN cDNA A930002H02 gene A930002H02Rik

1431530_a_at 0.85549 tetraspanin 5 Tspan5

1431531_at 0.88556 RIKEN cDNA 1700025D03 gene 1700025D03Rik

1431532_at 0.97405 RIKEN cDNA 4931400O07 gene 4931400O07Rik

1431533_at 0.88868 RIKEN cDNA 4921530D09 gene 4921530D09Rik

1431534_at 0.95036 protein kinase C and casein kinase substrate in neurons 3 Pacsin3

1431535_at 0.99958 phospholipase B1 Plb1

1431536_at 1.10426 RIKEN cDNA 4933432K03 gene 4933432K03Rik

1431537_at 0.86917 RIKEN cDNA 2010001E11 gene 2010001E11Rik

1431538_at 0.96769 RIKEN cDNA 4931415C17 gene 4931415C17Rik

1431539_at 0.91577 "SAM domain, SH3 domain and nuclear localization signals, 1" Samsn1

1431540_at 0.94944 RIKEN cDNA 4930511E03 gene 4930511E03Rik

1431541_at 0.94616 "STT3, subunit of the oligosaccharyltransferase complex, homolog B (S. cerevisiae)" Stt3b

1431542_at 0.98081 presenilin 2 Psen2

1431543_at 0.81193 RIKEN cDNA 9130001E16 gene 9130001E16Rik

1431544_at	1.20823	RIKEN cDNA 4930524B17	gene	4930524B17Rik
1431545_at	1.11257	RIKEN cDNA 4921532D01	gene	4921532D01Rik
1431546_at	1.02814	RIKEN cDNA 4930509K18	gene	4930509K18Rik
1431547_at	1.01781	RIKEN cDNA 4930435F18	gene	4930435F18Rik
1431548_at	0.98976	RIKEN cDNA 4933424G05	gene	4933424G05Rik
1431549_at	1.18940	---	---	
1431550_at	1.04409	RIKEN cDNA 4930488B22	gene	4930488B22Rik
1431551_at	1.09440	RIKEN cDNA 2610028D06	gene	2610028D06Rik
1431552_at	0.90680	RIKEN cDNA 1810006J02	gene	1810006J02Rik
1431553_at	0.91364	---	---	
1431554_a_at	1.18781	annexin A9	Anxa9	
1431555_at	0.91852	RIKEN cDNA 4933409F18	gene	4933409F18Rik
1431556_at	0.97825	RIKEN cDNA 4930435M08	gene	4930435M08Rik
1431557_at	1.02000	RIKEN cDNA 4930488N15	gene	4930488N15Rik
1431558_at	1.07164	RIKEN cDNA 2310016D03	gene	2310016D03Rik
1431559_at	1.06779	RIKEN cDNA 6430710C18	gene	6430710C18Rik
1431560_at	1.00806	dephospho-CoA kinase domain containing	Dcakd	
1431561_a_at	0.93511	DEAH (Asp-Glu-Ala-His) box polypeptide 34	Dhx34	
1431562_at	0.88355	RIKEN cDNA 4921513I03	gene	4921513I03Rik
1431563_at	1.02374	RIKEN cDNA 2010003K10	gene	2010003K10Rik
1431564_at	1.06032	breast cancer anti-estrogen resistance 3	Bcar3	
1431565_at	0.96291	RIKEN cDNA 4930511J24	gene	4930511J24Rik
1431566_at	0.98658	---	---	
1431567_at	0.98934	RIKEN cDNA 1700018A04	gene	1700018A04Rik
1431568_at	1.00766	RIKEN cDNA 4933427E13	gene	4933427E13Rik
1431569_a_at	1.07056	Ly6/Plaur domain containing 1	Lypd1	
1431570_at	1.23410	RIKEN cDNA 4930588G17	gene	4930588G17Rik
1431571_at	0.97356	RIKEN cDNA 4933437I04	gene	4933437I04Rik
1431572_at	0.96193	RIKEN cDNA 2310015K22	gene	2310015K22Rik
1431573_at	0.85593	spermidine/spermine N1-acetyl transferase-like 1	Satl1	
1431574_at	0.80953	RIKEN cDNA 4930465M20	gene	4930465M20Rik
1431575_at	0.98557	"C-type lectin domain family 4, member g"	Clec4g	
1431576_at	1.01359	RIKEN cDNA 4933404K08	gene	4933404K08Rik
1431577_at	1.01696	---	---	
1431578_at	1.00403	WBP2 N-terminal like	Wbp2nl	
1431579_at	0.81512	RIKEN cDNA 4933421D24	gene	4933421D24Rik
1431580_at	0.95259	RIKEN cDNA 4833421K07	gene	4833421K07Rik
1431581_at	1.13416	RIKEN cDNA 4922502B01	gene	4922502B01Rik
1431582_at	1.17567	integrin beta 4 binding protein	Itgb4bp	
1431583_at	1.05923	RIKEN cDNA 0610038L08	gene	0610038L08Rik
1431584_at	1.13156	RIKEN cDNA 1110028F18	gene	1110028F18Rik
1431585_at	1.02095	RIKEN cDNA 1700034P13	gene	1700034P13Rik
1431586_at	0.92890	RIKEN cDNA 4921529N20	gene	4921529N20Rik

1431587_at 0.93167 coiled-coil domain containing 7 Ccdc7
1431588_at 1.04914 RIKEN cDNA 2900045O20 gene 2900045O20Rik
1431589_at 0.98603 hydrocephalus inducing Hydin
1431590_at 1.00237 RIKEN cDNA 4930521C21 gene 4930521C21Rik
1431591_s_at 1.61175 hypothetical protein LOC677168 LOC677168
1431592_a_at 0.85860 SH3-domain kinase binding protein 1 Sh3kbp1
1431593_a_at 0.87312 "tRNA splicing endonuclease 34 homolog (SEN34, *S. cerevisiae*)" Tsen34
1431594_at 0.93611 dipeptidylpeptidase 3 Dpp3
1431595_at 0.99670 RIKEN cDNA 2900064B16 gene 2900064B16Rik
1431596_at 1.06601 --- ---
1431597_a_at 0.94488 nuclear receptor interacting protein 3 Nrip3
1431598_a_at 1.11508 LIM homeobox protein 9 Lhx9
1431599_at 1.00092 RIKEN cDNA 2310069B03 gene 2310069B03Rik
1431600_at 0.97314 RIKEN cDNA 4930564C03 gene 4930564C03Rik
1431601_at 0.83852 rhabdoid tumor deletion region gene 1 Rtdr1
1431602_a_at 0.80726 UDP-glucose ceramide glucosyltransferase-like 2 Ugcgl2
1431603_at 0.97970 RIKEN cDNA 4931407E12 gene 4931407E12Rik
1431604_a_at 0.95261 coiled-coil domain containing 97 Ccdc97
1431605_at 1.02305 RB1-inducible coiled-coil 1 Rb1cc1
1431606_a_at 0.98202 angel homolog 2 (*Drosophila*) Angel2
1431607_at 0.98213 lysophospholipase 1 Lypl1
1431608_at 1.03745 amyotrophic lateral sclerosis 2 (juvenile) homolog (human) Als2
1431609_a_at 0.84960 "acid phosphatase 5, tartrate resistant" Acp5
1431610_at 1.00494 RIKEN cDNA 5330439A09 gene 5330439A09Rik
1431611_a_at 0.84476 "immunoglobulin superfamily, member 4A" Igsf4a
1431612_at 1.02887 RIKEN cDNA 1700007J10 gene 1700007J10Rik
1431613_a_at 0.91227 RIKEN cDNA 1700024D23 gene 1700024D23Rik
1431614_at 0.93186 RIKEN cDNA 2810049E08 gene 2810049E08Rik
1431615_at 0.93024 RIKEN cDNA 4930558J22 gene 4930558J22Rik
1431616_at 1.02986 RIKEN cDNA 1700067C01 gene 1700067C01Rik
1431617_at 1.09086 RIKEN cDNA 4933405E24 gene 4933405E24Rik
1431618_a_at 0.83586 "DNA segment, Chr 14, ERATO Doi 581, expressed" D14Ertd581e
1431619_a_at 1.10881 dystrobrevin binding protein 1 Dtnbp1
1431620_at 0.96691 RIKEN cDNA 4930442J19 gene 4930442J19Rik
1431621_at 0.90029 RIKEN cDNA 4930420G21 gene 4930420G21Rik
1431622_at 0.97447 RIKEN cDNA 4933406K04 gene 4933406K04Rik
1431623_at 1.20376 RIKEN cDNA 1700023H06 gene 1700023H06Rik
1431624_a_at 0.95846 RIKEN cDNA 2610206B13 gene 2610206B13Rik
1431625_at 1.23621 "phosphodiesterase 1A, calmodulin-dependent" Pde1a
1431626_at 0.96025 RIKEN cDNA 4933429K18 gene 4933429K18Rik
1431627_at 0.88331 kinesin family member 27 Kif27
1431628_at 1.11498 RIKEN cDNA 4930435H24 gene 4930435H24Rik
1431629_at 0.94935 RIKEN cDNA 9130221D24 gene 9130221D24Rik

1431630_a_at	0.91419	Kruppel-like factor 3 (basic)	Klf3
1431631_at	0.90325	RIKEN cDNA 2900057B20	gene 2900057B20Rik
1431632_at	0.92280	RIKEN cDNA 4930526L06	gene 4930526L06Rik
1431633_x_at	0.96142	RIKEN cDNA 4930526L06	gene 4930526L06Rik
1431634_at	0.95964	RIKEN cDNA 4930455C13	gene 4930455C13Rik
1431635_at	1.03263	RIKEN cDNA 4930412F09	gene 4930412F09Rik
1431636_at	1.07611	RIKEN cDNA 4930558K02	gene 4930558K02Rik
1431637_at	1.03108	---	---
1431638_at	0.97717	RIKEN cDNA 4930592A05	gene 4930592A05Rik
1431639_at	1.05003	RIKEN cDNA 4930468A15	gene 4930468A15Rik
1431640_at	0.95326	RIKEN cDNA 4933431J24	gene 4933431J24Rik
1431641_at	0.90731	RIKEN cDNA 9330161L09	gene 9330161L09Rik
1431642_at	1.00327	"eukaryotic translation initiation factor 2, subunit 3, structural gene Y-linked"	Eif2s3y
1431643_at	1.05235	"spermatogenesis associated glutamate (E)-rich protein 6, pseudogene 1"	Speer6-ps1
1431644_a_at	0.90261	islet cell autoantigen 1	Ica1
1431645_a_at	0.89563	guanosine diphosphate (GDP) dissociation inhibitor 2	Gdi2
1431646_a_at	0.92786	syntaxin 6	Stx6
1431647_a_at	1.02119	CEA-related cell adhesion molecule 13	Ceacam13
1431648_at	1.04851	RIKEN cDNA 4930528F23	gene 4930528F23Rik
1431649_at	1.00714	RIKEN cDNA 1700108M19	gene 1700108M19Rik
1431650_at	1.00969	RIKEN cDNA 1110054P19	gene 1110054P19Rik
1431651_at	0.99422	RIKEN cDNA C330001K17	gene C330001K17Rik
1431652_at	1.09840	uridine monophosphate synthetase	Umps
1431653_at	1.06462	"T-cell receptor beta, joining region"	Tcrb-J
1431654_at	0.90030	RIKEN cDNA 1700112M01	gene 1700112M01Rik
1431655_a_at	0.91237	multiple substrate lipid kinase	Mulk
1431656_at	0.98001	RIKEN cDNA C630050I24	gene C630050I24Rik
1431657_at	0.87528	RIKEN cDNA 9230112D13	gene 9230112D13Rik
1431658_at	0.97546	"histone 1, H4j"	Hist1h4j
1431659_at	1.12528	transmembrane protein 86B	Tmem86b
1431660_at	0.95966	RIKEN cDNA 4930564D02	gene 4930564D02Rik
1431661_at	1.05333	RIKEN cDNA 4930562A09	gene 4930562A09Rik
1431662_at	0.99554	RIKEN cDNA 4930555F03	gene 4930555F03Rik
1431663_a_at	0.93498	ciliary neurotrophic factor receptor	Cntfr
1431664_at	1.23751	Luc7 homolog (S. cerevisiae)-like	Luc7l
1431665_a_at	1.08428	translocase of inner mitochondrial membrane 8 homolog b (yeast)	Timm8b
1431666_at	0.99362	RIKEN cDNA 2610021K21	gene 2610021K21Rik
1431667_s_at	1.01287	RIKEN cDNA 2610021K21	gene 2610021K21Rik
1431668_at	1.02243	"capping protein (actin filament) muscle Z-line, beta"	Capzb
1431669_at	0.86068	RIKEN cDNA 4930583I09	gene 4930583I09Rik
1431670_at	0.98887	"NADH dehydrogenase (ubiquinone) 1, subcomplex unknown, 2"	Ndufc2

1431671_at	0.88038	RIKEN cDNA 4930447A16 gene	4930447A16Rik
1431672_at	0.99657	RIKEN cDNA 9430069I07 gene	9430069I07Rik
1431673_at	0.99226	RIKEN cDNA 4930557A04 gene	4930557A04Rik
1431674_at	0.84559	RIKEN cDNA 2610303G11 gene	2610303G11Rik
1431675_a_at	0.80433	general transcription factor II I	Gtf2i
1431676_x_at	0.90755	general transcription factor II I	Gtf2i
1431677_at	1.31617	RIKEN cDNA 4631424J17 gene	4631424J17Rik
1431678_at	0.95769	RIKEN cDNA 4930556N08 gene	4930556N08Rik
1431679_at	0.90934	RIKEN cDNA 2510042H12 gene	2510042H12Rik
1431680_a_at	0.74485	"protein tyrosine phosphatase, receptor type, K"	Ptprk
1431681_at	0.86177	kallikrein 9	Klk9
1431682_at	1.00313	RIKEN cDNA 4930594C11 gene	4930594C11Rik
1431683_at	0.93551	RIKEN cDNA 4930473H19 gene	4930473H19Rik
1431684_at	1.00478	RIKEN cDNA 4933402J24 gene	4933402J24Rik
1431685_at	1.03855	RIKEN cDNA 4930549O18 gene	4930549O18Rik
1431686_a_at	1.14706	"glia maturation factor, beta"	Gmfb
1431687_at	0.97950	"glia maturation factor, beta"	Gmfb
1431688_at	0.95644	hypothetical LOC73899	LOC73899
1431689_at	1.27575	RIKEN cDNA 5830400J07 gene	5830400J07Rik
1431690_at	0.91400	"potassium voltage-gated channel, subfamily H (eag-related), member 3"	Kcnh3
1431691_a_at	0.88573	"RAB31, member RAS oncogene family"	Rab31
1431692_a_at	0.83295	Casitas B-lineage lymphoma c	Cblc
1431693_a_at	1.08423	interleukin 17B II17b	
1431694_a_at	0.99688	catenin beta interacting protein 1	Ctnnbip1
1431695_at	0.90475	rabphilin 3A-like (without C2 domains)	Rph3al
1431696_at	1.04771	RIKEN cDNA 4632413K17 gene	4632413K17Rik
1431697_at	0.96320	synaptojanin 2	Synj2
1431698_at	1.06024	"ubiquitin associated and SH3 domain containing, A"	Ubash3a
1431699_at	0.82630	"fucosyltransferase 4, pseudogene 1"	Fut4-ps1
1431700_at	1.07848	"glutamate receptor, ionotropic, NMDA2B (epsilon 2)"	Grin2b
1431701_a_at	1.02181	PDZ domain containing 1	Pdzk1
1431702_at	1.12112	---	---
1431703_at	1.03164	RIKEN cDNA 2900027M19 gene	2900027M19Rik
1431704_a_at	0.94582	Ral GEF with PH domain and SH3 binding motif 2	Ralgps2
1431705_a_at	1.00506	mucolipin 2	Mcoln2
1431706_at	1.07771	RAN binding protein 5	Ranbp5
1431707_a_at	1.01540	"pleckstrin homology, Sec7 and coiled-coil domains 3"	Pscd3
1431708_a_at	1.52890	cytotoxic granule-associated RNA binding protein 1	Tia1
1431709_at	0.90700	"metallothionein-like 5, testis-specific (tesmin)"	Mtl5
1431710_at	0.93757	CKLF-like MARVEL transmembrane domain containing 2A	Cmtm2a
1431711_a_at	0.94666	RIKEN cDNA 9030409G11 gene	9030409G11Rik
1431712_a_at	0.99337	RIKEN cDNA 2310022A10 gene	2310022A10Rik
1431713_at	1.04475	suppressor of fused homolog (Drosophila)	Sufu

1431714_at 1.02165 RIKEN cDNA 2310015D24 gene 2310015D24Rik
1431715_a_at 0.91588 "ligase III, DNA, ATP-dependent" Lig3
1431716_at 0.88800 hect domain and RLD 4 Herc4
1431717_at 1.09950 RIKEN cDNA 3526401B18 gene 3526401B18Rik
1431718_at 0.85754 kinesin family member 15 Kif15
1431719_a_at 0.89722 SLIT-ROBO Rho GTPase activating protein 1 Srgap1
1431720_at 1.06847 RIKEN cDNA 2810404F17 gene 2810404F17Rik
1431721_a_at 0.89990 "protein Z, vitamin K-dependent plasma glycoprotein" Proz
1431722_a_at 1.07362 arylformamidase Afmid
1431723_at 1.13823 RIKEN cDNA 1600022D10 gene 1600022D10Rik
1431724_a_at 1.06035 "purinergic receptor P2Y, G-protein coupled 12" P2ry12
1431725_at 1.06318 formin 2 Fmn2
1431726_a_at 0.92628 transmembrane protein 80 Tmem80
1431727_at 1.06759 submandibular gland protein C Smgc
1431728_at 1.13662 RIKEN cDNA 2310010I16 gene 2310010I16Rik
1431729_at 1.05008 spermatogenesis associated 17 Spata17
1431730_at 0.89871 RIKEN cDNA 4921506M07 gene 4921506M07Rik
1431731_at 1.00538 "BCL2/adenovirus E1B interacting protein 1, NIP2" Bnip2
1431732_at 1.10148 sperm associated antigen 16 Spag16
1431733_at 0.95932 alpha-kinase 1 Alpk1
1431734_a_at 0.81682 "DnaJ (Hsp40) homolog, subfamily B, member 4" Dnajb4
1431735_at 1.12714 RIKEN cDNA 4930469G21 gene 4930469G21Rik
1431736_at 0.95267 RIKEN cDNA 4933405D12 gene 4933405D12Rik
1431737_at 1.00285 hypothetical LOC629820 LOC629820
1431738_at 0.93111 RIKEN cDNA 4921531P14 gene 4921531P14Rik
1431739_at 1.00893 mitochondrial translation optimization 1 homolog (*S. cerevisiae*) Mto1
1431740_at 1.03916 "solute carrier family 7, (cationic amino acid transporter, y+ system) member 13" Slc7a13
1431741_a_at 0.89812 RIKEN cDNA 4930408O21 gene 4930408O21Rik
1431742_at 1.06697 RIKEN cDNA 1810053B23 gene 1810053B23Rik
1431743_a_at 0.93519 "solute carrier family 4 (anion exchanger), member 1" Slc4a1
1431744_a_at 0.98643 stromal membrane-associated protein 1 Smap1
1431745_a_at 0.96392 zinc finger CCCH type containing 14 Zc3h14
1431746_a_at 0.96277 ubiquitin-activating enzyme E1C Ube1c
1431747_at 1.12861 "ZW10 homolog (*Drosophila*), centromere/kinetochore protein" Zw10
1431748_a_at 0.84494 ring finger and FYVE like domain containing protein Rffl
1431749_a_at 1.09399 RAS guanyl releasing protein 1 Rasgrp1
1431750_at 0.89052 "Usher syndrome 2A (autosomal recessive, mild) homolog (human)" Ush2a
1431751_a_at 0.94695 metallophosphoesterase domain containing 2 Mpped2
1431752_a_at 0.96942 RIKEN cDNA 2900073H19 gene 2900073H19Rik
1431753_x_at 0.95789 RIKEN cDNA 2900073H19 gene 2900073H19Rik
1431754_at 1.04478 RIKEN cDNA 4930578N16 gene 4930578N16Rik
1431755_a_at 0.87831 coiled-coil domain containing 49 Ccdc49

1431756_at 1.08534 "Adult male testis cDNA, RIKEN full-length enriched library,
clone:4930516B21 product:unclassifiable, full insert sequence" ---

1431757_s_at 0.99141 RIKEN cDNA 5033417F24 gene 5033417F24Rik

1431758_at 0.91781 cDNA sequence BC051628 BC051628

1431759_at 0.91679 "zinc finger, BED domain containing 4 /// RIKEN cDNA 4930529C04 gene ///
similar to zinc finger, BED domain containing 4" Zbed4 /// 4930529C04Rik /// LOC669068

1431760_a_at 0.91291 serologically defined colon cancer antigen 3 Sdccag3

1431761_at 0.95754 ectonucleoside triphosphate diphosphohydrolase 4 Entpd4

1431762_at 0.84041 HtrA serine peptidase 3 Htra3

1431763_a_at 1.04354 chymotrypsin-like Ctrl

1431764_at 1.07241 RIKEN cDNA 2310067P03 gene 2310067P03Rik

1431765_a_at 1.04427 ribosomal protein S2 /// similar to 40S ribosomal protein S2 /// similar to 40S
ribosomal protein S2 /// similar to 40S ribosomal protein S2 /// similar to 40S ribosomal protein S2 ///
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ribosomal protein S2 /// similar to 40S ribosomal protein S2 /// similar to 40S ribosomal protein S2
Rps2 /// LOC243642 /// LOC544872 /// LOC545808 /// LOC546164 /// LOC546298 /// LOC620248 ///
LOC620559 /// LOC621842 /// LOC623242 /// LOC628962 /// LOC629408 /// LOC634678 ///
LOC635931 /// LOC637606 /// LOC639043 /// LOC639606 /// LOC665951 /// LOC666668 ///
LOC666682 /// LOC667093 /// LOC667116 /// LOC667279 /// LOC667847 /// LOC669028 ///
LOC669347 /// LOC669682 /// LOC670133 /// LOC670228 /// LOC670479 /// LOC670968 ///
LOC671469 /// LOC671914 /// LOC671929 /// LOC672225 /// LOC672467 /// LOC674219 ///
LOC674755 /// LOC675043 /// LOC676775 /// LOC676958 /// LOC677135

1431766_x_at 1.04941 ribosomal protein S2 Rps2

1431767_at 1.07365 RIKEN cDNA 4930442G15 gene 4930442G15Rik

1431768_a_at 1.08693 protein arginine N-methyltransferase 3 Prmt3

1431769_at 0.98200 RIKEN cDNA 2210406O10 gene 2210406O10Rik

1431770_at 0.98163 calcium homeostasis endoplasmic reticulum protein Cherp

1431771_a_at 1.00124 interleukin-1 receptor-associated kinase 1 binding protein 1 Irak1bp1

1431772_a_at 1.08106 intersectin 2 Itsn2

1431773_at 1.07927 chromobox homolog 1 (Drosophila HP1 beta) Cbx1

1431774_a_at 1.05224 RIKEN cDNA 4930404J24 gene 4930404J24Rik

1431775_at	0.84819	RIKEN cDNA 3100002H09 gene	3100002H09Rik	
1431776_at	0.87846	"transducer of ERBB2, 2"	Tob2	
1431777_a_at	0.98421	high mobility group nucleosomal binding domain 3		Hmgn3
1431778_at	1.01513	RIKEN cDNA 2310051N18 gene	2310051N18Rik	
1431779_at	0.95689	---	---	
1431780_at	1.08007	RIKEN cDNA 1700021K14 gene	1700021K14Rik	
1431781_at	1.09224	yippee-like 1 (Drosophila)	Ypell	
1431782_s_at	0.91446	yippee-like 1 (Drosophila)	Ypell	
1431783_at	1.02891	RIKEN cDNA 1700010L04 gene	1700010L04Rik	
1431784_a_at	1.12611	brix domain containing 5	Bxdc5	
1431785_at	0.99598	ribonuclease T2 Rnaset2		
1431786_s_at	0.91921	RIKEN cDNA 1190003J15 gene	1190003J15Rik	
1431787_at	0.97864	RIKEN cDNA 3300002A11 gene	3300002A11Rik	
1431788_at	1.10551	RIKEN cDNA 1700008G05 gene	1700008G05Rik	
1431789_s_at	1.24835	transmembrane emp24 protein transport domain containing 5		Tmed5
1431790_at	0.93968	RIKEN cDNA 4932412D23 gene	4932412D23Rik	
1431791_a_at	0.79892	"protein tyrosine phosphatase, non-receptor type 13"	Ptpn13	
1431792_a_at	0.74605	serine/threonine kinase 11 interacting protein	Stk11ip	
1431793_at	1.07683	RIKEN cDNA 3732407C23 gene	3732407C23Rik	
1431794_at	0.97956	RIKEN cDNA 4932434E15 gene	4932434E15Rik	
1431795_a_at	0.89345	"sema domain, immunoglobulin domain (Ig), short basic domain, secreted, (semaphorin) 3B"	Sema3b	
1431796_at	0.94702	RIKEN cDNA 2810430I11 gene	2810430I11Rik	
1431797_at	0.98299	RIKEN cDNA 4921518K17 gene	4921518K17Rik	
1431798_a_at	1.11429	"synapse defective 1, Rho GTPase, homolog 1 (C. elegans)"	Syde1	
1431799_at	1.05114	myeloid/lymphoid or mixed lineage-leukemia translocation to 4 homolog (Drosophila)	Mllt4	
1431800_at	0.97533	RIKEN cDNA 1300010F03 gene	1300010F03Rik	
1431801_at	1.02784	Down syndrome cell adhesion molecule	Dscam	
1431802_a_at	1.19720	"DNA segment, Chr 5, Wayne State University 178, expressed"	D5Wsu178e	
1431803_at	0.91892	"cytochrome P450, family 2, subfamily d, polypeptide 13"	Cyp2d13	
1431804_a_at	0.89224	trans-acting transcription factor 3	Sp3	
1431805_a_at	1.40813	"rhopilin, Rho GTPase binding protein 2"	Rhpn2	
1431806_at	0.94492	RIKEN cDNA 4931408D14 gene	4931408D14Rik	
1431807_at	0.89715	F-box and WD-40 domain protein 8	Fbxw8	
1431808_a_at	0.67011	"inter alpha-trypsin inhibitor, heavy chain 4"	Itih4	
1431809_at	1.00182	RIKEN cDNA 4932442L08 gene	4932442L08Rik	
1431810_a_at	0.93222	transmembrane and coiled-coil domains 4	Tmco4	
1431811_a_at	0.96734	F-box only protein 34	Fbxo34	
1431812_a_at	0.87009	"solute carrier family 6 (neurotransmitter transporter, glycine), member 9"	Slc6a9	
1431813_at	1.10479	myosin XVIIIb	Myo18b	
1431814_at	0.87443	RIKEN cDNA 4932443L11 gene	4932443L11Rik	

1431815_a_at	1.02580	MMS19 (MET18 <i>S. cerevisiae</i>)-like	Mms19l
1431816_at	1.05182	testis expressed gene 9	Tex9
1431817_at	0.87888	"alcohol dehydrogenase 6 (class V), pseudogene 1"	Adh6-ps1
1431818_at	0.75295	RIKEN cDNA 1700012B15 gene	1700012B15Rik
1431819_at	1.04447	---	---
1431820_at	1.12054	RIKEN cDNA 4632404H12 gene	4632404H12Rik
1431821_a_at	0.90686	EPS8-like 1	Eps8l1
1431822_a_at	0.96689	5-azacytidine induced gene 2	Azi2
1431823_at	0.93411	modulator of apoptosis 1	Moap1
1431824_at	0.96618	cadherin 13	Cdh13
1431825_at	0.95518	serine/arginine-rich protein specific kinase 3	Srpk3
1431826_a_at	0.99503	BR serine/threonine kinase 2	Brsk2
1431827_a_at	1.07107	tousled-like kinase 2 (<i>Arabidopsis</i>)	Tlk2
1431828_a_at	0.92058	synaptojanin 2	Synj2
1431829_a_at	0.78982	ral guanine nucleotide dissociation stimulator-like 3	Rgl3
1431830_at	0.96910	zinc finger protein 329	Zfp329
1431831_at	1.12703	RIKEN cDNA 2210009G21 gene	2210009G21Rik
1431832_x_at	1.11039	exosome component 10	Exosc10
1431833_a_at	0.96593	3-hydroxy-3-methylglutaryl-Coenzyme A synthase 2	Hmgcs2
1431834_a_at	0.92370	elastin microfibril interfacer 1	Emilin1
1431835_at	0.99712	transcription elongation regulator 1-like	Tcerg1l
1431836_x_at	0.96734	RIKEN cDNA 5430432N15 gene	5430432N15Rik
1431837_at	1.01412	RIKEN cDNA 1200007C13 gene	1200007C13Rik
1431838_at	0.91045	RIKEN cDNA 1700128E19 gene	1700128E19Rik
1431839_a_at	0.95580	coiled-coil domain containing 81	Ccdc81
1431840_at	0.97179	DENN/MADD domain containing 1C	Dennd1c
1431841_at	0.92100	RIKEN cDNA 4930434E21 gene	4930434E21Rik
1431842_at	0.98654	RIKEN cDNA 4930422C21 gene	4930422C21Rik
1431843_a_at	1.29766	"nuclear factor of kappa light polypeptide gene enhancer in B-cells inhibitor, epsilon"	Nfkbie
1431844_at	0.97814	"potassium large conductance calcium-activated channel, subfamily M, beta member 2"	Kcnmb2
1431845_at	0.97112	RIKEN cDNA 2810016G10 gene	2810016G10Rik
1431846_at	1.14102	RIKEN cDNA 4930546C10 gene	4930546C10Rik
1431847_at	1.06592	RIKEN cDNA 5330439K02 gene	5330439K02Rik
1431848_at	1.05232	angiopoietin-like 2	Angptl2
1431849_at	0.80050	RIKEN cDNA 4933432B09 gene	4933432B09Rik
1431850_at	0.97848	RIKEN cDNA 3200001D21 gene	3200001D21Rik
1431851_at	0.88823	RIKEN cDNA 4930534P07 gene	4930534P07Rik
1431852_at	1.00472	PR domain containing 8	Prdm8
1431853_at	0.90169	RIKEN cDNA 4933413C19 gene	4933413C19Rik
1431854_a_at	0.93705	RIKEN cDNA 4930452B06 gene	4930452B06Rik
1431855_at	0.95331	RIKEN cDNA 4930570E03 gene	4930570E03Rik

1431856_a_at 0.80230 C1q and tumor necrosis factor related protein 6 C1qtnf6
1431857_at 0.97592 PHD finger protein 19 Phf19
1431858_at 0.91408 RIKEN cDNA 4930480K15 gene 4930480K15Rik
1431859_at 0.96227 RIKEN cDNA 4930414F18 gene 4930414F18Rik
1431860_at 0.99382 cDNA sequence BC026513 BC026513
1431861_a_at 0.99304 outer dense fiber of sperm tails 2-like Odf2l
1431862_x_at 0.92728 outer dense fiber of sperm tails 2-like Odf2l
1431863_at 1.04316 RIKEN cDNA 4931430N09 gene 4931430N09Rik
1431864_at 0.96098 F-box only protein 24 Fbxo24
1431865_a_at 0.97174 RIKEN cDNA 4933405K07 gene 4933405K07Rik
1431866_at 0.91307 RIKEN cDNA 4930425K24 gene 4930425K24Rik
1431867_a_at 0.83675 RIKEN cDNA 1700007B13 gene 1700007B13Rik
1431868_at 0.91514 A kinase (PRKA) anchor protein 4 Akap4
1431869_at 0.91638 RIKEN cDNA 5730419F03 gene 5730419F03Rik
1431870_at 0.93976 RIKEN cDNA 4930463O16 gene 4930463O16Rik
1431871_at 1.09944 thioredoxin domain containing 3 (spermatozoa) Txndc3
1431872_at 0.97543 RIKEN cDNA 4930449A18 gene 4930449A18Rik
1431873_a_at 0.89463 epsilon-tubulin 1 Tube1
1431874_at 1.02915 RIKEN cDNA 4931429L15 gene 4931429L15Rik
1431875_a_at 0.91790 E2F transcription factor 1 E2f1
1431876_at 1.17352 RIKEN cDNA 4931429P17 gene 4931429P17Rik
1431877_a_at 1.00001 grainyhead-like 2 (Drosophila) Grhl2
1431878_at 0.98347 grainyhead-like 2 (Drosophila) Grhl2
1431879_at 0.87222 RIKEN cDNA 9030417H13 gene 9030417H13Rik
1431880_at 1.35025 RIKEN cDNA 3100003M19 gene 3100003M19Rik
1431881_at 1.00325 RIKEN cDNA 2410024F20 gene 2410024F20Rik
1431882_at 1.02988 "amyotrophic lateral sclerosis 2 (juvenile) chromosome region, candidate 12 (human)" Als2cr12
1431883_at 0.91226 tachykinin 1 Tac1
1431884_at 1.00549 RIKEN cDNA 1110019B22 gene 1110019B22Rik
1431885_a_at 0.93934 MUS81 endonuclease homolog (yeast) Mus81
1431886_at 0.98197 endoplasmic reticulum (ER) to nucleus signalling 1 Ern1
1431887_at 0.92250 "RNA binding motif 31, Y-linked" Rbm31y
1431888_s_at 0.88118 pregnancy-specific glycoprotein 21 Psg21
1431889_x_at 0.98661 pregnancy-specific glycoprotein 21 Psg21
1431890_a_at 0.78255 myeloid/lymphoid or mixed lineage-leukemia translocation to 3 homolog (Drosophila) Mllt3
1431891_at 0.76856 "zinc finger, SWIM domain containing 5" Zswim5
1431892_a_at 0.97083 "phospholipase C, delta 3" Plcd3
1431893_a_at 1.05886 "prenyl (solanesyl) diphosphate synthase, subunit 1" Pdss1
1431894_at 1.39247 RIKEN cDNA 4833424O12 gene 4833424O12Rik
1431895_at 0.98272 RIKEN cDNA 4933434M16 gene 4933434M16Rik
1431896_at 0.93946 Six6 opposite strand transcript 1 Six6os1

1431897_at	1.04424	carbohydrate (N-acetylgalactosamine 4-0) sulfotransferase 9	Chst9
1431898_at	0.92421	RIKEN cDNA 4930403J07 gene	4930403J07Rik
1431899_at	0.95167	"NK6 transcription factor related, locus 3 (Drosophila)"	Nkx6-3
1431900_a_at	0.84991	forkhead box A3	Foxa3
1431901_a_at	0.94068	"6-phosphofructo-2-kinase/fructose-2,6-biphosphatase 2"	Pfkfb2
1431902_at	0.93952	RIKEN cDNA 4930401A09 gene	4930401A09Rik
1431903_at	1.01627	RIKEN cDNA 4930524C18 gene	4930524C18Rik
1431904_at	0.94681	RIKEN cDNA 4933427G17 gene	4933427G17Rik
1431905_s_at	0.92868	RIKEN cDNA 4933427G17 gene	4933427G17Rik
1431906_at	0.88270	RIKEN cDNA 2700068H02 gene	2700068H02Rik
1431907_at	1.08145	RIKEN cDNA 4933404M02 gene	4933404M02Rik
1431908_at	1.05279	RIKEN cDNA 4933408J17 gene	4933408J17Rik
1431909_at	0.98053	"zinc finger CCCH type, antiviral 1"	Zc3hav1
1431910_at	1.00951	RIKEN cDNA 5730460C07 gene	5730460C07Rik
1431911_at	0.90777	RIKEN cDNA 4931423N10 gene	4931423N10Rik
1431912_at	1.11996	RIKEN cDNA 2700089I24 gene	2700089I24Rik
1431913_a_at	0.87813	"phosphodiesterase 3A, cGMP inhibited"	Pde3a
1431914_at	0.98499	"phosphodiesterase 3A, cGMP inhibited"	Pde3a
1431915_at	1.00900	RIKEN cDNA 4930442E04 gene	4930442E04Rik
1431916_at	0.97969	RIKEN cDNA 9030618K22 gene	9030618K22Rik
1431917_at	0.80743	"phosphatidylinositol glycan anchor biosynthesis, class Q"	Pigq
1431918_at	0.98839	RIKEN cDNA 1700036A12 gene	1700036A12Rik
1431919_at	0.97889	rotatin	Rtnn
1431920_a_at	1.09863	RIKEN cDNA 2010004A03 gene	2010004A03Rik
1431921_a_at	1.00281	stromal antigen 1	Stag1
1431922_at	1.05876	intestine specific homeobox	Isx
1431923_at	1.15021	RIKEN cDNA 2810401C16 gene	2810401C16Rik
1431924_at	0.92500	RIKEN cDNA 4930432N10 gene	4930432N10Rik
1431925_at	1.01326	RIKEN cDNA 4933433H22 gene	4933433H22Rik
1431926_a_at	0.87092	RIKEN cDNA 1300010M03 gene	1300010M03Rik
1431927_at	0.99727	RIKEN cDNA 5033417F24 gene	5033417F24Rik
1431928_at	0.99279	titin	Ttn
1431929_a_at	0.92609	syntaxin 17	Stx17
1431930_x_at	0.99750	cardiolipin synthase 1	Crls1
1431931_a_at	0.99942	"RAB, member of RAS oncogene family-like 2A"	Rabl2a
1431932_s_at	0.96077	tripartite motif-containing 44	Trim44
1431933_a_at	1.04008	decapping enzyme	Dcp1a
1431934_at	0.87674	RIKEN cDNA 4930505O20 gene	4930505O20Rik
1431935_at	1.10675	lipoic acid synthetase	Lias
1431936_a_at	0.70812	neuraminidase 2	Neu2
1431937_at	0.92367	kinesin family member 24	Kif24
1431938_a_at	1.02830	phosphomannomutase 2	Pmm2
1431939_a_at	0.95978	myc induced nuclear antigen	Mina

1431940_at	1.31261	hypothetical protein LOC74457	LOC74457
1431941_at	1.07746	"solute carrier family 35, member B4"	Slc35b4
1431942_at	0.95043	WW domain-containing oxidoreductase	Wwox
1431943_at	1.03021	RIKEN cDNA 2210011K15 gene	2210011K15Rik
1431944_at	0.99305	RIKEN cDNA 4930484H19 gene	4930484H19Rik
1431945_at	0.98462	ADP-ribosylation factor-like 13A	Arl13a
1431946_a_at	0.88036	"amyloid beta (A4) precursor protein-binding, family A, member 1 binding protein"	Apba2bp
1431947_at	2.17703	low density lipoprotein receptor	Ldlr
1431948_a_at	1.06965	pantothenate kinase 2 (Hallervorden-Spatz syndrome)	Pank2
1431949_at	1.08677	RIKEN cDNA 4930544F09 gene	4930544F09Rik
1431950_at	1.06638	RIKEN cDNA 4930528G23 gene	4930528G23Rik
1431951_a_at	0.88310	ubiquitin specific peptidase 16	Usp16
1431952_at	1.02520	RIKEN cDNA 4930532I03 gene	4930532I03Rik
1431953_at	0.97548	"ATPase, aminophospholipid transporter-like, class I, type 8A, member 2"	Atp8a2
1431954_x_at	0.91933	RIKEN cDNA 4930431C11 gene	4930431C11Rik
1431955_at	1.06048	RIKEN cDNA 4930453H23 gene	4930453H23Rik
1431956_at	0.94321	RIKEN cDNA 5830453K13 gene	5830453K13Rik
1431957_at	1.01705	leucine rich repeat containing 44	Lrrc44
1431958_at	0.97209	sperm associated antigen 16	Spag16
1431959_x_at	0.93146	pregnancy-specific glycoprotein 21	Psg21
1431960_at	1.70856	WW domain-containing oxidoreductase	Wwox
1431961_at	0.97645	dedicator of cytokinesis 8	Dock8
1431962_a_at	0.94617	Stam binding protein	Stampbp
1431963_at	0.93146	RIKEN cDNA 4930556A20 gene	4930556A20Rik
1431964_at	1.09221	RIKEN cDNA 4921518J05 gene	4921518J05Rik
1431965_at	0.81117	RIKEN cDNA 4933411E08 gene	4933411E08Rik
1431966_at	0.96802	ATP/GTP binding protein-like 3	Agbl3
1431967_at	1.06803	RIKEN cDNA 1700006A11 gene	1700006A11Rik
1431968_at	0.99619	spermatogenesis associated 16	Spata16
1431969_at	0.96745	RIKEN cDNA 4930402D18 gene	4930402D18Rik
1431970_at	1.04650	transmembrane 7 superfamily member 4	Tm7sf4
1431971_at	0.91626	---	---
1431972_a_at	0.99400	granule cell antiserum positive 14	Gcap14
1431973_at	0.99500	septin 6	6-Sep
1431974_at	0.91791	RIKEN cDNA 4930443G12 gene	4930443G12Rik
1431975_at	0.86775	RIKEN cDNA 4930521O11 gene	4930521O11Rik
1431976_at	0.96952	RIKEN cDNA 4930526F13 gene	4930526F13Rik
1431977_at	0.81875	RIKEN cDNA 4933402J15 gene	4933402J15Rik
1431978_at	0.97783	RIKEN cDNA 8430437L04 gene	8430437L04Rik
1431979_at	0.96657	RIKEN cDNA 4930444M15 gene	4930444M15Rik
1431980_a_at	0.98716	arsenic (+3 oxidation state) methyltransferase	As3mt

1431981_at	1.01928	"hypoxia inducible factor 1, alpha subunit"	Hif1a
1431982_at	1.01813	Rho GTPase activating protein 19	Arhgap19
1431983_at	0.92284	myosin IIIA	Myo3a
1431984_at	0.92023	RIKEN cDNA 1700007F19 gene	1700007F19Rik
1431985_at	0.85890	RIKEN cDNA 2210010B09 gene	2210010B09Rik
1431986_at	1.08586	RIKEN cDNA 4933421A08 gene	4933421A08Rik
1431987_at	1.04181	RIKEN cDNA 4930510E17 gene	4930510E17Rik
1431988_at	0.87329	RIKEN cDNA 1700112E06 gene	1700112E06Rik
1431989_at	0.89351	RIKEN cDNA 4930433E13 gene	4930433E13Rik
1431990_at	0.92077	RIKEN cDNA 4930429D17 gene	4930429D17Rik
1431991_at	0.97144	RIKEN cDNA 2410004P03 gene	2410004P03Rik
1431992_at	0.81745	RIKEN cDNA 4930568K20 gene	4930568K20Rik
1431993_a_at	0.82929	ring finger protein 38	Rnf38
1431994_at	0.86613	RIKEN cDNA 4930432B10 gene	4930432B10Rik
1431995_at	1.00386	anthrax toxin receptor 1	Antxr1
1431996_at	0.94201	---	---
1431997_at	1.12001	RIKEN cDNA 3000002C10 gene	3000002C10Rik
1431998_at	1.04737	RIKEN cDNA 4930432L08 gene	4930432L08Rik
1431999_at	0.94918	polymerase (RNA) III (DNA directed) polypeptide F	Polr3f
1432000_a_at	1.08205	death effector domain-containing	Dedd
1432001_at	1.02406	"zinc finger, MYND domain containing 17"	Zmynd17
1432002_at	1.05195	RIKEN cDNA 2810471M01 gene	2810471M01Rik
1432003_a_at	0.95689	ring finger protein 41	Rnf41
1432004_a_at	0.92753	dynamamin 2	Dnm2
1432005_at	0.94075	dynamamin 2	Dnm2
1432006_at	1.45332	"adaptor protein complex AP-2, alpha 2 subunit"	Ap2a2
1432007_s_at	2.48757	"adaptor protein complex AP-2, alpha 2 subunit"	Ap2a2
1432008_at	0.99043	transmembrane protein 86B	Tmem86b
1432009_at	0.91043	RIKEN cDNA 4933415D12 gene	4933415D12Rik
1432010_at	0.90965	RIKEN cDNA 4933408N05 gene	4933408N05Rik
1432011_at	0.94964	RIKEN cDNA 2900052L18 gene	2900052L18Rik
1432012_a_at	0.95880	NOL1/NOP2/Sun domain family 6	Nsun6
1432013_a_at	1.16963	RIKEN cDNA 2610016C23 gene	2610016C23Rik
1432014_at	1.09102	RIKEN cDNA 6530409C15 gene	6530409C15Rik
1432015_a_at	0.97223	RIKEN cDNA 4933428M03 gene	4933428M03Rik
1432016_a_at	0.90584	isocitrate dehydrogenase 3 (NAD+) alpha	Idh3a
1432017_at	1.06830	huntingtin interacting protein 1	Hip1
1432018_at	1.17811	achaete-scute complex homolog-like 2 (Drosophila)	Ascl2
1432019_at	1.01320	TruB pseudouridine (psi) synthase homolog 1 (E. coli)	Trub1
1432020_at	0.94347	RIKEN cDNA 6330411D24 gene	6330411D24Rik
1432021_at	0.88984	copine III	Cpne3
1432022_at	1.12328	Cdc42 GTPase-activating protein	Cdgap
1432023_a_at	0.97452	RIKEN cDNA 1700019H03 gene	1700019H03Rik

1432024_at	0.95478	RIKEN cDNA 2310015L07	gene	2310015L07Rik	
1432025_at	0.99181	RIKEN cDNA 4933430A20	gene	4933430A20Rik	
1432026_a_at	0.99924	hect domain and RLD 5		Herc5	
1432027_a_at	0.88724	"TBC1 domain family, member 14"		Tbc1d14	
1432028_at	1.13733	RAB11 family interacting protein 2 (class I)		Rab11fip2	
1432029_a_at	0.97351	stromal membrane-associated protein 1		Smap1	
1432030_at	0.89255	RIKEN cDNA 4930434J06	gene	4930434J06Rik	
1432031_at	0.86157	RIKEN cDNA 4930563E18	gene	4930563E18Rik	
1432032_a_at	0.91770	artemin		Artn	
1432033_at	0.88946	netrin G2		Ntng2	
1432034_at	0.96070	neurogenin 3		Neurog3	
1432035_at	0.89405	RIKEN cDNA 4933427J07	gene	4933427J07Rik	
1432036_at	0.86001	RIKEN cDNA 4933436H12	gene	4933436H12Rik	
1432037_at	1.07375	"protein phosphatase 1, regulatory (inhibitor) subunit 9A"		Ppp1r9a	
1432038_at	0.99891	RIKEN cDNA 4930509J09	gene	4930509J09Rik	
1432039_a_at	1.10664	Lrp2 binding protein		Lrp2bp	
1432040_at	0.98178	KCNQ1 overlapping transcript 1		Kcnq1ot1	
1432041_at	0.90090	RIKEN cDNA 4930519A11	gene	4930519A11Rik	
1432042_a_at	1.08806	smu-1 suppressor of mec-8 and unc-52 homolog (C. elegans)		Smu1	
1432043_at	0.92986	smu-1 suppressor of mec-8 and unc-52 homolog (C. elegans)		Smu1	
1432044_a_at	1.05575	RIKEN cDNA 4921504E06	gene	4921504E06Rik	
1432045_at	0.96168	testis-specific serine kinase 5		Tssk5	
1432046_at	0.90349	RIKEN cDNA 4930404K06	gene	4930404K06Rik	
1432047_at	0.94365	DEP domain containing 2		Depdc2	
1432048_at	0.91449	RIKEN cDNA 6330565B04	gene	6330565B04Rik	
1432049_at	1.02529	RIKEN cDNA 4930422I22	gene	4930422I22Rik	
1432050_at	1.01139	RIKEN cDNA 4930571B16	gene	4930571B16Rik	
1432051_at	1.01496	coiled-coil domain containing 7		Ccdc7	
1432052_at	1.04625	Exosome component 1		Exosc1	
1432053_at	0.90228	"regulatory factor X, 4 (influences HLA class II expression)"		Rfx4	
1432054_at	1.02259	"golgi autoantigen, golgin subfamily a, 1"		Golga1	
1432055_at	0.90860	SAM and SH3 domain containing 1		Sash1	
1432056_at	0.96552	"carboxypeptidase, vitellogenic-like"		Cpvl	
1432057_a_at	0.86284	PR domain containing 5		Prdm5	
1432058_at	0.91804	RIKEN cDNA 4930474H20	gene	4930474H20Rik	
1432059_x_at	0.88610	RIKEN cDNA 5031425E22	gene	5031425E22Rik	
1432060_at	0.98519	IQ motif containing with AAA domain		Iqca	
1432061_at	0.98341	RIKEN cDNA 4933406B17	gene	4933406B17Rik	
1432062_at	0.91338	MAM domain containing glycosylphosphatidylinositol anchor 1		Mdga1	
1432063_at	0.98990	RIKEN cDNA 4833427F10	gene	4833427F10Rik	
1432064_at	0.94342	U box domain containing 5		Ubox5	
1432065_at	1.19771	RIKEN cDNA 3100003L13	gene	3100003L13Rik	
1432066_at	0.95226	"DnaJ (Hsp40) homolog, subfamily A, member 3"		Dnaja3	

1432067_at	1.12134	RIKEN cDNA 4930473O22	gene	4930473O22Rik	
1432068_a_at	1.01723	ectonucleoside triphosphate diphosphohydrolase 7			Entpd7
1432069_at	0.88696	RIKEN cDNA 1700023G09	gene	1700023G09Rik	
1432070_at	1.03527	RIKEN cDNA 4921511C10	gene	4921511C10Rik	
1432071_at	0.93038	RIKEN cDNA 6330437I11	gene	6330437I11Rik	
1432072_at	1.04005	kinesin family member 2A		Kif2a	
1432073_at	0.91721	RIKEN cDNA 1700113I22	gene	1700113I22Rik	
1432074_at	0.93601	RIKEN cDNA 4930517E11	gene	4930517E11Rik	
1432075_a_at	0.95244	tektin 1		Tekt1	
1432076_at	1.09602	RIKEN cDNA 4933430H16	gene	4933430H16Rik	
1432077_at	0.95838	protection of telomeres 1		Pot1	
1432078_at	1.03034	RIKEN cDNA 4930453L07	gene	4930453L07Rik	
1432079_at	1.03129	RIKEN cDNA 4933436N17	gene	4933436N17Rik	
1432080_s_at	1.02083	MYST histone acetyltransferase 1		Myst1	
1432081_at	1.07787	"cytochrome P450, family 1, subfamily b, polypeptide 1"		Cyp1b1	
1432082_at	1.05873	RIKEN cDNA 4933436I20	gene	4933436I20Rik	
1432083_a_at	1.01040	leucine rich repeat containing 23		Lrrc23	
1432084_at	1.01877	RIKEN cDNA 4921515G04	gene	4921515G04Rik	
1432085_at	1.00468	RIKEN cDNA 4930425O10	gene	4930425O10Rik	
1432086_a_at	0.90662	RIB43A domain with coiled-coils 2		Ribc2	
1432087_at	0.88359	RIKEN cDNA 4933406G16	gene	4933406G16Rik	
1432088_at	0.91837	ventricular zone expressed PH domain homolog 1 (zebrafish)		Veph1	
1432089_at	0.99404	RIKEN cDNA 4930550C17	gene	4930550C17Rik	
1432090_at	0.83762	coiled-coil domain containing 83		Ccdc83	
1432091_a_at	0.98878	RUN and TBC1 domain containing 3		Rutbc3	
1432092_a_at	1.00503	"GULP, engulfment adaptor PTB domain containing 1"		Gulp1	
1432093_at	1.03678	RIKEN cDNA 2310005A03	gene	2310005A03Rik	
1432094_a_at	1.01228	RIKEN cDNA 1700034M03	gene	1700034M03Rik	
1432095_at	1.04296	RIKEN cDNA 4921534H16	gene	4921534H16Rik	
1432096_at	0.90773	small nuclear ribonucleoprotein N		Snrpn	
1432097_a_at	0.88514	"DNA cross-link repair 1A, PSO2 homolog (S. cerevisiae)"		Dclre1a	
1432098_a_at	0.98012	olfactory receptor 701		Olf701	
1432099_a_at	0.97653	proline dehydrogenase (oxidase) 2		Prodh2	
1432100_a_at	1.12782	RIKEN cDNA 4930478P22	gene	4930478P22Rik	
1432101_a_at	0.82120	testis-specific serine kinase 4		Tssk4	
1432102_at	1.08931	regulating synaptic membrane exocytosis 2		Rims2	
1432103_a_at	0.95033	SH3-domain GRB2-like 3		Sh3gl3	
1432104_a_at	0.93089	allantoicase		Allc	
1432105_at	1.00329	sirtuin 5 (silent mating type information regulation 2 homolog) 5 (S. cerevisiae)		Sirt5	
1432106_at	0.89019	RIKEN cDNA 6530403M18	gene	6530403M18Rik	
1432107_at	0.98570	RIKEN cDNA 2310010M20	gene	2310010M20Rik	
1432108_at	1.52832	polycomb group ring finger 6		Pcgf6	

1432109_at	1.01787	RIKEN cDNA 4930578I07	gene	4930578I07Rik	
1432110_at	0.99969	RIKEN cDNA 4930402F06	gene	4930402F06Rik	
1432111_at	1.00163	RIKEN cDNA 4930542D17	gene	4930542D17Rik	
1432112_at	1.05852	RIKEN cDNA 4930589L23	gene	4930589L23Rik	
1432113_at	0.94568	coiled-coil domain containing 18		Ccdc18	
1432114_at	0.91298	RIKEN cDNA 1110035E04	gene	1110035E04Rik	
1432115_a_at	0.94319	"phosphatidylinositol glycan anchor biosynthesis, class N"			Pign
1432116_at	1.16493	RIKEN cDNA 1700031F05	gene	1700031F05Rik	
1432117_at	0.96261	RIKEN cDNA 4933422A05	gene	4933422A05Rik	
1432118_at	1.01523	RIKEN cDNA 2410012E07	gene	2410012E07Rik	
1432119_at	1.00635	RIKEN cDNA 4933405I11	gene	4933405I11Rik	
1432120_at	0.85653	RIKEN cDNA 4921513D11	gene	4921513D11Rik	
1432121_a_at	0.84838	leucine rich repeat containing 44		Lrrc44	
1432122_at	0.82290	leucine rich repeat containing 44		Lrrc44	
1432123_at	1.01239	RIKEN cDNA 4930455H04	gene	4930455H04Rik	
1432124_at	1.03797	RIKEN cDNA 4921511I17	gene	4921511I17Rik	
1432125_at	1.08331	"adaptor-related protein complex 3, mu 2 subunit"			Ap3m2
1432126_at	1.00886	RIKEN cDNA 1700072H12	gene	1700072H12Rik	
1432127_at	0.97564	RIKEN cDNA 4933401L05	gene	4933401L05Rik	
1432128_at	1.01605	RIKEN cDNA 4930455B06	gene	4930455B06Rik	
1432129_a_at	1.00794	paired related homeobox 1		Prrx1	
1432130_a_at	0.79972	tetratricopeptide repeat domain 14		Ttc14	
1432131_at	0.98767	RIKEN cDNA 4930404N11	gene	4930404N11Rik	
1432132_at	1.18076	RIKEN cDNA 4933401D09	gene	4933401D09Rik	
1432133_at	1.00300	RIKEN cDNA 2210022J03	gene	2210022J03Rik	
1432134_at	1.06560	speedy homolog A (Drosophila)		Spdya	
1432135_s_at	0.92811	speedy homolog A (Drosophila) /// similar to speedy homolog 1		Spdya ///	
LOC638442					
1432136_s_at	0.86250	"zinc finger, DHHC domain containing 4"		Zdhhc4	
1432137_at	1.07395	RIKEN cDNA 4933434P08	gene	4933434P08Rik	
1432138_at	1.01875	RIKEN cDNA 1700080N15	gene	1700080N15Rik	
1432139_at	1.00111	serine/arginine-rich protein specific kinase 3		Srpk3	
1432140_at	1.02812	gasdermin 2		Gsdm2	
1432141_x_at	0.92315	gasdermin 2		Gsdm2	
1432142_at	0.99426	"solute carrier family 35, member F4"		Slc35f4	
1432143_a_at	0.90287	high mobility group box transcription factor 1		Hbp1	
1432144_a_at	1.17177	ring finger and CHY zinc finger domain containing 1			Rchy1
1432145_at	0.83248	RIKEN cDNA 2310026I22	gene	2310026I22Rik	
1432146_at	0.91783	"cation channel, sperm associated 3"		Catsper3	
1432147_at	1.27706	RIKEN cDNA 4930447F24	gene	4930447F24Rik	
1432148_at	1.05370	RIKEN cDNA 4930533K18	gene	4930533K18Rik	
1432149_at	0.98055	DPH1 homolog (S. cerevisiae)		Dph1	
1432150_at	1.05123	transmembrane protein 59		Tmem59	

1432151_at	1.31858	RIKEN cDNA 2810409H07	gene	2810409H07Rik
1432152_at	0.94686	RIKEN cDNA 2610109H07	gene	2610109H07Rik
1432153_at	0.99465	RIKEN cDNA 4933426B08	gene	4933426B08Rik
1432154_at	1.06824	RIKEN cDNA 4933407I08	gene	4933407I08Rik
1432155_at	0.94945	Wiskott-Aldrich syndrome-like (human)		Wasl
1432156_a_at	0.87312	ring finger protein 32		Rnf32
1432157_at	1.06656	RIKEN cDNA 4930515G13	gene	4930515G13Rik
1432158_a_at	0.90246	trafficking protein particle complex 2		Trappc2
1432159_a_at	0.96306	testis expressed gene 13		Tex13
1432160_at	1.09219	G protein-coupled receptor kinase-interactor 2		Git2
1432161_a_at	1.12755	protein prenyltransferase alpha subunit repeat containing 1		Ptar1
1432162_s_at	0.94780	protein prenyltransferase alpha subunit repeat containing 1		Ptar1
1432163_at	1.01086	RIKEN cDNA 4930567K12	gene	4930567K12Rik
1432164_a_at	1.13278	"glycine cleavage system protein H (aminomethyl carrier) /// similar to Glycine cleavage system H protein, mitochondrial precursor"		Gcsh /// LOC674321
1432165_at	0.91246	RIKEN cDNA 8430422H06	gene	8430422H06Rik
1432166_at	1.05021	RIKEN cDNA E130018N17	gene	E130018N17Rik
1432167_at	1.06088	---	---	
1432168_at	0.86670	RIKEN cDNA D730045A05	gene	D730045A05Rik
1432169_at	0.97178	RIKEN cDNA 4930523O13	gene	4930523O13Rik
1432170_at	0.94175	"Ts translation elongation factor, mitochondrial"		Tsfm
1432171_at	1.13938	RIKEN cDNA 4933409F18	gene	4933409F18Rik
1432172_at	0.93732	hypothetical protein LOC670300		LOC670300
1432173_at	0.90567	"serine (or cysteine) peptidase inhibitor, clade B (ovalbumin), member 12"		Serpinb12
1432174_a_at	0.99527	RIKEN cDNA 4933425K02	gene	4933425K02Rik
1432175_at	0.98908	RIKEN cDNA 5430434I15	gene	5430434I15Rik
1432176_a_at	0.93701	---	---	
1432177_a_at	1.06478	menage a trois 1		Mnat1
1432178_at	1.05093	RIKEN cDNA 4933431I19	gene	4933431I19Rik
1432179_x_at	1.01338	RIKEN cDNA 2810433K01	gene	2810433K01Rik
1432180_at	1.05567	RIKEN cDNA 4933403G17	gene	4933403G17Rik
1432181_s_at	1.18586	endothelial cell growth factor 1 (platelet-derived)		Ecgf1
1432182_at	1.04411	RIKEN cDNA 4922502B01	gene	4922502B01Rik
1432183_at	0.96832	transmembrane protein 116		Tmem116
1432184_a_at	1.15157	RIKEN cDNA 2610204M08	gene	2610204M08Rik
1432185_a_at	0.96137	speedy homolog B (Drosophila)		Spdyb
1432186_at	0.92567	RIKEN cDNA 1700028J19	gene	1700028J19Rik
1432187_at	1.01604	nucleoporin 43		Nup43
1432188_s_at	1.12440	nucleoporin 43		Nup43
1432189_a_at	0.80197	SRY-box containing gene 5		Sox5
1432190_at	1.02976	SRY-box containing gene 5		Sox5
1432191_at	1.03253	---	---	

1432192_at	1.06518	RIKEN cDNA 4930483K19	gene	4930483K19Rik
1432193_at	1.07233	RIKEN cDNA 4930412L05	gene	4930412L05Rik
1432194_at	0.93836	RIKEN cDNA 9530020I12	gene	9530020I12Rik
1432195_s_at	1.05144	cyclin L2	Ccnl2	
1432196_a_at	0.91598	Down syndrome cell adhesion molecule-like 1	Dscaml1	
1432197_at	1.01813	RIKEN cDNA 4933417D19	gene	4933417D19Rik
1432198_at	1.10414	---	---	
1432199_at	0.93290	RIKEN cDNA 4930568D16	gene	4930568D16Rik
1432200_at	1.19505	RIKEN cDNA C030044M21	gene	C030044M21Rik
1432201_a_at	0.94535	"zinc finger, SWIM domain containing 2"	Zswim2	
1432202_a_at	1.08508	WD repeat domain 51A	Wdr51a	
1432203_at	0.99287	RIKEN cDNA 4930563P21	gene	4930563P21Rik
1432204_at	0.80811	RIKEN cDNA 4930570B17	gene	4930570B17Rik
1432205_a_at	0.99799	RIKEN cDNA C130038G02	gene	C130038G02Rik
1432206_at	1.09374	RIKEN cDNA 9030404E10	gene	9030404E10Rik
1432207_a_at	0.95452	"target of EGR1, member 1 (nuclear)"	Toe1	
1432208_at	1.06419	RIKEN cDNA 4930459C07	gene	4930459C07Rik
1432209_at	1.10711	RIKEN cDNA 1810010K12	gene	1810010K12Rik
1432210_at	0.89970	RIKEN cDNA 4933401H06	gene	4933401H06Rik
1432211_a_at	0.97809	f-box only protein 9	Fbxo9	
1432212_at	1.01159	RIKEN cDNA 1810073H04	gene	1810073H04Rik
1432213_at	0.93093	RIKEN cDNA 9530047P18	gene	9530047P18Rik
1432214_at	1.08218	RIKEN cDNA 1700017D01	gene	1700017D01Rik
1432215_s_at	0.99489	RIKEN cDNA 4930417O22	gene	4930417O22Rik
1432216_s_at	1.12397	"membrane protein, palmitoylated 7 (MAGUK p55 subfamily member 7)"	Mpp7	
1432217_a_at	1.06721	WD repeat domain 16	Wdr16	
1432218_a_at	1.01258	nucleolar protein 9	Nol9	
1432219_at	1.06430	nucleolar protein 9	Nol9	
1432220_at	1.03133	RIKEN cDNA 4933430L12	gene	4933430L12Rik
1432221_at	0.99465	RIKEN cDNA 5330417H12	gene	5330417H12Rik
1432222_at	1.07249	RIKEN cDNA A930015G24	gene	A930015G24Rik
1432223_at	1.04098	RIKEN cDNA 4930552N02	gene	4930552N02Rik
1432224_at	0.99057	RIKEN cDNA 4831407H17	gene	4831407H17Rik
1432225_at	1.01420	---	---	
1432226_at	1.13968	RIKEN cDNA 4930548G14	gene	4930548G14Rik
1432227_at	0.90031	suppressor of variegation 3-9 homolog 1 (Drosophila)	Suv39h1	
1432228_at	1.06657	RIKEN cDNA 4933406B15	gene	4933406B15Rik
1432229_a_at	0.93545	"chromodomain protein, Y chromosome-like 2"	Cdyl2	
1432230_at	1.20494	huntingtin interacting protein 1	Hip1	
1432231_at	1.11926	RIKEN cDNA 4930526H09	gene	4930526H09Rik
1432232_at	0.96819	REST corepressor 3	Rcor3	
1432233_at	0.99931	RIKEN cDNA 1110020C17	gene	1110020C17Rik

1432234_at	1.07875	RIKEN cDNA 4933438B17 gene	4933438B17Rik
1432235_at	0.95379	epithelial stromal interaction 1 (breast)	Epsti1
1432236_a_at	0.91861	suppressor of variegation 3-9 homolog 1 (Drosophila)	Suv39h1
1432237_at	0.97687	RIKEN cDNA 5730522E02 gene	5730522E02Rik
1432238_at	1.05980	"spermatogenesis associated, serine-rich 1"	Spats1
1432239_at	1.07428	RIKEN cDNA 1700102J08 gene	1700102J08Rik
1432240_at	1.01661	RIKEN cDNA 4930533L02 gene	4930533L02Rik
1432241_at	0.94833	RIKEN cDNA 9030205G03 gene	9030205G03Rik
1432242_at	0.89409	RIKEN cDNA 4930447N08 gene	4930447N08Rik
1432243_a_at	0.99504	RIKEN cDNA 4933433G15 gene	4933433G15Rik
1432244_at	1.00787	RIKEN cDNA 4930507D10 gene	4930507D10Rik
1432245_s_at	1.17264	tetratricopeptide repeat domain 9C	Ttc9c
1432246_at	0.98490	RIKEN cDNA 4922502N22 gene	4922502N22Rik
1432247_at	0.92803	RIKEN cDNA 4633402D09 gene	4633402D09Rik
1432248_at	0.93814	RIKEN cDNA 5430402P08 gene	5430402P08Rik
1432249_a_at	1.04569	"excision repair cross-complementing rodent repair deficiency, complementation group 8"	Ercc8
1432250_at	0.94045	"UTP14, U3 small nucleolar ribonucleoprotein, homolog A (yeast)"	Utp14a
1432251_at	0.95665	RIKEN cDNA 4930403O15 gene	4930403O15Rik
1432252_a_at	1.14262	RIKEN cDNA 4933406K04 gene	4933406K04Rik
1432253_at	1.00564	RIKEN cDNA 4921523P09 gene	4921523P09Rik
1432254_at	1.00846	RIKEN cDNA C030013D06 gene	C030013D06Rik
1432255_at	0.99331	microtubule associated serine/threonine kinase 1	Mast1
1432256_at	0.95470	RIKEN cDNA 1700111N16 gene	1700111N16Rik
1432257_at	0.89703	gasdermin domain containing 2	Gsdmdc2
1432258_at	0.96931	RIKEN cDNA 9430014N10 gene	9430014N10Rik
1432259_s_at	0.83410	RIKEN cDNA 4931415C17 gene /// RIKEN cDNA 2810487C13 gene 4931415C17Rik /// 2810487C13Rik	
1432260_at	1.04807	G protein-coupled receptor 39	Gpr39
1432261_at	0.96748	RIKEN cDNA 2310039D24 gene	2310039D24Rik
1432262_at	0.89142	RIKEN cDNA 4930504E06 gene	4930504E06Rik
1432263_a_at	0.89056	cytochrome c oxidase subunit VIIa polypeptide 2-like	Cox7a2l
1432264_x_at	0.91146	cytochrome c oxidase subunit VIIa polypeptide 2-like	Cox7a2l
1432265_at	0.89488	RIKEN cDNA 4930445B16 gene	4930445B16Rik
1432266_at	1.08780	cDNA sequence BC026657	BC026657
1432267_at	0.85704	RIKEN cDNA 6720483E21 gene	6720483E21Rik
1432268_at	1.14628	RIKEN cDNA 2310068J16 gene	2310068J16Rik
1432269_a_at	0.84147	SH3-domain kinase binding protein 1	Sh3kbp1
1432270_a_at	0.91900	chromatin modifying protein 5	Chmp5
1432271_a_at	1.06230	"DCN1, defective in cullin neddylation 1, domain containing 5 (S. cerevisiae)"	Dcun1d5
1432272_a_at	0.83798	"excision repair cross-complementing rodent repair deficiency, complementation group 4"	Ercc4

1432273_a_at	1.02499	"Duffy blood group, chemokine receptor"	Darc
1432274_at	1.01865	RIKEN cDNA 4930543N07 gene	4930543N07Rik
1432275_at	1.03713	inhibitor of kappaB kinase beta	Ikbkb
1432276_at	0.92531	RIKEN cDNA 4930519F16 gene	4930519F16Rik
1432277_at	1.02432	RIKEN cDNA 4930565D16 gene	4930565D16Rik
1432278_at	1.06127	RIKEN cDNA 4930471M09 gene	4930471M09Rik
1432279_at	0.95623	RIKEN cDNA 4930452L02 gene	4930452L02Rik
1432280_at	1.05747	RIKEN cDNA 2310007L24 gene	2310007L24Rik
1432281_a_at	0.80530	integrin beta 6	Itgb6
1432282_a_at	0.86391	RIKEN cDNA 2010305C02 gene	2010305C02Rik
1432283_at	0.97641	RIKEN cDNA 4930417G10 gene	4930417G10Rik
1432284_at	1.03512	mirror-image polydactyly gene 1 homolog (human)	Mipol1
1432285_at	1.07285	RIKEN cDNA 1700128F08 gene	1700128F08Rik
1432286_at	1.03316	"polymerase (DNA-directed), epsilon 4 (p12 subunit)"	Pole4
1432287_a_at	1.04004	"syntrophin, gamma 1"	Sntg1
1432288_at	0.91557	RIKEN cDNA 5830468F06 gene	5830468F06Rik
1432289_a_at	0.93332	junctional sarcoplasmic reticulum protein 1	Jsrp1
1432290_at	0.92051	RIKEN cDNA 1700040G22 gene	1700040G22Rik
1432291_at	0.92235	RIKEN cDNA 0610033M10 gene	0610033M10Rik
1432292_at	1.16096	RIKEN cDNA 4930504C09 gene	4930504C09Rik
1432293_at	1.10927	RIKEN cDNA 4930579G18 gene	4930579G18Rik
1432294_at	1.02708	RIKEN cDNA 9330177L23 gene	9330177L23Rik
1432295_a_at	0.91031	RIKEN cDNA 2310035K24 gene	2310035K24Rik
1432296_a_at	1.08502	integrin alpha V	Itgav
1432297_at	1.01172	"zinc finger, imprinted 2"	Zim2
1432298_at	0.96252	RIKEN cDNA 4921508M14 gene	4921508M14Rik
1432299_at	0.92134	RIKEN cDNA 4921513I08 gene	4921513I08Rik
1432300_at	0.94350	RIKEN cDNA 1190003K10 gene	1190003K10Rik
1432301_a_at	1.03311	RIKEN cDNA 1700074P13 gene	1700074P13Rik
1432302_s_at	0.94836	RIKEN cDNA 2410024N13 gene	2410024N13Rik
1432303_at	0.84479	RIKEN cDNA 9330198I05 gene	9330198I05Rik
1432304_a_at	1.03497	RIKEN cDNA 9030624J02 gene	9030624J02Rik
1432305_at	1.09072	RIKEN cDNA 4930542C16 gene	4930542C16Rik
1432306_at	0.88867	RIKEN cDNA A930023M06 gene	A930023M06Rik
1432307_at	0.90641	RIKEN cDNA 1700086D15 gene	1700086D15Rik
1432308_at	0.92182	RIKEN cDNA 1700058C13 gene	1700058C13Rik
1432309_at	0.97368	RIKEN cDNA 9130410C08 gene	9130410C08Rik
1432310_at	0.89051	RIKEN cDNA 6530401F13 gene	6530401F13Rik
1432311_at	1.27646	transmembrane protein 168	Tmem168
1432312_a_at	0.95415	RIKEN cDNA 4931440B09 gene	4931440B09Rik
1432313_at	0.88383	RIKEN cDNA 4933433F19 gene	4933433F19Rik
1432314_at	1.04559	RIKEN cDNA 4930554C24 gene	4930554C24Rik
1432315_at	0.91135	RIKEN cDNA 4930483C13 gene	4930483C13Rik

1432316_at	1.05878	RIKEN cDNA 4933439N14	gene	4933439N14Rik
1432317_at	0.90901	RIKEN cDNA 9430078K24	gene	9430078K24Rik
1432318_at	1.05141	UDP-glucose ceramide glucosyltransferase-like 1	Ugcgl1	
1432319_at	1.19759	RIKEN cDNA 1700085C21	gene	1700085C21Rik
1432320_at	0.91750	RIKEN cDNA 9430065F17	gene	9430065F17Rik
1432321_at	0.97931	cellular nucleic acid binding protein 2	Cnbp2	
1432322_at	0.90841	RIKEN cDNA 2310016E02	gene	2310016E02Rik
1432323_at	0.96471	RIKEN cDNA 8430436N08	gene	8430436N08Rik
1432324_at	1.02112	RIKEN cDNA 4930432J09	gene	4930432J09Rik
1432325_at	0.99732	RIKEN cDNA 4930502M04	gene	4930502M04Rik
1432326_at	1.03127	RIKEN cDNA 1700042B14	gene	1700042B14Rik
1432327_at	0.96376	RIKEN cDNA 9530067L11	gene	9530067L11Rik
1432328_at	1.16904	RIKEN cDNA 4930516K23	gene	4930516K23Rik
1432329_a_at	1.01633	megakaryocyte-associated tyrosine kinase	Matk	
1432330_at	0.90551	complement factor H-like	LOC381294	
1432331_a_at	0.70566	paired related homeobox 2	Prrx2	
1432332_a_at	0.98337	nudix (nucleoside diphosphate linked moiety X)-type motif 19	Nudt19	
1432333_a_at	0.99728	RIKEN cDNA 1700027D21	gene	1700027D21Rik
1432334_at	0.88065	RIKEN cDNA 4930488P18	gene	4930488P18Rik
1432335_at	1.01287	RIKEN cDNA 4930551O13	gene	4930551O13Rik
1432336_at	0.98579	RIKEN cDNA 5033404E19	gene	5033404E19Rik
1432337_at	0.95214	"Fanconi anemia, complementation group L"	Fancl	
1432338_at	0.93946	RIKEN cDNA 4833419O12	gene	4833419O12Rik
1432339_at	1.00731	RIKEN cDNA 4933432I03	gene	4933432I03Rik
1432340_at	0.99914	RIKEN cDNA 1700121N20	gene	1700121N20Rik
1432341_at	0.96934	RIKEN cDNA 1700022A22	gene	1700022A22Rik
1432342_at	1.02490	nicotinamide nucleotide adenyltransferase 3	Nmnat3	
1432343_at	0.97514	RIKEN cDNA 4930419G24	gene	4930419G24Rik
1432344_a_at	0.75940	amyloid beta (A4) precursor-like protein 2	Aplp2	
1432345_at	0.89196	RIKEN cDNA 1700048F04	gene	1700048F04Rik
1432346_a_at	0.91564	cadherin 23 (otocadherin)	Cdh23	
1432347_at	1.01129	RIKEN cDNA 4931428L18	gene	4931428L18Rik
1432348_at	1.02730	RIKEN cDNA 4930524O07	gene	4930524O07Rik
1432349_a_at	1.02145	syncoilin	Sync	
1432350_at	0.93953	syncoilin	Sync	
1432351_at	0.99183	RIKEN cDNA 9030625G05	gene	9030625G05Rik
1432352_at	1.59311	RIKEN cDNA 5730405I09	gene	5730405I09Rik
1432353_at	0.86897	RIKEN cDNA 4933421B21	gene	4933421B21Rik
1432354_at	1.20448	RIKEN cDNA 1700003H04	gene	1700003H04Rik
1432355_at	0.95170	RIKEN cDNA 2210039B01	gene	2210039B01Rik
1432356_at	1.32244	RIKEN cDNA 1700056N10	gene	1700056N10Rik
1432357_at	0.86304	RIKEN cDNA 4930566D17	gene	4930566D17Rik
1432358_at	1.01361	RIKEN cDNA 1110008I14	gene	1110008I14Rik

1432359_at 1.14119 RIKEN cDNA 4930435F05 gene 4930435F05Rik
1432360_a_at 0.82032 transmembrane and tetratricopeptide repeat containing 4 Tmtc4
1432361_a_at 0.93347 centromere protein P Cenpp
1432362_at 1.32645 centromere protein P Cenpp
1432363_at 1.04202 RIKEN cDNA 2410018E23 gene 2410018E23Rik
1432364_at 1.11943 RIKEN cDNA 4930556N13 gene 4930556N13Rik
1432365_a_at 0.95065 RIKEN cDNA 4930556L07 gene 4930556L07Rik
1432366_at 0.86235 RIKEN cDNA 4930447K03 gene 4930447K03Rik
1432367_a_at 1.06938 ubiquitin fusion degradation 1 like Ufd1l
1432368_at 0.94107 RIKEN cDNA 4930594M17 gene 4930594M17Rik
1432369_at 1.02580 RIKEN cDNA 3010027C24 gene 3010027C24Rik
1432370_at 0.98933 RIKEN cDNA 4933411E02 gene 4933411E02Rik
1432371_a_at 1.07002 RIKEN cDNA 1700109K24 gene 1700109K24Rik
1432372_a_at 1.06909 sepiapterin reductase Spr
1432373_at 1.05329 RIKEN cDNA 4930533B01 gene 4930533B01Rik
1432374_a_at 1.09914 "DNA segment, Chr 5, Wayne State University 178, expressed" D5Wsu178e
1432375_a_at 1.09196 spermatogenesis associated 3 Spata3
1432376_at 0.93314 RIKEN cDNA 3830403N18 gene 3830403N18Rik
1432377_x_at 1.03503 shroom family member 3 Shroom3
1432378_at 1.00605 RIKEN cDNA C030004G16 gene C030004G16Rik
1432379_at 1.00116 RIKEN cDNA 4930590A17 gene 4930590A17Rik
1432380_s_at 0.86372 RIKEN cDNA 4933426G20 gene /// similar to gasdermin 1
4933426G20Rik /// LOC665769
1432381_a_at 0.97182 fibrous sheath-interacting protein 1 /// similar to fibrous sheath-interacting
protein 1 Fsip1 /// LOC640450
1432382_at 0.99898 RIKEN cDNA 4930434B07 gene 4930434B07Rik
1432383_a_at 1.13777 armadillo repeat containing 9 Armc9
1432384_a_at 1.04292 methyltransferase like 6 Mettl6
1432385_a_at 0.90502 ATP/GTP binding protein 1 Agtpbp1
1432386_a_at 0.90050 PHD finger protein 7 Phf7
1432387_at 0.95215 plasma glutamate carboxypeptidase Pgcp
1432388_at 1.03290 RIKEN cDNA 5430440P10 gene 5430440P10Rik
1432389_at 1.00033 RIKEN cDNA 4933438A12 gene 4933438A12Rik
1432390_at 1.06740 RIKEN cDNA 4933400F21 gene 4933400F21Rik
1432391_at 1.09755 coiled-coil domain containing 21 Ccdc21
1432392_at 1.02012 kinesin family member 2C Kif2c
1432393_a_at 1.16466 RIKEN cDNA 5730409G07 gene 5730409G07Rik
1432394_a_at 1.01770 apoptosis antagonizing transcription factor Aatf
1432395_at 0.89277 RIKEN cDNA 1700039E15 gene 1700039E15Rik
1432396_at 1.01262 RIKEN cDNA 3110018I06 gene 3110018I06Rik
1432397_at 0.94885 testis-specific serine kinase 3 Tssk3
1432398_at 0.91150 RIKEN cDNA 1700084J12 gene 1700084J12Rik
1432399_a_at 0.95284 Eph receptor A1 EphA1

1432400_at	0.91477 Eph receptor A1 EphA1	
1432401_a_at	0.98635 RIKEN cDNA 4930511M11 gene	4930511M11Rik
1432402_at	0.94298 RIKEN cDNA 4930402F11 gene	4930402F11Rik
1432403_at	1.00814 RIKEN cDNA 4933402C06 gene	4933402C06Rik
1432404_at	0.92899 RIKEN cDNA 4930505A04 gene	4930505A04Rik
1432405_a_at	1.04038 "phospholipase C, zeta 1" Plcz1	
1432406_at	0.97421 carboxylesterase 7 Ces7	
1432407_at	1.13391 RIKEN cDNA 4933426D04 gene	4933426D04Rik
1432408_a_at	1.10461 "DNA segment, KIST 2" D0Kist2	
1432409_at	0.94237 "protein kinase, cAMP dependent regulatory, type II alpha"	Prkar2a
1432410_a_at	0.83365 bone morphogenetic protein 7 Bmp7	
1432411_a_at	0.98716 F-box and WD-40 domain protein 2 Fbxw2	
1432412_at	0.98976 RIKEN cDNA 9530004P13 gene	9530004P13Rik
1432413_at	1.05078 RIKEN cDNA 1810044A24 gene	1810044A24Rik
1432414_at	0.92675 --- ---	
1432415_at	1.01988 "RAB3C, member RAS oncogene family" Rab3c	
1432416_a_at	1.12403 nucleophosmin 1 Npm1	
1432417_a_at	0.95370 tetraspanin 2 Tspan2	
1432418_a_at	0.96349 "creatine kinase, mitochondrial 1, ubiquitous" Ckmt1	
1432419_a_at	1.02340 RIKEN cDNA 2700078K21 gene	2700078K21Rik
1432420_a_at	1.03769 RIKEN cDNA 2310002L09 gene	2310002L09Rik
1432421_at	1.07933 postmeiotic segregation increased 2 (S. cerevisiae) Pms2	
1432422_at	1.00261 RIKEN cDNA 1700063K16 gene	1700063K16Rik
1432423_a_at	0.91513 RIKEN cDNA C530008M17 gene	C530008M17Rik
1432424_at	1.04798 RIKEN cDNA 4930505H01 gene	4930505H01Rik
1432425_at	1.00628 RIKEN cDNA 4930418G15 gene	4930418G15Rik
1432426_a_at	1.06982 ubiquitin-conjugating enzyme E2F (putative) Ube2f	
1432427_at	0.99146 NADH dehydrogenase (ubiquinone) 1 beta subcomplex 4	Ndufb4
1432428_at	0.97388 ankyrin repeat domain 55 Ankrd55	
1432429_at	0.98563 RIKEN cDNA 8430437N05 gene	8430437N05Rik
1432430_a_at	0.86054 RIKEN cDNA 1700081L11 gene	1700081L11Rik
1432431_s_at	0.80315 RIKEN cDNA 2900006F19 gene	2900006F19Rik
1432432_a_at	1.16029 "RAB3C, member RAS oncogene family" Rab3c	
1432433_at	1.03858 RIKEN cDNA 4930524N10 gene	4930524N10Rik
1432434_at	0.87358 RIKEN cDNA 4930522N08 gene	4930522N08Rik
1432435_s_at	1.05944 RIKEN cDNA C030004A17 gene	C030004A17Rik
1432436_a_at	0.91452 adenylate kinase 3 Ak3	
1432437_at	0.87520 RIKEN cDNA 4933413N12 gene	4933413N12Rik
1432438_at	0.87209 RIKEN cDNA 4930597L12 gene	4930597L12Rik
1432439_at	1.05056 zinc finger protein 715 Zfp715	
1432440_at	0.96910 RIKEN cDNA 1700111E14 gene	1700111E14Rik
1432441_at	0.96218 RIKEN cDNA 4933413J09 gene	4933413J09Rik
1432442_at	0.99381 RIKEN cDNA 4930481F22 gene	4930481F22Rik

1432443_at	1.03915	RIKEN cDNA 1700021P22	gene	1700021P22Rik	
1432444_a_at	1.09406	RIKEN cDNA 1810011O16	gene	1810011O16Rik	
1432445_at	0.91627	RIKEN cDNA 2310016G11	gene	2310016G11Rik	
1432446_at	1.12642	RIKEN cDNA 1700016K13	gene	1700016K13Rik	
1432447_a_at	0.98071	RIKEN cDNA 2310005N01	gene	2310005N01Rik	
1432448_at	0.87189	RIKEN cDNA 2600006K01	gene	2600006K01Rik	
1432449_at	0.95489	RIKEN cDNA 4930564B18	gene	4930564B18Rik	
1432450_at	0.91598	RIKEN cDNA 8030423J24	gene	8030423J24Rik	
1432451_at	1.00325	RIKEN cDNA 4930445E18	gene	4930445E18Rik	
1432452_at	0.95363	RIKEN cDNA 1700054O19	gene	1700054O19Rik	
1432453_a_at	1.00113	"membrane-spanning 4-domains, subfamily A, member 10"			Ms4a10
1432454_at	0.99602	RIKEN cDNA 2410137M14	gene	2410137M14Rik	
1432455_a_at	0.96993	prolactin-like protein C 4			Prlpc4
1432456_at	0.95850	RIKEN cDNA 4930533B18	gene	4930533B18Rik	
1432457_at	1.07196	RIKEN cDNA 4930448F12	gene	4930448F12Rik	
1432458_at	1.00194	RIKEN cDNA 1700011F14	gene	1700011F14Rik	
1432459_a_at	1.00984	zinc finger and BTB domain containing 32			Zbtb32
1432460_at	0.99661	RIKEN cDNA 1700125F08	gene	1700125F08Rik	
1432461_at	0.93422	eyes absent 4 homolog (Drosophila)			Eya4
1432462_a_at	1.00257	"cofactor required for Sp1 transcriptional activation, subunit 8"			Crsp8
1432463_at	0.95400	RIKEN cDNA 5730510P18	gene	5730510P18Rik	
1432464_a_at	0.77908	RIKEN cDNA 2310057J16	gene	2310057J16Rik	
1432465_at	1.16802	RIKEN cDNA 4933414I06	gene	4933414I06Rik	
1432466_a_at	1.02139	apolipoprotein E			Apoe
1432467_at	0.92897	leucine-rich repeats and calponin homology (CH) domain containing 3			Lrch3
1432468_at	1.12778	lipocalin 9			Lcn9
1432469_at	0.90449	RIKEN cDNA 1700094E07	gene	1700094E07Rik	
1432470_at	0.91256	RIKEN cDNA 1700016A09	gene	1700016A09Rik	
1432471_at	0.85506	RIKEN cDNA 5730419I09	gene	5730419I09Rik	
1432472_a_at	1.00784	methylcrotonoyl-Coenzyme A carboxylase 2 (beta)			Mccc2
1432473_a_at	1.11332	"ATP-binding cassette, sub-family A (ABC1), member 14"			Abca14
1432474_a_at	0.99357	keratinocyte associated protein 3			Krtcap3
1432475_at	0.89924	RIKEN cDNA 1700014B07	gene	1700014B07Rik	
1432476_at	1.13743	RIKEN cDNA 1700040F17	gene	1700040F17Rik	
1432477_at	0.83296	RIKEN cDNA 4930412D23	gene	4930412D23Rik	
1432478_a_at	1.01897	IBR domain containing 3			Ibrdc3
1432479_at	1.00108	"Mus musculus, clone IMAGE:6772209, mRNA"			---
1432480_a_at	1.00316	cyclin N-terminal domain containing 1			Cntd1
1432481_a_at	0.89514	lysozyme-like 6			Lyzl6
1432482_at	0.91233	lysozyme-like 6			Lyzl6
1432483_at	0.93065	RIKEN cDNA 1700124K17	gene	1700124K17Rik	
1432484_at	0.88645	RIKEN cDNA 2310061C15	gene	2310061C15Rik	
1432485_at	0.92185	RIKEN cDNA 3110057O12	gene	3110057O12Rik	

1432486_a_at 0.89696 testicular serine protease 2 *Tesp2*
1432487_at 0.86702 RIKEN cDNA 4930405H06 gene 4930405H06Rik
1432488_a_at 1.11986 "splicing factor 3a, subunit 3 /// similar to splicing factor 3a, subunit 3 (predicted)" *Sf3a3* /// LOC626830
1432489_a_at 1.35178 RIKEN cDNA 2410187C16 gene 2410187C16Rik
1432490_a_at 0.94626 phosphodiesterase 10A *Pde10a*
1432491_at 0.93672 protocadherin 8 *Pcdh8*
1432492_a_at 0.95257 "3-hydroxyanthranilate 3,4-dioxygenase" *Haa0*
1432493_at 1.10401 RIKEN cDNA 1700020M10 gene 1700020M10Rik
1432494_a_at 0.99457 RIKEN cDNA 1700019E19 gene 1700019E19Rik
1432495_at 1.16307 SRY-box containing gene 7 *Sox7*
1432496_at 0.94691 RIKEN cDNA 1700122O11 gene 1700122O11Rik
1432497_at 0.95186 coiled-coil domain containing 93 *Ccdc93*
1432498_at 1.01632 RIKEN cDNA 1700020C07 gene 1700020C07Rik
1432499_a_at 1.05041 "ubiquitination factor E4B, UFD2 homolog (*S. cerevisiae*)" *Ube4b*
1432500_at 0.98316 RIKEN cDNA 2410006F04 gene 2410006F04Rik
1432501_at 0.99394 RIKEN cDNA 1700092E16 gene 1700092E16Rik
1432502_at 1.21026 RIKEN cDNA 1700028N14 gene 1700028N14Rik
1432503_a_at 1.03131 phosducin-like 2 *Pdcl2*
1432504_at 0.94363 RIKEN cDNA 1700106N22 gene 1700106N22Rik
1432505_at 0.92315 RIKEN cDNA 1700009J07 gene 1700009J07Rik
1432506_at 0.94230 transmembrane protein 86B *Tmem86b*
1432507_at 0.80575 RIKEN cDNA B430319H21 gene B430319H21Rik
1432508_at 0.99252 RIKEN cDNA 1700113I22 gene 1700113I22Rik
1432509_at 0.94271 RIKEN cDNA 5033430I15 gene 5033430I15Rik
1432510_at 1.04844 RIKEN cDNA 4930512M02 gene 4930512M02Rik
1432511_s_at 1.02218 centrosomal protein 27 *Cep27*
1432512_at 1.03318 RIKEN cDNA 4931433A01 gene 4931433A01Rik
1432513_a_at 0.97300 RIKEN cDNA 1700001C02 gene 1700001C02Rik
1432514_at 1.08413 RIKEN cDNA 1700066J24 gene 1700066J24Rik
1432515_at 1.05541 RIKEN cDNA 2410124H12 gene 2410124H12Rik
1432516_at 0.96471 RIKEN cDNA 4930447F04 gene 4930447F04Rik
1432517_a_at 0.93493 nicotinamide N-methyltransferase *Nnmt*
1432518_at 1.02956 RIKEN cDNA 4930430J20 gene 4930430J20Rik
1432519_at 0.97329 RIKEN cDNA 1810059H22 gene 1810059H22Rik
1432520_at 0.85288 immunoglobulin heavy chain (J558 family) *Igh-VJ558*
1432521_at 1.01018 matrix-remodelling associated 8 *Mxra8*
1432522_s_at 1.03278 coiled-coil domain containing 46 *Ccdc46*
1432523_at 0.85340 homeodomain interacting protein kinase 2 *Hipk2*
1432524_at 1.01635 RIKEN cDNA 4930550G17 gene 4930550G17Rik
1432525_at 1.09351 RIKEN cDNA 1700001C02 gene 1700001C02Rik
1432526_a_at 1.00673 "SNF8, ESCRT-II complex subunit, homolog (*S. cerevisiae*)" *Snf8*
1432527_at 0.90197 RIKEN cDNA 1700108M19 gene 1700108M19Rik

1432528_at	1.01522	RIKEN cDNA 1700021P04 gene	1700021P04Rik	
1432529_at	0.87161	"bol, boule-like (Drosophila)" Boll		
1432530_s_at	0.92439	"bol, boule-like (Drosophila)" Boll		
1432531_at	1.11495	RIKEN cDNA 9230104L09 gene	9230104L09Rik	
1432532_at	0.98230	RIKEN cDNA 0610025J13 gene	0610025J13Rik	
1432533_a_at	0.84567	"solute carrier family 35 (UDP-galactose transporter), member 2"		Slc35a2
1432534_at	1.05547	RIKEN cDNA 3300002I08 gene	3300002I08Rik	
1432535_at	1.10346	RIKEN cDNA 5530400C23 gene	5530400C23Rik	
1432536_at	0.90262	RIKEN cDNA 1700055M20 gene	1700055M20Rik	
1432537_at	0.95145	RIKEN cDNA 4930465A12 gene	4930465A12Rik	
1432538_a_at	0.96390	replication factor C (activator 1) 3	Rfc3	
1432539_a_at	0.99678	nucleoporin 54	Nup54	
1432540_at	0.90030	RIKEN cDNA 5430433J05 gene	5430433J05Rik	
1432541_at	1.03152	RIKEN cDNA 4930408O17 gene	4930408O17Rik	
1432542_at	0.97925	RIKEN cDNA 2810474C18 gene	2810474C18Rik	
1432543_a_at	1.00251	Kruppel-like factor 13	Klf13	
1432544_at	0.99423	RIKEN cDNA 4930449E01 gene	4930449E01Rik	
1432545_at	0.93675	DC-STAMP domain containing 1	Dcst1	
1432546_at	1.01574	RIKEN cDNA 4930529I22 gene	4930529I22Rik	
1432547_at	0.98127	RIKEN cDNA C030005H24 gene	C030005H24Rik	
1432548_at	0.98474	"poly (ADP-ribose) polymerase family, member 14"		Parp14
1432549_s_at	0.89421	RIKEN cDNA 1700112M01 gene	1700112M01Rik	
1432550_at	0.99614	RIKEN cDNA 4930572D21 gene	4930572D21Rik	
1432551_at	1.03212	hypothetical LOC73317	LOC73317	
1432552_at	1.00122	transmembrane protein 89	Tmem89	
1432553_at	0.96442	RIKEN cDNA 1700069L16 gene	1700069L16Rik	
1432554_at	1.11987	RIKEN cDNA 2310040G07 gene	2310040G07Rik	
1432555_at	0.88846	whirlin	Whrn	
1432556_a_at	0.85165	RIKEN cDNA 3100002J23 gene	3100002J23Rik	
1432557_at	0.87995	RIKEN cDNA 1700045I11 gene	/// similar to RIKEN cDNA 1700045I11 gene	1700045I11Rik
1432558_a_at	0.90378	"myelin and lymphocyte protein, T-cell differentiation protein"	Mal	
1432559_at	0.89798	RIKEN cDNA 2210409D07 gene	2210409D07Rik	
1432560_at	1.04464	RIKEN cDNA 1700127D06 gene	1700127D06Rik	
1432561_at	0.99148	RIKEN cDNA 4930569F06 gene	4930569F06Rik	
1432562_at	0.91221	RIKEN cDNA 1110006G14 gene	1110006G14Rik	
1432563_at	1.09915	RIKEN cDNA 1700116B05 gene	1700116B05Rik	
1432564_at	1.03537	RIKEN cDNA 1700066C05 gene	1700066C05Rik	
1432565_at	1.10003	---	---	
1432566_at	0.99146	RIKEN cDNA 1700129I15 gene	1700129I15Rik	
1432567_at	1.15432	RIKEN cDNA A430105D02 gene	A430105D02Rik	
1432568_at	0.97164	RIKEN cDNA 1700121K02 gene	1700121K02Rik	
1432569_at	0.82706	RIKEN cDNA 1700121K02 gene	1700121K02Rik	

1432570_at	0.94908	RIKEN cDNA 6030458E02	gene	6030458E02Rik
1432571_at	1.03023	---	---	
1432572_at	1.16884	RIKEN cDNA 4931406H21	gene	4931406H21Rik
1432573_at	1.09251	RIKEN cDNA 4930506C02	gene	4930506C02Rik
1432574_at	0.95603	RIKEN cDNA 4930548F15	gene	4930548F15Rik
1432575_at	1.05936	RIKEN cDNA 4930548F15	gene	4930548F15Rik
1432576_at	0.95927	RIKEN cDNA 1700084C06	gene	1700084C06Rik
1432577_at	0.91557	---	---	
1432578_at	1.07930	RIKEN cDNA 4933440H19	gene	4933440H19Rik
1432579_at	1.14971	radial spokehead-like 2A	Rshl2a	
1432580_at	1.03185	radial spokehead-like 2A	Rshl2a	
1432581_at	1.07264	RIKEN cDNA 3110054G05	gene	3110054G05Rik
1432582_at	1.08807	RIKEN cDNA 3110054G05	gene	3110054G05Rik
1432583_at	0.96910	RIKEN cDNA 2900042E19	gene	2900042E19Rik
1432584_at	1.03466	RIKEN cDNA 1110004M10	gene	1110004M10Rik
1432585_at	0.92757	RIKEN cDNA 4931428A05	gene	4931428A05Rik
1432586_at	1.00060	RIKEN cDNA 5730596P11	gene	5730596P11Rik
1432587_at	0.99994	RIKEN cDNA 5730596P11	gene	5730596P11Rik
1432588_at	1.10463	RIKEN cDNA 5830468K08	gene	5830468K08Rik
1432589_at	0.80327	---	---	
1432590_at	1.04127	RIKEN cDNA 4930573O21	gene	4930573O21Rik
1432591_at	1.01554	pregnancy-associated plasma protein A	Pappa	
1432592_at	1.00044	pregnancy-associated plasma protein A	Pappa	
1432593_at	0.92538	RIKEN cDNA 4933428P19	gene	4933428P19Rik
1432594_at	0.88449	RIKEN cDNA 4933413I22	gene	4933413I22Rik
1432595_at	0.93853	---	---	
1432596_at	1.03336	RIKEN cDNA 4930599N24	gene	4930599N24Rik
1432597_at	0.86197	RIKEN cDNA 4930599N24	gene	4930599N24Rik
1432598_at	0.85573	germ cell-less homolog 1 (Drosophila)	Gmcl1	
1432599_at	1.21681	RIKEN cDNA 2610301H18	gene	2610301H18Rik
1432600_at	1.04976	RIKEN cDNA 2310061A09	gene	2310061A09Rik
1432601_at	1.05351	myeloid/lymphoid or mixed-lineage leukemia 5	Mll5	
1432602_at	0.92470	---	---	
1432603_at	0.95603	nuclear receptor interacting protein 1	Nrip1	
1432604_at	1.05250	retinoblastoma-like 1 (p107)	Rbl1	
1432605_at	0.91705	---	---	
1432606_at	0.97001	RIKEN cDNA 2610012C04	gene	2610012C04Rik
1432607_at	1.05740	RIKEN cDNA 2610012C04	gene	2610012C04Rik
1432608_at	0.87637	nicotinamide nucleotide transhydrogenase	Nnt	
1432609_at	1.11333	---	---	
1432610_at	0.96763	RIKEN cDNA 2900024J01	gene	2900024J01Rik
1432611_at	1.02911	RIKEN cDNA 4932430A15	gene	4932430A15Rik
1432612_at	1.05969	RIKEN cDNA 9030405F24	gene	9030405F24Rik

1432613_at	0.90659	RIKEN cDNA E130201N16 gene	E130201N16Rik
1432614_at	1.01539	WD repeat domain 37 Wdr37	
1432615_at	1.04332	WD repeat domain 37 Wdr37	
1432616_at	0.93153	RIKEN cDNA 4833422B07 gene	4833422B07Rik
1432617_at	0.87928	RIKEN cDNA 4933424C09 gene	4933424C09Rik
1432618_at	1.15146	RIKEN cDNA 3110037L02 gene	3110037L02Rik
1432619_at	0.98523	---	---
1432620_at	0.84868	---	---
1432621_at	1.04710	RIKEN cDNA 5730410E19 gene	5730410E19Rik
1432622_a_at	1.03815	RIKEN cDNA 4930507D05 gene	4930507D05Rik
1432623_at	0.94804	"suppressor of var1, 3-like 1 (S. cerevisiae)"	Supv311
1432624_at	1.03148	RIKEN cDNA 5830487K18 gene	5830487K18Rik
1432625_at	0.98706	RIKEN cDNA 5830487K18 gene	5830487K18Rik
1432626_at	1.23782	RIKEN cDNA 5730507A11 gene	5730507A11Rik
1432627_at	1.00927	RIKEN cDNA 5730507A11 gene	5730507A11Rik
1432628_at	0.93712	chromobox homolog 3 (Drosophila HP1 gamma)	Cbx3
1432629_at	0.93017	---	---
1432630_at	0.90092	"tankyrase, TRF1-interacting ankyrin-related ADP-ribose polymerase"	Tnks
1432631_at	0.96051	RIKEN cDNA 1190002C06 gene	1190002C06Rik
1432632_at	0.92931	RIKEN cDNA 9430063H18 gene	9430063H18Rik
1432633_at	0.90304	RIKEN cDNA 9430063H18 gene	9430063H18Rik
1432634_at	1.07718	RIKEN cDNA 4933429H19 gene	4933429H19Rik
1432635_a_at	0.87978	translocation protein 1 Tloc1	
1432636_at	1.05105	---	---
1432637_at	1.06813	similar to Hippocalcin-like protein 1 (Visinin-like protein 3) (VILIP-3) (Neural visinin-like protein 3) (NVL-3) (NVP-3) /// similar to Hippocalcin-like protein 1 (Visinin-like protein 3) (VILIP-3) (Neural visinin-like protein 3) (NVL-3) (NVP-3)	LOC675539 /// LOC675641
1432638_at	0.93142	BCL2-associated athanogene 5 Bag5	
1432639_at	0.95719	RIKEN cDNA 4633401L03 gene	4633401L03Rik
1432640_at	0.98837	RIKEN cDNA 4633401L03 gene	4633401L03Rik
1432641_at	0.94072	RIKEN cDNA 5830490A12 gene	5830490A12Rik
1432642_at	1.09859	RIKEN cDNA 4933411E06 gene	4933411E06Rik
1432643_at	0.94903	RIKEN cDNA 4933411E06 gene	4933411E06Rik
1432644_at	1.00809	RIKEN cDNA 6430711C07 gene	6430711C07Rik
1432645_at	0.91000	Attractin Atrn	
1432646_a_at	1.01748	hypothetical LOC640370 /// hypothetical protein LOC667653 /// hypothetical protein LOC672953	LOC640370 /// LOC667653 /// LOC672953
1432647_at	0.95424	epidermal growth factor receptor	Egfr
1432648_at	0.91974	RIKEN cDNA 4930466F19 gene	4930466F19Rik
1432649_at	0.90121	RIKEN cDNA 3300002A11 gene	3300002A11Rik
1432650_at	1.08741	RIKEN cDNA 2510019K15 gene	2510019K15Rik
1432651_at	1.03747	RIKEN cDNA 2510019K15 gene	2510019K15Rik
1432652_at	0.85800	RIKEN cDNA 1700008H02 gene	1700008H02Rik

1432653_at	1.09380	RIKEN cDNA 1700008H02	gene	1700008H02Rik
1432654_at	1.08385	RIKEN cDNA 5830426C09	gene	5830426C09Rik
1432655_at	1.05476	RIKEN cDNA 3222402N08	gene	3222402N08Rik
1432656_at	0.89544	RIKEN cDNA 3222402N08	gene	3222402N08Rik
1432657_at	1.00871	a disintegrin and metallopeptidase domain 1b	Adam1b	
1432658_at	0.95116	RIKEN cDNA 2810403D21	gene	2810403D21Rik
1432659_at	0.95077	RIKEN cDNA 4930402H05	gene	4930402H05Rik
1432660_at	0.95567	RIKEN cDNA 5830420C07	gene	5830420C07Rik
1432661_at	0.99027	RIKEN cDNA 0610042E11	gene	0610042E11Rik
1432662_at	0.84264	RIKEN cDNA 0610042E11	gene	0610042E11Rik
1432663_at	0.97637	RIKEN cDNA 1500005C15	gene	1500005C15Rik
1432664_at	0.91573	RIKEN cDNA 4933404M09	gene	4933404M09Rik
1432665_at	0.88848	RIKEN cDNA 2210416J07	gene	2210416J07Rik
1432666_at	1.06680	---	---	
1432667_at	0.98604	RIKEN cDNA 2210411G17	gene	2210411G17Rik
1432668_at	0.93783	RIKEN cDNA 2210411G17	gene	2210411G17Rik
1432669_at	1.09780	RIKEN cDNA 9030420N05	gene	9030420N05Rik
1432670_at	0.86808	RIKEN cDNA 9030420N05	gene	9030420N05Rik
1432671_at	0.90707	RIKEN cDNA 4933407A17	gene	4933407A17Rik
1432672_at	0.92159	RIKEN cDNA 4933407A17	gene	4933407A17Rik
1432673_at	1.00657	RIKEN cDNA 2300010F08	gene	2300010F08Rik
1432674_at	1.04805	RIKEN cDNA 2300010F08	gene	2300010F08Rik
1432675_at	0.94125	midasin homolog (yeast) Mdn1		
1432676_at	1.22246	---	---	
1432677_at	0.90373	Golgi-specific brefeldin A-resistance factor 1	Gbf1	
1432678_at	1.01663	integrin alpha V	Itgav	
1432679_at	0.89140	integrin alpha V	Itgav	
1432680_at	1.16647	RIKEN cDNA 9130009M17	gene	9130009M17Rik
1432681_at	0.98506	RIKEN cDNA 9130009M17	gene	9130009M17Rik
1432682_at	1.00320	RIKEN cDNA 5530400K19	gene	5530400K19Rik
1432683_at	1.03575	RIKEN cDNA 5530400K19	gene	5530400K19Rik
1432684_at	1.03563	RIKEN cDNA 4921521C08	gene	4921521C08Rik
1432685_at	1.06315	MON1 homolog b (yeast) Mon1b		
1432686_at	1.00517	RIKEN cDNA 4833406M21	gene	4833406M21Rik
1432687_at	0.93560	RIKEN cDNA 4833406M21	gene	4833406M21Rik
1432688_at	0.94998	RIKEN cDNA 4631422I05	gene	4631422I05Rik
1432689_at	0.97360	RIKEN cDNA 4631422I05	gene	4631422I05Rik
1432690_at	0.99206	RIKEN cDNA 9030407C09	gene	9030407C09Rik
1432691_at	0.94144	RIKEN cDNA 9030407C09	gene	9030407C09Rik
1432692_at	0.99882	RIKEN cDNA 4930517G19	gene	4930517G19Rik
1432693_at	1.00623	RIKEN cDNA 4930517G19	gene	4930517G19Rik
1432694_at	1.10929	RIKEN cDNA 4921527H02	gene	4921527H02Rik
1432695_at	0.97375	RIKEN cDNA 4921527H02	gene	4921527H02Rik

1432696_at	0.91522	RIKEN cDNA 5830431M20	gene	5830431M20Rik
1432697_at	1.09575	RIKEN cDNA 5830431M20	gene	5830431M20Rik
1432698_at	0.99401	RIKEN cDNA 2900024I21	gene	2900024I21Rik
1432699_at	1.02784	RIKEN cDNA 2900024I21	gene	2900024I21Rik
1432700_at	0.98903	RIKEN cDNA 2810047J09	gene	2810047J09Rik
1432701_at	1.11008	RIKEN cDNA 2810403G07	gene	2810403G07Rik
1432702_at	1.03847	RIKEN cDNA 2810403G07	gene	2810403G07Rik
1432703_at	1.15265	RIKEN cDNA 5730405A17	gene	5730405A17Rik
1432704_at	0.89754	RIKEN cDNA 5730405A17	gene	5730405A17Rik
1432705_at	0.95233	RIKEN cDNA 5330428N10	gene	5330428N10Rik
1432706_at	0.93841	RIKEN cDNA 5330428N10	gene	5330428N10Rik
1432707_at	0.98619	3-phosphoglycerate dehydrogenase /// similar to 3-phosphoglycerate dehydrogenase /// similar to 3-phosphoglycerate dehydrogenase /// similar to 3-phosphoglycerate dehydrogenase /// similar to 3-phosphoglycerate dehydrogenase /// similar to 3-phosphoglycerate dehydrogenase /// similar to 3-phosphoglycerate dehydrogenase /// similar to 3-phosphoglycerate dehydrogenase /// similar to 3-phosphoglycerate dehydrogenase /// similar to 3-phosphoglycerate dehydrogenase /// similar to 3-phosphoglycerate dehydrogenase /// similar to 3-phosphoglycerate dehydrogenase /// similar to 3-phosphoglycerate dehydrogenase Phgdh /// LOC382931 /// LOC384524 /// LOC385344 /// LOC547171 /// LOC627427 /// LOC666422 /// LOC666875 /// LOC668771 /// LOC669671 /// LOC669985 /// LOC670155 /// LOC670187 /// LOC671102 /// LOC671972 /// LOC673015 /// LOC675010		
1432708_at	0.92851	---	---	
1432709_at	1.12874	RIKEN cDNA 4930505O19	gene	4930505O19Rik
1432710_at	0.96130	RIKEN cDNA 4930505O19	gene	4930505O19Rik
1432711_at	1.08794	RIKEN cDNA 4933425M03	gene	4933425M03Rik
1432712_at	1.07415	RIKEN cDNA 4933425M03	gene	4933425M03Rik
1432713_at	0.94192	RIKEN cDNA 6430709C05	gene	6430709C05Rik
1432714_at	0.94613	RIKEN cDNA 6430709C05	gene	6430709C05Rik
1432715_at	0.94227	"leucine rich repeat protein 1, neuronal"		Lrrn1
1432716_at	0.91161	"leucine rich repeat protein 1, neuronal"		Lrrn1
1432717_at	0.96478	RIKEN cDNA 1810015A16	gene	1810015A16Rik
1432718_at	1.14264	---	---	
1432719_at	1.23566	RIKEN cDNA 4833412K13	gene	4833412K13Rik
1432720_at	0.91214	RIKEN cDNA 4833412K13	gene	4833412K13Rik
1432721_at	0.87701	RIKEN cDNA 5830424K16	gene	5830424K16Rik
1432722_at	0.87542	RIKEN cDNA 4933408K01	gene	4933408K01Rik
1432723_at	1.04477	RIKEN cDNA 4933408K01	gene	4933408K01Rik
1432724_at	0.98208	RIKEN cDNA 4930445G23	gene	4930445G23Rik
1432725_at	0.96952	RIKEN cDNA 4930445G23	gene	4930445G23Rik
1432726_at	1.06815	"proteasome (prosome, macropain) 26S subunit, non-ATPase, 11"		Psm11
1432727_at	1.10022	"proteasome (prosome, macropain) 26S subunit, non-ATPase, 11"		/// similar to

26S proteasome non-ATPase regulatory subunit 11 (26S proteasome regulatory subunit S9) (26S proteasome regulatory subunit p44.5) /// similar to 26S proteasome non-ATPase regulatory subunit 11 (26S proteasome regulatory subunit S9) (26S proteasome regulatory subunit p44.5)" Psm11 /// LOC627316 /// LOC671671

1432728_at	0.94247	RIKEN cDNA 5830433D23	gene	5830433D23Rik
1432729_at	1.04150	RIKEN cDNA 5830415G21	gene	5830415G21Rik
1432730_at	0.93603	RIKEN cDNA 5830415G21	gene	5830415G21Rik
1432731_at	0.86481	RIKEN cDNA 5830437K03	gene	5830437K03Rik
1432732_at	0.98500	---	---	
1432733_at	1.01626	RIKEN cDNA 4933402E15	gene	4933402E15Rik
1432734_at	1.10008	RIKEN cDNA 4933402E15	gene	4933402E15Rik
1432735_at	0.86538	RIKEN cDNA 1700017H01	gene	1700017H01Rik
1432736_at	1.02365	RIKEN cDNA 4933430H06	gene	4933430H06Rik
1432737_at	0.95133	RIKEN cDNA 2810026P18	gene	2810026P18Rik
1432738_at	0.88558	RIKEN cDNA 2810026P18	gene	2810026P18Rik
1432739_at	0.89729	RIKEN cDNA 2900060K15	gene	2900060K15Rik
1432740_at	1.01084	RIKEN cDNA 4833420L08	gene	4833420L08Rik
1432741_at	1.08192	RIKEN cDNA 9530071P10	gene	9530071P10Rik
1432742_at	1.05984	---	---	
1432743_at	0.99728	RIKEN cDNA 4930443G03	gene	4930443G03Rik
1432744_at	0.88964	RIKEN cDNA 4930443G03	gene	4930443G03Rik
1432745_at	1.17433	RIKEN cDNA 9430092D12	gene	9430092D12Rik
1432746_at	0.98040	RIKEN cDNA 6030442H21	gene	6030442H21Rik
1432747_at	1.01876	RIKEN cDNA 4933403J19	gene	4933403J19Rik
1432748_at	0.94867	RIKEN cDNA 4933403J19	gene	4933403J19Rik
1432749_at	0.78688	---	---	
1432750_at	0.97177	RIKEN cDNA 2810409C01	gene	2810409C01Rik
1432751_at	1.00056	RIKEN cDNA 4930403O18	gene	4930403O18Rik
1432752_at	1.05279	RIKEN cDNA 4930403O18	gene	4930403O18Rik
1432753_at	0.95487	RIKEN cDNA 9130214F15	gene	9130214F15Rik
1432754_at	1.01040	---	---	
1432755_at	1.06859	RIKEN cDNA 2210409O19	gene	2210409O19Rik
1432756_at	0.97383	RIKEN cDNA 2210409O19	gene	2210409O19Rik
1432757_at	2.63396	RIKEN cDNA 2900011L18	gene	2900011L18Rik
1432758_at	0.98232	RIKEN cDNA 2900011L18	gene	2900011L18Rik
1432759_at	1.27486	RIKEN cDNA 6330571C24	gene	6330571C24Rik
1432760_at	1.00223	BAT2 domain containing 1	Bat2d	
1432761_at	0.94151	RIKEN cDNA 4930551I15	gene	4930551I15Rik
1432762_at	0.96507	RIKEN cDNA 9530025H10	gene	9530025H10Rik
1432763_at	0.84001	proline-rich Gla (G-carboxyglutamic acid) polypeptide 2	Prrg2	
1432764_at	0.94104	RIKEN cDNA 4933404I11	gene	4933404I11Rik
1432765_at	1.00937	RIKEN cDNA C030007D22	gene	C030007D22Rik
1432766_at	0.90313	RIKEN cDNA E130112B07	gene	E130112B07Rik

1432767_at	0.93664	RIKEN cDNA E130112B07	gene	E130112B07Rik
1432768_at	0.99135	RIKEN cDNA 6430710M23	gene	6430710M23Rik
1432769_at	0.95406	"solute carrier family 44, member 1"		Slc44a1
1432770_at	0.95546	RIKEN cDNA 0610040A22	gene	0610040A22Rik
1432771_at	1.03766	"solute carrier family 9 (sodium/hydrogen exchanger), member 3"		Slc9a3
1432772_at	1.03530	RIKEN cDNA 1700081H04	gene	1700081H04Rik
1432773_at	0.97648	RIKEN cDNA 2610105M22	gene	2610105M22Rik
1432774_at	0.97449	RIKEN cDNA 2610105M22	gene	2610105M22Rik
1432775_at	0.95961	RIKEN cDNA 9430087J23	gene	9430087J23Rik
1432776_at	0.89598	RIKEN cDNA 9430087J23	gene	9430087J23Rik
1432777_at	0.88414	RIKEN cDNA 9430032J07	gene	9430032J07Rik
1432778_at	0.91083	---	---	
1432779_at	1.00675	RIKEN cDNA 1700037N05	gene	1700037N05Rik
1432780_at	0.98297	---	---	
1432781_at	0.89403	PDZ domain containing 6		Pdzd6
1432782_at	0.96068	RIKEN cDNA 4933435C09	gene	4933435C09Rik
1432783_at	1.11148	RIKEN cDNA 4930556A17	gene	4930556A17Rik
1432784_at	0.84384	RIKEN cDNA 4930556A17	gene	4930556A17Rik
1432785_at	0.96053	RIKEN cDNA A930028O11	gene	A930028O11Rik
1432786_at	1.09789	RIKEN cDNA 4833420D23	gene	4833420D23Rik
1432787_at	1.03046	RIKEN cDNA 6720420G18	gene	6720420G18Rik
1432788_at	0.97936	RIKEN cDNA 6720420G18	gene	6720420G18Rik
1432789_at	0.96659	RIKEN cDNA 8030476L19	gene	8030476L19Rik
1432790_at	1.14665	RIKEN cDNA 9030218A15	gene	9030218A15Rik
1432791_at	0.97357	RIKEN cDNA 9030218A15	gene	9030218A15Rik
1432792_at	1.04190	RIKEN cDNA 4930401B11	gene	4930401B11Rik
1432793_at	0.87752	RIKEN cDNA 4930401B11	gene	4930401B11Rik
1432794_at	1.07891	RIKEN cDNA 2310058F05	gene	2310058F05Rik
1432795_at	1.05850	RIKEN cDNA 2310058F05	gene	2310058F05Rik
1432796_at	1.08847	RIKEN cDNA 3110067C02	gene	3110067C02Rik
1432797_at	0.84768	RIKEN cDNA 2900060N12	gene	2900060N12Rik
1432798_at	1.05547	RIKEN cDNA 9530023I19	gene	9530023I19Rik
1432799_at	0.92777	RIKEN cDNA C030011G24	gene	C030011G24Rik
1432800_at	0.93308	RIKEN cDNA C030011G24	gene	C030011G24Rik
1432801_at	1.11535	RIKEN cDNA 4930418C01	gene	4930418C01Rik
1432802_at	1.00103	RIKEN cDNA 9230106L01	gene	9230106L01Rik
1432803_at	1.02124	RIKEN cDNA A230101C19	gene	A230101C19Rik
1432804_at	1.02412	RIKEN cDNA A230101C19	gene	A230101C19Rik
1432805_at	0.93501	RIKEN cDNA 8030431A06	gene	8030431A06Rik
1432806_at	0.89067	RIKEN cDNA 9430099H24	gene	9430099H24Rik
1432807_at	0.95421	RIKEN cDNA 5830426K05	gene	5830426K05Rik
1432808_at	1.05212	RIKEN cDNA 5830426K05	gene	5830426K05Rik
1432809_at	1.01234	RIKEN cDNA 1700113P08	gene	1700113P08Rik

1432810_at	1.00601	tetratricopeptide repeat domain 32	Ttc32
1432811_at	0.97299	---	---
1432812_at	1.09415	RIKEN cDNA 4930554P06 gene	4930554P06Rik
1432813_at	0.99796	RIKEN cDNA 2900064F13 gene	2900064F13Rik
1432814_at	0.99392	RIKEN cDNA 2900064F13 gene	2900064F13Rik
1432815_at	0.91030	RIKEN cDNA 4833419A21 gene	4833419A21Rik
1432816_s_at	1.06539	RIKEN cDNA 4930413G21 gene	4930413G21Rik
1432817_x_at	0.81393	RIKEN cDNA 4930413G21 gene	4930413G21Rik
1432818_at	0.92603	RIKEN cDNA 1190002C06 gene	1190002C06Rik
1432819_at	0.90628	RIKEN cDNA 1190002C06 gene	1190002C06Rik
1432820_at	1.09697	"proteasome (prosome, macropain) 26S subunit, non-ATPase, 7"	Psmc7
1432821_at	1.01643	"nuclear factor of activated T-cells, cytoplasmic, calcineurin-dependent 4"	Nfatc4
1432822_at	0.97484	RIKEN cDNA 4930557B21 gene	4930557B21Rik
1432823_at	1.01651	synaptophysin-like 2	Sypl2
1432824_at	0.96050	RIKEN cDNA 2900018N21 gene	2900018N21Rik
1432825_at	0.97895	RIKEN cDNA 2900018N21 gene	2900018N21Rik
1432826_a_at	0.91302	CD80 antigen	Cd80
1432827_x_at	0.89489	ubiquitin C	Ubc
1432828_at	1.17760	---	---
1432829_at	0.95250	"enolase 1, alpha non-neuron /// similar to enolase 1, alpha non-neuron /// similar to T21B10.2b /// similar to enolase 1, alpha non-neuron"	Eno1 /// LOC433182 /// LOC671462 /// LOC674877
1432830_at	1.05166	RIKEN cDNA 6230424H07 gene	6230424H07Rik
1432831_at	1.00166	RIKEN cDNA 6230424H07 gene	6230424H07Rik
1432832_at	0.99467	RIKEN cDNA 4933435G04 gene	4933435G04Rik
1432833_at	1.04338	RIKEN cDNA 4933435G04 gene	4933435G04Rik
1432834_at	0.96319	carboxypeptidase B2 (plasma)	Cpb2
1432835_at	1.04841	carboxypeptidase B2 (plasma)	Cpb2
1432836_at	1.02727	RIKEN cDNA 4833410I11 gene	4833410I11Rik
1432837_at	1.03923	RIKEN cDNA 2700080J24 gene	2700080J24Rik
1432838_at	0.99310	RIKEN cDNA 6430598H11 gene	6430598H11Rik
1432839_at	1.04066	RIKEN cDNA 6430598H11 gene	6430598H11Rik
1432840_at	0.96386	UDP-N-acetyl-alpha-D-galactosamine:polypeptide N-acetylgalactosaminyltransferase 11	Galnt11
1432841_at	0.99569	RIKEN cDNA E130116L18 gene	E130116L18Rik
1432842_s_at	1.00837	"tyrosine 3-monooxygenase/tryptophan 5-monooxygenase activation protein, theta polypeptide"	Ywhaq
1432843_at	1.20756	RIKEN cDNA 5730405O12 gene	5730405O12Rik
1432844_at	0.89187	phosphatase and actin regulator 4	Phactr4
1432845_at	0.90856	---	---
1432846_at	0.96259	RIKEN cDNA 5033428C03 gene	5033428C03Rik
1432847_at	0.78917	RIKEN cDNA 2810457G06 gene	2810457G06Rik

1432848_at	0.88835	RIKEN cDNA 1200004M23 gene	1200004M23Rik	
1432849_at	1.00419	---	---	
1432850_at	2.49113	RIKEN cDNA 5430434G16 gene	5430434G16Rik	
1432851_at	0.91244	phosphatase and actin regulator 1	Phactr1	
1432852_at	1.00274	phosphatase and actin regulator 1	Phactr1	
1432853_at	0.94815	RIKEN cDNA 2900057C01 gene	2900057C01Rik	
1432854_at	1.08957	RIKEN cDNA 1700018P08 gene	1700018P08Rik	
1432855_at	1.01053	potassium channel tetramerisation domain containing 3	Kctd3	
1432856_at	0.94762	---	---	
1432857_at	1.00048	RIKEN cDNA 6430500D05 gene	6430500D05Rik	
1432858_at	1.00852	RIKEN cDNA D730048M19 gene	D730048M19Rik	
1432859_at	1.41213	RIKEN cDNA D730048M19 gene	D730048M19Rik	
1432860_at	1.17244	pannexin 3	Panx3	
1432861_at	0.96329	RIKEN cDNA 2900046F13 gene	2900046F13Rik	
1432862_at	0.91643	"NK2 transcription factor related, locus 4 (Drosophila)"	Nkx2-4	
1432863_a_at	1.00507	"NK2 transcription factor related, locus 4 (Drosophila)"	Nkx2-4	
1432864_at	1.00471	---	---	
1432865_at	1.00696	RIKEN cDNA D730050C22 gene	D730050C22Rik	
1432866_at	1.10884	RIKEN cDNA 6030442K20 gene	6030442K20Rik	
1432867_at	0.95844	RIKEN cDNA 9230112J17 gene	9230112J17Rik	
1432868_at	0.95280	RIKEN cDNA 2310047C04 gene	2310047C04Rik	
1432869_at	0.94790	---	---	
1432870_at	0.90265	RIKEN cDNA 4932429P19 gene	4932429P19Rik	
1432871_at	1.10546	RIKEN cDNA 4932429P19 gene	4932429P19Rik	
1432872_at	0.81326	RIKEN cDNA 4932702M13 gene	4932702M13Rik	
1432873_at	0.99248	RIKEN cDNA 4932702M13 gene	4932702M13Rik	
1432874_at	0.97944	RIKEN cDNA 4632404M16 gene	4632404M16Rik	
1432875_at	0.93776	RIKEN cDNA 4632409D06 gene	4632409D06Rik	
1432876_at	0.91450	RIKEN cDNA 4632418H02 gene	4632418H02Rik	
1432877_at	1.02364	RIKEN cDNA 4930544N03 gene	4930544N03Rik	
1432878_at	0.93461	RIKEN cDNA 4930544N03 gene	4930544N03Rik	
1432879_at	0.91295	RIKEN cDNA 6820402A03 gene	6820402A03Rik	
1432880_at	0.99805	RIKEN cDNA 6820402A03 gene	6820402A03Rik	
1432881_at	0.92114	RIKEN cDNA 4932437C15 gene	4932437C15Rik	
1432882_at	0.93752	RIKEN cDNA 4932431P20 gene	4932431P20Rik	
1432883_at	1.02214	RIKEN cDNA 4932431P20 gene	4932431P20Rik	
1432884_at	0.97657	RIKEN cDNA 2010003H20 gene	2010003H20Rik	
1432885_at	0.99909	RIKEN cDNA 4632432E15 gene	4632432E15Rik	
1432886_at	1.03809	RIKEN cDNA 5730488B01 gene	5730488B01Rik	
1432887_at	1.00131	RIKEN cDNA 3110048L19 gene	3110048L19Rik	
1432888_at	1.07400	RIKEN cDNA 4930455M05 gene	4930455M05Rik	
1432889_at	0.87282	RIKEN cDNA 4930466I24 gene	4930466I24Rik	
1432890_at	1.08394	RIKEN cDNA 2610011E03 gene	2610011E03Rik	

1432891_at	1.07827	RIKEN cDNA 5730564L20	gene	5730564L20Rik
1432892_at	0.94943	RIKEN cDNA 5730564L20	gene	5730564L20Rik
1432893_at	1.03561	RIKEN cDNA 9130011L11	gene	9130011L11Rik
1432894_at	1.01338	RIKEN cDNA 9130011L11	gene	9130011L11Rik
1432895_at	0.96732	RIKEN cDNA 9030409K20	gene	9030409K20Rik
1432896_at	1.03109	RIKEN cDNA 9030409K20	gene	9030409K20Rik
1432897_at	0.89647	RIKEN cDNA 4930435N07	gene	4930435N07Rik
1432898_at	0.90743	RIKEN cDNA 4930544L18	gene	4930544L18Rik
1432899_at	0.92724	RIKEN cDNA 4833408G04	gene	4833408G04Rik
1432900_at	1.02808	cDNA sequence BC026657	BC026657	
1432901_at	1.29682	cDNA sequence BC026657	BC026657	
1432902_at	0.82832	RIKEN cDNA 9130414P19	gene	9130414P19Rik
1432903_at	0.87436	---	---	
1432904_at	0.96006	RIKEN cDNA 3110037C07	gene	3110037C07Rik
1432905_at	1.02867	RIKEN cDNA 3110037C07	gene	3110037C07Rik
1432906_at	0.96775	RIKEN cDNA 1700126A01	gene	1700126A01Rik
1432907_at	1.06000	RIKEN cDNA 1700126A01	gene	1700126A01Rik
1432908_at	0.89426	RIKEN cDNA 2810410P21	gene	2810410P21Rik
1432909_at	0.96171	RIKEN cDNA 4933424H11	gene	4933424H11Rik
1432910_at	1.66988	BTB (POZ) domain containing 7	Btbd7	
1432911_at	1.03610	scaffold attachment factor B	Safb	
1432912_at	0.95248	---	---	
1432913_at	0.91359	leucine rich repeat containing 52	Lrrc52	
1432914_at	1.01932	leucine rich repeat containing 52	Lrrc52	
1432915_at	0.97016	RIKEN cDNA 5730407I07	gene	5730407I07Rik
1432916_at	1.07318	RIKEN cDNA 5730407I07	gene	5730407I07Rik
1432917_at	0.97497	RIKEN cDNA 4933408M05	gene	4933408M05Rik
1432918_at	1.44204	RIKEN cDNA 4921511E18	gene	4921511E18Rik
1432919_at	0.92823	RIKEN cDNA 4921511E18	gene	4921511E18Rik
1432920_at	0.95398	RIKEN cDNA 5330421C15	gene	5330421C15Rik
1432921_at	0.95952	---	---	
1432922_at	0.86105	RIKEN cDNA 4933416E14	gene	4933416E14Rik
1432923_at	1.03276	RIKEN cDNA 2810404I24	gene	2810404I24Rik
1432924_at	1.39091	RIKEN cDNA 2810404I24	gene	2810404I24Rik
1432925_at	0.97156	RIKEN cDNA 4933416A02	gene	4933416A02Rik
1432926_at	0.98600	RIKEN cDNA 2900069M18	gene	2900069M18Rik
1432927_at	0.84013	RIKEN cDNA 4921507K24	gene	4921507K24Rik
1432928_at	1.04952	RIKEN cDNA 4921507K24	gene	4921507K24Rik
1432929_at	1.02028	RIKEN cDNA 4930453O09	gene	4930453O09Rik
1432930_at	1.04628	RIKEN cDNA 4930453O09	gene	4930453O09Rik
1432931_at	0.89092	neurexin III	Nrxn3	
1432932_at	1.01502	---	---	
1432933_at	0.96859	RIKEN cDNA 4930429N05	gene	4930429N05Rik

1432934_at	0.90120	RIKEN cDNA 4930429N05	gene	4930429N05Rik
1432935_at	0.84937	RIKEN cDNA 5330433J24	gene	5330433J24Rik
1432936_at	0.92157	RIKEN cDNA 5330433J24	gene	5330433J24Rik
1432937_at	1.12439	RIKEN cDNA 5430439C14	gene	5430439C14Rik
1432938_at	1.09016	RIKEN cDNA 5430439C14	gene	5430439C14Rik
1432939_at	1.06455	RIKEN cDNA 5730405N03	gene	5730405N03Rik
1432940_at	1.07748	RIKEN cDNA 5730405N03	gene	5730405N03Rik
1432941_at	0.88731	RIKEN cDNA 5730408A14	gene	5730408A14Rik
1432942_at	0.97287	RIKEN cDNA 5730408A14	gene	5730408A14Rik
1432943_at	0.91161	RIKEN cDNA 2900046L07	gene	2900046L07Rik
1432944_at	0.93354	RIKEN cDNA 2900046L07	gene	2900046L07Rik
1432945_at	0.78428	RIKEN cDNA 5230400M06	gene	5230400M06Rik
1432946_at	1.06380	RIKEN cDNA 5230400M06	gene	5230400M06Rik
1432947_at	0.95701	RIKEN cDNA 4921519G19	gene	4921519G19Rik
1432948_at	1.03107	RIKEN cDNA 4921519G19	gene	4921519G19Rik
1432949_at	0.94841	RIKEN cDNA 5330421F21	gene	5330421F21Rik
1432950_at	1.00218	RIKEN cDNA 5330421F21	gene	5330421F21Rik
1432951_at	1.05936	RIKEN cDNA 4930448E22	gene	4930448E22Rik
1432952_at	1.01437	RIKEN cDNA 4930448E22	gene	4930448E22Rik
1432953_at	1.00360	RIKEN cDNA 4921520E09	gene	4921520E09Rik
1432954_at	1.06915	RIKEN cDNA 4921520E09	gene	4921520E09Rik
1432955_at	0.93607	RIKEN cDNA 4930444K16	gene	4930444K16Rik
1432956_at	0.96215	RIKEN cDNA 4930444K16	gene	4930444K16Rik
1432957_at	0.94262	RIKEN cDNA 5730407O05	gene	5730407O05Rik
1432958_at	1.00472	RIKEN cDNA 5730407O05	gene	5730407O05Rik
1432959_at	1.07642	RIKEN cDNA 4930447I22	gene	4930447I22Rik
1432960_at	0.99541	RIKEN cDNA 4930447I22	gene	4930447I22Rik
1432961_at	0.98615	RIKEN cDNA 2610024D14	gene	2610024D14Rik
1432962_at	1.16936	RIKEN cDNA 2610024D14	gene	2610024D14Rik
1432963_at	0.91471	RIKEN cDNA 2410049M19	gene	2410049M19Rik
1432964_at	1.12497	RIKEN cDNA 2410049M19	gene	2410049M19Rik
1432965_at	0.96821	RIKEN cDNA 4921507H08	gene	4921507H08Rik
1432966_at	0.98804	RIKEN cDNA 4921507H08	gene	4921507H08Rik
1432967_at	0.95796	RIKEN cDNA 4930456A14	gene	4930456A14Rik
1432968_at	1.13880	RIKEN cDNA 4930456A14	gene	4930456A14Rik
1432969_at	1.02074	RIKEN cDNA 4933423K11	gene	4933423K11Rik
1432970_at	1.09439	RIKEN cDNA 4933423K11	gene	4933423K11Rik
1432971_at	0.76360	RIKEN cDNA 4921518B13	gene	4921518B13Rik
1432972_at	0.98864	RIKEN cDNA 4921518B13	gene	4921518B13Rik
1432973_at	1.06073	RIKEN cDNA 4933421H12	gene	4933421H12Rik
1432974_at	1.01758	RIKEN cDNA 4933421H12	gene	4933421H12Rik
1432975_at	1.11448	RIKEN cDNA 2310038E17	gene	2310038E17Rik
1432976_at	1.09231	RIKEN cDNA 2310038E17	gene	2310038E17Rik

1432977_at	0.94250	RIKEN cDNA 9030607L02	gene	9030607L02Rik
1432978_at	1.35425	RIKEN cDNA 9030607L02	gene	9030607L02Rik
1432979_at	1.01694	GATA zinc finger domain containing 2B		Gatad2b
1432980_at	0.87269	RIKEN cDNA 5730420F10	gene	5730420F10Rik
1432981_at	1.02847	mitogen-activated protein kinase kinase kinase 9		Map3k9
1432982_at	0.99843	RIKEN cDNA 4930456J16	gene	4930456J16Rik
1432983_at	1.10549	RIKEN cDNA 1700026H06	gene	1700026H06Rik
1432984_at	1.04568	RIKEN cDNA 1700026H06	gene	1700026H06Rik
1432985_at	1.08119	RIKEN cDNA 5430402M21	gene	5430402M21Rik
1432986_at	0.98783	RIKEN cDNA 5430402M21	gene	5430402M21Rik
1432987_at	0.89557	---	---	
1432988_at	1.08874	RIKEN cDNA 2610040L17	gene	2610040L17Rik
1432989_at	0.98100	RIKEN cDNA 2900011F02	gene	2900011F02Rik
1432990_at	0.94746	RIKEN cDNA 5830495A06	gene	5830495A06Rik
1432991_at	0.89043	RIKEN cDNA 4930519E07	gene	4930519E07Rik
1432992_at	0.99050	RIKEN cDNA 4930403L11	gene	4930403L11Rik
1432993_at	1.19143	RIKEN cDNA 4930403L11	gene	4930403L11Rik
1432994_at	1.01184	RIKEN cDNA 2900057E15	gene	2900057E15Rik
1432995_at	1.14462	RIKEN cDNA 2900057E15	gene	2900057E15Rik
1432996_at	1.01177	---	---	
1432997_at	1.03985	RIKEN cDNA 5830462P14	gene	5830462P14Rik
1432998_at	0.99363	RIKEN cDNA 5830462O15	gene	5830462O15Rik
1432999_at	0.99991	phosphodiesterase 11A		Pde11a
1433000_at	0.97244	---	---	
1433001_at	0.94539	RIKEN cDNA 1700019L13	gene	1700019L13Rik
1433002_at	0.92635	RIKEN cDNA 9030613N10	gene	9030613N10Rik
1433003_at	0.92012	RIKEN cDNA 4933423N03	gene	4933423N03Rik
1433004_at	0.98333	RIKEN cDNA 4833446E11	gene	4833446E11Rik
1433005_at	1.05533	RIKEN cDNA 4833446E11	gene	4833446E11Rik
1433006_at	0.92858	RIKEN cDNA 5430430B14	gene	5430430B14Rik
1433007_at	0.98862	RIKEN cDNA 5430430B14	gene	5430430B14Rik
1433008_at	0.95657	RIKEN cDNA 6330526H18	gene	6330526H18Rik
1433009_at	1.10181	RIKEN cDNA 9030607J07	gene	9030607J07Rik
1433010_at	0.84929	RIKEN cDNA 5530400N10	gene	5530400N10Rik
1433011_at	1.03079	RIKEN cDNA 4921506L19	gene	4921506L19Rik
1433012_at	0.97052	MAX gene associated		Mga
1433013_at	0.95748	RIKEN cDNA 4930485G23	gene	4930485G23Rik
1433014_at	1.01076	RIKEN cDNA 6330436F06	gene	6330436F06Rik
1433015_at	0.99225	RIKEN cDNA 6330436F06	gene	6330436F06Rik
1433016_s_at	1.14605	RIKEN cDNA 4933412F11	gene	RIKEN cDNA 4930529H12 gene
hypothetical LOC622653	///	hypothetical LOC624958	///	hypothetical LOC625130
///		///		/// similar to RIKEN
cDNA 4930529H12	gene	4933412F11Rik	///	4930529H12Rik
///		///		/// LOC622653
///		///		/// LOC624958
///		///		/// LOC625130
///		///		/// LOC665740

1433017_at	0.99825	RIKEN cDNA 8430420C20	gene	8430420C20Rik
1433018_at	0.95807	RIKEN cDNA 4930560O18	gene	4930560O18Rik
1433019_at	0.85004	RIKEN cDNA 4833419G08	gene	4833419G08Rik
1433020_at	0.95193	RIKEN cDNA 5830413G11	gene	5830413G11Rik
1433021_at	0.89207	RIKEN cDNA 5830413G11	gene	5830413G11Rik
1433022_at	0.85621	RIKEN cDNA 2310068G24	gene	2310068G24Rik
1433023_at	0.90272	RIKEN cDNA 2310068G24	gene	2310068G24Rik
1433024_at	1.09972	RIKEN cDNA 4930421C12	gene	4930421C12Rik
1433025_x_at	0.97287	"BCL2/adenovirus E1B interacting protein 1, NIP2"		Bnip2
1433026_at	0.95193	---	---	
1433027_at	1.19551	RIKEN cDNA 5830485P09	gene	5830485P09Rik
1433028_at	0.91758	RIKEN cDNA 4833411I10	gene	4833411I10Rik
1433029_at	1.01759	"potassium inwardly-rectifying channel, subfamily J, member 9"		Kcnj9
1433030_at	1.06784	RIKEN cDNA 5730406E14	gene	5730406E14Rik
1433031_at	0.94391	RIKEN cDNA 2900064K03	gene	2900064K03Rik
1433032_at	1.02927	RIKEN cDNA 2900064K03	gene	2900064K03Rik
1433033_at	0.95739	RIKEN cDNA 2310007H11	gene	2310007H11Rik
1433034_at	0.99409	RIKEN cDNA 2310007H11	gene	2310007H11Rik
1433035_at	0.87154	RIKEN cDNA 5430430K15	gene	5430430K15Rik
1433036_at	0.94474	RIKEN cDNA 4930564B12	gene	4930564B12Rik
1433037_at	0.95580	RIKEN cDNA 5830434F19	gene	5830434F19Rik
1433038_at	0.89566	"Visual cortex cDNA, RIKEN full-length enriched library, clone:K230311O21 product:unclassifiable, full insert sequence"	---	
1433039_at	1.02207	RIKEN cDNA 4930555B11	gene	4930555B11Rik
1433040_at	0.84525	RIKEN cDNA 2900018E21	gene	2900018E21Rik
1433041_at	1.01446	Rab9 effector protein with kelch motifs		Rabepk
1433042_at	0.95376	---	---	
1433043_at	1.14299	chondroitin sulfate proteoglycan 2		Cspg2
1433044_at	1.09760	chondroitin sulfate proteoglycan 2		Cspg2
1433045_at	1.01572	RIKEN cDNA 4930535F04	gene	4930535F04Rik
1433046_at	0.97062	RIKEN cDNA 4930535F04	gene	4930535F04Rik
1433047_at	1.28300	RIKEN cDNA 5330430B06	gene	5330430B06Rik
1433048_at	1.09424	RIKEN cDNA 4933428L01	gene	4933428L01Rik
1433049_at	1.05413	RIKEN cDNA 5730478J17	gene	5730478J17Rik
1433050_at	1.24097	RIKEN cDNA 5730478J17	gene	5730478J17Rik
1433051_at	1.02780	RIKEN cDNA 2610011I18	gene	2610011I18Rik
1433052_at	0.98517	"ELAV (embryonic lethal, abnormal vision, Drosophila)-like 2 (Hu antigen B)"		Elavl2
1433053_at	1.10446	integrin beta 2-like		Itgb2l
1433054_at	1.15667	integrin beta 2-like		Itgb2l
1433055_at	1.10887	RIKEN cDNA 4931407J08	gene	4931407J08Rik
1433056_at	1.00769	RIKEN cDNA 4931407J08	gene	4931407J08Rik
1433057_at	0.99632	RIKEN cDNA 4933411O13	gene	4933411O13Rik

1433058_at	1.08030	RIKEN cDNA 4933411O13 gene	4933411O13Rik
1433059_at	1.02776	RIKEN cDNA 4930439G18 gene	4930439G18Rik
1433060_at	1.06301	RIKEN cDNA 6230426I18 gene	6230426I18Rik
1433061_at	1.03068	RIKEN cDNA 4933423L19 gene	4933423L19Rik
1433062_at	1.04332	WD repeat domain 66 Wdr66	
1433063_at	0.99611	WD repeat domain 66 Wdr66	
1433064_at	1.04633	acyl-CoA thioesterase 12 Acot12	
1433065_at	0.86145	RIKEN cDNA 6330510M09 gene	6330510M09Rik
1433066_at	1.04136	RIKEN cDNA 6330510M09 gene	6330510M09Rik
1433067_at	1.04121	RIKEN cDNA 6330582A15 gene	6330582A15Rik
1433068_at	1.06763	RIKEN cDNA 6330582A15 gene	6330582A15Rik
1433069_at	1.04817	---	---
1433070_at	1.02848	RIKEN cDNA 5830433M15 gene	5830433M15Rik
1433071_at	0.89839	RIKEN cDNA 6430605C03 gene	6430605C03Rik
1433072_at	1.19848	RIKEN cDNA 4933425E08 gene	4933425E08Rik
1433073_at	1.03727	RIKEN cDNA 4933425E08 gene	4933425E08Rik
1433074_at	0.95059	RIKEN cDNA 4931412I15 gene	4931412I15Rik
1433075_at	1.02134	RIKEN cDNA 4931412I15 gene	4931412I15Rik
1433076_at	0.97788	RIKEN cDNA 5430402O13 gene	5430402O13Rik
1433077_at	0.85606	RIKEN cDNA 2700078F05 gene	2700078F05Rik
1433078_at	0.93099	RIKEN cDNA 5830435N06 gene	5830435N06Rik
1433079_at	1.06729	RIKEN cDNA 4933436P19 gene	4933436P19Rik
1433080_at	1.03409	RIKEN cDNA 4933436P19 gene	4933436P19Rik
1433081_at	0.95340	RIKEN cDNA 4933407I18 gene	4933407I18Rik
1433082_at	1.18292	RIKEN cDNA 4930448K20 gene	4930448K20Rik
1433083_at	1.02940	RIKEN cDNA 4930448K20 gene	4930448K20Rik
1433084_at	1.05962	RIKEN cDNA 4930402C16 gene	4930402C16Rik
1433085_at	1.14799	RIKEN cDNA 5930427J20 gene	5930427J20Rik
1433086_at	1.02779	RIKEN cDNA 5930427J20 gene	5930427J20Rik
1433087_at	1.01142	RIKEN cDNA 5330430C04 gene	5330430C04Rik
1433088_at	0.93990	RIKEN cDNA 5330430C04 gene	5330430C04Rik
1433089_at	1.06387	---	---
1433090_at	1.10468	---	---
1433091_at	1.08587	RIKEN cDNA 6430514K02 gene	6430514K02Rik
1433092_at	1.00755	RIKEN cDNA 3110062G12 gene	3110062G12Rik
1433093_at	0.96930	RIKEN cDNA 2900092N22 gene	2900092N22Rik
1433094_at	1.02555	"solute carrier family 1 (glial high affinity glutamate transporter), member 2" Slc1a2	
1433095_at	1.07594	---	---
1433096_at	0.97659	RIKEN cDNA 4930473M17 gene	4930473M17Rik
1433097_at	1.03830	RIKEN cDNA 4930473M17 gene	4930473M17Rik
1433098_at	1.09281	---	---
1433099_at	0.80675	RIKEN cDNA 4933428L12 gene	4933428L12Rik

1433100_at	1.12369	transmembrane protein 153	Tmem153
1433101_at	1.20827	RIKEN cDNA 9030419F21	gene 9030419F21Rik
1433102_at	0.98454	RIKEN cDNA 3100002H20	gene 3100002H20Rik
1433103_at	0.96947	hydrolethalus syndrome 1	Hyls1
1433104_at	0.98404	RIKEN cDNA 5730412F04	gene 5730412F04Rik
1433105_at	1.00893	RIKEN cDNA 5830438M01	gene 5830438M01Rik
1433106_at	1.08285	RIKEN cDNA 5830438M01	gene 5830438M01Rik
1433107_at	0.99893	---	---
1433108_at	1.03743	"T-cell receptor beta, joining region /// T-cell receptor beta, variable 8.2 /// T-cell receptor beta, variable 13 /// similar to T-cell receptor beta-1 chain C region" Tcrb-J /// Tcrb-V8.2 /// Tcrb-V13 /// LOC669847	
1433109_at	0.91602	macrophage migration inhibitory factor	Mif
1433110_at	1.06854	RIKEN cDNA 5830474E16	gene 5830474E16Rik
1433111_at	1.13780	NHS-like 1	Nhsl1
1433112_at	1.02344	RIKEN cDNA 4933424L07	gene 4933424L07Rik
1433113_at	0.92987	RIKEN cDNA 4921504P20	gene 4921504P20Rik
1433114_at	0.99099	RIKEN cDNA 4921504P20	gene 4921504P20Rik
1433115_at	1.01790	RIKEN cDNA 5730552O08	gene 5730552O08Rik
1433116_at	0.93102	RIKEN cDNA 5730552O08	gene 5730552O08Rik
1433117_at	1.05773	RIKEN cDNA 2300004M11	gene 2300004M11Rik
1433118_at	0.83355	RIKEN cDNA 2300004M11	gene 2300004M11Rik
1433119_at	0.96456	RIKEN cDNA 4930456G14	gene 4930456G14Rik
1433120_at	1.00203	RIKEN cDNA 9530002O20	gene 9530002O20Rik
1433121_at	1.11205	RIKEN cDNA 5830490A04	gene 5830490A04Rik
1433122_at	0.89812	RIKEN cDNA 2900069G24	gene 2900069G24Rik
1433123_at	0.86853	RIKEN cDNA 2900069G24	gene 2900069G24Rik
1433124_at	1.07498	RIKEN cDNA 4930534I15	gene 4930534I15Rik
1433125_at	1.02965	RIKEN cDNA 5330422M15	gene 5330422M15Rik
1433126_at	0.99702	RIKEN cDNA 5330422M15	gene 5330422M15Rik
1433127_at	0.87101	RIKEN cDNA 4930513L20	gene 4930513L20Rik
1433128_at	1.02083	RIKEN cDNA 4930423C22	gene 4930423C22Rik
1433129_at	1.07821	RIKEN cDNA 2900040J22	gene 2900040J22Rik
1433130_at	0.95857	RIKEN cDNA 8430418B16	gene 8430418B16Rik
1433131_at	0.98169	RIKEN cDNA 5033425B01	gene 5033425B01Rik
1433132_at	0.96369	EDAR (ectodysplasin-A receptor)-associated death domain Edaradd	
1433133_at	1.05075	EDAR (ectodysplasin-A receptor)-associated death domain Edaradd	
1433134_at	1.02498	RIKEN cDNA 5830448L01	gene 5830448L01Rik
1433135_at	1.02431	"golgi autoantigen, golgin subfamily b, macrogolgin 1" Golgb1	
1433136_at	0.96036	RIKEN cDNA 4930432H08	gene 4930432H08Rik
1433137_at	0.91077	RIKEN cDNA 5031415H12	gene 5031415H12Rik
1433138_at	0.91647	---	---
1433139_at	0.88297	BTB (POZ) domain containing 14B Btbd14b	
1433140_a_at	0.98666	BTB (POZ) domain containing 14B Btbd14b	

1433141_at	1.03947	RIKEN cDNA 4921504P05	gene	4921504P05Rik
1433142_at	1.15723	RIKEN cDNA 4921504P05	gene	4921504P05Rik
1433143_at	0.91197	RIKEN cDNA 9230118N17	gene	9230118N17Rik
1433144_at	0.97889	"asparagine-linked glycosylation 2 homolog (yeast, alpha-1,3-mannosyltransferase)"	Alg2	
1433145_at	0.80585	RIKEN cDNA 4933431C10	gene	4933431C10Rik
1433146_at	0.91999	caldesmon 1	Cald1	
1433147_at	2.05215	caldesmon 1	Cald1	
1433148_at	1.18991	RIKEN cDNA 4930513N20	gene	4930513N20Rik
1433149_at	1.02281	RIKEN cDNA 4930513N20	gene	4930513N20Rik
1433150_at	0.98866	RIKEN cDNA 4933415B22	gene	4933415B22Rik
1433151_at	0.90112	RIKEN cDNA 4933415B22	gene	4933415B22Rik
1433152_at	1.03812	RIKEN cDNA 2900022B07	gene	2900022B07Rik
1433153_at	0.96309	RIKEN cDNA 9030201C23	gene	9030201C23Rik
1433154_at	0.96790	RIKEN cDNA B230204H03	gene	B230204H03Rik
1433155_at	1.05042	RIKEN cDNA B230204H03	gene	B230204H03Rik
1433156_at	0.86427	Kv channel-interacting protein 1		Kcnip1
1433157_at	1.03398	RIKEN cDNA 6330562C20	gene	6330562C20Rik
1433158_at	1.00025	kinesin family member 13B		Kif13b
1433159_at	1.05823	kinesin family member 13B		Kif13b
1433160_at	0.88679	RIKEN cDNA 1700008N17	gene	1700008N17Rik
1433161_at	0.91233	RIKEN cDNA 1700008N17	gene	1700008N17Rik
1433162_at	1.04013	RIKEN cDNA 6230414M07	gene	6230414M07Rik
1433163_at	0.91612	RIKEN cDNA 6230414M07	gene	6230414M07Rik
1433164_at	1.01459	RIKEN cDNA 4930570E01	gene	4930570E01Rik
1433165_at	1.04758	RIKEN cDNA 4930570E01	gene	4930570E01Rik
1433166_at	1.02738	RIKEN cDNA 4930485E13	gene	4930485E13Rik
1433167_at	1.00550	RIKEN cDNA 4930444E06	gene	4930444E06Rik
1433168_x_at	1.02194	RIKEN cDNA 5830456J23	gene	5830456J23Rik
1433169_at	0.87093	RIKEN cDNA 5830456J23	gene	5830456J23Rik
1433170_at	1.17670	RIKEN cDNA 2900034C19	gene	2900034C19Rik
1433171_at	0.95993	RIKEN cDNA 3110049I03	gene	3110049I03Rik
1433172_at	0.99922	RIKEN cDNA 3110049I03	gene	3110049I03Rik
1433173_at	0.88748	RIKEN cDNA 5430440L12	gene	5430440L12Rik
1433174_a_at	1.20709	RIKEN cDNA 5430440L12	gene	5430440L12Rik
1433175_at	0.97157	---	---	
1433176_at	0.87658	RIKEN cDNA 4930563F15	gene	4930563F15Rik
1433177_at	1.59811	RIKEN cDNA 5830411G16	gene	5830411G16Rik
1433178_at	1.01531	RIKEN cDNA 4933427C19	gene	4933427C19Rik
1433179_at	0.97022	MORN repeat containing 1		Morn1
1433180_at	1.25839	MORN repeat containing 1		Morn1
1433181_at	1.03747	RIKEN cDNA 4930431N21	gene	4930431N21Rik
1433182_at	0.80420	RIKEN cDNA 4930431N21	gene	4930431N21Rik

1433183_at	0.96571	RIKEN cDNA 6720477C19	gene	6720477C19Rik
1433184_at	1.08658	RIKEN cDNA 6720477C19	gene	6720477C19Rik
1433185_at	1.06424	RIKEN cDNA 2610019A05	gene	2610019A05Rik
1433186_at	1.21133	RIKEN cDNA 2610019A05	gene	2610019A05Rik
1433187_at	1.02980	RIKEN cDNA B230112I24	gene	B230112I24Rik
1433188_at	1.00939	RIKEN cDNA B230112I24	gene	B230112I24Rik
1433189_at	0.89362	RIKEN cDNA 4933433N18	gene	4933433N18Rik
1433190_at	0.98172	RIKEN cDNA 4933433N18	gene	4933433N18Rik
1433191_at	1.10328	RIKEN cDNA A430105P17	gene	A430105P17Rik
1433192_at	1.06812	RIKEN cDNA A430105P17	gene	A430105P17Rik
1433193_at	1.03511	RIKEN cDNA 4930433M22	gene	4930433M22Rik
1433194_at	0.96835	RIKEN cDNA 4930433M22	gene	4930433M22Rik
1433195_at	0.88670	RIKEN cDNA 9530006O14	gene	9530006O14Rik
1433196_at	0.95450	RIKEN cDNA 9530006O14	gene	9530006O14Rik
1433197_at	1.08240	RIKEN cDNA 4930445B03	gene	4930445B03Rik
1433198_at	1.02871	RIKEN cDNA 4930445B03	gene	4930445B03Rik
1433199_at	0.91278	RIKEN cDNA B230114A03	gene	B230114A03Rik
1433200_at	1.00366	RIKEN cDNA B230114A03	gene	B230114A03Rik
1433201_at	1.10219	RIKEN cDNA 2310079F09	gene	2310079F09Rik
1433202_at	0.87778	RIKEN cDNA 6030400A10	gene	6030400A10Rik
1433203_at	1.12281	RIKEN cDNA 6030400A10	gene	6030400A10Rik
1433204_at	0.94737	RIKEN cDNA 2810436B12	gene	2810436B12Rik
1433205_at	1.46773	RIKEN cDNA 2810436B12	gene	2810436B12Rik
1433206_at	1.01116	RIKEN cDNA 5033430J17	gene	5033430J17Rik
1433207_at	0.98917	RIKEN cDNA 5033430J17	gene	5033430J17Rik
1433208_at	0.91419	RIKEN cDNA 2210017G18	gene	2210017G18Rik
1433209_at	0.88377	RIKEN cDNA 2210017G18	gene	2210017G18Rik
1433210_at	0.95841	repetin Rptn		
1433211_at	1.12531	repetin Rptn		
1433212_at	0.96352	RIKEN cDNA 2700008E08	gene	2700008E08Rik
1433213_at	1.00156	RIKEN cDNA 2700008E08	gene	2700008E08Rik
1433214_x_at	1.01542	RIKEN cDNA 2700008E08	gene	2700008E08Rik
1433215_at	0.92119	---	---	
1433216_at	0.88419	RIKEN cDNA 9430024C03	gene	9430024C03Rik
1433217_at	1.16925	RIKEN cDNA 4930566N20	gene	4930566N20Rik
1433218_at	0.90455	RIKEN cDNA 4930566N20	gene	4930566N20Rik
1433219_at	0.95438	RIKEN cDNA 6720482G16	gene	6720482G16Rik
1433220_at	0.97494	RIKEN cDNA 6720482G16	gene	6720482G16Rik
1433221_at	0.89927	RIKEN cDNA 2610311E24	gene	2610311E24Rik
1433222_at	1.05570	RIKEN cDNA 2610311E24	gene	2610311E24Rik
1433223_at	0.94798	RIKEN cDNA 5830442K09	gene	5830442K09Rik
1433224_at	1.04040	RIKEN cDNA 5830442K09	gene	5830442K09Rik
1433225_at	1.04827	RIKEN cDNA 6330403N20	gene	6330403N20Rik

1433226_at	0.98924	RIKEN cDNA 6330403N20	gene	6330403N20Rik
1433227_at	1.05825	RIKEN cDNA 9430081G06	gene	9430081G06Rik
1433228_at	1.09973	RIKEN cDNA 9430081G06	gene	9430081G06Rik
1433229_at	1.09929	RIKEN cDNA 9430031J08	gene	9430031J08Rik
1433230_at	1.03406	RIKEN cDNA 5730415C11	gene	5730415C11Rik
1433231_at	1.05343	RIKEN cDNA 4933426I03	gene	4933426I03Rik
1433232_at	1.07712	RIKEN cDNA 4933426I03	gene	4933426I03Rik
1433233_at	0.95074	RIKEN cDNA 5430425E15	gene	5430425E15Rik
1433234_at	0.95832	RIKEN cDNA 4930424E08	gene	4930424E08Rik
1433235_at	0.99133	RIKEN cDNA C030011I16	gene	C030011I16Rik
1433236_at	1.00547	RIKEN cDNA 3830422I06	gene	3830422I06Rik
1433237_at	0.96580	RIKEN cDNA 3830422I06	gene	3830422I06Rik
1433238_at	1.19697	RIKEN cDNA 4930448E06	gene	4930448E06Rik
1433239_at	0.90355	RIKEN cDNA 4930448E06	gene	4930448E06Rik
1433240_at	1.00687	RIKEN cDNA 9430013L17	gene	9430013L17Rik
1433241_at	1.11732	RIKEN cDNA 9430013L17	gene	9430013L17Rik
1433242_at	1.08759	RIKEN cDNA 5830415B17	gene	5830415B17Rik
1433243_at	1.05042	RIKEN cDNA 2810421E14	gene	2810421E14Rik
1433244_at	1.05502	RIKEN cDNA 4930573G07	gene	4930573G07Rik
1433245_at	0.88917	RIKEN cDNA 6720475M21	gene	6720475M21Rik
1433246_at	0.97850	RIKEN cDNA 6720475M21	gene	6720475M21Rik
1433247_at	0.96022	RIKEN cDNA C530007A02	gene	C530007A02Rik
1433248_at	0.91836	RIKEN cDNA 2310075M01	gene	2310075M01Rik
1433249_at	1.00842	RIKEN cDNA 4931409D07	gene	4931409D07Rik
1433250_at	1.22263	RIKEN cDNA 4931409D07	gene	4931409D07Rik
1433251_at	1.02195	"heat shock protein 90kDa alpha (cytosolic), class A member 1 /// similar to heat shock protein 1, alpha" Hsp90aa1 /// LOC670148		
1433252_at	0.87734	RIKEN cDNA 4932703K07	gene	4932703K07Rik
1433253_at	0.74403	RIKEN cDNA 5033423K11	gene	5033423K11Rik
1433254_at	0.97941	RIKEN cDNA 5033423K11	gene	5033423K11Rik
1433255_at	1.22150	RIKEN cDNA 8430437B07	gene	8430437B07Rik
1433256_at	0.94583	RIKEN cDNA A930011E06	gene	A930011E06Rik
1433257_at	0.99653	RIKEN cDNA A330102K18	gene	A330102K18Rik
1433258_at	1.06456	RIKEN cDNA A330102K18	gene	A330102K18Rik
1433259_at	0.94813	RIKEN cDNA 4930518C04	gene	4930518C04Rik
1433260_at	1.10974	RIKEN cDNA 4930500E03	gene	4930500E03Rik
1433261_at	0.93070	RIKEN cDNA 9430052A13	gene	9430052A13Rik
1433262_at	0.93353	RIKEN cDNA 9430052A13	gene	9430052A13Rik
1433263_at	1.02562	---	---	
1433264_at	0.83681	---	---	
1433265_at	0.99926	RIKEN cDNA 2810416A17	gene	2810416A17Rik
1433266_at	0.98108	RIKEN cDNA 2810416A17	gene	2810416A17Rik
1433267_at	0.96130	"inhibitor of growth family, member 1" Ing1		

1433268_at	0.89316	RIKEN cDNA 4930553M12	gene	4930553M12Rik
1433269_at	0.97035	RIKEN cDNA 4930553M12	gene	4930553M12Rik
1433270_at	1.04600	RIKEN cDNA 9530004M16	gene	9530004M16Rik
1433271_at	1.06855	RIKEN cDNA 9530004M16	gene	9530004M16Rik
1433272_at	0.97414	RIKEN cDNA A430110A21	gene	A430110A21Rik
1433273_at	0.99850	RIKEN cDNA 2610200G18	gene	2610200G18Rik
1433274_at	0.93113	RIKEN cDNA 2610200G18	gene	2610200G18Rik
1433275_at	0.99475	RIKEN cDNA 4933401B01	gene	4933401B01Rik
1433276_at	0.98693	RIKEN cDNA 4930556A12	gene	4930556A12Rik
1433277_at	1.07178	RIKEN cDNA D730048A03	gene	D730048A03Rik
1433278_at	0.91026	RIKEN cDNA 5830440H09	gene	5830440H09Rik
1433279_at	1.16995	RIKEN cDNA 5830440H09	gene	5830440H09Rik
1433280_at	0.90804	RIKEN cDNA 4933417G07	gene	4933417G07Rik
1433281_at	1.07811	prune homolog (Drosophila)	Prune	
1433282_at	0.91390	RIKEN cDNA 4930529H12	gene	4930529H12Rik
1433283_s_at	0.89074	---	---	
1433284_at	0.99977	RIKEN cDNA 4921522E08	gene	4921522E08Rik
1433285_at	1.01064	RIKEN cDNA 9330179C17	gene	9330179C17Rik
1433286_at	0.98309	RIKEN cDNA 4833417J16	gene	4833417J16Rik
1433287_at	1.02430	RIKEN cDNA 4833417J16	gene	4833417J16Rik
1433288_at	0.95925	RIKEN cDNA 1520401O13	gene	1520401O13Rik
1433289_at	0.99182	RIKEN cDNA 9530078B04	gene	9530078B04Rik
1433290_at	1.13168	RIKEN cDNA C030045M22	gene	C030045M22Rik
1433291_at	1.00576	RIKEN cDNA C030045M22	gene	C030045M22Rik
1433292_at	0.91361	RIKEN cDNA 1500032O14	gene	1500032O14Rik
1433293_at	1.00452	RIKEN cDNA 1500032O14	gene	1500032O14Rik
1433294_at	0.93631	RIKEN cDNA 4930449I04	gene	4930449I04Rik
1433295_at	1.15046	RIKEN cDNA 4930449I04	gene	4930449I04Rik
1433296_at	0.99612	RIKEN cDNA 2310045N14	gene	2310045N14Rik
1433297_at	1.10074	RIKEN cDNA 1700039M15	gene	1700039M15Rik
1433298_at	1.07416	RIKEN cDNA 5830461L22	gene	5830461L22Rik
1433299_at	0.94660	RIKEN cDNA 5830461L22	gene	5830461L22Rik
1433300_at	0.94598	RIKEN cDNA 2900005I04	gene	2900005I04Rik
1433301_at	0.92509	RIKEN cDNA 4930478M09	gene	4930478M09Rik
1433302_at	0.98413	cadherin 10	Cdh10	
1433303_at	0.85339	RIKEN cDNA 4930483C01	gene	4930483C01Rik
1433304_at	1.03909	RIKEN cDNA 4930483C01	gene	4930483C01Rik
1433305_at	0.96480	RIKEN cDNA 9530034D02	gene	9530034D02Rik
1433306_at	0.86486	RIKEN cDNA 4930456K20	gene	4930456K20Rik
1433307_at	0.89728	RIKEN cDNA 4930509E22	gene	4930509E22Rik
1433308_at	1.00591	RIKEN cDNA 9330159N22	gene	9330159N22Rik
1433309_at	0.82629	RIKEN cDNA 4930472F24	gene	4930472F24Rik
1433310_at	0.84898	RIKEN cDNA D530031A16	gene	D530031A16Rik

1433311_at	1.01709	RIKEN cDNA D530031A16	gene	D530031A16Rik
1433312_at	1.21720	RIKEN cDNA 2700057C20	gene	2700057C20Rik
1433313_at	0.94912	RIKEN cDNA 6720454L07	gene	6720454L07Rik
1433314_at	0.95287	RIKEN cDNA 4930486A15	gene	4930486A15Rik
1433315_at	0.94368	RIKEN cDNA 4930486A15	gene	4930486A15Rik
1433316_at	0.95470	RIKEN cDNA 4930477O15	gene	4930477O15Rik
1433317_at	0.96362	RIKEN cDNA 4930477O15	gene	4930477O15Rik
1433318_at	0.94681	RIKEN cDNA 5430416B10	gene	5430416B10Rik
1433319_at	0.97758	RIKEN cDNA 5430437A18	gene	5430437A18Rik
1433320_at	1.04679	RIKEN cDNA 4930519N06	gene	4930519N06Rik
1433321_at	0.92217	RIKEN cDNA 9430087B13	gene	9430087B13Rik
1433322_at	1.00670	RIKEN cDNA 4930529F21	gene	4930529F21Rik
1433323_at	1.10146	RIKEN cDNA 4930529F21	gene	4930529F21Rik
1433324_at	1.07619	RIKEN cDNA 9330154F10	gene	9330154F10Rik
1433325_at	0.96385	RIKEN cDNA 9330154F10	gene	9330154F10Rik
1433326_at	1.03562	RIKEN cDNA 4930488N24	gene	4930488N24Rik
1433327_at	0.97073	---	---	
1433328_at	0.85465	RIKEN cDNA 1700095A21	gene	1700095A21Rik
1433329_at	0.96151	RIKEN cDNA 4930573C08	gene	4930573C08Rik
1433330_at	0.93005	RIKEN cDNA 4933401J01	gene	4933401J01Rik
1433331_at	0.95776	RIKEN cDNA C030046M01	gene	C030046M01Rik
1433332_at	1.15498	RIKEN cDNA C030004M13	gene	C030004M13Rik
1433333_at	1.11977	RIKEN cDNA C030004M13	gene	C030004M13Rik
1433334_at	0.94768	RIKEN cDNA 1700024P12	gene	1700024P12Rik
1433335_at	1.06684	RIKEN cDNA 4930405G09	gene	4930405G09Rik
1433336_at	1.06183	RIKEN cDNA 4930405G09	gene	4930405G09Rik
1433337_at	0.97699	RIKEN cDNA 6720460K10	gene	6720460K10Rik
1433338_at	1.74395	RIKEN cDNA 6720460K10	gene	6720460K10Rik
1433339_at	1.06666	RIKEN cDNA 6030460B20	gene	6030460B20Rik
1433340_at	0.96505	RIKEN cDNA 4930480K02	gene	4930480K02Rik
1433341_at	0.99123	RIKEN cDNA 5730416F02	gene	5730416F02Rik
1433342_at	1.00858	RIKEN cDNA 5730416F02	gene	5730416F02Rik
1433343_at	0.82494	RIKEN cDNA 4930448K20	gene	4930448K20Rik
1433344_at	0.88810	RIKEN cDNA 4930448K20	gene	4930448K20Rik
1433345_s_at	0.94755	RIKEN cDNA 4930448K20	gene	4930448K20Rik
1433346_at	0.98239	RIKEN cDNA 5830432F11	gene	5830432F11Rik
1433347_at	1.09405	RIKEN cDNA A930036A04	gene	A930036A04Rik
1433348_at	0.96667	RIKEN cDNA 2810414N06	gene	2810414N06Rik
1433349_at	0.92308	RIKEN cDNA 4930553P13	gene	4930553P13Rik
1433350_at	0.90430	RIKEN cDNA 4930553P13	gene	4930553P13Rik
1433351_at	1.04187	RIKEN cDNA 4933412L11	gene	4933412L11Rik
1433352_at	0.92021	RIKEN cDNA C430049A07	gene	C430049A07Rik
1433353_at	0.98281	RIKEN cDNA D530015H24	gene	D530015H24Rik

1433354_at	1.05134 RIKEN cDNA D530015H24 gene	D530015H24Rik	
1433355_at	1.04415 RIKEN cDNA 2610021J01 gene	2610021J01Rik	
1433356_at	0.97712 RIKEN cDNA 2610021J01 gene	2610021J01Rik	
1433357_at	0.97225 RIKEN cDNA 5730422E09 gene	5730422E09Rik	
1433358_at	0.85369 RIKEN cDNA A230102O21 gene	A230102O21Rik	
1433359_at	0.98727 --- ---		
1433360_at	0.90723 RIKEN cDNA 1700023B13 gene	1700023B13Rik	
1433361_at	1.06418 RIKEN cDNA 4930527F18 gene	4930527F18Rik	
1433362_at	0.95631 RIKEN cDNA 4930527F18 gene	4930527F18Rik	
1433363_at	1.00786 RIKEN cDNA A930036I15 gene	A930036I15Rik	
1433364_at	0.89575 RIKEN cDNA A930036I15 gene	A930036I15Rik	
1433365_at	0.95283 RIKEN cDNA 4930563N14 gene	4930563N14Rik	
1433366_at	1.02084 RIKEN cDNA 6030458A19 gene	6030458A19Rik	
1433367_at	1.22175 RIKEN cDNA 8430401P03 gene	8430401P03Rik	
1433368_x_at	1.04262 RIKEN cDNA 8430401P03 gene	8430401P03Rik	
1433369_at	1.09589 RIKEN cDNA 8430401P03 gene	8430401P03Rik	
1433370_at	1.13775 RIKEN cDNA 4930474A20 gene	4930474A20Rik	
1433371_at	1.02825 RIKEN cDNA 5430427N15 gene	5430427N15Rik	
1433372_at	1.10647 RIKEN cDNA 5330425B07 gene	5330425B07Rik	
1433373_at	1.12969 suppressor of Ty 7 (<i>S. cerevisiae</i>)-like Supt7l		
1433374_at	1.07586 suppressor of Ty 7 (<i>S. cerevisiae</i>)-like Supt7l		
1433375_at	0.98027 RIKEN cDNA D530037P16 gene	D530037P16Rik	
1433376_at	1.10828 RIKEN cDNA 5830445D09 gene	5830445D09Rik	
1433377_at	1.04924 RIKEN cDNA 5830445D09 gene	5830445D09Rik	
1433378_at	1.01046 RIKEN cDNA 9430019H13 gene	9430019H13Rik	
1433379_at	0.89250 RIKEN cDNA 9430019H13 gene	9430019H13Rik	
1433380_at	0.96351 RIKEN cDNA 4921509A06 gene	4921509A06Rik	
1433381_at	1.03733 RIKEN cDNA 4921509A06 gene	4921509A06Rik	
1433382_at	0.97324 RIKEN cDNA 1110065P19 gene	1110065P19Rik	
1433383_at	1.11829 RIKEN cDNA 1500002K03 gene	1500002K03Rik	
1433384_at	1.04740 RIKEN cDNA 9330199C07 gene	9330199C07Rik	
1433385_at	0.96857 RIKEN cDNA 4933428C20 gene	4933428C20Rik	
1433386_at	1.03182 RIKEN cDNA 4933428C20 gene	4933428C20Rik	
1433387_at	1.11050 RIKEN cDNA 2900022M07 gene	2900022M07Rik	
1433388_at	0.89253 RIKEN cDNA 2900022M07 gene	2900022M07Rik	
1433389_at	0.98514 "cytochrome P450, family 4, subfamily f, polypeptide 18"		Cyp4f18
1433390_at	0.89069 RIKEN cDNA A230105L22 gene	A230105L22Rik	
1433391_at	1.02967 --- ---		
1433392_at	0.98618 --- ---		
1433393_at	0.92606 RIKEN cDNA 3300002I10 gene	3300002I10Rik	
1433394_at	0.92960 microtubule associated serine/threonine kinase family member 4		Mast4
1433395_at	0.90470 RIKEN cDNA 4921504P13 gene	4921504P13Rik	
1433396_at	1.02066 RIKEN cDNA 4921504P13 gene	4921504P13Rik	

1433397_at 1.03353 RIKEN cDNA 9430012M22 gene 9430012M22Rik
1433398_at 1.87318 "FYVE, RhoGEF and PH domain containing 3" Fgd3
1433399_at 1.10371 RIKEN cDNA 2900078I11 gene 2900078I11Rik
1433400_at 0.93748 RIKEN cDNA 5033405D04 gene 5033405D04Rik
1433401_at 1.00260 RIKEN cDNA 5033405D04 gene 5033405D04Rik
1433402_at 1.04321 RIKEN cDNA 4930451E06 gene 4930451E06Rik
1433403_at 1.01702 RIKEN cDNA 2810489J07 gene 2810489J07Rik
1433404_at 1.07690 RIKEN cDNA 4930423O20 gene 4930423O20Rik
1433405_at 1.01567 --- ---
1433406_at 1.00510 hypothetical protein LOC668408 /// hypothetical protein LOC670057
LOC668408 /// LOC670057
1433407_at 0.91352 minichromosome maintenance deficient 10 (S. cerevisiae) Mcm10
1433408_a_at 1.08068 minichromosome maintenance deficient 10 (S. cerevisiae) Mcm10
1433409_at 1.18826 RIKEN cDNA C030014O09 gene C030014O09Rik
1433410_at 0.89685 RIKEN cDNA C030014O09 gene C030014O09Rik
1433411_at 0.84400 --- ---
1433412_at 1.19093 RIKEN cDNA 5730446C15 gene 5730446C15Rik
1433413_at 1.00328 neurexin I Nrnx1
1433414_at 1.05193 RIKEN cDNA 9530086P17 gene 9530086P17Rik
1433415_at 1.12780 RIKEN cDNA 5430432H19 gene 5430432H19Rik
1433416_at 0.96152 RIKEN cDNA 5430432H19 gene 5430432H19Rik
1433417_at 0.79743 RIKEN cDNA 8030497O21 gene 8030497O21Rik
1433418_at 1.10419 --- ---
1433419_at 0.98997 RIKEN cDNA 4930405A07 gene 4930405A07Rik
1433420_at 0.83235 RIKEN cDNA 4930527J03 gene 4930527J03Rik
1433421_at 1.06684 RIKEN cDNA 4930527J03 gene 4930527J03Rik
1433422_at 0.89659 hypothetical protein LOC626748 LOC626748
1433423_at 0.86562 RIKEN cDNA 2210008N01 gene 2210008N01Rik
1433424_at 1.05825 --- ---
1433425_at 0.91441 RIKEN cDNA 1700101I19 gene 1700101I19Rik
1433426_at 0.94024 RIKEN cDNA 1700111A04 gene 1700111A04Rik
1433427_at 1.00695 RIKEN cDNA 1700111A04 gene 1700111A04Rik
1433428_x_at 0.95480 "transglutaminase 2, C polypeptide" Tgm2
1433429_at 0.94510 "phosphatidylinositol glycan anchor biosynthesis, class S" Pigs
1433430_s_at 0.99779 "CDC23 (cell division cycle 23, yeast, homolog)" Cdc23
1433431_at 0.91749 pancreatic lipase Pnlip
1433432_x_at 1.03213 ribosomal protein S12 /// similar to ribosomal protein S12 /// similar to
ribosomal protein S12 Rps12 /// LOC670751 /// LOC672959
1433433_at 0.92021 MYST histone acetyltransferase 2 Myst2
1433434_at 1.09140 expressed sequence AW551984 AW551984
1433435_at 1.10349 expressed sequence AW551984 AW551984
1433436_s_at 0.95182 "thiamine triphosphatase /// zinc finger homeobox 2, antisense" Thtpa ///
Zfhx2as

1433437_at	0.98484	dedicator of cytokinesis 8	Dock8
1433438_x_at	0.96281	melanoma antigen	Mela
1433439_at	1.10121	copine I	Cpne1
1433440_x_at	0.95812	ubiquitin-like 1 (sentrin) activating enzyme E1B	Uble1b
1433441_at	0.96411	F-box and leucine-rich repeat protein 5	Fbxl5
1433442_at	1.01222	Kelch-like 9 (Drosophila)	Klhl9
1433443_a_at	1.40268	3-hydroxy-3-methylglutaryl-Coenzyme A synthase 1	Hmgcs1
1433444_at	1.43825	3-hydroxy-3-methylglutaryl-Coenzyme A synthase 1	Hmgcs1
1433445_x_at	1.42824	3-hydroxy-3-methylglutaryl-Coenzyme A synthase 1	Hmgcs1
1433446_at	1.63936	3-hydroxy-3-methylglutaryl-Coenzyme A synthase 1	Hmgcs1
1433447_x_at	1.03525	chaperonin subunit 4 (delta)	Cct4
1433448_at	0.98960	RIKEN cDNA B430110G05 gene	B430110G05Rik
1433449_at	0.97314	RIKEN cDNA B930037P14 gene	B930037P14Rik
1433450_at	0.88289	"cyclin-dependent kinase 5, regulatory subunit (p35) 1"	Cdk5r1
1433451_at	0.90218	"cyclin-dependent kinase 5, regulatory subunit (p35) 1"	Cdk5r1
1433452_at	0.88614	RIKEN cDNA B630019K06 gene	B630019K06Rik
1433453_a_at	0.92209	ankyrin repeat and BTB (POZ) domain containing 2	Abtb2
1433454_at	0.99085	ankyrin repeat and BTB (POZ) domain containing 2	Abtb2
1433455_at	0.97053	SH2B adaptor protein 3	Sh2b3
1433456_at	1.02136	expressed sequence AU022751	AU022751
1433457_s_at	1.01182	G-rich RNA sequence binding factor 1	Grsf1
1433458_at	0.89602	RIKEN cDNA E330021D16 gene	E330021D16Rik
1433459_x_at	0.93718	"protease, serine, 2"	Prss2
1433460_at	1.31241	tetratricopeptide repeat domain 7B	Ttc7b
1433461_at	0.98207	"splicing factor 3b, subunit 2"	Sf3b2
1433462_a_at	0.97420	phosphatidylinositol 4-kinase type 2 alpha	Pi4k2a
1433463_at	0.94309	RIKEN cDNA 0610010K06 gene	0610010K06Rik
1433464_at	0.89032	importin 13	Ipo13
1433465_a_at	1.22129	expressed sequence AI467606	AI467606
1433466_at	0.97101	expressed sequence AI467606	AI467606
1433467_at	1.07345	"solute carrier family 7 (cationic amino acid transporter, y+ system), member 6"	Slc7a6
1433468_at	0.89436	RIKEN cDNA 6430527G18 gene	6430527G18Rik
1433469_at	1.03102	"leucine rich repeat protein 2, neuronal"	Lrrn2
1433470_a_at	1.06140	"inner membrane protein, mitochondrial"	Immt
1433471_at	1.12048	"transcription factor 7, T-cell specific"	Tcf7
1433472_x_at	0.91211	ribosomal protein L38 /// similar to ribosomal protein L38 /// similar to ribosomal protein L38 /// similar to ribosomal protein L38 /// similar to ribosomal protein L38	Rpl38 /// LOC625646 /// LOC638514 /// LOC664868 /// LOC672023 /// LOC674951
1433473_x_at	0.88267	TM2 domain containing 2	Tm2d2
1433474_at	0.85543	EGF-like repeats and discoidin I-like domains 3	Edil3
1433475_a_at	0.91804	expressed sequence C78339	C78339

1433476_at	0.89665	expressed sequence C78339	C78339
1433477_at	0.97302	active BCR-related gene Abr	
1433478_at	1.17307	"presenilin associated, rhomboid-like" Parl	
1433479_at	0.94864	RIKEN cDNA 5730410I19 gene	5730410I19Rik
1433480_at	1.11114	RIKEN cDNA 2900010J23 gene	2900010J23Rik
1433481_at	0.79824	FK506 binding protein 14	Fkbp14
1433482_a_at	0.97508	far upstream element (FUSE) binding protein 1	Fubp1
1433483_s_at	0.90649	expressed sequence C86187	C86187
1433484_at	1.07294	"DNA segment, Chr 6, ERATO Doi 527, expressed"	D6Ertd527e
1433485_x_at	0.56181	G protein-coupled receptor 56	Gpr56
1433486_at	1.10031	chloride channel 3	Clcn3
1433487_at	0.99697	chloride channel 3	Clcn3
1433488_x_at	0.95396	glucosamine (N-acetyl)-6-sulfatase	Gns
1433489_s_at	0.63720	fibroblast growth factor receptor 2	Fgfr2
1433490_s_at	1.02098	erythrocyte protein band 4.1-like 2	Epb4.112
1433491_at	1.03479	erythrocyte protein band 4.1-like 2	Epb4.112
1433492_at	0.87475	erythrocyte protein band 4.1-like 2	Epb4.112
1433493_at	0.84677	RIKEN cDNA 5730472N09 gene	5730472N09Rik
1433494_at	0.87166	downstream of Stk11	Dos
1433495_at	0.93360	glycosyltransferase 25 domain containing 1	Glt25d1
1433496_at	0.93715	glycosyltransferase 25 domain containing 1	Glt25d1
1433497_at	1.00164	aquarius	Aqr
1433498_at	0.87892	RIKEN cDNA 2010005J08 gene	2010005J08Rik
1433499_at	0.83098	RIKEN cDNA 2010005J08 gene	2010005J08Rik
1433500_at	0.86759	DENN/MADD domain containing 2A	Dennd2a
1433501_at	0.87146	cathepsin O	Ctso
1433502_s_at	1.15801	"TSR1, 20S rRNA accumulation, homolog (yeast)"	Tsr1
1433503_at	0.81796	"zinc binding alcohol dehydrogenase, domain containing 1"	Zadh1
1433504_at	0.78168	brain glycogen phosphorylase	Pygb
1433505_a_at	0.89919	leucine rich repeat containing 8D	Lrrc8d
1433506_at	1.01227	leucine rich repeat containing 8D	Lrrc8d
1433507_a_at	0.91222	high mobility group nucleosomal binding domain 2	Hmgn2
1433508_at	1.03557	Kruppel-like factor 6	Klf6
1433509_s_at	1.03005	receptor accessory protein 1	Reep1
1433510_x_at	1.07470	similar to ribosomal protein L36	LOC622534
1433511_at	1.05010	RIKEN cDNA 6330549H03 gene	6330549H03Rik
1433512_at	1.06166	Friend leukemia integration 1	Fli1
1433513_x_at	0.98847	"NADH dehydrogenase (ubiquinone) 1 alpha subcomplex, 12"	Ndufa12
1433514_at	1.04994	ethanolamine kinase 1	Etnk1
1433515_s_at	1.02554	ethanolamine kinase 1	Etnk1
1433516_a_at	0.88251	myeloma overexpressed 2	Myeov2
1433517_at	0.97538	myeloma overexpressed 2	Myeov2
1433518_at	0.89360	leucine carboxyl methyltransferase 2	Lcmt2

1433519_at	1.07869	nuclear casein kinase and cyclin-dependent kinase substrate 1	Nucks1
1433520_at	0.93147	SREBP cleavage activating protein	Scap
1433521_at	0.98207	ankyrin repeat domain 13c	Ankrd13c
1433522_at	0.90656	protein serine kinase H1	Pskh1
1433523_at	1.09365	RIKEN cDNA D930005D10 gene	D930005D10Rik
1433524_at	0.84766	expressed sequence AU042671	AU042671
1433525_at	0.95937	endothelin receptor type A	Ednra
1433526_at	0.88712	kelch-like 8 (Drosophila)	Klhl8
1433527_at	0.95250	iron responsive element binding protein 2	Ireb2
1433528_at	1.05196	"general transcription factor II A, 2"	Gtf2a2
1433529_at	0.87920	RIKEN cDNA E430002G05 gene	E430002G05Rik
1433530_at	1.63717	RIKEN cDNA 2210411K19 gene	2210411K19Rik
1433531_at	1.34927	acyl-CoA synthetase long-chain family member 4	Acsl4
1433532_a_at	1.12795	myelin basic protein	Mbp
1433533_x_at	0.94540	"protein kinase, AMP-activated, gamma 1 non-catalytic subunit"	Prkag1
1433534_a_at	1.03207	chaperonin subunit 2 (beta)	Cct2
1433535_x_at	0.98988	chaperonin subunit 2 (beta)	Cct2
1433536_at	0.98550	low density lipoprotein receptor-related protein 11	Lrp11
1433537_at	1.01996	RIKEN cDNA 4833408C14 gene	4833408C14Rik
1433538_at	0.86901	MARVEL (membrane-associating) domain containing 1	Marveld1
1433539_at	1.02321	COMM domain containing 3	Commd3
1433540_x_at	0.89817	"protein phosphatase 1, catalytic subunit, beta isoform"	Ppp1cb
1433541_a_at	1.02119	ubiquitin associated protein 2-like	Ubap2l
1433542_at	0.93022	inositol polyphosphate-5-phosphatase F	Inpp5f
1433543_at	1.11612	"anillin, actin binding protein (scraps homolog, Drosophila)"	Anln
1433544_at	0.95570	"amyotrophic lateral sclerosis 2 (juvenile) chromosome region, candidate 2 (human)"	Als2cr2
1433545_s_at	0.83378	"acyl-Coenzyme A dehydrogenase family, member 11"	Acad11
1433546_at	1.01480	glucosamine (N-acetyl)-6-sulfatase	Gns
1433547_s_at	1.12612	NudC domain containing 1	Nudcd1
1433548_at	0.99649	alpha globin regulatory element containing gene	Mare
1433549_x_at	0.94813	ribosomal protein S21	Rps21
1433550_at	0.99114	checkpoint with forkhead and ring finger domains	Chfr
1433551_at	0.98669	expressed sequence AI427515	AI427515
1433552_a_at	0.94316	polymerase (RNA) II (DNA directed) polypeptide B	Polr2b
1433553_at	0.83368	GTPase activating RANGAP domain-like 3	Garnl3
1433554_at	0.97088	expressed sequence AU022870	AU022870
1433555_at	1.05267	ELL associated factor 1	Eaf1
1433556_at	0.87294	"centaurin, alpha 1"	Centa1
1433557_at	1.08440	chromobox homolog 7	Cbx7
1433558_at	0.95257	disabled homolog 2 (Drosophila) interacting protein	Dab2ip
1433559_at	0.92640	"solute carrier family 45, member 4"	Slc45a4
1433560_at	0.89957	"solute carrier family 45, member 4"	Slc45a4

1433561_at 1.00083 "centaurin, beta 2" Centb2
1433562_s_at 0.98641 "ATP synthase, H⁺ transporting, mitochondrial F0 complex, subunit b, isoform 1" Atp5f1
1433563_s_at 0.95149 "Der1-like domain family, member 1" Der1l
1433564_at 0.96535 "diacylglycerol kinase, delta" Dgkd
1433565_at 1.04440 PRP38 pre-mRNA processing factor 38 (yeast) domain containing A Prpf38a
1433566_at 1.01405 "RAS-like, family 10, member B" Rasl10b
1433567_at 1.05454 guanine monophosphate synthetase Gmps
1433568_at 0.70061 PAP associated domain containing 4 Papd4
1433569_x_at 1.05871 "RAN, member RAS oncogene family /// similar to RAN, member RAS oncogene family /// similar to RAN, member RAS oncogene family" Ran /// LOC625094 /// LOC671251
1433570_s_at 0.87412 "MAK10 homolog, amino-acid N-acetyltransferase subunit, (S. cerevisiae)" Mak10
1433571_at 0.50088 serine incorporator 5 Serinc5
1433572_a_at 1.00237 cDNA sequence BC010304 BC010304
1433573_x_at 0.92018 "protease, serine, 2" Prss2
1433574_at 1.16650 cell division cycle 37 homolog (S. cerevisiae)-like 1 Cdc3711
1433575_at 0.75850 similar to Transcription factor SOX-4 LOC672274
1433576_at 0.99407 "methionine adenosyltransferase II, alpha" Mat2a
1433577_at 1.00011 RIKEN cDNA A730017C20 gene A730017C20Rik
1433578_at 1.03343 "solute carrier family 10 (sodium/bile acid cotransporter family), member 4" Slc10a4
1433579_at 1.02802 transmembrane protein 30B Tmem30b
1433580_at 1.02893 nucleoporin 54 Nup54
1433581_at 0.70937 RIKEN cDNA 1190002N15 gene 1190002N15Rik
1433582_at 0.70602 RIKEN cDNA 1190002N15 gene 1190002N15Rik
1433583_at 1.08749 zinc finger protein 365 /// similar to zinc finger protein 365 Zfp365 /// LOC674611
1433584_at 0.89018 "tubulin, alpha 1 /// similar to tubulin, alpha 1" Tuba1 /// LOC544863
1433585_at 1.09208 transportin 1 Tnpol
1433586_at 0.84664 "RGM domain family, member B" Rgmb
1433587_at 0.98893 "RGM domain family, member B" Rgmb
1433588_at 1.18868 "DNA segment, Chr 6, Wayne State University 116, expressed" D6Wsu116e
1433589_at 1.03598 "DNA segment, Chr 6, Wayne State University 116, expressed" D6Wsu116e
1433590_at 0.69677 hect domain and RLD 3 Herc3
1433591_at 1.06255 "protein phosphatase 3, regulatory subunit B, alpha isoform (calcineurin B, type I)" Ppp3r1
1433592_at 0.93184 calmodulin 1 Calm1
1433593_at 0.87252 yippee-like 5 (Drosophila) Ypel5
1433594_at 0.91735 COMM domain containing 2 Commd2
1433595_at 1.01662 --- ---
1433596_at 0.99269 "DnaJ (Hsp40) homolog, subfamily C, member 6" Dnajc6
1433597_at 1.02368 RIKEN cDNA 9430010O03 gene 9430010O03Rik

1433598_at 1.30120 RIKEN cDNA 9430010O03 gene 9430010O03Rik
1433599_at 1.18621 bromodomain adjacent to zinc finger domain 1A Baz1a
1433600_at 1.06334 "adrenergic receptor, alpha 2a" Adra2a
1433601_at 0.94542 "adrenergic receptor, alpha 2a" Adra2a
1433602_at 1.01308 "gamma-aminobutyric acid (GABA-A) receptor, subunit alpha 5" Gabra5
1433603_at 1.03220 NADH dehydrogenase (ubiquinone) Fe-S protein 6 Ndufs6
1433604_x_at 1.13529 "aldolase 1, A isoform" Aldoa
1433605_at 0.82290 inositol polyphosphate-5-phosphatase A Inpp5a
1433606_at 0.99433 decapping enzyme Dcp1a
1433607_at 1.01978 cerebellin 4 precursor protein Cbln4
1433608_at 0.94814 sec1 family domain containing 2 Scfd2
1433609_s_at 1.11806 surfet gene 2 Surf2
1433610_at 0.79752 expressed sequence AA986860 AA986860
1433611_s_at 1.07056 BUD31 homolog (yeast) Bud31
1433612_at 0.99357 "adaptor-related protein complex 2, sigma 1 subunit" Ap2s1
1433613_at 1.10255 --- ---
1433614_at 0.98042 sorting nexin family member 27 Snx27
1433615_at 0.69288 transmembrane protein 117 Tmem117
1433616_a_at 0.93803 RIKEN cDNA 2310028O11 gene 2310028O11Rik
1433617_s_at 1.05164 "UDP-Gal:betaGlcNAc beta 1,4-galactosyltransferase, polypeptide 5"
B4galt5
1433618_at 1.00604 RIKEN cDNA C330006A16 gene C330006A16Rik
1433619_at 0.84778 expressed sequence AI894139 AI894139
1433620_at 0.90392 WD repeat domain 41 Wdr41
1433621_at 0.83931 WD repeat domain 41 Wdr41
1433622_at 1.09226 gem (nuclear organelle) associated protein 4 Gemin4
1433623_at 0.93332 zinc finger protein 367 Zfp367
1433624_at 0.94590 RIKEN cDNA 5830434P21 gene 5830434P21Rik
1433625_at 1.02831 RIKEN cDNA 5830434P21 gene 5830434P21Rik
1433626_at 1.17431 phospholipid scramblase 4 Plscr4
1433627_at 1.02661 Sec23 interacting protein Sec23ip
1433628_at 0.95035 coenzyme Q10 homolog A (yeast) Coq10a
1433629_s_at 1.15438 expressed sequence AI316787 AI316787
1433630_at 0.94021 MAP6 domain containing 1 Map6d1
1433631_at 1.06111 eukaryotic translation initiation factor 5 Eif5
1433632_at 0.86498 interferon regulatory factor 2 binding protein 2 /// similar to interferon
regulatory factor 2 binding protein 2 Irf2bp2 /// LOC672960
1433633_at 0.93605 interferon regulatory factor 2 binding protein 2 /// similar to interferon
regulatory factor 2 binding protein 2 Irf2bp2 /// LOC672960
1433634_at 0.86796 interferon regulatory factor 2 binding protein 2 /// similar to interferon
regulatory factor 2 binding protein 2 Irf2bp2 /// LOC672960
1433635_at 1.04134 WD repeat domain 18 Wdr18
1433636_at 1.10220 "BMS1-like, ribosome assembly protein (yeast)" Bms11

1433637_at 0.76709 aarF domain containing kinase 2 Adck2
1433638_s_at 0.99933 homeo box D8 Hoxd8
1433639_at 1.02115 RIKEN cDNA 5730593F17 gene 5730593F17Rik
1433640_at 0.92767 Far upstream element (FUSE) binding protein 1 Fubp1
1433641_at 0.88183 MAD homolog 5 (Drosophila) Smad5
1433642_at 0.96002 ADP-ribosylation factor-like 15 Arl15
1433643_at 0.93557 "calcium channel, voltage-dependent, alpha2/delta subunit 1" Cacna2d1
1433644_at 1.04823 protein-O-mannosyltransferase 2 Pomt2
1433645_at 0.87216 RIKEN cDNA 2210409B22 gene 2210409B22Rik
1433646_at 1.00115 mitochondrial ribosomal protein S27 Mrps27
1433647_s_at 0.95317 Rho-related BTB domain containing 3 Rhobtb3
1433648_at 1.05184 sperm associated antigen 9 Spag9
1433649_at 0.82135 "amine oxidase, flavin containing 1" Aof1
1433650_at 1.04258 hydroxyacylglutathione hydrolase-like Hagh1
1433651_at 0.88706 WT1-interacting protein Wtip
1433652_at 0.99892 "immunoglobulin superfamily, member 1" Igsf1
1433653_at 0.94533 cDNA sequence BC029169 BC029169
1433654_at 0.96459 meningioma expressed antigen 5 (hyaluronidase) Mgea5
1433655_at 1.07011 ring finger protein 141 Rnf141
1433656_a_at 1.25772 guanine nucleotide binding protein-like 3 (nucleolar) Gnl3
1433657_at 0.89993 RIKEN cDNA A130092J06 gene A130092J06Rik
1433658_x_at 0.99920 poly(rC) binding protein 4 Pcbp4
1433659_at 0.93612 Transformation related protein 53 binding protein 1 Trp53bp1
1433660_at 1.01384 "DNA segment, Chr 2, ERATO Doi 435, expressed" D2Ertd435e
1433661_at 1.08905 cDNA sequence BC034204 BC034204
1433662_s_at 0.86256 tissue inhibitor of metalloproteinase 2 Timp2
1433663_s_at 1.13672 expressed sequence AU014645 AU014645
1433664_at 0.96393 ubiquitin-conjugating enzyme E2Q (putative) 2 Ube2q2
1433665_at 1.01202 vacuolar protein sorting 41 (yeast) Vps41
1433666_s_at 0.92542 vacuolar protein sorting 41 (yeast) Vps41
1433667_at 0.92744 "leucine-rich repeat LGI family, member 3" Lgi3
1433668_at 0.92814 proline-rich nuclear receptor coactivator 1 Pnrc1
1433669_at 1.04144 A kinase (PRKA) anchor protein 8 Akap8
1433670_at 0.90750 epithelial membrane protein 2 Emp2
1433671_at 1.03270 RIKEN cDNA A130022J15 gene A130022J15Rik
1433672_at 0.85317 RIKEN cDNA 4732479N06 gene 4732479N06Rik
1433673_at 0.85140 RIKEN cDNA E130309D14 gene E130309D14Rik
1433674_a_at 1.11418 "small nucleolar RNA, C/D box 22" Snord22
1433675_at 1.06156 "small nucleolar RNA, C/D box 22" Snord22
1433676_at 0.93732 WNK lysine deficient protein kinase 1 Wnk1
1433677_at 0.97091 "splicing factor, arginine/serine-rich 8" Sfrs8
1433678_at 0.86504 "phospholipase D family, member 4" Pld4
1433679_at 1.04907 far upstream element (FUSE) binding protein 3 Fubp3

1433680_x_at 1.06686 Cd27 binding protein (Hindu God of destruction) Siva
1433681_x_at 0.90609 calpain 3 Capn3
1433682_at 0.99240 Rho guanine nucleotide exchange factor (GEF) 17 Arhgef17
1433683_at 0.96232 RNA binding motif protein 35b Rbm35b
1433684_at 1.17676 chromatin modifying protein 6 Chmp6
1433685_a_at 0.93372 RIKEN cDNA C330011F01 gene /// RIKEN cDNA 6430706D22 gene
C330011F01Rik /// 6430706D22Rik
1433686_at 0.94619 calcineurin binding protein 1 Cabin1
1433687_at 0.81247 expressed sequence AI662250 AI662250
1433688_x_at 1.13182 ribosomal protein L14 Rpl14
1433689_s_at 1.04423 ribosomal protein S9 Rps9
1433690_at 0.91925 RIKEN cDNA 2210016L21 gene 2210016L21Rik
1433691_at 0.96062 "protein phosphatase 1, regulatory (inhibitor) subunit 3C" Ppp1r3c
1433692_at 1.12674 N-acetyltransferase 10 Nat10
1433693_x_at 0.88769 vesicle-associated membrane protein 3 Vamp3
1433694_at 0.79548 "phosphodiesterase 3B, cGMP-inhibited" Pde3b
1433695_at 0.98192 RIKEN cDNA 1500041B16 gene 1500041B16Rik
1433696_at 0.92189 "DNA segment, Chr 17, ERATO Doi 441, expressed" D17Ertd441e
1433697_at 1.04886 expressed sequence AV312086 AV312086
1433698_a_at 1.08553 thioredoxin-like 4 Txnl4
1433699_at 1.48027 "tumor necrosis factor, alpha-induced protein 3" Tnfaip3
1433700_at 0.99953 RIKEN cDNA 4933433P14 gene 4933433P14Rik
1433701_at 0.97446 metallophosphoesterase domain containing 1 Mpped1
1433702_at 0.91917 "DNA segment, Chr 19, Wayne State University 12, expressed" D19Wsu12e
1433703_s_at 1.02302 bromo adjacent homology domain containing 1 Bahd1
1433704_s_at 0.87470 translocation protein 1 Tloc1
1433705_at 0.82652 zinc finger protein 213 Zfp213
1433706_a_at 0.96588 protein tyrosine phosphatase-like A domain containing 1 Ptplad1
1433707_at 1.00014 "gamma-aminobutyric acid (GABA-A) receptor, subunit alpha 4" Gabra4
1433708_at 1.01024 signal recognition particle 68 Srp68
1433709_at 0.96471 calcium activated nucleotidase 1 Cant1
1433710_at 0.93546 enhancer of mRNA decapping 3 homolog (S. cerevisiae) Edc3
1433711_s_at 0.86198 sestrin 1 Sesn1
1433712_at 0.96693 expressed sequence AW555464 AW555464
1433713_at 0.97418 GCN1 general control of amino-acid synthesis 1-like 1 (yeast) Gcn111
1433714_at 0.95284 "sulfotransferase family 4A, member 1" Sult4a1
1433715_at 0.93958 copine VII Cpne7
1433716_x_at 1.03696 glial cell line derived neurotrophic factor family receptor alpha 2 Gfra2
1433717_at 0.96503 "DNA segment, Chr 19, Wayne State University 162, expressed" D19Wsu162e
1433718_a_at 0.95100 chromobox homolog 1 (Drosophila HP1 beta) Cbx1
1433719_at 0.92723 "solute carrier family 9 (sodium/hydrogen exchanger), isoform 9" Slc9a9
1433720_s_at 0.80132 Nur77 downstream gene 2 Ndg2
1433721_x_at 0.96180 ribosomal protein S21 Rps21

1433722_at 1.01056 A kinase (PRKA) anchor protein 13 Akap13
1433723_s_at 0.89779 small EDRK-rich factor 2 Serf2
1433724_at 0.96871 "DNA segment, Chr 15, ERATO Doi 621, expressed" D15Ertd621e
1433725_at 0.89631 "activin A receptor, type 1B" Acvr1b
1433726_at 0.92888 --- ---
1433727_at 0.96590 cDNA sequence BC038479 BC038479
1433728_at 0.98458 cDNA sequence BC038479 BC038479
1433729_x_at 0.99104 peptidase (mitochondrial processing) beta Pmpcb
1433730_at 0.82762 ELMO domain containing 2 Elmod2
1433731_at 0.96832 insulin-like growth factor 2 mRNA binding protein 3 Igf2bp3
1433732_x_at 1.03943 insulin-like growth factor 2 mRNA binding protein 3 Igf2bp3
1433733_a_at 1.00086 cryptochrome 1 (photolyase-like) Cry1
1433734_at 0.87111 "solute carrier family 13 (sodium/sulfate symporters), member 4" Slc13a4
1433735_a_at 0.71082 transmembrane protein 64 Tmem64
1433736_at 0.96699 host cell factor C1 Hcfc1
1433737_at 1.02250 similar to U2AF homology motif (UHM) kinase 1 LOC677213
1433738_at 1.08795 RIKEN cDNA 5730445M16 gene 5730445M16Rik
1433739_at 1.04094 nucleolar protein 10 Nol10
1433740_at 0.91870 RIKEN cDNA 2610301K12 gene 2610301K12Rik
1433741_at 0.90844 CD38 antigen Cd38
1433742_at 0.60065 ankyrin repeat domain 15 Ankrd15
1433743_at 0.93985 dachshund 1 (Drosophila) Dach1
1433744_at 1.17903 expressed sequence AI841794 AI841794
1433745_at 0.99580 triple functional domain (PTPRF interacting) Trio
1433746_at 0.98351 WD repeat domain 3 Wdr3
1433747_at 1.05755 leucyl/cystinyl aminopeptidase Lnpep
1433748_at 0.99146 "zinc finger, DHHC domain containing 18" Zdhhc18
1433749_at 1.06778 "guanine nucleotide binding protein, alpha 13" Gna13
1433750_at 1.03718 "solute carrier family 31, member 1" Slc31a1
1433751_at 0.98587 "solute carrier family 39 (zinc transporter), member 10" Slc39a10
1433752_s_at 0.89705 RIKEN cDNA D030016E14 gene D030016E14Rik
1433753_x_at 0.93495 Era (G-protein)-like 1 (E. coli) Era1
1433754_at 0.97459 muscleblind-like 2 Mbnl2
1433755_at 1.19158 mesoderm induction early response 1 homolog (Xenopus laevis) Mier1
1433756_at 1.12097 S100P binding protein S100pbp
1433757_a_at 1.12804 nischarin Nisch
1433758_at 1.23831 nischarin Nisch
1433759_at 1.09928 dpy-19-like 1 (C. elegans) Dpy191
1433760_a_at 1.06006 rhomboid domain containing 3 Rhbdd3
1433761_at 0.85211 phosphodiesterase 4D interacting protein (myomegalin) Pde4dip
1433762_at 0.83800 RIKEN cDNA C630043F03 gene C630043F03Rik
1433763_at 0.86691 ectonucleoside triphosphate diphosphohydrolase 5 Entpd5
1433764_at 1.04028 "C-type lectin domain family, member 1 /// similar to CD69 antigen (p60, early

T-cell activation antigen)" Clec2l /// LOC665180

1433765_at 0.96846 ubiquitin-conjugating enzyme E2o Ube2o

1433766_at 1.04685 RIKEN cDNA C330023M02 gene C330023M02Rik

1433767_at 1.12325 RIKEN cDNA 1110018G07 gene 1110018G07Rik

1433768_at 0.84505 "palladin, cytoskeletal associated protein" Palld

1433769_at 1.43274 ALS2 C-terminal like Als2cl

1433770_at 0.95486 dihydropyrimidinase-like 2 Dpysl2

1433771_at 0.90635 RIKEN cDNA 5730446C15 gene 5730446C15Rik

1433772_at 0.96663 "stress 70 protein chaperone, microsomal-associated, human homolog" Stch

1433773_at 0.86205 ribonucleotide reductase M2 B (TP53 inducible) Rrm2b

1433774_x_at 1.02795 component of oligomeric golgi complex 1 Cog1

1433775_at 1.19076 expressed sequence C77080 C77080

1433776_at 1.02084 lipoma HMGIC fusion partner Lhfp

1433777_at 0.97909 l(3)mbt-like 2 (Drosophila) L3mbtl2

1433778_at 0.93691 "tankyrase, TRF1-interacting ankyrin-related ADP-ribose polymerase" Tnks

1433779_at 1.00685 cancer susceptibility candidate 4 Casc4

1433780_at 1.00570 ubinuclein 1 Ubn1

1433781_a_at 1.04150 claudin 12 Cldn12

1433782_at 1.03817 claudin 12 Cldn12

1433783_at 1.01276 LIM domain binding 3 Ldb3

1433784_at 0.97351 expressed sequence AI265322 AI265322

1433785_at 1.02456 myelin-associated oligodendrocytic basic protein Mobp

1433786_x_at 0.96958 small EDRK-rich factor 2 Serf2

1433787_at 0.92798 NEL-like 1 (chicken) Nell1

1433788_at 1.01736 neurexin III Nrnx3

1433789_at 1.12427 "small nucleolar RNA host gene (non-protein coding) 3 /// similar to RNA, U17d small nucleolar" Snhg3 /// LOC670438

1433790_at 0.99145 trophinin associated protein Troap

1433791_at 0.93276 "RAB9B, member RAS oncogene family" Rab9b

1433792_at 0.84690 nuclear receptor interacting protein 2 Nrip2

1433793_s_at 0.89029 nuclear receptor interacting protein 2 Nrip2

1433794_at 0.97570 senataxin Setx

1433795_at 0.95702 "transforming growth factor, beta receptor III" Tgfbr3

1433796_at 1.00565 endonuclease domain containing 1 Endod1

1433797_at 1.00062 RIKEN cDNA E130309D02 gene E130309D02Rik

1433798_a_at 1.07126 RIKEN cDNA E330034G19 gene E330034G19Rik

1433799_at 0.92037 retinol dehydrogenase 13 (all-trans and 9-cis) Rdh13

1433800_a_at 0.94773 pro-opiomelanocortin-alpha Pomc1

1433801_at 1.08226 RIKEN cDNA 9930012K11 gene 9930012K11Rik

1433802_at 0.84185 transmembrane protein 151 Tmem151

1433803_at 0.92450 Janus kinase 1 Jak1

1433804_at 0.88564 Janus kinase 1 Jak1

1433805_at 1.00316 Janus kinase 1 Jak1

1433806_x_at 0.96494 calreticulin Calr
1433807_at 0.93678 RIKEN cDNA 6720463M24 gene 6720463M24Rik
1433808_at 0.93110 RIKEN cDNA D330001F17 gene D330001F17Rik
1433809_at 0.86798 DEAD (Asp-Glu-Ala-Asp) box polypeptide 5 Ddx5
1433810_x_at 0.83048 DEAD (Asp-Glu-Ala-Asp) box polypeptide 5 Ddx5
1433811_at 0.90105 similar to myeloid/lymphoid or mixed lineage-leukemia translocation to 6 homolog /// similar to myeloid/lymphoid or mixed lineage-leukemia translocation to 6 homolog
LOC639490 /// LOC669933
1433812_at 1.01139 Lix1-like Lix1l
1433813_at 0.89096 transmembrane protein 48 Tmem48
1433814_at 0.91246 Wingless-related MMTV integration site 5B Wnt5b
1433815_at 0.79796 janus kinase and microtubule interacting protein 1 Jakmip1
1433816_at 0.99540 mitochondrial carrier triple repeat 1 Mcart1
1433817_at 0.96872 1-acylglycerol-3-phosphate O-acyltransferase 3 Agpat3
1433818_at 0.83193 1-acylglycerol-3-phosphate O-acyltransferase 3 Agpat3
1433819_s_at 0.96023 1-acylglycerol-3-phosphate O-acyltransferase 3 Agpat3
1433820_a_at 0.93634 RIKEN cDNA 1110012D08 gene 1110012D08Rik
1433821_at 0.98040 RIKEN cDNA 1110012D08 gene 1110012D08Rik
1433822_x_at 0.91336 anaphase-promoting complex subunit 5 Anapc5
1433823_at 0.91465 expressed sequence AW456874 AW456874
1433824_x_at 0.97351 G-rich RNA sequence binding factor 1 Grsf1
1433825_at 1.03042 "Neurotrophic tyrosine kinase, receptor, type 3" Ntrk3
1433826_at 0.90205 TSPY-like 3 Tspyl3
1433827_at 0.88028 "ATPase, aminophospholipid transporter (APLT), class I, type 8A, member 1" Atp8a1
1433828_at 0.93958 "ATPase, aminophospholipid transporter (APLT), class I, type 8A, member 1" Atp8a1
1433829_a_at 1.13863 heterogeneous nuclear ribonucleoprotein A2/B1 Hnrpa2b1
1433830_at 0.97318 heterogeneous nuclear ribonucleoprotein A2/B1 Hnrpa2b1
1433831_at 1.10285 RIKEN cDNA 4833418A01 gene 4833418A01Rik
1433832_at 0.97999 unc-84 homolog B (C. elegans) Unc84b
1433833_at 0.89305 fibronectin type III domain containing 3B Fndc3b
1433834_at 0.88803 membrane-associated ring finger (C3HC4) 6 6-Mar
1433835_at 0.92006 "protein phosphatase 3, catalytic subunit, beta isoform" Ppp3cb
1433836_a_at 0.98729 RIKEN cDNA 8430408G22 gene 8430408G22Rik
1433837_at 1.01672 RIKEN cDNA 8430408G22 gene 8430408G22Rik
1433838_at 0.99937 aspartyl-tRNA synthetase 2 (mitochondrial) Dars2
1433839_at 1.09498 asparaginyl-tRNA synthetase 2 (mitochondrial)(putative) Nars2
1433840_a_at 0.98073 KIAA0415 protein KIAA0415
1433841_at 0.98158 KIAA0415 protein KIAA0415
1433842_at 1.06849 leucine rich repeat (in FLII) interacting protein 1 Lrrfip1
1433843_at 1.08772 HS1 binding protein 3 Hs1bp3
1433844_a_at 1.03892 dual specificity phosphatase 9 Dusp9

1433845_x_at 1.25387 dual specificity phosphatase 9 Dusp9
1433846_s_at 1.03881 RIKEN cDNA C430003P19 gene C430003P19Rik
1433847_at 0.93344 RIKEN cDNA D330017J20 gene D330017J20Rik
1433848_at 0.99129 cell division cycle 27 homolog (S. cerevisiae) Cdc27
1433849_at 0.92512 cell division cycle 27 homolog (S. cerevisiae) Cdc27
1433850_at 1.08414 "protein phosphatase 4, regulatory subunit 2 /// similar to protein phosphatase 4, regulatory subunit 2" Ppp4r2 /// LOC627908
1433851_at 0.95735 "protein phosphatase 4, regulatory subunit 2" Ppp4r2
1433852_at 0.87768 RIKEN cDNA C330002I19 gene C330002I19Rik
1433853_at 0.96913 mindbomb homolog 1 (Drosophila) Mib1
1433854_at 0.90390 transmembrane protein 164 Tmem164
1433855_at 1.00996 4-aminobutyrate aminotransferase Abat
1433856_at 0.99903 histidine acid phosphatase domain containing 1 Hisppd1
1433857_at 0.95947 fat tumor suppressor homolog (Drosophila) /// similar to fat tumor suppressor homolog Fath /// LOC672128
1433858_at 1.13865 leucine rich repeat containing 28 Lrrc28
1433859_at 0.99918 RIKEN cDNA D030070L09 gene D030070L09Rik
1433860_at 1.02020 RIKEN cDNA 6030458C11 gene 6030458C11Rik
1433861_at 1.07156 coiled-coil and C2 domain containing 1B Cc2d1b
1433862_at 1.03039 extra spindle poles-like 1 (S. cerevisiae) Esp11
1433863_at 0.89031 basic transcription factor 3 /// similar to basic transcription factor 3 /// similar to basic transcription factor 3 /// similar to basic transcription factor 3 Btf3 /// LOC640665 /// LOC665385 /// LOC677433
1433864_at 0.86091 low density lipoprotein-related protein 12 Lrp12
1433865_at 1.08023 RIKEN cDNA E330016L19 gene /// RIKEN cDNA E330010L02 gene E330016L19Rik /// E330010L02Rik
1433866_x_at 1.00271 peroxiredoxin 1 Prdx1
1433867_at 1.00508 RIKEN cDNA 1810030O07 gene 1810030O07Rik
1433868_at 1.00437 BTB (POZ) domain containing 3 Btbd3
1433869_at 1.04774 ZXD family zinc finger C Zxdc
1433870_at 0.97558 "ATPase family, AAA domain containing 4" Atad4
1433871_at 1.00785 R3H domain 1 (binds single-stranded nucleic acids) R3hdm1
1433872_at 0.98614 RIKEN cDNA 2410042D21 gene 2410042D21Rik
1433873_s_at 0.96718 pericentrin (kendrin) Pcnt
1433874_at 1.00898 slingshot homolog 1 (Drosophila) Ssh1
1433875_at 0.94407 RIKEN cDNA 4732418C07 gene 4732418C07Rik
1433876_at 0.99466 leucine rich repeat containing 24 Lrrc24
1433877_at 1.11690 RIKEN cDNA 4732473B16 gene 4732473B16Rik
1433878_at 1.02095 mitochondrial ribosomal protein S26 Mrps26
1433879_a_at 1.03386 RIKEN cDNA C130032J12 gene C130032J12Rik
1433880_at 1.13092 "DnaJ (Hsp40) homolog, subfamily C, member 11" Dnajc11
1433881_at 0.91726 "DnaJ (Hsp40) homolog, subfamily C, member 11" Dnajc11
1433882_at 1.02357 "CCR4-NOT transcription complex, subunit 10" Cnot10

1433883_at 0.88411 tropomyosin 4 Tpm4
1433884_at 1.01431 synaptotagmin I Syt1
1433885_at 0.86849 IQ motif containing GTPase activating protein 2 Iqgap2
1433886_at 1.11724 "eukaryotic translation initiation factor 2B, subunit 5 epsilon" Eif2b5
1433887_at 0.84364 "DnaJ (Hsp40) homolog, subfamily C, member 3" Dnajc3
1433888_at 0.88389 "ATPase, Ca++ transporting, plasma membrane 2" Atp2b2
1433889_at 1.01663 SRY-box containing gene 9 Sox9
1433890_a_at 1.00748 HLA-B-associated transcript 3 Bat3
1433891_at 0.80428 leucine-rich repeat-containing G protein-coupled receptor 4 /// similar to
Leucine-rich repeat-containing G-protein coupled receptor 4 precursor Lgr4 /// LOC669588
1433892_at 1.04151 sperm associated antigen 5 Spag5
1433893_s_at 1.10088 sperm associated antigen 5 Spag5
1433894_at 0.80116 expressed sequence AI591476 AI591476
1433895_at 0.99209 transmembrane protein 127 Tmem127
1433896_at 0.94299 transmembrane protein 127 Tmem127
1433897_at 0.94784 expressed sequence AI597468 AI597468
1433898_at 1.02089 Expressed sequence AV025504 AV025504
1433899_x_at 0.86741 "TSC22 domain family, member 1" Tsc22d1
1433900_at 1.07268 RIKEN cDNA 0610007L01 gene 0610007L01Rik
1433901_at 0.91377 GPI-anchored membrane protein 1 Gpiap1
1433902_at 1.11780 kelch repeat and BTB (POZ) domain containing 8 Kbtbd8
1433903_at 0.94723 expressed sequence AU021838 AU021838
1433904_at 1.04023 gametogenetin binding protein 1 Ggnbp1
1433905_at 0.95692 A kinase (PRKA) anchor protein 7 Akap7
1433906_at 1.00013 RIKEN cDNA 4933402J24 gene 4933402J24Rik
1433907_at 0.95782 Pbx/knotted 1 homeobox 2 Pknox2
1433908_a_at 0.97504 cortactin Ctn
1433909_at 0.91800 synaptotagmin XVII Syt17
1433910_at 0.96902 "zinc finger, CCHC domain containing 6" Zcchc6
1433911_at 0.87572 "Smg-6 homolog, nonsense mediated mRNA decay factor (C. elegans)"
Smg6
1433912_at 0.99963 "TIP41, TOR signalling pathway regulator-like (S. cerevisiae)" Tipr1
1433913_at 0.87824 expressed sequence C80913 C80913
1433914_at 1.07446 expressed sequence AI747699 AI747699
1433915_s_at 1.14352 epsin 2 Epn2
1433916_at 0.93137 vesicle-associated membrane protein 3 Vamp3
1433917_x_at 0.77914 "DNA segment, Chr 2, Brigham & Women's Genetics 1423 expressed"
D2Bwg1423e
1433918_at 0.99152 autophagy-related 4D (yeast) Atg4d
1433919_at 1.02786 ankyrin repeat and SOCS box-containing protein 4 Asb4
1433920_at 0.87324 "sema domain, immunoglobulin domain (Ig), transmembrane domain (TM) and
short cytoplasmic domain, (semaphorin) 4C" Sema4c
1433921_s_at 0.98923 "zinc finger, CSL domain containing 2" Zcs12

1433922_at	0.98126	"RAB35, member RAS oncogene family"	Rab35
1433923_at	1.07476	keratin 77	Krt77
1433924_at	0.99754	paternally expressed 3	Peg3
1433925_at	0.84752	"dynein, cytoplasmic 1 light intermediate chain 2"	Dync1li2
1433926_at	0.98178	"dynein, cytoplasmic 1 light intermediate chain 2"	Dync1li2
1433927_at	1.05266	ubiquitin specific peptidase like 1	Usp11
1433928_a_at	1.07553	ribosomal protein L13a	Rpl13a
1433929_at	0.93370	NHL repeat containing 2	Nhlrc2
1433930_at	1.43590	Heparanase	Hpse
1433931_at	0.93230	RIKEN cDNA C030046I01 gene	C030046I01Rik
1433932_x_at	0.94725	RIKEN cDNA C030046I01 gene	C030046I01Rik
1433933_s_at	0.95372	"solute carrier organic anion transporter family, member 2b1"	Slco2b1
1433934_at	1.03860	"SEC24 related gene family, member A (S. cerevisiae)"	Sec24a
1433935_at	1.18119	expressed sequence AU020206	AU020206
1433936_at	1.02036	RIKEN cDNA 0610010E21 gene	0610010E21Rik
1433937_at	0.94923	transformation related protein 53 binding protein 2	Trp53bp2
1433938_at	0.82946	transformation related protein 53 binding protein 2	Trp53bp2
1433939_at	0.73250	hypothetical protein A730046J16	A730046J16
1433940_at	1.15210	sperm associated antigen 7	Spag7
1433941_at	0.96295	"phosphatidylinositol (4,5) bisphosphate 5-phosphatase, A"	Pib5pa
1433942_at	0.96335	myosin VI	Myo6
1433943_at	0.95053	cDNA sequence BC063749	BC063749
1433944_at	1.16606	HECT domain containing 2	Hectd2
1433945_at	1.04496	RIKEN cDNA 5730507A09 gene	5730507A09Rik
1433946_at	0.86857	zinc finger protein interacting with K protein 1	Zik1
1433947_at	1.04476	"RAB37, member of RAS oncogene family"	Rab37
1433948_at	1.21337	"Phospholipase A2, group IB, pancreas"	Pla2g1b
1433949_x_at	0.94332	"phospholipase A2, group IB, pancreas"	Pla2g1b
1433950_at	1.06241	"immunoglobulin superfamily, member 21"	Igsf21
1433951_at	0.89842	ADP-ribosylation factor-like 5A	Arl5a
1433952_at	1.01353	"Tu translation elongation factor, mitochondrial"	Tufm
1433953_at	0.90161	zinc finger protein 277	Zfp277
1433954_at	0.94067	RIKEN cDNA 4632419I22 gene	4632419I22Rik
1433955_at	0.98679	bromodomain and WD repeat domain containing 1	Brwd1
1433956_at	1.11538	cadherin 5	Cdh5
1433957_at	0.78702	RIKEN cDNA C030048B08 gene	C030048B08Rik
1433958_at	0.84810	"centrobin, centrosomal BRCA2 interacting protein"	Cntrob
1433959_at	0.91207	"zinc finger, matrin type 4"	Zmat4
1433960_at	1.06630	interferon stimulated exonuclease gene 20-like 2	Isg20l2
1433961_at	0.95729	cDNA sequence BC023814	BC023814
1433962_at	1.05131	RIKEN cDNA 6720458F09 gene	6720458F09Rik
1433963_a_at	1.04442	cDNA sequence BC032204	BC032204
1433964_s_at	1.08030	cDNA sequence BC032204	BC032204

1433965_at 0.84131 "ATPase, aminophospholipid transporter (APLT), class I, type 8A, member 1" Atp8a1
 1433966_x_at 1.01231 asparagine synthetase Asns
 1433967_at 1.07861 similar to RIKEN cDNA 5430413K10 LOC433492
 1433968_a_at 0.91563 --- ---
 1433969_at 1.04215 expressed sequence AU067824 AU067824
 1433970_at 1.02005 bolA-like 3 (E. coli) Bola3
 1433971_at 0.84533 calmodulin binding transcription activator 1 Camta1
 1433972_at 0.86057 calmodulin binding transcription activator 1 Camta1
 1433973_at 1.05016 selenophosphate synthetase 1 Sephs1
 1433974_at 1.06452 selenophosphate synthetase 1 Sephs1
 1433975_at 0.90313 cyclin-dependent kinase (CDC2-like) 10 Cdk10
 1433976_at 1.08837 "DNA segment, Chr 10, University of California at Los Angeles 1" D10Ucla1
 1433977_at 1.04407 heparan sulfate (glucosamine) 3-O-sulfotransferase 3B1 Hs3st3b1
 1433978_at 0.96819 RIKEN cDNA 4930430F08 gene 4930430F08Rik
 1433979_at 0.97344 "RNA binding motif, single stranded interacting protein 2" Rbms2
 1433980_at 0.99409 "HIV-1 tat interactive protein, homolog (human)" Htatip
 1433981_s_at 0.91082 "HIV-1 tat interactive protein, homolog (human)" Htatip
 1433982_at 0.97978 ubiquitin specific peptidase 28 Usp28
 1433983_at 0.92216 Cnksr family member 3 Cnksr3
 1433984_a_at 0.98119 "malate dehydrogenase 2, NAD (mitochondrial)" Mdh2
 1433985_at 0.93453 abl-interactor 2 Abi2
 1433986_at 0.97642 cDNA sequence BC024659 BC024659
 1433987_at 1.12313 hippocalcin-like 4 Hpcal4
 1433988_s_at 0.92855 RIKEN cDNA C230098O21 gene C230098O21Rik
 1433989_at 1.12201 "solute carrier family 6 (neurotransmitter transporter, GABA), member 11" Slc6a11
 1433990_at 0.97307 lipoma HMGIC fusion partner-like 3 Lhfpl3
 1433991_x_at 1.09534 diazepam binding inhibitor Dbi
 1433992_at 1.06690 shroom family member 2 /// similar to Apical-like protein (APXL protein) Shroom2 /// LOC670546
 1433993_at 0.93352 RIKEN cDNA 4931406P16 gene 4931406P16Rik
 1433994_at 1.11152 RIKEN cDNA 4931406P16 gene 4931406P16Rik
 1433995_s_at 1.00105 "DNA segment, Chr 16, Brigham & Women's Genetics 1543 expressed" D16Bwg1543e
 1433996_at 1.02874 suppressor of variegation 3-9 homolog 2 (Drosophila) Suv39h2
 1433997_at 0.90510 RIKEN cDNA 2410127E18 gene 2410127E18Rik
 1433998_at 1.02875 RIKEN cDNA 4933427D14 gene 4933427D14Rik
 1433999_at 1.01072 STE20-like kinase (yeast) Slk
 1434000_at 0.99531 v-Ki-ras2 Kirsten rat sarcoma viral oncogene homolog Kras
 1434001_at 1.11491 tetratricopeptide repeat domain 9C Ttc9c
 1434002_at 0.79028 checkpoint suppressor 1 Ches1

1434003_a_at 1.02865 deoxyhypusine synthase Dhps
1434004_at 1.07436 deoxyhypusine synthase Dhps
1434005_at 1.01662 "RNA binding motif, single stranded interacting protein 1" Rbms1
1434006_at 0.94129 cDNA sequence BC051227 BC051227
1434007_at 1.01170 glycosyltransferase-like 1B Gylt1b
1434008_at 0.95713 "sodium channel, type IV, beta" Scn4b
1434009_at 1.04157 glucocorticoid receptor DNA binding factor 1 Grfl1
1434010_at 1.10301 "amyotrophic lateral sclerosis 2 (juvenile) chromosome region, candidate 13 (human)" Als2cr13
1434011_a_at 0.96665 integrator complex subunit 5 Ints5
1434012_at 0.93946 integrator complex subunit 5 Ints5
1434013_at 0.88535 "actin binding LIM protein family, member 3" Ablim3
1434014_at 0.95684 autophagy-related 4C (yeast) Atg4c
1434015_at 1.30424 "solute carrier family 2 (facilitated glucose transporter), member 6" Slc2a6
1434016_at 1.05560 zinc and ring finger 2 Znrf2
1434017_at 0.92229 --- ---
1434018_at 0.98454 cDNA sequence BC043098 BC043098
1434019_at 0.99519 PDGFA associated protein 1 Pdap1
1434020_at 1.11331 PDGFA associated protein 1 Pdap1
1434021_at 0.97709 RIKEN cDNA C230096C10 gene C230096C10Rik
1434022_at 0.86362 zinc finger and BTB domain containing 33 Zbtb33
1434023_at 0.84973 expressed sequence AU016693 AU016693
1434024_at 0.89492 nephronophthisis 4 (juvenile) homolog (human) Nphp4
1434025_at 1.25272 --- ---
1434026_at 0.86837 "Atpase, class I, type 8B, member 2" Atp8b2
1434027_at 0.84851 Down syndrome critical region gene 1-like 2 Dscr112
1434028_at 0.87185 aryl hydrocarbon receptor nuclear translocator 2 Arnt2
1434029_at 0.93028 RIKEN cDNA 2410025L10 gene 2410025L10Rik
1434030_at 0.97500 cDNA sequence BC034507 BC034507
1434031_at 1.20831 zinc finger protein 692 Zfp692
1434032_at 0.83324 --- ---
1434033_at 0.86225 "transducin-like enhancer of split 1, homolog of Drosophila E(spl)" Tle1
1434034_at 0.73468 ceramide kinase CerK
1434035_at 0.96019 "DnaJ (Hsp40) homolog, subfamily B, member 6" Dnajb6
1434036_at 0.95522 metastasis suppressor 1 Mtss1
1434037_s_at 0.98696 p300/CBP-associated factor Pcaf
1434038_at 1.23445 "DnaJ (Hsp40) homolog, subfamily C, member 13" Dnajc13
1434039_at 0.92440 amyloid beta precursor protein (cytoplasmic tail) binding protein 2 Appbp2
1434040_at 0.98945 amyloid beta precursor protein (cytoplasmic tail) binding protein 2 Appbp2
1434041_at 1.02909 amyloid beta precursor protein (cytoplasmic tail) binding protein 2 Appbp2
1434042_s_at 1.01281 myotubularin related protein 3 Mtmr3
1434043_a_at 0.90421 --- ---
1434044_at 0.81526 replication initiator 1 Repin1

1434045_at 0.93281 cyclin-dependent kinase inhibitor 1B Cdkn1b
1434046_at 0.88778 expressed sequence AA467197 AA467197
1434047_x_at 1.07504 heterogeneous nuclear ribonucleoprotein A2/B1 Hnrpa2b1
1434048_at 0.84682 RIKEN cDNA 4930471M23 gene 4930471M23Rik
1434049_at 0.91650 ectonucleoside triphosphate diphosphohydrolase 3 Entpd3
1434050_at 0.93794 expressed sequence AI315068 AI315068
1434051_s_at 0.86394 heat shock protein 12A Hspa12a
1434052_at 1.10309 expressed sequence AI593442 AI593442
1434053_x_at 0.75657 "ATP synthase, H+ transporting, mitochondrial F1F0 complex, subunit e"
Atp5k
1434054_at 1.18421 "v-maf musculoaponeurotic fibrosarcoma oncogene family, protein G
(avian)" Mafg
1434055_at 1.01334 UDP-N-acetyl-alpha-D-galactosamine:polypeptide N-
acetylgalactosaminyltransferase 9 Galnt9
1434056_a_at 0.91461 "NADH dehydrogenase (ubiquinone) 1 beta subcomplex, 6" Ndufb6
1434057_at 0.96188 "NADH dehydrogenase (ubiquinone) 1 beta subcomplex, 6" Ndufb6
1434058_at 1.10293 Vmyotubularin related protein 12 Mtmr12
1434059_at 0.89622 RIKEN cDNA B230312A22 gene B230312A22Rik
1434060_at 1.03373 hect (homologous to the E6-AP (UBE3A) carboxyl terminus) domain and
RCC1 (CHC1)-like domain (RLD) 1 Herc1
1434061_at 1.01540 retinitis pigmentosa 2 homolog (human) Rp2h
1434062_at 0.99648 RAB GTPase activating protein 1-like Rabgap11
1434063_at 1.02335 zinc finger protein 664 Zfp664
1434064_at 0.94101 transmembrane protein 142C Tmem142c
1434065_at 1.03804 "CWF19-like 1, cell cycle control (S. pombe)" Cwf1911
1434066_at 0.97287 general transcription factor III C 1 Gtf3c1
1434067_at 1.01836 expressed sequence AI662270 AI662270
1434068_s_at 0.83198 expressed sequence AI662270 AI662270
1434069_at 0.93860 SET domain containing 6 Setd6
1434070_at 0.88332 jagged 1 Jag1
1434071_a_at 0.99494 pelota homolog (Drosophila) Pelo
1434072_at 0.89546 "Smith-Magenis syndrome chromosome region, candidate 7 homolog (human)"
Smcr7
1434073_at 0.99667 G protein-coupled receptor associated sorting protein 2 Gprasp2
1434074_x_at 0.98919 ADP-ribosylation factor 4 Arf4
1434075_at 0.88646 cDNA sequence BC030336 BC030336
1434076_at 1.05400 WD repeat domain 37 Wdr37
1434077_at 1.10374 WD repeat domain 37 Wdr37
1434078_at 1.00065 "DNA segment, Chr 7, Wayne State University 128, expressed" D7Wsu128e
1434079_s_at 0.92080 minichromosome maintenance deficient 2 mitotin (S. cerevisiae) Mcm2
1434080_at 1.09933 AE binding protein 2 Aebp2
1434081_at 1.06421 "adaptor protein complex AP-1, gamma 1 subunit" Ap1g1
1434082_at 0.95261 --- ---

1434083_a_at	0.70232	ELMO domain containing 1	Elmod1
1434084_at	0.91861	RIKEN cDNA 5730601F06 gene	5730601F06Rik
1434085_at	0.93616	zinc finger protein 523	Zfp523
1434086_at	1.00045	expressed sequence AI790205	AI790205
1434087_at	1.07268	"5,10-methylenetetrahydrofolate reductase"	Mthfr
1434088_at	0.94425	zinc finger protein 496	Zfp496
1434089_at	0.57408	synaptopodin	Synpo
1434090_at	0.98499	RIKEN cDNA 4930429A22 gene	4930429A22Rik
1434091_at	0.86083	fatty acid amide hydrolase	Faah
1434092_at	1.32710	nitric oxide synthase 3 antisense	Nos3as
1434093_at	0.83405	"LysM, putative peptidoglycan-binding, domain containing 4"	Lysmd4
1434094_at	0.82888	RIKEN cDNA 6330530A05 gene	6330530A05Rik
1434095_at	1.05095	unc-5 homolog A (C. elegans)	Unc5a
1434096_at	0.96726	"solute carrier family 4 (anion exchanger), member 4"	Slc4a4
1434097_at	0.99157	Transcribed locus	---
1434098_at	0.94350	"glycine receptor, alpha 2 subunit"	Gla2
1434099_at	0.68403	"Peroxisome proliferative activated receptor, gamma, coactivator 1 alpha"	Ppargc1a
1434100_x_at	0.79849	"Peroxisome proliferative activated receptor, gamma, coactivator 1 alpha"	Ppargc1a
1434101_at	1.18142	nuclear factor I/B	Nfib
1434102_at	0.90842	nuclear factor I/B	Nfib
1434103_at	0.91318	expressed sequence AA408278	AA408278
1434104_at	0.98770	hypothetical protein 6030458H05	6030458H05
1434105_at	0.90936	EPM2A (laforin) interacting protein 1	Epm2aip1
1434106_at	0.88663	EPM2A (laforin) interacting protein 1	Epm2aip1
1434107_at	1.02383	spermatogenesis associated 2	Spata2
1434108_at	0.98132	F-box protein 11	Fbxo11
1434109_at	0.97997	SH3 domain binding glutamic acid-rich protein like 2	Sh3bgrl2
1434110_x_at	0.94532	major urinary protein 1	Mup1
1434111_at	0.76494	latrophilin 2	Lphn2
1434112_at	0.81159	latrophilin 2	Lphn2
1434113_a_at	1.05752	"REX4, RNA exonuclease 4 homolog (S. cerevisiae)"	Rexo4
1434114_at	1.21646	"REX4, RNA exonuclease 4 homolog (S. cerevisiae)"	Rexo4
1434115_at	1.18266	---	---
1434116_at	0.95101	chromobox homolog 2 (Drosophila Pc class)	Cbx2
1434117_at	0.95122	"transcription elongation factor B (SIII), polypeptide 3"	Tceb3
1434118_at	0.73415	RIKEN cDNA 0610009K11 gene	0610009K11Rik
1434119_at	0.91327	"DNA segment, Chr 2, Wayne State University 81, expressed"	D2Wsu81e
1434120_a_at	1.05256	methionine aminopeptidase 2	Metap2
1434121_at	0.87699	"leucine-rich repeat LGI family, member 4"	Lgi4
1434122_at	0.90663	F-box protein 27	Fbxo27
1434123_at	0.87316	fucosyltransferase 11	Fut11

1434124_x_at 1.02482 RIKEN cDNA 2400001E08 gene 2400001E08Rik
1434125_at 1.16621 "UTP15, U3 small nucleolar ribonucleoprotein, homolog (yeast)" Utp15
1434126_at 0.94120 RIKEN cDNA 4930402H24 gene 4930402H24Rik
1434127_a_at 0.96252 "H3 histone, family 3A /// similar to H3 histone, family 3B /// similar to H3 histone, family 3B /// similar to H3 histone, family 3B /// similar to H3 histone, family 3B /// similar to H3 histone, family 3B" H3f3a /// LOC544973 /// LOC625328 /// LOC669889 /// LOC670535 /// LOC675719
1434128_a_at 0.82247 zinc finger protein 574 Zfp574
1434129_s_at 1.18438 lipoma HMGIC fusion partner-like 2 Lhfpl2
1434130_at 1.31245 lipoma HMGIC fusion partner-like 2 Lhfpl2
1434131_at 0.91231 RUN and FYVE domain containing 1 Rufy1
1434132_at 1.09627 RIKEN cDNA E430025E21 gene E430025E21Rik
1434133_s_at 0.83396 WD repeat domain 42A Wdr42a
1434134_at 0.97161 WD repeat domain 42A Wdr42a
1434135_at 1.12210 "UDP-GalNAc:betaGlcNAc beta 1,3-galactosaminyltransferase, polypeptide 2" B3galnt2
1434136_at 1.08488 RIKEN cDNA 6332401O19 gene 6332401O19Rik
1434137_x_at 1.06748 RIKEN cDNA 1810010M01 gene 1810010M01Rik
1434138_at 0.83293 prune homolog (Drosophila) Prune
1434139_at 1.10479 "poly (ADP-ribose) polymerase family, member 11" Parp11
1434140_at 0.59012 mcf.2 transforming sequence-like Mcf2l
1434141_at 1.08387 "guanylate cyclase 1, soluble, alpha 3" Gucyl1a3
1434142_at 0.84660 "protein phosphatase 1D magnesium-dependent, delta isoform" Ppm1d
1434143_at 0.92494 cDNA sequence BC060631 BC060631
1434144_s_at 1.42371 RIKEN cDNA 2410187C16 gene 2410187C16Rik
1434145_s_at 0.84955 serine hydrolase-like Serhl
1434146_at 1.06255 "glutamate receptor, ionotropic, AMPA2 (alpha 2)" Gria2
1434147_at 0.99381 RIKEN cDNA 4631403P03 gene 4631403P03Rik
1434148_at 0.89359 transcription factor 4 Tcf4
1434149_at 0.83352 transcription factor 4 Tcf4
1434150_a_at 0.86388 methyltransferase like 7A /// UbiE-YGHL1 fusion protein Mettl7a /// Ubie
1434151_at 0.95280 methyltransferase like 7A Mettl7a
1434152_at 0.94356 "RIKEN cDNA 2210421G13 gene /// hypothetical protein LOC193676 /// similar to apolipoprotein L, 3 (predicted)" 2210421G13Rik /// LOC193676 /// LOC666661
1434153_at 1.12869 src homology 2 domain-containing transforming protein B Shb
1434154_at 0.93894 potassium channel tetramerisation domain containing 13 Kctd13
1434155_a_at 1.04097 RIKEN cDNA 2310061I04 gene 2310061I04Rik
1434156_at 0.90130 --- ---
1434157_at 0.89813 taxilin alpha Txlna
1434158_at 0.92476 "GDP-mannose 4, 6-dehydratase" Gmds
1434159_at 1.03201 serine/threonine kinase 4 Stk4
1434160_at 1.10219 zinc finger protein 592 Zfp592
1434161_at 1.01815 RIKEN cDNA 5830457H20 gene 5830457H20Rik

1434162_at 0.99722 RIKEN cDNA 2700078E11 gene 2700078E11Rik
1434163_at 1.03926 expressed sequence AU024076 AU024076
1434164_s_at 1.07503 expressed sequence AU024076 AU024076
1434165_at 0.96865 chloride intracellular channel 6 Clic6
1434166_at 1.04536 RIKEN cDNA 9330151L19 gene 9330151L19Rik
1434167_at 1.05842 "solute carrier family 35, member E4" Slc35e4
1434168_at 1.12421 progressive external ophthalmoplegia 1 (human) Peo1
1434169_at 0.91326 RIKEN cDNA 9030409G11 gene 9030409G11Rik
1434170_at 0.96463 WD repeat domain 40B Wdr40b
1434171_at 0.92414 RIKEN cDNA C330011K17 gene C330011K17Rik
1434172_at 0.96555 cannabinoid receptor 1 (brain) Cnr1
1434173_s_at 1.05037 "DNA segment, Chr 19, Brigham & Women's Genetics 1357 expressed /// SWI/SNF related, matrix associated, actin dependent regulator of chromatin, subfamily a, member 2" D19Bwg1357e /// Smarca2
1434174_at 1.06224 "LysM, putative peptidoglycan-binding, domain containing 3" Lysmd3
1434175_s_at 0.92783 RIKEN cDNA 2210010N04 gene 2210010N04Rik
1434176_x_at 1.04782 "polymerase (DNA-directed), delta interacting protein 3" Poldip3
1434177_at 1.04949 endothelin converting enzyme 1 Ece1
1434178_at 0.89629 myeloid/lymphoid or mixed-lineage leukemia 3 Mll3
1434179_at 0.75585 myeloid/lymphoid or mixed-lineage leukemia 3 Mll3
1434180_at 0.97912 "pleckstrin homology domain containing, family C (with FERM domain) member 1" Plekhc1
1434181_at 0.95702 "pleckstrin homology domain containing, family C (with FERM domain) member 1" Plekhc1
1434182_at 1.01714 DEAD/H (Asp-Glu-Ala-Asp/His) box polypeptide 31 Ddx31
1434183_at 0.98061 similar to sterile alpha motif domain containing 1 /// similar to sterile alpha motif domain containing 1 /// similar to sterile alpha motif domain containing 1 LOC546078 /// LOC666704 /// LOC671345
1434184_s_at 1.11730 RIKEN cDNA 9430080K19 gene 9430080K19Rik
1434185_at 1.06686 similar to acetyl-coenzyme A carboxylase alpha LOC672245
1434186_at 1.08091 G protein-coupled receptor 23 Gpr23
1434187_at 1.08858 --- ---
1434188_at 0.92472 "solute carrier family 16 (monocarboxylic acid transporters), member 12" Slc16a12
1434189_at 1.06968 stromal antigen 1 Stag1
1434190_at 1.07281 spermine synthase Sms
1434191_at 1.02894 RIKEN cDNA A530016O06 gene A530016O06Rik
1434192_at 1.12931 "zinc finger, ZZ-type with EF hand domain 1" Zzef1
1434193_at 1.02785 "zinc finger, MYM-type 6" Zmym6
1434194_at 0.66868 microtubule-associated protein 2 Mtap2
1434195_at 1.06741 "protease, serine, 35" Prss35
1434196_at 1.05253 "DnaJ (Hsp40) homolog, subfamily A, member 4" Dnaja4
1434197_at 0.95328 attractin Atrn

1434198_at 1.05673 ATPase inhibitory factor 1 Atpif1
1434199_at 0.98444 "gene model 323, (NCBI)" Gm323
1434200_at 0.94892 cDNA sequence BC010981 BC010981
1434201_at 0.90407 Chordin-like 1 Chrdl1
1434202_a_at 0.86403 cDNA sequence BC055107 BC055107
1434203_at 0.77536 cDNA sequence BC055107 BC055107
1434204_x_at 1.03030 Serine hydroxymethyl transferase 2 (mitochondrial) Shmt2
1434205_at 0.83515 "protein phosphatase 2, regulatory subunit B (B56), gamma isoform" Ppp2r5c
1434206_s_at 0.68477 "protein phosphatase 2, regulatory subunit B (B56), gamma isoform"
Ppp2r5c
1434207_at 0.96585 RIKEN cDNA 2900057K09 gene 2900057K09Rik
1434208_at 0.85592 RIKEN cDNA 2900057K09 gene 2900057K09Rik
1434209_at 0.94136 protein arginine N-methyltransferase 6 Prmt6
1434210_s_at 1.21435 leucine-rich repeats and immunoglobulin-like domains 1 Lrig1
1434211_at 0.95192 SH3 domain binding glutamic acid-rich protein like 2 Sh3bgrl2
1434212_at 0.98941 NADH dehydrogenase (ubiquinone) Fe-S protein 8 Ndufs8
1434213_x_at 0.98691 NADH dehydrogenase (ubiquinone) Fe-S protein 8 Ndufs8
1434214_at 0.99758 RIKEN cDNA 0910001L09 gene 0910001L09Rik
1434215_at 1.03109 RIKEN cDNA B230308N11 gene B230308N11Rik
1434216_a_at 0.92286 nudix (nucleoside diphosphate linked moiety X)-type motif 19 Nudt19
1434217_at 0.99062 RIKEN cDNA C330019G07 gene C330019G07Rik
1434218_at 0.90724 RIKEN cDNA C330019G07 gene C330019G07Rik
1434219_at 0.99964 stromal interaction molecule 2 Stim2
1434220_at 1.12306 nucleoporin 98 Nup98
1434221_at 0.86362 cDNA sequence BC030863 BC030863
1434222_at 0.94533 signal-induced proliferation-associated 1 like 1 Sipal11
1434223_at 0.90428 RIKEN cDNA 1810007P19 gene 1810007P19Rik
1434224_at 1.27394 transducin (beta)-like 2 Tbl2
1434225_at 1.00498 SWA-70 protein Swap70
1434226_at 0.88366 cytochrome b-561 domain containing 1 Cyb561d1
1434227_at 1.07738 keratinocyte differentiation associated protein Krtdap
1434228_at 0.98206 "protein phosphatase 2C, magnesium dependent, catalytic subunit" Ppm2c
1434229_a_at 1.04765 "polymerase (DNA directed), beta" Polb
1434230_at 0.93532 "polymerase (DNA directed), beta /// RIKEN cDNA A430088C08 gene"
Polb /// A430088C08Rik
1434231_x_at 1.00026 ribosomal protein L35 Rpl35
1434232_a_at 0.96222 RIKEN cDNA 2610030H06 gene 2610030H06Rik
1434233_at 1.06635 RIKEN cDNA 2610030H06 gene 2610030H06Rik
1434234_at 0.97411 zinc finger protein 341 Zfp341
1434235_at 1.06360 "solute carrier family 20, member 2" Slc20a2
1434236_at 0.94037 "zinc finger, DHHC domain containing 20" Zdhhc20
1434237_at 1.17490 uroplakin 3B Upk3b
1434238_at 1.01073 "TAF2 RNA polymerase II, TATA box binding protein (TBP)-associated

factor" Taf2

1434239_at	1.28587	expressed sequence AA408556	AA408556
1434240_at	0.91581	RIKEN cDNA 4632434I11 gene	4632434I11Rik
1434241_at	1.01257	WD repeat domain 67	Wdr67
1434242_at	0.96281	ubiquitin specific peptidase 37	Usp37
1434243_s_at	1.13441	---	---
1434244_x_at	0.99449	HpaII tiny fragments locus 9c	Htf9c
1434245_a_at	0.93574	"cytochrome b, ascorbate dependent 3"	Cybasec3
1434246_at	1.01288	l(3)mbt-like 3 (Drosophila)	L3mbtl3
1434247_at	0.91314	lactate dehydrogenase A-like 6B	Ldhal6b
1434248_at	0.84224	"protein kinase C, eta"	Prkch
1434249_s_at	1.03480	---	---
1434250_at	0.96018	p21 (CDKN1A)-activated kinase 2	Pak2
1434251_at	1.05660	"CCR4-NOT transcription complex, subunit 1"	Cnot1
1434252_at	0.81162	transmembrane and coiled coil domains 3	Tmcc3
1434253_s_at	0.92022	transmembrane and coiled coil domains 3	Tmcc3
1434254_at	0.97729	"guanine nucleotide binding protein, alpha 11"	Gna11
1434255_at	0.91163	phosphofurin acidic cluster sorting protein 2	Pacs2
1434256_s_at	1.01085	CDP-diacylglycerol synthase (phosphatidate cytidyltransferase) 2	Cds2
1434257_s_at	1.00772	"phosphatidylinositol transfer protein, beta"	Pitpnb
1434258_s_at	1.06472	phosphatase and actin regulator 4	Phactr4
1434259_at	1.08197	"similar to H2A histone family, member J"	LOC666811
1434260_at	0.79877	FCH and double SH3 domains 2	Fchsd2
1434261_at	0.73447	signal-induced proliferation-associated 1 like 2	Sipa1l2
1434262_at	1.05787	---	---
1434263_at	0.96028	RIKEN cDNA F630110N24 gene	F630110N24Rik
1434264_at	1.06960	"ankyrin 2, brain"	Ank2
1434265_s_at	0.94078	"ankyrin 2, brain"	Ank2
1434266_at	1.01971	expressed sequence AI847670	AI847670
1434267_at	0.92294	NIMA (never in mitosis gene a)-related expressed kinase 1	Nek1
1434268_at	1.04043	"adenosine deaminase, RNA-specific"	Adar
1434269_at	0.73296	expressed sequence AA536717	AA536717
1434270_at	0.76940	neuronal pentraxin receptor	Nptxr
1434271_at	0.90939	glucosidase beta 2	Gba2
1434272_at	0.97845	cytoplasmic polyadenylation element binding protein 2	Cpeb2
1434273_at	1.06946	RIKEN cDNA A830073O21 gene	A830073O21Rik
1434274_at	1.02372	"pam, highwire, rpm 1"	Phr1
1434275_at	0.85026	Transcribed locus	---
1434276_x_at	0.98821	cDNA sequence BC004004	BC004004
1434277_a_at	0.78594	Hypothetical protein 6430570G24	6430570G24
1434278_at	1.18770	X-linked myotubular myopathy gene 1	Mtm1
1434279_at	1.13791	X-linked myotubular myopathy gene 1	Mtm1
1434280_at	1.14632	Widely-interspaced zinc finger motifs	Wiz

1434281_at 0.85687 RIKEN cDNA 1500034J01 gene 1500034J01Rik
1434282_at 0.89112 inhibitor of Bruton agammaglobulinemia tyrosine kinase Ibtk
1434283_at 0.84203 AT rich interactive domain 5B (Mrf1 like) Arid5b
1434284_at 1.12197 "B double prime 1, subunit of RNA polymerase III transcription initiation factor IIIB" Bdp1
1434285_at 0.94818 FERM domain containing 4A Frmd4a
1434286_at 0.52547 trichorhinophalangeal syndrome I (human) Trps1
1434287_at 0.98558 "1-acylglycerol-3-phosphate O-acyltransferase 5 (lysophosphatidic acid acyltransferase, epsilon)" Apat5
1434288_at 0.97773 "basic, immunoglobulin-like variable motif containing" Bivm
1434289_at 0.96872 "NOL1/NOP2/Sun domain family, member 5" Nsun5
1434290_at 1.00888 glycosyltransferase-like domain containing 1 Gtdc1
1434291_a_at 0.83184 small EDRK-rich factor 1 Serf1
1434292_at 0.98419 RIKEN cDNA E130013N09 gene E130013N09Rik
1434293_at 0.88327 HECT domain containing 3 Hectd3
1434294_at 0.82977 Transcribed locus ---
1434295_at 1.07269 RAS guanyl releasing protein 1 Rasgrp1
1434296_at 1.05369 cDNA sequence BC049349 BC049349
1434297_at 1.27815 RIKEN cDNA E130304F04 gene E130304F04Rik
1434298_at 0.92086 zinc finger homeobox 1b Zfhx1b
1434299_x_at 1.03098 "RAB, member of RAS oncogene family-like 4" Rabl4
1434300_at 1.02121 RIKEN cDNA 2610101N10 gene 2610101N10Rik
1434301_at 0.79059 RIKEN cDNA D330050I23 gene D330050I23Rik
1434302_at 1.69048 Ras association (RalGDS/AF-6) and pleckstrin homology domains 1 Raph1
1434303_at 1.79657 Ras association (RalGDS/AF-6) and pleckstrin homology domains 1 Raph1
1434304_s_at 1.02267 "DNA segment, Chr 10, ERATO Doi 438, expressed" D10Erd438e
1434305_at 0.97754 U2 small nuclear RNA auxiliary factor 1-like 4 U2af114
1434306_at 0.96104 RAB3A interacting protein Rab3ip
1434307_at 0.82108 transmembrane protein 64 Tmem64
1434308_at 1.01770 "solute carrier family 43, member 2" Slc43a2
1434309_at 0.80283 "farnesyltransferase, CAAX box, beta" Fntb
1434310_at 0.96586 "bone morphogenic protein receptor, type II (serine/threonine kinase)" Bmpr2
1434311_at 0.83627 "CCR4-NOT transcription complex, subunit 6-like" Cnot6l
1434312_at 0.97659 expressed sequence AI788669 AI788669
1434313_at 0.95718 RIKEN cDNA 6330407D12 gene 6330407D12Rik
1434314_s_at 0.85659 RAB11 family interacting protein 5 (class I) Rab11fip5
1434315_at 0.75783 NIPA-like domain containing 3 Npal3
1434316_at 1.07690 carbohydrate (chondroitin) synthase 1 Chsy1
1434317_s_at 1.09610 testis expressed gene 10 Tex10
1434318_a_at 0.98123 transcription factor E3 Tcf3
1434319_at 1.03578 "malate dehydrogenase 1, NAD (soluble)" Mdh1
1434320_at 1.00538 "general transcription factor IIIC, polypeptide 4" Gtf3c4
1434321_at 0.88255 RIKEN cDNA 6620401K05 gene 6620401K05Rik

1434322_at 1.49514 RIKEN cDNA A930021H16 gene A930021H16Rik
1434323_at 0.86444 "eukaryotic translation initiation factor 3, subunit 3 (gamma) /// similar to eukaryotic translation initiation factor 3, subunit 3 gamma, 40kDa" Eif3s3 /// LOC625074
1434324_x_at 0.90543 "eukaryotic translation initiation factor 3, subunit 3 (gamma) /// similar to eukaryotic translation initiation factor 3, subunit 3 gamma, 40kDa" Eif3s3 /// LOC625074
1434325_x_at 1.01317 "protein kinase, cAMP dependent regulatory, type I beta" Prkar1b
1434326_x_at 0.71469 "coronin, actin binding protein, 2B" Coro2b
1434327_at 0.92306 --- ---
1434328_at 0.92100 ribosomal protein L15 Rpl15
1434329_s_at 1.14372 adiponectin receptor 2 Adipor2
1434330_at 1.07253 leucine rich repeat containing 35 Lrrc35
1434331_at 1.03636 "eukaryotic translation initiation factor 2C, 1" Eif2c1
1434332_at 1.02501 "zinc finger, ZZ domain containing 3" Zzz3
1434333_a_at 0.97355 protein kinase D2 Prkd2
1434334_at 0.86584 protein kinase D2 Prkd2
1434335_at 0.92827 expressed sequence AI317237 AI317237
1434336_s_at 0.87822 REST corepressor 1 Rcor1
1434337_at 0.91604 Transcribed locus ---
1434338_at 1.04549 RIKEN cDNA 1110029E03 gene 1110029E03Rik
1434339_at 0.93575 formin binding protein 1-like Fnbp11
1434340_at 1.13343 --- ---
1434341_x_at 1.01358 RIKEN cDNA 1110020P15 gene /// similar to ubiquinol-cytochrome c reductase complex 7.2kDa protein isoform a 1110020P15Rik /// LOC622178
1434342_at 0.89137 "S100 protein, beta polypeptide, neural" S100b
1434343_at 1.03055 RIKEN cDNA 5730403M16 gene 5730403M16Rik
1434344_at 0.93421 G patch domain and KOW motifs Gpkow
1434345_at 1.00277 transmembrane protein 12 Tmem12
1434346_at 0.87993 --- ---
1434347_s_at 1.03820 --- ---
1434348_at 1.00275 fasciculation and elongation protein zeta 2 (zygin II) Fez2
1434349_at 0.95505 valyl-tRNA synthetase 2-like Vars21
1434350_at 0.87492 AXIN1 up-regulated 1 Axud1
1434351_at 1.04670 nucleoporin 214 Nup214
1434352_at 1.11310 RIKEN cDNA B630005N14 gene B630005N14Rik
1434353_at 0.87035 Scm-like with four mbt domains 2 Sfm2
1434354_at 0.97426 monoamine oxidase B Maob
1434355_at 0.99669 zinc finger protein 617 Zfp617
1434356_a_at 1.05959 "proteasome (prosome, macropain) subunit, alpha type 5" Psma5
1434357_a_at 1.12195 karyopherin (importin) beta 1 Kpnb1
1434358_x_at 0.99778 ribosomal protein S21 Rps21
1434359_at 1.08044 RIKEN cDNA 6330500D04 gene 6330500D04Rik
1434360_s_at 1.02376 "protein tyrosine phosphatase, receptor type, G /// similar to protein tyrosine phosphatase, receptor type, G" Ptpg /// LOC632664

1434361_at 0.90556 SH3 and PX domain containing 3 Sh3px3
1434362_at 1.00799 expressed sequence AW550831 AW550831
1434363_x_at 0.99378 PAX interacting (with transcription-activation domain) protein 1 Paxip1
1434364_at 0.98589 mitogen-activated protein kinase kinase kinase 14 Map3k14
1434365_a_at 0.96988 cDNA sequence BC055324 BC055324
1434366_x_at 0.99274 "complement component 1, q subcomponent, beta polypeptide" C1qb
1434367_s_at 1.02856 nuclear transport factor 2 Nutf2
1434368_at 0.90855 RIKEN cDNA 1700008K24 gene 1700008K24Rik
1434369_a_at 0.87574 "crystallin, alpha B" Cryab
1434370_s_at 0.98085 Fas-associated factor 1 Faf1
1434371_x_at 1.10992 ubiquitin specific peptidase 7 Usp7
1434372_at 1.74218 expressed sequence AW112010 AW112010
1434373_at 0.85141 RIKEN cDNA B930006L02 gene B930006L02Rik
1434374_at 0.90910 RIKEN cDNA B930006L02 gene B930006L02Rik
1434375_at 0.98802 RIKEN cDNA B930006L02 gene B930006L02Rik
1434376_at 1.13961 CD44 antigen Cd44
1434377_x_at 1.06324 ribosomal protein S6 /// similar to 40S ribosomal protein S6 /// similar to 40S ribosomal protein S6 Rps6 /// LOC667739 /// LOC672620
1434378_a_at 0.73815 Max dimerization protein 4 Mxd4
1434379_at 0.71383 Max dimerization protein 4 Mxd4
1434380_at 1.00425 guanylate binding protein 7 Gbp7
1434381_at 0.83784 cDNA sequence BC060631 BC060631
1434382_at 0.69565 serine incorporator 2 Serinc2
1434383_at 0.97196 "praja 2, RING-H2 motif containing" Pja2
1434384_at 0.82682 nuclear receptor interacting protein 1 Nrip1
1434385_at 0.85383 target of myb1-like 2 (chicken) Tom112
1434386_at 0.91628 "ATPase, Ca⁺⁺-sequestering" Atp2c1
1434387_at 0.85657 integrin alpha FG-GAP repeat containing 3 Itfg3
1434388_at 1.01802 "MOB1, Mps One Binder kinase activator-like 2A (yeast)" Mobk12a
1434389_at 0.97089 expressed sequence AI449023 AI449023
1434390_at 0.95520 Heterogeneous nuclear ribonucleoprotein U Hnrpu
1434391_at 0.90163 Heterogeneous nuclear ribonucleoprotein U Hnrpu
1434392_at 1.02064 ubiquitin specific peptidase 34 Usp34
1434393_at 1.05497 ubiquitin specific peptidase 34 Usp34
1434394_at 0.90799 Bcl3 binding protein B3bp
1434395_at 0.89796 "mannosidase, alpha, class 1A, member 2" Man1a2
1434396_a_at 0.95064 "myosin, light polypeptide 6, alkali, smooth muscle and non-muscle /// similar to myosin, light polypeptide 6, alkali, smooth muscle and non-muscle /// similar to Myosin light polypeptide 6 (Myosin light chain alkali 3) (Myosin light chain 3) (MLC-3) (LC17) /// similar to Myosin light polypeptide 6 (Myosin light chain alkali 3) (Myosin light chain 3) (MLC-3) (LC17)" Myl6 /// LOC433297 /// LOC667952 /// LOC671953
1434397_at 1.00687 "zinc finger, DHHC domain containing 17" Zdhhc17
1434398_at 1.05885 NF-kappaB repressing factor Nkrf

1434399_at 0.94489 UDP-N-acetyl-alpha-D-galactosamine:polypeptide N-acetylgalactosaminyltransferase 6 Galnt6
1434400_at 0.98091 TGFβ-induced factor 2 Tgif2
1434401_at 0.99257 "zinc finger, CCHC domain containing 2" Zcchc2
1434402_at 0.94561 sterile alpha motif domain containing 8 Samd8
1434403_at 1.16913 "sprouty-related, EVH1 domain containing 2" Spred2
1434404_at 0.93974 RIKEN cDNA C030011O14 gene C030011O14Rik
1434405_at 1.13336 RIKEN cDNA A730024A03 gene A730024A03Rik
1434406_at 1.00630 SLIT-ROBO Rho GTPase activating protein 2 Srgap2
1434407_at 0.93036 SLIT-ROBO Rho GTPase activating protein 2 Srgap2
1434408_at 0.86268 "Machado-Joseph disease (spinocerebellar ataxia 3, olivopontocerebellar ataxia 3, autosomal dominant, ataxin 3) homolog (human)" Mjd
1434409_at 0.79129 "armadillo repeat containing, X-linked 6" Armcx6
1434410_at 0.90437 cDNA sequence BC043118 BC043118
1434411_at 0.76548 --- ---
1434412_x_at 0.99756 STIP1 homology and U-Box containing protein 1 Stub1
1434413_at 0.67428 --- ---
1434414_at 1.09829 FAD-dependent oxidoreductase domain containing 2 Foxred2
1434415_at 0.84945 dact 3 Dact3
1434416_a_at 1.04592 Small optic lobes homolog (Drosophila) Solh
1434417_at 0.91641 --- ---
1434418_at 1.08440 longevity assurance homolog 6 (S. cerevisiae) Lass6
1434419_s_at 1.07629 TAR DNA binding protein Tardbp
1434420_x_at 1.05218 translocase of outer mitochondrial membrane 22 homolog (yeast) Tomm22
1434421_at 0.96303 immunoglobulin superfamily containing leucine-rich repeat 2 Islr2
1434422_at 1.06690 --- ---
1434423_at 0.97049 --- ---
1434424_at 1.03584 RIKEN cDNA 9630055N22 gene /// similar to Feline leukemia virus subgroup C receptor-related protein 1 (Feline leukemia virus subgroup C receptor) (hFLVCR) 9630055N22Rik /// LOC670389
1434425_at 1.03874 trichohyalin Tchh
1434426_at 1.19003 RIKEN cDNA B130055D15 gene B130055D15Rik
1434427_a_at 0.86281 ring finger protein 157 Rnf157
1434428_at 0.69557 RIKEN cDNA D330028D13 gene D330028D13Rik
1434429_at 1.01148 synaptotagmin XVI Syt16
1434430_s_at 0.90668 adenosine A2b receptor Adora2b
1434431_x_at 1.03944 adenosine A2b receptor Adora2b
1434432_at 0.83133 ring finger and FYVE like domain containing protein Rffl
1434433_x_at 1.07797 WD repeat domain 61 Wdr61
1434434_s_at 0.99923 transcription elongation regulator 1 (CA150) Tcerg1
1434435_s_at 0.90860 "cytochrome c oxidase, subunit XVII assembly protein homolog (yeast)" Cox17
1434436_at 0.95721 microrchidia 4 Morc4

1434437_x_at 0.91050 ribonucleotide reductase M2 Rrm2
1434438_at 1.00341 "SAM domain and HD domain, 1" Samhd1
1434439_at 1.04209 glycogen synthase kinase 3 beta Gsk3b
1434440_at 0.49752 "guanine nucleotide binding protein, alpha inhibiting 1" Gnai1
1434441_at 1.03161 RIKEN cDNA 1110018J18 gene 1110018J18Rik
1434442_at 0.79994 "DNA segment, Chr 5, ERATO Doi 593, expressed" D5Ertd593e
1434443_at 0.90611 --- ---
1434444_s_at 0.91822 --- ---
1434445_at 0.95123 "DNA segment, Chr 15, Wayne State University 169, expressed" D15Wsu169e
1434446_at 0.91595 RIKEN cDNA D630014A15 gene D630014A15Rik
1434447_at 1.05006 expressed sequence AI838057 AI838057
1434448_at 0.94592 taxilin alpha Txlna
1434449_at 1.01849 aquaporin 4 Aqp4
1434450_s_at 1.14287 "adrenergic receptor kinase, beta 2" Adrbk2
1434451_at 0.65630 "NOD-derived CD11c +ve dendritic cells cDNA, RIKEN full-length enriched library, clone:F630119A15 product:hypothetical protein, full insert sequence" ---
1434452_x_at 1.06642 eukaryotic translation initiation factor 2a Eif2a
1434453_at 1.04753 polymerase (RNA) III (DNA directed) polypeptide A Polr3a
1434454_at 0.84013 "DNA segment, Chr 16, Wayne State University 65, expressed" D16Wsu65e
1434455_at 0.83636 : F-box protein 44 Fbxo44
1434456_at 0.80351 "gene model 440, (NCBI)" Gm440
1434457_at 1.01266 nuclear antigen Sp100 Sp100
1434458_at 0.85547 Follistatin Fst
1434459_at 0.92776 cDNA sequence BC057627 BC057627
1434460_at 0.92177 Bardet-Biedl syndrome 4 homolog (human) Bbs4
1434461_at 0.95476 zinc finger protein 715 Zfp715
1434462_at 0.96944 cDNA sequence BC088983 BC088983
1434463_at 0.84491 "beaded filament structural protein 2, phakinin" Bfsp2
1434464_at 1.07016 aquaporin 12 Aqp12
1434465_x_at 1.00401 very low density lipoprotein receptor Vldlr
1434466_at 0.97966 "ataxia, cerebellar, Cayman type homolog (human)" Atcay
1434467_at 1.07966 "ataxia, cerebellar, Cayman type homolog (human)" Atcay
1434468_at 1.08203 OTU domain containing 4 Otud4
1434469_at 1.08376 OTU domain containing 4 Otud4
1434470_at 0.86434 synaptotagmin XIII Syt13
1434471_at 1.02330 cDNA sequence BC003331 BC003331
1434472_at 1.04305 dual specificity phosphatase 3 (vaccinia virus phosphatase VH1-related) Dusp3
1434473_at 0.98110 "solute carrier family 16 (monocarboxylic acid transporters), member 5" Slc16a5
1434474_at 0.85901 "ATP-binding cassette, sub-family A (ABC1), member 5" Abca5
1434475_at 0.92067 peptidyl-prolyl isomerase G (cyclophilin G) Ppig
1434476_at 0.92680 CREB regulated transcription coactivator 1 Crtc1
1434477_at 0.85461 headcase homolog (Drosophila) Heca
1434478_at 0.91839 headcase homolog (Drosophila) Heca

1434479_at 0.86240 "Procollagen, type V, alpha 1" Col5a1
1434480_at 0.99810 RIKEN cDNA 4930402E16 gene 4930402E16Rik
1434481_at 0.89352 RIKEN cDNA 4121402D02 gene 4121402D02Rik
1434482_at 1.02500 "DNA segment, Chr 4, ERATO Doi 22, expressed" D4Ertd22e
1434483_at 1.00839 ubiquitin specific peptidase 12 Usp12
1434484_at 1.02550 RIKEN cDNA 1100001G20 gene 1100001G20Rik
1434485_a_at 0.98401 UDP-glucose pyrophosphorylase 2 Ugp2
1434486_x_at 1.01942 UDP-glucose pyrophosphorylase 2 Ugp2
1434487_at 0.86349 myocyte enhancer factor 2D Mef2d
1434488_at 1.10068 ADP-ribosylation factor related protein 1 Arfrp1
1434489_at 0.93033 "engulfment and cell motility 3, ced-12 homolog (C. elegans)" Elmo3
1434490_at 0.94310 "scavenger receptor class F, member 1" Scarf1
1434491_a_at 0.99646 "cytochrome c oxidase, subunit VIc" Cox6c
1434492_at 0.97465 RIKEN cDNA A130022J15 gene A130022J15Rik
1434493_at 1.14126 RIKEN cDNA 1810022K09 gene 1810022K09Rik
1434494_at 0.90602 zygote arrest 1 Zar1
1434495_at 0.83587 zinc finger protein 278 Zfp278
1434496_at 1.12690 polo-like kinase 3 (Drosophila) Plk3
1434497_at 1.05724 --- ---
1434498_at 0.98110 "gene model 1673, (NCBI)" Gm1673
1434499_a_at 1.16649 lactate dehydrogenase B Ldhb
1434500_at 0.92099 tweety homolog 2 (Drosophila) Ttyh2
1434501_at 0.95247 --- ---
1434502_x_at 0.95384 "solute carrier family 4 (anion exchanger), member 1" Slc4a1
1434503_s_at 0.93249 lysosomal membrane glycoprotein 2 Lamp2
1434504_at 0.90345 "zinc finger, FYVE domain containing 28" Zfyve28
1434505_a_at 0.95006 --- ---
1434506_at 0.89057 AT rich interactive domain 2 (Arid-rfx like) Arid2
1434507_at 0.85955 aminopeptidase-like 1 Npepl1
1434508_at 1.05810 ubiquitin-conjugating enzyme E2Q (putative) Ube2q
1434509_at 0.97562 Rap guanine nucleotide exchange factor (GEF) 6 Rapgef6
1434510_at 0.92174 3'-phosphoadenosine 5'-phosphosulfate synthase 2 Paps2
1434511_at 0.93584 phosphorylase kinase beta Phkb
1434512_x_at 0.95611 "splicing factor, arginine/serine-rich 3 (SRp20) /// similar to Splicing factor, arginine/serine-rich 3 (Pre-mRNA splicing factor SRP20) (X16 protein)" Sfrs3 /// LOC632248
1434513_at 1.08365 ATPase type 13A3 Atp13a3
1434514_at 1.01318 RNA binding motif protein 15 Rbm15
1434515_at 0.81764 nuclear receptor coactivator 1 Ncoal
1434516_at 0.99018 phosphoserine-tRNA kinase Pstk
1434517_at 1.15026 WD repeat and FYVE domain containing 2 Wdfy2
1434518_at 1.03874 phosphorylase kinase alpha 2 Phka2
1434519_at 0.89554 dimethylarginine dimethylaminohydrolase 1 Ddah1
1434520_at 1.40818 "sterol-C5-desaturase (fungal ERG3, delta-5-desaturase) homolog (S.

cerevisae)" Sc5d
 1434521_at 1.03395 regulatory factor X domain containing 2 homolog (human) Rfxdc2
 1434522_at 0.90441 WD repeat domain 4 Wdr4
 1434523_x_at 1.01480 "eukaryotic translation initiation factor 3, subunit 6" Eif3s6
 1434524_at 1.09742 "eukaryotic translation initiation factor 2B, subunit 3" Eif2b3
 1434525_at 0.99944 protein kinase N3 Pkn3
 1434526_at 0.99570 abhydrolase domain containing 7 Abhd7
 1434527_at 0.97301 "NACHT, leucine rich repeat and PYD containing 4B" Nalp4b
 1434528_at 0.95339 alanine and arginine rich domain containing protein Aard
 1434529_x_at 0.93493 checkpoint with forkhead and ring finger domains Chfr
 1434530_at 1.10895 odd Oz/ten-m homolog 4 (Drosophila) Odz4
 1434531_at 1.04717 "mannoside acetylglucosaminyltransferase 5, isoenzyme B" Mgat5b
 1434532_at 0.82632 cDNA sequence BC035295 BC035295
 1434533_x_at 1.01284 CEA-related cell adhesion molecule 11 Ceacam11
 1434534_at 1.01454 --- ---
 1434535_at 1.09330 keratin 222 Krt222
 1434536_at 0.94630 G protein-coupled receptor 75 Gpr75
 1434537_at 0.92213 "solute carrier organic anion transporter family, member 3a1" Slco3a1
 1434538_x_at 1.03349 "eukaryotic translation initiation factor 2B, subunit 2 beta" Eif2b2
 1434539_at 0.97060 "leucine rich repeat protein 3, neuronal" Lrrn3
 1434540_a_at 1.09912 "clathrin, light polypeptide (Lca)" Clta
 1434541_x_at 1.01440 --- ---
 1434542_at 1.03129 glutamic pyruvate transaminase (alanine aminotransferase) 2 Gpt2
 1434543_a_at 0.98832 bolA-like 2 (E. coli) Bola2
 1434544_at 1.00954 bolA-like 2 (E. coli) Bola2
 1434545_x_at 0.96150 bolA-like 2 (E. coli) Bola2
 1434546_at 0.97149 "Smg-5 homolog, nonsense mediated mRNA decay factor (C. elegans)"
 Smg5
 1434547_at 0.96708 carboxypeptidase D Cpd
 1434548_at 0.91897 serine incorporator 3 Serinc3
 1434549_at 0.93660 --- ---
 1434550_at 0.96049 RIKEN cDNA 3830406C13 gene 3830406C13Rik
 1434551_at 0.89362 heterogeneous nuclear ribonucleoprotein U-like 2 Hnrpul2
 1434552_at 0.96283 WD repeat domain 77 Wdr77
 1434553_at 0.98070 transmembrane protein 56 Tmem56
 1434554_at 0.89129 tripartite motif protein 37 Trim37
 1434555_at 0.90175 "acidic (leucine-rich) nuclear phosphoprotein 32 family, member A" Anp32a
 1434556_at 1.34811 "Bone marrow macrophage cDNA, RIKEN full-length enriched library, clone:
 I830003K21 product:unclassifiable, full insert sequence" ---
 1434557_at 0.79329 huntingtin interacting protein 1 Hip1
 1434558_at 1.06162 WD repeat domain 47 Wdr47
 1434559_at 1.06203 syntaxin 3 Stx3
 1434560_at 0.98200 WD and tetratricopeptide repeats 1 Wdtd1

1434561_at 0.92462 additional sex combs like 1 (Drosophila) Asxl1
1434562_at 0.97032 microfibrillar-associated protein 3 Mfap3
1434563_at 0.94191 ribosomal protein S6 kinase polypeptide 1 Rps6kc1
1434564_at 1.19087 E2F transcription factor 3 E2f3
1434565_at 0.84983 cell growth regulator with ring finger domain 1 Cgrrf1
1434566_a_at 0.77288 RIKEN cDNA 4732496O08 gene 4732496O08Rik
1434567_at 0.87174 RIKEN cDNA 4732496O08 gene 4732496O08Rik
1434568_at 1.07031 Transcribed locus ---
1434569_at 0.93016 expressed sequence AA474455 AA474455
1434570_at 1.06177 cDNA sequence AK122525 AK122525
1434571_at 0.91190 vacuolar protein sorting 13B (yeast) /// similar to vacuolar protein sorting 13B isoform 4 /// region containing RIKEN cDNA C330002D13 gene; RIKEN cDNA 2310042E16 gene; vacuolar protein sorting 13B (yeast) Vps13b /// LOC666173 /// LOC669978
1434572_at 1.02762 histone deacetylase 9 Hdac9
1434573_at 0.96083 TRAF3 interacting protein 3 Traf3ip3
1434574_at 1.04186 RIKEN cDNA 9430008C03 gene 9430008C03Rik
1434575_at 0.94985 erythrocyte protein band 4.1-like 1 Epb4.111
1434576_at 0.94201 testis specific gene A14 Tsga14
1434577_at 1.03758 cDNA sequence BC052040 BC052040
1434578_x_at 1.06896 "RAN, member RAS oncogene family /// similar to RAN, member RAS oncogene family /// similar to RAN, member RAS oncogene family" Ran /// LOC625094 /// LOC671251
1434579_x_at 0.94543 NADH dehydrogenase (ubiquinone) Fe-S protein 8 Ndufs8
1434580_at 0.97117 ectonucleotide pyrophosphatase/phosphodiesterase 4 Enpp4
1434581_at 0.67813 --- ---
1434582_at 0.82823 "DNA segment, Chr 14, ERATO Doi 171, expressed" D14Ertd171e
1434583_at 0.98197 transmembrane protein 26 Tmem26
1434584_a_at 0.99293 preferentially expressed antigen in melanoma like 5 /// Unknown (protein for IMAGE:30033377) /// similar to preferentially expressed antigen in melanoma like 5 Pramel5 /// LOC384077 /// LOC674184
1434585_at 1.10346 RIKEN cDNA 2210038L17 gene 2210038L17Rik
1434586_a_at 0.88105 phosphatidylserine synthase 2 Ptdss2
1434587_x_at 0.97581 phosphatidylserine synthase 2 Ptdss2
1434588_x_at 0.97655 tubulin cofactor a /// similar to Tubulin-specific chaperone A (Tubulin-folding cofactor A) (CFA) (TCP1-chaperonin cofactor A) Tbca /// LOC620313
1434589_x_at 0.91882 surfeit gene 4 Surf4
1434590_at 0.82826 RIKEN cDNA B230209C24 gene B230209C24Rik
1434591_at 0.81402 RIKEN cDNA 4732460K03 gene 4732460K03Rik
1434592_at 0.89378 "solute carrier family 16 (monocarboxylic acid transporters), member 10" Slc16a10
1434593_at 0.96413 eukaryotic translation initiation factor 5A2 Eif5a2
1434594_at 0.97423 "dynein, axonemal, heavy chain 1" Dnahc1
1434595_at 0.99470 tripartite motif protein 9 Trim9
1434596_at 0.95224 "PREDICTED: Mus musculus hypothetical protein LOC625568 (LOC625568),

mRNA" ---

1434597_at 1.01075 "La ribonucleoprotein domain family, member 5" Larp5
 1434598_at 1.03318 "La ribonucleoprotein domain family, member 5" Larp5
 1434599_a_at 0.86049 tight junction protein 2 Tjp2
 1434600_at 0.89221 tight junction protein 2 Tjp2
 1434601_at 0.93539 adhesion molecule with Ig like domain 2 Amigo2
 1434602_at 0.96175 thyroid hormone receptor associated protein 2 Thrap2
 1434603_at 0.94619 thyroid hormone receptor associated protein 2 Thrap2
 1434604_at 1.09908 eukaryotic translation initiation factor 5B Eif5b
 1434605_at 1.08487 eukaryotic translation initiation factor 5B Eif5b
 1434606_at 0.83898 v-erb-b2 erythroblastic leukemia viral oncogene homolog 3 (avian) Erbb3
 1434607_at 0.98512 DEAD (Asp-Glu-Ala-Asp) box polypeptide 52 Ddx52
 1434608_at 1.05891 --- ---
 1434609_at 1.08721 RIKEN cDNA B930007L02 gene B930007L02Rik
 1434610_at 1.06100 plectin 1 Plec1
 1434611_at 1.01263 ring finger protein 123 Rnf123
 1434612_s_at 1.01983 "sno, strawberry notch homolog 1 (Drosophila)" Sbnol
 1434613_at 0.98514 RIKEN cDNA 1810013L24 gene 1810013L24Rik
 1434614_at 1.00168 --- ---
 1434615_x_at 0.85793 --- ---
 1434616_at 1.07521 RIKEN cDNA 1810073N04 gene 1810073N04Rik
 1434617_x_at 0.91022 RIKEN cDNA 1810073N04 gene 1810073N04Rik
 1434618_at 1.04074 --- ---
 1434619_at 0.93356 --- ---
 1434620_s_at 0.91155 RIKEN cDNA 2610024E20 gene 2610024E20Rik
 1434621_at 0.86583 cDNA sequence BC054438 BC054438
 1434622_at 0.91637 IQ motif containing E Iqce
 1434623_at 0.97996 ubiquitin A-52 residue ribosomal protein fusion product 1 /// similar to ubiquitin
 A-52 residue ribosomal protein fusion product 1 /// similar to ubiquitin A-52 residue ribosomal protein
 fusion product 1 /// similar to ubiquitin A-52 residue ribosomal protein fusion product 1 /// similar to
 ubiquitin A-52 residue ribosomal protein fusion product 1 /// similar to ubiquitin A-52 residue ribosomal
 protein fusion product 1 /// similar to ubiquitin A-52 residue ribosomal protein fusion product 1 ///
 similar to ubiquitin A-52 residue ribosomal protein fusion product 1 Uba52 /// LOC383341 ///
 LOC665823 /// LOC666586 /// LOC669193 /// LOC669699 /// LOC676355 /// LOC676687
 1434624_x_at 1.04957 ribosomal protein S9 Rps9
 1434625_at 0.84256 RIKEN cDNA 4930432O21 gene 4930432O21Rik
 1434626_at 0.96778 RNA pseudouridylate synthase domain containing 3 Rpusd3
 1434627_at 1.56810 Nuclear respiratory factor 1 Nrf1
 1434628_a_at 1.19667 "rhophilin, Rho GTPase binding protein 2" Rhpn2
 1434629_at 1.21433 zinc finger and BTB domain containing 26 Zbtb26
 1434630_at 1.05394 ankyrin repeat domain 28 Ankrd28
 1434631_at 0.90309 "DNA segment, Chr 3, ERATO Doi 751, expressed" D3Ert751e
 1434632_at 1.00716 Transcribed locus ---

1434633_at 0.83990 CREB binding protein Crebbp
1434634_at 0.89394 "gene model 166, (NCBI)" Gm166
1434635_at 0.87228 rabphilin 3A Rph3a
1434636_at 0.97248 homeodomain leucine zipper-encoding gene Homez
1434637_x_at 1.06869 "transcriptional regulator, SIN3B (yeast)" Sin3b
1434638_at 1.01513 "Adult male corpora quadrigemina cDNA, RIKEN full-length enriched library, clone:B230036G05 product:unclassifiable, full insert sequence" ---
1434639_at 0.91100 kelch repeat and BTB (POZ) domain containing 9 Kbtbd9
1434640_at 1.08415 neurensin 2 Nrsn2
1434641_x_at 1.00988 seizure related 6 homolog like 2 Sez6l2
1434642_at 0.94897 dehydrogenase/reductase (SDR family) member 8 Dhrs8
1434643_at 1.02889 transducin (beta)-like 1 X-linked Tbl1x
1434644_at 0.94359 transducin (beta)-like 1 X-linked Tbl1x
1434645_at 1.00230 RIKEN cDNA C530008M17 gene C530008M17Rik
1434646_s_at 1.08245 Sin3-associated polypeptide 18 Sap18
1434647_at 0.97451 expressed sequence AU040377 AU040377
1434648_a_at 0.96834 cerebral cavernous malformation 2 homolog (human) Ccm2
1434649_at 0.91332 cerebral cavernous malformation 2 homolog (human) Ccm2
1434650_at 0.88097 pogo transposable element with ZNF domain Pogz
1434651_a_at 0.87219 claudin 3 Cldn3
1434652_at 0.99580 Cdc42 binding protein kinase beta Cdc42bpb
1434653_at 0.82596 PTK2 protein tyrosine kinase 2 beta Ptk2b
1434654_at 1.01628 component of oligomeric golgi complex 3 Cog3
1434655_at 1.12522 forkhead box K1 Foxk1
1434656_at 1.02673 RIKEN cDNA B230339M05 gene B230339M05Rik
1434657_at 1.14104 glutaminase Glu
1434658_at 0.97893 RIKEN cDNA 3110056O03 gene 3110056O03Rik
1434659_at 0.95941 RIKEN cDNA 5830411G16 gene 5830411G16Rik
1434660_at 1.29684 "alkB, alkylation repair homolog 1 (E. coli)" Alkbh1
1434661_at 0.96301 synaptogyrin 1 Syng1
1434662_at 0.90268 autophagy-related 4A (yeast) /// similar to Cysteine protease ATG4A (Autophagy-related protein 4 homolog A) (Autophagin-2) (Autophagy-related cysteine endopeptidase 2) (AUT-like 2 cysteine endopeptidase) /// similar to Cysteine protease ATG4A (Autophagy-related protein 4 homolog A) (Autophagin-2) (Autophagy-related cysteine endopeptidase 2) (AUT-like 2 cysteine endopeptidase) Atg4a /// LOC666476 /// LOC675173
1434663_at 0.91415 RIKEN cDNA 2410129H14 gene 2410129H14Rik
1434664_at 0.95825 RIKEN cDNA 2410129H14 gene 2410129H14Rik
1434665_at 1.07626 aspartylglucosaminidase Aga
1434666_at 1.08166 polycomb group ring finger 5 Pcgf5
1434667_at 0.77848 "procollagen, type VIII, alpha 2" Col8a2
1434668_at 1.03453 nuclear casein kinase and cyclin-dependent kinase substrate 1 Nucks1
1434669_at 0.94997 Ral GEF with PH domain and SH3 binding motif 1 Ralgps1
1434670_at 0.94626 kinesin family member 5A Kif5a

1434671_at 0.91019 RIKEN cDNA B230337E12 gene B230337E12Rik
1434672_at 1.05455 G protein-coupled receptor 22 Gpr22
1434673_at 1.07444 G protein-coupled receptor 22 Gpr22
1434674_at 0.82826 lysosomal trafficking regulator Lyst
1434675_at 1.01337 RIKEN cDNA 1700065O13 gene 1700065O13Rik
1434676_at 0.94864 myotubularin related protein 9 Mtmr9
1434677_at 0.90178 Hermansky-Pudlak syndrome 5 homolog (human) Hps5
1434678_at 1.03124 muscleblind-like 3 (Drosophila) Mbnl3
1434679_at 0.98926 chondroitin sulfate proteoglycan 3 Cspg3
1434680_at 0.99802 "pleckstrin homology domain containing, family G (with RhoGef domain)
member 3" Plekhg3
1434681_at 1.13701 RIKEN cDNA 4932441K18 gene 4932441K18Rik
1434682_at 1.13091 RIKEN cDNA 6430601A21 gene 6430601A21Rik
1434683_at 0.74528 Cut-like 1 (Drosophila) Cutl1
1434684_at 0.96227 Ras and Rab interactor 3 Rin3
1434685_at 0.90711 "DNA segment, Chr 3, Brigham & Women's Genetics 0562 expressed"
D3Bwg0562e
1434686_at 1.00372 --- ---
1434687_at 0.88829 hypothetical protein C730026J16 C730026J16
1434688_x_at 0.88648 developmentally regulated GTP binding protein 1 Drg1
1434689_at 0.86162 zinc finger protein 637 Zfp637
1434690_at 0.89458 lysocardiolipin acyltransferase Lypcat
1434691_at 0.93284 "splicing factor, arginine/serine-rich 2, interacting protein" Sfrs2ip
1434692_at 1.11507 RIKEN cDNA 1110034B05 gene 1110034B05Rik
1434693_at 1.06478 "Adult male medulla oblongata cDNA, RIKEN full-length enriched library,
clone:6330441D15 product:hypothetical protein, full insert sequence" ---
1434694_at 0.98541 leucine rich repeat containing 8A Lrrc8a
1434695_at 0.98705 denticleless homolog (Drosophila) Dtl
1434696_at 1.00391 intraflagellar transport 52 homolog (Chlamydomonas) Ift52
1434697_at 0.87830 RIKEN cDNA 1110001P04 gene 1110001P04Rik
1434698_at 0.95389 RIKEN cDNA 9130229H14 gene 9130229H14Rik
1434699_at 1.03245 RIKEN cDNA 6030408C04 gene 6030408C04Rik
1434700_at 1.07414 RIKEN cDNA 6030408C04 gene 6030408C04Rik
1434701_at 1.12584 expressed sequence AW215868 AW215868
1434702_at 1.05347 RIKEN cDNA 2600009E05 gene 2600009E05Rik
1434703_at 1.01169 exostoses (multiple)-like 3 Extl3
1434704_at 0.93658 myeloid/lymphoid or mixed-lineage leukemia 5 Mll5
1434705_at 0.96204 "C-terminal binding protein 2 /// zinc finger, RAN-binding domain containing
1" Ctbp2 /// Zranb1
1434706_at 0.99013 valosin containing protein (p97)/p47 complex interacting protein 1 Vcpip1
1434707_at 1.00336 SET binding factor 1 Sbf1
1434708_at 1.10521 von Hippel-Lindau syndrome homolog Vhlh
1434709_at 1.02762 RIKEN cDNA C130076O07 gene C130076O07Rik

1434710_at 1.05268 DEAH (Asp-Glu-Ala-His) box polypeptide 29 Dhx29
1434711_at 0.96381 cDNA sequence BC030867 BC030867
1434712_at 0.97798 "Smith-Magenis syndrome chromosome region, candidate 7-like (human)"
Smcr7l
1434713_at 0.96158 "Smith-Magenis syndrome chromosome region, candidate 7-like (human)"
Smcr7l
1434714_at 1.04488 ERO1-like beta (*S. cerevisiae*) Ero1lb
1434715_at 1.04234 RIKEN cDNA 1600014C10 gene 1600014C10Rik
1434716_at 1.03959 hepatitis A virus cellular receptor 1 Havcr1
1434717_at 0.97996 cullin 3 Cul3
1434718_at 1.01638 cullin 3 Cul3
1434719_at 1.01762 alpha-2-macroglobulin /// hypothetical protein LOC677369 A2m ///
LOC677369
1434720_at 1.00913 RIKEN cDNA 9530033F24 gene 9530033F24Rik
1434721_at 1.01016 "DNA segment, Chr 5, ERATO Doi 585, expressed" D5Ert585e
1434722_at 0.88508 adenosine monophosphate deaminase 1 (isoform M) Ampd1
1434723_at 0.75800 expressed sequence AI426748 AI426748
1434724_at 1.03133 ubiquitin specific peptidase 31 Usp31
1434725_at 0.88712 GRAM domain containing 1C Gramd1c
1434726_at 0.99069 transmembrane protein 146 Tmem146
1434727_at 0.98250 RIKEN cDNA 2900042E01 gene 2900042E01Rik
1434728_at 0.92209 "glutamate receptor, ionotropic, AMPA3 (alpha 3)" Gria3
1434729_at 0.86883 cDNA sequence AK129302 AK129302
1434730_at 1.00183 expressed sequence AI854517 AI854517
1434731_x_at 1.01539 peroxiredoxin 1 Prdx1
1434732_x_at 0.86864 Translocase of outer mitochondrial membrane 7 homolog (yeast) Tomm7
1434733_at 0.82857 "serine/threonine kinase 36 (fused homolog, *Drosophila*)" Stk36
1434734_at 1.00370 RIKEN cDNA E130016E03 gene E130016E03Rik
1434735_at 0.92600 hepatic leukemia factor Hlf
1434736_at 0.72542 hepatic leukemia factor Hlf
1434737_at 0.97108 oligonucleotide/oligosaccharide-binding fold containing 1 Obfc1
1434738_at 0.86320 threonyl-tRNA synthetase-like 2 Tarsl2
1434739_at 1.13097 fragile X mental retardation 1 neighbor Fmr1nb
1434740_at 0.98136 "scavenger receptor class F, member 2" Scarf2
1434741_at 1.87641 ras responsive element binding protein 1 Rreb1
1434742_s_at 0.88435 RIKEN cDNA 2810401C16 gene 2810401C16Rik
1434743_x_at 0.92009 RUN and SH3 domain containing 1 Rusc1
1434744_at 1.00750 yrdC domain containing (*E.coli*) Yrdc
1434745_at 0.98045 cyclin D2 Ccnd2
1434746_at 1.09158 MAX gene associated Mga
1434747_at 0.95759 chymotrypsin C (caldecrin) Ctrc
1434748_at 1.07562 cytoskeleton associated protein 2 Ckap2
1434749_at 1.06576 cDNA sequence BC067068 BC067068

1434750_at	0.88067	exocyst complex component 3	Exoc3
1434751_at	0.75172	iduronate 2-sulfatase	Ids
1434752_at	1.84943	zinc finger protein 207	Zfp207
1434753_at	1.08596	nuclear factor related to kappa B binding protein	Nfrkb
1434754_at	0.60760	GTPase activating RANGAP domain-like 4	Garnl4
1434755_at	0.79258	"coronin, actin binding protein, 2B"	Coro2b
1434756_at	0.83165	hypothetical protein 5430421B17	5430421B17
1434757_at	0.89221	"core-binding factor, runt domain, alpha subunit 2, translocated to, 2 homolog (human)"	Cbfa2t2h
1434758_at	0.97027	cysteine-rich secretory protein LCCL domain containing 2	Crispld2
1434759_at	1.06437	leucine rich repeat transmembrane neuronal 3	Lrrtm3
1434760_at	0.92834	leucine rich repeat transmembrane neuronal 3	Lrrtm3
1434761_at	1.03200	leucine rich repeat transmembrane neuronal 3	Lrrtm3
1434762_at	1.02848	transmembrane protein 142B	Tmem142b
1434763_at	0.88701	transmembrane protein 142B	Tmem142b
1434764_at	0.93344	A kinase (PRKA) anchor protein 11	Akap11
1434765_at	1.00056	E1A binding protein p300	Ep300
1434766_at	0.70728	"Mus musculus, clone IMAGE:1512359, mRNA"	---
1434767_at	0.97705	expressed sequence C79407	C79407
1434768_at	1.03164	tripeptidyl peptidase I	Tpp1
1434769_at	0.92793	BTB (POZ) domain containing 9	Btbd9
1434770_at	0.81510	IQ calmodulin-binding motif containing 1	Iqcb1
1434771_at	0.95576	RIKEN cDNA 0610011F06 gene	0610011F06Rik
1434772_at	1.07593	adenosine A2b receptor	Adora2b
1434773_a_at	1.38598	"solute carrier family 2 (facilitated glucose transporter), member 1"	Slc2a1
1434774_at	0.97171	rhomboid 5 homolog 2 (Drosophila)	Rhbdf2
1434775_at	0.84128	par-3 (partitioning defective 3) homolog (C. elegans)	Pard3
1434776_at	0.96027	---	---
1434777_at	0.92367	"v-myc myelocytomatosis viral oncogene homolog 1, lung carcinoma derived (avian)"	Mycl1
1434778_at	1.00992	Transcribed locus	---
1434779_at	0.95338	cerebellin 2 precursor protein	Cbln2
1434780_at	0.91965	RIKEN cDNA C130038G02 gene	C130038G02Rik
1434781_at	0.97613	---	---
1434782_at	1.02720	ubiquitin specific peptidase 42	Usp42
1434783_at	0.86435	transmembrane protein 106C	Tmem106c
1434784_s_at	0.83415	transmembrane protein 106C	Tmem106c
1434785_at	1.00200	"calcium channel, voltage-dependent, gamma subunit 5"	Cacng5
1434786_at	1.01395	"protein phosphatase 1, regulatory (inhibitor) subunit 12B"	Ppp1r12b
1434787_at	1.07046	ADP-ribosylation factor 3	Arf3
1434788_at	1.06149	RIKEN cDNA D930050A07 gene	D930050A07Rik
1434789_at	1.04390	DEP domain containing 1B	Depdc1b
1434790_a_at	0.91336	leukotriene A4 hydrolase	Lta4h

1434791_at	0.89197	"ATPase, H ⁺ transporting, lysosomal V0 subunit A2"	Atp6v0a2
1434792_at	0.91938	RIKEN cDNA 2010320M18 gene	2010320M18Rik
1434793_at	0.95130	WD repeat domain 78	Wdr78
1434794_at	0.91401	"ras homolog gene family, member f"	Rhof
1434795_at	0.95480	dispatched homolog 1 (Drosophila)	Disp1
1434796_at	1.00499	vesicle-associated membrane protein 4	Vamp4
1434797_at	0.95510	RIKEN cDNA 6720469N11 gene	6720469N11Rik
1434798_at	1.06440	"ATPase, H ⁺ transporting, lysosomal V0 subunit D2"	Atp6v0d2
1434799_x_at	1.13239	"aldolase 1, A isoform"	Aldoa
1434800_at	0.88628	synaptic vesicle glycoprotein 2 b	Sv2b
1434801_x_at	1.04704	"solute carrier family 25 (mitochondrial carrier, adenine nucleotide translocator), member 5"	Slc25a5
1434802_s_at	0.97370	neurotrophin 3	Ntf3
1434803_a_at	0.86438	syncollin	Sycn
1434804_at	0.97521	SEC15-like 2 (S. cerevisiae)	Sec15l2
1434805_at	0.91095	myeloid/lymphoid or mixed lineage-leukemia translocation to 1 homolog (Drosophila)	Mllt1
1434806_at	0.84431	metaxin 3	Mtx3
1434807_s_at	0.85607	metaxin 3	Mtx3
1434808_at	1.19374	cDNA sequence BC066140	BC066140
1434809_at	1.05932	Rho GTPase activating protein 28	Arhgap28
1434810_a_at	1.06959	cancer antigen 1	Cage1
1434811_at	1.01785	cancer antigen 1	Cage1
1434812_s_at	0.88277	docking protein 7	Dok7
1434813_x_at	1.07845	tryptophanyl-tRNA synthetase	Wars
1434814_x_at	1.04225	glucose phosphate isomerase 1	Gpi1
1434815_a_at	1.05986	mitogen-activated protein kinase-activated protein kinase 3	Mapkapk3
1434816_at	0.97932	vacuolar protein sorting 33A (yeast)	Vps33a
1434817_s_at	0.98750	RIKEN cDNA 4930535B03 gene	4930535B03Rik
1434818_at	0.94712	hypothetical protein 9330159K06	9330159K06
1434819_at	0.98652	"beta galactoside alpha 2,6 sialyltransferase 2"	St6gal2
1434820_s_at	0.84605	"protein kinase inhibitor, gamma"	Pkig
1434821_at	0.97341	bromodomain containing 1	Brd1
1434822_at	0.99348	periphilin 1	Pphln1
1434823_x_at	0.87047	myeloma overexpressed 2	Myeov2
1434824_at	0.97267	"bromodomain adjacent to zinc finger domain, 1B"	Baz1b
1434825_at	0.77803	similar to CAGL79	LOC381742
1434826_at	0.92363	expressed sequence AI256775	AI256775
1434827_at	1.13064	THO complex 6 homolog (Drosophila)	Thoc6
1434828_at	0.99184	RIKEN cDNA B430201A12 gene	B430201A12Rik
1434829_at	1.07653	Casitas B-lineage lymphoma	Cbl
1434830_at	0.84665	MAX dimerization protein 1	Mxd1
1434831_a_at	0.99352	forkhead box O3a	Foxo3a

1434832_at 0.98803 forkhead box O3a Foxo3a
1434833_at 0.91083 Mitogen activated protein kinase kinase kinase 2 Map4k2
1434834_at 0.82854 suppressor of cytokine signaling 7 Socs7
1434835_at 1.01825 wings apart-like homolog (Drosophila) Wapal
1434836_at 0.94884 "nuclear factor of activated T-cells, cytoplasmic, calcineurin-dependent 2 interacting protein" Nfatc2ip
1434837_at 0.91570 mediator of DNA damage checkpoint 1 Mdc1
1434838_at 1.00595 similar to Potassium voltage-gated channel subfamily G member 2 (Voltage-gated potassium channel subunit Kv6.2) (Cardiac potassium channel subunit) LOC240444
1434839_s_at 0.92488 transducin (beta)-like 1X-linked receptor 1 Tbl1xr1
1434840_at 0.90219 HIV-1 Rev binding protein Hrb
1434841_at 1.17788 CDNA sequence BC010584 BC010584
1434842_s_at 1.03182 UPF3 regulator of nonsense transcripts homolog B (yeast) Upf3b
1434843_at 1.04230 RIKEN cDNA A430041B07 gene A430041B07Rik
1434844_at 0.92492 "hexosaminidase (glycosyl hydrolase family 20, catalytic domain) containing" Hexdc
1434845_at 1.01599 Expressed sequence AI195381 AI195381
1434846_at 1.01838 RIKEN cDNA 1700065A05 gene 1700065A05Rik
1434847_at 0.93672 --- ---
1434848_at 0.94690 Transcribed locus ---
1434849_at 1.00686 TSPY-like 2 Tspyl2
1434850_at 0.95388 IQ motif containing GTPase activating protein 3 Iqgap3
1434851_s_at 0.88509 crumbs homolog 3 (Drosophila) Crb3
1434852_at 1.06528 "phospholipase A2, group IIF" Pla2g2f
1434853_x_at 0.86588 "makorin, ring finger protein, 1" Mkrn1
1434854_a_at 0.98774 ribosomal protein S10 Rps10
1434855_at 1.16677 amidohydrolase domain containing 2 Amdhd2
1434856_at 0.90252 ankyrin repeat domain 44 Ankrd44
1434857_at 0.90442 RIKEN cDNA A230106D06 gene A230106D06Rik
1434858_x_at 1.13129 zinc finger protein 511 Zfp511
1434859_at 0.94651 uridine monophosphate synthetase Umps
1434860_at 0.83761 NMDA receptor-regulated gene 3 Narg3
1434861_at 0.96350 Transcribed locus ---
1434862_at 1.09376 fucosyltransferase 2 Fut2
1434863_at 0.97625 expressed sequence AU041129 AU041129
1434864_at 0.88587 non imprinted in Prader-Willi/Angelman syndrome 1 homolog (human) Nipa1
1434865_a_at 0.93606 exocyst complex component 7 Exoc7
1434866_x_at 0.86671 "carnitine palmitoyltransferase 1a, liver" Cpt1a
1434867_at 1.05787 "solute carrier family 4, sodium bicarbonate transporter-like, member 11" Slc4a11
1434868_at 0.99739 RIKEN cDNA 4933431E20 gene 4933431E20Rik
1434869_at 0.86767 tudor domain containing 3 Tdrd3

1434870_at 1.02614 RIKEN cDNA 2810004N23 gene 2810004N23Rik
 1434871_at 1.02138 "cytochrome c oxidase, subunit XVII assembly protein homolog (yeast)"
 Cox17
 1434872_x_at 0.95177 ribosomal protein L37 /// similar to ribosomal protein L37 Rpl37 ///
 LOC383438
 1434873_a_at 0.98706 "centaurin, beta 1" Centb1
 1434874_x_at 1.00742 "centaurin, beta 1" Centb1
 1434875_a_at 1.07637 high mobility group nucleosomal binding domain 3 Hmgn3
 1434876_at 1.05537 glycosyltransferase 8 domain containing 3 Glt8d3
 1434877_at 1.08325 neuronal pentraxin 1 Nptx1
 1434878_at 1.09414 "SLIT and NTRK-like family, member 4" Slitrk4
 1434879_at 1.07609 cell division cycle 34 homolog (S. cerevisiae) Cdc34
 1434880_at 1.02529 ets variant gene 6 (TEL oncogene) Etv6
 1434881_s_at 0.84594 potassium channel tetramerisation domain containing 12 Kctd12
 1434882_at 1.06508 Metadherin Mtdh
 1434883_at 1.11103 Metadherin Mtdh
 1434884_at 0.99187 Metadherin Mtdh
 1434885_at 0.79943 "SPT2, Suppressor of Ty, domain containing 1 (S. cerevisiae)" Spty2d1
 1434886_at 1.16037 --- ---
 1434887_at 0.94511 frequenin homolog (Drosophila) Freq
 1434888_a_at 0.86756 matrin 3 Matr3
 1434889_at 0.96786 "pleckstrin homology domain containing, family A member 7" Plekha7
 1434890_at 0.94154 optic atrophy 1 homolog (human) Opa1
 1434891_at 0.80851 prostaglandin F2 receptor negative regulator Ptgfrn
 1434892_x_at 0.90866 retinoblastoma binding protein 4 Rbbp4
 1434893_at 0.99154 "ATPase, Na⁺/K⁺ transporting, alpha 2 polypeptide" Atp1a2
 1434894_at 1.17129 zinc finger CCCH type containing 13 Zc3h13
 1434895_s_at 0.75115 "protein phosphatase 1, regulatory (inhibitor) subunit 13B" Ppp1r13b
 1434896_at 0.95201 "zinc finger protein 422, related sequence 1" Zfp422-rs1
 1434897_a_at 0.99651 "solute carrier family 25 (mitochondrial carrier, adenine nucleotide
 translocator), member 4" Slc25a4
 1434898_at 0.87693 trinucleotide repeat containing 6a Tnrc6a
 1434899_s_at 0.97999 trinucleotide repeat containing 6a Tnrc6a
 1434900_at 0.96387 MKL (megakaryoblastic leukemia)/myocardin-like 1 Mk11
 1434901_at 0.89544 "gene model 1103, (NCBI)" Gm1103
 1434902_at 0.77012 ring finger protein 157 Rnf157
 1434903_s_at 1.02505 interleukin 1 receptor-like 2 Il1rl2
 1434904_at 0.99058 human immunodeficiency virus type I enhancer binding protein 2 Hivep2
 1434905_at 0.93396 cDNA sequence BC064011 BC064011
 1434906_at 0.85817 RIKEN cDNA 0610005C13 gene 0610005C13Rik
 1434907_at 0.95288 TATA box binding protein-like 1 Tbp1l
 1434908_at 1.01514 expressed sequence AI480556 AI480556
 1434909_at 0.95267 Ras-related GTP binding D Rragd

1434910_at 0.88304 --- ---
1434911_s_at 1.02224 Rho GTPase activating protein 19 Arhgap19
1434912_at 1.12782 resistance to inhibitors of cholinesterase 3 homolog (C. elegans) Ric3
1434913_at 1.11784 3-hydroxymethyl-3-methylglutaryl-Coenzyme A lyase-like 1 Hmgcll1
1434914_at 0.77077 "RAB6B, member RAS oncogene family" Rab6b
1434915_s_at 1.06190 leucine rich repeat containing 19 Lrrc19
1434916_at 0.89608 "vitamin K epoxide reductase complex, subunit 1-like 1" Vkorc111
1434917_at 0.84156 cordon-bleu Cobl
1434918_at 0.96877 SRY-box containing gene 6 Sox6
1434919_at 1.06530 Transcribed locus ---
1434920_a_at 0.82506 Ena-vasodilator stimulated phosphoprotein Evl
1434921_at 0.93752 "nuclear receptor subfamily 2, group E, member 1" Nr2e1
1434922_at 1.04386 PHD finger protein 12 Phf12
1434923_at 0.89000 RIKEN cDNA 2810437L13 gene 2810437L13Rik
1434924_at 0.98940 PHD finger protein 2 Phf2
1434925_at 0.92099 RIKEN cDNA 4833424O15 gene 4833424O15Rik
1434926_at 0.99160 Rho guanine nucleotide exchange factor (GEF) 11 /// similar to Rho guanine nucleotide exchange factor (GEF) 11 Arhgef11 /// LOC673478
1434927_at 0.94089 "heat shock protein family, member 7 (cardiovascular)" Hspb7
1434928_at 1.03706 growth arrest-specific 2 like 1 Gas2l1
1434929_at 0.97975 cDNA sequence BC035044 BC035044
1434930_at 0.85395 two pore channel 1 Tpcn1
1434931_at 0.78269 neogenin Neol
1434932_at 0.83313 "adenosine deaminase, RNA-specific, B1" Adarb1
1434933_at 1.09357 RIKEN cDNA 5730557L09 gene 5730557L09Rik
1434934_at 0.91079 ATP synthase mitochondrial F1 complex assembly factor 1 Atpaf1
1434935_at 1.01141 hypothetical protein LOC637741 LOC637741
1434936_at 1.15036 HIRA interacting protein 3 Hirip3
1434937_at 0.90589 "pam, highwire, rpm 1" Phr1
1434938_at 0.97082 RNA binding motif protein 9 Rbm9
1434939_at 0.95171 forkhead box F1a Foxf1a
1434940_x_at 1.13132 regulator of G-protein signaling 19 Rgs19
1434941_s_at 1.13078 RIKEN cDNA 2610101J03 gene 2610101J03Rik
1434942_at 1.07638 RIKEN cDNA 2610101J03 gene 2610101J03Rik
1434943_at 0.97930 cDNA sequence BC023055 BC023055
1434944_at 0.80830 dystrophin myotonic-protein kinase Dmpk
1434945_at 0.96666 acyltransferase like 1 Ayt1l
1434946_at 1.09503 RIKEN cDNA C330021A05 gene C330021A05Rik
1434947_at 0.81216 kinesin family member 3C Kif3c
1434948_at 0.87524 RAB11 family interacting protein 2 (class I) Rab11fip2
1434949_at 0.89981 armadillo repeat containing 8 Armc8
1434950_a_at 0.94909 armadillo repeat containing 8 Armc8
1434951_at 0.85543 armadillo repeat containing 8 Armc8

1434952_at 0.81249 COX4 neighbor Cox4nb
1434953_at 0.91632 "Adult male pituitary gland cDNA, RIKEN full-length enriched library, clone:5330439J20 product:unclassifiable, full insert sequence" ---
1434954_at 1.09183 --- ---
1434955_at 1.00442 membrane-associated ring finger (C3HC4) 1 1-Mar
1434956_at 0.92198 similar to ring finger protein 170 /// similar to ring finger protein 170
LOC668047 /// LOC676342
1434957_at 0.79371 cell adhesion molecule-related/down-regulated by oncogenes Cdon
1434958_at 0.96603 saccin Sacs
1434959_at 0.89535 desert hedgehog Dhh
1434960_at 0.97566 "TAF9B RNA polymerase II, TATA box binding protein (TBP)-associated factor" Taf9b
1434961_at 0.88387 ankyrin repeat and SOCS box-containing protein 1 Asb1
1434962_x_at 0.90358 chemokine (C-C motif) ligand 27 Ccl27
1434963_at 1.16383 suppressor of Ty 3 homolog (S. cerevisiae) Supt3h
1434964_at 0.90990 expressed sequence AA408420 AA408420
1434965_at 1.02020 "DNA segment, Chr 19, ERATO Doi 386, expressed" D19Ertd386e
1434966_at 1.04112 "splicing factor, arginine/serine-rich 8" Sfrs8
1434967_at 1.21648 "zinc finger, SWIM domain containing 6" Zswim6
1434968_a_at 0.96125 ARP3 actin-related protein 3 homolog (yeast) Actr3
1434969_at 0.83130 "bruno-like 5, RNA binding protein (Drosophila)" Brunol5
1434970_a_at 0.97564 --- ---
1434971_x_at 0.97942 --- ---
1434972_x_at 0.98694 "splicing factor, arginine/serine-rich 1 (ASF/SF2)" Sfrs1
1434973_at 0.94518 carbonic anhydrase 7 Car7
1434974_at 2.29609 "pyruvate dehydrogenase kinase, isoenzyme 1" Pdk1
1434975_x_at 0.86176 RIKEN cDNA 4933439C20 gene 4933439C20Rik
1434976_x_at 1.12831 eukaryotic translation initiation factor 4E binding protein 1 Eif4ebp1
1434977_at 0.85243 RIKEN cDNA 4933403F05 gene 4933403F05Rik
1434978_at 0.96734 RIKEN cDNA 4933403F05 gene 4933403F05Rik
1434979_at 0.99424 RIKEN cDNA 4933403F05 gene 4933403F05Rik
1434980_at 0.99179 "phosphoinositide-3-kinase, regulatory subunit 5, p101" Pik3r5
1434981_at 0.99808 RIKEN cDNA E130303B06 gene E130303B06Rik
1434982_at 0.77148 Ring finger protein 182 Rnf182
1434983_at 0.86309 RIKEN cDNA B230363K08 gene B230363K08Rik
1434984_at 0.91105 RIKEN cDNA 6330514A18 gene 6330514A18Rik
1434985_a_at 1.03508 eukaryotic translation initiation factor 4A1 Eif4a1
1434986_a_at 0.95728 Sec61 alpha 1 subunit (S. cerevisiae) Sec61a1
1434987_at 0.77668 "aldehyde dehydrogenase 2, mitochondrial" Aldh2
1434988_x_at 0.95397 "aldehyde dehydrogenase 2, mitochondrial" Aldh2
1434989_at 1.03443 cDNA sequence BC042720 BC042720
1434990_at 0.96668 protein phosphatase 1E (PP2C domain containing) Ppm1e
1434991_at 0.82074 F-box and WD-40 domain protein 17 Fbxw17

1434992_at	0.87286	RIKEN cDNA 9130206N08 gene	9130206N08Rik
1434993_at	1.02203	RIKEN cDNA B830045N13 gene	B830045N13Rik
1434994_at	0.88547	death effector domain-containing	Dedd
1434995_s_at	0.95656	death effector domain-containing	Dedd
1434996_at	0.98457	"solute carrier family 25 (mitochondrial carrier, Graves disease autoantigen), member 16"	Slc25a16
1434997_at	0.90960	cell division cycle 2-like 6 (CDK8-like)	Cdc216
1434998_at	1.01867	IQ motif containing GTPase activating protein 1	Iqgap1
1434999_at	1.00153	suppressor of variegation 4-20 homolog 1 (Drosophila)	Suv420h1
1435000_at	1.05339	G1 to S phase transition 1	Gspt1
1435001_at	0.97075	"phospholipase A2, activating protein"	Plaa
1435002_at	1.08183	zinc finger protein 507	zfp507
1435003_at	1.07057	"phosphatidylinositol 4-kinase, catalytic, alpha polypeptide"	Pik4ca
1435004_at	0.90344	pantothenate kinase 4	Pank4
1435005_at	1.00925	centromere protein E	Cenpe
1435006_s_at	0.98877	"ATP-binding cassette, sub-family B (MDR/TAP), member 7"	Abcb7
1435007_s_at	1.02148	expressed sequence AI132487	AI132487
1435008_at	0.87851	"solute carrier family 9 (sodium/hydrogen exchanger), isoform 6"	Slc9a6
1435009_at	1.00150	"solute carrier family 9 (sodium/hydrogen exchanger), isoform 6"	Slc9a6
1435010_at	0.86564	ankyrin repeat and SOCS box-containing protein 7	Asb7
1435011_x_at	0.99083	"aldo-keto reductase family 1, member A4 (aldehyde reductase)"	Akr1a4
1435012_x_at	1.10710	"elastase 3, pancreatic /// similar to elastase 3B, pancreatic"	Ela3 ///
LOC638418			
1435013_at	1.03440	HEAT repeat containing 2	Heatr2
1435014_at	1.03139	"RAB39B, member RAS oncogene family"	Rab39b
1435015_at	1.02259	RIKEN cDNA 2210018M03 gene	2210018M03Rik
1435016_at	0.91795	"trafficking protein, kinesin binding 2"	Trak2
1435017_at	0.99041	melanoma nuclear protein 13	Mel13
1435018_at	0.91014	RIKEN cDNA 5930434B04 gene	5930434B04Rik
1435019_at	0.94204	ataxin 7-like 3	Atxn7l3
1435020_at	0.94993	serologically defined colon cancer antigen 1	Sdccag1
1435021_at	0.97761	"gamma-aminobutyric acid (GABA-A) receptor, subunit beta 3"	Gabrb3
1435022_at	0.85260	cartilage homeo protein 1	Cart1
1435023_at	1.02645	intersectin 2	Itsn2
1435024_at	0.83526	RIKEN cDNA 4931428F04 gene	4931428F04Rik
1435025_at	0.99974	2-oxoglutarate and iron-dependent oxygenase domain containing 1	Ogfod1
1435026_at	0.87267	"sparc/osteonectin, cwcv and kazal-like domains proteoglycan 2"	Spock2
1435027_at	0.88793	"golgi autoantigen, golgin subfamily a, 1"	Golga1
1435028_at	0.91854	WD repeat domain 7	Wdr7
1435029_at	0.87101	---	---
1435030_at	1.05925	UPF2 regulator of nonsense transcripts homolog (yeast)	Upf2
1435031_at	0.98513	RIKEN cDNA 2010310D06 gene	2010310D06Rik
1435032_at	0.88470	"golgi autoantigen, golgin subfamily b, macrogolgin 1"	Golgb1

1435033_at 1.07772 Rho guanine nucleotide exchange factor (GEF) 4 Arhgef4
1435034_at 1.16425 expressed sequence AW060207 AW060207
1435035_at 1.09356 RNA (guanine-9-) methyltransferase domain containing 2 Rg9mtd2
1435036_at 0.94416 RIKEN cDNA A530050D06 gene A530050D06Rik
1435037_at 1.06988 per1-like domain containing 1 Perld1
1435038_s_at 1.01114 AP2 associated kinase 1 Aak1
1435039_a_at 0.96081 "Phosphatidylinositol-4-phosphate 5-kinase, type 1 beta" Pip5k1b
1435040_at 1.00047 interleukin-1 receptor-associated kinase 3 Irak3
1435041_at 0.91386 "myosin, light polypeptide 6, alkali, smooth muscle and non-muscle" Myl6
1435042_at 1.00228 RIKEN cDNA 9130004C02 gene 9130004C02Rik
1435043_at 1.01246 "phospholipase C, beta 1" Plcb1
1435044_at 0.94766 early B-cell factor 4 Ebf4
1435045_s_at 0.92911 mitogen-activated protein kinase 8 interacting protein 2 Mapk8ip2
1435046_at 1.12246 Myosin ID Myo1d
1435047_at 0.98408 Transcribed locus ---
1435048_at 0.95438 expressed sequence AI854703 AI854703
1435049_s_at 0.95250 expressed sequence AI854703 AI854703
1435050_at 0.70339 "DNA segment, Chr 10, Brigham & Women's Genetics 1379 expressed"
D10Bwg1379e
1435051_at 0.99660 WD repeat domain 44 Wdr44
1435052_at 0.88101 RIKEN cDNA 4930455F23 gene 4930455F23Rik
1435053_s_at 1.12960 "pleckstrin homology domain containing, family H (with MyTH4 domain)
member 1" Plekhh1
1435054_at 1.04164 essential meiotic endonuclease 1 homolog 1 (S. pombe) Eme1
1435055_a_at 1.09475 target of myb1 homolog (chicken) /// similar to TOM1 Tom1 /// LOC545878
1435056_x_at 0.82327 protein O-fucosyltransferase 2 Pofut2
1435057_x_at 0.90681 polymerase (RNA) I polypeptide E Polr1e
1435058_x_at 1.02371 syntaxin binding protein 3A Stxbp3a
1435059_at 0.98727 development and differentiation enhancing Ddef1
1435060_at 0.96965 tropomodulin 2 Tmod2
1435061_at 0.96709 nudix (nucleoside diphosphate linked moiety X)-type motif 10 Nudt10
1435062_at 0.88153 RIKEN cDNA 1500001A10 gene 1500001A10Rik
1435063_at 1.00937 "similar to CG8312-PA, isoform A" LOC625638
1435064_a_at 1.05293 transmembrane protein 27 Tmem27
1435065_x_at 1.02280 Vav2 oncogene Vav2
1435066_at 0.89664 "phosphatidylinositol transfer protein, cytoplasmic 1" Pitpnc1
1435067_at 0.97521 RIKEN cDNA B230208H17 gene B230208H17Rik
1435068_at 1.04167 "phosphatidylinositol-4-phosphate 5-kinase, type II, beta" Pip5k2b
1435069_at 0.95072 cDNA sequence BC064078 BC064078
1435070_at 0.95594 AE binding protein 2 Aebp2
1435071_at 0.84289 "zinc finger, FYVE domain containing 1" Zfyve1
1435072_at 0.94494 "zinc finger, FYVE domain containing 1" Zfyve1
1435073_a_at 0.95908 armadillo repeat containing 9 Armc9

1435074_at 0.90108 transmembrane protein 106B Tmem106b
1435075_at 0.82507 transmembrane protein 106B Tmem106b
1435076_at 1.11152 RIKEN cDNA 2310047D13 gene 2310047D13Rik
1435077_at 0.98125 additional sex combs like 1 (Drosophila) Asx11
1435078_at 0.84555 "tetratricopeptide repeat, ankyrin repeat and coiled-coil containing 2" Tanc2
1435079_at 1.01571 RIKEN cDNA 5730406M06 gene 5730406M06Rik
1435080_x_at 1.13042 RIKEN cDNA 5730406M06 gene 5730406M06Rik
1435081_at 1.06398 synaptophysin-like protein Sypl
1435082_at 1.07961 synaptophysin-like protein Sypl
1435083_at 0.78781 cortexin 1 Ctxn1
1435084_at 0.99896 RIKEN cDNA C730049O14 gene C730049O14Rik
1435085_at 0.90281 cAMP responsive element binding protein-like 2 Crebl2
1435086_s_at 0.92389 kelch domain containing 2 Klhdc2
1435087_at 0.87865 cDNA sequence BC039093 BC039093
1435088_at 0.99027 nuclear receptor-binding SET-domain protein 1 Nsd1
1435089_at 1.09944 RIKEN cDNA 2010111I01 gene 2010111I01Rik
1435090_at 0.95598 "tubulin, gamma complex associated protein 6" Tubgcp6
1435091_at 1.06984 zinc finger protein 568 Zfp568
1435092_at 0.99234 ADP-ribosylation factor-like 4A Arl4a
1435093_at 0.88370 "zinc finger, FYVE domain containing 20" Zfyve20
1435094_at 0.95174 "potassium inwardly-rectifying channel, subfamily J, member 16" Kcnj16
1435095_at 0.93677 RIKEN cDNA C030009O12 gene C030009O12Rik
1435096_at 0.91718 resistance to inhibitors of cholinesterase 8 homolog B (C. elegans) Ric8b
1435097_at 1.10835 methylmalonic aciduria (cobalamin deficiency) type B homolog (human)
Mmab
1435098_at 1.09067 similar to Na+ dependent glucose transporter 1 LOC213332
1435099_at 1.09625 "UTP14, U3 small nucleolar ribonucleoprotein, homolog A (yeast)" Utp14a
1435100_at 1.02794 RIKEN cDNA A130042E20 gene A130042E20Rik
1435101_at 0.94683 "Der1-like domain family, member 2" Derl2
1435102_a_at 0.94324 syntaxin 18 Stx18
1435103_x_at 1.03627 "phenylalanine-tRNA synthetase-like, beta subunit" Farslb
1435104_at 1.08998 RIKEN cDNA D130004H04 gene D130004H04Rik
1435105_at 0.94901 RIKEN cDNA 1110061N23 gene 1110061N23Rik
1435106_at 0.83584 RIKEN cDNA 3732412D22 gene 3732412D22Rik
1435107_at 1.01371 expressed sequence AW456874 AW456874
1435108_at 1.06429 Rho GTPase activating protein 22 Arhgap22
1435109_at 0.78717 RIKEN cDNA 0710007G10 gene /// RIKEN cDNA 3010001K23 gene
0710007G10Rik /// 3010001K23Rik
1435110_at 0.84824 unc-5 homolog B (C. elegans) Unc5b
1435111_at 0.97930 RIKEN cDNA 2310011E23 gene 2310011E23Rik
1435112_a_at 0.96283 "ATP synthase, H+ transporting, mitochondrial F0 complex, subunit d" Atp5h
1435113_x_at 0.95670 stathmin-like 3 Stmn3
1435114_at 0.92977 WD repeat and HMG-box DNA binding protein 1 Wdhd1

1435115_at 1.12337 fibronectin type III domain containing 5 Fndc5
1435116_at 0.93603 RIKEN cDNA 4933403G14 gene 4933403G14Rik
1435117_a_at 0.88825 RIKEN cDNA C330021A05 gene C330021A05Rik
1435118_at 0.96184 "tRNA splicing endonuclease 2 homolog (SEN2, *S. cerevisiae*)" Tsen2
1435119_at 1.11745 "Adult male medulla oblongata cDNA, RIKEN full-length enriched library, clone:6330578N16 product:unclassifiable, full insert sequence" ---
1435120_at 1.04287 Transcribed locus ---
1435121_at 1.02223 "deiodinase, iodothyronine type III, antisense" Dio3as
1435122_x_at 0.99412 DNA methyltransferase (cytosine-5) 1 Dnmt1
1435123_at 1.03121 RIKEN cDNA C030014M07 gene /// similar to RIKEN cDNA C920006C10 C030014M07Rik /// LOC668212
1435124_at 1.04188 hypothetical gene supported by AK045595 LOC328644
1435125_at 0.98516 expressed sequence BB120497 BB120497
1435126_at 0.92463 Dual specificity phosphatase-like 15 Dusp15
1435127_a_at 1.05323 O-sialoglycoprotein endopeptidase-like 1 Osgepl1
1435128_at 1.05275 brain-specific angiogenesis inhibitor 1-associated protein 2 Baiap2
1435129_at 0.88106 Protein tyrosine phosphatase 4a2 Ptp4a2
1435130_at 1.02203 similar to RIKEN cDNA 1810036I24 LOC383103
1435131_at 0.96261 zinc finger protein 13 Zfp13
1435132_at 1.13849 dispatched homolog 1 (*Drosophila*) Disp1
1435133_at 1.07307 UDP-glucose ceramide glucosyltransferase Ugcg
1435134_at 0.95020 arylacetamide deacetylase-like 1 Aadacl1
1435135_at 0.90880 arylacetamide deacetylase-like 1 Aadacl1
1435136_at 1.04790 Wolf-Hirschhorn syndrome candidate 1 (human) Whsc1
1435137_s_at 1.32287 RIKEN cDNA 1200015M12 gene /// RIKEN cDNA 1200016E24 gene /// RIKEN cDNA A130040M12 gene /// RIKEN cDNA E430024C06 gene 1200015M12Rik /// 1200016E24Rik /// A130040M12Rik /// E430024C06Rik
1435138_at 0.92102 transmembrane protein 28 Tmem28
1435139_at 1.08208 NMDA receptor-regulated gene 1 Narg1
1435140_at 0.99501 insulin degrading enzyme Ide
1435141_at 0.94398 SFT2 domain containing 2 Sft2d2
1435142_at 0.98972 SFT2 domain containing 2 Sft2d2
1435143_at 1.08926 PCTAIRE-motif protein kinase 2 Pctk2
1435144_at 1.00124 Prohibitin Phb
1435145_at 0.85228 "immunoglobulin superfamily, member 4" Igsf4d
1435146_s_at 0.79017 "immunoglobulin superfamily, member 4" Igsf4d
1435147_x_at 0.93548 "immunoglobulin superfamily, member 4" Igsf4d
1435148_at 0.95283 "ATPase, Na⁺/K⁺ transporting, beta 2 polypeptide" Atp1b2
1435149_at 0.91136 "phospholipase C, gamma 1" Plcg1
1435150_at 0.90150 "2 days neonate sympathetic ganglion cDNA, RIKEN full-length enriched library, clone:7120492P10 product:hypothetical protein, full insert sequence" ---
1435151_a_at 1.03195 ribosomal protein S3 Rps3
1435152_at 1.29799 leukocyte receptor cluster (LRC) member 8 Leng8

1435153_at 0.83510 BTB (POZ) domain containing 6 Btbd6
 1435154_at 1.03228 "similar to solute carrier family 7 (cationic amino acid transporter, y+ system), member 3" LOC245128
 1435155_at 1.28799 cingulin Cgn
 1435156_at 0.88256 cDNA sequence BC046331 BC046331
 1435157_at 0.87346 RIKEN cDNA 5830454D03 gene 5830454D03Rik
 1435158_at 0.93202 RNA binding motif protein 12B Rbm12b
 1435159_at 1.20010 RIKEN cDNA 9430038I01 gene 9430038I01Rik
 1435160_at 0.97607 RIKEN cDNA 1110064P04 gene 1110064P04Rik
 1435161_at 0.96014 "DNA segment, Chr 5, ERATO Doi 135, expressed" D5Ertd135e
 1435162_at 1.74801 "protein kinase, cGMP-dependent, type II" Prkg2
 1435163_at 1.18283 RIKEN cDNA 9030612M13 gene 9030612M13Rik
 1435164_s_at 0.98761 ubiquitin-activating enzyme E1C Ube1c
 1435165_at 0.98045 contactin 2 Cntn2
 1435166_at 1.12814 contactin 2 Cntn2
 1435167_at 0.76218 RAN binding protein 6 Ranbp6
 1435168_at 0.98420 zinc finger protein 710 Zfp710
 1435169_at 1.02965 RIKEN cDNA A930001N09 gene A930001N09Rik
 1435170_at 0.96539 "TSR2, 20S rRNA accumulation, homolog (*S. cerevisiae*)" Tsr2
 1435171_at 0.62077 RIKEN cDNA 2810416G20 gene 2810416G20Rik
 1435172_at 1.01598 eomesodermin homolog (*Xenopus laevis*) Eomes
 1435173_at 0.89165 arginine-tRNA-protein transferase 1 Ate1
 1435174_at 1.23711 "rosbin, round spermatid basic protein 1" Rsb1
 1435175_at 1.10699 cDNA sequence BC034090 BC034090
 1435176_a_at 0.71745 inhibitor of DNA binding 2 Id2
 1435177_a_at 0.93020 anaphase-promoting complex subunit 5 Anapc5
 1435178_x_at 0.91734 anaphase-promoting complex subunit 5 Anapc5
 1435179_at 0.89989 RIKEN cDNA C130071C03 gene C130071C03Rik
 1435180_at 1.13655 podocan Podn
 1435181_at 1.04415 expressed sequence AI461788 AI461788
 1435182_at 0.90036 RIKEN cDNA C530028I08 gene C530028I08Rik
 1435183_at 0.96559 TBK1 binding protein 1 Tbkbp1
 1435184_at 0.96337 natriuretic peptide receptor 3 Npr3
 1435185_at 0.93906 leucine rich repeat and coiled-coil domain containing 1 Lrrcc1
 1435186_s_at 0.89992 leucine rich repeat and coiled-coil domain containing 1 Lrrcc1
 1435187_at 0.85553 Translocase of outer mitochondrial membrane 20 homolog (yeast) Tomm20
 1435188_at 1.99016 "gene model 129, (NCBI)" Gm129
 1435189_at 1.06249 FERM and PDZ domain containing 1 /// similar to FERM and PDZ domain containing 1 /// similar to FERM and PDZ domain containing 1 Frmpd1 /// LOC669360 /// LOC674439
 1435190_at 0.87520 cell adhesion molecule with homology to L1CAM Chl1
 1435191_at 0.74408 corneodesmosin Cdsn
 1435192_at 0.95144 similar to SRY (sex determining region Y)-box 3 LOC631901
 1435193_at 0.94065 RIKEN cDNA A230050P20 gene A230050P20Rik

1435194_at 0.82128 --- ---
1435195_at 0.89612 vasohibin 1 Vash1
1435196_at 0.88108 "neurotrophic tyrosine kinase, receptor, type 2" Ntrk2
1435197_at 0.98796 "POU domain, class 3, transcription factor 3" Pou3f3
1435198_at 1.05176 hypothetical protein LOC545228 LOC545228
1435199_at 0.87570 --- ---
1435200_at 1.02317 RIKEN cDNA 6330419J24 gene 6330419J24Rik
1435201_at 0.93620 RIKEN cDNA 0610009B10 gene 0610009B10Rik
1435202_at 0.93218 zinc finger protein 574 Zfp574
1435203_at 1.02148 "mannosidase 2, alpha 2" Man2a2
1435204_at 0.99062 protein arginine N-methyltransferase 8 Prmt8
1435205_at 0.89096 "transcription factor AP-2, epsilon" Tcfap2e
1435206_at 1.05343 "solute carrier family 24 (sodium/potassium/calcium exchanger), member 4"
Slc24a4
1435207_at 0.81317 DIX domain containing 1 Dixdc1
1435208_at 1.11243 deltex 3-like (Drosophila) Dtx3l
1435209_at 1.01760 cDNA sequence BC057079 BC057079
1435210_s_at 0.97857 syntaphilin Snph
1435211_at 1.00516 tetratricopeptide repeat domain 12 /// similar to tetratricopeptide repeat domain
12 Ttc12 /// LOC640389
1435212_at 1.15765 "purinergic receptor P2X, ligand-gated ion channel, 2" P2rx2
1435213_at 0.97107 NHL repeat containing 1 Nhlrc1
1435214_at 0.95630 gap junction membrane channel protein alpha 12 Gja12
1435215_at 0.81755 AP1 gamma subunit binding protein 1 Ap1gbp1
1435216_a_at 1.07900 outer dense fiber of sperm tails 2 Odf2
1435217_at 0.92918 hypothetical protein LOC666185 /// hypothetical protein LOC666222 ///
hypothetical protein LOC673030 LOC666185 /// LOC666222 /// LOC673030
1435218_at 0.99149 "RasGEF domain family, member 1A" Rasgef1a
1435219_x_at 0.93895 "beclin 1 (coiled-coil, myosin-like BCL2-interacting protein)" Becn1
1435220_s_at 1.05099 CDC42 small effector 2 Cdc42se2
1435221_at 0.89149 "Adult male corpora quadrigemina cDNA, RIKEN full-length enriched library,
clone:B230341P20 product:inferred: forkhead box P1, full insert sequence" ---
1435222_at 0.89175 forkhead box P1 Foxp1
1435223_at 0.90155 "SPFH domain family, member 2" Spfh2
1435224_at 0.95928 CREB binding protein Crebbp
1435225_s_at 0.91180 "bromodomain and PHD finger containing, 3" Brpf3
1435226_at 1.03062 IBR domain containing 3 Ibrdc3
1435227_at 1.04090 --- ---
1435228_at 0.79101 cDNA sequence BC023829 BC023829
1435229_at 0.86850 GRAM domain containing 1B Gramd1b
1435230_at 1.12074 ankyrin repeat domain 12 Ankrd12
1435231_at 1.03791 coenzyme Q4 homolog (yeast) Coq4
1435232_x_at 0.97269 --- ---

1435233_at	0.90051	nuclear receptor coactivator 2	Ncoa2
1435234_at	0.97915	nuclear receptor coactivator 2	Ncoa2
1435235_at	0.93134	thioredoxin-like 1	Txn1l
1435236_at	0.92492	RIKEN cDNA A630018P17 gene	A630018P17Rik
1435237_at	0.98149	RIKEN cDNA 2310009A05 gene	2310009A05Rik
1435238_x_at	1.04188	RIKEN cDNA 2310009A05 gene	2310009A05Rik
1435239_at	1.02235	"glutamate receptor, ionotropic, AMPA1 (alpha 1)"	Gria1
1435240_at	0.81809	"bromodomain adjacent to zinc finger domain, 2B"	Baz2b
1435241_at	1.00430	RIKEN cDNA D930036F22 gene	D930036F22Rik
1435242_at	0.87237	androgen-induced proliferation inhibitor	Aprin
1435243_at	0.92459	zinc finger protein 746	Zfp746
1435244_at	1.01343	Vav2 oncogene	Vav2
1435245_at	0.86740	"glutaminase 2 (liver, mitochondrial)"	Gls2
1435246_at	0.89044	progesterin and adipoQ receptor family member VIII	Paqr8
1435247_at	0.84566	ubiquitin-activating enzyme E1-domain containing 1	Ube1dc1
1435248_a_at	1.07810	"BTAF1 RNA polymerase II, B-TFIID transcription factor-associated, (Mot1 homolog, S. cerevisiae)"	Btaf1
1435249_at	1.03506	"BTAF1 RNA polymerase II, B-TFIID transcription factor-associated, (Mot1 homolog, S. cerevisiae)"	Btaf1
1435250_at	1.01771	integrator complex subunit 8	Ints8
1435251_at	0.93586	sorting nexin 13	Snx13
1435252_at	1.04406	"UDP-Gal:betaGal beta 1,3-galactosyltransferase, polypeptide 6"	B3galt6
1435253_at	1.18227	"RAB11B, member RAS oncogene family"	Rab11b
1435254_at	0.81371	plexin B1	Plxnb1
1435255_at	0.91124	plexin B1	Plxnb1
1435256_at	0.96981	RIKEN cDNA 1500005P14 gene	1500005P14Rik
1435257_at	0.78136	kelch-like 17 (Drosophila)	Klh17
1435258_at	0.96777	transmembrane protein 141	Tmem141
1435259_s_at	1.04082	transmembrane protein 141	Tmem141
1435260_at	0.91133	---	---
1435261_at	1.08407	RIKEN cDNA 4732416N19 gene	4732416N19Rik
1435262_at	0.93491	Transcribed locus	---
1435263_at	0.96765	transmembrane protein 106A	Tmem106a
1435264_at	1.04530	elastin microfibril interfacier 2	Emilin2
1435265_at	0.80264	"Adult male cecum cDNA, RIKEN full-length enriched library, clone:9130204C03 product:unclassifiable, full insert sequence"	---
1435266_at	0.84512	RIKEN cDNA A930041I02 gene	A930041I02Rik
1435267_at	1.96562	RIKEN cDNA A430108E01 gene	A430108E01Rik
1435268_at	0.99886	"guanine nucleotide binding protein, alpha z subunit"	Gnaz
1435269_at	1.18767	RIKEN cDNA 2510005D08 gene	2510005D08Rik
1435270_x_at	1.03686	RIKEN cDNA 2510005D08 gene	2510005D08Rik
1435271_at	0.99235	interferon regulatory factor 3	Irf3
1435272_at	0.84036	"inositol 1,4,5-trisphosphate 3-kinase B"	Itpkb

1435273_at 0.96973 tryptophanyl tRNA synthetase 2 (mitochondrial) Wars2
1435274_at 0.97610 RIKEN cDNA 4921517N04 gene 4921517N04Rik
1435275_at 1.05239 cytochrome c oxidase subunit VIb polypeptide 2 Cox6b2
1435276_a_at 0.86665 DiGeorge syndrome critical region gene 2 Dgcr2
1435277_x_at 1.06626 "expressed in non-metastatic cells 1, protein" Nme1
1435278_at 0.75480 SFT2 domain containing 3 Sft2d3
1435279_at 0.89819 cDNA sequence BC059842 BC059842
1435280_at 0.83035 expressed sequence AI452195 AI452195
1435281_at 0.92561 carnitine palmitoyltransferase 1c Cpt1c
1435282_at 0.79631 "gene model 967, (NCBI)" Gm967
1435283_s_at 0.71452 "gene model 967, (NCBI)" Gm967
1435284_at 2.05182 reticulon 4 Rtn4
1435285_at 0.84331 metallophosphoesterase domain containing 2 Mpped2
1435286_at 1.13525 expressed sequence AW125296 AW125296
1435287_at 1.13983 adducin 2 (beta) Add2
1435288_at 1.02625 "coronin, actin binding protein 1A" Coro1a
1435289_at 0.88600 RIKEN cDNA D230014K01 gene D230014K01Rik
1435290_x_at 0.82325 "histocompatibility 2, class II antigen A, alpha" H2-Aa
1435291_at 0.92119 LEM domain containing 3 Lemd3
1435292_at 0.78022 "TBC1 domain family, member 4" Tbc1d4
1435293_at 1.11174 RIKEN cDNA 2900022I03 gene 2900022I03Rik
1435294_at 0.95164 myotubularin related protein 11 Mtmr11
1435295_at 0.88251 dopey family member 1 Dopey1
1435296_at 1.02360 "adrenergic receptor, alpha 2c" Adra2c
1435297_at 0.81722 gap junction membrane channel protein alpha 9 Gja9
1435298_at 1.06742 component of oligomeric golgi complex 5 Cog5
1435299_at 0.86363 hypothetical protein C230069N13 C230069N13
1435300_at 0.94575 --- ---
1435301_at 0.94476 RIKEN cDNA 1110004E09 gene 1110004E09Rik
1435302_at 1.11507 "TAF4B RNA polymerase II, TATA box binding protein (TBP)-associated factor" Taf4b
1435303_at 1.25150 "TAF4B RNA polymerase II, TATA box binding protein (TBP)-associated factor" Taf4b
1435304_at 0.91703 "superoxide dismutase 1, soluble" Sod1
1435305_at 0.92824 "neurotrophic tyrosine kinase, receptor, type 2" Ntrk2
1435306_a_at 1.02339 kinesin family member 11 Kif11
1435307_at 0.98028 RIKEN cDNA 6430502M16 gene 6430502M16Rik
1435308_at 1.01398 fucosyltransferase 9 Fut9
1435309_at 0.99542 cDNA sequence BC019943 BC019943
1435310_at 1.09986 F-box only protein 7 Fbxo7
1435311_s_at 0.91370 F-box only protein 7 Fbxo7
1435312_at 0.87830 Progesterin and adipoQ receptor family member VII Paqr7
1435313_at 1.13409 Cd200 receptor 4 Cd200r4

1435314_at 0.96886 tryptophan hydroxylase 2 Tph2
1435315_s_at 0.94750 RIKEN cDNA 2900034E22 gene 2900034E22Rik
1435316_at 1.08374 "Proteasome (prosome, macropain) subunit, alpha type 6" Psma6
1435317_x_at 1.11777 "proteasome (prosome, macropain) subunit, alpha type 6" Psma6
1435318_at 0.96556 RIKEN cDNA E130218I03 gene E130218I03Rik
1435319_at 0.83869 inositol hexaphosphate kinase 2 Ihpk2
1435320_at 0.99386 "CTD (carboxy-terminal domain, RNA polymerase II, polypeptide A) small phosphatase like 2" Ctdspl2
1435321_at 0.82616 RIKEN cDNA 3732412D22 gene 3732412D22Rik
1435322_at 0.89824 RIKEN cDNA 1110011C06 gene 1110011C06Rik
1435323_a_at 1.03047 membrane bound O-acyltransferase domain containing 1 Mboat1
1435324_x_at 0.98510 high mobility group box 1 Hmgb1
1435325_at 0.80544 ubiquitin specific peptidase 46 Usp46
1435326_at 0.81865 expressed sequence AW112037 AW112037
1435327_at 0.85200 expressed sequence AW112037 AW112037
1435328_at 0.92115 cysteine and histidine rich 1 Cyhr1
1435329_at 1.02219 F-box and leucine-rich repeat protein 11 Fbx11
1435330_at 0.88482 --- ---
1435331_at 0.92551 expressed sequence AI447904 AI447904
1435332_at 0.91094 5-hydroxytryptamine (serotonin) receptor 7 Htr7
1435333_at 0.94961 RIKEN cDNA 1110007M04 gene 1110007M04Rik
1435334_at 0.95967 tetratricopeptide repeat domain 7 Ttc7
1435335_a_at 0.98931 "N-acetylglucosamine-1-phosphate transferase, alpha and beta subunits" Gnptab
1435336_at 0.69867 cadherin EGF LAG seven-pass G-type receptor 2 Celsr2
1435337_at 1.04035 teashirt zinc finger family member 3 Tshz3
1435338_at 1.08713 cyclin-dependent kinase 6 Cdk6
1435339_at 0.88105 potassium channel tetramerisation domain containing 15 Kctd15
1435340_at 0.98769 jumonji domain containing 2A Jmjd2a
1435341_at 1.00586 peptidyl-prolyl isomerase G (cyclophilin G) Ppig
1435342_at 1.00474 "potassium inwardly-rectifying channel, subfamily K, member 6" Kcnk6
1435343_at 1.00868 dedicator of cytokinesis 10 /// similar to Dedicator of cytokinesis protein 10 (Protein zizimin 3) Dock10 /// LOC630691
1435344_at 0.90014 transcription factor Dp 2 Tfdp2
1435345_at 0.78392 cerebral endothelial cell adhesion molecule 1 Ceecam1
1435346_at 1.25776 coiled-coil domain containing 82 Ccdc82
1435347_at 1.04883 staufen (RNA binding protein) homolog 1 (Drosophila) Stau1
1435348_at 0.83449 RIKEN cDNA D930009K15 gene D930009K15Rik
1435349_at 1.54407 neuropilin 2 Nrp2
1435350_at 0.93218 Tnf receptor-associated factor 6 Traf6
1435351_at 1.02589 RIKEN cDNA 2310026E23 gene 2310026E23Rik
1435352_at 0.97796 "phosphatidylinositol glycan anchor biosynthesis, class K" Pigk
1435353_a_at 0.82102 RIKEN cDNA 4933439C20 gene 4933439C20Rik

1435354_at 0.94711 "potassium inwardly-rectifying channel, subfamily J, member 15" Kcnj15
1435355_at 1.03737 nebulin Neb
1435356_at 1.02801 F-box protein 42 Fbxo42
1435357_at 0.86068 --- ---
1435358_at 1.00943 CUE domain containing 1 Cuedc1
1435359_at 1.06283 cDNA sequence BC060632 BC060632
1435360_at 0.99905 zinc finger protein 651 Zfp651
1435361_at 1.00712 --- ---
1435362_at 1.12851 forkhead box J3 Foxj3
1435363_at 0.75311 "pleckstrin homology domain containing, family G (with RhoGef domain) member 1" Plekhg1
1435364_at 1.03325 cysteine and histidine rich 1 Cyhr1
1435365_at 0.97794 RIKEN cDNA 4732415M23 gene 4732415M23Rik
1435366_at 0.92018 RIKEN cDNA D430042O09 gene D430042O09Rik
1435367_at 0.90655 mitogen-activated protein kinase 4 Mapk4
1435368_a_at 0.93443 "poly (ADP-ribose) polymerase family, member 1" Parp1
1435369_at 1.00480 FAST kinase domains 5 Fastkd5
1435370_a_at 0.92760 carboxylesterase 3 Ces3
1435371_x_at 1.01614 carboxylesterase 3 Ces3
1435372_a_at 1.15081 proliferation-associated 2G4 Pa2g4
1435373_at 0.87812 "casein kinase 1, epsilon" Csnk1e
1435374_at 0.89567 Transcribed locus ---
1435375_at 0.85197 cDNA sequence BC052328 BC052328
1435376_at 1.09631 DDHD domain containing 2 Ddhd2
1435377_at 1.01452 --- ---
1435378_at 1.06142 RIKEN cDNA 2210020M01 gene 2210020M01Rik
1435379_at 1.07369 cDNA sequence AK122209 AK122209
1435380_at 1.01428 "COX10 homolog, cytochrome c oxidase assembly protein, heme A: farnesyltransferase (yeast)" Cox10
1435381_at 1.29196 RIKEN cDNA 2610110G12 gene 2610110G12Rik
1435382_at 1.00153 necdin Ndn
1435383_x_at 0.90631 necdin Ndn
1435384_at 0.78761 ubiquitin-conjugating enzyme E2N Ube2n
1435385_at 0.90852 teashirt zinc finger family member 2 Tshz2
1435386_at 0.99318 Von Willebrand factor homolog Vwf
1435387_at 0.90525 "solute carrier family 2 (facilitated glucose transporter), member 13" Slc2a13
1435388_at 1.09829 NADPH dependent diflavin oxidoreductase 1 Ndor1
1435389_at 1.01762 RALBP1 associated Eps domain containing protein 2 Reps2
1435390_at 1.04716 exonuclease domain containing 1 Exod1
1435391_at 1.16583 RIKEN cDNA 3200001K10 gene 3200001K10Rik
1435392_at 1.04925 WD repeat domain 17 Wdr17
1435393_at 0.98500 "CDNA, clone:Y0G0149N04, strand:unspecified" ---
1435394_s_at 0.94674 "ras homolog gene family, member C" Rhoc

1435395_s_at 0.95113 "ATP synthase, H⁺ transporting, mitochondrial F0 complex, subunit f, isoform 2" Atp5j2
 1435396_at 0.90010 syntaxin binding protein 6 (amisyn) Stxbp6
 1435397_at 0.76332 cDNA sequence BC038156 BC038156
 1435398_at 1.03658 syntaxin binding protein 5 (tomosyn) Stxbp5
 1435399_at 1.01768 RIKEN cDNA 2310068J10 gene 2310068J10Rik
 1435400_at 0.90669 RIKEN cDNA 2900075B16 gene 2900075B16Rik
 1435401_at 0.89578 tetratricopeptide repeat domain 26 Ttc26
 1435402_at 0.90824 GRAM domain containing 1B Gramd1b
 1435403_at 1.05532 RIKEN cDNA 1700007L15 gene 1700007L15Rik
 1435404_at 1.06268 dispatched homolog 2 (Drosophila) Disp2
 1435405_at 0.98762 SET domain containing 4 Setd4
 1435406_at 0.92135 coiled-coil alpha-helical rod protein 1 Cchcr1
 1435407_at 0.90400 Zinc finger protein 483 Zfp483
 1435408_at 1.19153 RIKEN cDNA 2700092H06 gene 2700092H06Rik
 1435409_at 0.93946 Transcribed locus ---
 1435410_at 1.00050 testicular cell adhesion molecule 1 Tcam1
 1435411_at 1.01649 Transcribed locus ---
 1435412_at 0.96068 RIKEN cDNA 1700007E06 gene 1700007E06Rik
 1435413_x_at 1.04034 RIKEN cDNA 2700060E02 gene 2700060E02Rik
 1435414_s_at 0.90078 dynactin 1 Dctn1
 1435415_x_at 0.87779 MARCKS-like 1 Marcksl1
 1435416_x_at 0.94064 "phosphatidylinositol glycan anchor biosynthesis, class Q" Pigq
 1435417_at 0.87449 expressed sequence AI464131 AI464131
 1435418_at 0.86217 "solute carrier family 22 (organic anion transporter), member 8" Slc22a8
 1435419_at 0.91115 retrotransposon gag domain containing 4 Rgag4
 1435420_at 0.98805 RIKEN cDNA C330016K18 gene C330016K18Rik
 1435421_at 0.97843 fibronectin type 3 and SPRY domain-containing protein Fsd1
 1435422_at 0.99581 RIKEN cDNA 4933433P14 gene 4933433P14Rik
 1435423_x_at 1.00430 RIKEN cDNA 4933433P14 gene 4933433P14Rik
 1435424_x_at 0.99567 Transcribed locus ---
 1435425_at 0.93735 RIKEN cDNA 4933439C20 gene /// phosphatidylserine decarboxylase 4933439C20Rik /// Pisd
 1435426_s_at 0.75017 RIKEN cDNA 4933439C20 gene /// phosphatidylserine decarboxylase 4933439C20Rik /// Pisd
 1435427_x_at 0.93201 cDNA sequence BC037112 BC037112
 1435428_at 0.96055 cDNA sequence BC037112 BC037112
 1435429_x_at 1.04004 ribosomal protein S27-like Rps27l
 1435430_at 1.00639 transmembrane protein 1 Tmem1
 1435431_at 1.07934 RIKEN cDNA 2310047M15 gene 2310047M15Rik
 1435432_at 1.04654 "centaurin, gamma 2" Centg2
 1435433_at 1.02872 "centaurin, gamma 2" Centg2
 1435434_at 0.99448 Braf transforming gene Braf

1435435_at 1.00195 cortactin binding protein 2 Ctnbp2
1435436_at 0.92856 Endothelial PAS domain protein 1 Epas1
1435437_at 0.99829 SET domain containing (lysine methyltransferase) 7 Setd7
1435438_at 1.10050 SRY-box containing gene 8 Sox8
1435439_at 0.96643 DiGeorge syndrome critical region gene 8 Dgcr8
1435440_at 0.96145 PDZ domain containing 8 Pdzd8
1435441_at 0.96751 actin-binding LIM protein 2 Ablim2
1435442_at 1.10038 WD repeats and SOF domain containing 1 Wdsof1
1435443_at 1.00410 eyes absent 3 homolog (Drosophila) Eya3
1435444_at 1.03020 activating transcription factor 6 Atf6
1435445_at 0.94085 cyclin T2 Ccnt2
1435446_a_at 0.90934 choline phosphotransferase 1 Chpt1
1435447_at 1.12769 major intrinsic protein of eye lens fiber Mip
1435448_at 0.80807 BCL2-like 11 (apoptosis facilitator) Bcl2l11
1435449_at 0.78903 BCL2-like 11 (apoptosis facilitator) Bcl2l11
1435450_at 1.01388 copine III Cpne3
1435451_at 1.08002 "helicase, mus308-like (Drosophila)" Hel308
1435452_at 1.34291 transmembrane protein 20 Tmem20
1435453_at 0.95074 RIKEN cDNA A930011O12 gene A930011O12Rik
1435454_a_at 1.06474 cDNA sequence BC006779 BC006779
1435455_at 1.13257 expressed sequence C79267 C79267
1435456_at 0.80521 tetratricopeptide repeat domain 28 Ttc28
1435457_at 0.99413 PDZ domain containing 4 Pdzd4
1435458_at 1.14118 proviral integration site 1 Pim1
1435459_at 0.90720 flavin containing monooxygenase 2 Fmo2
1435460_at 1.84324 "protein kinase, cGMP-dependent, type II" Prkg2
1435461_at 0.86729 "membrane associated guanylate kinase, WW and PDZ domain containing 3"
Magi3
1435462_at 1.07261 hypothetical LOC433022 LOC433022
1435463_s_at 1.02715 myosin ID Myo1d
1435464_at 0.95891 RIKEN cDNA 1110003E01 gene 1110003E01Rik
1435465_at 0.93193 kelch repeat and BTB (POZ) domain containing 11 /// similar to Protein
KIAA0711 Kbtbd11 /// LOC632344
1435466_at 0.89609 RIKEN cDNA A830091I15 gene A830091I15Rik
1435467_at 1.11608 "FYVE, RhoGEF and PH domain containing 6" Fgd6
1435468_at 1.00822 expressed sequence C76566 C76566
1435469_at 1.05669 quiescin Q6-like 1 Qscn6l1
1435470_at 0.93016 similar to p47 protein isoform a LOC547150
1435471_at 0.88204 zinc finger protein 708 Zfp708
1435472_at 1.03521 kringle containing transmembrane protein 1 Kremen1
1435473_at 0.83136 "gene model 347, (NCBI)" Gm347
1435474_at 0.94431 "TAF5 RNA polymerase II, TATA box binding protein (TBP)-associated
factor" Taf5

1435475_at	0.87439	"lectin, mannose-binding 2-like"	Lman2l
1435476_a_at	0.93750	"Fc receptor, IgG, low affinity IIb"	Fcgr2b
1435477_s_at	0.88574	"Fc receptor, IgG, low affinity IIb"	Fcgr2b
1435478_at	0.95358	WD repeat domain 26	Wdr26
1435479_at	0.92787	---	---
1435480_at	0.98231	Braf transforming gene	Braf
1435481_at	0.98208	zinc finger protein 653	Zfp653
1435482_at	1.10654	fibrinogen C domain containing 1	Fibcd1
1435483_x_at	0.84078	"solute carrier family 25, member 32"	Slc25a32
1435484_at	1.11534	expressed sequence BF642829	BF642829
1435485_at	0.96020	RIKEN cDNA C230096C10 gene	C230096C10Rik
1435486_at	1.18755	---	---
1435487_at	1.01808	"glutamate receptor, ionotropic, delta 2"	Grid2
1435488_at	0.99374	RIKEN cDNA C630010D07 gene /// similar to methylenetetrahydrofolate dehydrogenase (NADP+ dependent) 2-like /// region containing RIKEN cDNA C630010D07 gene; RIKEN cDNA 1110019K23 gene	C630010D07Rik /// LOC665563 /// LOC674367
1435489_at	0.75452	hypothetical LOC552912	LOC552912
1435490_at	0.99894	hexokinase 3	Hk3
1435491_at	0.88700	RIKEN cDNA 9830167H18 gene	9830167H18Rik
1435492_at	1.05132	suppressor of cytokine signaling 6	Socs6
1435493_at	0.91975	desmoplakin	Dsp
1435494_s_at	0.86367	desmoplakin	Dsp
1435495_at	0.85434	adenosine A1 receptor	Adora1
1435496_at	1.09177	RIKEN cDNA 5730469M10 gene	5730469M10Rik
1435497_at	1.06344	RIKEN cDNA 5730590G19 gene	5730590G19Rik
1435498_at	0.79980	ADP-ribosylation factor interacting protein 2	Arfip2
1435499_at	0.91445	RIKEN cDNA D030041N04 gene	D030041N04Rik
1435500_at	0.88647	"RAB26, member RAS oncogene family"	Rab26
1435501_at	0.92674	RIKEN cDNA C030004M05 gene	C030004M05Rik
1435502_a_at	1.22449	"bromodomain, testis-specific"	Brdt
1435503_at	1.06700	KIAA0415 protein	KIAA0415
1435504_at	1.05757	Transcribed locus	---
1435505_at	0.93304	dystrophia myotonica-containing WD repeat motif	Dmwd
1435506_at	0.83905	---	---
1435507_x_at	1.03659	"protease, serine, 2"	Prss2
1435508_x_at	1.04518	RIKEN cDNA 0610009D07 gene	0610009D07Rik
1435509_x_at	0.91770	CDK2 (cyclin-dependent kinase 2)-associated protein 1	Cdk2ap1
1435510_at	1.28007	protein phosphatase 1H (PP2C domain containing)	Ppm1h
1435511_at	1.03921	synapsin II	Syn2
1435512_at	0.95306	expressed sequence AI836003	AI836003
1435513_at	0.91819	5-hydroxytryptamine (serotonin) receptor 2C	Htr2c
1435514_at	0.94278	leucine zipper transcription factor-like 1	Lztfl1
1435515_at	0.90623	Transcribed locus	---

1435516_x_at 1.00172 regulator of telomere elongation helicase 1 Rtel1
1435517_x_at 0.87737 v-ral simian leukemia viral oncogene homolog B (ras related) Ralb
1435518_at 0.96662 RAS related protein 1b Rap1b
1435519_at 0.96850 RAS related protein 1b Rap1b
1435520_at 0.91840 Musashi homolog 2 (Drosophila) Msi2
1435521_at 0.82090 Musashi homolog 2 (Drosophila) Msi2
1435522_a_at 1.01642 RIKEN cDNA 2310016E02 gene 2310016E02Rik
1435523_s_at 0.92547 RIKEN cDNA 2700089E24 gene 2700089E24Rik
1435524_at 0.99649 RIKEN cDNA 2010109N14 gene 2010109N14Rik
1435525_at 0.80846 potassium channel tetramerisation domain containing 17 Kctd17
1435526_at 1.00998 torsin A interacting protein 2 Tor1aip2
1435527_at 0.95353 nuclear factor I/C Nfic
1435528_at 0.99769 "DNA segment, Chr 2, Brigham & Women's Genetics 1335 expressed"
D2Bwg1335e
1435529_at 0.90084 similar to interferon-induced protein with tetratricopeptide repeats 1 /// similar
to interferon-induced protein with tetratricopeptide repeats 1 LOC667373 /// LOC669129
1435530_at 1.01398 calmodulin regulated spectrin-associated protein 1 Camsap1
1435531_at 0.86277 ubiquitin specific peptidase 3 Usp3
1435532_at 0.99087 RIKEN cDNA 4933426K21 gene 4933426K21Rik
1435533_s_at 0.99271 RIKEN cDNA 4933426K21 gene 4933426K21Rik
1435534_a_at 1.09853 translocase of outer mitochondrial membrane 20 homolog (yeast) Tomm20
1435535_at 1.00335 DEP domain containing 5 Depdc5
1435536_at 0.81156 "regulatory solute carrier protein, family 1, member 1" Rsc1a1
1435537_at 0.99505 "protein tyrosine phosphatase, receptor type, D" Ptpnd
1435538_at 1.10526 RIKEN cDNA 0610030E20 gene 0610030E20Rik
1435539_at 0.89660 Huntington disease gene homolog Hdh
1435540_at 0.89592 "immunity-related GTPase family, Q" Irgq
1435541_at 1.01116 "betacellulin, epidermal growth factor family member" Btc
1435542_s_at 0.92298 CTTNBP2 N-terminal like Ctnnbp2nl
1435543_at 0.96875 adenomatosis polyposis coli Apc
1435544_at 1.11023 exosome component 6 Exosc6
1435545_at 0.80496 cDNA sequence BC032203 BC032203
1435546_a_at 1.10663 RIKEN cDNA 1810013L24 gene 1810013L24Rik
1435547_at 0.89971 "17 days embryo heart cDNA, RIKEN full-length enriched library, clone:
I920018H16 product:hypothetical protein, full insert sequence" ---
1435548_at 1.11007 "MRS2-like, magnesium homeostasis factor (S. cerevisiae)" Mrs2l
1435549_at 0.82668 "transient receptor potential cation channel, subfamily M, member 4" Trpm4
1435550_at 0.91941 RIKEN cDNA C430014K11 gene C430014K11Rik
1435551_at 0.86748 formin homology 2 domain containing 3 Fhod3
1435552_at 1.00174 OTU domain containing 7B Otud7b
1435553_at 1.07779 PDZ domain containing 2 Pdzd2
1435554_at 0.71713 transmembrane and coiled coil domains 3 Tmcc3
1435555_at 1.11436 --- ---

1435556_at 0.89128 zinc finger protein 597 Zfp597
1435557_at 1.24799 formin homology 2 domain containing 1 Fhod1
1435558_at 0.65503 brain-specific angiogenesis inhibitor 2 Bai2
1435559_at 0.75728 myosin VI Myo6
1435560_at 0.88634 integrin alpha L /// similar to Integrin alpha-L precursor (Leukocyte adhesion glycoprotein LFA-1 alpha chain) (LFA-1A) (Leukocyte function-associated molecule 1 alpha chain) (Lymphocyte antigen Ly-15) (CD11a antigen) Itgal /// LOC671163
1435561_at 1.01964 Ets2 repressor factor Erf
1435562_at 0.85340 PDZ domain containing 8 Pdzd8
1435563_at 0.82394 mitochondrial ribosomal protein S5 Mrps5
1435564_at 0.88865 RIKEN cDNA C230078M08 gene C230078M08Rik
1435565_at 1.06991 RIKEN cDNA 1500011H22 gene 1500011H22Rik
1435566_s_at 0.95453 v-raf murine sarcoma 3611 viral oncogene homolog Araf
1435567_at 1.20333 phosphorylase kinase alpha 1 Phka1
1435568_at 0.94975 cDNA sequence AK129128 AK129128
1435569_at 1.00740 RIKEN cDNA D630029K05 gene D630029K05Rik
1435570_s_at 1.09040 RIKEN cDNA D630029K05 gene D630029K05Rik
1435571_at 0.92674 RIKEN cDNA A530065I17 gene A530065I17Rik
1435572_at 0.93421 RIKEN cDNA 2310014L17 gene 2310014L17Rik
1435573_at 0.91323 OTU domain containing 5 Otud5
1435574_at 1.09481 "Cysteine and histidine-rich domain (CHORD)-containing, zinc-binding protein 1" Chordc1
1435575_at 0.88050 kinetochore associated 1 Kntc1
1435576_at 1.02615 expressed sequence AW413774 AW413774
1435577_at 1.05448 disabled homolog 1 (Drosophila) Dab1
1435578_s_at 1.01745 disabled homolog 1 (Drosophila) Dab1
1435579_at 1.11191 "Opioid receptor, sigma 1" Oprs1
1435580_at 0.92306 RIKEN cDNA C230081A13 gene C230081A13Rik
1435581_at 0.84762 beta-site APP-cleaving enzyme 2 Bace2
1435582_at 0.95762 WD repeat and FYVE domain containing 4 Wdfy4
1435583_at 1.00922 RIKEN cDNA 9330182L06 gene 9330182L06Rik
1435584_at 1.02156 expressed sequence AI662791 AI662791
1435585_at 1.00527 transcription elongation factor A (SII)-like 7 Tceal7
1435586_at 1.03837 PCI domain containing 2 Pcid2
1435587_at 1.12643 PCI domain containing 2 Pcid2
1435588_at 0.96347 WD repeat and FYVE domain containing 1 Wdfy1
1435589_at 1.12647 coiled-coil domain containing 85B Ccdc85b
1435590_at 0.97105 RIKEN cDNA D430047L21 gene D430047L21Rik
1435591_at 0.77909 expressed sequence AI426330 AI426330
1435592_at 1.03060 eukaryotic translation initiation factor 5B Eif5b
1435593_x_at 1.03605 ribosomal protein S7 Rps7
1435594_at 0.76171 ADP-ribosylation factor-like 6 interacting protein 2 Arl6ip2
1435595_at 1.27511 RIKEN cDNA 1810011O10 gene 1810011O10Rik

1435596_at 0.94428 RIKEN cDNA A530088I07 gene A530088I07Rik
1435597_at 1.10669 RIKEN cDNA C130052G03 gene C130052G03Rik
1435598_at 1.01281 Transcribed locus ---
1435599_at 0.92140 cDNA sequence BC020535 BC020535
1435600_s_at 0.94513 cDNA sequence BC020535 BC020535
1435601_at 1.24483 expressed sequence AI481772 AI481772
1435602_at 0.93115 Transcribed locus ---
1435603_at 0.89470 "sushi, nidogen and EGF-like domains 1" Sned1
1435604_at 0.95144 tripartite motif protein 37 Trim37
1435605_at 0.86675 ARP3 actin-related protein 3 homolog B (yeast) Actr3b
1435606_at 1.05956 galactose-3-O-sulfotransferase 4 Gal3st4
1435607_at 1.03263 "G protein-coupled receptor, family C, group 1, member B" Grm2
1435608_at 0.93107 zinc and ring finger 3 /// similar to Goliath homolog precursor (Ring finger protein 130) (R-goliath) Znrf3 /// LOC631806
1435609_at 0.96211 --- ---
1435610_at 0.90334 RIKEN cDNA C130034I18 gene C130034I18Rik
1435611_x_at 1.07473 "elastase 3, pancreatic" Ela3
1435612_at 1.05291 opioid binding protein/cell adhesion molecule-like Opcml
1435613_x_at 0.94586 "cytochrome c oxidase, subunit Vb" Cox5b
1435614_s_at 0.89164 RAS protein-specific guanine nucleotide-releasing factor 1 Rasgrf1
1435615_at 1.01131 zinc finger protein 365 Zfp365
1435616_at 0.95616 "cytochrome P450, family 20, subfamily A, polypeptide 1" Cyp20a1
1435617_at 0.88704 expressed sequence AI480743 AI480743
1435618_at 0.90497 paraneoplastic antigen MA2 Pnma2
1435619_at 0.77428 Transcribed locus ---
1435620_at 1.07302 RIKEN cDNA 1190002J23 gene 1190002J23Rik
1435621_at 0.93319 male sterility domain containing 1 Mlstd1
1435622_at 1.02632 heparan sulfate (glucosamine) 3-O-sulfotransferase 3A1 Hs3st3a1
1435623_at 0.97532 transmembrane protein 118 Tmem118
1435624_at 0.97991 RIKEN cDNA 9030409G11 gene 9030409G11Rik
1435625_at 1.01266 ectonucleoside triphosphate diphosphohydrolase 7 Entpd7
1435626_a_at 0.92214 "homocysteine-inducible, endoplasmic reticulum stress-inducible, ubiquitin-like domain member 1" Herpud1
1435627_x_at 0.79848 MARCKS-like 1 /// similar to MARCKS-related protein (MARCKS-like protein 1) (Macrophage myristoylated alanine-rich C kinase substrate) (Mac-MARCKS) (MacMARCKS) (Brain protein F52) Marcks11 /// LOC673071
1435628_x_at 1.25493 RIKEN cDNA 2610305J24 gene /// cDNA sequence BC005512 /// hypothetical protein LOC629242 /// similar to melanoma antigen /// similar to melanoma antigen /// similar to melanoma antigen /// hypothetical ENV polyprotein /// similar to melanoma antigen /// similar to melanoma antigen /// hypothetical protein LOC670270 /// similar to melanoma antigen /// similar to melanoma antigen /// hypothetical protein LOC672969 /// similar to melanoma antigen /// similar to melanoma antigen /// similar to melanoma antigen /// similar to melanoma antigen /// similar to melanoma antigen /// similar to melanoma antigen

2610305J24Rik /// BC005512 /// LOC629242 /// LOC630525 /// LOC631386 /// LOC637482 ///
 LOC641366 /// LOC668904 /// LOC669885 /// LOC670270 /// LOC672061 /// LOC672939 ///
 LOC672969 /// LOC673876 /// LOC674927 /// LOC675168 /// LOC675706 /// LOC676247 ///
 LOC676718 /// LOC677506
 1435629_at 0.92135 Transcribed locus ---
 1435630_s_at 1.49157 acetyl-Coenzyme A acetyltransferase 2 Acat2
 1435631_x_at 1.06412 exocyst complex component 6 Exoc6
 1435632_at 1.08754 nuclear fragile X mental retardation protein interacting protein 2 Nufip2
 1435633_at 1.08702 RIKEN cDNA 4631424J17 gene 4631424J17Rik
 1435634_at 0.91746 protein-L-isoaspartate (D-aspartate) O-methyltransferase domain containing 1
 Pcmt1
 1435635_at 0.82016 protein-L-isoaspartate (D-aspartate) O-methyltransferase domain containing 1
 Pcmt1
 1435636_at 1.03071 RIKEN cDNA 2310051F07 gene 2310051F07Rik
 1435637_at 0.85970 integrin alpha FG-GAP repeat containing 1 Itfg1
 1435638_at 0.87909 glycogen synthase kinase 3 alpha Gsk3a
 1435639_at 1.15972 RIKEN cDNA 2610528A11 gene 2610528A11Rik
 1435640_x_at 1.36543 RIKEN cDNA A130040M12 gene A130040M12Rik
 1435641_at 0.96714 RIKEN cDNA 9530018I07 gene 9530018I07Rik
 1435642_at 0.84763 RIKEN cDNA 9630019K15 gene 9630019K15Rik
 1435643_x_at 0.97458 "ubiquitin B /// gene model 1821, (NCBI)" Ubb /// Gm1821
 1435644_at 0.90127 SH3 and PX domains 2B Sh3pxd2b
 1435645_at 0.89116 monocyte to macrophage differentiation-associated /// similar to monocyte to
 macrophage differentiation-associated Mmd /// LOC676546
 1435646_at 0.90898 inhibitor of kappaB kinase gamma Ikbkg
 1435647_at 0.88390 inhibitor of kappaB kinase gamma Ikbkg
 1435648_at 1.02534 RIKEN cDNA B430119L13 gene B430119L13Rik
 1435649_at 0.58376 nexilin Nexn
 1435650_at 0.87184 hyaluronan and proteoglycan link protein 4 Hapln4
 1435651_a_at 0.95207 "SWI/SNF-related, matrix-associated actin-dependent regulator of chromatin,
 subfamily a, containing DEAD/H box 1`" Smarcd1
 1435652_a_at 0.91379 "guanine nucleotide binding protein, alpha inhibiting 2" Gnai2
 1435653_at 0.95341 "Adult male liver tumor cDNA, RIKEN full-length enriched library, clone:
 C730024P13 product:unclassifiable, full insert sequence" ---
 1435654_at 0.94567 RIKEN cDNA G630039H03 gene G630039H03Rik
 1435655_at 1.34479 "small nucleolar RNA, H/ACA box 65" Snora65
 1435656_at 0.88486 guanine monphosphate synthetase Gmps
 1435657_at 0.96434 expressed sequence AI425999 AI425999
 1435658_at 0.91989 "solute carrier family 27 (fatty acid transporter), member 1" Slc27a1
 1435659_a_at 1.17264 triosephosphate isomerase 1 Tpi1
 1435660_at 1.11110 RIKEN cDNA C130026I21 gene /// RIKEN cDNA A530032D15Rik gene ///
 similar to SP140 nuclear body protein (predicted) C130026I21Rik /// A530032D15Rik ///
 LOC620078

1435661_at 0.87109 "amyotrophic lateral sclerosis 2 (juvenile) chromosome region, candidate 4"
Als2cr4

1435662_at 0.93465 RIKEN cDNA 2610020O08 gene 2610020O08Rik

1435663_at 0.96531 estrogen receptor 1 (alpha) Esr1

1435664_at 1.10400 zinc finger protein 397 Zfp397

1435665_at 0.98105 expressed sequence AI451617 AI451617

1435666_at 1.00389 microtubule associated serine/threonine kinase 3 Mast3

1435667_at 1.08058 regulating synaptic membrane exocytosis 1 Rims1

1435668_at 0.96486 RIKEN cDNA 4932442K08 gene 4932442K08Rik

1435669_at 0.86789 zinc finger protein 532 Zfp532

1435670_at 0.80952 transcription factor AP-2 beta Tcfap2b

1435671_at 0.96129 mirror-image polydactyly gene 1 homolog (human) Mipol1

1435672_at 0.85731 hypothetical protein 3830612M24 3830612M24

1435673_at 1.01565 expressed sequence AI448005 AI448005

1435674_at 0.98454 Rho-related BTB domain containing 2 Rhobtb2

1435675_at 0.96269 "TBC1D12: TBC1 domain family, member 12" Tbc1d12

1435676_at 0.97109 centrosomal protein 164 Cep164

1435677_at 0.97329 expressed sequence C85492 C85492

1435678_at 0.96114 RIKEN cDNA 2610017I09 gene 2610017I09Rik

1435679_at 1.02300 optineurin Optn

1435680_a_at 1.03884 dipeptidylpeptidase 7 Dpp7

1435681_s_at 0.88187 DEAD (Asp-Glu-Ala-Asp) box polypeptide 49 Ddx49

1435682_at 0.89933 "leucyl-tRNA synthetase, mitochondrial" Lars2

1435683_a_at 0.96961 "ATP-binding cassette, sub-family C (CFTR/MRP), member 5" Abcc5

1435684_at 0.89615 "ATP-binding cassette, sub-family C (CFTR/MRP), member 5" Abcc5

1435685_x_at 1.05402 "ATP-binding cassette, sub-family C (CFTR/MRP), member 5" Abcc5

1435686_at 0.81985 expressed sequence AA536717 AA536717

1435687_at 0.98289 synaptic vesicle glycoprotein 2 b Sv2b

1435688_at 0.75152 Chromodomain helicase DNA binding protein 9 Chd9

1435689_at 1.01707 RIKEN cDNA 9030025P20 gene 9030025P20Rik

1435690_at 1.15758 RIKEN cDNA 2310008H09 gene 2310008H09Rik

1435691_at 0.92796 RIKEN cDNA C630028N24 gene C630028N24Rik

1435692_at 1.06326 "similar to retinoic acid, EGF, and NGF upregulated" LOC622320

1435693_at 0.87041 "mal, T-cell differentiation protein-like" Mall

1435694_at 1.01799 Rho GTPase activating protein 26 Arhgap26

1435695_a_at 1.28633 RIKEN cDNA A030007L17 gene A030007L17Rik

1435696_s_at 0.88717 RIKEN cDNA 3110031B13 gene 3110031B13Rik

1435697_a_at 0.77238 "pleckstrin homology, Sec7 and coiled-coil domains, binding protein" Pscdbp

1435698_at 1.11601 RIKEN cDNA 4921505C17 gene 4921505C17Rik

1435699_at 0.94705 protein phosphatase 1 (formerly 2C)-like Ppm1l

1435700_at 0.83715 talin 2 Tln2

1435701_at 1.04140 "DnaJ (Hsp40) homolog, subfamily B, member 6" Dnajb6

1435702_s_at 0.91348 "tyrosine 3-monooxygenase/tryptophan 5-monooxygenase activation protein,

epsilon polypeptide" Ywhae

1435703_at 0.99487 Transcribed locus ---
 1435704_at 0.97611 RIKEN cDNA C920006O11 gene C920006O11Rik
 1435705_at 0.97408 similar to hypothetical protein FLJ12895 LOC232875
 1435706_at 0.79351 BTB (POZ) domain containing 9 Btbd9
 1435707_at 1.12235 asparaginyl-tRNA synthetase 2 (mitochondrial)(putative) Nars2
 1435708_at 1.54128 Glutaminase Gls
 1435709_at 1.12013 "signal sequence receptor, delta" Ssr4
 1435710_at 0.92941 expressed sequence AI661384 AI661384
 1435711_at 0.89155 --- ---
 1435712_a_at 1.03179 ribosomal protein S18 Rps18
 1435713_at 1.01852 methyltransferase like 2 Mettl2
 1435714_x_at 1.10315 interleukin 17D Il17d
 1435715_x_at 1.03112 thymine DNA glycosylase /// similar to thymine DNA glycosylase isoform 2 ///
 similar to G/T mismatch-specific thymine DNA glycosylase (C-JUN leucine zipper interactive protein
 JZA-3) /// similar to thymine DNA glycosylase isoform 2 /// similar to G/T mismatch-specific thymine
 DNA glycosylase (C-JUN leucine zipper interactive protein JZA-3) Tdg /// LOC545124 ///
 LOC624784 /// LOC670176 /// LOC672659
 1435716_x_at 1.07048 small nuclear ribonucleoprotein N Snrpn
 1435717_at 1.04875 RIKEN cDNA 4833428C12 gene 4833428C12Rik
 1435718_at 0.90228 "adaptor-related protein complex 3, sigma 2 subunit" Ap3s2
 1435719_at 0.90329 expressed sequence AI448984 AI448984
 1435720_at 0.89292 "potassium voltage-gated channel, Shal-related family, member 3" Kcnd3
 1435721_at 1.00106 "potassium voltage-gated channel, subfamily Q, member 4" Kcnq4
 1435722_at 1.02663 "glutamate receptor, ionotropic, AMPA4 (alpha 4)" Gria4
 1435723_at 0.91521 hypothetical protein A530089L17 A530089L17
 1435724_at 1.11730 RIKEN cDNA 4933400N17 gene 4933400N17Rik
 1435725_x_at 1.06823 similar to 60S ribosomal protein L12 /// ribosomal protein L12 /// similar to
 60S ribosomal protein L12 /// similar to 60S ribosomal protein L12 /// similar to 60S ribosomal protein
 L12 /// similar to 60S ribosomal protein L12 /// similar to 60S ribosomal protein L12 /// similar to 60S
 ribosomal protein L12 /// similar to 60S ribosomal protein L12 /// similar to 60S ribosomal protein
 L12 /// similar to 60S ribosomal protein L12 /// similar to 60S ribosomal protein
 L12 /// similar to 60S ribosomal protein L12 /// similar to 60S ribosomal protein
 L12 /// similar to 60S ribosomal protein L12 LOC241053 /// Rpl12 /// LOC546695 /// LOC622589 ///
 LOC630855 /// LOC633363 /// LOC633570 /// LOC639471 /// LOC639477 /// LOC668706 ///
 LOC668844 /// LOC671640 /// LOC672158 /// LOC672173 /// LOC672854 /// LOC676326
 1435726_at 1.08172 LIM domain and actin binding 1 Lima1
 1435727_s_at 0.98656 LIM domain and actin binding 1 Lima1
 1435728_at 0.99063 tRNA-yW synthesizing protein 3 homolog (S. cerevisiae) Tyw3
 1435729_at 1.02174 mitogen activated protein binding protein interacting protein Mapbpip
 1435730_at 0.77209 RIKEN cDNA D930026N18 gene D930026N18Rik
 1435731_x_at 0.90384 Stromal antigen 1 Stag1
 1435732_x_at 1.05440 "ATPase, H+ transporting, lysosomal V0 subunit C" Atp6v0c

1435733_x_at 0.97153 RIKEN cDNA 1500026D16 gene 1500026D16Rik
1435734_x_at 1.03776 dihydrouridine synthase 1-like (*S. cerevisiae*) Dus11
1435735_x_at 0.85700 histocompatibility 47 H47
1435736_x_at 0.96551 "RAB, member of RAS oncogene family-like 4" Rabl4
1435737_a_at 0.96629 nuclear distribution gene E homolog 1 (*A nidulans*) Nde1
1435738_x_at 0.97993 small EDRK-rich factor 2 Serf2
1435739_at 1.02071 expressed sequence AW208599 AW208599
1435740_at 0.79991 Retinoic acid induced 17 Rai17
1435741_at 1.06366 phosphodiesterase 8B Pde8b
1435742_at 1.01080 RIKEN cDNA 1110034C04 gene 1110034C04Rik
1435743_at 0.97390 kelch-like 23 (*Drosophila*) Khl23
1435744_at 0.97620 RIKEN cDNA 6720401G13 gene 6720401G13Rik
1435745_at 0.74571 RIKEN cDNA 5031439G07 gene 5031439G07Rik
1435746_at 1.08855 serine/arginine-rich protein specific kinase 2 Srpk2
1435747_at 0.96989 fibroblast growth factor 14 Fgf14
1435748_at 1.09342 guanine deaminase Gda
1435749_at 1.00207 guanine deaminase Gda
1435750_at 1.34841 GTP cyclohydrolase I feedback regulator Gchfr
1435751_at 1.08836 "ATP-binding cassette, sub-family C (CFTR/MRP), member 9" Abcc9
1435752_s_at 0.86692 "ATP-binding cassette, sub-family C (CFTR/MRP), member 9" Abcc9
1435753_a_at 1.02855 nuclear casein kinase and cyclin-dependent kinase substrate 1 Nucks1
1435754_at 0.96078 RIKEN cDNA 2810482G21 gene 2810482G21Rik
1435755_at 0.95902 RIKEN cDNA 1110001A16 gene 1110001A16Rik
1435756_at 0.89232 sterile alpha motif domain containing 10 Samd10
1435757_a_at 0.93665 ubiquinol cytochrome c reductase core protein 2 Uqcrc2
1435758_at 1.08353 "similar to Beta-1,4-galactosyltransferase 6 (Beta-1,4-GalTase 6) (Beta4Gal-T6) (b4Gal-T6) (UDP-galactose:beta-N-acetylglucosamine beta-1,4-galactosyltransferase 6) (UDP-Gal:beta-GlcNAc beta-1,4-galactosyltransferase 6)" LOC675709
1435759_at 0.92629 cytidine 5'-triphosphate synthase 2 Ctps2
1435760_at 0.95961 cystatin A Csta
1435761_at 0.92305 stefin A3 Stfa3
1435762_at 0.88862 phosphofurin acidic cluster sorting protein 1 Pacs1
1435763_at 0.93635 "TBC1 domain family, member 16" Tbc1d16
1435764_a_at 0.98657 gem (nuclear organelle) associated protein 7 Gemin7
1435765_at 0.94766 RIKEN cDNA E130114P18 gene E130114P18Rik
1435766_at 0.99618 periphilin 1 Pphln1
1435767_at 1.13344 "sodium channel, voltage-gated, type III, beta" Scn3b
1435768_at 0.93436 AT rich interactive domain 4B (*Rbp1* like) Arid4b
1435769_at 0.86913 A kinase (PRKA) anchor protein (*yotiao*) 9 Akap9
1435770_at 0.90343 thioredoxin domain containing 13 Txndc13
1435771_at 0.85104 "phospholipase C, beta 4" Plcb4
1435772_at 0.93781 kinesin family member 21B Kif21b
1435773_at 1.09032 RIKEN cDNA 4930547N16 gene 4930547N16Rik

1435774_at 1.13722 expressed sequence AV024533 AV024533
1435775_at 0.84657 circadian locomoter output cycles kaput Clock
1435776_at 1.02850 RIKEN cDNA 2310009B15 gene 2310009B15Rik
1435777_at 0.98342 RIKEN cDNA E030018N11 gene E030018N11Rik
1435778_at 0.93930 ankyrin repeat domain 11 Ankrd11
1435779_at 1.13259 centrosomal protein 110 Cep110
1435780_at 0.98591 pleckstrin and Sec7 domain containing Psd
1435781_at 0.71015 cullin associated and neddylation disassociated 1 Cand1
1435782_at 1.02301 similar to ubiquitin A-52 residue ribosomal protein fusion product 1
LOC668206
1435783_at 1.09370 RIKEN cDNA B230112C05 gene B230112C05Rik
1435784_at 0.98375 autophagy-related 9A (yeast) Atg9a
1435785_at 0.82521 similar to EH-domain containing 2 LOC673251
1435786_at 0.95662 kelch-like 12 (Drosophila) Klhl12
1435787_at 0.96620 protein phosphatase 1 (formerly 2C)-like Ppm11
1435788_at 0.83283 RIKEN cDNA 2900086B20 gene 2900086B20Rik
1435789_x_at 0.95150 Transcribed locus ---
1435790_at 0.88336 olfactomedin 2 Olfm2
1435791_x_at 1.03006 ribosomal protein L17 Rpl17
1435792_at 1.01766 component of Sp100-rs /// similar to component of Sp100-rs Csprs ///
LOC665338
1435793_at 0.81980 anterior pharynx defective 1b homolog (C. elegans) Aph1b
1435794_at 0.92377 cDNA sequence BC050254 BC050254
1435795_at 0.90916 "galactosidase, beta 1" Glb1
1435796_at 1.00427 RIKEN cDNA C530024P05 gene C530024P05Rik
1435797_at 1.00940 "DNA segment, Chr 5, Wayne State University 178, expressed" D5Wsu178e
1435798_a_at 0.87282 "splicing factor, arginine/serine-rich 14" Sfrs14
1435799_at 0.94099 "splicing factor, arginine/serine-rich 14" Sfrs14
1435800_a_at 1.03059 cold shock domain protein A Csda
1435801_at 0.89532 Fukuyama type congenital muscular dystrophy homolog (human) Fcmd
1435802_at 0.92584 zinc finger protein 499 Zfp499
1435803_a_at 1.10541 eukaryotic translation initiation factor 4E member 2 Eif4e2
1435804_at 0.91756 eukaryotic translation initiation factor 4E member 2 Eif4e2
1435805_at 0.98255 lin-7 homolog A (C. elegans) Lin7a
1435806_at 1.02282 --- ---
1435807_at 1.20783 cell division cycle 42 homolog (S. cerevisiae) Cdc42
1435808_at 0.97172 RIKEN cDNA A230051G13 gene A230051G13Rik
1435809_at 0.88780 ankyrin repeat domain 34 Ankrd34
1435810_at 1.09144 RIKEN cDNA 5730455O13 gene 5730455O13Rik
1435811_a_at 1.02281 unc-50 homolog (C. elegans) Unc50
1435812_at 0.99933 unc-50 homolog (C. elegans) Unc50
1435813_at 1.07569 myopalladin Mypn
1435814_at 1.11273 exportin 7 Xpo7

1435815_at 0.95535 "leucine zipper, down-regulated in cancer 1" Ldoc1
 1435816_at 1.17295 ribosomal protein S6 /// similar to 40S ribosomal protein S6 /// similar to 40S
 ribosomal protein S6 /// similar to 40S ribosomal protein S6 /// similar to 40S ribosomal protein S6 ///
 similar to 40S ribosomal protein S6 /// similar to 40S ribosomal protein S6 /// similar to 40S ribosomal
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 ribosomal protein S6 /// similar to 40S ribosomal protein S6 /// similar to 40S ribosomal protein S6 /// similar to 40S
 Rps6 /// LOC214738 /// LOC216036 /// LOC384383 /// LOC545640 /// LOC626259 /// LOC629575 ///
 LOC638497 /// LOC639124 /// LOC639593 /// LOC639750 /// LOC667717 /// LOC667739 ///
 LOC667922 /// LOC671432 /// LOC672620 /// LOC675985 /// LOC676999
 1435817_x_at 1.09296 ribosomal protein S6 /// similar to 40S ribosomal protein S6 /// similar to 40S
 ribosomal protein S6 /// similar to 40S ribosomal protein S6 /// similar to 40S ribosomal protein S6 ///
 similar to 40S ribosomal protein S6 /// similar to 40S ribosomal protein S6 /// similar to 40S ribosomal
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 ribosomal protein S6 /// similar to 40S ribosomal protein S6 /// similar to 40S ribosomal protein S6 /// similar to 40S
 ribosomal protein S6 /// similar to 40S ribosomal protein S6 /// similar to 40S ribosomal protein S6 /// similar to 40S
 Rps6 /// LOC214738 /// LOC216036 /// LOC384383 /// LOC545640 /// LOC626259 /// LOC629575 ///
 LOC638497 /// LOC639124 /// LOC639593 /// LOC639750 /// LOC667717 /// LOC667739 ///
 LOC667922 /// LOC671432 /// LOC672620 /// LOC675985 /// LOC676999
 1435818_at 1.00299 "Adult male corpus striatum cDNA, RIKEN full-length enriched library, clone:
 C030020B06 product:unclassifiable, full insert sequence" ---
 1435819_at 1.37830 RIKEN cDNA 4922503N01 gene 4922503N01Rik
 1435820_x_at 0.90749 "discoidin domain receptor family, member 1" Ddr1
 1435821_s_at 1.00187 "protein phosphatase 1, regulatory (inhibitor) subunit 8" Ppp1r8
 1435822_at 0.96241 RIKEN cDNA D830012I24 gene D830012I24Rik
 1435823_x_at 1.06256 EGF-like domain 7 Egfl7
 1435824_at 0.97962 YY1 transcription factor Yy1
 1435825_at 1.05464 "activin A receptor, type II-like 1" Acvrl1
 1435826_at 0.99264 RAD18 homolog (*S. cerevisiae*) Rad18
 1435827_at 0.99553 RIKEN cDNA 4933404O12 gene 4933404O12Rik
 1435828_at 1.06208 avian musculoaponeurotic fibrosarcoma (v-maf) AS42 oncogene homolog
 Maf
 1435829_at 0.91448 RIKEN cDNA B930008K04 gene B930008K04Rik
 1435830_a_at 0.84176 RIKEN cDNA 5430435G22 gene 5430435G22Rik
 1435831_at 1.40608 uroplakin 1B Upk1b
 1435832_at 1.08473 leucine rich repeat containing 4 Lrrc4
 1435833_at 1.18121 --- ---
 1435834_at 0.86267 growth arrest specific 2 Gas2
 1435835_at 0.93364 "Meckel syndrome, type 1" Mks1
 1435836_at 0.89156 "pyruvate dehydrogenase kinase, isoenzyme 1" Pdk1
 1435837_at 1.05154 "crystallin, alpha B" Cryab

1435838_at 1.03116 alanine-glyoxylate aminotransferase 2 Agxt2
1435839_at 1.02508 Transcribed locus ---
1435840_x_at 0.95798 "similar to 2-cell-stage, variable group, member 3" LOC625360
1435841_s_at 0.91836 "succinate-Coenzyme A ligase, GDP-forming, beta subunit" Suclg2
1435842_at 0.98395 RIKEN cDNA 1110038O08 gene 1110038O08Rik
1435843_x_at 1.03082 mitochondrial ribosomal protein S9 Mrps9
1435844_at 0.94704 RIKEN cDNA A330009N23 gene A330009N23Rik
1435845_at 0.90469 "B16 F10Y cells cDNA, RIKEN full-length enriched library, clone:
G370036C05 product:unclassifiable, full insert sequence" ---
1435846_x_at 0.93761 --- ---
1435847_at 1.06895 Cdc42 binding protein kinase alpha Cdc42bpa
1435848_at 0.98396 RIKEN cDNA D430041D05 gene D430041D05Rik
1435849_at 0.93716 RIKEN cDNA 6330417G02 gene 6330417G02Rik
1435850_at 0.96503 Transcribed locus ---
1435851_at 0.91655 "leucine-rich repeat LGI family, member 1" Lgi1
1435852_at 1.37964 "sprouty-related, EVH1 domain containing 3" Spred3
1435853_at 1.00937 RIKEN cDNA 9030605E09 gene 9030605E09Rik
1435854_at 1.13113 transmembrane protein 10 Tmem10
1435855_x_at 0.99699 "aldehyde dehydrogenase 18 family, member A1" Aldh18a1
1435856_x_at 0.93684 "SWI/SNF related, matrix associated, actin dependent regulator of chromatin,
subfamily b, member 1" Smarcb1
1435857_s_at 0.77480 amyloid beta (A4) precursor-like protein 1 Aplp1
1435858_at 0.92291 Transcribed locus ---
1435859_x_at 0.99004 "proteasome (prosome, macropain) 26S subunit, ATPase 2" Psmc2
1435860_at 0.96581 "solute carrier family 5 (sodium-dependent vitamin transporter), member 6"
Slc5a6
1435861_at 0.97064 "protein tyrosine phosphatase, receptor type, f polypeptide (PTPRF), interacting
protein, alpha 1" Ppfial
1435862_at 0.94218 Son cell proliferation protein Son
1435863_at 1.09162 COMM domain containing 6 Commd6
1435864_a_at 1.05630 RIKEN cDNA 1810063B05 gene 1810063B05Rik
1435865_at 1.01625 "histone 3, H2a" Hist3h2a
1435866_s_at 0.85530 "histone 3, H2a" Hist3h2a
1435867_at 0.98146 "Visual cortex cDNA, RIKEN full-length enriched library, clone:K230008J08
product:hypothetical protein, full insert sequence" ---
1435868_at 0.93002 ankyrin repeat domain 13c Ankrd13c
1435869_s_at 1.12350 "adaptor protein complex AP-2, alpha 2 subunit" Ap2a2
1435870_at 1.02874 Synaptonemal complex protein 3 Sycp3
1435871_at 0.93428 RIKEN cDNA 2600009P04 gene 2600009P04Rik
1435872_at 0.95743 Transcribed locus ---
1435873_a_at 0.99877 ribosomal protein L13a /// zinc finger protein 526 Rpl13a /// Zfp526
1435874_at 0.94977 --- ---
1435875_at 0.90989 "protein kinase, AMP-activated, beta 2 non-catalytic subunit" Prkab2

1435876_at 0.99019 expressed sequence C86350 C86350
1435877_at 0.79304 serine/threonine kinase 38 like Stk381
1435878_at 1.03339 serine/threonine kinase 38 like Stk381
1435879_at 0.83860 thymoma viral proto-oncogene 3 Akt3
1435880_at 0.89118 ankrin repeat domain 50 Ankrd50
1435881_at 1.06912 poly(rC) binding protein 2 Pcbp2
1435882_at 0.97562 ubiquitin associated protein 2-like Ubap2l
1435883_at 1.09767 expressed sequence AW413431 AW413431
1435884_at 0.95464 intersectin 1 (SH3 domain protein 1A) Itsn1
1435885_s_at 0.94929 intersectin 1 (SH3 domain protein 1A) Itsn1
1435886_at 0.98333 "DnaJ (Hsp40) homolog, subfamily B, member 12" Dnajb12
1435887_at 1.01595 "serine (or cysteine) peptidase inhibitor, clade A (alpha-1 antiproteinase, antitrypsin), member 11" Serpina11
1435888_at 1.06288 epidermal growth factor receptor Egfr
1435889_at 1.03058 MAP/microtubule affinity-regulating kinase 2 Mark2
1435890_at 1.02981 RIKEN cDNA 5730596K20 gene 5730596K20Rik
1435891_x_at 0.98818 RIKEN cDNA 2610021A01 gene /// similar to zinc finger protein 347 2610021A01Rik /// LOC668572
1435892_at 0.99728 additional sex combs like 3 (Drosophila) Asxl3
1435893_at 0.92951 very low density lipoprotein receptor Vldlr
1435894_at 0.99380 hypothetical protein C030014L02 C030014L02
1435895_at 0.97574 RIKEN cDNA D930023J12 gene D930023J12Rik
1435896_at 1.01127 sideroflexin 2 /// similar to Sideroflexin-2 Sfxn2 /// LOC677143
1435897_at 1.02214 ribosomal protein L32 Rpl32
1435898_x_at 0.95823 guanosine diphosphate (GDP) dissociation inhibitor 2 Gdi2
1435899_at 1.01803 RIKEN cDNA 9430079B08 gene 9430079B08Rik
1435900_at 0.89532 zinc finger protein 297B Zfp297b
1435901_at 0.94041 ubiquitin specific peptidase 40 Usp40
1435902_at 1.00187 nudix (nucleoside diphosphate linked moiety X)-type motif 18 Nudt18
1435903_at 0.93527 CD300A antigen Cd300a
1435904_at 1.08292 "eukaryotic translation initiation factor 2C, 3" Eif2c3
1435905_at 0.85379 "DNA segment, Chr 3, ERATO Doi 300, expressed" D3Ert300e
1435906_x_at 1.16110 guanylate nucleotide binding protein 2 Gbp2
1435907_at 0.79251 neurexin II Nrnx2
1435908_at 0.98731 neurexin II Nrnx2
1435909_at 1.01489 RIKEN cDNA C030034I22 gene C030034I22Rik
1435910_at 0.80297 fatty acid desaturase 3 Fads3
1435911_s_at 0.97964 "solute carrier family 2 (facilitated glucose transporter), member 12" Slc2a12
1435912_at 1.01853 UBX domain containing 7 Ubx7
1435913_at 0.77337 "beta-1,4-N-acetyl-galactosaminyl transferase 4" B4galnt4
1435914_at 1.03205 nuclear receptor co-repressor 1 Ncor1
1435915_at 1.07198 jumonji domain containing 4 Jmjd4
1435916_at 1.00212 RIKEN cDNA 2210410P13 gene 2210410P13Rik

1435917_at 0.76004 Transcribed locus ---
1435918_at 0.97301 cDNA sequence BC055107 BC055107
1435919_at 1.01168 "ATPase, Na⁺/K⁺ transporting, alpha 1 polypeptide" Atp1a1
1435920_x_at 1.05299 "ATPase, Na⁺/K⁺ transporting, alpha 1 polypeptide" Atp1a1
1435921_at 0.89427 "DNA segment, Chr 8, ERATO Doi 587, expressed" D8ErtD587e
1435922_at 1.00577 F-box and leucine-rich repeat protein 19 Fbxl19
1435923_at 0.98219 "gene model 237, (NCBI)" Gm237
1435924_at 1.15393 "transcription factor B1, mitochondrial" Tfb1m
1435925_at 0.97748 G protein-coupled receptor kinase-interactor 2 Git2
1435926_at 1.11093 RIKEN cDNA E030003F13 gene E030003F13Rik
1435927_at 0.88185 DENN/MADD domain containing 3 Dennd3
1435928_at 1.01299 --- ---
1435929_at 0.97024 similar to RIKEN cDNA 9630033F20 gene LOC677429
1435930_at 0.90046 zinc finger protein 291 Zfp291
1435931_at 1.15959 RIKEN cDNA E130308A19 gene E130308A19Rik
1435932_at 1.00382 Hermansky-Pudlak syndrome 6 Hps6
1435933_at 0.90258 "sodium channel, voltage-gated, type II, alpha 1" Scn2a1
1435934_at 1.24863 "NADH dehydrogenase (ubiquinone) 1, alpha/beta subcomplex, 1" Ndufab1
1435935_at 0.91403 Transcribed locus ---
1435936_at 0.86191 "Solute carrier family 13 (sodium-dependent citrate transporter), member 5" Slc13a5
1435937_at 0.85349 "serine palmitoyltransferase, long chain base subunit 2" Sptlc2
1435938_at 1.05124 cytoskeleton associated protein 2-like Ckap2l
1435939_s_at 0.96541 expressed sequence AI987662 AI987662
1435940_at 0.72764 expressed sequence AI836758 AI836758
1435941_at 0.80600 "rhomboid, veinlet-like 3 (Drosophila)" Rhbdl3
1435942_at 1.02216 "potassium voltage-gated channel, subfamily Q, member 2" Kcnq2
1435943_at 1.05413 dipeptidase 1 (renal) Dpep1
1435944_s_at 1.16211 centromere protein B Cenpb
1435945_a_at 1.14481 "potassium intermediate/small conductance calcium-activated channel, subfamily N, member 4" Kcnn4
1435946_at 0.89983 "DNA segment, Chr 5, ERATO Doi 135, expressed" D5ErtD135e
1435947_at 1.03382 RIKEN cDNA 2810455D13 gene 2810455D13Rik
1435948_at 1.13635 G protein-coupled receptor 178 Gpr178
1435949_at 0.99291 zinc finger CCCH type containing 3 Zc3h3
1435950_at 0.65833 hairless Hr
1435951_at 0.91261 glutamate receptor interacting protein 1 Grip1
1435952_at 0.76390 --- ---
1435953_at 1.33151 "BTAF1 RNA polymerase II, B-TFIID transcription factor-associated, (Mot1 homolog, S. cerevisiae)" Btaf1
1435954_at 0.87640 zinc finger protein 710 Zfp710
1435955_at 1.25038 sialic acid binding Ig-like lectin G Siglecg
1435956_at 0.92852 RIKEN cDNA A930004K21 gene A930004K21Rik

1435957_at 1.02176 hypothetical protein B830032F12 B830032F12
1435958_at 1.01276 "phosphatidylinositol glycan anchor biosynthesis, class B" Pigb
1435959_at 0.98955 Rho GTPase activating protein 15 Arhgap15
1435960_at 0.99679 Similar to KIAA1183 protein LOC434128
1435961_at 0.89191 cDNA sequence BC047219 BC047219
1435962_at 1.10190 ribosomal protein S6 /// similar to 40S ribosomal protein S6 /// similar to 40S
ribosomal protein S6 /// similar to 40S ribosomal protein S6 /// similar to 40S ribosomal protein S6 ///
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protein S6 /// similar to 40S ribosomal protein S6 /// similar to 40S ribosomal protein S6 /// similar to
40S ribosomal protein S6 /// similar to 40S ribosomal protein S6 /// similar to 40S ribosomal protein
S6 /// similar to 40S ribosomal protein S6 Rps6 /// LOC214738 /// LOC384383 /// LOC545640 ///
LOC626259 /// LOC629575 /// LOC638497 /// LOC639593 /// LOC639750 /// LOC667739 ///
LOC667922 /// LOC671432 /// LOC672620 /// LOC675985
1435963_at 1.00442 --- ---
1435964_a_at 1.30965 TAO kinase 3 Taok3
1435965_at 1.00342 "CCR4-NOT transcription complex, subunit 3" Cnot3
1435966_x_at 1.01472 mitochondrial ribosomal protein L13 Mrpl13
1435967_s_at 0.88717 3-hydroxyisobutyrate dehydrogenase Hibadh
1435968_at 0.86855 "Protein tyrosine phosphatase, receptor type, N polypeptide 2" Ptpn2
1435969_at 1.07530 BTB (POZ) domain containing 12 Btbd12
1435970_at 0.98961 nemo like kinase Nlk
1435971_at 0.93633 regulating synaptic membrane exocytosis 3 Rims3
1435972_at 0.92013 calpastatin Cast
1435973_at 1.05192 Transcribed locus ---
1435974_at 0.94113 Cdc42 guanine nucleotide exchange factor (GEF) 9 Arhgef9
1435975_at 0.92062 expressed sequence AI115600 AI115600
1435976_at 0.87724 --- ---
1435977_at 0.88083 "hepatoma-derived growth factor, related protein 3" Hdgfrp3
1435978_at 1.14046 GDP-mannose pyrophosphorylase A Gmppa
1435979_a_at 1.08234 myosin XVB Myo15b
1435980_x_at 0.96176 wingless-related MMTV integration site 6 Wnt6
1435981_at 0.69467 RIKEN cDNA E430014L09 gene E430014L09Rik
1435982_at 0.96628 syntaxin 12 Stx12
1435983_at 1.06082 RIKEN cDNA 1700019P01 gene 1700019P01Rik
1435984_at 1.17237 RIKEN cDNA 1110033F14 gene 1110033F14Rik
1435985_at 1.08552 "FERM, RhoGEF and pleckstrin domain protein 2" Farp2
1435986_x_at 0.94152 "succinate dehydrogenase complex, subunit C, integral membrane protein"
Sdhc
1435987_x_at 0.91486 RIKEN cDNA 1110059G02 gene 1110059G02Rik
1435988_x_at 0.99148 IK cytokine Ik
1435989_x_at 0.97608 "similar to Keratin, type II cytoskeletal 8 (Cytokeratin 8) (Cytokeratin endo
A) /// similar to Keratin, type II cytoskeletal 8 (Cytokeratin-8) (CK-8) (Keratin-8) (K8) (Cytokeratin
endo A)" LOC434261 /// LOC675884

1435990_at 0.45066 "a disintegrin-like and metallopeptidase (reprolysin type) with thrombospondin type 1 motif, 2" Adamts2

1435991_at 0.84226 "nuclear receptor subfamily 3, group C, member 2" Nr3c2

1435992_at 0.84086 IQ motif containing with AAA domain Iqca

1435993_at 0.95141 zinc finger protein 306-like RP23-298F22.2

1435994_at 1.10669 "potassium voltage-gated channel, subfamily H (eag-related), member 1" Kcnh1

1435995_at 1.14914 mitochondrial ribosomal protein L22 Mrpl22

1435996_at 0.97342 "caspase recruitment domain family, member 11" Card11

1435997_at 0.98657 sperm associated antigen 8 Spag8

1435998_at 0.97136 "gene model 288, (NCBI)" Gm288

1435999_at 0.94576 RIKEN cDNA C630041L24 gene C630041L24Rik

1436000_a_at 1.06500 S-phase kinase-associated protein 2 (p45) Skp2

1436001_at 1.00480 RIKEN cDNA 2310028O11 gene 2310028O11Rik

1436002_at 0.73157 Transcribed locus ---

1436003_at 1.07715 vascular cell adhesion molecule 1 Vcam1

1436004_at 0.98336 "ubiquitin specific peptidase 27, X chromosome" Usp27x

1436005_at 0.95219 "splicing factor, arginine/serine-rich 14" Sfrs14

1436006_at 0.91782 eukaryotic translation initiation factor 2 alpha kinase 1 Eif2ak1

1436007_a_at 0.97264 THUMP domain containing 1 Thumpd1

1436008_at 1.07104 Tumor protein D52 Tpd52

1436009_at 0.84689 ubiquitin specific peptidase 30 Usp30

1436010_at 1.08041 cDNA sequence BC036313 BC036313

1436011_at 1.08631 "engulfment and cell motility 2, ced-12 homolog (C. elegans)" Elmo2

1436012_s_at 0.92668 secernin 2 Scrn2

1436013_at 0.95953 GSG1-like Gsg1l

1436014_a_at 0.98663 RUN and SH3 domain containing 1 Rusc1

1436015_s_at 1.00406 serine/threonine kinase 4 Stk4

1436016_x_at 0.93728 guanosine diphosphate (GDP) dissociation inhibitor 2 Gdi2

1436017_at 1.07919 alanyl-tRNA synthetase like Aarsl

1436018_at 0.95058 RIKEN cDNA 2700083E18 gene 2700083E18Rik

1436019_a_at 1.09444 trafficking protein particle complex 1 Trappc1

1436020_at 1.04570 "DNA segment, Chr 8, ERATO Doi 457, expressed" D8Ertd457e

1436021_at 1.06493 RIKEN cDNA A930031D07 gene A930031D07Rik

1436022_at 0.91097 endonuclease G-like 1 Endogl1

1436023_at 1.02429 BCL2-associated transcription factor 1 Bclaf1

1436024_at 0.88914 RIKEN cDNA A930025D01 gene A930025D01Rik

1436025_at 0.90156 RIKEN cDNA A430106J12 gene A430106J12Rik

1436026_at 0.98287 zinc finger protein 703 Zfp703

1436027_at 1.03807 oxysterol binding protein-like 11 Osbpl11

1436028_at 0.93501 transmembrane protein 33 Tmem33

1436029_at 0.99343 bicaudal C homolog 1 (Drosophila) Biccl

1436030_at 0.91771 cache domain containing 1 Cachd1

1436031_at 0.91420 cache domain containing 1 Cachd1

1436032_at 0.94093 "Bone marrow macrophage cDNA, RIKEN full-length enriched library, clone:
 I830083F22 product:unclassifiable, full insert sequence" ---
 1436033_at 0.84492 cDNA sequence BC031353 BC031353
 1436034_at 0.92936 centrosomal protein 68 Cep68
 1436035_at 0.98849 RIKEN cDNA 3830431G21 gene 3830431G21Rik
 1436036_at 0.94777 Wolf-Hirschhorn syndrome candidate 1 (human) Whsc1
 1436037_at 0.83404 ceramide kinase-like Cerkl
 1436038_a_at 0.96528 "phosphatidylinositol glycan anchor biosynthesis, class P" Pigp
 1436039_at 1.06590 Transcribed locus ---
 1436040_at 1.08131 RIKEN cDNA 2310005L22 gene 2310005L22Rik
 1436041_at 0.94966 --- ---
 1436042_at 0.98757 talin 1 Tln1
 1436043_at 0.95616 "sodium channel, voltage-gated, type VII, alpha" Scn7a
 1436044_at 0.98275 "sodium channel, voltage-gated, type VII, alpha" Scn7a
 1436045_at 0.72026 testis specific 10 Tsga10
 1436046_x_at 1.11855 ribosomal protein L29 /// similar to 60S ribosomal protein L29 /// similar to
 60S ribosomal protein L29 /// similar to 60S ribosomal protein L29 /// similar to 60S ribosomal protein
 L29 /// similar to 60S ribosomal protein L29 /// similar to 60S ribosomal protein L29 /// similar to 60S
 ribosomal protein L29 /// hypothetical protein LOC669999 /// similar to 60S ribosomal protein L29 ///
 similar to 60S ribosomal protein L29 Rpl29 /// LOC433350 /// LOC433941 /// LOC622707 ///
 LOC626772 /// LOC665032 /// LOC666642 /// LOC667466 /// LOC669999 /// LOC670334 ///
 LOC675846
 1436047_at 0.91072 "gene model 672, (NCBI)" Gm672
 1436048_at 0.95537 exocyst complex component 8 Exoc8
 1436049_at 0.88447 RIKEN cDNA 5730589K01 gene 5730589K01Rik
 1436050_x_at 0.79135 hairy and enhancer of split 6 (Drosophila) Hes6
 1436051_at 1.12406 RIKEN cDNA 9630007J19 gene 9630007J19Rik
 1436052_at 0.89206 RIKEN cDNA 1700020O03 gene 1700020O03Rik
 1436053_at 1.01014 "TBC1 domain family, member 22B" Tbc1d22b
 1436054_at 0.89748 RIKEN cDNA 9130227C08Rik gene 9130227C08Rik
 1436055_at 1.11164 leucine rich repeat containing 15 Lrrc15
 1436056_at 0.85650 kinesin family member 13B Kif13b
 1436057_at 0.99856 "GTL2, imprinted maternally expressed untranslated mRNA" Gtl2
 1436058_at 1.43577 radical S-adenosyl methionine domain containing 2 Rsad2
 1436059_at 0.97023 "regulatory factor X, 1 (influences HLA class II expression)" Rfx1
 1436060_at 0.91812 RIKEN cDNA 0710005M24 gene 0710005M24Rik
 1436061_at 0.84431 --- ---
 1436062_at 1.01868 archain 1 Arcn1
 1436063_at 1.04813 lysyl oxidase-like 1 Lox11
 1436064_x_at 1.03447 ribosomal protein S24 Rps24
 1436065_at 0.98819 RIKEN cDNA A130091K22 gene A130091K22Rik
 1436066_at 0.99711 "kalirin, RhoGEF kinase" Kalrn
 1436067_at 0.96199 zinc finger and BTB domain containing 10 /// similar to zinc finger and BTB

domain containing 10 Zbtb10 /// LOC669642

1436068_at 0.89327 zinc finger and BTB domain containing 10 /// similar to zinc finger and BTB domain containing 10 Zbtb10 /// LOC669642

1436069_at 1.17039 "inhibitor of growth family, member 5" Ing5

1436070_at 0.98520 glyoxalase 1 Glo1

1436071_at 0.86249 Ankyrin repeat domain 26 Ankrd26

1436072_at 0.90810 --- ---

1436073_at 0.94559 RIKEN cDNA A630007B06 gene /// similar to oocyte-testis gene 1 A630007B06Rik /// LOC671029

1436074_at 0.96701 cDNA sequence AY078069 AY078069

1436075_at 1.03917 secreted frizzled-related sequence protein 5 Sfrp5

1436076_at 1.09905 "discs, large (Drosophila) homolog-associated protein 1" Dlgap1

1436077_a_at 0.77079 FCH domain only 1 Fcho1

1436078_at 0.90341 FCH domain only 1 Fcho1

1436079_s_at 0.80932 "vesicle-associated membrane protein, associated protein B and C" Vapb

1436080_at 1.02299 expressed sequence AW011738 AW011738

1436081_a_at 1.00734 zinc finger protein 414 Zfp414

1436082_at 0.90034 "Myelin basic protein expression factor 2, repressor" Myef2

1436083_at 0.89157 low density lipoprotein receptor-related protein 3 Lrp3

1436084_at 0.95554 "scratch homolog 1, zinc finger protein (Drosophila)" Sqrt1

1436085_at 0.99779 zinc finger and BTB domain containing 34 Zbtb34

1436086_at 0.92623 RIKEN cDNA 4921526F01 gene 4921526F01Rik

1436087_at 0.98644 dipeptidylpeptidase 10 Dpp10

1436088_at 1.01331 RIKEN cDNA 0910001A06 gene 0910001A06Rik

1436089_at 1.05768 integrator complex subunit 6 Ints6

1436090_at 1.06737 ectonucleotide pyrophosphatase/phosphodiesterase 6 Enpp6

1436091_at 1.10195 RIKEN cDNA 2810022L02 gene 2810022L02Rik

1436092_at 0.77051 "Transcribed locus, moderately similar to XP_574723.1 PREDICTED: similar to LRRGT00097 [Rattus norvegicus]" ---

1436093_at 1.03351 Eph receptor A10 Epha10

1436094_at 1.02977 VGF nerve growth factor inducible Vgf

1436095_at 1.05752 chromodomain helicase DNA binding protein 5 Chd5

1436096_at 1.19829 "Adult male corpora quadrigemina cDNA, RIKEN full-length enriched library, clone:B230209O07 product:unclassifiable, full insert sequence" ---

1436097_x_at 0.85399 Rho GTPase activating protein 9 Arhgap9

1436098_at 1.01350 butyrylcholinesterase Bche

1436099_at 1.03354 expressed sequence AI836003 AI836003

1436100_at 0.90229 SH2 domain containing 5 Sh2d5

1436101_at 1.02358 ring finger protein 24 Rnf24

1436102_at 0.99740 SEC22 vesicle trafficking protein-like C (S. cerevisiae) Sec22c

1436103_at 0.91729 RAB3A interacting protein Rab3ip

1436104_a_at 0.98823 RIKEN cDNA 2310015A05 gene 2310015A05Rik

1436105_at 0.79897 RIKEN cDNA 2310015A05 gene 2310015A05Rik

1436106_x_at 0.89079 RIKEN cDNA 2310015A05 gene 2310015A05Rik
1436107_at 0.83177 "LSM8 homolog, U6 small nuclear RNA associated (*S. cerevisiae*)" Lsm8
1436108_at 0.94018 thioredoxin domain containing 9 Txndc9
1436109_at 0.92361 expressed sequence AI317395 AI317395
1436110_at 0.94556 "10 days neonate medulla oblongata cDNA, RIKEN full-length enriched library, clone:B830012K24 product:unclassifiable, full insert sequence" ---
1436111_at 1.01563 RIKEN cDNA E030011K20 gene E030011K20Rik
1436112_at 0.93367 expressed sequence AI118078 AI118078
1436113_a_at 1.01730 suppression of tumorigenicity 13 St13
1436114_at 1.06223 "gene model 96, (NCBI)" Gm96
1436115_at 1.03714 "gene model 266, (NCBI)" Gm266
1436116_x_at 1.04415 RIKEN cDNA 2900057D21 gene 2900057D21Rik
1436117_at 1.11918 RIKEN cDNA A830010M20 gene A830010M20Rik
1436118_at 0.88086 "vang-like 2 (van gogh, *Drosophila*)" Vangl2
1436119_at 1.06301 "aldehyde dehydrogenase 1 family, member L2" Aldh112
1436120_at 1.05827 "SET domain, bifurcated 2" Setdb2
1436121_a_at 1.01475 non-SMC element 1 homolog (*S. cerevisiae*) Nsmc1
1436122_at 0.91277 zinc finger protein 667 Zfp667
1436123_at 0.99854 bassoon Bsn
1436124_at 1.29914 "phosphate cytidylyltransferase 1, choline, beta isoform" Pcyt1b
1436125_at 1.03282 "DNA segment, Chr 16, ERATO Doi 472, expressed" D16Ertd472e
1436126_at 1.16023 "dynein, axonemal, light intermediate polypeptide 1" Dnali1
1436127_at 0.91941 corticotropin releasing hormone binding protein Crhbp
1436128_at 1.10850 "pleckstrin homology domain containing, family A (phosphoinositide binding specific) member 8" Plekha8
1436129_at 0.89985 RIKEN cDNA 4930523C11 gene /// a disintegrin and metallopeptidase domain 6 4930523C11Rik /// Adam6
1436130_s_at 0.91383 RIKEN cDNA 4930523C11 gene /// a disintegrin and metallopeptidase domain 6 4930523C11Rik /// Adam6
1436131_at 0.88671 sialic acid binding Ig-like lectin H Siglech
1436132_at 0.88159 RIKEN cDNA 9830001H06 gene 9830001H06Rik
1436133_at 1.14045 RIKEN cDNA 0610011N22 gene 0610011N22Rik
1436134_at 0.94515 "sodium channel, voltage-gated, type II, beta" Scn2b
1436135_at 1.11793 RIKEN cDNA 2900075N08 gene 2900075N08Rik
1436136_at 1.59393 nucleolar complex associated 2 homolog (*S. cerevisiae*) Noc21
1436137_at 0.97598 "solute carrier family 6 (neurotransmitter transporter), member 17" Slc6a17
1436138_at 0.96962 --- ---
1436139_at 0.95549 "Adult male medulla oblongata cDNA, RIKEN full-length enriched library, clone:6330445K22 product:unclassifiable, full insert sequence" ---
1436140_at 1.00952 RIKEN cDNA 1700027J05 gene 1700027J05Rik
1436141_at 0.94096 "zinc finger, matrin type 5" Zmat5
1436142_at 0.96536 RIKEN cDNA 3526401B18 gene 3526401B18Rik
1436143_at 0.89094 RIKEN cDNA 4933425L03 gene 4933425L03Rik

1436144_at 0.90459 ribosomal protein L31 /// similar to ribosomal protein L31 /// similar to ribosomal protein L31 /// similar to ribosomal protein L31 /// similar to ribosomal protein L31 /// similar to ribosomal protein L31 /// similar to ribosomal protein L31 /// similar to ribosomal protein L31 /// similar to ribosomal protein L31 /// similar to ribosomal protein L31 /// similar to ribosomal protein L31 /// similar to ribosomal protein L31 /// similar to ribosomal protein L31 /// similar to ribosomal protein L31 /// similar to ribosomal protein L31 /// similar to ribosomal protein L31 Rpl31 /// LOC432681 /// LOC625917 /// LOC626367 /// LOC631731 /// LOC637860 /// LOC638399 /// LOC667682 /// LOC668409 /// LOC669190 /// LOC672957 /// LOC675096 /// LOC676592
1436145_at 1.27421 "Insulin-like growth factor 2, binding protein 3" Igf2bp3
1436146_at 1.05743 RIKEN cDNA 4930481A15 gene 4930481A15Rik
1436147_at 1.04574 Sorting nexin 19 Snx19
1436148_at 0.96594 "Adult male olfactory brain cDNA, RIKEN full-length enriched library, clone:6430531K17 product:unclassifiable, full insert sequence" ---
1436149_at 0.95922 "cytochrome c oxidase, subunit Vb" Cox5b
1436150_at 0.89352 RIKEN cDNA 1700066J24 gene 1700066J24Rik
1436151_x_at 0.97190 cDNA sequence BC031781 BC031781
1436152_a_at 1.03996 hepatitis B virus x interacting protein Hbxip
1436153_a_at 0.93320 "zinc finger, MYND domain containing 11" Zmynd11
1436154_at 1.11058 "0 day neonate thymus cDNA, RIKEN full-length enriched library, clone: A430066P15 product:hypothetical protein, full insert sequence" ---
1436155_at 1.06825 nicotinamide nucleotide adenyltransferase 2 Nmnat2
1436156_at 1.14217 cell division cycle and apoptosis regulator 1 Ccar1
1436157_at 1.09304 cell division cycle and apoptosis regulator 1 Ccar1
1436158_at 1.13099 eukaryotic translation initiation factor 4E binding protein 2 Eif4ebp2
1436159_at 1.01511 ubiquitin specific peptidase 32 Usp32
1436160_at 0.93513 keratin 26 Krt26
1436161_at 0.83655 androgen-induced proliferation inhibitor Aprin
1436162_at 0.99788 RIKEN cDNA C730048C13 gene /// similar to integral membrane transport protein UST1R /// similar to integral membrane transport protein UST1R /// similar to integral membrane transport protein UST1R C730048C13Rik /// LOC434674 /// LOC623195 /// LOC674021
1436163_at 1.13637 "potassium inwardly-rectifying channel, subfamily J, member 16" Kcnj16
1436164_at 0.86922 "solute carrier family 30 (zinc transporter), member 1" Slc30a1
1436165_at 1.26252 LUC7-like 2 (S. cerevisiae) Luc7l2
1436166_at 0.96674 storkhead box 2 Stox2
1436167_at 0.96209 Src homology 2 domain containing F Shf
1436168_at 0.77961 RIKEN cDNA C730029A08 gene C730029A08Rik
1436169_at 0.88533 RIKEN cDNA C730029A08 gene C730029A08Rik
1436170_a_at 1.13705 casein alpha s2-like A Csn1s2a
1436171_at 0.99162 Rho GTPase activating protein 30 Arhgap30
1436172_at 1.08395 hypothetical protein 9530028C05 9530028C05
1436173_at 0.99637 deleted in liver cancer 1 Dlc1
1436174_at 1.05171 "ATPase family, AAA domain containing 2" Atad2
1436175_at 1.08934 RIKEN cDNA A430107N12 gene A430107N12Rik
1436176_at 1.12620 synaptotagmin XI Syt11

1436177_at 0.97635 Transcribed locus ---
1436178_at 0.79392 leprecan-like 1 Leprel1
1436179_a_at 1.10727 "DnaJ (Hsp40) homolog, subfamily C, member 5" Dnajc5
1436180_at 1.04483 "DnaJ (Hsp40) homolog, subfamily C, member 5" Dnajc5
1436181_at 0.92391 development and differentiation enhancing factor 2 Ddef2
1436182_at 0.75046 "10 days embryo whole body cDNA, RIKEN full-length enriched library, clone:2610306G12 product:unclassifiable, full insert sequence" ---
1436183_at 1.08076 RIKEN cDNA 9830115L13 gene 9830115L13Rik
1436184_at 1.14153 "gene model 632, (NCBI)" Gm632
1436185_at 0.78273 expressed sequence AI314180 AI314180
1436186_at 0.94343 E2F transcription factor 8 E2f8
1436187_at 0.91997 RIKEN cDNA 1110054M08 gene 1110054M08Rik
1436188_a_at 0.93027 N-myc downstream regulated gene 4 NdrG4
1436189_at 1.10695 "NAD(P)H dehydrogenase, quinone 2" Nqo2
1436190_at 0.88092 "neural precursor cell expressed, developmentally down-regulated gene 10" Nedd10
1436191_at 0.91379 AT rich interactive domain 4A (Rbp1 like) Arid4a
1436192_at 1.06325 ADP-ribosylation factor guanine nucleotide-exchange factor 2 (brefeldin A-inhibited) /// similar to ADP-ribosylation factor guanine nucleotide-exchange factor 2 (brefeldin A-inhibited) Arfgef2 /// LOC670796
1436193_at 0.87031 "mannosidase, alpha, class 1C, member 1" Man1c1
1436194_at 1.17262 RIKEN cDNA C330008K14 gene C330008K14Rik
1436195_at 0.80833 cDNA sequence BC046404 BC046404
1436196_at 1.26151 hypothetical protein C030046G05 C030046G05
1436197_at 1.00663 CDC42 binding protein kinase gamma (DMPK-like) Cdc42bpg
1436198_at 0.89452 prolyl endopeptidase Prep
1436199_at 1.05821 --- ---
1436200_at 1.04126 LON peptidase N-terminal domain and ring finger 3 Lonrf3
1436201_x_at 1.05295 myelin basic protein Mbp
1436202_at 1.24702 --- ---
1436203_a_at 0.86012 RIKEN cDNA 1110059G02 gene 1110059G02Rik
1436204_at 0.85628 RIKEN cDNA 1110059G02 gene 1110059G02Rik
1436205_at 0.93185 Transcribed locus ---
1436206_at 1.06960 F-box protein 10 Fbxo10
1436207_at 0.79631 transcription factor 3 Tcf3
1436208_at 0.95124 ankyrin repeat and SOCS box-containing protein 1 Asb1
1436209_at 1.07672 "DnaJ (Hsp40) homolog, subfamily C, member 16" Dnajc16
1436210_at 0.87996 RIKEN cDNA C330018K18 gene C330018K18Rik
1436211_at 1.13679 THO complex 4 Thoc4
1436212_at 0.84181 transmembrane protein 71 Tmem71
1436213_a_at 0.88210 RIKEN cDNA 1110028C15 gene 1110028C15Rik
1436214_at 0.88355 RIKEN cDNA 1110028C15 gene 1110028C15Rik
1436215_at 1.08508 inositol polyphosphate multikinase Ipmk

1436216_s_at 1.10650 RIKEN cDNA 2610204M08 gene 2610204M08Rik
1436217_at 0.99165 zinc finger protein 148 Zfp148
1436218_at 0.82466 similar to leucine-rich repeat-containing G protein-coupled receptor 6 isoform 1 LOC676300
1436219_at 0.94137 RIKEN cDNA 4933403F05 gene 4933403F05Rik
1436220_at 0.85103 zinc finger protein 287 Zfp287
1436221_at 1.19115 "DNA segment, Chr 1, ERATO Doi 471, expressed" D1Ert471e
1436222_at 1.22208 growth arrest specific 5 Gas5
1436223_at 1.02462 integrin beta 8 Itgb8
1436224_at 0.85457 "18-day embryo whole body cDNA, RIKEN full-length enriched library, clone:1110061F19 product:unclassifiable, full insert sequence" ---
1436225_at 0.97266 "transient receptor potential cation channel, subfamily M, member 2" Trpm2
1436226_at 0.97680 "transcription elongation factor B (SIII), polypeptide 1" Tceb1
1436227_at 1.10678 Left-right determination factor 2 Lefty2
1436228_at 0.87426 --- ---
1436229_at 1.10726 CDNA sequence BC049806 BC049806
1436230_at 0.91022 G protein-coupled receptor 114 Gpr114
1436231_at 0.93777 RIKEN cDNA 2900052N01 gene 2900052N01Rik
1436232_a_at 0.92128 "GA repeat binding protein, beta 1" Gabpb1
1436233_at 0.97328 expressed sequence AI117581 AI117581
1436234_at 0.93156 RIKEN cDNA 4732471D19 gene 4732471D19Rik
1436235_x_at 0.87120 RIKEN cDNA 4732471D19 gene 4732471D19Rik
1436236_x_at 0.91696 coactosin-like 1 (Dictyostelium) Cot11
1436237_at 0.95951 tetratricopeptide repeat domain 9 Ttc9
1436238_at 0.87158 "leucine-rich repeat LGI family, member 3" Lgi3
1436239_at 1.00393 "solute carrier family 5 (sodium iodide symporter), member 5" Slc5a5
1436240_at 0.87350 sclerostin Sost
1436241_s_at 0.99812 histone cell cycle regulation defective homolog A (S. cerevisiae) Hira
1436242_a_at 0.98827 chemokine-like factor Cklf
1436243_at 0.97627 RIKEN cDNA A930004K21 gene A930004K21Rik
1436244_a_at 0.69211 "transducin-like enhancer of split 2, homolog of Drosophila E(spl)" Tle2
1436245_at 0.95687 ubiquitin specific peptidase 20 Usp20
1436246_at 1.01718 "Adult male medulla oblongata cDNA, RIKEN full-length enriched library, clone:6330566A22 product:unclassifiable, full insert sequence" ---
1436247_at 1.09066 integrator complex subunit 2 Ints2
1436248_at 0.98868 "Adult male spinal cord cDNA, RIKEN full-length enriched library, clone:A330033L11 product:unclassifiable, full insert sequence" ---
1436249_at 1.01231 "tubulin tyrosine ligase-like family, member 4" Ttl4
1436250_at 0.85731 RIKEN cDNA 5430405G05 gene 5430405G05Rik
1436251_at 1.00533 phosphodiesterase 1C Pde1c
1436252_at 0.92685 "solute carrier family 45, member 1" Slc45a1
1436253_at 0.95758 peroxisome biogenesis factor 16 Pex16
1436254_at 1.24800 Transcribed locus ---

1436255_at	0.94196	RIKEN cDNA D830044I16 gene	D830044I16Rik
1436256_at	0.87626	"GrpE-like 2, mitochondrial"	Grpel2
1436257_at	1.41616	"synovial sarcoma translocation, Chromosome 18"	Ss18
1436258_at	1.04980	Cdc42 guanine nucleotide exchange factor (GEF) 9	Arhgef9
1436259_at	1.07231	Ankyrin repeat and SOCS box-containing protein 2	Asb2
1436260_at	0.96694	Transcribed locus	---
1436261_at	0.94690	Dysferlin	Dysf
1436262_x_at	0.85046	OTU domain containing 5	Otud5
1436263_at	1.00503	myelin-associated oligodendrocytic basic protein	Mobp
1436264_at	1.00622	cDNA sequence BC025920	BC025920
1436265_at	1.00968	hypothetical protein 6330405H19	6330405H19
1436266_x_at	0.99990	chromobox homolog 1 (Drosophila HP1 beta)	Cbx1
1436267_a_at	1.00201	FK506 binding protein 12-rapamycin associated protein 1	Frap1
1436268_at	1.11693	dendrin	Ddn
1436269_s_at	1.30986	HtrA serine peptidase 2	Htra2
1436270_at	0.88136	"gene model 740, (NCBI)"	Gm740
1436271_at	0.94408	"cDNA sequence BC085284 /// similar to apolipoprotein L, 3 (predicted)"	BC085284 /// LOC666348
1436272_at	0.77794	RAB3 GTPase activating protein subunit 2	Rab3gap2
1436273_at	0.96111	postmeiotic segregation increased 2 (S. cerevisiae)	Pms2
1436274_at	0.92506	Transcribed locus	---
1436275_at	0.91707	---	---
1436276_at	0.94359	Transcribed locus	---
1436277_at	1.05382	RIKEN cDNA 330010C22 gene	D330010C22Rik
1436278_at	0.94412	---	---
1436279_at	0.96932	Transcribed locus	---
1436280_at	0.87791	zinc finger protein 31	Zfp31
1436281_at	0.96936	Similar to hypothetical protein FLJ38281	LOC245305
1436282_at	1.00237	RIKEN cDNA 4732440D04 gene	4732440D04Rik
1436283_at	0.91550	RIKEN cDNA D630042F21 gene	D630042F21Rik
1436284_s_at	0.72679	zinc finger protein 319	Zfp319
1436285_at	0.90447	coiled-coil domain containing 113	Ccdc113
1436286_at	0.98278	RIKEN cDNA B830017H08 gene	B830017H08Rik
1436287_at	0.93931	"13 days embryo head cDNA, RIKEN full-length enriched library, clone:3110033I20 product:hypothetical protein, full insert sequence"	---
1436288_at	0.97051	RIKEN cDNA 1700049M11 gene	1700049M11Rik
1436289_x_at	1.10907	RIKEN cDNA 1600014C10 gene	1600014C10Rik
1436290_at	0.99771	---	---
1436291_a_at	0.97867	dihydropyrimidinase	Dpys
1436292_a_at	0.96401	ornithine decarboxylase antizyme	Oaz1
1436293_x_at	1.06223	"DNA segment, Chr 1, ERATO Doi 471, expressed"	D1Ertd471e
1436294_at	0.83556	ankyrin repeat domain 29	Ankrd29
1436295_at	1.03917	hypocretin (orexin) receptor 1	Hcrtr1

1436296_x_at 1.01545 "Similar to dynein, axonemal, intermediate polypeptide 2" LOC432611
 1436297_a_at 0.94354 "glutamate receptor, ionotropic, N-methyl D-aspartate-associated protein 1
 (glutamate binding)" Grina
 1436298_x_at 0.97775 "phosphoribosylaminoimidazole carboxylase,
 phosphoribosylaminoribosylaminoimidazole, succinocarboxamide synthetase" Paics
 1436299_at 1.05512 glutaminase Gls
 1436300_at 0.84495 receptor interacting protein kinase 5 Ripk5
 1436301_at 0.87868 receptor interacting protein kinase 5 Ripk5
 1436302_at 0.90150 RIKEN cDNA 2410193C02 gene 2410193C02Rik
 1436303_at 1.01505 myeloid/lymphoid or mixed lineage-leukemia translocation to 4 homolog
 (Drosophila) Mllt4
 1436304_at 1.04698 RIKEN cDNA C030003D03 gene C030003D03Rik
 1436305_at 1.00709 IBR domain containing 1 Ibrdc1
 1436306_at 1.00029 "SAPS domain family, member 1" Saps1
 1436307_at 0.95647 myosin IXa Myo9a
 1436308_at 1.06642 zinc finger protein 292 Zfp292
 1436309_at 0.90043 neuropilin (NRP) and tolloid (TLL)-like 2 Neto2
 1436310_at 1.00027 gem (nuclear organelle) associated protein 5 Gemin5
 1436311_at 0.82074 gem (nuclear organelle) associated protein 5 Gemin5
 1436312_at 0.95449 "zinc finger protein, subfamily 1A, 1 (Ikaros)" Zfpn1a1
 1436313_at 1.05713 SCY1-like 2 (S. cerevisiae) Scyl2
 1436314_at 0.98827 SCY1-like 2 (S. cerevisiae) Scyl2
 1436315_at 0.96773 MYST histone acetyltransferase (monocytic leukemia) 3 Myst3
 1436316_at 0.84955 RIKEN cDNA 9430029L20 gene 9430029L20Rik
 1436317_at 0.97097 RIKEN cDNA D230012E17 gene D230012E17Rik
 1436318_at 1.00722 TAR DNA binding protein Tardbp
 1436319_at 0.96374 sulfatase 1 Sulf1
 1436320_at 0.88022 CDNA clone IMAGE:1328649 ---
 1436321_at 0.92716 "UDP-GlcNAc:betaGal beta-1,3-N-acetylglucosaminyltransferase 7" B3gnt7
 1436322_a_at 0.94589 RIKEN cDNA 2810001A02 gene 2810001A02Rik
 1436323_at 1.00678 RIKEN cDNA 2810001A02 gene 2810001A02Rik
 1436324_at 1.06616 START domain containing 9 /// similar to StAR-related lipid transfer protein 9
 (StARD9) (START domain-containing protein 9) /// similar to StAR-related lipid transfer protein 9
 (StARD9) (START domain-containing protein 9) Stard9 /// LOC668856 /// LOC670354
 1436325_at 1.22973 RAR-related orphan receptor alpha Rora
 1436326_at 0.88293 RAR-related orphan receptor alpha Rora
 1436327_a_at 1.06913 vacuolar protein sorting 53 (yeast) Vps53
 1436328_at 1.02987 RNA (guanine-7-) methyltransferase Rnmt
 1436329_at 1.00989 early growth response 3 Egr3
 1436330_x_at 0.89522 similar to reduced expression 2 /// similar to reduced expression 2
 LOC278757 /// LOC631624
 1436331_at 0.93566 vacuolar protein sorting 13 D (yeast) Vps13d
 1436332_at 0.91339 "heat shock protein, alpha-crystallin-related, B6" Hspb6

1436333_at 1.09083 synaptojanin 1 Synj1
1436334_at 1.03429 synaptojanin 1 Synj1
1436335_at 0.93906 phospholipase C-like 4 Plcl4
1436336_at 1.10704 peptidyl-tRNA hydrolase 1 homolog (*S. cerevisiae*) Pth1
1436337_at 0.90883 cyclin D binding myb-like transcription factor 1 /// RIKEN cDNA 4930420K17
gene Dmtf1 /// 4930420K17Rik
1436338_at 1.27683 --- ---
1436339_at 0.80653 RIKEN cDNA 1810058I24 gene 1810058I24Rik
1436340_at 0.99773 RIKEN cDNA 9630041G16 gene 9630041G16Rik
1436341_at 1.03377 homeobox containing 1 Hmbox1
1436342_a_at 0.98963 "DNA segment, Chr 19, ERATO Doi 721, expressed" D19Ert721e
1436343_at 1.00749 chromodomain helicase DNA binding protein 4 Chd4
1436344_at 1.07793 --- ---
1436345_at 0.77045 RIKEN cDNA 5730559C18 gene 5730559C18Rik
1436346_at 0.92358 CD109 antigen Cd109
1436347_a_at 1.06514 RIKEN cDNA 5530601H04 gene 5530601H04Rik
1436348_at 0.86539 Transcribed locus ---
1436349_at 0.83115 RIKEN cDNA 2700094K13 gene 2700094K13Rik
1436350_at 0.99799 RIKEN cDNA D430039N05 gene D430039N05Rik
1436351_at 1.14840 "coenzyme Q3 homolog, methyltransferase (yeast)" Coq3
1436352_at 1.00901 centrosomal protein 78 Cep78
1436353_at 1.16839 RIKEN cDNA A230046K03 gene A230046K03Rik
1436354_at 0.98790 DAZ interacting protein 1-like Dzip11
1436355_at 1.00373 "fatty acid desaturase domain family, member 6" Fads6
1436356_at 1.26082 "lectin, galactose binding, soluble 7" Lgals7
1436357_at 1.42311 "6 days neonate spleen cDNA, RIKEN full-length enriched library, clone:
F430010L17 product:hypothetical protein, full insert sequence" ---
1436358_at 1.24270 ATP synthase mitochondrial F1 complex assembly factor 1 Atpaf1
1436359_at 0.88705 "16 days neonate cerebellum cDNA, RIKEN full-length enriched library,
clone:9630013F18 product:unclassifiable, full insert sequence" ---
1436360_at 0.92182 GLI-Kruppel family member HKR2 Hkr2
1436361_at 0.85673 vestigial like 2 homolog (*Drosophila*) Vgl12
1436362_x_at 1.41914 RIKEN cDNA 2700079J08 gene 2700079J08Rik
1436363_a_at 0.77005 nuclear factor I/X Nfix
1436364_x_at 0.76702 nuclear factor I/X Nfix
1436365_at 0.89247 zinc finger and BTB domain containing 7C Zbtb7c
1436366_at 0.91924 "protein phosphatase 1, regulatory (inhibitor) subunit 15b" Ppp1r15b
1436367_at 1.40086 "Protein tyrosine phosphatase, receptor type, B" Ptpb
1436368_at 0.79745 "solute carrier family 16 (monocarboxylic acid transporters), member 10"
Slc16a10
1436369_at 0.95761 RIKEN cDNA 2900076A07 gene 2900076A07Rik
1436370_at 0.88488 guanylate cyclase 2c Gucy2c
1436371_at 0.95333 RecQ protein-like Recql

1436372_a_at 0.92298 RIKEN cDNA 2210010A19 gene 2210010A19Rik
1436373_at 1.15080 mitogen activated protein kinase kinase kinase 10 Map3k10
1436374_x_at 1.00890 F11 receptor F11r
1436375_at 0.97302 S1 RNA binding domain 1 Srbd1
1436376_s_at 0.98478 S1 RNA binding domain 1 Srbd1
1436377_at 0.86973 G protein-coupled receptor 137 Gpr137
1436378_at 0.88763 leucine rich repeat (in FLII) interacting protein 2 Lrrfip2
1436379_at 0.78811 CREB regulated transcription coactivator 3 Crtc3
1436380_at 1.23403 Cdc42 binding protein kinase alpha Cdc42bpa
1436381_at 1.04056 "discs, large (Drosophila) homolog-associated protein 3" Dlgap3
1436382_at 1.04576 zinc finger and BTB domain containing 12 Zbtb12
1436383_at 0.90257 complexin 2 Cplx2
1436384_at 1.05939 ribosomal protein S10 /// similar to ribosomal protein S10 /// similar to
ribosomal protein S10 /// similar to ribosomal protein S10 /// similar to ribosomal protein S10 Rps10 ///
LOC621772 /// LOC636539 /// LOC666274 /// LOC668457
1436385_at 0.91532 "glucosidase, alpha; neutral C" Ganc
1436386_x_at 1.03817 Similar to reduced expression 2 LOC195531
1436387_at 1.09467 RIKEN cDNA C330006P03 gene C330006P03Rik
1436388_a_at 0.85133 RIKEN cDNA 3830406C13 gene 3830406C13Rik
1436389_at 0.94336 Transcribed locus ---
1436390_a_at 0.90988 chloride channel CLIC-like 1 Clcc1
1436391_s_at 0.93936 chloride channel CLIC-like 1 Clcc1
1436392_s_at 1.13820 "transcription factor AP-2, gamma" Tcfap2c
1436393_a_at 0.96703 tripartite motif protein 37 Trim37
1436394_at 1.02485 tripartite motif protein 37 Trim37
1436395_at 0.86592 "caspase recruitment domain family, member 6" Card6
1436396_at 0.99168 RIKEN cDNA D430033N04 gene D430033N04Rik
1436397_at 0.95631 cDNA sequence BC027057 BC027057
1436398_at 0.86738 Lymphoid enhancer binding factor 1 Lef1
1436399_s_at 0.93121 Nik related kinase Nrk
1436400_at 0.84414 --- ---
1436401_at 0.90429 --- ---
1436402_at 0.89667 deoxyhypusine hydroxylase/monooxygenase Dohh
1436403_at 0.64978 cDNA sequence BC025575 BC025575
1436404_at 0.85271 TLC domain containing 1 Tlcd1
1436405_at 0.82995 dedicator of cytokinesis 4 Dock4
1436406_at 0.90793 RIKEN cDNA A930007I19 gene A930007I19Rik
1436407_at 1.07544 RIKEN cDNA 9430069J07 gene 9430069J07Rik
1436408_at 1.14109 reprimin-like Rprml
1436409_at 1.22709 "cytochrome c oxidase, subunit VIIIa" Cox8a
1436410_at 0.95894 "expressed sequence AI449023 /// gene model 941, (NCBI)" AI449023 ///
Gm941
1436411_at 1.03110 ATPase type 13A5 Atp13a5

1436412_at 1.06455 "7 days neonate cerebellum cDNA, RIKEN full-length enriched library, clone: A730031J22 product:unclassifiable, full insert sequence" ---

1436413_at 1.16542 fyn-related kinase Frk

1436414_at 0.89164 expressed sequence AW822216 AW822216

1436415_at 0.90301 Transcribed locus ---

1436416_x_at 1.04945 fractured callus expressed transcript 1 Fxc1

1436417_at 0.94030 --- ---

1436418_at 0.98710 RIKEN cDNA 1600027N09 gene 1600027N09Rik

1436419_a_at 1.09377 RIKEN cDNA 1700097N02 gene 1700097N02Rik

1436420_a_at 1.01716 importin 4 Ipo4

1436421_s_at 1.03715 "actin related protein 2/3 complex, subunit 5-like" Arpc51

1436422_at 0.99460 cDNA sequence BC026590 BC026590

1436423_at 0.95363 RIKEN cDNA E430004N04 gene E430004N04Rik

1436424_at 0.87402 RIKEN cDNA 1600020E01 gene 1600020E01Rik

1436425_at 0.99763 ankyrin repeat domain 38 Ankrd38

1436426_at 0.76887 RIKEN cDNA 5730509K17 gene 5730509K17Rik

1436427_at 1.04538 PRP4 pre-mRNA processing factor 4 homolog B (yeast) Prpf4b

1436428_at 0.99216 "cholinergic receptor, nicotinic, beta polypeptide 2 (neuronal)" Chrn2

1436429_at 1.08341 zinc finger protein 606 Zfp606

1436430_at 0.94262 RIKEN cDNA B430105G09 gene B430105G09Rik

1436431_at 1.20909 RIKEN cDNA 1700025G04 gene 1700025G04Rik

1436432_at 0.90845 RIKEN cDNA B230343J05 gene B230343J05Rik

1436433_at 0.99632 cDNA sequence BC049762 BC049762

1436434_at 1.13325 hypothetical protein 9230110J10 9230110J10

1436435_at 0.90977 RIKEN cDNA 9530026P05 gene 9530026P05Rik

1436436_at 0.79058 cornichon homolog 4 (Drosophila) Cnih4

1436437_at 0.96983 "gene model 97, (NCBI)" Gm97

1436438_s_at 0.74433 --- ---

1436439_at 0.91584 hypothetical protein LOC666145 /// hypothetical protein LOC672141 /// hypothetical protein LOC675886 LOC666145 /// LOC672141 /// LOC675886

1436440_at 0.77950 RIKEN cDNA B230107K20 gene B230107K20Rik

1436441_at 0.99571 Prostaglandin E receptor 4 (subtype EP4) Ptger4

1436442_at 1.20400 "protein tyrosine phosphatase, non-receptor type 14" Ptpn14

1436443_a_at 0.91147 KDEL (Lys-Asp-Glu-Leu) containing 1 Kdelc1

1436444_at 1.02819 RIKEN cDNA 6030405A18 gene 6030405A18Rik

1436445_at 1.06519 "gene model 996, (NCBI)" Gm996

1436446_at 1.12167 RIKEN cDNA 2310007O11 gene 2310007O11Rik

1436447_at 1.06853 RIKEN cDNA A630026N12 gene A630026N12Rik

1436448_a_at 0.65489 prostaglandin-endoperoxide synthase 1 Ptgs1

1436449_at 0.94889 "15 days embryo head cDNA, RIKEN full-length enriched library, clone: D930025C07 product:unclassifiable, full insert sequence" ---

1436450_at 0.88801 "DNA segment, Chr 11, Brigham & Women's Genetics 0517 expressed" D11Bwg0517e

1436451_a_at 1.01169 transmembrane emp24 domain trafficking protein 2 Tmed2
1436452_x_at 0.92210 transmembrane emp24 domain trafficking protein 2 Tmed2
1436453_at 0.92763 expressed sequence BB144871 BB144871
1436454_x_at 0.93871 flap structure specific endonuclease 1 Fen1
1436455_at 0.96024 aspartate-beta-hydroxylase Asph
1436456_at 1.07065 RIKEN cDNA 9130023D20 gene 9130023D20Rik
1436457_at 1.37492 RIKEN cDNA F830028O17 gene F830028O17Rik
1436458_at 0.92008 "sema domain, transmembrane domain (TM), and cytoplasmic domain,
(semaphorin) 6A" Sema6a
1436459_at 0.91021 G protein-coupled receptor 161 Gpr161
1436460_at 1.17115 cDNA sequence BC030440 BC030440
1436461_at 1.02519 "Adult male corpora quadrigemina cDNA, RIKEN full-length enriched library,
clone:B230207H11 product:unclassifiable, full insert sequence" ---
1436462_at 1.15057 similar to gonadotropin inducible ovarian transcription factor 1 /// similar to zinc
finger protein 709 /// similar to gonadotropin inducible ovarian transcription factor 1 /// similar to
gonadotropin inducible ovarian transcription factor 1 LOC626832 /// LOC628147 /// LOC668030 ///
LOC668066
1436463_at 1.04055 Janus kinase and microtubule interacting protein 2 Jakmip2
1436464_at 1.03544 "13 days embryo heart cDNA, RIKEN full-length enriched library, clone:
D330003D13 product:unclassifiable, full insert sequence" ---
1436465_at 0.92451 kelch-like 1 (Drosophila) Klhl1
1436466_at 0.95668 RIKEN cDNA 9130227C08Rik gene 9130227C08Rik
1436467_at 0.98403 RIKEN cDNA D230004N17 gene D230004N17Rik
1436468_at 1.00132 "zinc finger, DHHC domain containing 8" Zdhhc8
1436469_at 1.14142 similar to bromodomain containing 7 LOC634327
1436470_at 0.98530 regulating synaptic membrane exocytosis 2 Rims2
1436471_at 0.90916 "RAB36, member RAS oncogene family" Rab36
1436472_at 1.08902 schlafen 9 Slfn9
1436473_at 1.11750 zinc finger protein 248 Zfp248
1436474_at 0.89552 mannose receptor-like precursor Mrcl
1436475_at 0.72295 Transcribed locus ---
1436476_at 0.83983 "DAN domain family, member 5" Dand5
1436477_x_at 0.94679 "RAB2, member RAS oncogene family" Rab2
1436478_at 0.89569 RIKEN cDNA 1810013L24 gene 1810013L24Rik
1436479_a_at 1.12102 dipeptidylpeptidase 7 Dpp7
1436480_at 1.10508 dipeptidylpeptidase 7 Dpp7
1436481_at 1.00977 Transcribed locus ---
1436482_a_at 0.77202 syndecan 3 Sdc3
1436483_at 0.94428 myelin transcription factor 1-like Myt11
1436484_at 1.04752 RIKEN cDNA C030019I05 gene C030019I05Rik
1436485_s_at 0.87361 whirlin Whrn
1436486_x_at 1.03525 Whirlin Whrn
1436487_x_at 1.58366 F-box and WD-40 domain protein 2 Fbxw2

1436488_at 0.93066 --- ---
1436489_x_at 0.92040 expressed sequence C85492 C85492
1436490_x_at 1.02623 "RAN, member RAS oncogene family /// similar to RAN, member RAS
oncogene family /// similar to RAN, member RAS oncogene family /// similar to RAN, member RAS
oncogene family" Ran /// LOC625094 /// LOC626393 /// LOC671251
1436491_at 1.09304 RIKEN cDNA 5830431A10 gene 5830431A10Rik
1436492_x_at 0.96920 RIKEN cDNA 6430514M23 gene 6430514M23Rik
1436493_at 1.03134 similar to Cortixin-1 LOC626281
1436494_x_at 1.02618 TRM1 tRNA methyltransferase 1 homolog (S. cerevisiae) Trmt1
1436495_s_at 0.87991 zinc finger protein 260 Zfp260
1436496_at 1.10148 RIKEN cDNA 1700012O15 gene 1700012O15Rik
1436497_at 0.98181 Insulin-like growth factor 1 Igf1
1436498_at 0.93249 ariadne ubiquitin-conjugating enzyme E2 binding protein homolog 1
(Drosophila) Arih1
1436499_at 0.91726 transmembrane protein 23 Tmem23
1436500_at 0.92701 ribosomal protein S24 Rps24
1436501_at 0.53995 mitochondrial tumor suppressor 1 Mtus1
1436502_at 0.49530 mitochondrial tumor suppressor 1 Mtus1
1436503_at 1.02499 cDNA sequence BC048546 BC048546
1436504_x_at 0.91350 apolipoprotein A-IV Apoa4
1436505_at 1.00180 peptidyl-prolyl isomerase G (cyclophilin G) Ppig
1436506_a_at 1.06267 RIKEN cDNA 1110008H02 gene 1110008H02Rik
1436507_at 0.94267 interleukin-1 receptor-associated kinase 2 Irak2
1436508_at 0.95298 RIKEN cDNA 2410014A08 gene 2410014A08Rik
1436509_at 0.97004 RIKEN cDNA 2410014A08 gene 2410014A08Rik
1436510_a_at 0.98078 leucine rich repeat (in FLII) interacting protein 2 Lrrfip2
1436511_at 1.03285 cDNA sequence BC031781 BC031781
1436512_at 0.94672 ADP-ribosylation factor-like 4C /// similar to ADP-ribosylation factor-like
protein 7 Arl4c /// LOC632433
1436513_at 0.87695 "tetratricopeptide repeat, ankyrin repeat and coiled-coil containing 2" Tanc2
1436514_at 0.77940 glypican 4 Gpc4
1436515_at 0.93880 RIKEN cDNA E030004N02 gene E030004N02Rik
1436516_at 0.93772 abhydrolase domain containing 13 Abhd13
1436517_at 0.97162 "H1 histone family, member X" H1fx
1436518_at 0.94707 Ubiquitin specific peptidase 46 Usp46
1436519_a_at 1.00856 RIKEN cDNA 1110057K04 gene 1110057K04Rik
1436520_at 1.06772 expressed sequence AI450948 AI450948
1436521_at 0.99694 "solute carrier family 36 (proton/amino acid symporter), member 2" Slc36a2
1436522_at 1.02666 mitogen activated protein kinase kinase kinase 3 Map3k3
1436523_s_at 1.05685 RIKEN cDNA 1810022K09 gene 1810022K09Rik
1436524_at 1.36520 RIKEN cDNA 4833438C02 gene 4833438C02Rik
1436525_at 0.97892 "adaptor-related protein complex 3, sigma 2 subunit" Ap3s2
1436526_at 1.12267 "N-acetylglucosamine-1-phosphotransferase, gamma subunit" Gnptg

1436527_at 1.14568 --- ---
1436528_at 0.68710 Kazal-type serine peptidase inhibitor domain 1 Kazald1
1436529_at 1.03843 G protein-coupled receptor 123 Gpr123
1436530_at 0.94232 CDNA clone MGC:107680 IMAGE:6766535 ---
1436531_at 1.04978 methionine aminopeptidase 2 Metap2
1436532_at 0.59348 doublecortin and CaM kinase-like 3 Dcamk13
1436533_at 1.06219 "TROVE domain family, member 2" Trove2
1436534_at 1.06184 "TROVE domain family, member 2" Trove2
1436535_at 1.10859 "TROVE domain family, member 2" Trove2
1436536_at 0.91278 RIKEN cDNA C730015A04 gene C730015A04Rik
1436537_at 0.76098 zinc finger protein 629 Zfp629
1436538_at 2.17121 ankyrin repeat domain 37 Ankrd37
1436539_at 0.79377 calmin Clmn
1436540_at 0.69001 RIKEN cDNA 6720427I07 gene 6720427I07Rik
1436541_at 1.07348 RIKEN cDNA 2310008H09 gene 2310008H09Rik
1436542_at 0.94237 Prostaglandin E receptor 1 (subtype EP1) Ptger1
1436543_at 0.91534 cDNA sequence BC034507 BC034507
1436544_at 0.89721 "ATPase, Class V, type 10D" Atp10d
1436545_at 1.00204 deltex 4 homolog (Drosophila) Dtx4
1436546_at 0.98832 Lix1-like Lix1l
1436547_at 1.02069 "diacylglycerol kinase, epsilon" Dgke
1436548_at 1.11720 RIKEN cDNA 1810012P15 gene 1810012P15Rik
1436549_a_at 1.09729 heterogeneous nuclear ribonucleoprotein A1 Hnrpa1
1436550_at 1.08371 F-box protein 30 Fbxo30
1436551_at 1.06954 fibroblast growth factor receptor 1 Fgfr1
1436552_at 1.07713 janus kinase and microtubule interacting protein 2 Jakmip2
1436553_at 0.98812 lipoma HMGIC fusion partner-like protein 4 Lhfpl4
1436554_at 1.03799 "zinc finger protein, subfamily 1A, 5" Zfpn1a5
1436555_at 1.08139 "solute carrier family 7 (cationic amino acid transporter, y+ system), member 2" Slc7a2
1436556_at 1.02113 transmembrane protein 139 Tmem139
1436557_at 1.01886 keratin 73 Krt73
1436558_at 1.30511 Josephin domain containing 3 Josd3
1436559_a_at 1.24422 "proteasome (prosome, macropain) 26S subunit, non-ATPase, 10" Psmd10
1436560_at 0.83391 expressed sequence AW124694 AW124694
1436561_at 1.02687 suppressor of variegation 3-9 homolog 2 (Drosophila) Suv39h2
1436562_at 1.15232 DEAD (Asp-Glu-Ala-Asp) box polypeptide 58 Ddx58
1436563_at 0.91315 RIKEN cDNA 4932441J04 gene 4932441J04Rik
1436564_at 0.88714 Regulating synaptic membrane exocytosis 4 Rims4
1436565_at 0.94824 CEA-related cell adhesion molecule 10 Ceacam10
1436566_at 0.95607 "Rab40b, member RAS oncogene family" Rab40b
1436567_a_at 1.05289 "NADH dehydrogenase (ubiquinone) 1 alpha subcomplex, 7 (B14.5a)"
Ndufa7

1436568_at 1.01768 junction adhesion molecule 2 Jam2
1436569_at 0.82451 RIKEN cDNA C030045D06 gene C030045D06Rik
1436570_at 0.80055 Transcribed locus ---
1436571_at 1.04807 "Adult male testis cDNA, RIKEN full-length enriched library,
clone:4930525M22 product:unclassifiable, full insert sequence" ---
1436572_at 0.88188 coiled-coil domain containing 45 Ccdc45
1436573_at 0.99071 secernin 3 Scrn3
1436574_at 1.08190 RIKEN cDNA 1700029I01 gene /// similar to zinc finger protein 665 /// similar
to zinc finger protein 665 1700029I01Rik /// LOC666532 /// LOC671566
1436575_at 1.02715 "glutamate receptor ionotropic, NMDA3A" Grin3a
1436576_at 1.14300 RIKEN cDNA A630077B13 gene A630077B13Rik
1436577_at 0.96486 Cdc42 guanine nucleotide exchange factor (GEF) 9 Arhgef9
1436578_at 1.08275 RIKEN cDNA A330104H05 gene A330104H05Rik
1436579_s_at 0.99709 expressed sequence AI839402 AI839402
1436580_at 0.87908 --- ---
1436581_at 0.91440 coiled-coil domain containing 39 Ccdc39
1436582_at 0.80078 "zinc finger, DHHC domain containing 15" Zdhhc15
1436583_at 2.02610 RIKEN cDNA 2600005O03 gene 2600005O03Rik
1436584_at 0.92696 sprouty homolog 2 (Drosophila) Spry2
1436585_at 0.93670 expressed sequence BB182297 BB182297
1436586_x_at 1.10849 ribosomal protein S14 Rps14
1436587_at 1.02963 RIKEN cDNA 1700123D08 gene 1700123D08Rik
1436588_at 0.92153 rotatin Rttm
1436589_x_at 1.00094 protein kinase D2 Prkd2
1436590_at 1.55332 "protein phosphatase 1, regulatory (inhibitor) subunit 3B" Ppp1r3b
1436591_at 0.86992 cDNA sequence BC023744 BC023744
1436592_at 0.91446 "12 days embryo head cDNA, RIKEN full-length enriched library,
clone:3010015L05 product:unclassifiable, full insert sequence" ---
1436593_at 0.95321 RIKEN cDNA 1700016K19 gene 1700016K19Rik
1436594_at 0.90205 zinc finger protein 719 Zfp719
1436595_at 1.05073 RNA binding motif protein 34 Rbm34
1436596_at 0.84079 "H2A histone family, member V" H2afv
1436597_at 0.77488 ankyrin repeat and KH domain containing 1 Ankhd1
1436598_at 1.05820 --- ---
1436599_at 0.92003 Expressed sequence AW493225 AW493225
1436600_at 0.97817 trinucleotide repeat containing 9 Tnrc9
1436601_at 0.88269 F-box protein 46 Fbxo46
1436602_x_at 1.07630 "calcium channel, voltage-dependent, N type, alpha 1B subunit" Cacna1b
1436603_at 0.98347 transducin (beta)-like 2 Tbl2
1436604_at 1.21078 "tubulin tyrosine ligase-like family, member 3" Ttl13
1436605_at 0.95379 transketolase Tkt
1436606_at 1.01260 "Adult male colon cDNA, RIKEN full-length enriched library,
clone:9030003C19 product:unclassifiable, full insert sequence" ---

1436607_a_at 0.84638 RIKEN cDNA C330002I19 gene /// similar to kinase D-interacting substance
220 C330002I19Rik /// LOC633081

1436608_at 0.99075 cDNA sequence BC038822 BC038822

1436609_a_at 0.77380 low density lipoprotein receptor-related protein associated protein 1 Lrpap1

1436610_at 1.00613 ankyrin repeat domain 12 Ankrd12

1436611_at 1.03027 "solute carrier family 39 (zinc transporter), member 12" Slc39a12

1436612_at 1.06120 RIKEN cDNA 2410004I01 gene 2410004I01Rik

1436613_at 0.91014 "Coronin, actin binding protein 6" Coro6

1436614_at 0.92233 expressed sequence AI843639 AI843639

1436615_a_at 0.91112 ornithine transcarbamylase Otc

1436616_at 0.83112 expressed sequence R74740 R74740

1436617_at 0.78709 centrin 4 Ctn4

1436618_at 0.96976 sideroflexin 5 Sfxn5

1436619_at 1.09077 RIKEN cDNA D630045M09 gene D630045M09Rik

1436620_at 0.95386 coiled-coil domain containing 45 Ccdc45

1436621_at 0.78467 P140 gene P140

1436622_at 0.95535 IQ motif and Sec7 domain 2 Iqsec2

1436623_at 0.90392 ectonucleoside triphosphate diphosphohydrolase 7 Entpd7

1436624_at 0.93327 dynamin 3 Dnm3

1436625_at 1.05255 "Fc receptor, IgG, high affinity I" Fcgr1

1436626_at 0.98768 tetraspanin 11 Tspan11

1436627_at 0.92680 "DNA segment, Chr 17, ERATO Doi 663, expressed" D17Ertd663e

1436628_at 0.79270 unc-51-like kinase 4 (C. elegans) Ulk4

1436629_at 0.98456 RIKEN cDNA A830041P22 gene A830041P22Rik

1436630_at 1.03571 Aquaporin 11 Aqp11

1436631_at 1.91058 RIKEN cDNA 2010010M04 gene 2010010M04Rik

1436632_at 1.16610 RIKEN cDNA D130059P03 gene D130059P03Rik

1436633_at 1.05706 BAH domain and coiled-coil containing 1 Bahcc1

1436634_at 0.99846 roundabout homolog 3 (Drosophila) Robo3

1436635_at 0.97731 ubiquitin-conjugating enzyme E2Z (putative) Ube2z

1436636_at 0.96518 --- ---

1436637_at 1.62479 Williams-Beuren syndrome chromosome region 1 homolog (human) Wbscr1

1436638_at 0.82158 RIKEN cDNA 6720401G13 gene 6720401G13Rik

1436639_at 0.99667 bolA-like 2 (E. coli) Bola2

1436640_x_at 0.99649 "1-acylglycerol-3-phosphate O-acyltransferase 1 (lysophosphatidic acid
acyltransferase, delta)" Agpat4

1436641_at 1.03295 expressed sequence AI415730 AI415730

1436642_x_at 1.09843 Myocyte enhancer factor 2D Mef2d

1436643_x_at 1.15333 hepcidin antimicrobial peptide 2 Hamp2

1436644_x_at 0.91658 transmembrane protein 25 Tmem25

1436645_a_at 1.02178 "CCR4-NOT transcription complex, subunit 4" Cnot4

1436646_at 1.08480 Transcribed locus ---

1436647_at 1.11021 tau tubulin kinase 2 Ttbk2

1436648_at	0.91807	nanos homolog 1 (Drosophila)	Nanos1	
1436649_at	1.01200	"zinc finger protein, subfamily 1A, 3 (Aiolos)"	Zfpn1a3	
1436650_at	0.86364	filamin A interacting protein 1	Filip1	
1436651_at	0.96053	Transcribed locus	---	
1436652_at	0.90486	RIKEN cDNA 5830418K08	gene	5830418K08Rik
1436653_at	1.08800	RIKEN cDNA 4833401D15	gene	4833401D15Rik
1436654_at	1.14955	RIKEN cDNA 5830483C08	gene	5830483C08Rik
1436655_at	0.99208	RIKEN cDNA 1300002A08	gene	1300002A08Rik
1436656_at	0.95885	cDNA sequence BC062109	BC062109	
1436657_at	0.89235	hypothetical protein LOC629972	LOC629972	
1436658_at	0.90036	RIKEN cDNA B130024G19	gene	B130024G19Rik
1436659_at	1.17516	Double cortin and calcium/calmodulin-dependent protein kinase-like 1		
Dcamk1l				
1436660_at	1.11793	ribosome binding protein 1	Rrbp1	
1436661_at	1.01594	dipeptidylpeptidase 10	Dpp10	
1436662_at	1.14866	VPS10 domain receptor protein SORCS 1	Sorcs1	
1436663_at	0.97432	---	---	
1436664_a_at	1.01417	"solute carrier family 35 (UDP-galactose transporter), member 2"		Slc35a2
1436665_a_at	0.76581	latent transforming growth factor beta binding protein 4	Ltbp4	
1436666_at	2.68294	Flotillin 2	Flot2	
1436667_at	1.07310	"solute carrier family 6 (neurotransmitter transporter), member 20"		Slc6a20
1436668_at	1.09104	RIKEN cDNA 5730409G15	gene	5730409G15Rik
1436669_at	0.88478	RIKEN cDNA 1700019G17	gene	1700019G17Rik
1436670_x_at	1.04415	RIKEN cDNA 1700019G17	gene	1700019G17Rik
1436671_at	0.86198	hypothetical protein A030011M19	A030011M19	
1436672_at	1.18095	RIKEN cDNA 5830428H23	gene	5830428H23Rik
1436673_at	0.95066	RIKEN cDNA 2610020H08	gene	2610020H08Rik
1436674_at	0.96384	Rap2 interacting protein	Rap2ip	
1436675_at	0.86582	WD repeat domain 63	Wdr63	
1436676_at	0.81106	mitogen-activated protein kinase 8 interacting protein 3		Mapk8ip3
1436677_at	1.41431	RIKEN cDNA 1810032O08	gene	1810032O08Rik
1436678_at	0.78577	---	---	
1436679_at	0.93777	RIKEN cDNA 2210010N04	gene	2210010N04Rik
1436680_s_at	0.95728	damage specific DNA binding protein 2	Ddb2	
1436681_x_at	0.99886	RIKEN cDNA 0610009D07	gene	0610009D07Rik
1436682_at	0.91510	"thymosin, beta 10"	Tmsb10	
1436683_at	1.12766	RIKEN cDNA 1700011M02	gene	1700011M02Rik
1436684_a_at	0.91961	RIO kinase 2 (yeast)	Riok2	
1436685_at	1.16525	hypothetical LOC544888	LOC544888	
1436686_at	0.98370	zinc finger protein 706	Zfp706	
1436687_x_at	0.93664	zinc finger protein 706	Zfp706	
1436688_x_at	1.11888	ribosomal protein L14	Rpl14	
1436689_a_at	0.90680	"aldehyde dehydrogenase 9, subfamily A1"		Aldh9a1

1436690_at 0.94521 LPS-responsive beige-like anchor Lrba
1436691_x_at 1.03804 peroxiredoxin 1 Prdx1
1436692_at 1.03760 "12 days embryo head cDNA, RIKEN full-length enriched library, clone:3000003N14 product:hypothetical protein, full insert sequence" ---
1436693_x_at 0.99687 "solute carrier family 35, member E4" Slc35e4
1436694_s_at 1.00732 neurogenic differentiation 4 Neurod4
1436695_x_at 0.93942 RNA binding motif and ELMO domain 1 Rbed1
1436696_x_at 1.02449 CEA-related cell adhesion molecule 11 Ceacam11
1436697_at 0.89635 Transcribed locus ---
1436698_x_at 1.16244 cDNA sequence BC054438 BC054438
1436699_x_at 1.08381 ribosomal protein L18 /// similar to 60S ribosomal protein L18 Rpl18 /// LOC672898
1436700_a_at 1.11235 DEAD (Asp-Glu-Ala-Asp) box polypeptide 48 Ddx48
1436701_at 1.00836 tripartite motif-containing 61 Trim61
1436702_at 0.88109 RIKEN cDNA 1110038O08 gene 1110038O08Rik
1436703_x_at 0.90275 "small nuclear RNA activating complex, polypeptide 2" Snapc2
1436704_x_at 1.03208 "methylenetetrahydrofolate dehydrogenase (NADP+ dependent), methenyltetrahydrofolate cyclohydrolase, formyltetrahydrofolate synthase" Mthfd1
1436705_at 0.97612 transmembrane protein 32 Tmem32
1436706_at 0.97544 transmembrane protein 32 Tmem32
1436707_x_at 0.97822 barren homolog (Drosophila) Brn1
1436708_x_at 0.94560 minichromosome maintenance deficient 4 homolog (S. cerevisiae) Mcm4
1436709_at 0.90827 RIKEN cDNA C230096C10 gene C230096C10Rik
1436710_at 0.92941 "zinc finger, SWIM domain containing 4" Zswim4
1436711_at 0.95728 ARP5 actin-related protein 5 homolog (yeast) Actr5
1436712_at 0.94325 "phospholipase A2, group IVC (cytosolic, calcium-independent)" Pla2g4c
1436713_s_at 1.02766 "GTL2, imprinted maternally expressed untranslated mRNA" Gtl2
1436714_at 1.04279 LIM domain containing preferred translocation partner in lipoma Lpp
1436715_s_at 0.93362 CDP-diacylglycerol--inositol 3-phosphatidyltransferase (phosphatidylinositol synthase) Cdpt
1436716_at 0.96521 "protein phosphatase 1, regulatory (inhibitor) subunit 14B" Ppp1r14b
1436717_x_at 0.92211 "hemoglobin Y, beta-like embryonic chain" Hbb-y
1436718_at 1.02799 neurexophilin 1 Nxph1
1436719_at 1.07593 "solute carrier family 35, member F1" Slc35f1
1436720_s_at 0.97525 oogenesis 3 Oog3
1436721_x_at 0.98399 oogenesis 3 Oog3
1436722_a_at 1.00070 "actin, beta, cytoplasmic" Actb
1436723_at 1.00092 centromere protein I Cenpi
1436724_a_at 0.90249 ligatin Lgtn
1436725_at 0.85056 RIKEN cDNA E130306D19 gene E130306D19Rik
1436726_s_at 0.86357 "serine palmitoyltransferase, long chain base subunit 1" Sptlc1
1436727_x_at 0.97572 "serine palmitoyltransferase, long chain base subunit 1" Sptlc1
1436728_s_at 1.06285 regulator of telomere elongation helicase 1 Rtel1

1436729_at 0.86942 RIKEN cDNA 2600003E23 gene 2600003E23Rik
1436730_at 1.02371 chloride channel CLIC-like 1 Clcc1
1436731_at 0.88198 zinc finger protein 533 Zfp533
1436732_s_at 0.86766 F-box and WD-40 domain protein 8 Fbxw8
1436733_at 0.90018 RIKEN cDNA E130309F12 gene E130309F12Rik
1436734_at 0.89610 RIKEN cDNA E130309F12 gene E130309F12Rik
1436735_at 0.92539 NOL1/NOP2/Sun domain family 3 Nsun3
1436736_x_at 0.48360 "DNA segment, human D4S114" D0H4S114
1436737_a_at 0.61346 sorbin and SH3 domain containing 1 Sorbs1
1436738_at 1.09737 expressed sequence AI449441 AI449441
1436739_at 1.04685 "angiotensin II receptor, type 1a" Agtr1a
1436740_at 1.48466 RIKEN cDNA A430108E01 gene A430108E01Rik
1436741_at 0.99508 oocyte specific homeobox 1 /// similar to oocyte specific homeobox 3 /// similar
to oocyte specific homeobox 3 /// similar to oocyte specific homeobox 2 /// similar to oocyte specific
homeobox 1 Obox1 /// MGC117731 /// LOC545921 /// LOC666466 /// LOC674153
1436742_a_at 1.00127 "gene model 1967, (NCBI)" Gm1967
1436743_at 1.09140 "immunoglobulin superfamily, member 4" Igsf4d
1436744_x_at 1.05771 F-box and WD-40 domain protein 14 /// similar to F-box and WD-40 domain
protein 14 /// similar to F-box and WD-40 domain protein 14 Fbxw14 /// LOC668758 /// LOC673189
1436745_at 0.84032 SCY1-like 1 binding protein 1 Scyl1bp1
1436746_at 0.95532 WNK lysine deficient protein kinase 1 Wnk1
1436747_at 0.92481 RIKEN cDNA 1110014K08 gene /// hypothetical protein LOC664786 ///
hypothetical protein LOC664849 /// hypothetical protein LOC669054 /// hypothetical protein
LOC672175 1110014K08Rik /// LOC664786 /// LOC664849 /// LOC669054 /// LOC672175
1436748_at 1.01299 RIKEN cDNA 1110014K08 gene 1110014K08Rik
1436749_at 1.11733 mesoderm development candiate 2 Mesdc2
1436750_a_at 0.86749 3-oxoacid CoA transferase 1 Oxct1
1436751_at 0.99258 cDNA sequence BC052883 BC052883
1436752_at 0.92682 TBCC domain containing 1 Tbccd1
1436753_at 0.98052 aarF domain containing kinase 5 Adck5
1436754_at 0.77171 expressed sequence AI839735 /// hypothetical protein LOC638311
AI839735 /// LOC638311
1436755_at 1.05694 inter-alpha (globulin) inhibitor H5 Itih5
1436756_x_at 0.89778 "L-3-hydroxyacyl-Coenzyme A dehydrogenase, short chain" Hadhsc
1436757_a_at 0.94065 "cytochrome c oxidase, subunit VIb polypeptide 1" Cox6b1
1436758_at 1.04679 histone deacetylase 4 Hdac4
1436759_x_at 0.84773 "calponin 3, acidic" Cnn3
1436760_a_at 1.04381 ribosomal protein S8 Rps8
1436761_s_at 1.00343 RIKEN cDNA 1200015N20 gene 1200015N20Rik
1436762_x_at 0.88039 Elongation protein 3 homolog (S. cerevisiae) Elp3
1436763_a_at 0.78402 Kruppel-like factor 9 Klf9
1436764_at 1.09632 par-3 (partitioning defective 3) homolog (C. elegans) Pard3
1436765_at 1.08648 par-3 (partitioning defective 3) homolog (C. elegans) Pard3

1436766_at 1.88066 LUC7-like 2 (S. cerevisiae) Luc712
1436767_at 1.94994 LUC7-like 2 (S. cerevisiae) Luc712
1436768_x_at 0.95889 RIKEN cDNA E130112L23 gene /// similar to quattro E130112L23Rik ///
LOC671456
1436769_at 1.15065 "proteasome (prosome, macropain) subunit, alpha type 1" Psmal1
1436770_x_at 1.07169 "proteasome (prosome, macropain) subunit, alpha type 1" Psmal1
1436771_x_at 1.23441 phosphogluconate dehydrogenase Pgd
1436772_at 1.02718 "Glutamate receptor, ionotropic, AMPA4 (alpha 4)" Gria4
1436773_x_at 0.91674 MARVEL (membrane-associating) domain containing 1 Marveld1
1436774_at 0.92895 Sell (suppressor of lin-12) 1 homolog (C. elegans) Sellh
1436775_a_at 0.98030 ankyrin repeat domain 17 Ankrd17
1436776_x_at 0.80266 "solute carrier family 7 (cationic amino acid transporter, y+ system), member
4" Slc7a4
1436777_at 1.01835 --- ---
1436778_at 0.98370 "cytochrome b-245, beta polypeptide" Cybb
1436779_at 0.87357 "cytochrome b-245, beta polypeptide" Cybb
1436780_at 1.08760 O-linked N-acetylglucosamine (GlcNAc) transferase (UDP-N-
acetylglucosamine:polypeptide-N-acetylglucosaminyl transferase) Ogt
1436781_at 1.51557 "mannosidase 2, alpha B1" Man2b1
1436782_at 0.99467 --- ---
1436783_x_at 0.86161 "tyrosine 3-monooxygenase/tryptophan 5-monooxygenase activation protein,
beta polypeptide" Ywhab
1436784_x_at 0.86037 "splicing factor 3b, subunit 4 /// similar to splicing factor 3b, subunit 4
(predicted) /// similar to splicing factor 3b, subunit 4 (predicted)" Sf3b4 /// LOC634414 /// LOC666113
1436785_a_at 1.06778 RIKEN cDNA 1110069O07 gene 1110069O07Rik
1436786_at 0.79609 RIKEN cDNA 1110069O07 gene 1110069O07Rik
1436787_x_at 1.03002 RIKEN cDNA 1110069O07 gene 1110069O07Rik
1436788_at 1.06306 --- ---
1436789_at 1.01528 cyclin J-like Ccnj1
1436790_a_at 1.11731 SRY-box containing gene 11 Sox11
1436791_at 0.92133 wingless-related MMTV integration site 5A Wnt5a
1436792_at 1.23495 WAS protein homology region 2 domain containing 1 Whdc1
1436793_at 0.97502 suppression of tumorigenicity 18 St18
1436794_at 1.03121 "NACHT, leucine rich repeat and PYD containing 4F" Nalp4f
1436795_at 0.83302 RIKEN cDNA 9630058J23 gene 9630058J23Rik
1436796_at 1.28738 RIKEN cDNA 1110061A14 gene 1110061A14Rik
1436797_a_at 0.94532 surfeit gene 4 Surf4
1436798_at 0.87357 ribosomal protein L9 /// similar to ribosomal protein L9 /// similar to ribosomal
protein L9 /// similar to 60S ribosomal protein L9 /// similar to ribosomal protein L9 /// similar to
ribosomal protein L9 /// similar to 60S ribosomal protein L9 /// similar to ribosomal protein L9
Rpl9 /// LOC432768 /// LOC433546 /// LOC667290 /// LOC670203 /// LOC671349 /// LOC672532 ///
LOC674636
1436799_at 1.08780 RIKEN cDNA D230005D02 gene D230005D02Rik

1436800_at 0.96101 follistatin-like 5 Fstl5
1436801_x_at 0.99328 CDC42 effector protein (Rho GTPase binding) 4 Cdc42ep4
1436802_at 0.99336 interleukin enhancer binding factor 3 Ilf3
1436803_a_at 1.00090 "NADH dehydrogenase (ubiquinone) 1 beta subcomplex, 9" Ndufb9
1436804_s_at 1.06285 SCY1-like 1 (S. cerevisiae) Scyl1
1436805_at 1.14523 RIKEN cDNA 2810457I06 gene 2810457I06Rik
1436806_at 0.89835 tripartite motif-containing 62 Trim62
1436807_x_at 0.87141 tripartite motif-containing 62 Trim62
1436808_x_at 0.93693 "minichromosome maintenance deficient 5, cell division cycle 46 (S. cerevisiae)" Mcm5
1436809_a_at 0.94952 spindlin Spin
1436810_x_at 0.98306 RIKEN cDNA 2900010M23 gene 2900010M23Rik
1436811_at 0.98447 potassium channel tetramerisation domain containing 3 Kctd3
1436812_at 0.96239 fukutin related protein Fkrp
1436813_x_at 1.14739 KH-type splicing regulatory protein Khgrp
1436814_at 0.99103 Two pore segment channel 2 Tpcn2
1436815_x_at 0.98437 Two pore segment channel 2 Tpcn2
1436816_at 1.01467 Nucleoporin 133 Nup133
1436817_at 1.02647 exocyst complex component 5 Exoc5
1436818_a_at 0.91374 Musashi homolog 2 (Drosophila) Msi2
1436819_at 0.92549 septin 6 6-Sep
1436820_at 0.95542 potassium channel tetramerisation domain containing 11 Kctd11
1436821_at 0.98909 "phosphatidylinositol-specific phospholipase C, X domain containing 3" Plcxd3
1436822_x_at 1.05962 hexokinase 1 /// ribosomal protein L17 /// similar to 60S ribosomal protein L17 (L23) (Amino acid starvation-induced protein) (ASI) /// similar to 60S ribosomal protein L17 (L23) (Amino acid starvation-induced protein) (ASI) /// similar to 60S ribosomal protein L17 (L23) (Amino acid starvation-induced protein) (ASI) Hk1 /// Rpl17 /// LOC620155 /// LOC620303 /// LOC676276
1436823_x_at 0.95289 "hemoglobin Y, beta-like embryonic chain" Hbb-y
1436824_x_at 0.95014 ring finger protein 26 /// similar to ring finger protein 26 /// similar to ring finger protein 26 Rnf26 /// LOC668155 /// LOC669536
1436825_a_at 1.04756 RIKEN cDNA A630095E13 gene A630095E13Rik
1436826_at 1.10675 transmembrane and tetratricopeptide repeat containing 3 Tmtc3
1436827_at 0.85127 "gene model 944, (NCBI)" Gm944
1436828_a_at 0.98092 tumor protein D52-like 2 Tpd52l2
1436829_at 1.05825 tripartite motif-containing 67 Trim67
1436830_at 1.10480 MARVEL (membrane-associating) domain containing 1 Marveld1
1436831_at 0.95919 RIKEN cDNA 4932408B21 gene 4932408B21Rik
1436832_at 1.10320 RIKEN cDNA 1110067B18 gene 1110067B18Rik
1436833_x_at 0.94834 tubulin tyrosine ligase-like 1 Ttll1
1436834_x_at 0.97941 "malate dehydrogenase 1, NAD (soluble)" Mdh1
1436835_at 0.94754 v-crk sarcoma virus CT10 oncogene homolog (avian) Crk
1436836_x_at 0.93379 "calponin 3, acidic" Cnn3
1436837_at 1.03487 maelstrom homolog (Drosophila) Mael

1436838_x_at 0.84476 coactosin-like 1 (Dictyostelium) Cot11
1436839_at 0.89522 --- ---
1436840_x_at 0.99509 ribosomal protein L35 Rpl35
1436841_at 1.25873 RIKEN cDNA B230380D07 gene B230380D07Rik
1436842_at 1.06327 RIKEN cDNA B230380D07 gene B230380D07Rik
1436843_at 0.95293 RIKEN cDNA C430048L16 gene C430048L16Rik
1436844_at 1.27924 RIKEN cDNA 6330403M23 gene 6330403M23Rik
1436845_at 1.16064 axin2 Axin2
1436846_x_at 0.94223 "tyrosine 3-monooxygenase/tryptophan 5-monooxygenase activation protein, theta polypeptide /// similar to 14-3-3 protein theta (14-3-3 protein tau) /// similar to 14-3-3 protein theta (14-3-3 protein tau)" Ywhaq /// LOC546165 /// LOC636369
1436847_s_at 1.07647 cell division cycle associated 8 Cdca8
1436848_x_at 1.00225 inositol (myo)-1(or 4)-monophosphatase 1 Impa1
1436849_x_at 0.93851 "glucosidase, alpha, acid" Gaa
1436850_at 1.07527 cellular repressor of E1A-stimulated genes 2 Creg2
1436851_at 0.99430 prostaglandin E receptor 1 (subtype EP1) /// protein kinase N1 Ptger1 /// Pkn1
1436852_at 1.05510 RIKEN cDNA E130308A19 gene E130308A19Rik
1436853_a_at 0.98809 "synuclein, alpha" Snca
1436854_at 1.37818 "transient receptor potential cation channel, subfamily C, member 2" Trpc2
1436855_at 1.23308 "protein tyrosine phosphatase, receptor type, S" Ptpns
1436856_x_at 0.93791 threonyl-tRNA synthetase-like 1 Tarsl1
1436857_at 1.04229 defensin related cryptdin 21 /// defensin related cryptdin 22 Defcr21 /// Defcr22
1436858_at 0.95049 muscleblind-like 2 Mbnl2
1436859_at 1.01855 RIKEN cDNA 2700007P21 gene 2700007P21Rik
1436860_at 1.08609 SUMO1/sentrin specific peptidase 7 Senp7
1436861_at 0.99665 interleukin 7 Il7
1436862_at 0.78314 "double C2, alpha" Doc2a
1436863_at 1.04193 RIKEN cDNA 1700010I14 gene 1700010I14Rik
1436864_at 1.12119 RIKEN cDNA A430090L17 gene A430090L17Rik
1436865_at 1.05189 "solute carrier family 26, member 11" Slc26a11
1436866_at 0.97027 ephrin A5 Efna5
1436867_at 0.77982 sarcalumenin /// similar to sarcalumenin Srl /// LOC669709
1436868_at 0.67973 reticulon 4 receptor-like 1 Rtn4rl1
1436869_at 0.87096 sonic hedgehog Shh
1436870_s_at 0.97095 expressed sequence AU041783 AU041783
1436871_at 0.99326 "splicing factor, arginine/serine-rich 7" Sfrs7
1436872_at 1.27988 "transforming, acidic coiled-coil containing protein 3" Tacc3
1436873_at 1.00777 RIKEN cDNA 4732454E20 gene 4732454E20Rik
1436874_x_at 1.07056 "solute carrier family 25 (mitochondrial carrier, adenine nucleotide translocator), member 5" Slc25a5
1436875_at 0.95973 dynamin 3 Dnm3
1436876_at 1.07421 "DNA segment, Chr 13, Brigham & Women's Genetics 1146 expressed" D13Bwg1146e

1436877_at 1.05061 leucine-rich repeats and calponin homology (CH) domain containing 2 Lrch2
1436878_at 1.06969 "brain and acute leukemia, cytoplasmic" Baalc
1436879_x_at 1.12743 alpha fetoprotein Afp
1436880_at 1.07788 Alpha fetoprotein Afp
1436881_x_at 1.09270 alpha fetoprotein Afp
1436882_at 1.02958 ubiquitin-like 5 /// similar to ubiquitin-like 5 /// similar to ubiquitin-like 5
Ub15 /// LOC435336 /// LOC673926
1436883_at 0.97740 "membrane-bound transcription factor peptidase, site 2" Mbtps2
1436884_x_at 0.89669 Ewing sarcoma breakpoint region 1 Ewsr1
1436885_a_at 0.91503 calcium homeostasis endoplasmic reticulum protein Cherp
1436886_x_at 0.96302 XPA binding protein 2 Xab2
1436887_x_at 0.87277 glutamate-rich WD repeat containing 1 Grwd1
1436888_at 0.88997 nescient helix loop helix 2 Nhlh2
1436889_at 0.99957 "gamma-aminobutyric acid (GABA-A) receptor, subunit alpha 1" Gabra1
1436890_at 1.23192 UDP-N-acteylglucosamine pyrophosphorylase 1-like 1 Uap111
1436891_at 1.08673 ubiquitin specific peptidase 15 Usp15
1436892_at 1.43486 "sprouty-related, EVH1 domain containing 2" Spred2
1436893_a_at 0.96964 membrane-associated ring finger (C3HC4) 7 7-Mar
1436894_at 1.17804 "DNA segment, Chr 1, ERATO Doi 471, expressed" D1Ertd471e
1436895_at 0.96130 "centaurin, delta 1" Centd1
1436896_at 0.93699 "DNA segment, Chr 14, ERATO Doi 231, expressed" D14Ertd231e
1436897_at 0.99920 malignant fibrous histiocytoma amplified sequence 1 Mfhas1
1436898_at 1.38226 splicing factor proline/glutamine rich (polypyrimidine tract binding protein associated) Sfpq
1436899_at 0.89700 RIKEN cDNA 2700019D07 gene 2700019D07Rik
1436900_x_at 0.87628 leptin receptor overlapping transcript Leprot
1436901_at 1.07153 "Mammary gland RCB-0526 Jyg-MC(A) cDNA, RIKEN full-length enriched library, clone:G830022P11 product:unclassifiable, full insert sequence /// Notch gene homolog 4 (Drosophila)" Notch4
1436902_x_at 1.09601 "thymosin, beta 10" Tmsb10
1436903_at 1.00168 ubiquilin 3 Ubqln3
1436904_at 1.10611 thyroid hormone receptor associated protein 1 Thrapp1
1436905_x_at 1.06087 lysosomal-associated protein transmembrane 5 Laptm5
1436906_at 0.95074 ring finger protein 166 Rnf166
1436907_at 0.97002 neuron navigator 1 Nav1
1436908_at 0.89330 pericentriolar material 1 Pcm1
1436909_at 0.96177 RIKEN cDNA B430110G05 gene B430110G05Rik
1436910_at 1.06948 RAS protein activator like 2 Rasal2
1436911_at 0.94982 synovial sarcoma translocation gene on chromosome 18-like 1 Ss1811
1436912_at 0.89390 "calcium channel, voltage-dependent, beta 4 subunit" Cacnb4
1436913_at 1.04156 CDC14 cell division cycle 14 homolog A (S. cerevisiae) Cdc14a
1436914_at 1.08061 "CTD (carboxy-terminal domain, RNA polymerase II, polypeptide A) small phosphatase like 2" Ctdspl2

1436915_x_at 0.91202 lysosomal-associated protein transmembrane 4B Laptm4b
1436916_at 1.04196 transmembrane protein 108 Tmem108
1436917_s_at 0.84436 "G-protein signalling modulator 1 (AGS3-like, C. elegans)" Gpsm1
1436918_at 1.07211 dual-specificity tyrosine-(Y)-phosphorylation regulated kinase 2 Dyrk2
1436919_at 1.00126 Trp53 inducible protein 11 Trp53i11
1436920_at 1.08564 protocadherin 17 Pcdh17
1436921_at 0.92113 "ATPase, Cu++ transporting, alpha polypeptide" Atp7a
1436922_at 0.95552 peptidylprolyl isomerase (cyclophilin) like 5 Ppil5
1436923_at 1.18110 "RAB2B, member RAS oncogene family" Rab2b
1436924_x_at 0.99746 ERO1-like (S. cerevisiae) /// ribosomal protein L31 /// similar to ribosomal
protein L31 /// similar to ribosomal protein L31 /// similar to ribosomal protein L31 /// similar to
ribosomal protein L31 /// similar to ribosomal protein L31 /// similar to ribosomal protein L31 /// similar
to ribosomal protein L31 /// similar to ribosomal protein L31 /// similar to ribosomal protein L31 ///
similar to ribosomal protein L31 /// similar to ribosomal protein L31 /// similar to ribosomal protein
L31 /// similar to ribosomal protein L31 /// similar to ribosomal protein L31 Ero11 /// Rpl31 ///
LOC544868 /// LOC632618 /// LOC638193 /// LOC638399 /// LOC665533 /// LOC665562 ///
LOC667682 /// LOC668850 /// LOC670050 /// LOC670381 /// LOC671205 /// LOC672957 ///
LOC673582 /// LOC676592
1436925_at 0.92765 checkpoint suppressor 1 Ches1
1436926_at 0.98895 "estrogen related receptor, beta" Esrrb
1436927_at 0.94586 cysteine-rich perinuclear theca 4 /// cysteine-rich perinuclear theca 9 /// cysteine-
rich perinuclear theca 10 Cypt4 /// Cypt9 /// Cypt10
1436928_s_at 0.95479 adenylate cyclase 3 Adcy3
1436929_x_at 0.77456 adenylate cyclase 3 Adcy3
1436930_x_at 1.02362 hydroxymethylbilane synthase /// similar to hydroxymethylbilane synthase ///
similar to hydroxymethylbilane synthase Hmbs /// LOC623818 /// LOC639157
1436931_at 0.97237 "regulatory factor X, 4 (influences HLA class II expression)" Rfx4
1436932_at 0.89566 grainyhead-like 3 (Drosophila) Grhl3
1436933_at 2.29424 A kinase (PRKA) anchor protein 13 Akap13
1436934_s_at 0.84711 "aconitase 2, mitochondrial" Aco2
1436935_x_at 1.06701 "chloride channel, nucleotide-sensitive, 1A" Clns1a
1436936_s_at 1.16079 inactive X specific transcripts Xist
1436937_at 0.88123 "RNA binding motif, single stranded interacting protein" Rbms3
1436938_at 0.90402 "RNA binding motif, single stranded interacting protein" Rbms3
1436939_at 0.92535 unc-45 homolog B (C. elegans) Unc45b
1436940_at 0.89051 Purine rich element binding protein B Purb
1436941_at 1.00643 "gene model 1752, (NCBI)" Gm1752
1436942_at 0.93531 RIKEN cDNA A930035D04 gene A930035D04Rik
1436943_at 0.77135 cytochrome b5 domain containing 2 Cyb5d2
1436944_x_at 0.85380 RIKEN cDNA 4933439C20 gene 4933439C20Rik
1436945_x_at 0.98627 stromal interaction molecule 1 Stim1
1436946_s_at 0.95935 "guanine nucleotide binding protein (G protein), gamma 5 subunit" Gng5
1436947_a_at 1.01628 thioredoxin-like 1 Txnl1

1436948_a_at 0.93555 RIKEN cDNA 6430550H21 gene 6430550H21Rik
 1436949_a_at 0.96541 "transcription elongation factor B (SIII), polypeptide 2 /// similar to transcription elongation factor B (SIII), polypeptide 2 /// similar to transcription elongation factor B (SIII), polypeptide 2" Tceb2 /// LOC668554 /// LOC671613
 1436950_at 1.03007 v-crk sarcoma virus CT10 oncogene homolog (avian)-like Crkl
 1436951_x_at 1.03417 thioredoxin domain containing 9 Txndc9
 1436952_at 0.88131 Kruppel-like factor 9 Klf9
 1436953_at 0.78490 Wiskott-Aldrich syndrome protein interacting protein Waspip
 1436954_at 0.82303 Wiskott-Aldrich syndrome protein interacting protein Waspip
 1436955_at 0.95601 tumor suppressing subtransferable candidate 1 Tssc1
 1436956_at 1.00575 Beta-site APP cleaving enzyme 1 Bace1
 1436957_at 0.96777 "Gamma-aminobutyric acid (GABA-A) receptor, subunit alpha 3" Gabra3
 1436958_x_at 0.98126 "Tropomyosin 3, gamma" Tpm3
 1436959_x_at 0.83434 nasal embryonic LHRH factor Nelf
 1436960_at 0.87460 bromodomain containing 3 Brd3
 1436961_at 0.89650 heat shock protein 12A Hspa12a
 1436962_at 0.99119 PR domain containing 6 Prdm6
 1436963_x_at 0.88269 RIKEN cDNA 1810059G22 gene 1810059G22Rik
 1436964_at 0.98753 "DNA segment, Chr 7, ERATO Doi 715, expressed" D7Ertd715e
 1436965_at 0.93181 elastin microfibril interfacier 3 Emilin3
 1436966_at 0.97790 pellino 2 Peli2
 1436967_at 2.03524 ankyrin repeat domain 11 Ankrd11
 1436968_x_at 0.80364 kelch-like 24 (Drosophila) Klhl24
 1436969_at 1.02500 RIKEN cDNA 1700013E18 gene 1700013E18Rik
 1436970_a_at 0.93011 "platelet derived growth factor receptor, beta polypeptide" Pdgfrb
 1436971_x_at 1.09709 "tyrosine 3-monooxygenase/tryptophan 5-monooxygenase activation protein, zeta polypeptide" Ywhaz
 1436972_at 0.95176 G protein-coupled receptor 61 Gpr61
 1436973_at 1.03266 chaperonin subunit 8 (theta) Cct8
 1436974_at 1.14625 RIKEN cDNA A230069A22 gene A230069A22Rik
 1436975_at 0.97223 "Gene model 24, (NCBI)" Gm24
 1436976_a_at 1.10067 RIKEN cDNA 9930028C20 gene 9930028C20Rik
 1436977_at 1.26976 Epidermal growth factor receptor pathway substrate 8 Eps8
 1436978_at 1.11090 wingless-type MMTV integration site 9A Wnt9a
 1436979_x_at 1.08334 RNA binding motif protein 14 Rbm14
 1436980_x_at 1.03527 "CCR4-NOT transcription complex, subunit 2" Cnot2
 1436981_a_at 0.87813 "tyrosine 3-monooxygenase/tryptophan 5-monooxygenase activation protein, zeta polypeptide" Ywhaz
 1436982_at 0.96889 trinucleotide repeat containing 6b Tnrc6b
 1436983_at 0.93891 CREB binding protein Crebbp
 1436984_at 0.99641 abl-interactor 2 Abi2
 1436985_at 0.93790 zinc finger protein 644 Zfp644
 1436986_at 1.04530 "syntrophin, basic 2" Sntb2

1436987_at 0.91612 RIKEN cDNA 5430433G21 gene 5430433G21Rik
1436988_at 1.08589 RIKEN cDNA 5430433G21 gene 5430433G21Rik
1436989_s_at 0.91605 "solute carrier family 12, member 6" Slc12a6
1436990_s_at 0.77539 Nur77 downstream gene 2 Ndg2
1436991_x_at 0.86990 gelsolin Gsn
1436992_x_at 0.96216 voltage-dependent anion channel 1 Vdac1
1436993_x_at 0.82141 profilin 2 Pfn2
1436994_a_at 1.36289 "histone 1, H1c" Hist1h1c
1436995_a_at 1.05476 ribosomal protein L26 Rpl26
1436996_x_at 0.94201 P lysozyme structural Lzp-s
1436997_x_at 0.82302 SH3-binding domain glutamic acid-rich protein like Sh3bgrl
1436998_at 1.03594 ankyrin repeat domain 43 Ankrd43
1436999_at 0.67876 RIKEN cDNA 5033414K04 gene 5033414K04Rik
1437000_at 1.00044 "diacylglycerol kinase, theta" Dgkq
1437001_at 1.01590 glycogen synthase kinase 3 beta Gsk3b
1437002_at 0.94964 RIKEN cDNA C030011O14 gene C030011O14Rik
1437003_at 1.00029 --- ---
1437004_at 0.95797 similar to tripartite motif-containing 43 LOC547109
1437005_a_at 1.06933 ribosomal protein L18 Rpl18
1437006_x_at 0.95375 "beclin 1 (coiled-coil, myosin-like BCL2-interacting protein)" Becn1
1437007_x_at 1.00064 ubiquitin specific peptidase 39 Usp39
1437008_x_at 1.00160 transmembrane protein 109 Tmem109
1437009_a_at 0.93577 zinc finger protein 364 Zfp364
1437010_a_at 1.02255 ornithine decarboxylase antizyme 3 Oaz3
1437011_x_at 0.89919 ornithine decarboxylase antizyme 3 Oaz3
1437012_x_at 1.26243 Rap guanine nucleotide exchange factor (GEF) 3 Rapgef3
1437013_x_at 0.96061 "ATPase, H+ transporting, lysosomal V0 subunit B" Atp6v0b
1437014_x_at 0.98453 similar to Peroxiredoxin 1 (Thioredoxin peroxidase 2) (Thioredoxin-dependent peroxide reductase 2) (Osteoblast specific factor 3) (OSF-3) (Macrophage 23 kDa stress protein) /// similar to Peroxiredoxin-1 (Thioredoxin peroxidase 2) (Thioredoxin-dependent peroxide reductase 2) (Osteoblast-specific factor 3) (OSF-3) (Macrophage 23 kDa stress protein) LOC545161 /// LOC637273
1437015_x_at 1.00854 "phospholipase A2, group IB, pancreas" Pla2g1b
1437016_x_at 0.96367 "RAP2C, member of RAS oncogene family" Rap2c
1437017_at 0.97604 expressed sequence AI480653 AI480653
1437018_at 1.08757 paraneoplastic antigen MA2 Pnma2
1437019_at 0.91634 RIKEN cDNA 2200001I15 gene 2200001I15Rik
1437020_at 1.03565 E1A binding protein p400 Ep400
1437021_at 1.04496 ADP-ribosylation factor-like 13B Arl13b
1437022_at 1.13003 RIKEN cDNA D130059P03 gene D130059P03Rik
1437023_at 1.05608 zinc finger protein 12 Zfp12
1437024_at 0.84772 sphingomyelin phosphodiesterase 4 Smpd4
1437025_at 0.92531 CD28 antigen Cd28

1437026_at 0.95616 cDNA sequence BC057893 BC057893
1437027_x_at 1.01201 ribonucleic acid binding protein S1 Rnps1
1437028_at 0.99898 surfactant associated protein B Sftpb
1437029_at 1.06743 tachykinin receptor 3 Tacr3
1437030_at 1.06423 "phospholipase C, delta 4" Plcd4
1437031_at 1.05651 Acyl-CoA synthetase long-chain family member 6 Acsl6
1437032_x_at 0.99954 RNA binding motif protein 14 Rbm14
1437033_a_at 1.11551 S-phase kinase-associated protein 2 (p45) Skp2
1437034_x_at 0.95075 myristoylated alanine rich protein kinase C substrate Marcks
1437035_x_at 1.09672 ring finger protein 14 Rnf14
1437036_at 0.95573 Expressed sequence AL033314 AL033314
1437037_x_at 1.06037 expressed sequence AL033314 AL033314
1437038_x_at 1.04185 RIKEN cDNA 1700029I08 gene 1700029I08Rik
1437039_at 1.09782 "COP9 (constitutive photomorphogenic) homolog, subunit 2 (Arabidopsis thaliana)" Cops2
1437040_at 0.93090 ethanolamine kinase 2 Etnk2
1437041_at 0.74601 RIKEN cDNA 5730406M06 gene 5730406M06Rik
1437042_at 0.96342 aryl hydrocarbon receptor nuclear translocator Arnt
1437043_a_at 1.03418 RIKEN cDNA 1110012M11 gene 1110012M11Rik
1437044_a_at 1.11246 "glucosidase, beta, acid" Gba
1437045_at 1.46899 mitogen activated protein kinase 8 Mapk8
1437046_x_at 0.89183 RIKEN cDNA 4930504E06 gene 4930504E06Rik
1437047_at 0.97526 zinc finger protein 664 Zfp664
1437048_x_at 0.98041 syntaxin binding protein 2 Stxbp2
1437049_at 1.00895 Transcribed locus ---
1437050_s_at 0.89276 angel homolog 2 (Drosophila) Angel2
1437051_at 0.94786 "DNA fragmentation factor, beta subunit" Dffb
1437052_s_at 1.08266 "solute carrier family 2 (facilitated glucose transporter), member 3" Slc2a3
1437053_x_at 1.07054 apoptosis inhibitor 5 Api5
1437054_x_at 0.99530 protamine 1 Prm1
1437055_x_at 1.13170 Major facilitator superfamily domain containing 1 Mfsd1
1437056_x_at 1.02519 cysteine-rich secretory protein LCCL domain containing 2 Crisp1d2
1437057_at 0.68338 multiple EGF-like-domains 6 Megf6
1437058_at 0.71382 multiple EGF-like-domains 6 Megf6
1437059_at 1.06134 SRY-box containing gene 21 Sox21
1437060_at 1.02564 olfactomedin 4 Olfm4
1437061_at 1.03908 methyl-CpG binding domain protein 1 Mbd1
1437062_s_at 1.06230 phytanoyl-CoA hydroxylase interacting protein-like Phyh1pl
1437063_at 0.88680 feminization 1 homolog a (C. elegans) Fem1a
1437064_at 0.96313 androgen receptor Ar
1437065_at 0.95612 zinc finger and BTB domain containing 20 Zbtb20
1437066_at 1.07108 zinc finger and BTB domain containing 20 Zbtb20
1437067_at 0.95563 putative homeodomain transcription factor 2 Phtf2

1437068_at 1.18314 SoxLZ/Sox6 leucine zipper binding protein in testis Solt
1437069_at 1.09201 oxysterol binding protein-like 8 Osbp18
1437070_at 1.11048 CDC14 cell division cycle 14 homolog B (S. cerevisiae) Cdc14b
1437071_at 0.88995 "eukaryotic translation initiation factor 1A, Y-linked" Eif1ay
1437072_at 0.88371 Rho GTPase activating protein 25 Arhgap25
1437073_x_at 0.89792 expressed sequence AV025504 AV025504
1437074_at 1.76592 sorting nexin 5 Snx5
1437075_at 1.03081 FERM domain containing 3 Frmd3
1437076_at 1.03023 RIKEN cDNA A930017M01 gene A930017M01Rik
1437077_at 0.96964 expressed sequence AW495713 AW495713
1437078_at 0.92535 similar to vacuolar protein sorting 52 LOC631721
1437079_at 0.89786 "solute carrier family 18 (vesicular monoamine), member 2" Slc18a2
1437080_s_at 1.11923 "proteasome (prosome, macropain) 26S subunit, non-ATPase, 11 /// similar to
26S proteasome non-ATPase regulatory subunit 11 (26S proteasome regulatory subunit S9) (26S
proteasome regulatory subunit p44.5) /// similar to 26S proteasome non-ATPase regulatory subunit 11
(26S proteasome regulatory subunit S9) (26S proteasome regulatory subunit p44.5)" Psm11 ///
LOC627316 /// LOC671671
1437081_at 0.93877 Tissue inhibitor of metalloproteinase 2 Timp2
1437082_at 0.89212 A kinase (PRKA) anchor protein (yotiao) 9 Akap9
1437083_at 1.05020 RIKEN cDNA A830020B06 gene A830020B06Rik
1437084_at 1.00109 "peptidyl arginine deiminase, type VI" Padi6
1437085_at 0.96042 RIKEN cDNA D630039A03 gene D630039A03Rik
1437086_at 1.11655 achaete-scute complex homolog-like 1 (Drosophila) Ascl1
1437087_at 0.98501 --- ---
1437088_at 1.19297 SDA1 domain containing 1 Sdad1
1437089_at 0.94418 RIKEN cDNA 4833409A17 gene 4833409A17Rik
1437090_at 1.12187 coiled-coil domain containing 63 Ccdc63
1437091_at 1.12426 "amiloride-sensitive cation channel 4, pituitary" Accn4
1437092_at 1.16826 restin-like 2 Rsnl2
1437093_at 0.96283 "dynein, axonemal, intermediate chain 1" Dnaic1
1437094_x_at 1.02841 "dynein, axonemal, intermediate chain 1" Dnaic1
1437095_at 0.94178 tetraspanin 18 Tspan18
1437096_at 1.02189 tetratricopeptide repeat domain 29 Ttc29
1437097_at 0.84040 "zinc finger, BED domain containing 3" Zbed3
1437098_x_at 1.06955 Transcribed locus ---
1437099_x_at 1.04515 heterogeneous nuclear ribonucleoprotein F /// similar to heterogeneous nuclear
ribonucleoprotein F /// similar to heterogeneous nuclear ribonucleoprotein F /// similar to heterogeneous
nuclear ribonucleoprotein F Hnrpf /// LOC637008 /// LOC665008 /// LOC672616
1437100_x_at 1.05849 proviral integration site 3 Pim3
1437101_at 1.12645 large tumor suppressor 2 Lats2
1437102_at 0.94145 YTH domain family 1 Ythdf1
1437103_at 1.26803 insulin-like growth factor 2 mRNA binding protein 2 Igf2bp2
1437104_at 0.94608 ADP-ribosylation factor guanine nucleotide-exchange factor 1(brefeldin A-

inhibited) Arfgef1

1437105_at 0.94739 "jumonji, AT rich interactive domain 1A (Rbp2 like)" Jarid1a

1437106_at 0.88597 "jumonji, AT rich interactive domain 1A (Rbp2 like)" Jarid1a

1437107_at 1.00957 "RAB6B, member RAS oncogene family" Rab6b

1437108_at 0.97354 "LSM6 homolog, U6 small nuclear RNA associated (S. cerevisiae)" Lsm6

1437109_s_at 1.05855 "LSM6 homolog, U6 small nuclear RNA associated (S. cerevisiae)" Lsm6

1437110_at 0.99201 RIKEN cDNA 2810474O19 gene 2810474O19Rik

1437111_at 1.11769 zinc finger CCCH type containing 12C Zc3h12c

1437112_at 0.95496 phospholipase D1 Pld1

1437113_s_at 0.96987 phospholipase D1 Pld1

1437114_at 1.10723 RIKEN cDNA A930001N09 gene A930001N09Rik

1437115_at 1.09916 expressed sequence AA673488 AA673488

1437116_at 1.13906 LETM1 domain containing 1 Letmd1

1437117_at 0.89776 "centaurin, beta 1" Centb1

1437118_at 0.89909 ubiquitin specific peptidase 7 Usp7

1437119_at 1.23136 Endoplasmic reticulum (ER) to nucleus signalling 1 Ern1

1437120_at 1.02471 sorting nexin family member 30 Snx30

1437121_at 0.90323 DENN/MADD domain containing 1C Dennd1c

1437122_at 0.90548 B-cell leukemia/lymphoma 2 Bcl2

1437123_at 0.96251 multimerin 2 Mmrn2

1437124_at 1.02223 RIKEN cDNA A630052C17 gene A630052C17Rik

1437125_at 1.01959 calcium/calmodulin-dependent protein kinase II alpha Camk2a

1437126_at 1.45811 "Inner membrane protein, mitochondrial" Immt

1437127_at 1.08500 RIKEN cDNA A630033E08 gene A630033E08Rik

1437128_a_at 0.84902 RIKEN cDNA A630033E08 gene A630033E08Rik

1437129_at 1.18242 RIKEN cDNA E330018D03 gene E330018D03Rik

1437130_at 1.01365 similar to Kelch repeat and BTB (POZ) domain containing 7 (predicted)

LOC432879

1437131_x_at 0.98220 mitochondrial ribosomal protein L11 Mrpl11

1437132_x_at 0.84330 "neural precursor cell expressed, developmentally down-regulated gene 9" Nedd9

1437133_x_at 1.04449 "aldo-keto reductase family 1, member B3 (aldose reductase)" Akr1b3

1437134_at 0.98915 "Ubiquinol-cytochrome c reductase, complex III subunit VII" Uqcrcq

1437135_at 1.01165 F-box and WD-40 domain protein 10 Fbxw10

1437136_at 1.30868 RIKEN cDNA 5830436I19 gene 5830436I19Rik

1437137_at 1.07172 RIKEN cDNA 6430550H21 gene 6430550H21Rik

1437138_at 0.98663 forkhead box M1 Foxm1

1437139_at 1.01230 "glycine receptor, alpha 1 subunit" Glra1

1437140_at 1.04311 RIKEN cDNA 4930412F15 gene 4930412F15Rik

1437141_x_at 0.92569 dymeclin Dym

1437142_a_at 0.94586 "phosphatidylinositol glycan anchor biosynthesis, class O" Pigo

1437143_a_at 0.90724 thioredoxin domain containing 1 Txndc1

1437144_x_at 1.11729 "proteasome (prosome, macropain) subunit, alpha type 6" Psma6

1437145_s_at 1.01306 RIKEN cDNA 2310002J15 gene 2310002J15Rik
 1437146_x_at 1.01083 coronin 7 Coro7
 1437147_at 0.99190 "gamma-aminobutyric acid (GABA-A) receptor, subunit gamma 2" Gabrg2
 1437148_at 0.97736 "actin related protein 2/3 complex, subunit 2" Arpc2
 1437149_at 1.09358 "solute carrier family 6 (neurotransmitter transporter, taurine), member 6" Slc6a6
 1437150_at 1.18427 RIKEN cDNA 1700012H17 gene 1700012H17Rik
 1437151_at 0.96781 ubiquitin specific peptidase 22 Usp22
 1437152_at 0.64636 ring finger and KH domain containing 3 Rkhd3
 1437153_at 1.16367 similar to Oogenesis 2 /// oogenesis 2 MGC91195 /// Oog2
 1437154_at 1.12094 centrosomal protein 170 Cep170
 1437155_a_at 0.87858 WW domain containing transcription regulator 1 Wwtr1
 1437156_at 0.97507 EF hand calcium binding protein 1 Efcfbp1
 1437157_at 0.96402 "ATPase, Na⁺/K⁺ transporting, beta 1 polypeptide" Atp1b1
 1437158_at 1.13267 Nipped-B homolog (Drosophila) Nipbl
 1437159_at 1.01929 --- ---
 1437160_at 1.02692 neuroligin 1 Nlgn1
 1437161_x_at 0.92389 RNA binding protein gene with multiple splicing Rbpms
 1437162_at 2.42987 GPI-anchored membrane protein 1 Gpiap1
 1437163_x_at 0.96902 "general transcription factor II H, polypeptide 4" Gtf2h4
 1437164_x_at 0.86153 "ATP synthase, H⁺ transporting, mitochondrial F1 complex, O subunit /// similar to ATP synthase, H⁺ transporting, mitochondrial F1 complex, O subunit" Atp5o /// LOC432676
 1437165_a_at 0.92036 procollagen C-endopeptidase enhancer protein Pcolce
 1437166_at 0.85616 Terf1 (TRF1)-interacting nuclear factor 2 Tinf2
 1437167_at 0.99361 tropomodulin 2 Tmod2
 1437168_at 1.03896 serine-arginine repressor protein Srrp
 1437169_at 1.07488 PDZ and LIM domain 1 (elfin) /// similar to PDZ and LIM domain protein 1 (Elfin) (LIM domain protein CLP-36) (C-terminal LIM domain protein 1) Pdlim1 /// LOC545743
 1437170_x_at 0.99843 PDZ and LIM domain 1 (elfin) /// similar to PDZ and LIM domain protein 1 (Elfin) (LIM domain protein CLP-36) (C-terminal LIM domain protein 1) Pdlim1 /// LOC545743
 1437171_x_at 0.86202 gelsolin Gsn
 1437172_x_at 0.85460 "hydroxyacyl-Coenzyme A dehydrogenase/3-ketoacyl-Coenzyme A thiolase/enoyl-Coenzyme A hydratase (trifunctional protein), beta subunit" Hadhb
 1437173_at 0.95926 "endothelial differentiation, sphingolipid G-protein-coupled receptor, 3" Edg3
 1437174_at 0.99437 transcription factor Dp 2 Tfdp2
 1437175_at 0.98883 PDLIM1 interacting kinase 1 like Pdik11
 1437176_at 1.46318 expressed sequence AI451557 AI451557
 1437177_at 1.25123 RIKEN cDNA D330037H05 gene /// similar to La-related protein 4 (La ribonucleoprotein domain family member 4) /// similar to La-related protein 4 (La ribonucleoprotein domain family member 4) /// similar to La-related protein 4 (La ribonucleoprotein domain family member 4) D330037H05Rik /// LOC669298 /// LOC673040 /// LOC677019
 1437178_at 0.91903 hypothetical LOC631105 LOC631105

1437179_at 1.16076 Rap1 interacting factor 1 homolog (yeast) /// similar to Telomere-associated protein RIF1 (Rap1-interacting factor 1 homolog) (mRif1) Rif1 /// LOC671598
1437180_at 1.43111 RIKEN cDNA 6530403A03 gene 6530403A03Rik
1437181_at 0.57541 pellino 2 Peli2
1437182_at 1.42806 death inducer-obliterator 1 Dido1
1437183_at 1.10632 leucine rich repeat containing 4B Lrrc4b
1437184_at 1.10140 GUF1 GTPase homolog (S. cerevisiae) Guf1
1437185_s_at 1.01276 "thymosin, beta 10" Tmsb10
1437186_at 1.16833 cDNA sequence BC055324 BC055324
1437187_at 0.97856 E2F transcription factor 7 E2f7
1437188_at 0.86414 "gamma-aminobutyric acid (GABA-B) receptor, 1" Gabbr1
1437189_x_at 1.08156 dendritic cell protein GA17 Ga17
1437190_at 1.10247 serine/threonine/tyrosine kinase 1 Styk1
1437191_at 1.01173 cDNA sequence BC050789 BC050789
1437192_x_at 0.94161 voltage-dependent anion channel 1 Vdac1
1437193_s_at 1.02703 small nuclear ribonucleoprotein B Snrpb
1437194_x_at 0.97886 RIKEN cDNA 1200011O22 gene 1200011O22Rik
1437195_x_at 0.87296 mitogen activated protein kinase 10 Mapk10
1437196_x_at 0.99642 ribosomal protein S16 Rps16
1437197_at 0.83179 sorbin and SH3 domain containing 2 Sorbs2
1437198_at 0.92281 "ligase III, DNA, ATP-dependent" Lig3
1437199_at 0.75012 dual specificity phosphatase 5 Dusp5
1437200_at 0.95796 FCH domain only 2 Fcho2
1437201_at 1.06745 leucine rich repeat containing 4C Lrrc4c
1437202_at 1.06517 "myb-like, SWIRM and MPN domains 1" Mysm1
1437203_at 1.10513 Casitas B-lineage lymphoma-like 1 Cbl1
1437204_a_at 1.31916 transcription factor 25 (basic helix-loop-helix) Tcf25
1437205_at 1.27377 transcription factor 25 (basic helix-loop-helix) Tcf25
1437206_at 1.09987 SET domain containing 5 Setd5
1437207_at 0.95554 ring finger protein 170 Rnf170
1437208_at 0.93713 septin 10 10-Sep
1437209_at 1.03643 centrosomal protein 76 Cep76
1437210_a_at 0.97117 bromodomain containing 2 Brd2
1437211_x_at 0.97916 "ELOVL family member 5, elongation of long chain fatty acids (yeast)" Elov15
1437212_at 1.06575 RIKEN cDNA B230312I18 gene B230312I18Rik
1437213_at 0.95662 nudix (nucleoside diphosphate linked moiety X)-type motif 21 Nudt21
1437214_at 0.99344 leucine rich repeat transmembrane neuronal 4 Lrrtm4
1437215_at 1.14078 similar to MTH2 protein /// similar to MTH2 protein LOC667026 ///
LOC671874
1437216_at 1.08626 RIKEN cDNA A430106J12 gene A430106J12Rik
1437217_at 0.92973 ankyrin repeat domain 6 Ankrd6
1437218_at 1.74331 Fibronectin 1 Fn1

1437219_at 3.25841 Insulin-like growth factor I receptor Igf1r
1437220_x_at 1.10068 "DnaJ (Hsp40) homolog, subfamily A, member 1" Dnaja1
1437221_at 1.14808 ribonucleotide reductase M2 B (TP53 inducible) Rrm2b
1437222_x_at 0.95994 ribonucleotide reductase M2 B (TP53 inducible) Rrm2b
1437223_s_at 0.88261 X-box binding protein 1 Xbp1
1437224_at 0.72423 reticulon 4 Rtn4
1437225_x_at 0.89846 "guanine nucleotide binding protein, alpha inhibiting 3" Gnai3
1437226_x_at 0.82832 MARCKS-like 1 Marcksl1
1437227_at 0.82313 RIKEN cDNA 2900073H19 gene 2900073H19Rik
1437228_at 1.19406 cDNA sequence BC048651 BC048651
1437229_at 0.94481 "EF hand domain family, member B" Efhb
1437230_at 1.16118 "Adult male corpus striatum cDNA, RIKEN full-length enriched library, clone:7630403E16 product:unclassifiable, full insert sequence" ---
1437231_at 0.92949 "SLIT and NTRK-like family, member 6" Slitrk6
1437232_at 0.78062 bactericidal/permeability-increasing protein-like 2 Bpil2
1437233_x_at 0.95430 RIKEN cDNA 1110049F12 gene 1110049F12Rik
1437234_x_at 0.96999 protein arginine N-methyltransferase 2 Prmt2
1437235_x_at 1.10042 LIM domain containing preferred translocation partner in lipoma Lpp
1437236_a_at 0.76074 zinc finger protein 110 Zfp110
1437237_x_at 0.73556 zinc finger protein 110 Zfp110
1437238_x_at 1.05476 NMD3 homolog (S. cerevisiae) Nmd3
1437239_x_at 1.01025 polyhomeotic-like 2 (Drosophila) Phc2
1437240_at 1.45620 phosphoglucomutase 2 Pgm2
1437241_at 0.67811 TGFB inducible early growth response 3 Tieg3
1437242_at 1.04458 "tubulin tyrosine ligase-like family, member 12" Ttl12
1437243_at 1.05845 "D-2-hydroxyglutarate dehydrogenase /// similar to CG3835-PA, isoform A" D2hgdh /// LOC673662
1437244_at 1.32340 growth arrest-specific 2 like 3 Gas2l3
1437245_at 1.03057 transmembrane protein 154 Tmem154
1437246_x_at 1.09623 ribosomal protein S6 /// similar to 40S ribosomal protein S6 /// similar to 40S ribosomal protein S6 /// similar to 40S ribosomal protein S6 /// similar to 40S ribosomal protein S6 /// similar to 40S ribosomal protein S6 /// similar to 40S ribosomal protein S6 /// similar to 40S ribosomal protein S6 /// similar to 40S ribosomal protein S6 /// similar to 40S ribosomal protein S6 /// similar to 40S ribosomal protein S6 /// similar to 40S ribosomal protein S6 /// similar to 40S ribosomal protein S6 /// similar to 40S ribosomal protein S6 /// similar to 40S ribosomal protein S6 /// similar to 40S ribosomal protein S6 /// similar to 40S ribosomal protein S6 /// similar to 40S ribosomal protein S6 /// similar to 40S ribosomal protein S6 Rps6 /// LOC214738 /// LOC384383 /// LOC545640 /// LOC626259 /// LOC629575 /// LOC638497 /// LOC639593 /// LOC639750 /// LOC666577 /// LOC667739 /// LOC667922 /// LOC671306 /// LOC671432 /// LOC672620 /// LOC675985
1437247_at 1.16687 fos-like antigen 2 /// similar to fos-like antigen 2 Fosl2 /// LOC634417
1437248_at 0.76921 RIKEN cDNA 2700049A03 gene 2700049A03Rik
1437249_at 1.03318 src family associated phosphoprotein 1 Scap1
1437250_at 0.82672 melanoregulin Mreg
1437251_at 0.99298 cell division cycle associated 2 Cdca2

1437252_at	0.76456	opposite strand transcription unit to Stag3	Gats
1437253_at	1.01771	RIKEN cDNA A630054L15 gene	A630054L15Rik
1437254_at	0.90243	centrosomal protein 76	Cep76
1437255_at	1.10553	Zinc finger and BTB domain containing 7a	Zbtb7a
1437256_at	0.98634	---	---
1437257_at	1.02182	WD repeat domain 47	Wdr47
1437258_at	0.93296	small proline-rich protein 2A	Sprr2a
1437259_at	0.97177	---	---
1437260_at	1.05971	multimerin 1	Mmrn1
1437261_at	1.29092	RIKEN cDNA 2900024O10 gene	2900024O10Rik
1437262_x_at	0.86486	breast carcinoma amplified sequence 2	Bcas2
1437263_at	1.18902	RIKEN cDNA A730089K16 gene	A730089K16Rik
1437264_at	0.82767	cDNA sequence BC051142	BC051142
1437265_at	0.86689	RIKEN cDNA 5330438D12 gene	5330438D12Rik
1437266_at	1.00340	thioredoxin domain containing 2 (spermatzoa)	Txndc2
1437267_x_at	0.99906	Heterogeneous nuclear ribonucleoprotein H1	Hnrph1
1437268_at	0.95450	LanC lantibiotic synthetase component C-like 3 (bacterial)	Lancl3
1437269_at	1.04319	expressed sequence AW124722	AW124722
1437270_a_at	0.90367	cardiotrophin-like cytokine factor 1	Clcf1
1437271_at	0.89115	cardiotrophin-like cytokine factor 1	Clcf1
1437272_at	1.02548	RIKEN cDNA 4932443J21 gene	4932443J21Rik
1437273_at	0.96129	HtrA serine peptidase 4	Htra4
1437274_at	0.97478	coatomer protein complex subunit alpha	Copa
1437275_at	0.99152	"Catenin (cadherin associated protein), alpha 1"	Ctnna1
1437276_at	1.26212	Transcribed locus	---
1437277_x_at	0.90403	"transglutaminase 2, C polypeptide"	Tgm2
1437278_a_at	0.89689	ubiquitin-like 1 (sentrin) activating enzyme E1B	Uble1b
1437279_x_at	0.85422	syndecan 1	Sdc1
1437280_s_at	1.06463	Serpine1 mRNA binding protein 1	Serbp1
1437281_x_at	0.99288	XPA binding protein 2	Xab2
1437282_at	0.98048	Dedicator of cyto-kinesis 2	Dock2
1437283_at	1.08197	"transportin 2 (importin 3, karyopherin beta 2b)"	Tnpo2
1437284_at	1.02130	frizzled homolog 1 (Drosophila)	Fzd1
1437285_at	1.25894	---	---
1437286_x_at	1.03355	---	---
1437287_at	1.06161	RIKEN cDNA 1110020G09 gene	1110020G09Rik
1437288_at	0.93053	inositol monophosphatase domain containing 1	Impad1
1437289_at	1.01792	inositol monophosphatase domain containing 1	Impad1
1437290_at	0.91123	inositol monophosphatase domain containing 1	Impad1
1437291_at	0.89006	RIKEN cDNA 2700081O15 gene	2700081O15Rik
1437292_at	1.06300	RIKEN cDNA A330019N05 gene	A330019N05Rik
1437293_x_at	0.95290	uronyl-2-sulfotransferase	Ust
1437294_at	1.03227	exosome component 10	Exosc10

1437295_at 1.04903 protein kinase N2 Pkn2
1437296_at 1.15538 protein kinase N2 Pkn2
1437297_at 0.98219 chromodomain helicase DNA binding protein 8 Chd8
1437298_at 0.94625 RIKEN cDNA 5730509C05 gene 5730509C05Rik
1437299_at 1.06234 "NACHT, leucine rich repeat and PYD containing 4F" Nalp4f
1437300_at 1.05256 myocyte enhancer factor 2D Mef2d
1437301_a_at 1.28570 "dishevelled, dsh homolog 1 (Drosophila)" Dvl1
1437302_at 0.95267 "adrenergic receptor, beta 2" Adrb2
1437303_at 1.10386 interleukin 6 signal transducer Il6st
1437304_at 0.84445 Casitas B-lineage lymphoma b Cblb
1437305_at 1.14636 RIKEN cDNA 6430601A21 gene 6430601A21Rik
1437306_at 0.95614 RIKEN cDNA C130092O11 gene C130092O11Rik
1437307_at 0.85720 SUMO/sentrin specific peptidase 8 Senp8
1437308_s_at 0.94514 coagulation factor II (thrombin) receptor F2r
1437309_a_at 0.98167 replication protein A1 Rpa1
1437310_at 0.82057 Bardet-Biedl syndrome 1 homolog (human) Bbs1
1437311_at 1.03566 RIKEN cDNA A930034L06 gene A930034L06Rik
1437312_at 0.95952 "bone morphogenetic protein receptor, type 1B" Bmpr1b
1437313_x_at 1.01580 high mobility group box 2 /// high mobility group box 2 /// high mobility group box 2 Hmgb2 /// LOC678110 /// LOC678341
1437314_a_at 1.04233 TRM1 tRNA methyltransferase 1 homolog (S. cerevisiae) Trmt1
1437315_at 0.90119 G protein-coupled receptor 172B Gpr172b
1437316_at 1.06785 F-box and WD-40 domain protein 19 Fbxw19
1437317_at 1.01616 ubiquitin-activating enzyme E1-like Ube1l
1437318_at 1.00970 p21 (CDKN1A)-activated kinase 3 Pak3
1437319_at 0.91558 unc-13 homolog C (C. elegans) Unc13c
1437320_s_at 1.20567 "xeroderma pigmentosum, complementation group A" Xpa
1437321_at 0.90475 --- ---
1437322_at 1.70591 RNA binding motif protein 4 Rbm4
1437323_a_at 0.98696 islet amyloid polypeptide Iapp
1437324_x_at 0.59765 fibromodulin Fmod
1437325_x_at 0.98405 "aldehyde dehydrogenase 18 family, member A1" Aldh18a1
1437326_x_at 1.05306 "elastase 3, pancreatic /// similar to elastase 3B, pancreatic" Ela3 /// LOC638418
1437327_x_at 1.00562 RIKEN cDNA 2310057D15 gene 2310057D15Rik
1437328_x_at 0.97405 cleavage and polyadenylation specificity factor 3 Cpsf3
1437329_at 1.01371 "protein tyrosine phosphatase-like (proline instead of catalytic arginine), member b" Ptplb
1437330_at 0.80765 leucine-rich repeat kinase 1 Lrrk1
1437331_a_at 1.00518 ADP-ribosylation factor 3 Arf3
1437332_at 0.97484 "Adult male corpora quadrigemina cDNA, RIKEN full-length enriched library, clone:B230334A10 product:unclassifiable, full insert sequence" ---
1437333_x_at 0.96581 "aldehyde dehydrogenase 18 family, member A1" Aldh18a1

1437334_x_at 0.84681 poly(A)-specific ribonuclease (deadenylation nuclease) Parn
1437335_x_at 0.95816 "polymerase (DNA-directed), delta interacting protein 3" Poldip3
1437336_x_at 1.10474 RIKEN cDNA 1110002E23 gene 1110002E23Rik
1437337_x_at 0.95460 fucokinase Fuk
1437338_x_at 0.90147 Elongation protein 3 homolog (*S. cerevisiae*) Elp3
1437339_s_at 0.94425 proprotein convertase subtilisin/kexin type 5 Pcsk5
1437340_x_at 1.06036 gastroke 1 Gkn1
1437341_x_at 0.84547 cyclic nucleotide phosphodiesterase 1 Cnp1
1437342_x_at 1.05060 pituitary tumor-transforming 1 interacting protein Pttg1ip
1437343_x_at 1.12987 "ATPase family, AAA domain containing 3A" Atad3a
1437344_x_at 1.00258 --- ---
1437345_a_at 0.96225 Bernardinelli-Seip congenital lipodystrophy 2 homolog (human) Bscl2
1437346_x_at 0.90858 Immunoglobulin heavy chain (J558 family) Igh-VJ558
1437347_at 1.00754 endothelin receptor type B Ednrb
1437348_at 1.07193 F-box protein 28 Fbxo28
1437349_at 1.01879 cytoskeleton associated protein 5 Ckap5
1437350_at 0.92357 candidate tumor suppressor OVCA2 RP23-143A14.5
1437351_at 0.57307 CXXC finger 4 Cxxc4
1437352_at 0.93771 "acyl-Coenzyme A oxidase 3, pristanoyl" Acox3
1437353_at 0.91241 SET domain containing 1B Setd1b
1437354_at 0.91688 RIKEN cDNA C230091D08 gene C230091D08Rik
1437355_at 0.95828 "zinc finger, CCHC domain containing 5" Zcchc5
1437356_at 1.00135 Epstein-Barr virus induced gene 2 Ebi2
1437357_at 1.09240 YTH domain containing 2 Ythdc2
1437358_at 0.83487 WD repeat and FYVE domain containing 1 Wdfy1
1437359_at 1.52567 Ribonucleic acid binding protein S1 Rnps1
1437360_at 0.84958 protocadherin 19 Pcdh19
1437361_at 0.99768 RIKEN cDNA A430060F13 gene A430060F13Rik
1437362_at 0.98920 RIKEN cDNA A930028L21 gene A930028L21Rik
1437363_at 1.17174 homer homolog 1 (*Drosophila*) Homer1
1437364_at 1.00305 "coenzyme Q3 homolog, methyltransferase (yeast)" Coq3
1437365_at 1.04645 NOL1/NOP2/Sun domain family 3 Nsun3
1437366_at 0.98909 expressed sequence AI608492 AI608492
1437367_at 1.03979 "Histocompatibility 2, K1, K region" H2-K1
1437368_at 1.02811 "general transcription factor II H, polypeptide 4 /// similar to general
transcription factor II H, polypeptide 4" Gtf2h4 /// LOC640308
1437369_at 0.91707 --- ---
1437370_at 0.91033 shugoshin-like 2 (*S. pombe*) Sgol2
1437371_at 0.74868 RIKEN cDNA 9930021J17 gene 9930021J17Rik
1437372_at 1.43939 cleavage and polyadenylation specific factor 6 Cpsf6
1437373_at 1.08401 interferon stimulated exonuclease gene 20-like 1 Isg20l1
1437374_at 0.97295 RIKEN cDNA 6430526N21 gene 6430526N21Rik
1437375_at 0.93219 "0 day neonate thymus cDNA, RIKEN full-length enriched library, clone:

A430010E21 product:unclassifiable, full insert sequence" ---

1437376_at 1.04491 a disintegrin and metallopeptidase domain 32 Adam32

1437377_a_at 0.94629 polymerase (RNA) mitochondrial (DNA directed) Polrmt

1437378_x_at 0.91995 "scavenger receptor class B, member 1" Scarb1

1437379_x_at 0.92258 TNF receptor-associated protein 1 Trap1

1437380_x_at 1.28477 phosphogluconate dehydrogenase Pgd

1437381_x_at 1.02182 G protein-coupled receptor 172B Gpr172b

1437382_at 0.83623 Activin receptor IIA Acvr2a

1437383_at 0.95626 Cyclin L1 Ccnl1

1437384_at 0.90206 RIKEN cDNA 1700015E13 gene 1700015E13Rik

1437385_at 0.97369 collagen and calcium binding EGF domains 1 Ccbe1

1437386_at 1.03271 leucine rich repeat neuronal 6A Lrn6a

1437387_at 0.94352 sushi domain containing 5 Susd5

1437388_at 0.98579 fucosyltransferase 10 Fut10

1437389_x_at 0.97144 --- ---

1437390_x_at 1.01576 syntaxin 1A (brain) Stx1a

1437391_x_at 0.98071 mitochondrial ribosomal protein L44 Mrpl44

1437392_at 0.97448 hypothetical LOC433485 LOC433485

1437393_at 0.81874 expressed sequence AI875142 AI875142

1437394_at 0.96781 "centaurin, gamma 2" Centg2

1437395_at 0.99346 "zinc finger, CCHC domain containing 11" Zcchc11

1437396_at 0.87681 cAMP responsive element binding protein 3-like 2 Creb3l2

1437397_at 0.90938 prolactin receptor Prlr

1437398_a_at 1.04549 "aldehyde dehydrogenase 9, subfamily A1" Aldh9a1

1437399_at 0.81567 claudin domain containing 1 Clnd1

1437400_at 0.98379 "Adult male corpus striatum cDNA, RIKEN full-length enriched library, clone: C030044L04 product:unclassifiable, full insert sequence" ---

1437401_at 0.78029 --- ---

1437402_x_at 0.98353 transmembrane protein 41B Tmem41b

1437403_at 1.08433 RIKEN cDNA E130306M17 gene E130306M17Rik

1437404_at 0.82607 microtubule associated serine/threonine kinase family member 4 Mast4

1437405_a_at 0.55145 insulin-like growth factor binding protein 4 Igfbp4

1437406_x_at 0.61242 insulin-like growth factor binding protein 4 Igfbp4

1437407_at 1.03408 RIKEN cDNA 9930039A11 gene 9930039A11Rik

1437408_at 1.00798 G protein-coupled receptor 126 Gpr126

1437409_s_at 0.82366 G protein-coupled receptor 126 Gpr126

1437410_at 0.99661 "aldehyde dehydrogenase 2, mitochondrial" Aldh2

1437411_at 1.11588 cDNA sequence BC010584 BC010584

1437412_at 1.00731 "peroxiredoxin 6, related sequence 1" Prdx6-rs1

1437413_x_at 0.90559 ribosomal protein S29 /// similar to 40S ribosomal protein S29 /// similar to 40S ribosomal protein S29 Rps29 /// LOC629957 /// LOC637599

1437414_at 0.99808 zinc finger protein 217 Zfp217

1437415_at 1.04747 RIKEN cDNA 4933427D06 gene 4933427D06Rik

1437416_at 0.94798 cDNA sequence BC034902 BC034902
 1437417_s_at 0.76173 glypican 6 Gpc6
 1437418_at 1.05625 --- ---
 1437419_at 0.94968 BMP2 inducible kinase Bmp2k
 1437420_at 0.92756 "bromodomain adjacent to zinc finger domain, 1B" Baz1b
 1437421_at 0.97992 RIKEN cDNA 6330509M05 gene 6330509M05Rik
 1437422_at 0.95710 "sema domain, seven thrombospondin repeats (type 1 and type 1-like),
 transmembrane domain (TM) and short cytoplasmic domain, (semaphorin) 5A" Sema5a
 1437423_a_at 0.99083 steroid receptor RNA activator 1 Sra1
 1437424_at 1.11224 "synapse defective 1, Rho GTPase, homolog 2 (C. elegans) /// similar to synapse
 defective 1, Rho GTPase, homolog 1" Syde2 /// LOC639654
 1437425_at 0.90860 ganglioside-induced differentiation-associated-protein 1 Gdap1
 1437426_at 1.67433 WW domain containing adaptor with coiled-coil Wac
 1437427_at 1.06982 RIKEN cDNA 4933405O20 gene 4933405O20Rik
 1437428_x_at 1.06447 "eukaryotic translation initiation factor 2B, subunit 2 beta" Eif2b2
 1437429_at 0.84383 RIKEN cDNA 5930437A14 gene 5930437A14Rik
 1437430_at 0.94797 "solute carrier family 45, member 2" Slc45a2
 1437431_at 1.62510 --- ---
 1437432_a_at 1.04412 tripartite motif protein 12 Trim12
 1437433_at 1.05469 "UDP-Gal:betaGlcNAc beta 1,3-galactosyltransferase, polypeptide 2"
 B3galt2
 1437434_a_at 0.86080 G protein-coupled receptor 177 Gpr177
 1437435_at 0.87812 RIKEN cDNA 1700061G19 gene /// similar to gonadotropin-regulated long
 chain acyl-CoA synthetase 1700061G19Rik /// LOC669384
 1437436_s_at 0.97263 G protein-coupled receptor kinase 6 Gprk6
 1437437_x_at 0.89227 aspartyl aminopeptidase Dnpep
 1437438_x_at 1.16711 pancreatic lipase-related protein 2 Pnliprp2
 1437439_at 0.96551 RIKEN cDNA 2410089E03 gene 2410089E03Rik
 1437440_at 0.91249 Transcribed locus ---
 1437441_at 1.00269 expressed sequence AA388235 AA388235
 1437442_at 0.73638 --- ---
 1437443_at 1.18226 "PREDICTED: Mus musculus hypothetical protein LOC620382 (LOC620382),
 mRNA" ---
 1437444_at 1.02702 "protocadherin gamma subfamily C, 3" Pcdhgc3
 1437445_at 0.90854 "transient receptor potential cation channel, subfamily M, member 1" Trpm1
 1437446_at 1.17380 "RAB5B, member RAS oncogene family" Rab5b
 1437447_s_at 0.99843 "excision repair cross-complementing rodent repair deficiency,
 complementation group 1" Ercc1
 1437448_s_at 0.98513 "catenin (cadherin associated protein), delta 1" Ctnnd1
 1437449_at 1.15279 radical S-adenosyl methionine domain containing 1 Rsad1
 1437450_x_at 0.99943 RIKEN cDNA 2700060E02 gene 2700060E02Rik
 1437451_at 1.37640 RIKEN cDNA 1110006O17 gene 1110006O17Rik
 1437452_x_at 1.09827 voltage-dependent anion channel 1 Vdac1

1437453_s_at 1.33597 proprotein convertase subtilisin/kexin type 9 Pcsk9
 1437454_a_at 0.99955 thioredoxin domain containing 14 /// similar to thioredoxin-related
 transmembrane protein 2 Txndc14 /// LOC433144
 1437455_a_at 0.94191 "B-cell translocation gene 1, anti-proliferative" Btg1
 1437456_x_at 0.96988 YTH domain family 1 Ythdf1
 1437457_a_at 1.01621 myotrophin Mtpn
 1437458_x_at 0.84254 clusterin Clu
 1437459_x_at 0.94394 "solute carrier family 25 (mitochondrial deoxynucleotide carrier), member
 19" Slc25a19
 1437460_x_at 1.08390 Ras and Rab interactor 1 Rin1
 1437461_s_at 0.77267 "RNA-binding region (RNP1, RRM) containing 3" Rnpc3
 1437462_x_at 0.72836 matrix metalloproteinase 15 Mmp15
 1437463_x_at 0.98794 "transforming growth factor, beta induced" Tgfb1
 1437464_at 0.91457 spermatogenesis associated 7 Spata7
 1437465_a_at 0.94896 "prolyl 4-hydroxylase, beta polypeptide" P4hb
 1437466_at 0.69113 activated leukocyte cell adhesion molecule Alcam
 1437467_at 0.84916 activated leukocyte cell adhesion molecule Alcam
 1437468_x_at 0.94706 F-box and WD-40 domain protein 11 Fbxw11
 1437469_at 0.97031 zinc finger protein 750 Zfp750
 1437470_at 0.91799 Pbx/knotted 1 homeobox Pknox1
 1437471_at 0.96783 leucine rich repeat containing 45 Lrrc45
 1437472_at 0.79846 unc-13 homolog A (C. elegans) Unc13a
 1437473_at 0.95088 avian musculoaponeurotic fibrosarcoma (v-maf) AS42 oncogene homolog
 Maf
 1437474_at 1.00169 GATA zinc finger domain containing 2B Gatad2b
 1437475_at 0.92782 cDNA sequence BC028454 BC028454
 1437476_at 0.82444 ribonucleotide reductase M2 B (TP53 inducible) Rrm2b
 1437477_at 1.04090 leucine rich repeat (in FLII) interacting protein 1 Lrrfip1
 1437478_s_at 1.02214 EF hand domain containing 2 Efhd2
 1437479_x_at 0.89997 T-box 3 Tbx3
 1437480_at 1.05688 --- ---
 1437481_at 1.09352 hypothetical LOC623451 LOC623451
 1437482_at 1.03095 steroid 5 alpha-reductase 2-like 2 Srd5a2l2
 1437483_at 1.00323 zinc finger protein 513 Zfp513
 1437484_at 0.99535 zinc finger and BTB domain containing 5 Zbtb5
 1437485_at 1.30650 nitric oxide synthase 1 (neuronal) adaptor protein Nos1ap
 1437486_at 1.06203 "G protein-coupled receptor, family C, group 5, member A" Gprc5a
 1437487_at 0.99627 MYST histone acetyltransferase (monocytic leukemia) 3 Myst3
 1437488_at 1.02687 "solute carrier family 9 (sodium/hydrogen exchanger), isoform 9" Slc9a9
 1437489_x_at 1.01694 "succinate dehydrogenase complex, subunit D, integral membrane protein"
 Sdhb
 1437490_x_at 1.02113 UDP-N-acetylglucosamine pyrophosphorylase 1 /// similar to UDP-N-
 acetylhexosamine pyrophosphorylase Uap1 /// LOC640502

1437491_at	1.26660 bicaudal D homolog 2 (Drosophila)	Bicd2
1437492_at	0.89884 mohawk	Mkx
1437493_at	1.88914 RIKEN cDNA 5230400M03 gene	5230400M03Rik
1437494_at	1.04000 mitogen-activated protein kinase-activated protein kinase 3	Mapkapk3
1437495_at	1.15448 "membrane-bound transcription factor peptidase, site 2"	Mbtps2
1437496_at	1.01581 RIKEN cDNA 9330129D05 gene	9330129D05Rik
1437497_a_at	1.08614 "heat shock protein 90kDa alpha (cytosolic), class A member 1"	Hsp90aa1
1437498_at	0.97598 ---	---
1437499_at	1.48349 ankyrin repeat domain 39	Ankrd39
1437500_at	1.15486 "Solute carrier family 39 (zinc transporter), member 12"	Slc39a12
1437501_at	0.89384 cDNA sequence AF529169	AF529169
1437502_x_at	0.86776 CD24a antigen	Cd24a
1437503_a_at	0.93703 scotin gene	Scotin
1437504_at	0.85235 Src-like-adaptor 2	Sla2
1437505_at	1.80642 Triple functional domain (PTPRF interacting)	Trio
1437506_at	0.68199 ---	---
1437507_at	0.93320 GATA zinc finger domain containing 2B	Gatad2b
1437508_at	0.85010 trans-acting transcription factor 4	Sp4
1437509_x_at	0.92143 protein kinase D2	Prkd2
1437510_x_at	1.02909 ribosomal protein S17	Rps17
1437511_x_at	0.94089 chloride channel CLIC-like 1	Clcc1
1437512_x_at	0.94010 EBNA1 binding protein 2	Ebna1bp2
1437513_a_at	0.96221 serine incorporator 1	Serinc1
1437514_at	1.08324 RIKEN cDNA B430306N03 gene	B430306N03Rik
1437515_at	0.97616 ---	---
1437516_at	1.12373 Transcribed locus	---
1437517_x_at	0.94907 "serine (or cysteine) peptidase inhibitor, clade B (ovalbumin), member 3A"	Serpib3a
1437518_at	1.00761 "10 days neonate cerebellum cDNA, RIKEN full-length enriched library, clone:6530404G16 product:hypothetical protein, full insert sequence"	---
1437519_x_at	1.09476 hydroxyacyl glutathione hydrolase	Hagh
1437520_a_at	1.00291 nucleoporin 85	Nup85
1437521_s_at	1.00473 RIKEN cDNA E230022H04 gene	E230022H04Rik
1437522_x_at	0.93041 growth hormone	Gh
1437523_s_at	1.02305 "sarcoglycan, gamma (dystrophin-associated glycoprotein)"	Sgcg
1437524_x_at	0.94683 coronin 7	Coro7
1437525_a_at	1.01657 polymerase (RNA) III (DNA directed) polypeptide A	Polr3a
1437526_x_at	0.95805 heterogeneous nuclear ribonucleoprotein R /// similar to heterogeneous nuclear ribonucleoprotein R	Hnrpr /// LOC669321
1437527_x_at	0.86373 myeloid cell leukemia sequence 1	Mcl1
1437528_x_at	0.89315 RIKEN cDNA A730017C20 gene	A730017C20Rik
1437529_at	1.00263 "MAS-related GPR, member B1"	Mrgprb1
1437530_at	0.96859 Programmed cell death 10	Pdcd10

1437531_at 0.99847 "transient receptor potential cation channel, subfamily M, member 1" Trpm1
1437532_at 0.93077 RIKEN cDNA 2810055G22 gene /// similar to ubiquitin conjugating enzyme 7
interacting protein 1 2810055G22Rik /// LOC673825
1437533_at 1.08820 baculoviral IAP repeat-containing 4 Birc4
1437534_at 1.08817 "PREDICTED: Mus musculus similar to Rap guanine nucleotide exchange
factor 2 (PDZ domain containing guanine nucleotide exchange factor 1) (PDZ-GEF1) (LOC620996),
mRNA" ---
1437535_at 0.98157 "protein phosphatase 3, regulatory subunit B, alpha isoform (calcineurin B, type
II)" Ppp3r2
1437536_at 0.86457 fukutin related protein Fkrp
1437537_at 0.92077 caspase 9 Casp9
1437538_at 1.18729 RIKEN cDNA 2610002F03 gene 2610002F03Rik
1437539_at 0.95942 "protein kinase, AMP-activated, alpha 1 catalytic subunit" Prkaa1
1437540_at 0.92945 --- ---
1437541_at 1.00250 --- ---
1437542_at 0.92413 RIKEN cDNA A730095J18 gene A730095J18Rik
1437543_at 1.72683 far upstream element (FUSE) binding protein 1 Fubp1
1437544_at 1.16168 far upstream element (FUSE) binding protein 1 Fubp1
1437545_at 1.03892 REST corepressor 1 Rcor1
1437546_at 0.91929 "DnaJ (Hsp40) homolog, subfamily C, member 14" Dnajc14
1437547_s_at 0.88725 "DnaJ (Hsp40) homolog, subfamily C, member 14" Dnajc14
1437548_at 1.13138 bicaudal D homolog 1 (Drosophila) Bicd1
1437549_at 1.07690 RIKEN cDNA 2810408I11 gene 2810408I11Rik
1437550_at 0.94081 DEAH (Asp-Glu-Ala-His) box polypeptide 36 Dhx36
1437551_at 1.29216 RIKEN cDNA 4930504E06 gene 4930504E06Rik
1437552_at 0.96904 "RIKEN cDNA 2410127L17 gene /// similar to RIKEN cDNA 2410127L17 ///
similar to RIKEN cDNA 2410127L17 /// hypothetical protein LOC665401 /// hypothetical protein
LOC665714 /// similar to CG11596-PA, isoform A /// similar to CG11596-PA, isoform A /// similar to
RIKEN cDNA 2410127L17 gene" 2410127L17Rik /// LOC624038 /// LOC626107 /// LOC665401 ///
LOC665714 /// LOC666456 /// LOC677519 /// LOC677553
1437553_at 1.02427 "BRCA1/BRCA2-containing complex, subunit 3" Brcc3
1437554_at 0.95797 plectin 1 Plec1
1437555_at 1.04819 RIKEN cDNA E130309B19 gene E130309B19Rik
1437556_at 0.97489 zinc finger homeodomain 4 Zfhx4
1437557_at 0.91504 "DNA segment, Chr 3, ERATO Doi 254, expressed" D3Ertd254e
1437558_at 0.55992 RIKEN cDNA B130021B11 gene B130021B11Rik
1437559_at 0.99908 "DNA segment, Chr 13, Brigham & Women's Genetics 1146 expressed"
D13Bwg1146e
1437560_at 0.91792 "neurotrophic tyrosine kinase, receptor, type 2" Ntrk2
1437561_at 0.97400 T cell receptor associated transmembrane adaptor 1 Trat1
1437562_at 0.97430 RIKEN cDNA 4932422M17 gene 4932422M17Rik
1437563_at 0.96509 PHD finger protein 20-like 1 Phf2011
1437564_at 1.27011 "polymerase (DNA directed), gamma" Polg

1437565_a_at 1.09487 similar to guanine nucleotide binding protein-like 2 (nucleolar)
 LOC633966
 1437566_at 1.04429 similar to guanine nucleotide binding protein-like 2 (nucleolar) LOC633966
 1437567_at 1.02972 CDNA sequence BC023179 BC023179
 1437568_at 0.93920 matrix metalloproteinase 16 Mmp16
 1437569_at 0.95063 transmembrane protein 132E Tmem132e
 1437570_at 1.40021 expressed sequence AI503301 AI503301
 1437571_at 1.25620 hypermethylated in cancer 2 Hic2
 1437572_at 0.90922 DCP1 decapping enzyme homolog b (S. cerevisiae) Dcp1b
 1437573_at 1.08766 "Mus musculus, clone IMAGE:3493537, mRNA" ---
 1437574_at 1.03322 "a disintegrin-like and metalloproteinase (reprolysin type) with thrombospondin
 type 1 motif, 18" Adamts18
 1437575_at 1.08988 minichromosome maintenance deficient domain containing 1 Mcmdc1
 1437576_at 1.36038 RIKEN cDNA 2810427A07 gene 2810427A07Rik
 1437577_at 2.31810 --- ---
 1437578_at 1.11042 chloride channel calcium activated 1 /// chloride channel calcium activated 2
 Clca1 /// Clca2
 1437579_at 0.92110 NIMA (never in mitosis gene a)-related expressed kinase 2 Nek2
 1437580_s_at 1.08639 NIMA (never in mitosis gene a)-related expressed kinase 2 Nek2
 1437581_at 1.04416 EST AA407452 AA407452
 1437582_at 1.06042 acyl-CoA synthetase short-chain family member 1 Acss1
 1437583_x_at 1.14325 acyl-CoA synthetase short-chain family member 1 Acss1
 1437584_at 1.07037 Cyclin D3 Cnd3
 1437585_x_at 0.84098 zinc finger protein 161 Zfp161
 1437586_at 1.01675 "CCR4-NOT transcription complex, subunit 4" Cnot4
 1437587_at 1.11549 Transcribed locus ---
 1437588_at 0.97642 "POU domain, class 4, transcription factor 2" Pou4f2
 1437589_x_at 0.88580 Lck interacting transmembrane adaptor 1 Lime1
 1437590_at 0.96909 RIKEN cDNA 4833409A17 gene 4833409A17Rik
 1437591_a_at 0.93557 WD repeat domain 1 Wdr1
 1437592_x_at 0.98325 coiled-coil domain containing 101 Ccdc101
 1437593_x_at 0.96440 apoptosis inhibitor 5 Api5
 1437594_x_at 0.88389 "phosphatidylinositol glycan anchor biosynthesis, class T" Pigt
 1437595_at 0.98627 RIKEN cDNA E030010A14 gene E030010A14Rik
 1437596_at 0.96661 Ly6/Plaur domain containing 4 Lypd4
 1437597_at 0.94262 Transcribed locus ---
 1437598_at 0.93460 zinc finger and BTB domain containing 20 Zbtb20
 1437599_at 1.41784 Coiled-coil-helix-coiled-coil-helix domain containing 3 Chchd3
 1437600_at 0.96375 "Atpase, class VI, type 11C" Atp11c
 1437601_at 0.95238 orthodenticle homolog 1 (Drosophila) Otx1
 1437602_at 0.99281 similar to Hypothetical protein MGC26988 RP23-27307.4
 1437603_at 1.09320 cDNA sequence BC050777 BC050777
 1437604_x_at 0.89273 adenomatous polyposis coli down-regulated 1 Apcdd1

1437605_at 0.94676 "nephrosis 2 homolog, podocin (human)" Nphs2
1437606_at 0.99760 "Transcribed locus, weakly similar to NP_808467.1 hypothetical protein
LOC327860 [Mus musculus]" ---
1437607_at 1.01167 "Glucosaminyl (N-acetyl) transferase 2, I-branching enzyme" Gcnt2
1437608_x_at 0.90875 "tyrosine 3-monooxygenase/tryptophan 5-monooxygenase activation protein,
theta polypeptide" Ywhaq
1437609_at 0.96960 ubiquitin-conjugating enzyme E2U (putative) Ube2u
1437610_x_at 1.04856 ribosomal protein S8 Rps8
1437611_x_at 1.08377 kinesin family member 2C Kif2c
1437612_at 0.92161 Nucleoporin 62 Nup62
1437613_s_at 0.75490 expressed sequence AW456874 AW456874
1437614_x_at 0.97942 "zinc finger, DHHC domain containing 14" Zdhhc14
1437615_s_at 0.96174 Vacuolar protein sorting 37C (yeast) Vps37c
1437616_x_at 1.01182 "zinc finger, DHHC domain containing 14" Zdhhc14
1437617_x_at 0.99783 RIKEN cDNA 1110034G24 gene 1110034G24Rik
1437618_x_at 0.94019 G protein-coupled receptor 85 Gpr85
1437619_x_at 1.05630 "discoidin domain receptor family, member 1" Ddr1
1437620_x_at 1.04947 "Aldehyde dehydrogenase 18 family, member A1" Aldh18a1
1437621_x_at 1.16164 3-phosphoglycerate dehydrogenase /// similar to 3-phosphoglycerate
dehydrogenase /// similar to 3-phosphoglycerate dehydrogenase /// similar to 3-phosphoglycerate
dehydrogenase Phgdh /// LOC668771 /// LOC671972 /// LOC673015
1437622_x_at 1.03737 mitochondrial ribosomal protein L28 Mrpl28
1437623_x_at 1.09404 X-ray repair complementing defective repair in Chinese hamster cells 3 Xrcc3
1437624_x_at 0.91185 nudix (nucleoside diphosphate linked moiety X)-type motif 16-like 1
Nudt16l1
1437625_at 0.97431 "Ribosomal protein S6 kinase, polypeptide 5" Rps6ka5
1437626_at 1.00514 "zinc finger protein 36, C3H type-like 2" Zfp3612
1437627_at 0.96909 ring finger (C3HC4 type) and KH domain containing 1 Rkhd1
1437628_s_at 0.98653 "ras homolog gene family, member A" Rhoa
1437629_at 0.84634 Rho guanine nucleotide exchange factor (GEF) 19 Arhgef19
1437630_at 0.97246 large subunit GTPase 1 homolog (S. cerevisiae) Lsg1
1437631_at 1.05826 Kv channel interacting protein 4 Kcnip4
1437632_at 0.98236 "Adult male cecum cDNA, RIKEN full-length enriched library,
clone:9130409M07 product:hypothetical protein, full insert sequence" ---
1437633_at 1.08325 ankyrin repeat domain 11 Ankrd11
1437634_at 1.00134 THO complex 2 Thoc2
1437635_at 1.12627 "discoidin, CUB and LCCL domain containing 2" Dcbl2
1437636_at 0.91661 Similar to Interferon-activatable protein 203 (Ifi-203) (Interferon-inducible
protein p203) LOC623121
1437637_at 0.97397 putative homeodomain transcription factor 2 Phtf2
1437638_at 2.01974 serine/arginine repetitive matrix 2 Srrm2
1437639_at 0.64076 RIKEN cDNA 3321401G04 gene 3321401G04Rik
1437640_at 1.01319 RIKEN cDNA 6430704M03 gene 6430704M03Rik

1437641_at 0.76918 RIKEN cDNA 4930535B03 gene 4930535B03Rik
1437642_at 1.21376 HIV-1 Rev binding protein-like Hrbl
1437643_at 1.06008 centromere protein J Cenpj
1437644_at 1.03114 "UDP-Gal:betaGlcNAc beta 1,3-galactosyltransferase, polypeptide 2"
B3galt2
1437645_at 1.05165 activating transcription factor 7 Atf7
1437646_at 0.87185 zinc and ring finger 3 /// similar to Goliath homolog precursor (Ring finger
protein 130) (R-goliath) Znrf3 /// LOC631806
1437647_at 1.21089 death inducer-obliterator 1 Dido1
1437648_at 1.17024 "phosphate cytidylyltransferase 1, choline, beta isoform" Pcyt1b
1437649_x_at 0.95832 peptidylprolyl isomerase B Ppib
1437650_at 1.08839 hypothetical protein C730026J16 C730026J16
1437651_a_at 0.87964 "dystrobrevin, beta" Dtnb
1437652_at 1.03486 "sema domain, transmembrane domain (TM), and cytoplasmic domain,
(semaphorin) 6C" Sema6c
1437653_at 1.06472 "immunity-related GTPase family, Q" Irgq
1437654_at 1.09894 RIKEN cDNA 3110001K24 gene 3110001K24Rik
1437655_at 0.89653 RIKEN cDNA D430020J02 gene D430020J02Rik
1437656_at 0.97759 Rho guanine nucleotide exchange factor (GEF) 4 Arhgef4
1437657_at 0.87472 zinc finger protein 291 Zfp291
1437658_a_at 1.15237 "small nucleolar RNA, C/D box 22" Snord22
1437659_at 0.96940 "amyotrophic lateral sclerosis 2 (juvenile) chromosome region, candidate 11
(human)" Als2cr11
1437660_at 1.42515 natural killer tumor recognition sequence Nktr
1437661_at 0.84874 expressed sequence AU021092 AU021092
1437662_at 1.09186 RIKEN cDNA C730027J19 gene C730027J19Rik
1437663_at 1.11072 "phosphatidylinositol glycan anchor biosynthesis, class V" Pigv
1437664_at 1.04101 homeo box D3 Hoxd3
1437665_at 1.12099 "Interleukin 22 receptor, alpha 2" Il22ra2
1437666_x_at 0.91356 ubiquitin C Ubc
1437667_a_at 1.24738 BTB and CNC homology 2 Bach2
1437668_at 0.89921 Chemokine (C-C motif) receptor-like 1 Ccr1l
1437669_x_at 0.89121 Chemokine (C-C motif) receptor-like 1 Ccr1l
1437670_x_at 0.89717 CD151 antigen Cd151
1437671_x_at 1.03500 "protease, serine, 23" Prss23
1437672_at 1.04742 insulin receptor substrate 3 Irs3
1437673_at 1.01191 Transcribed locus ---
1437674_at 1.22619 RAS-related C3 botulinum substrate 1 Rac1
1437675_at 0.96240 "solute carrier family 8 (sodium/calcium exchanger), member 1" Slc8a1
1437676_at 1.08441 sperm associated antigen 9 Spag9
1437677_at 1.14522 EST AI449595 AI449595
1437678_at 0.93954 "gene model 1564, (NCBI)" Gm1564
1437679_a_at 0.98472 glutaredoxin 2 (thioltransferase) Glrx2

1437680_x_at 1.10221 glutaredoxin 2 (thioltransferase) Glrx2
1437681_at 1.04423 "glutamate receptor, ionotropic, kainate 4" Grik4
1437682_x_at 1.03077 RIKEN cDNA 1110004E09 gene 1110004E09Rik
1437683_x_at 1.02122 small EDRK-rich factor 2 Serf2
1437684_at 0.96161 CDNA sequence BC025546 BC025546
1437685_x_at 0.66058 fibromodulin Fmod
1437686_x_at 0.90239 cut-like 1 (Drosophila) Cutl1
1437687_x_at 0.88238 FK506 binding protein 9 Fkbp9
1437688_x_at 0.98381 "ATPase, H+ transporting, lysosomal accessory protein 2" Atp6ap2
1437689_x_at 0.85299 clusterin Clu
1437690_x_at 0.94082 "casein kinase 1, delta" Csnk1d
1437691_at 0.90866 annexin A2 Anxa2
1437692_x_at 1.04168 annexin A2 Anxa2
1437693_at 1.03371 "DNA segment, Chr 1, Pasteur Institute 1" D1Pas1
1437694_at 1.01153 Expressed sequence BB114266 BB114266
1437695_at 0.92233 prokineticin receptor 2 Prokr2
1437696_at 1.04980 cDNA sequence BC049807 BC049807
1437697_at 0.86491 RAD23a homolog (S. cerevisiae) Rad23a
1437698_at 1.09273 myosin Vb Myo5b
1437699_at 0.98668 RIKEN cDNA E430014B02 gene E430014B02Rik
1437700_at 2.53823 Schwannomin interacting protein 1 Schip1
1437701_at 0.95461 "Smith-Magenis syndrome chromosome region, candidate 8 homolog (human)"
Smcr8
1437702_at 0.99622 transglutaminase 6 Tgm6
1437703_at 0.94227 similar to F-box- and WD40-repeat-containing protein LOC382156
1437704_at 1.21384 RIKEN cDNA 2900024O10 gene 2900024O10Rik
1437705_at 1.01106 zinc finger protein 551 Zfp551
1437706_x_at 1.09909 ribosomal protein S14 /// similar to ribosomal protein S14 Rps14 ///
LOC672818
1437707_at 0.93222 cDNA sequence BC019561 BC019561
1437708_x_at 0.99713 vesicle-associated membrane protein 3 Vamp3
1437709_x_at 0.89785 tetratricopeptide repeat domain 13 Ttc13
1437710_x_at 0.90140 RIKEN cDNA 1700021P22 gene 1700021P22Rik
1437711_x_at 1.00533 "ornithine decarboxylase, structural 1 /// similar to Ornithine decarboxylase
(ODC) /// similar to Ornithine decarboxylase (ODC) /// similar to Ornithine decarboxylase (ODC) ///
hypothetical protein LOC671973 /// similar to Ornithine decarboxylase (ODC)" Odc1 ///
LOC666231 /// LOC668343 /// LOC670279 /// LOC671973 /// LOC673302
1437712_x_at 1.02163 exosome component 4 Exosc4
1437713_x_at 0.91727 RIKEN cDNA 1500010J02 gene 1500010J02Rik
1437714_x_at 0.96911 Ubiquitin specific peptidase 14 Usp14
1437715_x_at 1.18694 apurinic/aprimidinic endonuclease 1 Apex1
1437716_x_at 1.01936 kinesin family member 22 Kif22
1437717_x_at 1.41776 RIKEN cDNA 2610005L07 gene /// RIKEN cDNA A430108E01 gene ///

similar to p47 protein isoform a 2610005L07Rik /// A430108E01Rik /// LOC547150

1437718_x_at 0.70663 fibromodulin Fmod

1437719_x_at 1.03390 RIKEN cDNA A230046K03 gene A230046K03Rik

1437720_at 1.03383 Ligatin Lgtn

1437721_at 1.18346 "coronin, actin binding protein 1C" Coro1c

1437722_x_at 0.92102 Poly(rC) binding protein 3 Pcbp3

1437723_s_at 0.99163 "Der1-like domain family, member 1" Der1l

1437724_x_at 0.90651 phosphatidylinositol membrane-associated 1 Pitpnm1

1437725_x_at 0.96198 TM2 domain containing 2 Tm2d2

1437726_x_at 1.16555 "complement component 1, q subcomponent, beta polypeptide" C1qb

1437727_at 0.98682 Odd Oz/ten-m homolog 3 (Drosophila) Odz3

1437728_at 1.21096 "alkB, alkylation repair homolog 5 (E. coli)" Alkbh5

1437729_at 0.95944 ribosomal protein L27a Rpl27a

1437730_at 1.05593 "protein phosphatase 2 (formerly 2A), regulatory subunit B (PR 52), alpha isoform" Ppp2r2a

1437731_at 0.86597 RIKEN cDNA 5830404H04 gene 5830404H04Rik

1437732_at 0.98664 "oocyte maturation, alpha /// similar to oocyte maturation, beta" Omt2a /// LOC640894

1437733_at 1.04455 eukaryotic translation initiation factor 4E binding protein 2 Eif4ebp2

1437734_at 0.97719 "protein phosphatase 1, regulatory (inhibitor) subunit 12A" Ppp1r12a

1437735_at 0.95549 "protein phosphatase 1, regulatory (inhibitor) subunit 12A" Ppp1r12a

1437736_at 0.95557 AKT1 substrate 1 (proline-rich) Akt1s1

1437737_at 0.91720 expressed sequence AV340375 AV340375

1437738_at 0.93137 "ATPase, Ca⁺⁺-sequestering" Atp2c1

1437739_a_at 0.93357 RIKEN cDNA 5430432N15 gene 5430432N15Rik

1437740_at 0.86437 "pleckstrin homology domain containing, family M (with RUN domain) member 2" Plekhm2

1437741_at 0.99514 "RAB21, member RAS oncogene family" Rab21

1437742_at 0.98364 "RAB21, member RAS oncogene family" Rab21

1437743_at 1.13576 AE binding protein 2 Aebp2

1437744_at 1.00298 "SLIT and NTRK-like family, member 4" Slitrk4

1437745_at 1.04570 chromodomain helicase DNA binding protein 7 Chd7

1437746_at 1.00193 leucine rich repeat transmembrane neuronal 1 Lrrtm1

1437747_at 1.03455 "ubiquitination factor E4A, UFD2 homolog (S. cerevisiae)" Ube4a

1437748_at 0.90582 fucosyltransferase 11 Fut11

1437749_s_at 0.96437 mitochondrial ribosomal protein L9 Mrpl9

1437750_at 1.08911 Transcribed locus ---

1437751_at 0.64629 "peroxisome proliferative activated receptor, gamma, coactivator 1 alpha" Ppargc1a

1437752_at 1.03985 lin-28 homolog (C. elegans) Lin28

1437753_at 1.01449 Expressed sequence AW742931 AW742931

1437754_at 0.94143 expressed sequence AW146299 AW146299

1437755_at 0.95177 "solute carrier family 5 (sodium/glucose cotransporter), member 12" Slc5a12

1437756_at 0.90459 "GTPase, IMAP family member 9" Gimap9
1437757_at 1.24161 --- ---
1437758_a_at 1.00504 "solute carrier family 4 (anion exchanger), member 1, adaptor protein"
Slc4a1ap
1437759_at 0.59282 "Phosphofructokinase, platelet" Pfkp
1437760_at 1.08722 UDP-N-acetyl-alpha-D-galactosamine:polypeptide N-
acetylgalactosaminyltransferase 12 Galnt12
1437761_at 1.01514 LUC7-like 2 (S. cerevisiae) Luc7l2
1437762_at 1.02023 "RAB39, member RAS oncogene family" Rab39
1437763_at 1.05504 "DCN1, defective in cullin neddylation 1, domain containing 3 (S. cerevisiae)"
Dcun1d3
1437764_at 0.93875 RIKEN cDNA A730011L01 gene A730011L01Rik
1437765_at 1.07576 cytoplasmic polyadenylation element binding protein 3 Cpeb3
1437766_at 0.87866 Transcribed locus ---
1437767_s_at 0.77992 fused toes Fts
1437768_at 1.19277 ankyrin repeat and IBR domain containing 1 Ankib1
1437769_at 1.18865 Epidermal growth factor receptor pathway substrate 15-like 1 Eps15l1
1437770_at 0.86140 expressed sequence AW557046 AW557046
1437771_at 1.06899 RIKEN cDNA 1110067I12 gene 1110067I12Rik
1437772_s_at 0.90863 "fucosidase, alpha-L- 1, tissue" Fuca1
1437773_x_at 0.91150 DEAD (Asp-Glu-Ala-Asp) box polypeptide 17 Ddx17
1437774_at 0.91786 RIKEN cDNA 1700020I14 gene 1700020I14Rik
1437775_at 1.41715 dihydrolipoamide S-succinyltransferase (E2 component of 2-oxo-glutarate
complex) Dlst
1437776_at 1.25679 transmembrane and coiled coil domains 1 Tmcc1
1437777_at 1.24885 activating transcription factor 2 Atf2
1437778_at 0.92157 RNA binding motif protein 15B Rbm15b
1437779_at 1.09473 forkhead box H1 Foxh1
1437780_at 1.17963 "Fanconi anemia, complementation group B" Fancb
1437781_at 0.96663 Transcribed locus ---
1437782_at 0.98021 contactin associated protein-like 2 Cntnap2
1437783_x_at 0.99977 transmembrane protein 4 Tmem4
1437784_at 1.11785 "runt-related transcription factor 1; translocated to, 1 (cyclin D-related)"
Runx1t1
1437785_at 0.96345 "a disintegrin-like and metallopeptidase (reprolysin type) with thrombospondin
type 1 motif, 9" Adamts9
1437786_at 0.97355 expressed sequence C80008 C80008
1437787_at 0.82888 Leucine rich repeat transmembrane neuronal 2 Lrrtm2
1437788_at 0.80537 trans-acting transcription factor 6 Sp6
1437789_at 2.30779 baculoviral IAP repeat-containing 6 Birc6
1437790_at 1.09088 echinoderm microtubule associated protein like 5 Eml5
1437791_s_at 1.00603 echinoderm microtubule associated protein like 5 Eml5
1437792_at 0.93763 RIKEN cDNA 1700013D24 gene 1700013D24Rik

1437793_at 1.94141 "Tankyrase, TRF1-interacting ankyrin-related ADP-ribose polymerase 2"
 Tnks2
 1437794_at 0.97105 calcineurin binding protein 1 Cabin1
 1437795_at 1.23832 --- ---
 1437796_at 1.05549 Transcribed locus ---
 1437797_at 1.83269 "ATPase, Ca++ transporting, cardiac muscle, slow twitch 2" Atp2a2
 1437798_at 1.16320 RIKEN cDNA 6720422M22 gene 6720422M22Rik
 1437799_at 1.13258 RIKEN cDNA 4933407H18 gene 4933407H18Rik
 1437800_at 0.95153 EDAR (ectodysplasin-A receptor)-associated death domain Edaradd
 1437801_at 1.02920 mortality factor 4 like 1 /// similar to mortality factor 4 like 1 isoform b ///
 similar to mortality factor 4 like 1 /// similar to mortality factor 4 like 1 (predicted) Morf411 ///
 LOC433955 /// LOC627352 /// LOC635663
 1437802_x_at 0.95172 mortality factor 4 like 1 /// similar to mortality factor 4 like 1 isoform b ///
 similar to mortality factor 4 like 1 /// similar to mortality factor 4 like 1 (predicted) Morf411 ///
 LOC433955 /// LOC627352 /// LOC635663
 1437803_at 0.96553 stem-loop binding protein /// similar to stem-loop binding protein /// similar to
 stem-loop binding protein /// similar to stem-loop binding protein Slbp /// LOC666976 ///
 LOC670567 /// LOC674607
 1437804_at 0.99227 Insulin-like growth factor binding protein 7 Igfbp7
 1437805_at 0.94816 --- ---
 1437806_x_at 0.94636 mitochondrial carrier homolog 2 (C. elegans) Mtch2
 1437807_x_at 0.97954 "catenin (cadherin associated protein), alpha 1" Ctnna1
 1437808_x_at 0.88651 "Interleukin 10 receptor, alpha" Il10ra
 1437809_x_at 1.01256 "E26 avian leukemia oncogene 2, 3' domain" Ets2
 1437810_a_at 0.97292 "hemoglobin Z, beta-like embryonic chain" Hbb-bh1
 1437811_x_at 0.87324 Coactosin-like 1 (Dictyostelium) Cotl1
 1437812_x_at 0.93208 alpha glucosidase 2 alpha neutral subunit Ganab
 1437813_at 1.01937 absent in melanoma 1-like Aim11
 1437814_at 0.93676 "DnaJ (Hsp40) homolog, subfamily C, member 5 gamma" Dnajc5g
 1437815_at 0.88710 RIKEN cDNA 1700111A04 gene 1700111A04Rik
 1437816_at 0.97814 paired box gene 6 Pax6
 1437817_at 1.00184 synaptosomal-associated protein 91 Snap91
 1437818_at 1.03718 RIKEN cDNA 9430016H08 gene 9430016H08Rik
 1437819_s_at 1.16395 RIKEN cDNA 9530020G05 gene 9530020G05Rik
 1437820_at 0.92590 forkhead-like 18 (Drosophila) Fkhl18
 1437821_at 1.84766 Diaphanous homolog 1 (Drosophila) Diap1
 1437822_at 1.22255 YME1-like 1 (S. cerevisiae) Yme11
 1437823_at 0.85219 RIKEN cDNA 2310040A07 gene 2310040A07Rik
 1437824_at 1.06380 "glutamate receptor, ionotropic, delta 2" Grid2
 1437825_at 0.91976 leucine rich repeat containing 4C Lrrc4c
 1437826_at 0.89569 Pore forming protein-like Pfpl
 1437827_s_at 0.95534 Pore forming protein-like Pfpl
 1437828_s_at 1.13804 WD repeat domain 46 Wdr46

1437829_s_at 0.71452 eukaryotic elongation factor-2 kinase Eef2k
1437830_x_at 0.92711 "zinc finger, BED domain containing 3" Zbed3
1437831_at 0.93021 "zinc finger, CCHC domain containing 4" Zcchc4
1437832_x_at 0.94470 tryptophanyl-tRNA synthetase Wars
1437833_at 1.50846 latent transforming growth factor beta binding protein 3 Ltbp3
1437834_s_at 0.96134 protein kinase C and casein kinase substrate in neurons 3 Pacsin3
1437835_a_at 0.93768 RIKEN cDNA 0610011L14 gene 0610011L14Rik
1437836_x_at 0.91414 RIKEN cDNA 0610011L14 gene 0610011L14Rik
1437837_x_at 0.89578 "polymerase (DNA-directed), delta interacting protein 3" Poldip3
1437838_x_at 0.96947 G-rich RNA sequence binding factor 1 Grsf1
1437839_x_at 0.96645 mitochondrial ribosomal protein L11 Mrpl11
1437840_s_at 1.03681 "similar to CG3835-PA, isoform A" LOC673662
1437841_x_at 0.83914 "cold shock domain containing C2, RNA binding" Csdc2
1437842_at 0.95265 "phosphatidylinositol-specific phospholipase C, X domain containing 1" Plcx1
1437843_s_at 1.06348 Nucleoporin like 1 Nupl1
1437844_x_at 0.97096 component of oligomeric golgi complex 1 Cog1
1437845_x_at 0.78764 protein O-fucosyltransferase 2 Pofut2
1437846_x_at 1.00163 beta-site APP-cleaving enzyme 2 Bace2
1437847_x_at 1.03549 "RNA binding motif protein, X chromosome" Rbmx
1437848_x_at 0.81453 ADP-dependent glucokinase Adpgk
1437849_x_at 1.08444 "armadillo repeat containing, X-linked 2" Armcx2
1437850_a_at 1.01019 cellular nucleic acid binding protein Cnbp
1437851_x_at 1.01768 ribonucleic acid binding protein S1 Rnps1
1437852_x_at 1.12075 cleavage and polyadenylation specificity factor 3 Cpsf3
1437853_x_at 0.90391 necdin Ndn
1437854_at 1.05693 Leucine rich repeat containing 18 Lrrc18
1437855_at 1.06594 microtubule-associated protein 4 Mtap4
1437856_at 1.13335 inositol polyphosphate multikinase Ipmk
1437857_at 0.87423 dpy-19-like 3 (C. elegans) Dpy193
1437858_at 1.02995 dpy-19-like 3 (C. elegans) Dpy193
1437859_x_at 1.21417 eukaryotic translation initiation factor 5A Eif5a
1437860_at 1.01431 "protein kinase C, epsilon" Prkce
1437861_s_at 0.90565 --- ---
1437862_at 1.79447 RIKEN cDNA 2600011C06 gene 2600011C06Rik
1437863_at 0.86342 butyrylcholinesterase Bche
1437864_at 1.03851 adiponectin receptor 2 Adipor2
1437865_at 0.94260 spermatogenesis associated 13 Spata13
1437866_at 1.11938 dual specificity phosphatase 18 Dusp18
1437867_at 1.10365 "2 cells egg cDNA, RIKEN full-length enriched library, clone:B020014N01
product:hypothetical protein, full insert sequence" ---
1437868_at 0.70564 cDNA sequence BC023892 BC023892
1437869_at 0.99408 RIKEN cDNA 3222402P14 gene 3222402P14Rik
1437870_at 1.01735 "solute carrier organic anion transporter family, member 4C1" Slco4c1

1437871_at 0.85319 phosphoglucomutase 5 Pgm5
1437872_at 1.14233 cDNA sequence AB112350 AB112350
1437873_at 0.75507 RIKEN cDNA 6030490I01 gene 6030490I01Rik
1437874_s_at 0.88284 hexosaminidase B Hexb
1437875_at 1.06439 bicaudal D homolog 2 (Drosophila) Bicd2
1437876_at 1.06892 fibronectin type III domain containing 6 Fndc6
1437877_at 1.00107 coiled-coil domain containing 39 /// tetratricopeptide repeat domain 14
Ccdc39 /// Ttc14
1437878_s_at 1.31114 coiled-coil domain containing 39 /// tetratricopeptide repeat domain 14
Ccdc39 /// Ttc14
1437879_at 1.01599 expressed sequence AU045404 AU045404
1437880_at 1.06892 ladybird homeobox 1 homolog (Drosophila) corepressor 1 Lbxcor1
1437881_at 1.09421 RIKEN cDNA 6330416L07 gene 6330416L07Rik
1437882_s_at 0.95671 RIKEN cDNA 3110048L19 gene /// RIKEN cDNA 6330416L07 gene
3110048L19Rik /// 6330416L07Rik
1437883_s_at 2.22842 --- ---
1437884_at 0.83603 ADP-ribosylation factor-like 5B Arl5b
1437885_at 0.90855 RIKEN cDNA D030029J20 gene D030029J20Rik
1437886_at 1.05022 kelch-like 6 (Drosophila) Klhl6
1437887_at 0.91521 RIKEN cDNA E130306M17 gene E130306M17Rik
1437888_at 0.90371 WNK lysine deficient protein kinase 1 Wnk1
1437889_x_at 0.98085 biglycan Bgn
1437890_at 0.98328 RIKEN cDNA 1500005I02 gene 1500005I02Rik
1437891_at 1.03772 fibroblast growth factor receptor substrate 2 Frs2
1437892_at 1.37330 zinc finger protein 306 Zfp306
1437893_at 0.89902 phospholipase B1 /// cDNA sequence BC033606 /// similar to phospholipase
B Pib1 /// BC033606 /// LOC665270
1437894_at 0.92631 prospero-related homeobox 1 Prox1
1437895_at 0.74108 transmembrane protein 16H Tmem16h
1437896_at 1.27326 Zinc finger CCHC-type and RNA binding motif 1 Zcrb1
1437897_at 0.98961 kelch-like 21 (Drosophila) Klhl21
1437898_at 0.87952 RIKEN cDNA C920009B18 gene C920009B18Rik
1437899_at 0.88798 "gene model 776, (NCBI)" Gm776
1437900_at 1.12166 RIKEN cDNA 4930523C07 gene 4930523C07Rik
1437901_a_at 0.98188 vacuolar protein sorting 41 (yeast) Vps41
1437902_s_at 0.90776 retinoic acid receptor responder (tazarotene induced) 2 Rarres2
1437903_at 0.84852 lysyl oxidase Lox
1437904_at 0.93394 developmentally regulated RNA binding protein 1 Drbp1
1437905_at 1.09767 thioredoxin-like 1 Txnl1
1437906_x_at 1.04367 thioredoxin-like 1 Txnl1
1437907_a_at 1.04254 tubulin cofactor a Tbca
1437908_a_at 0.87698 RIKEN cDNA 1200007D18 gene 1200007D18Rik
1437909_at 1.01298 RIKEN cDNA D030022P06 gene D030022P06Rik

1437910_at 0.99733 transmembrane protein 39b Tmem39b
1437911_at 0.95920 RIKEN cDNA 6330416L07 gene 6330416L07Rik
1437912_at 0.95760 IQ motif and Sec7 domain 3 Iqsec3
1437913_at 0.97593 B-cell leukemia/lymphoma 2 related protein A1a /// B-cell leukemia/lymphoma
2 related protein A1b /// B-cell leukemia/lymphoma 2 related protein A1d Bcl2a1a /// Bcl2a1b ///
Bcl2a1d
1437914_at 1.07810 E2F transcription factor 6 E2f6
1437915_at 1.15990 target of myb1-like 2 (chicken) Tom112
1437916_at 1.17893 --- ---
1437917_at 1.43934 RIKEN cDNA D530037H12 gene D530037H12Rik
1437918_at 1.06393 RIKEN cDNA 4930539E08 gene 4930539E08Rik
1437919_at 1.44196 "B double prime 1, subunit of RNA polymerase III transcription initiation factor
IIIB" Bdp1
1437920_at 0.92318 Eph receptor A5 Epha5
1437921_x_at 0.87235 zinc finger protein 516 Zfp516
1437922_at 1.05709 "DNA segment, Chr 10, ERATO Doi 322, expressed /// similar to mitochondrial
ribosomal protein L42 /// similar to mitochondrial ribosomal protein L42" D10Ert322e ///
LOC623346 /// LOC630452
1437923_at 1.18079 expressed sequence AI314760 AI314760
1437924_at 0.88467 --- ---
1437925_at 1.07142 RIKEN cDNA 2210018M03 gene 2210018M03Rik
1437926_at 1.06465 "X-prolyl aminopeptidase (aminopeptidase P) 3, putative" Xpnpep3
1437927_at 1.00213 "discs, large homolog 2 (Drosophila)" Dlgh2
1437928_at 0.88625 protocadherin 12 Pcdh12
1437929_at 0.87550 "dapper homolog 2, antagonist of beta-catenin (xenopus)" Dact2
1437930_at 1.00114 glycosyltransferase 25 domain containing 2 Glt25d2
1437931_at 0.97237 toll-like receptor 12 Tlr12
1437932_a_at 0.95147 claudin 1 Cldn1
1437933_at 0.89056 Hedgehog-interacting protein Hhip
1437934_at 1.04800 cytotoxic granule-associated RNA binding protein 1 Tia1
1437935_at 0.95663 RIKEN cDNA 4930486G11 gene 4930486G11Rik
1437936_at 1.04633 RIKEN cDNA 6330534C20 gene 6330534C20Rik
1437937_at 0.97864 chemokine binding protein 2 Ccbp2
1437938_x_at 1.09346 dynamin 2 Dnm2
1437939_s_at 0.98068 cathepsin C Ctsc
1437940_at 0.98309 "amyloid beta (A4) precursor protein binding, family A, member 1" Apba1
1437941_at 0.94048 phosphorylase kinase alpha 2 Phka2
1437942_x_at 1.11362 epsilon-tubulin 1 Tube1
1437943_s_at 1.10023 male enhanced antigen 1 Mea1
1437944_at 1.09205 Src homology 2 domain-containing transforming protein C2 Shc2
1437945_x_at 0.96710 nucleosome assembly protein 1-like 1 Nap111
1437946_x_at 1.02319 Rap guanine nucleotide exchange factor (GEF) 3 Rapgef3
1437947_x_at 1.00214 voltage-dependent anion channel 1 Vdac1

1437948_x_at 0.98910 "eukaryotic translation initiation factor 3, subunit 6 interacting protein"
Eif3s6ip

1437949_x_at 0.98846 zinc finger protein 313 Zfp313

1437950_at 0.89937 cDNA sequence BC035537 BC035537

1437951_at 1.02517 RIKEN cDNA 4930429A22 gene 4930429A22Rik

1437952_at 1.01690 expressed sequence AU040096 AU040096

1437953_at 1.02327 preimplantation protein 4 Prei4

1437954_at 1.00051 --- ---

1437955_at 1.04266 RIKEN cDNA 4922504M18 gene 4922504M18Rik

1437956_at 0.94032 expressed sequence BB220380 BB220380

1437957_at 1.23328 RIKEN cDNA 7030407O06 gene 7030407O06Rik

1437958_at 0.82747 xenotropic and polytropic retrovirus receptor 1 Xpr1

1437959_at 1.12466 zinc finger protein 324 Zfp324

1437960_at 1.01349 calpain 13 Capn13

1437961_x_at 0.83454 Expressed sequence AA407659 AA407659

1437962_at 1.04950 "gene model 443, (NCBI)" Gm443

1437963_at 1.07481 RIKEN cDNA C530045E16 gene C530045E16Rik

1437964_at 0.94189 neurexophilin 2 Nxph2

1437965_at 1.25962 HEAT repeat containing 1 Heatr1

1437966_at 0.95402 proline-rich transmembrane protein 3 Prrt3

1437967_at 0.95315 Transcribed locus ---

1437968_at 0.94779 "glutamate receptor, ionotropic, NMDA1 (zeta 1)" Grin1

1437969_s_at 1.12806 RIKEN cDNA 0610007P22 gene 0610007P22Rik

1437970_at 0.86435 CDNA sequence BC053401 BC053401

1437971_at 0.97886 Superkiller viralicidic activity 2-like 2 (S. cerevisiae) Skiv2l2

1437972_s_at 1.12713 "splicing factor 3b, subunit 5" Sf3b5

1437973_at 0.94478 leucine rich repeat neuronal 6C Lrrn6c

1437974_a_at 1.02622 hexokinase 1 Hk1

1437975_a_at 1.02150 ribosomal protein L23a /// hypothetical gene supported by BC029892 ///
similar to 60S ribosomal protein L23a /// similar to 60S ribosomal protein L23a /// similar to 60S
ribosomal protein L23a /// similar to 60S ribosomal protein L23a Rpl23a /// LOC385336 ///
LOC666448 /// LOC666501 /// LOC666548 /// LOC666738

1437976_x_at 1.01135 ribosomal protein L23a Rpl23a

1437977_at 1.20010 "Small glutamine-rich tetratricopeptide repeat (TPR)-containing, beta" Sgtb

1437978_at 0.97582 Slit homolog 2 (Drosophila) Slit2

1437979_at 1.15771 "Zinc finger, CCHC domain containing 2" Zcchc2

1437980_at 0.93224 RIKEN cDNA 9130230N09 gene 9130230N09Rik

1437981_x_at 1.00856 RIKEN cDNA 1110057K04 gene 1110057K04Rik

1437982_x_at 1.00152 "COX15 homolog, cytochrome c oxidase assembly protein (yeast)" Cox15

1437983_at 0.82419 --- ---

1437984_x_at 0.99342 HLA-B-associated transcript 1A Bat1a

1437985_a_at 1.07549 RIKEN cDNA 2310061I04 gene 2310061I04Rik

1437986_x_at 0.91351 fucokinase Fuk

1437987_at 0.92940 --- ---
1437988_x_at 1.05399 RIKEN cDNA 1700003E24 gene 1700003E24Rik
1437989_at 0.95642 phosphodiesterase 8B Pde8b
1437990_x_at 1.03773 "hemoglobin Z, beta-like embryonic chain" Hbb-bh1
1437991_x_at 1.00034 RUN and SH3 domain containing 1 Rusc1
1437992_x_at 0.92240 gap junction membrane channel protein alpha 1 Gja1
1437993_x_at 1.03952 quinoid dihydropteridine reductase Qdpr
1437994_x_at 1.00222 "mesoderm induction early response 1, family member 2" Mier2
1437995_x_at 0.96993 septin 7 7-Sep
1437996_s_at 1.01549 RIKEN cDNA 1500012D20 gene 1500012D20Rik
1437997_x_at 1.06515 mitochondrial ribosomal protein L48 Mrpl48
1437998_at 0.99300 "Adaptor-related protein complex 1, sigma 2 subunit" Ap1s2
1437999_x_at 0.96126 "phosphatidylinositol glycan anchor biosynthesis, class Q" Pigq
1438000_x_at 0.99771 zinc finger protein 622 Zfp622
1438001_x_at 0.93865 receptor accessory protein 5 Reep5
1438002_at 0.90774 --- ---
1438003_at 1.03148 polymerase (DNA directed) sigma Pols
1438004_at 1.01377 polymerase (DNA directed) sigma Pols
1438005_at 0.92962 RIKEN cDNA B230315F11 gene B230315F11Rik
1438006_at 1.02162 RIKEN cDNA 4933439F18 gene 4933439F18Rik
1438007_at 0.89254 expressed sequence AI851790 AI851790
1438008_at 0.88527 "Golgi associated, gamma adaptin ear containing, ARF binding protein 3" Gga3
1438009_at 1.10414 similar to histone 2a MGC73635
1438010_at 0.96498 "DNA segment, Chr 4, ERATO Doi 429, expressed" D4Ertd429e
1438011_at 0.98575 "phosphate cytidylyltransferase 1, choline, alpha isoform" Pcyt1a
1438012_at 1.03990 protein phosphatase 1 (formerly 2C)-like Ppm1l
1438013_x_at 0.89881 DEAH (Asp-Glu-Ala-His) box polypeptide 32 Dhx32
1438014_at 0.79326 mitochondrial ribosomal protein L34 Mrpl34
1438015_at 1.17263 cDNA sequence BC068171 BC068171
1438016_at 1.24173 cDNA sequence BC068171 BC068171
1438017_at 0.99392 RUN and SH3 domain containing 1 Rusc1
1438018_at 0.95735 hook homolog 1 (Drosophila) Hook1
1438019_at 1.03796 "inositol 1,3,4,5,6-pentakisphosphate 2-kinase" Ippk
1438020_at 1.14410 hyaluronan and proteoglycan link protein 1 Hapln1
1438021_at 0.92354 cDNA sequence BC013481 BC013481
1438022_at 0.96413 RAB11 family interacting protein 3 (class II) Rab11fip3
1438023_at 0.86485 spermatogenesis associated 6 Spata6
1438024_at 0.85808 RIKEN cDNA 6230416A05 gene 6230416A05Rik
1438025_at 1.05505 mitochondrial translational release factor 1-like Mtrf1l
1438026_at 0.92627 zinc finger protein 560 Zfp560
1438027_at 1.30641 --- ---
1438028_at 1.04913 RIKEN cDNA 4930535B03 gene 4930535B03Rik
1438029_at 0.90403 RIKEN cDNA 4930535B03 gene 4930535B03Rik

1438030_at	0.87982	"RAS, guanyl releasing protein 3"	Rasgrp3
1438031_at	1.05264	"RAS, guanyl releasing protein 3"	Rasgrp3
1438032_at	0.86003	leucine-rich repeats and calponin homology (CH) domain containing 1	Lrch1
1438033_at	1.03801	thyrotroph embryonic factor	Tef
1438034_at	0.90895	RIKEN cDNA 2410005O16 gene	2410005O16Rik
1438035_at	0.85723	expressed sequence AW061290	AW061290
1438036_x_at	0.88348	expressed sequence AW061290	AW061290
1438037_at	1.02457	hect domain and RLD 5	Herc5
1438038_at	0.85071	RIKEN cDNA 4930402H24 gene	4930402H24Rik
1438039_at	0.97646	HECT domain containing 1	Hectd1
1438040_a_at	0.90944	"heat shock protein 90kDa beta (Grp94), member 1"	Hsp90b1
1438041_at	1.20659	---	---
1438042_at	0.76603	short stature homeobox 2	Shox2
1438043_at	1.04158	"eukaryotic translation initiation factor 2C, 2"	Eif2c2
1438044_at	1.02043	RIKEN cDNA 1700047M11 gene	1700047M11Rik
1438045_at	0.89620	early endosome antigen 1	Eea1
1438046_at	0.99116	expressed sequence AU019823	AU019823
1438047_at	1.02228	zinc finger protein 384	Zfp384
1438048_at	1.02250	myc target 1	Myct1
1438049_at	1.99117	RIKEN cDNA A430108E01 gene	A430108E01Rik
1438050_x_at	0.97955	---	---
1438051_at	0.85507	tetratricopeptide repeat domain 14	Ttc14
1438052_at	1.02086	RIKEN cDNA A130071D04 gene	A130071D04Rik
1438053_at	0.95877	Trk-fused gene	Tfg
1438054_x_at	1.01965	"protein phosphatase 2 (formerly 2A), regulatory subunit B (PR 52), gamma isoform"	Ppp2r2c
1438055_at	1.04319	retinoic acid receptor responder (tazarotene induced) 1	Rarres1
1438056_x_at	1.00063	"ATP-binding cassette, sub-family C (CFTR/MRP), member 5"	Abcc5
1438057_at	0.99744	Transcribed locus	---
1438058_s_at	0.80434	prostate tumor over expressed gene 1	Ptov1
1438059_at	1.13474	similar to Cortexin-1	LOC629147
1438060_at	0.89764	neuronal PAS domain protein 3	Npas3
1438061_at	1.00081	RIKEN cDNA 4930523C07 gene	4930523C07Rik
1438062_at	0.98123	RIKEN cDNA 4832420A03 gene	4832420A03Rik
1438063_at	0.99201	M-phase phosphoprotein 9	Mphosph9
1438064_at	1.44910	Y box protein 1	Ybx1
1438065_at	1.14635	---	---
1438066_at	1.02914	General transcription factor IIB	Gtf2b
1438067_at	1.18525	neurofibromatosis 1	Nf1
1438068_at	0.81506	"Transcribed locus, strongly similar to XP_579813.1 PREDICTED: hypothetical protein XP_579813 [Rattus norvegicus]"	---
1438069_a_at	1.06734	RNA binding motif protein 5	Rbm5
1438070_at	0.83979	PHD finger protein 3	Phf3

1438071_at 1.03437 postmeiotic segregation increased 1 (S. cerevisiae) Pms1
1438072_at 2.96479 Nuclear factor I/B Nfib
1438073_at 0.86917 sprouty homolog 3 (Drosophila) Spry3
1438074_at 0.94721 RIKEN cDNA 2210010C17 gene 2210010C17Rik
1438075_at 0.88273 fem-1 homolog c (C.elegans) Fem1c
1438076_at 1.09334 ribosomal protein L30 Rpl30
1438077_at 0.93848 "NACHT, leucine rich repeat and PYD containing 4A" Nalp4a
1438078_at 1.07111 Transcribed locus ---
1438079_at 1.26008 cDNA sequence BC050078 BC050078
1438080_at 0.92890 mitochondrial ribosomal protein L11 Mrpl11
1438081_at 0.92416 mutated in colorectal cancers Mcc
1438082_at 0.87629 RIKEN cDNA 2310028N02 gene 2310028N02Rik
1438083_at 1.05405 Hedgehog-interacting protein Hhip
1438084_at 1.01636 A disintegrin and metallopeptidase domain 23 Adam23
1438085_at 0.80202 "RIKEN cDNA A230048G03 gene /// similar to CG2747-PB, isoform B"
A230048G03Rik /// LOC670614
1438086_at 0.94856 neuropeptide Y receptor Y6 Npy6r
1438087_at 0.87811 thiopurine methyltransferase Tpm1
1438088_at 0.89042 "Adult male cecum cDNA, RIKEN full-length enriched library,
clone:9130222J03 product:unclassifiable, full insert sequence" ---
1438089_a_at 1.15884 BCL2-associated transcription factor 1 Bclaf1
1438090_x_at 0.95879 ankyrin repeat domain 54 Ankrd54
1438091_a_at 1.02489 "H2A histone family, member Z /// similar to H2A histone family, member
Z /// similar to H2A histone family, member Z" H2afz /// LOC631864 /// LOC675899
1438092_x_at 1.02065 "H2A histone family, member Z /// similar to H2A histone family, member
Z /// similar to H2A histone family, member Z /// similar to H2A histone family, member Z /// similar to
H2A histone family, member Z" H2afz /// LOC631864 /// LOC666634 /// LOC669445 /// LOC675899
1438093_x_at 1.21777 diazepam binding inhibitor Dbi
1438094_x_at 1.10908 RIKEN cDNA 2810409H07 gene 2810409H07Rik
1438095_x_at 0.89641 Nucleolar complex associated 4 homolog (S. cerevisiae) Noc4l
1438096_a_at 1.00509 deoxythymidylate kinase Dtymk
1438097_at 1.33399 "RAB20, member RAS oncogene family" Rab20
1438098_at 0.84435 "discs, large (Drosophila) homolog-associated protein 1" Dlgap1
1438099_at 0.88560 triple functional domain (PTPRF interacting) Trio
1438100_at 1.00118 RIKEN cDNA B230208H17 gene B230208H17Rik
1438101_at 1.00917 Frizzled homolog 3 (Drosophila) Fzd3
1438102_at 0.98221 SUMO/sentrin specific peptidase 8 Senp8
1438103_at 1.03120 "protocadherin gamma subfamily C, 4" Pcdhgc4
1438104_at 1.24721 --- ---
1438105_at 0.97660 hypothetical protein LOC632686 LOC632687
1438106_at 0.91300 protocadherin beta 22 Pcdhb22
1438107_x_at 0.97067 expressed sequence BB236558 BB236558
1438108_at 1.12215 RIKEN cDNA 9430067K14 gene 9430067K14Rik

1438109_at	0.96589 chloride channel calcium activated 5	Clca5
1438110_at	1.20625 Zinc finger and BTB domain containing 1	Zbtb1
1438111_at	1.02648 Poliovirus receptor-related 1	Pvr1
1438112_at	0.84207 RIKEN cDNA 9430021M05 gene	9430021M05Rik
1438113_at	0.83679 "zinc finger, matrin type 4"	Zmat4
1438114_x_at	0.95994 Embryonal Fyn-associated substrate	Efs
1438115_a_at	0.90859 "solute carrier family 9 (sodium/hydrogen exchanger), isoform 3 regulator 1"	Slc9a3r1
1438116_x_at	0.91441 "solute carrier family 9 (sodium/hydrogen exchanger), isoform 3 regulator 1"	Slc9a3r1
1438117_x_at	0.96255 transmembrane protein 41B	Tmem41b
1438118_x_at	1.01334 vimentin	Vim
1438119_at	1.15625 Integrin beta 1 (fibronectin receptor beta)	Itgb1
1438120_x_at	1.05663 ---	---
1438121_at	0.97002 "SEC24 related gene family, member A (<i>S. cerevisiae</i>)"	Sec24a
1438122_at	1.10218 RIKEN cDNA 2900006K08 gene	2900006K08Rik
1438123_at	1.07380 ---	---
1438124_at	0.88961 similar to putative pheromone receptor (Go-VN5) /// similar to putative pheromone receptor (Go-VN5) /// similar to putative pheromone receptor (Go-VN5) /// similar to putative pheromone receptor (Go-VN5) /// similar to putative pheromone receptor (Go-VN5) /// similar to putative pheromone receptor (Go-VN5) /// similar to putative pheromone receptor (Go-VN5) /// similar to putative pheromone receptor (Go-VN5) /// similar to putative pheromone receptor (Go-VN5) /// similar to putative pheromone receptor (Go-VN5) /// similar to putative pheromone receptor (Go-VN5) /// similar to putative pheromone receptor (Go-VN5) /// similar to putative pheromone receptor (Go-VN5) ///	LOC195691 /// LOC224552 /// LOC224572 /// LOC433070 /// LOC627476 /// LOC637519 /// LOC639943 /// LOC665253 /// LOC665376 /// LOC670484 /// LOC676392
1438125_at	1.02391 RIKEN cDNA C230085N15 gene	C230085N15Rik
1438126_at	1.21571 exocyst complex component 5	Exoc5
1438127_at	1.76814 hypothetical LOC552906	LOC552906
1438128_at	1.04209 "Guanine nucleotide binding protein, alpha transducing 2"	Gnat2
1438129_at	1.36319 WD repeat and SOCS box-containing 2	Wsb2
1438130_at	2.14654 "TAF15 RNA polymerase II, TATA box binding protein (TBP)-associated factor"	Taf15
1438131_at	1.18356 F-box and WD-40 domain protein 2	Fbxw2
1438132_at	1.09370 RIKEN cDNA C030005G22 gene	C030005G22Rik
1438133_a_at	0.69729 cysteine rich protein 61	Cyr61
1438134_at	1.01689 protocadherin 10	Pcdh10
1438135_at	1.06004 RIKEN cDNA E130114P18 gene	E130114P18Rik
1438136_at	1.20749 astacin-like metalloendopeptidase (M12 family)	Astl
1438137_at	1.02679 RIKEN cDNA 2810055G22 gene	2810055G22Rik
1438138_a_at	0.94901 peroxisomal biogenesis factor 6	Pex6
1438139_at	0.93590 Rho GTPase activating protein 28	Arhgap28
1438140_a_at	0.93182 zinc finger protein 64	Zfp64
1438141_at	0.91998 cDNA sequence BC029684	BC029684

1438142_s_at 1.00655 RIKEN cDNA 1700021K14 gene 1700021K14Rik
1438143_s_at 0.99458 ataxin 2 Atxn2
1438144_x_at 0.98725 ataxin 2 Atxn2
1438145_at 1.01184 "SPT2, Suppressor of Ty, domain containing 1 (S. cerevisiae)" Spty2d1
1438146_x_at 0.88018 "SPT2, Suppressor of Ty, domain containing 1 (S. cerevisiae)" Spty2d1
1438147_at 1.14757 "scavenger receptor cysteine rich domain containing, group B (4 domains)" Srcrb4d
1438148_at 0.92347 "gene model 1960, (NCBI)" Gm1960
1438149_at 1.04856 Expressed sequence AI449441 AI449441
1438150_at 1.02566 farnesyl diphosphate synthetase /// similar to farnesyl diphosphate synthetase /// similar to farnesyl diphosphate synthetase Fdps /// LOC544723 /// LOC671145
1438151_x_at 0.91187 "zinc finger, DHHC domain containing 14" Zdhhc14
1438152_at 0.96674 GPI anchor attachment protein 1 Gpaal
1438153_x_at 0.86452 coenzyme Q10 homolog A (yeast) Coq10a
1438154_x_at 1.06871 RIKEN cDNA 2610002J02 gene 2610002J02Rik
1438155_x_at 0.90721 "phosphatidylinositol glycan anchor biosynthesis, class O" Pigo
1438156_x_at 0.85646 "carnitine palmitoyltransferase 1a, liver" Cpt1a
1438157_s_at 1.49977 "nuclear factor of kappa light chain gene enhancer in B-cells inhibitor, alpha" Nfkbia
1438158_at 0.96644 --- ---
1438159_x_at 0.93528 NADH dehydrogenase (ubiquinone) flavoprotein 2 Ndufv2
1438160_x_at 1.19866 "solute carrier organic anion transporter family, member 4a1" Slco4a1
1438161_s_at 0.95507 replication factor C (activator 1) 4 Rfc4
1438162_x_at 0.99788 hypothetical LOC231885 /// hypothetical protein LOC670882
LOC231885 /// LOC670882
1438163_x_at 0.90803 Rhomboid domain containing 2 Rhbdd2
1438164_x_at 0.87944 flotillin 2 Flot2
1438165_x_at 1.02573 vesicle amine transport protein 1 homolog (T californica) Vat1
1438166_x_at 0.88278 NADH dehydrogenase (ubiquinone) Fe-S protein 4 Ndufs4
1438167_x_at 1.05516 Folliculin Flcn
1438168_x_at 1.15406 DEAD (Asp-Glu-Ala-Asp) box polypeptide 39 Ddx39
1438169_a_at 0.84103 FERM domain containing 4B Frmd4b
1438170_x_at 0.97561 adhesion regulating molecule 1 Adrm1
1438171_x_at 1.02266 methyltransferase like 9 Mettl9
1438172_x_at 1.00372 exonuclease domain containing 1 Exod1
1438173_x_at 0.99821 Polyamine-modulated factor 1 Pmf1
1438174_x_at 0.92465 "protein phosphatase 2 (formerly 2A), regulatory subunit A (PR 65), alpha isoform" Ppp2r1a
1438175_x_at 1.87810 myomesin 2 Myom2
1438176_x_at 0.97741 RIKEN cDNA 1110031B06 gene 1110031B06Rik
1438177_x_at 0.98003 ectonucleoside triphosphate diphosphohydrolase 4 Entpd4
1438178_x_at 1.02841 "ATPase family, AAA domain containing 3A" Atad3a
1438179_s_at 0.99158 Signal transducer and activator of transcription interacting protein 1 Statip1

1438180_x_at	0.96921 silica-induced gene 111 /// HCLS1 associated X-1	Silg111 /// Hax1
1438181_x_at	0.89056 TM2 domain containing 2 Tm2d2	
1438182_x_at	0.94970 ankyrin repeat and SOCS box-containing protein 6	Asb6
1438183_x_at	0.94664 sorbitol dehydrogenase Sord	
1438184_a_at	1.00354 ankyrin repeat domain 5 Ankrd5	
1438185_at	1.05341 --- ---	
1438186_at	1.38813 PDZ and LIM domain 5 Pdlim5	
1438187_at	1.12052 "solute carrier family 25 (mitochondrial carrier, palmitoylcarnitine transporter), member 29"	Slc25a29
1438188_x_at	0.95078 "solute carrier family 25 (mitochondrial carrier, palmitoylcarnitine transporter), member 29"	Slc25a29
1438189_s_at	1.11560 erythrocyte protein band 4.9 Epb4.9	
1438190_x_at	1.03241 trophoblast specific protein alpha Tpbpa	
1438191_a_at	0.93783 ring finger protein 40 Rnf40	
1438192_s_at	1.10569 "bromodomain adjacent to zinc finger domain, 2A"	Baz2a
1438193_at	0.89314 neurexin III Nrnx3	
1438194_at	1.16625 RIKEN cDNA 2900019G14 gene	2900019G14Rik
1438195_at	0.84930 glycerol-3-phosphate dehydrogenase 1-like /// similar to glycerol-3-phosphate dehydrogenase 1-like Gpd11 /// LOC672508	
1438196_at	0.78447 glycerol-3-phosphate dehydrogenase 1-like	Gpd11
1438197_at	0.96551 "vezatin, adherens junctions transmembrane protein"	Vezt
1438198_at	0.95183 Bri3 binding protein Bri3bp	
1438199_at	0.87153 expressed sequence AI316807	AI316807
1438200_at	0.94512 sulfatase 1 Sulf1	
1438201_at	0.91795 "protein phosphatase 2C, magnesium dependent, catalytic subunit"	Ppm2c
1438202_at	1.05247 RIKEN cDNA C920005C14 gene	C920005C14Rik
1438203_at	0.90937 "scavenger receptor class F, member 2"	Scarf2
1438204_at	1.00584 "histone 1, H1c"	Hist1h1c
1438205_at	0.92768 RIKEN cDNA 1110014N23 gene	1110014N23Rik
1438206_a_at	0.96021 RIKEN cDNA 2610042O14 gene	2610042O14Rik
1438207_at	1.12337 golgi-specific brefeldin A-resistance factor 1 Gbf1	
1438208_at	0.85403 TAO kinase 2 Taok2	
1438209_at	1.03337 RIKEN cDNA 4632404H12 gene	4632404H12Rik
1438210_at	1.03011 G protein-coupled receptor 149 Gpr149	
1438211_s_at	0.65431 D site albumin promoter binding protein Dbp	
1438212_at	0.99095 ubiquitin specific peptidase 37 Usp37	
1438213_at	0.99012 RIKEN cDNA A830018L16 gene	A830018L16Rik
1438214_at	0.81699 trichorhinophalangeal syndrome I (human)	Trps1
1438215_at	0.88249 --- ---	
1438216_at	0.90971 ras responsive element binding protein 1	Rreb1
1438217_at	0.91811 ataxin 2 binding protein 1 A2bp1	
1438218_at	0.80870 zinc finger and BTB domain containing 24	Zbtb24
1438219_at	0.81295 Purine rich element binding protein A Pura	

1438220_at	1.18604 forkhead box J3 Foxj3	
1438221_at	1.14630 RIKEN cDNA C130065N10 gene	C130065N10Rik
1438222_at	2.24137 "DNA segment, Chr 2, ERATO Doi 612, expressed"	D2Ertd612e
1438223_at	1.07610 Vacuolar protein sorting 54 (yeast)	Vps54
1438224_at	0.83914 "zinc finger, SWIM domain containing 5"	Zswim5
1438225_x_at	1.05220 translocating chain-associating membrane protein 1	Tram1
1438226_at	1.04918 expressed sequence AU022252	AU022252
1438227_at	0.97900 src homology 2 domain-containing transforming protein E	She
1438228_at	0.94985 RIKEN cDNA 4930452B06 gene	4930452B06Rik
1438229_at	1.08882 "protein geranylgeranyltransferase type I, beta subunit"	Pggt1b
1438230_at	1.09690 "protein geranylgeranyltransferase type I, beta subunit"	Pggt1b
1438231_at	0.59932 forkhead box P2 Foxp2	
1438232_at	0.58125 forkhead box P2 Foxp2	
1438233_at	1.27119 RIKEN cDNA 6030443O07 gene	6030443O07Rik
1438234_at	1.93569 WD repeat domain 26	Wdr26
1438235_at	1.71056 --- ---	
1438236_at	0.70476 nuclear factor I/A	Nfia
1438237_at	0.95501 reduced expression 2	Rex2
1438238_at	0.93291 RIKEN cDNA 2010315B03 gene	2010315B03Rik
1438239_at	2.61001 midline 1	Mid1
1438240_at	0.92360 calicin Ccin	
1438241_at	0.85715 "RGM domain family, member A"	Rgma
1438242_at	0.90792 ubiquitin specific peptidase 40	Usp40
1438243_at	0.93310 --- ---	
1438244_at	2.03192 nuclear factor I/B	Nfib
1438245_at	1.84362 --- ---	
1438246_at	1.03156 "casein kinase 1, gamma 1"	Csnk1g1
1438247_at	0.78445 kelch-like 15 (Drosophila)	Klhl15
1438248_at	0.92615 proprotein convertase subtilisin/kexin type 5	Pcsk5
1438249_at	1.44225 Ubiquitin specific peptidase 7	Usp7
1438250_s_at	1.01464 "TAF9 RNA polymerase II, TATA box binding protein (TBP)-associated factor /// similar to TAF9 RNA polymerase II, TATA box binding protein (TBP)-associated factor isoform 1 /// similar to TAF9 RNA polymerase II, TATA box binding protein (TBP)-associated factor isoform 1"	Taf9 /// LOC545600 /// LOC671012
1438251_x_at	0.70637 HtrA serine peptidase 1	Htra1
1438252_at	1.05754 quiescin Q6	Qscn6
1438253_at	1.00756 slingshot homolog 1 (Drosophila)	Ssh1
1438254_at	1.08616 RIKEN cDNA 1110007A13 gene	1110007A13Rik
1438255_at	0.82002 checkpoint suppressor 1	Ches1
1438256_at	1.04666 eukaryotic translation initiation factor 5A2	Eif5a2
1438257_at	1.32672 Zinc finger protein 313	Zfp313
1438258_at	1.03544 very low density lipoprotein receptor	Vldlr
1438259_at	1.60389 --- ---	

1438260_at 0.90547 "potassium voltage-gated channel, subfamily Q, member 2" Kcnq2
 1438261_at 1.05506 "Cbp/p300-interacting transactivator, with Glu/Asp-rich carboxy-terminal domain, 4" Cited4
 1438262_at 0.98281 "solute carrier family 8 (sodium/calcium exchanger), member 2" Slc8a2
 1438263_at 0.82372 RIKEN cDNA 9430020K01 gene 9430020K01Rik
 1438264_a_at 1.10179 tripeptidyl peptidase II Tpp2
 1438265_at 1.71636 --- ---
 1438266_at 0.99692 ADAMTS-like 5 Adamtsl5
 1438267_x_at 1.02659 ribonucleic acid binding protein S1 Rnps1
 1438268_at 1.24482 membrane associated DNA binding protein Mnab
 1438269_at 1.27557 zinc finger and BTB domain containing 38 Zbtb38
 1438270_at 0.95605 expressed sequence AI846148 AI846148
 1438271_at 0.90097 LIM domain containing preferred translocation partner in lipoma Lpp
 1438272_at 0.89754 CUB and Sushi multiple domains 3 Csm3
 1438273_at 0.91070 --- ---
 1438274_at 1.03410 "zinc finger protein, subfamily 1A, 4 (Eos)" Zfpn1a4
 1438275_at 0.94044 RIKEN cDNA C130046K22 gene C130046K22Rik
 1438276_at 1.10666 Nuclear receptor co-repressor 1 Ncor1
 1438277_at 1.24233 RIKEN cDNA E130308A19 gene E130308A19Rik
 1438278_a_at 1.20193 cDNA sequence BC003993 BC003993
 1438279_at 0.91647 Dipeptidylpeptidase 4 Dpp4
 1438280_at 1.09641 RIKEN cDNA 1700023L04 gene 1700023L04Rik
 1438281_x_at 1.00698 RIKEN cDNA 1700023L04 gene 1700023L04Rik
 1438282_at 0.95162 synaptotagmin I Syt1
 1438283_at 1.06072 RIKEN cDNA 3110057O12 gene 3110057O12Rik
 1438284_at 1.16060 oculocerebrorenal syndrome of Lowe Ocl1
 1438285_at 0.98965 RIKEN cDNA 2210015D19 gene 2210015D19Rik
 1438286_at 1.03118 Thyroid hormone receptor associated protein 3 Thrap3
 1438287_x_at 1.06060 DEAD (Asp-Glu-Ala-Asp) box polypeptide 39 Ddx39
 1438288_x_at 0.97894 RIKEN cDNA 1110059G02 gene 1110059G02Rik
 1438289_a_at 1.00911 SMT3 suppressor of mif two 3 homolog 1 (yeast) Sumo1
 1438290_x_at 0.97368 Surfactant associated protein C Sftpc
 1438291_x_at 0.99526 ribosomal protein L37 /// similar to ribosomal protein L37 Rpl37 ///
 LOC383438
 1438292_x_at 1.02956 adenosine kinase Adk
 1438293_at 1.17888 "trichoplein, keratin filament binding" Tchp
 1438294_at 0.87713 ataxin 1 Atxn1
 1438295_at 0.98431 Glucocorticoid induced transcript 1 Glcci1
 1438296_at 1.07135 hypothetical protein C430010C01 C430010C01
 1438297_at 1.20716 EST AA545190 AA545190
 1438298_a_at 1.07584 --- ---
 1438299_at 1.34114 RIKEN cDNA 3100002L24 gene /// RIKEN cDNA 9230108I15 gene /// similar to gonadotropin inducible ovarian transcription factor 1 /// similar to zinc finger protein 709 /// similar to

zinc finger protein 709 /// similar to zinc finger protein 709 /// similar to zinc finger protein 97
3100002L24Rik /// 9230108I15Rik /// LOC626832 /// LOC628147 /// LOC665043 /// LOC665078 ///
LOC668039

1438300_at 0.94308 RIKEN cDNA 4921511I16 gene 4921511I16Rik
1438301_at 1.68897 Development and differentiation enhancing Ddef1
1438302_at 1.01141 HCF-binding transcription factor Zhangfei Zf
1438303_at 0.84353 "transforming growth factor, beta 2" Tgfb2
1438304_at 1.00680 CXXC finger 4 Cxxc4
1438305_at 0.92692 regulating synaptic membrane exocytosis 1 Rims1
1438306_at 1.02589 ring finger protein 180 Rnf180
1438307_at 0.94649 High mobility group box 2 Hmgb2
1438308_at 0.99097 RIKEN cDNA 2310026E23 gene 2310026E23Rik
1438309_at 0.99159 Transcribed locus ---
1438310_at 1.21145 "10 days neonate cerebellum cDNA, RIKEN full-length enriched library, clone:
B930095L19 product:unclassifiable, full insert sequence" ---
1438311_at 1.06305 RIKEN cDNA 2010002M12 gene 2010002M12Rik
1438312_s_at 0.71683 latent transforming growth factor beta binding protein 3 Ltbp3
1438313_at 0.99190 Transcribed locus ---
1438314_at 1.00588 PCTAIRE-motif protein kinase 1 Pctk1
1438315_x_at 0.92971 "aldo-keto reductase family 7, member A5 (aflatoxin aldehyde reductase)"
Akr7a5
1438316_a_at 0.94943 coiled-coil domain containing 102A /// similar to CG31638-PA Ccdc102a ///
LOC672218

1438317_a_at 1.05593 endonuclease G Endog
1438318_x_at 0.98715 "neuroguidin, EIF4E binding protein" Ngdn
1438319_x_at 0.95644 FAST kinase domains 2 Fastkd2
1438320_s_at 0.90853 minichromosome maintenance deficient 7 (S. cerevisiae) Mcm7
1438321_x_at 0.91082 RIKEN cDNA 4930504E06 gene 4930504E06Rik
1438322_x_at 1.35460 farnesyl diphosphate farnesyl transferase 1 Fdft1
1438323_at 0.90093 lens intrinsic membrane protein 2 Lim2
1438324_at 0.94691 RIKEN cDNA 9330182L06 gene 9330182L06Rik
1438325_at 0.80604 ecotropic viral integration site 1 Evi1
1438326_at 1.69108 RIKEN cDNA 3300001M20 gene 3300001M20Rik
1438327_at 1.05452 Zinc finger protein 533 Zfp533
1438328_at 1.01050 host cell factor C2 Hcfc2
1438329_at 0.89250 toll-like receptor 12 Tlr12
1438330_at 0.94468 cDNA sequence BC065397 BC065397
1438331_at 1.13324 Yippee-like 2 (Drosophila) Ypel2
1438332_at 1.10165 "solute carrier family 22 (organic anion transporter), member 6" Slc22a6
1438333_at 0.97518 protogenin homolog (Gallus gallus) Prtg
1438334_at 0.96519 Dedicator of cyto-kinesis 2 Dock2
1438335_at 1.13592 RIKEN cDNA 6030413G23 gene 6030413G23Rik
1438336_at 1.13183 F-box and WD-40 domain protein 11 Fbxw11

1438337_x_at 1.00138 RIKEN cDNA 9930032O22 gene 9930032O22Rik
1438338_at 1.10044 "malate dehydrogenase 1, NAD (soluble)" Mdh1
1438339_at 0.92378 "Fanconi anemia, complementation group D2" Fancd2
1438340_at 0.89602 RIKEN cDNA A930006D11Rik gene A930006D11Rik
1438341_at 0.95439 transglutaminase 4 (prostate) Tgm4
1438342_at 0.98009 RIKEN cDNA A930025D01 gene A930025D01Rik
1438343_at 0.97857 RIKEN cDNA 0610037L13 gene 0610037L13Rik
1438344_at 1.07463 RIKEN cDNA 4833424O15 gene 4833424O15Rik
1438345_at 1.83980 --- ---
1438346_at 0.96461 RIKEN cDNA 4930525G20 gene 4930525G20Rik
1438347_at 0.97350 Usher syndrome 3A homolog (human) Ush3a
1438348_x_at 1.07532 cysteine conjugate-beta lyase 1 Ccb1l
1438349_at 0.81314 cDNA sequence BC043476 BC043476
1438350_at 0.98445 G protein-coupled receptor 64 Gpr64
1438351_at 1.04868 "homeo box, msh-like 2" Msx2
1438352_at 0.99987 "Microtubule-associated protein, RP/EB family, member 2" Mapre2
1438353_at 0.87689 --- ---
1438354_x_at 0.89487 "Calponin 3, acidic" Cnn3
1438355_at 1.09304 transmembrane protein 90a Tmem90a
1438356_x_at 0.99959 RIKEN cDNA 4933432K03 gene 4933432K03Rik
1438357_at 1.11468 prefoldin 5 /// prefoldin 5 pseudogene Pfdn5 /// LOC434860
1438358_x_at 0.99348 prefoldin 5 /// prefoldin 5 pseudogene Pfdn5 /// LOC434860
1438359_at 1.04490 RIKEN cDNA 2010003K15 gene 2010003K15Rik
1438360_x_at 0.92314 "solute carrier family 25 (mitochondrial carrier, adenine nucleotide translocator), member 5 /// similar to ADP/ATP translocase 2 (Adenine nucleotide translocator 2) (ANT 2) (ADP,ATP carrier protein 2) (Solute carrier family 25 member 5) /// similar to ADP/ATP translocase 2 (Adenine nucleotide translocator 2) (ANT 2) (ADP,ATP carrier protein 2) (Solute carrier family 25 member 5) /// similar to ADP/ATP translocase 2 (Adenine nucleotide translocator 2) (ANT 2) (ADP,ATP carrier protein 2) (Solute carrier family 25 member 5) /// similar to solute carrier family 25, member A6" Slc25a5 /// LOC433326 /// LOC630624 /// LOC669381 /// LOC672115
1438361_at 0.98900 RIKEN cDNA 2310035C23 gene 2310035C23Rik
1438362_x_at 0.82406 RIKEN cDNA 2310035C23 gene 2310035C23Rik
1438363_at 0.93906 similar to KIAA1183 protein LOC434128
1438364_x_at 0.92667 "angiogenin, ribonuclease A family, member 4" Ang4
1438365_x_at 0.86910 lysosomal-associated protein transmembrane 4B Laptm4b
1438366_x_at 1.11447 chloride channel 3 Clcn3
1438367_x_at 0.94787 "discoidin domain receptor family, member 1" Ddr1
1438368_a_at 0.93515 matrin 3 Matr3
1438369_x_at 1.05638 transmembrane protein 85 Tmem85
1438370_x_at 0.88843 downstream of Stk11 Dos
1438371_x_at 0.95049 DEAD (Asp-Glu-Ala-Asp) box polypeptide 5 Ddx5
1438372_at 0.80542 myomesin 2 Myom2
1438373_at 1.05760 Amyloid beta (A4) precursor protein App

1438374_x_at 1.08807 amyloid beta (A4) precursor protein App
1438375_at 0.97490 Fibulin 2 Fbln2
1438376_s_at 0.94374 tripartite motif protein 27 Trim27
1438377_x_at 0.93015 "solute carrier family 13 (sodium-dependent dicarboxylate transporter), member 3" Slc13a3
1438378_at 0.84562 RIKEN cDNA 1700013G23 gene 1700013G23Rik
1438379_x_at 0.99357 RIKEN cDNA 2310007F21 gene 2310007F21Rik
1438380_at 0.99298 DEAD (Asp-Glu-Ala-Asp) box polypeptide 47 Ddx47
1438381_x_at 0.99580 DEAD (Asp-Glu-Ala-Asp) box polypeptide 47 Ddx47
1438382_x_at 0.93222 RIKEN cDNA 2010005J08 gene 2010005J08Rik
1438383_x_at 0.92244 "protein phosphatase 2 (formerly 2A), regulatory subunit A (PR 65), alpha isoform" Ppp2r1a
1438384_at 1.04162 RIKEN cDNA 1700047L15 gene 1700047L15Rik
1438385_s_at 1.01272 glutamic pyruvate transaminase (alanine aminotransferase) 2 Gpt2
1438386_x_at 0.96497 "methionine adenosyltransferase II, alpha" Mat2a
1438387_x_at 0.91421 topoisomerase (DNA) III beta Top3b
1438388_at 0.98696 Neurexophilin 1 Nxph1
1438389_x_at 1.11173 ATP citrate lyase Acly
1438390_s_at 0.98101 pituitary tumor-transforming 1 Pttg1
1438391_x_at 1.05153 hydroxyacyl-Coenzyme A dehydrogenase type II Hadh2
1438392_at 0.99335 CD247 antigen Cd247
1438393_at 0.94707 "In vitro fertilized eggs cDNA, RIKEN full-length enriched library, clone:7420416P19 product:unclassifiable, full insert sequence" ---
1438394_x_at 1.03267 keratin 4 Krt4
1438395_at 0.97998 --- ---
1438396_at 0.99128 oculocerebrorenal syndrome of Lowe Oclrl
1438397_a_at 1.04569 RNA binding motif protein 39 Rbm39
1438398_at 2.17065 "RNA-binding region (RNP1, RRM) containing 2" Rnpc2
1438399_at 0.99625 peroxin 2 Pex2
1438400_at 1.03859 RIKEN cDNA 4632411B12 gene 4632411B12Rik
1438401_at 1.02983 ubinuclein 1 Ubn1
1438402_at 0.96523 RIKEN cDNA 9630050M13 gene 9630050M13Rik
1438403_s_at 1.68933 --- ---
1438404_at 0.92159 ring finger protein 144 Rnf144
1438405_at 1.07116 fibroblast growth factor 7 Fgf7
1438406_at 0.93526 "scavenger receptor class F, member 2" Scarf2
1438407_at 0.95958 RIKEN cDNA 9330132E09 gene 9330132E09Rik
1438408_at 1.07782 ankyrin repeat domain 56 Ankrd56
1438409_at 1.08960 centrosomal protein 63 Cep63
1438410_at 0.96077 protogenin homolog (Gallus gallus) Prtg
1438411_at 1.18262 G protein-coupled receptor 81 Gpr81
1438412_at 0.70664 --- ---
1438413_at 0.76030 SUMO1/sentrin specific peptidase 7 Senp7

1438414_at 0.93195 fukutin related protein Fkrp
1438415_s_at 0.82266 "Yip1 domain family, member 2" Yipf2
1438416_at 0.95677 thyroid hormone receptor associated protein 5 Thrap5
1438417_at 1.02053 PWWP domain containing 2 Pwwp2
1438418_at 0.96522 RIKEN cDNA 4932432K03 gene 4932432K03Rik
1438419_at 1.07808 RNA binding motif protein 16 Rbm16
1438420_at 1.21252 RNA binding motif protein 39 Rbm39
1438421_at 0.85755 poliovirus receptor-related 1 Pvr11
1438422_at 0.94670 leucine rich repeat containing 20 Lrrc20
1438423_at 0.67516 single-stranded DNA binding protein 2 Ssbp2
1438424_at 0.83884 "TBC1 domain family, member 24" Tbc1d24
1438425_at 1.08629 general transcription factor III C 1 Gtf3c1
1438426_at 0.99098 transmembrane protein 58 Tmem58
1438427_at 1.02287 RIKEN cDNA 4932442K08 gene 4932442K08Rik
1438428_at 1.17951 Junctophilin 1 Jph1
1438429_at 1.61260 RIKEN cDNA 2610319H10 gene 2610319H10Rik
1438430_at 1.05348 high mobility group box transcription factor 1 Hbp1
1438431_at 0.92633 "ATP-binding cassette, sub-family D (ALD), member 2" Abcd2
1438432_at 1.03021 leucyl/cystinyl aminopeptidase Lnpep
1438433_at 1.29329 WAS protein homology region 2 domain containing 1 Whdc1
1438434_at 1.10871 Rho GTPase activating protein 11A Arhgap11a
1438435_at 0.89087 "phytoceramidase, alkaline" Phca
1438436_at 0.99365 RIKEN cDNA 6330407J23 gene 6330407J23Rik
1438437_a_at 0.98956 RIKEN cDNA 4933439C10 gene 4933439C10Rik
1438438_at 0.89669 RIKEN cDNA 2900002H16 gene 2900002H16Rik
1438439_at 0.95846 G protein-coupled receptor 171 Gpr171
1438440_at 0.96393 "ribonuclease III, nuclear" Rnasen
1438441_at 0.81619 Inhibitor of DNA binding 4 Id4
1438442_at 0.92750 expressed sequence AI450236 AI450236
1438443_at 1.03244 zinc finger and BTB domain containing 20 Zbtb20
1438444_at 0.90990 RIKEN cDNA A230091H23 gene A230091H23Rik
1438445_at 0.83087 "leucine rich repeat containing 8 family, member E" Lrrc8e
1438446_x_at 1.01803 putative phosphatase RP23-136K12.4
1438447_at 1.05743 "DEAD/H (Asp-Glu-Ala-Asp/His) box polypeptide 11 (CHL1-like helicase homolog, *S. cerevisiae*)" Ddx11
1438448_at 1.01662 otopetrin 1 Otop1
1438449_at 1.08779 Cdc42 binding protein kinase alpha Cdc42bpa
1438450_at 0.86857 lin-7 homolog A (*C. elegans*) Lin7a
1438451_at 0.81105 Rho GTPase-activating protein Grit
1438452_at 0.79003 nebulin Nebl
1438453_at 0.85847 Rad51 homolog c (*S. cerevisiae*) Rad51c
1438454_at 1.67994 RIKEN cDNA B430203M17 gene B430203M17Rik
1438455_at 0.99749 RIKEN cDNA C330050A14 gene C330050A14Rik

1438456_at 0.95097 histocompatibility 13 H13
1438457_at 0.95166 "Transcribed locus, strongly similar to XP_574375.1 PREDICTED: similar to MGC15476 protein [Rattus norvegicus]" ---
1438458_a_at 1.18160 splicing factor proline/glutamine rich (polypyrimidine tract binding protein associated) Sfpq
1438459_x_at 1.18957 splicing factor proline/glutamine rich (polypyrimidine tract binding protein associated) Sfpq
1438460_at 1.26484 "zinc finger, MYM-type 4" Zmym4
1438461_at 1.00922 --- ---
1438462_x_at 0.98185 --- ---
1438463_x_at 0.90970 "zinc finger, DHHC domain containing 6" Zdhhc6
1438464_at 1.07393 AT rich interactive domain 4A (Rbp1 like) Arid4a
1438465_at 1.08667 RIKEN cDNA 5830428H23 gene 5830428H23Rik
1438466_at 0.78827 "dynein, axonemal, heavy chain 12 /// dynein, axonemal, heavy chain 7b /// dynein heavy chain-related" Dnahc12 /// Dnahc7b /// LOC433299
1438467_at 0.94165 macrophage galactose N-acetyl-galactosamine specific lectin 2 Mgl2
1438468_at 1.02273 preferentially expressed antigen in melanoma like 4 /// preferentially expressed antigen in melanoma like 5 Pramel4 /// Pramel5
1438469_at 0.87799 BCL6 co-repressor-like 1 Bcorl1
1438470_at 0.61346 suppressor of cytokine signaling 2 Socs2
1438471_at 1.12320 Wilms' tumour 1-associating protein Wtap
1438472_at 0.86274 --- ---
1438473_at 1.03186 ADP-ribosylation factor-like 15 Arl15
1438474_at 1.06019 ankyrin repeat domain 35 Ankrd35
1438475_at 1.02287 cDNA sequence AB124611 AB124611
1438476_a_at 0.94781 chromodomain helicase DNA binding protein 4 Chd4
1438477_a_at 0.81933 methylmalonyl CoA epimerase Mcee
1438478_a_at 0.92583 "protein phosphatase 3, catalytic subunit, alpha isoform" Ppp3ca
1438479_at 0.98644 zinc finger protein 213 Zfp213
1438480_a_at 1.01700 thymocyte nuclear protein 1 Thyn1
1438481_at 1.05231 PRP4 pre-mRNA processing factor 4 homolog (yeast) Prpf4
1438482_at 1.20281 WW domain containing E3 ubiquitin protein ligase 2 Wwp2
1438483_at 1.06937 "nitric oxide synthase 1, neuronal" Nos1
1438484_at 0.90610 "gene model 237, (NCBI)" Gm237
1438485_at 0.92739 RIKEN cDNA E130112L23 gene E130112L23Rik
1438486_at 1.13379 testis-specific serine kinase 4 Tssk4
1438487_s_at 0.87354 "zinc finger, ZZ domain containing 3" Zzz3
1438488_at 1.00281 esterase D/formylglutathione hydrolase Esd
1438489_at 1.17419 --- ---
1438490_at 1.01210 "solute carrier family 39 (zinc transporter), member 14" Slc39a14
1438491_x_at 0.91444 RIKEN cDNA A530054K11 gene A530054K11Rik
1438492_at 0.88171 Suppressor of cytokine signaling 7 Socs7
1438493_at 0.95839 RIKEN cDNA 4933433K01 gene 4933433K01Rik

1438494_at 1.01222 histamine receptor H 1 Hrh1
1438495_at 1.57825 topoisomerase (DNA) I Top1
1438496_a_at 0.88914 RIKEN cDNA 6330505F04 gene 6330505F04Rik
1438497_at 1.05773 RIKEN cDNA 2810423E13 gene 2810423E13Rik
1438498_at 1.15473 "zinc finger, MYND-type containing 15" Zmynd15
1438499_at 1.02907 RIKEN cDNA 1700029I15 gene 1700029I15Rik
1438500_at 0.98534 proline-rich transmembrane protein 3 Prrt3
1438501_at 0.99887 Ribosomal protein S17 Rps17
1438502_x_at 1.03112 ribosomal protein S17 /// similar to 40S ribosomal protein S17 Rps17 ///
LOC383032
1438503_x_at 1.04897 endothelial differentiation-related factor 1 /// similar to Endothelial
differentiation-related factor 1 (EDF-1) (Multiprotein bridging factor 1) (MBF1) (Calmodulin-associated
peptide 19) (CAP-19) /// similar to Endothelial differentiation-related factor 1 (EDF-1) (Multiprotein
bridging factor 1) (MBF1) (Calmodulin-associated peptide 19) (CAP-19) /// similar to Endothelial
differentiation-related factor 1 (EDF-1) (Multiprotein bridging factor 1) (MBF1) (Calmodulin-associated
peptide 19) (CAP-19) Edf1 /// LOC665181 /// LOC671865 /// LOC674653
1438504_x_at 0.96813 Transmembrane 7 superfamily member 3 Tm7sf3
1438505_s_at 1.00801 "ribonuclease III, nuclear" Rnasen
1438506_s_at 1.10475 "Prenyl (solanesyl) diphosphate synthase, subunit 1" Pdss1
1438507_x_at 1.11281 ribosomal protein L14 Rpl14
1438508_at 1.01464 Jumonji domain containing 1B Jmjd1b
1438509_at 0.93260 "Proteasome (prosome, macropain) 28 subunit, 3" Psme3
1438510_a_at 0.97225 histidyl-tRNA synthetase Hars
1438511_a_at 1.03475 RIKEN cDNA 1190002H23 gene 1190002H23Rik
1438512_at 1.02426 cDNA sequence BC048679 BC048679
1438513_at 1.77402 TatD DNase domain containing 1 Tatdn1
1438514_at 1.09911 NPC1-like 1 Npc11
1438515_at 1.58248 zinc finger protein 207 Zfp207
1438516_at 1.89683 Rap1 interacting factor 1 homolog (yeast) Rif1
1438517_at 0.94560 WW domain-containing oxidoreductase Wwox
1438518_at 0.89538 RIKEN cDNA 9930021J03 gene 9930021J03Rik
1438519_at 2.07292 kelch-like 24 (Drosophila) Khl24
1438520_at 0.96571 "solute carrier family 25, member 36" Slc25a36
1438521_at 0.90197 Osteopetrosis associated transmembrane protein 1 Ostm1
1438522_at 1.12865 RIKEN cDNA 1700054H16 gene 1700054H16Rik
1438523_x_at 1.00982 RIKEN cDNA 1500001A10 gene 1500001A10Rik
1438524_x_at 0.89551 proline-rich nuclear receptor coactivator 1 Pnrc1
1438525_at 1.08029 Myeloid/lymphoid or mixed lineage-leukemia translocation to 10 homolog
(Drosophila) Mllt10
1438526_at 0.95923 G protein-coupled receptor 158 Gpr158
1438527_at 1.50561 ribosomal protein L3 /// similar to 60S ribosomal protein L3 (J1 protein) ///
similar to 60S ribosomal protein L3 (L4) /// similar to 60S ribosomal protein L3 (J1 protein) /// similar to
60S ribosomal protein L3 (L4) /// similar to ribosomal protein L3 isoform b /// similar to 60S ribosomal

protein L3 (J1 protein) /// similar to 60S ribosomal protein L3 (J1 protein) /// similar to 60S ribosomal
 protein L3 (J1 protein) /// similar to 60S ribosomal protein L3 (J1 protein) /// similar to ribosomal
 protein L3 isoform b Rpl3 /// LOC242809 /// LOC433745 /// LOC545864 /// LOC635340 ///
 LOC668219 /// LOC669672 /// LOC670435 /// LOC670753 /// LOC674810 /// LOC674874
 1438528_at 2.30925 Pericentriolar material 1 Pcm1
 1438529_at 0.98861 "DNA segment, Chr 4, ERATO Doi 429, expressed" D4Ertd429e
 1438530_at 1.05977 tissue factor pathway inhibitor Tfpi
 1438531_at 0.79977 RIKEN cDNA A730054J21 gene A730054J21Rik
 1438532_at 1.03457 hemicentin 1 Hmcn1
 1438533_at 1.30258 myosin IXb Myo9b
 1438534_x_at 0.99319 cDNA sequence BC004004 BC004004
 1438535_at 1.73479 pleckstrin homology domain interacting protein Phip
 1438536_at 1.01504 RIKEN cDNA 5830411N06 gene 5830411N06Rik
 1438537_at 1.15932 --- ---
 1438538_at 1.13416 RIKEN cDNA D630021H01 gene D630021H01Rik
 1438539_at 1.06906 Activating signal cointegrator 1 complex subunit 1 Ascc1
 1438540_at 1.02323 --- ---
 1438541_at 1.06161 RIKEN cDNA 4933439F18 gene 4933439F18Rik
 1438542_at 1.68033 --- ---
 1438543_at 1.10611 Spermatogenesis associated 13 Spata13
 1438544_at 0.83169 RIKEN cDNA 1700001G11 gene 1700001G11Rik
 1438545_at 1.13775 "solute carrier family 25 (mitochondrial carrier, adenine nucleotide
 translocator), member 5 /// similar to ADP/ATP translocase 2 (Adenine nucleotide translocator 2) (ANT
 2) (ADP,ATP carrier protein 2) (Solute carrier family 25 member 5) /// similar to ADP/ATP translocase
 2 (Adenine nucleotide translocator 2) (ANT 2) (ADP,ATP carrier protein 2) (Solute carrier family 25
 member 5) /// similar to solute carrier family 25, member 5 /// similar to ADP/ATP translocase 2
 (Adenine nucleotide translocator 2) (ANT 2) (ADP,ATP carrier protein 2) (Solute carrier family 25
 member 5) /// similar to solute carrier family 25, member 5 /// similar to ADP/ATP translocase 2
 (Adenine nucleotide translocator 2) (ANT 2) (ADP,ATP carrier protein 2) (Solute carrier family 25
 member 5) /// similar to solute carrier family 25, member 5" Slc25a5 /// LOC433326 /// LOC433923 ///
 LOC623402 /// LOC630624 /// LOC633283 /// LOC669381 /// LOC670371
 1438546_x_at 1.08527 "solute carrier family 25 (mitochondrial carrier, adenine nucleotide
 translocator), member 5 /// similar to ADP/ATP translocase 2 (Adenine nucleotide translocator 2) (ANT
 2) (ADP,ATP carrier protein 2) (Solute carrier family 25 member 5) /// similar to ADP/ATP translocase
 2 (Adenine nucleotide translocator 2) (ANT 2) (ADP,ATP carrier protein 2) (Solute carrier family 25
 member 5) /// similar to solute carrier family 25, member 5 /// similar to ADP/ATP translocase 2
 (Adenine nucleotide translocator 2) (ANT 2) (ADP,ATP carrier protein 2) (Solute carrier family 25
 member 5) /// similar to solute carrier family 25, member 5 /// similar to ADP/ATP translocase 2
 (Adenine nucleotide translocator 2) (ANT 2) (ADP,ATP carrier protein 2) (Solute carrier family 25
 member 5) /// similar to solute carrier family 25, member 5" Slc25a5 /// LOC433326 /// LOC433923 ///
 LOC623402 /// LOC630624 /// LOC633283 /// LOC669381 /// LOC670371
 1438547_x_at 1.00316 "torsin family 2, member A" Tor2a
 1438548_x_at 0.91200 rhomboid domain containing 3 Rhbdd3

1438549_a_at 0.76045 serine racemase Srr
1438550_x_at 0.97794 serine racemase Srr
1438551_at 1.07494 neurogenin 1 Neurog1
1438552_x_at 0.86509 zyxin Zyx
1438553_x_at 0.91674 RIKEN cDNA 4930453N24 gene 4930453N24Rik
1438554_x_at 0.94658 Williams-Beuren syndrome chromosome region 1 homolog (human) Wbscr1
1438555_x_at 0.82547 RIKEN cDNA 4933405I11 gene /// mucin 4 4933405I11Rik /// Muc4
1438556_a_at 0.84618 tropomodulin 3 Tmod3
1438557_x_at 0.84927 aspartyl aminopeptidase Dnpep
1438558_x_at 0.74588 Forkhead box Q1 Foxq1
1438559_x_at 0.87136 "solute carrier family 44, member 2" Slc44a2
1438560_x_at 1.01878 chaperonin subunit 4 (delta) Cct4
1438561_x_at 0.97317 RIKEN cDNA 4930538D17 gene 4930538D17Rik
1438562_a_at 1.15992 "protein tyrosine phosphatase, non-receptor type 2" Ptpn2
1438563_s_at 0.96052 mitochondrial ribosomal protein S24 Mrps24
1438564_at 1.16666 Growth arrest specific 2 Gas2
1438565_at 1.07625 RIKEN cDNA A830010M20 gene A830010M20Rik
1438566_at 1.04104 "ST8 alpha-N-acetyl-neuraminide alpha-2,8-sialyltransferase 6" St8sia6
1438567_at 1.04027 von Willebrand factor A domain containing 2 Vwa2
1438568_at 0.81800 "MAS-related GPR, member E" Mrgpre
1438569_at 0.98217 coiled-coil domain containing 38 Ccdc38
1438570_at 0.96070 Transcribed locus ---
1438571_at 1.06608 Budding uninhibited by benzimidazoles 1 homolog (S. cerevisiae) Bub1
1438572_at 0.92924 CUB and Sushi multiple domains 3 Csm3
1438573_at 0.94841 "RMI1, RecQ mediated genome instability 1, homolog (S. cerevisiae)" Rmi1
1438574_at 0.87344 G protein-coupled receptor 152 Gpr152
1438575_a_at 0.96348 RIKEN cDNA 2900056M20 gene 2900056M20Rik
1438576_x_at 0.88169 RIKEN cDNA 2900056M20 gene 2900056M20Rik
1438577_at 0.94772 Transcribed locus ---
1438578_a_at 0.98545 BTB (POZ) domain containing 10 Btbd10
1438579_at 1.13980 "UTP14, U3 small nucleolar ribonucleoprotein, homolog B (yeast)" Utp14b
1438580_at 1.38915 "zinc finger, CCHC domain containing 7" Zcchc7
1438581_at 1.50588 RIKEN cDNA 4932439K10 gene 4932439K10Rik
1438582_at 0.92013 ARP5 actin-related protein 5 homolog (yeast) Actr5
1438583_at 0.96315 Endoplasmic reticulum (ER) to nucleus signalling 1 Ern1
1438584_at 0.94169 RIKEN cDNA 4932411G14 gene 4932411G14Rik
1438585_at 1.00603 synaptic nuclear envelope 2 Syne2
1438586_at 0.90048 T-box 22 Tbx22
1438587_at 0.96985 RIKEN cDNA 4933417M04 gene 4933417M04Rik
1438588_at 1.05052 Pleiomorphic adenoma gene-like 1 Plagl1
1438589_at 1.06080 "immunoglobulin superfamily, member 1" Igsf1
1438590_at 1.45002 Rap guanine nucleotide exchange factor (GEF) 3 Rapgef3
1438591_at 1.25036 RIKEN cDNA 2610019A05 gene 2610019A05Rik

1438592_at 1.05780 NIMA (never in mitosis gene a)-related expressed kinase 1 Nek1
1438593_at 1.01558 hypothetical LOC622461 LOC622461
1438594_at 0.95344 "PREDICTED: Mus musculus similar to poly(A) binding protein, cytoplasmic 1 (LOC245545), mRNA" ---
1438595_at 1.03112 "gene model 546, (NCBI)" Gm546
1438596_at 1.07030 HECT domain containing 2 Hectd2
1438597_x_at 0.94054 mortality factor 4 like 1 /// similar to mortality factor 4 like 1 isoform b /// similar to mortality factor 4 like 1 /// similar to mortality factor 4 like 1 (predicted) Morf411 /// LOC433955 /// LOC627352 /// LOC635663
1438598_at 0.88072 src homology 2 domain-containing transforming protein C2 Shc2
1438599_at 0.96218 --- ---
1438600_at 0.96422 similar to Cyclin-dependent kinases regulatory subunit 1 (CKS-1) (Sid 1334) LOC624855
1438601_at 1.15990 "protein kinase, membrane associated tyrosine/threonine 1" Pkmyt1
1438602_s_at 0.66039 mannan-binding lectin serine peptidase 1 Masp1
1438603_x_at 0.63224 Mannan-binding lectin serine peptidase 1 Masp1
1438604_at 1.01081 "ATPase, Class I, type 8B, member 3" Atp8b3
1438605_at 0.88664 Cell division cycle 26 Cdc26
1438606_a_at 0.95120 chloride intracellular channel 4 (mitochondrial) Clic4
1438607_at 0.98497 "zinc finger, DHHC domain containing 19" Zdhhc19
1438608_at 1.00638 "Troponin I, skeletal, fast 2" Tnni2
1438609_x_at 0.92904 "troponin I, skeletal, fast 2" Tnni2
1438610_a_at 0.86222 "Crystallin, zeta" Cryz
1438611_at 0.95981 "Tubulin, alpha 3" Tuba3
1438612_a_at 1.02896 "colipase, pancreatic" Clps
1438613_at 1.03928 "potassium voltage-gated channel, shaker-related subfamily, member 4" Kcna4
1438614_x_at 1.12814 Oxysterol binding protein-like 9 Osbp19
1438615_x_at 0.97978 RIKEN cDNA 2010317E24 gene 2010317E24Rik
1438616_x_at 0.96404 seminal vesicle secretion 5 Svs5
1438617_at 0.91478 "Serine (or cysteine) peptidase inhibitor, clade A (alpha-1 antiproteinase, antitrypsin), member 7" Serpina7
1438618_at 1.07794 expressed sequence AI507597 AI507597
1438619_x_at 0.94470 "zinc finger, DHHC domain containing 14" Zdhhc14
1438620_x_at 1.02498 secreted frizzled-related sequence protein 1 Sfrp1
1438621_x_at 1.06969 AXL receptor tyrosine kinase Axl
1438622_x_at 1.05496 "tubulin, beta 2c" Tubb2c
1438623_x_at 0.96287 ring-box 1 Rbx1
1438624_x_at 1.05723 heparan sulfate (glucosamine) 3-O-sulfotransferase 2 Hs3st2
1438625_s_at 0.93699 necdin /// PCTAIRE-motif protein kinase 1 Ndn /// Pctk1
1438626_x_at 1.11665 ribosomal protein L14 Rpl14
1438627_x_at 1.22024 phosphogluconate dehydrogenase Pgd
1438628_x_at 1.05469 contactin 3 Cntn3
1438629_x_at 0.93983 granulin Grn

1438630_x_at 0.98431 "methionine adenosyltransferase II, alpha" Mat2a
1438631_x_at 0.89332 tetratricopeptide repeat domain 13 Ttc13
1438632_x_at 0.93188 transition protein 1 Tnp1
1438633_x_at 1.00734 LIM and SH3 protein 1 Lasp1
1438634_x_at 1.09674 LIM and SH3 protein 1 Lasp1
1438635_x_at 1.04266 RIKEN cDNA B930041F14 gene B930041F14Rik
1438636_s_at 1.01007 sushi domain containing 5 Susd5
1438637_x_at 1.02665 "splicing factor 3b, subunit 2" Sf3b2
1438638_x_at 0.91722 RIKEN cDNA 1700027J05 gene 1700027J05Rik
1438639_x_at 1.09109 RIKEN cDNA 4933417E01 gene 4933417E01Rik
1438640_x_at 1.19893 phosphoglycerate kinase 1 /// phosphoglycerate kinase 1-like ///
phosphoglycerate kinase 1 pseudogene /// phosphoglycerate kinase 1-like Pgk1 /// LOC433594 ///
LOC665053 /// LOC668435
1438641_x_at 0.96148 RIKEN cDNA 1500016O10 gene 1500016O10Rik
1438642_at 1.03368 --- ---
1438643_at 1.19658 Calcium/calmodulin-dependent protein kinase ID Camk1d
1438644_x_at 0.90900 COMM domain containing 9 Commd9
1438645_x_at 1.06803 beta-site APP-cleaving enzyme 2 Bace2
1438646_x_at 0.93635 RIKEN cDNA 2510039O18 gene 2510039O18Rik
1438647_x_at 0.91420 centrin 2 Cctn2
1438648_x_at 0.97375 RIKEN cDNA 1190003M12 gene 1190003M12Rik
1438649_x_at 0.91752 phosphatidylethanolamine binding protein 1 Pebp1
1438650_x_at 0.96897 gap junction membrane channel protein alpha 1 Gja1
1438651_a_at 0.93821 angiotensin receptor-like 1 Agtrl1
1438652_x_at 0.91567 "phosphatidylinositol glycan anchor biosynthesis, class Q" Pigq
1438653_x_at 0.92561 ataxin 10 Atn10
1438654_x_at 1.00744 monocyte to macrophage differentiation-associated 2 Mmd2
1438655_a_at 1.09604 RIKEN cDNA 1100001I22 gene 1100001I22Rik
1438656_x_at 1.01093 translocase of inner mitochondrial membrane 17b Timm17b
1438657_x_at 0.96102 protein tyrosine phosphatase 4a1 /// protein tyrosine phosphatase 4a1-like ///
protein tyrosine phosphatase 4a1 pseudogene /// protein tyrosine phosphatase 4a1-like /// similar to
protein tyrosine phosphatase 4a1 Ptp4a1 /// LOC433406 /// LOC627166 /// LOC627200 ///
LOC667723
1438658_a_at 0.93812 "endothelial differentiation, sphingolipid G-protein-coupled receptor, 3"
Edg3
1438659_x_at 0.99180 coiled-coil-helix-coiled-coil-helix domain containing 6 Chchd6
1438660_at 1.39471 "glucosaminyl (N-acetyl) transferase 2, I-branching enzyme" Gcnt2
1438661_a_at 0.74766 ADP-ribosylation factor 2 Arf2
1438662_at 0.94361 similar to transmembrane protein SHREW1 LOC433810
1438663_at 1.12561 BAT2 domain containing 1 Bat2d
1438664_at 1.01322 "protein kinase, cAMP dependent regulatory, type II beta" Prkar2b
1438665_at 1.09491 "sphingomyelin phosphodiesterase 3, neutral" Smpd3
1438666_at 0.76655 low density lipoprotein receptor class A domain containing 3 Ldlrad3

1438667_at 0.98224 RIKEN cDNA 5730410E15 gene 5730410E15Rik
1438668_x_at 1.06688 ataxin 2-like Atxn2l
1438669_at 0.91314 WD repeat domain 40A Wdr40a
1438670_at 0.97055 "protein tyrosine phosphatase, non-receptor type 1" Ptpn1
1438671_at 1.01606 "protein phosphatase 2 (formerly 2A), regulatory subunit B (PR 52), gamma isoform" Ppp2r2c
1438672_at 0.96183 "parvin, beta" Parvb
1438673_at 1.10510 --- ---
1438674_a_at 1.20603 "splicing factor, arginine/serine-rich 8" Sfrs8
1438675_at 1.75105 "splicing factor, arginine/serine-rich 8" Sfrs8
1438676_at 1.68641 macrophage activation 2 like /// similar to macrophage activation 2 like /// similar to macrophage activation 2 like Mpa2l /// LOC626578 /// LOC673101
1438677_at 0.89823 plakophilin 4 Pkp4
1438678_at 0.89291 RIKEN cDNA 1500011K16 gene 1500011K16Rik
1438679_at 0.98612 Tripartite motif protein 8 Trim8
1438680_at 0.94408 autism susceptibility candidate 2 Aut2
1438681_at 0.90650 "alkB, alkylation repair homolog 1 (E. coli)" Alkbl1
1438682_at 0.93692 "phosphatidylinositol 3-kinase, regulatory subunit, polypeptide 1 (p85 alpha)" Pik3r1
1438683_at 0.98546 "WAS protein family, member 2" Wasf2
1438684_at 0.82714 "NUAK family, SNF1-like kinase, 1" Nuak1
1438685_at 0.91001 "zinc finger, MYM-type 6" Zmym6
1438686_at 1.00580 "eukaryotic translation initiation factor 4, gamma 1" Eif4g1
1438687_at 1.03002 lymphocyte transmembrane adaptor 1 Lax1
1438688_at 1.13240 serine/arginine repetitive matrix 2 Srrm2
1438689_at 0.92777 RIKEN cDNA C730049F21 gene C730049F21Rik
1438690_at 1.83250 Thymidylate synthase Tysm
1438691_at 0.86827 "zinc finger, ZZ-type with EF hand domain 1" Zzef1
1438692_at 1.05421 "general transcription factor IIIC, polypeptide 4" Gtf3c4
1438693_at 0.76109 transmembrane protein 110 Tmem110
1438694_at 1.08323 casein alpha s2-like A Csn1s2a
1438695_at 0.85321 RIKEN cDNA C230091D08 gene C230091D08Rik
1438696_at 0.98151 endothelin 3 Edn3
1438697_at 1.00694 RIKEN cDNA 4632425D07 gene 4632425D07Rik
1438698_at 0.98880 RIKEN cDNA 4632425D07 gene 4632425D07Rik
1438699_at 0.96448 steroid 5 alpha-reductase 1 Srd5a1
1438700_at 1.88254 formin binding protein 4 Fnbp4
1438701_at 1.17503 bicaudal D homolog 1 (Drosophila) Bicd1
1438702_at 1.04553 --- ---
1438703_at 0.91954 ankyrin repeat domain 26 Ankrd26
1438704_at 0.91812 tripartite motif-containing 63 Trim63
1438705_at 0.69472 expressed sequence AI465270 AI465270
1438706_at 0.92060 RIKEN cDNA 6430517E21 gene 6430517E21Rik

1438707_at 0.91670 ATPase type 13A4 Atp13a4
1438708_x_at 0.90300 "tyrosine 3-monooxygenase/tryptophan 5-monooxygenase activation protein, beta polypeptide" Ywhab
1438709_at 0.98249 "WD repeat domain, phosphoinositide interacting 1" Wipi1
1438710_at 0.96259 5-hydroxytryptamine (serotonin) receptor 1A Htr1a
1438711_at 0.92712 pyruvate kinase liver and red blood cell Pklr
1438712_at 1.08068 DENN/MADD domain containing 2D Dennd2d
1438713_at 0.79749 Ras association (RalGDS/AF-6) domain family 8 Rassf8
1438714_at 1.39219 zinc finger protein 207 Zfp207
1438715_at 0.86138 coiled-coil domain containing 102A Ccdc102a
1438716_at 1.01554 expressed sequence AI451617 AI451617
1438717_a_at 1.00829 oxysterol binding protein-like 6 Osbp16
1438718_at 1.08782 fibroblast growth factor 9 Fgf9
1438719_at 1.00116 expressed sequence AI585793 AI585793
1438720_at 0.92776 RIKEN cDNA 9330159F19 gene 9330159F19Rik
1438721_a_at 1.23472 interferon regulatory factor 3 Irf3
1438722_at 0.84930 RIKEN cDNA 2610014I16 gene 2610014I16Rik
1438723_a_at 0.99890 ribosomal protein S10 /// similar to ribosomal protein S10 /// similar to ribosomal protein S10 Rps10 /// LOC636539 /// LOC666274
1438724_at 1.08099 oxysterol binding protein-like 3 Osbp13
1438725_at 1.09381 thyroid hormone receptor associated protein 1 Thrapl1
1438726_at 0.86578 "microtubule associated monooxygenase, calponin and LIM domain containing 2" Mical2
1438727_at 0.89922 WD repeat domain 32 Wdr32
1438728_at 0.90794 Ngg1 interacting factor 3-like 1 (S. pombe) Nif311
1438729_at 1.01546 Expressed sequence BB176347 BB176347
1438730_at 1.00435 cDNA sequence BC028801 BC028801
1438731_at 0.99093 --- ---
1438732_at 0.99958 hypothetical protein 6430511F03 6430511F03
1438733_at 0.96120 zinc finger protein 689 Zfp689
1438734_at 1.03740 RIKEN cDNA A530016L24 gene A530016L24Rik
1438735_at 0.96061 remodeling and spacing factor 1 Rsf1
1438736_at 1.10471 THO complex 2 Thoc2
1438737_at 1.00670 zinc finger protein of the cerebellum 3 Zic3
1438738_at 1.16816 CDNA sequence BC010584 BC010584
1438739_at 2.15942 cellular nucleic acid binding protein Cnbp
1438740_at 0.97578 N-myristoyltransferase 2 Nmt2
1438741_at 0.88409 cDNA sequence BC019943 BC019943
1438742_at 0.82641 zinc finger protein 629 Zfp629
1438743_at 1.07578 "cytochrome P450, family 7, subfamily a, polypeptide 1" Cyp7a1
1438744_at 0.91331 ankyrin repeat and SOCS box-containing protein 7 Asb7
1438745_at 0.97610 hypothetical protein LOC666945 LOC666945
1438746_at 0.99154 RIKEN cDNA A530058N18 gene A530058N18Rik

1438747_at 0.94845 RIKEN cDNA A930006K02 gene A930006K02Rik
 1438748_at 1.22380 RIKEN cDNA 9830127L17 gene 9830127L17Rik
 1438749_at 1.07574 fuzzy homolog (Drosophila) Fuz
 1438750_at 0.98913 Alpha thalassemia/mental retardation syndrome X-linked homolog (human)
 Atrx
 1438751_at 1.04175 "solute carrier family 30, member 10" Slc30a10
 1438752_at 1.09927 RIKEN cDNA A230058F20 gene A230058F20Rik
 1438753_at 0.96381 Porcupine homolog (Drosophila) Porcn
 1438754_at 0.98774 --- ---
 1438755_at 1.06074 expressed sequence C80068 C80068
 1438756_at 0.92588 ankyrin repeat domain 29 Ankrd29
 1438757_at 1.11238 RIKEN cDNA C130069I09 gene C130069I09Rik
 1438758_at 0.90988 --- ---
 1438759_x_at 0.94972 --- ---
 1438760_x_at 0.94209 a disintegrin and metallopeptidase domain 15 (metargidin) Adam15
 1438761_a_at 1.12430 "ornithine decarboxylase, structural 1 /// similar to Ornithine decarboxylase
 (ODC) /// similar to Ornithine decarboxylase (ODC) /// similar to Ornithine decarboxylase (ODC) ///
 hypothetical protein LOC671973 /// similar to Ornithine decarboxylase (ODC)" Odc1 ///
 LOC666231 /// LOC668343 /// LOC670279 /// LOC671973 /// LOC673302
 1438762_at 1.43252 Transcription factor 12 Tcf12
 1438763_at 0.83428 "dynein, axonemal, heavy chain 2 /// dynein heavy chain domain 3"
 Dnahc2 /// Dnhd3
 1438764_at 1.00264 annexin A7 Anxa7
 1438765_at 1.07453 DEAH (Asp-Glu-Ala-His) box polypeptide 33 Dhx33
 1438766_at 1.32621 proline-rich nuclear receptor coactivator 2 Pnrc2
 1438767_at 0.99501 oncostatin M Osm
 1438768_at 1.00321 hypothetical protein LOC665225 /// hypothetical protein LOC669601 ///
 hypothetical protein LOC676250 LOC665225 /// LOC669601 /// LOC676250
 1438769_a_at 1.04799 thymocyte nuclear protein 1 Thyn1
 1438770_at 1.08568 "gene model 266, (NCBI)" Gm266
 1438771_at 1.09224 bromodomain containing 1 Brd1
 1438772_at 0.97077 zinc finger protein 367 Zfp367
 1438773_at 1.08246 six transmembrane epithelial antigen of prostate 2 Steap2
 1438774_s_at 0.88523 phosphoglucomutase 2-like 1 Pgm2l1
 1438775_at 1.09821 "Peroxisome proliferative activated receptor, gamma, coactivator 1 beta"
 Ppargc1b
 1438776_x_at 1.02881 ribosomal protein S17 /// similar to ribosomal protein S17 /// similar to 40S
 ribosomal protein S17 Rps17 /// LOC268650 /// LOC383032
 1438777_a_at 1.12216 EST AA881470 AA881470
 1438778_at 1.03372 "Adult male diencephalon cDNA, RIKEN full-length enriched library,
 clone:9330183L06 product:unclassifiable, full insert sequence" ---
 1438779_at 0.47395 "procollagen, type IV, alpha 3" Col4a3
 1438780_at 1.08829 --- ---

1438781_at 2.32391 RIKEN cDNA E130014J05 gene E130014J05Rik
 1438782_at 0.99096 contactin 4 Cntn4
 1438783_at 0.98816 similar to Nedd4 WW binding protein 4 LOC637870
 1438784_at 1.10172 --- ---
 1438785_at 1.02685 ectonucleotide pyrophosphatase/phosphodiesterase 6 Enpp6
 1438786_a_at 1.09222 RIKEN cDNA 2610021A01 gene /// similar to zinc finger protein 347
 2610021A01Rik /// LOC668572
 1438787_at 0.91493 zinc finger protein 206 Zfp206
 1438788_at 2.49853 "DNA segment, Chr 5, Wayne State University 152, expressed" D5Wsu152e
 1438789_s_at 0.86009 dihydropyrimidinase-like 3 Dpysl3
 1438790_x_at 0.94909 transmembrane protein 41B Tmem41b
 1438791_at 0.97781 RIKEN cDNA 1700016P04 gene 1700016P04Rik
 1438792_at 0.89821 RIKEN cDNA 1200007D18 gene 1200007D18Rik
 1438793_x_at 1.00641 RIKEN cDNA 1200007D18 gene 1200007D18Rik
 1438794_x_at 1.03351 ribosomal protein S13 /// similar to ribosomal protein S13 /// similar to
 ribosomal protein S13 /// similar to ribosomal protein S13 /// similar to ribosomal protein S13 /// similar
 to ribosomal protein S13 /// similar to ribosomal protein S13 /// similar to ribosomal protein S13
 Rps13 /// LOC625298 /// LOC628061 /// LOC633920 /// LOC637251 /// LOC668239 /// LOC671723 ///
 LOC671887
 1438795_x_at 1.04513 FK506 binding protein 1a Fkbp1a
 1438796_at 1.13239 --- ---
 1438797_at 0.88101 RIKEN cDNA 6330509M05 gene 6330509M05Rik
 1438798_at 1.13069 RIKEN cDNA 4931406P16 gene 4931406P16Rik
 1438799_at 0.90131 Dlx6 opposite strand transcript 1 Dlx6os1
 1438800_at 1.20539 N-acetylglucosamine kinase Nagk
 1438801_at 0.97884 dynamin 3 Dnm3
 1438802_at 1.76286 Forkhead box P1 Foxp1
 1438803_s_at 0.98141 sorting nexin 16 Snx16
 1438804_at 1.07046 septin 10 10-Sep
 1438805_at 1.09067 RIKEN cDNA 9230106B05 gene 9230106B05Rik
 1438806_at 0.92867 B-cell CLL/lymphoma 9-like Bcl9l
 1438807_at 1.32039 Heterogeneous nuclear ribonucleoprotein R Hnrpr
 1438808_at 1.01850 transformation related protein 53 Trp53
 1438809_at 0.78443 "ATP synthase, H+ transporting, mitochondrial F1 complex, gamma
 polypeptide 1" Atp5c1
 1438810_at 1.07333 "DNA segment, Chr 10, ERATO Doi 755, expressed" D10Ert755e
 1438811_at 1.23694 "discs, large homolog 7 (Drosophila)" Dlg7
 1438812_x_at 0.89285 ubiquitin specific peptidase 19 Usp19
 1438813_at 1.02849 RIKEN cDNA 1700003O08 gene 1700003O08Rik
 1438814_at 1.65898 --- ---
 1438815_at 0.71022 "histone 2, H2aa2" Hist2h2aa2
 1438816_at 1.34902 AT hook containing transcription factor 1 Ahctf1
 1438817_at 1.47391 DNA2 DNA replication helicase 2-like (yeast) Dna2l

1438818_at 1.06231 NMD3 homolog (*S. cerevisiae*) Nmd3
1438819_at 1.17272 Transcribed locus ---
1438820_at 1.09993 ring finger protein 17 Rnf17
1438821_at 1.27086 Ring finger and WD repeat domain 2 Rfwd2
1438822_at 0.82799 dorso-medial telencephalon gene 2 Dmt2
1438823_at 1.00955 Transcribed locus ---
1438824_at 1.67416 "solute carrier family 20, member 1" Slc20a1
1438825_at 0.96559 Calmodulin 3 Calm3
1438826_x_at 0.96743 calmodulin 3 Calm3
1438827_at 1.00643 Glutaminase Glu
1438828_at 1.52911 Rap guanine nucleotide exchange factor (GEF) 6 Rapgef6
1438829_at 0.81401 "gene model 96, (NCBI)" Gm96
1438830_at 0.91643 "In vitro fertilized eggs cDNA, RIKEN full-length enriched library, clone:7420422E11 product:hypothetical protein, full insert sequence" ---
1438831_at 1.76904 "Cdc2-related kinase, arginine/serine-rich" Crkrs
1438832_x_at 0.93796 DEAH (Asp-Glu-Ala-His) box polypeptide 30 Dhx30
1438833_at 0.96823 cancer susceptibility candidate 5 /// similar to cancer susceptibility candidate 5 isoform 2 Casc5 /// LOC677208
1438834_at 1.99605 motile sperm domain containing 2 Mospd2
1438835_a_at 1.03554 elongation factor Tu GTP binding domain containing 2 Eftud2
1438836_at 0.96794 Elongation factor Tu GTP binding domain containing 2 Eftud2
1438837_at 1.00877 "Cholinergic receptor, nicotinic, alpha polypeptide 9" Chrna9
1438838_at 1.06907 RIKEN cDNA B230206F22 gene B230206F22Rik
1438839_a_at 1.05010 "tyrosine 3-monooxygenase/tryptophan 5-monooxygenase activation protein, epsilon polypeptide" Ywhae
1438840_x_at 1.04608 apolipoprotein A-I Apoa1
1438841_s_at 1.00594 arginase type II Arg2
1438842_at 1.07587 mitochondrial carrier homolog 2 (*C. elegans*) Mtch2
1438843_x_at 1.09307 mitochondrial carrier homolog 2 (*C. elegans*) Mtch2
1438844_x_at 1.01755 spermatogenesis associated 5 Spata5
1438845_at 0.91340 inositol hexaphosphate kinase 1 Ihpk1
1438846_x_at 0.94745 inositol hexaphosphate kinase 1 Ihpk1
1438847_at 0.93938 Max dimerization protein 3 Mxd3
1438848_at 1.35247 Recombining binding protein suppressor of hairless (*Drosophila*) Rbpsuh
1438849_at 0.93153 --- ---
1438850_at 1.03523 claudin 1 Cldn1
1438851_x_at 0.86991 claudin 1 Cldn1
1438852_x_at 0.86891 "minichromosome maintenance deficient 6 (MIS5 homolog, *S. pombe*) (*S. cerevisiae*)" Mcm6
1438853_x_at 1.07069 DEAD (Asp-Glu-Ala-Asp) box polypeptide 54 Ddx54
1438854_x_at 0.99231 phosphatidylinositol membrane-associated 1 Pitpm1
1438855_x_at 1.37289 "tumor necrosis factor, alpha-induced protein 2" Tnfaip2
1438856_x_at 1.03729 "serine (or cysteine) peptidase inhibitor, clade B, member 5" Serpinb5

1438857_x_at 1.16776 --- ---
1438858_x_at 1.00906 "Histocompatibility 2, class II antigen A, alpha" H2-Aa
1438859_x_at 0.88558 ribosomal protein S29 /// similar to 40S ribosomal protein S29 /// similar to
40S ribosomal protein S29 Rps29 /// LOC629957 /// LOC637599
1438860_a_at 1.02107 "solute carrier family 44, member 2" Slc44a2
1438861_at 1.03385 basonuclin 2 Bnc2
1438862_at 1.43783 RIKEN cDNA A630005I04 gene A630005I04Rik
1438863_at 0.99640 RIKEN cDNA A530079E22 gene A530079E22Rik
1438864_at 2.02402 RIKEN cDNA C920006C10 gene C920006C10Rik
1438865_at 0.91516 histocompatibility 13 H13
1438866_at 1.06309 "glutamate receptor ionotropic, NMDA3A" Grin3a
1438867_at 0.92124 "gene model 1070, (NCBI)" Gm1070
1438868_at 1.02196 "DNA segment, Chr 14, ERATO Doi 668, expressed" D14Ertd668e
1438869_at 1.12023 RIKEN cDNA 4930515G01 gene 4930515G01Rik
1438870_at 1.01460 --- ---
1438871_at 1.11847 Son cell proliferation protein Son
1438872_at 1.04287 cDNA sequence BC050188 BC050188
1438873_at 1.04535 --- ---
1438874_at 1.16635 "non-metastatic cells 7, protein expressed in" Nme7
1438875_at 1.03859 KRAB box containing protein RP23-221M24.3
1438876_at 0.90285 RIKEN cDNA A130014A01 gene A130014A01Rik
1438877_at 1.13810 "transient receptor potential cation channel, subfamily M, member 6" Trpm6
1438878_at 2.45797 RIKEN cDNA 6430537K16 gene 6430537K16Rik
1438879_at 1.00726 Dimethylarginine dimethylaminohydrolase 1 Ddah1
1438880_at 1.14338 RIKEN cDNA 1700012D14 gene 1700012D14Rik
1438881_at 0.90935 "vezatin, adherens junctions transmembrane protein" Vezt
1438882_at 2.72249 Rho GTPase activating protein 18 Arhgap18
1438883_at 1.01016 fibroblast growth factor 5 Fgf5
1438884_at 0.92267 RIKEN cDNA D830007B15 gene D830007B15Rik
1438885_at 0.97417 expressed sequence BB182387 BB182387
1438886_at 1.03869 hairy/enhancer-of-split related with YRPW motif-like Heyl
1438887_a_at 1.01395 germ cell-less homolog 1 (Drosophila) Gmcl1
1438888_at 1.09851 germ cell-less homolog 1 (Drosophila) Gmcl1
1438889_at 1.13345 "15 days embryo head cDNA, RIKEN full-length enriched library, clone:
D930025C07 product:unclassifiable, full insert sequence" ---
1438890_at 1.34643 --- ---
1438891_at 1.03329 "Catenin (cadherin associated protein), delta 2" Ctnnd2
1438892_at 1.32946 Quaking Qk
1438893_at 0.98848 RIKEN cDNA 5530601H04 gene 5530601H04Rik
1438894_at 1.05529 "16 days embryo head cDNA, RIKEN full-length enriched library, clone:
C130060L08 product:unclassifiable, full insert sequence" ---
1438895_at 2.26036 RIKEN cDNA A430102J17 gene A430102J17Rik
1438896_at 1.11471 "DnaJ (Hsp40) homolog, subfamily C, member 6" Dnajc6

1438897_at 1.08094 "Zinc finger, DHHC domain containing 3" Zdhhc3
1438898_at 0.98836 "Adult male corpora quadrigemina cDNA, RIKEN full-length enriched library, clone:B230310B10 product:unclassifiable, full insert sequence" ---
1438899_at 0.98174 Kruppel-like factor 17 Klf17
1438900_at 1.86752 "SAC1 (suppressor of actin mutations 1, homolog)-like (S. cerevisiae)" Sacm11
1438901_at 0.89631 cDNA sequence BC024814 BC024814
1438902_a_at 1.04924 "heat shock protein 90kDa alpha (cytosolic), class A member 1" Hsp90aa1
1438903_at 1.07527 Macrophage erythroblast attacher Maea
1438904_at 0.73509 NHS-like 1 Nhs11
1438905_x_at 0.97432 RIKEN cDNA C030046I01 gene C030046I01Rik
1438906_at 3.45713 --- ---
1438907_at 0.94281 "3 days neonate thymus cDNA, RIKEN full-length enriched library, clone: A630033C13 product:unclassifiable, full insert sequence" ---
1438908_at 0.84199 mitogen activated protein kinase kinase kinase 12 Map3k12
1438909_at 0.95962 saccharopine dehydrogenase (putative) Sccpdh
1438910_a_at 0.92597 stomatin Stom
1438911_at 0.98692 Septin 7 7-Sep
1438912_at 0.90225 "hepatoma-derived growth factor, related protein 2" Hdgfrp2
1438913_x_at 1.04118 "hepatoma-derived growth factor, related protein 2" Hdgfrp2
1438914_at 1.09589 RIKEN cDNA 2310028N02 gene 2310028N02Rik
1438915_at 0.93806 RIKEN cDNA 6720401G13 gene 6720401G13Rik
1438916_x_at 0.86724 RIKEN cDNA 6720401G13 gene 6720401G13Rik
1438917_x_at 0.98848 nucleoporin 62 Nup62
1438918_at 0.96973 farnesyl diphosphate farnesyl transferase 1 /// similar to farnesyl diphosphate farnesyl transferase 1 Fdft1 /// LOC627717
1438919_x_at 1.00469 farnesyl diphosphate farnesyl transferase 1 /// similar to farnesyl diphosphate farnesyl transferase 1 Fdft1 /// LOC627717
1438920_x_at 1.05201 Complement receptor related protein Crry
1438921_at 1.05059 Ataxia telangiectasia and Rad3 related Atr
1438922_x_at 1.12206 "solute carrier family 25 (mitochondrial carrier, adenine nucleotide translocator), member 5 /// similar to ADP/ATP translocase 2 (Adenine nucleotide translocator 2) (ANT 2) (ADP,ATP carrier protein 2) (Solute carrier family 25 member 5) /// similar to ADP/ATP translocase 2 (Adenine nucleotide translocator 2) (ANT 2) (ADP,ATP carrier protein 2) (Solute carrier family 25 member 5) /// similar to ADP/ATP translocase 2 (Adenine nucleotide translocator 2) (ANT 2) (ADP, ATP carrier protein 2) (Solute carrier family 25 member 5) /// similar to solute carrier family 25, member A6" Slc25a5 /// LOC433326 /// LOC630624 /// LOC669381 /// LOC672115
1438923_at 0.90390 Fibroblast growth factor (acidic) intracellular binding protein Fibp
1438924_x_at 0.91707 Fibroblast growth factor (acidic) intracellular binding protein Fibp
1438925_x_at 1.08685 "ATPase, H+ transporting, lysosomal V0 subunit C" Atp6v0c
1438926_at 1.06711 "Transcribed locus, strongly similar to NP_063918.1 ribosomal protein L36a [Mus musculus]" ---
1438927_x_at 0.98053 similar to 60S ribosomal protein L23a /// ribosomal protein L23a /// hypothetical gene supported by BC029892 /// hypothetical gene supported by BC029892 /// similar to

60S ribosomal protein L23a /// similar to 60S ribosomal protein L23a /// similar to 60S ribosomal protein L23a /// hypothetical LOC631470 /// similar to 60S ribosomal protein L23a /// similar to 60S ribosomal protein L23a /// similar to 60S ribosomal protein L23a /// similar to 60S ribosomal protein L23a /// similar to 60S ribosomal protein L23a LOC227054 /// Rpl23a /// LOC384840 /// LOC385336 /// LOC619975 /// LOC620750 /// LOC621498 /// LOC631470 /// LOC666448 /// LOC666501 /// LOC666548 /// LOC666738 /// LOC666790
 1438928_x_at 0.97216 ninjurin 1 Ninj1
 1438929_at 1.20019 --- ---
 1438930_s_at 0.73965 methyl CpG binding protein 2 Mecp2
 1438931_s_at 0.85971 sestrin 1 Sesn1
 1438932_at 1.01356 "RAS, guanyl releasing protein 2" Rasgrp2
 1438933_x_at 0.93936 "RAS, guanyl releasing protein 2" Rasgrp2
 1438934_x_at 0.95046 "sema domain, immunoglobulin domain (Ig), transmembrane domain (TM) and short cytoplasmic domain, (semaphorin) 4A" Sema4a
 1438935_at 0.95552 expressed sequence BB234005 BB234005
 1438936_s_at 0.99513 "angiogenin, ribonuclease A family, member 1" Ang1
 1438937_x_at 1.02637 "angiogenin, ribonuclease A family, member 1" Ang1
 1438938_x_at 1.07114 prohibitin 2 Phb2
 1438939_x_at 1.04118 necdin Ndn
 1438940_x_at 0.92364 high mobility group nucleosomal binding domain 1 Hmgn1
 1438941_x_at 1.03471 adenosine monophosphate deaminase 2 (isoform L) Ampd2
 1438942_x_at 1.03623 "transglutaminase 2, C polypeptide" Tgm2
 1438943_x_at 0.97388 ribophorin I Rpn1
 1438944_at 0.98254 RIKEN cDNA 1700007J24 gene 1700007J24Rik
 1438945_x_at 0.94851 gap junction membrane channel protein alpha 1 Gjal
 1438946_at 0.99703 "platelet derived growth factor receptor, alpha polypeptide" Pdgfra
 1438947_x_at 0.89488 "sema domain, immunoglobulin domain (Ig), short basic domain, secreted, (semaphorin) 3 F" Sema3f
 1438948_x_at 0.90622 translocator protein Tspo
 1438949_at 1.00865 resistance to inhibitors of cholinesterase 8 homolog (C. elegans) Ric8
 1438950_x_at 1.05313 resistance to inhibitors of cholinesterase 8 homolog (C. elegans) Ric8
 1438951_x_at 0.98161 nucleoporin 54 Nup54
 1438952_x_at 1.03894 TAR (HIV) RNA binding protein 2 Tarbp2
 1438953_at 1.08394 --- ---
 1438954_x_at 1.08134 --- ---
 1438955_x_at 1.02065 peptidylprolyl isomerase F (cyclophilin F) Ppif
 1438956_x_at 1.06072 proviral integration site 3 Pim3
 1438957_x_at 1.08122 CDP-diacylglycerol synthase (phosphatidate cytidyltransferase) 2 Cds2
 1438958_x_at 1.04772 FK506 binding protein 1a Fkbp1a
 1438959_x_at 0.97398 Transcribed locus ---
 1438960_at 0.91263 --- ---
 1438961_s_at 0.97493 bleomycin hydrolase Blmh
 1438962_s_at 1.04897 DEAD/H (Asp-Glu-Ala-Asp/His) box polypeptide 31 Ddx31

1438963_s_at 0.99069 TCF3 (E2A) fusion partner Tfpt
1438964_x_at 0.94536 TCF3 (E2A) fusion partner Tfpt
1438965_x_at 0.97272 RIKEN cDNA A230056P14 gene A230056P14Rik
1438966_x_at 0.66852 fibromodulin Fmod
1438967_x_at 0.94364 anti-Mullerian hormone type 2 receptor Amhr2
1438968_x_at 1.01778 "serine protease inhibitor, Kunitz type 2" Spint2
1438969_x_at 0.92886 DEAH (Asp-Glu-Ala-His) box polypeptide 30 Dhx30
1438970_x_at 0.96292 RIKEN cDNA 1700034F02 gene 1700034F02Rik
1438971_x_at 0.78211 ubiquitin-conjugating enzyme E2H Ube2h
1438972_x_at 0.95604 RIKEN cDNA 2810410L24 gene 2810410L24Rik
1438973_x_at 0.93458 gap junction membrane channel protein alpha 1 Gja1
1438974_x_at 0.88949 phosphatidylinositol membrane-associated 1 Pitpm1
1438975_x_at 0.93744 "zinc finger, DHHC domain containing 14" Zdhhc14
1438976_x_at 1.01690 "methionine adenosyltransferase II, alpha" Mat2a
1438977_x_at 1.04732 "RAN, member RAS oncogene family /// similar to RAN, member RAS oncogene family" Ran /// LOC671251
1438978_x_at 0.95010 necdin Ndn
1438979_s_at 0.98886 RIKEN cDNA 1700029I15 gene 1700029I15Rik
1438980_x_at 0.91695 RIKEN cDNA 4732466D17 gene 4732466D17Rik
1438981_at 1.01904 "Protein kinase C, beta 1" Prkcb1
1438982_s_at 0.83657 RIKEN cDNA 2810417J12 gene 2810417J12Rik
1438983_x_at 0.82767 expressed sequence AI413782 AI413782
1438984_x_at 0.99998 "proteasome (prosome, macropain) subunit, beta type 4" Psmb4
1438985_x_at 1.00953 OTU domain containing 5 Otud5
1438986_x_at 1.05531 ribosomal protein S17 Rps17
1438987_at 0.93007 --- ---
1438988_x_at 1.01085 hematological and neurological expressed sequence 1 Hn1
1438989_s_at 0.58003 RIKEN cDNA B130021B11 gene B130021B11Rik
1438990_x_at 0.92626 IQ motif containing F3 Iqcf3
1438991_x_at 0.91722 "protein phosphatase 2 (formerly 2A), regulatory subunit A (PR 65), alpha isoform" Ppp2r1a
1438992_x_at 0.93780 activating transcription factor 4 Atf4
1438993_a_at 1.01066 "ATPase, H+ transporting, lysosomal V1 subunit D" Atp6v1d
1438994_at 0.94861 hyaluronoglucosaminidase 5 Hyal5
1438995_at 1.10202 RIKEN cDNA 6330500D04 gene 6330500D04Rik
1438996_at 0.93716 "gene model 118, (NCBI)" Gm118
1438997_at 1.09275 Endoplasmic reticulum (ER) to nucleus signalling 1 Ern1
1438998_at 1.00927 RAS p21 protein activator 1 Rasa1
1438999_a_at 1.00224 nuclear factor of activated T-cells 5 Nfat5
1439000_at 1.09844 acyl-CoA synthetase bubblegum family member 2 Acsbg2
1439001_at 1.04863 hypothetical protein LOC665286 /// hypothetical protein LOC677425
LOC665286 /// LOC677425
1439002_s_at 1.07525 chorionic somatomammotropin hormone 1 Csh1

1439003_s_at 0.87484 RIKEN cDNA 1700008F21 gene /// similar to coiled-coil domain containing
7 1700008F21Rik /// LOC672965

1439004_at 0.84786 "ribosomal protein S6 kinase, polypeptide 5" Rps6ka5

1439005_x_at 1.06874 "tyrosine 3-monooxygenase/tryptophan 5-monooxygenase activation protein,
zeta polypeptide" Ywhaz

1439006_x_at 0.97295 RIKEN cDNA 6430550H21 gene 6430550H21Rik

1439007_at 0.93691 "asparagine-linked glycosylation 6 homolog (yeast, alpha-1,3,-
glucosyltransferase)" Alg6

1439008_at 0.80225 zinc finger protein 319 Zfp319

1439009_at 1.01658 guanine nucleotide binding protein-like 3 (nucleolar)-like Gnl3l

1439010_at 1.16857 RIKEN cDNA D330037H05 gene /// similar to La-related protein 4 (La
ribonucleoprotein domain family member 4) /// similar to La-related protein 4 (La ribonucleoprotein
domain family member 4) D330037H05Rik /// LOC673040 /// LOC677019

1439011_at 1.01515 Transcribed locus ---

1439012_a_at 0.93324 deoxycytidine kinase Dck

1439013_x_at 0.90596 "acylphosphatase 1, erythrocyte (common) type" Acyp1

1439014_at 0.86162 RIKEN cDNA 1600021P15 gene 1600021P15Rik

1439015_at 0.96889 glial cell line derived neurotrophic factor family receptor alpha 1 Gfra1

1439016_x_at 1.08844 small proline-rich protein 2A Sprr2a

1439017_x_at 0.89033 adiponectin receptor 1 Adipor1

1439018_at 0.68392 RIKEN cDNA 6330505N24 gene 6330505N24Rik

1439019_at 0.51752 Fraser syndrome 1 homolog (human) Fras1

1439020_at 1.01682 expressed sequence AW146020 AW146020

1439021_at 0.88337 "centaurin, beta 5" Centb5

1439022_at 0.78129 phosphatase and actin regulator 1 Phactr1

1439023_at 1.02971 DEAD (Asp-Glu-Ala-Asp) box polypeptide 55 Ddx55

1439024_at 1.00270 BCL2-associated athanogene 4 Bag4

1439025_at 0.88426 zinc finger protein 446 Zfp446

1439026_at 1.03120 "transient receptor potential cation channel, subfamily M, member 3" Trpm3

1439027_at 1.04249 RIKEN cDNA C330023M02 gene C330023M02Rik

1439028_at 0.87077 ubiquitin-fold modifier 1 Ufm1

1439029_at 1.02053 glutamic pyruvate transaminase (alanine aminotransferase) 2 Gpt2

1439030_at 0.96513 GDP-mannose pyrophosphorylase B Gmppb

1439031_at 0.99756 junctophilin 4 Jph4

1439032_at 0.94913 centrosomal protein 27 Cep27

1439033_at 1.20991 "zinc finger, CCHC domain containing 7" Zcchc7

1439034_at 0.97066 sialophorin Spn

1439035_at 1.05904 "12 days embryo spinal cord cDNA, RIKEN full-length enriched library, clone:
C530050B12 product:unclassifiable, full insert sequence" ---

1439036_a_at 0.72867 "ATPase, Na⁺/K⁺ transporting, beta 1 polypeptide" Atp1b1

1439037_at 0.90095 DEAD (Asp-Glu-Ala-Asp) box polypeptide 17 Ddx17

1439038_at 1.01491 RIKEN cDNA 9130227C08Rik gene 9130227C08Rik

1439039_at 0.92458 transmembrane protein 104 Tmem104

1439040_at 0.94957 centromere protein E Cenpe
 1439041_at 1.01729 "solute carrier family 39 (zinc transporter), member 10" Slc39a10
 1439042_at 0.91465 adenylate cyclase activating polypeptide 1 receptor 1 Adcyap1r1
 1439043_at 1.22560 transformer 2 alpha homolog (Drosophila) Tra2a
 1439044_at 0.69787 zinc finger protein 354C Zfp354c
 1439045_x_at 0.95134 membrane targeting (tandem) C2 domain containing 1 Mtac2d1
 1439046_at 1.08238 coiled-coil domain containing 55 Ccdc55
 1439047_s_at 0.89310 RecQ protein-like Recql
 1439048_at 0.74305 RIKEN cDNA 2900078E11 gene 2900078E11Rik
 1439049_at 1.27745 DPH5 homolog (S. cerevisiae) Dph5
 1439050_at 0.86248 --- ---
 1439051_a_at 1.06833 MAP/microtubule affinity-regulating kinase 4 Mark4
 1439052_at 1.07082 RIKEN cDNA 5430433G21 gene 5430433G21Rik
 1439053_at 1.08340 "DNA segment, Chr 10, Brigham & Women's Genetics 1070 expressed"
 D10Bwg1070e
 1439054_at 0.97854 "zinc finger protein 36, C3H type-like 2" Zfp3612
 1439055_at 0.91318 hypothetical LOC217066 LOC217066
 1439056_at 1.05600 hypothetical protein LOC636791 LOC636791
 1439057_x_at 0.97299 "zinc finger, DHHC domain containing 6" Zdhhc6
 1439058_at 0.91499 splicing factor proline/glutamine rich (polypyrimidine tract binding protein
 associated) Sfpq
 1439059_at 1.03758 CDNA sequence BC031748 BC031748
 1439060_s_at 0.89353 "WD repeat domain, phosphoinositide interacting 1" Wipi1
 1439061_at 1.22787 coiled-coil domain containing 84 Ccdc84
 1439062_at 0.97080 kelch repeat and BTB (POZ) domain containing 3 Kbtbd3
 1439063_at 1.30060 additional sex combs like 2 (Drosophila) Asxl2
 1439064_at 0.92765 cDNA sequence BC030046 BC030046
 1439065_x_at 1.02558 Similar to zinc finger protein 665 MGC117846
 1439066_at 1.03871 angiopoietin 1 Angpt1
 1439067_at 1.02984 leukocyte-associated Ig-like receptor 1 Lair1
 1439068_at 1.24821 type 1 tumor necrosis factor receptor shedding aminopeptidase regulator Arts1
 1439069_a_at 0.94014 RIKEN cDNA 4933439C20 gene 4933439C20Rik
 1439070_x_at 0.94577 RIKEN cDNA 4933439C20 gene /// RIKEN cDNA D030013I16 gene
 4933439C20Rik /// D030013I16Rik
 1439071_at 0.99949 RIKEN cDNA 5430416N02 gene 5430416N02Rik
 1439072_at 0.94107 "solute carrier family 1 (glial high affinity glutamate transporter), member 3"
 Slc1a3
 1439073_at 1.13738 similar to Afadin (Af-6 protein) LOC627912
 1439074_a_at 1.01993 Son cell proliferation protein Son
 1439075_at 1.09510 polymerase (RNA) III (DNA directed) polypeptide F Polr3f
 1439076_at 0.97348 DEAH (Asp-Glu-Ala-His) box polypeptide 29 Dhx29
 1439077_at 0.90398 "zinc finger, X-linked, duplicated A /// similar to Zinc finger X-linked protein
 ZXDA" Zxda /// LOC668171

1439078_at	1.11143	kelch-like 4 (Drosophila)	Klhl4
1439079_a_at	1.53988	ErbB2 interacting protein	ErbB2ip
1439080_at	1.36817	ErbB2 interacting protein	ErbB2ip
1439081_at	2.43054	meningioma expressed antigen 5 (hyaluronidase)	Mgea5
1439082_at	1.11093	DEAD (Asp-Glu-Ala-Asp) box polypeptide 50	Ddx50
1439083_at	0.89545	Abelson helper integration site	Ahi1
1439084_at	0.93669	chemokine (C-X-C motif) ligand 12	Cxcl12
1439085_at	0.95810	RIKEN cDNA C130039O16 gene	C130039O16Rik
1439086_at	0.98965	RIKEN cDNA A930009L07 gene	A930009L07Rik
1439087_a_at	0.76985	RIKEN cDNA 1500004A08 gene	1500004A08Rik
1439088_at	1.06971	PDZ domain containing 8	Pdzd8
1439089_at	0.94480	zinc finger and BTB domain containing 41 homolog	Zbtb41
1439090_at	1.34029	RIKEN cDNA D030022P07 gene	D030022P07Rik
1439091_at	1.03178	"Fanconi anemia, complementation group D2"	Fancd2
1439092_at	1.12445	nuclear RNA export factor 1 homolog (S. cerevisiae)	Nxf1
1439093_at	1.37289	---	---
1439094_at	1.41323	"clathrin, heavy polypeptide (Hc)"	Cltc
1439095_at	1.04960	"splicing factor, arginine/serine-rich 11"	Sfrs11
1439096_at	0.87252	D-aspartate oxidase	Ddo
1439097_at	1.14043	"DNA segment, Chr 10, Wayne State University 52, expressed"	D10Wsu52e
1439098_at	1.06413	RIKEN cDNA E230013L22 gene	E230013L22Rik
1439099_at	1.02673	neuro-oncological ventral antigen 2	Nova2
1439100_s_at	1.08297	---	---
1439101_at	0.96661	RIKEN cDNA D830007F02 gene	D830007F02Rik
1439102_at	1.04969	fetal Alzheimer antigen	Falz
1439103_at	0.99015	"Vcell division cycle 73, Paf1/RNA polymerase II complex component, homolog (S. cerevisiae)"	Cdc73
1439104_at	0.92966	RIKEN cDNA C230081A13 gene	C230081A13Rik
1439105_at	0.92285	cytidine and dCMP deaminase domain containing 1	Cdadc1
1439106_at	0.96667	zinc finger protein 462	Zfp462
1439107_a_at	1.06963	myeloid/lymphoid or mixed-lineage leukemia 5	Mll5
1439108_at	0.98237	myeloid/lymphoid or mixed-lineage leukemia 5	Mll5
1439109_at	0.89807	coiled-coil domain containing 68	Ccdc68
1439110_at	0.98859	RIKEN cDNA A930012O16 gene	A930012O16Rik
1439111_at	0.98303	---	---
1439112_at	0.87608	RIKEN cDNA 2810442I21 gene	2810442I21Rik
1439113_at	0.90330	RIKEN cDNA 2410018L13 gene /// RIKEN cDNA 6030426L16 gene /// similar to RIKEN cDNA 6030426L16 /// similar to gonadotropin inducible ovarian transcription factor 1 /// similar to gonadotropin inducible ovarian transcription factor 1 /// similar to gonadotropin inducible ovarian transcription factor 1 /// similar to gonadotropin inducible ovarian transcription factor 1 /// similar to Hippocalcin-like protein 1 (Visinin-like protein 3) (VILIP-3) (Neural visinin-like protein 3) (NVL-3) (NVP-3) /// similar to Hippocalcin-like protein 1 (Visinin-like protein 3) (VILIP-3) (Neural visinin-like protein 3) (NVL-3)	

(NVP-3) 2410018L13Rik /// 6030426L16Rik /// LOC245297 /// LOC668553 /// LOC668645 ///
 LOC668662 /// LOC668701 /// LOC668710 /// LOC675539 /// LOC675641
 1439114_at 0.98161 cDNA sequence BC013672 BC013672
 1439115_at 1.02678 "microcephaly, primary autosomal recessive 1" Mcph1
 1439116_at 0.92250 "Cofactor required for Sp1 transcriptional activation, subunit 2" Crsp2
 1439117_at 0.74687 calmin Clmn
 1439118_at 1.04170 "solute carrier family 35, member E4" Slc35e4
 1439119_a_at 0.97685 cDNA sequence BC010304 BC010304
 1439120_at 0.98551 cDNA sequence BC010304 BC010304
 1439121_at 0.97491 "histocompatibility 2, T region locus 22" H2-T22
 1439122_at 1.06714 --- ---
 1439123_at 2.66248 PHD finger protein 21A Phf21a
 1439124_at 0.96374 RIKEN cDNA 9530020G05 gene 9530020G05Rik
 1439125_at 1.43232 Adducin 1 (alpha) Add1
 1439126_at 1.72116 RIKEN cDNA 1110007A13 gene 1110007A13Rik
 1439127_at 0.91246 expressed sequence AI314180 AI314180
 1439128_at 1.04634 zinc finger and BTB domain containing 20 Zbtb20
 1439129_at 1.07209 RIKEN cDNA 1110060D06 gene /// dedicator of cytokinesis 5
 1110060D06Rik /// Dock5
 1439130_at 1.03790 autophagy-related 4D (yeast) Atg4d
 1439131_at 1.18152 RIKEN cDNA 4932441K18 gene 4932441K18Rik
 1439132_at 1.01822 PHD finger protein 8 Phf8
 1439133_at 0.97538 RIKEN cDNA 4930524O07 gene 4930524O07Rik
 1439134_s_at 0.94393 RIKEN cDNA 4930524O07 gene 4930524O07Rik
 1439135_at 0.99743 "Adult male spinal cord cDNA, RIKEN full-length enriched library, clone:
 A330034O07 product:unclassifiable, full insert sequence" ---
 1439136_at 2.06244 Single-stranded DNA binding protein 3 Ssbp3
 1439137_at 1.04295 "solute carrier family 4, sodium bicarbonate cotransporter, member 9" Slc4a9
 1439138_at 1.72087 RIKEN cDNA 2310035C23 gene 2310035C23Rik
 1439139_at 1.29639 "DNA segment, Chr 2, ERATO Doi 640, expressed" D2Ertd640e
 1439140_at 1.08466 RIKEN cDNA 1700019P01 gene 1700019P01Rik
 1439141_at 1.03675 G protein-coupled receptor 18 Gpr18
 1439142_at 0.99572 "Swi/SNF related matrix associated, actin dependent regulator of chromatin,
 subfamily a-like 1" Smarcal1
 1439143_at 0.94865 hypothetical protein 9330167E06 9.33E+12
 1439144_at 1.00487 "CWF19-like 1, cell cycle control (S. pombe)" Cwf19l1
 1439145_at 1.01835 lymphocyte protein tyrosine kinase Lck
 1439146_s_at 0.96209 lymphocyte protein tyrosine kinase Lck
 1439147_at 1.02491 RIKEN cDNA 1110049B09 gene 1110049B09Rik
 1439148_a_at 1.27125 "phosphofructokinase, liver, B-type" Pfk1
 1439149_s_at 0.95142 hypothetical protein LOC629141 LOC629141
 1439150_x_at 0.91850 GH regulated TBC protein 1 Grtp1
 1439151_at 0.82713 methionine sulfoxide reductase B3 Msrb3

1439152_at	0.93354	methenyltetrahydrofolate synthetase domain containing	Mthfsd
1439153_at	0.91944	IBR domain containing 2	Ibrdc2
1439154_at	1.07315	nucleoporin 98	Nup98
1439155_at	1.14911	methyltransferase-like 1	Mettl1
1439156_at	1.03363	"gene model 962, (NCBI)"	Gm962
1439157_at	1.06152	RIKEN cDNA D330012D11 gene	D330012D11Rik
1439158_at	1.03203	tousled-like kinase 1	Tlk1
1439159_at	1.59390	---	---
1439160_at	0.90390	RIKEN cDNA 4732496O08 gene	4732496O08Rik
1439161_at	1.93352	"SAPS domain family, member 3"	Saps3
1439162_at	0.96276	RIKEN cDNA 1700011J10 gene	1700011J10Rik
1439163_at	0.93257	zinc finger and BTB domain containing 16	Zbtb16
1439164_at	0.88749	"cyclin-dependent kinase inhibitor 2C (p18, inhibits CDK4)"	Cdkn2c
1439165_at	1.23932	---	---
1439166_at	1.16233	nucleolar protein 11	Nol11
1439167_at	0.95823	peroxisomal trans-2-enoyl-CoA reductase	Pecr
1439168_at	1.83200	"calcium/calmodulin-dependent protein kinase II, delta"	Camk2d
1439169_at	1.39738	Eukaryotic translation initiation factor 4E nuclear import factor 1	Eif4enif1
1439170_at	1.00395	---	---
1439171_at	0.88038	RIKEN cDNA C530008M17 gene	C530008M17Rik
1439172_at	1.02857	ATPase type 13A5	Atp13a5
1439173_at	1.06717	hook homolog 1 (Drosophila)	Hook1
1439174_at	0.93445	Unc-5 homolog C (C. elegans)	Unc5c
1439175_at	1.00628	"Protein tyrosine phosphatase, receptor type, D"	Ptprd
1439176_at	0.97286	RIKEN cDNA E130120F12 gene	E130120F12Rik
1439177_at	0.93899	Transcribed locus	---
1439178_at	1.06897	"adrenergic receptor kinase, beta 2"	Adrbk2
1439179_a_at	0.84618	RIKEN cDNA 5830405N20 gene	5830405N20Rik
1439180_at	1.20363	RIKEN cDNA A430093A21 gene	A430093A21Rik
1439181_at	1.04054	cDNA sequence BC043301	BC043301
1439182_at	0.85895	"DNA segment, Chr 17, Wayne State University 92, expressed"	D17Wsu92e
1439183_at	0.94727	N-acylsphingosine amidohydrolase (alkaline ceramidase) 3	Asah3
1439184_s_at	1.07385	thioredoxin-like 5	Txn15
1439185_x_at	0.89955	RIKEN cDNA D430028G21 gene	D430028G21Rik
1439186_at	1.01106	RIKEN cDNA E130319B15 gene	E130319B15Rik
1439187_at	0.97068	vacuolar protein sorting 13 D (yeast)	Vps13d
1439188_at	1.10464	cleavage and polyadenylation specific factor 6	Cpsf6
1439189_at	1.09083	RIKEN cDNA D630023B12 gene	D630023B12Rik
1439190_at	1.01764	forkhead-associated (FHA) phosphopeptide binding domain 1	Fhad1
1439191_at	1.03154	---	---
1439192_at	0.88530	neuro-oncological ventral antigen 2	Nova2
1439193_at	0.91004	Paired related homeobox protein-like 1	Prrxl1
1439194_at	0.98228	RIKEN cDNA C030048H21 gene	C030048H21Rik

1439195_at 1.10333 --- ---
1439196_at 1.04066 hook homolog 3 (Drosophila) Hook3
1439197_at 0.89880 "phosphatidylinositol 4-kinase, catalytic, beta polypeptide" Pik4cb
1439198_at 2.59156 PTK2 protein tyrosine kinase 2 Ptk2
1439199_at 1.05125 "Adult male corpora quadrigemina cDNA, RIKEN full-length enriched library, clone:B230317E19 product:unclassifiable, full insert sequence" ---
1439200_x_at 1.24344 Reproductive homeobox 4B Rhox4b
1439201_at 1.37829 ubiquitin specific peptidase 14 Usp14
1439202_at 0.97854 --- ---
1439203_at 0.83617 Special AT-rich sequence binding protein 1 Satb1
1439204_at 0.46704 "sodium channel, voltage-gated, type III, alpha" Scn3a
1439205_at 0.80592 "nuclear factor of activated T-cells, cytoplasmic, calcineurin-dependent 2" Nfatc2
1439206_at 1.01296 "Guanine nucleotide binding protein, alpha 12" Gna12
1439207_at 1.05078 paraneoplastic antigen family 5 Pnma5
1439208_at 0.93970 checkpoint kinase 1 homolog (S. pombe) Chek1
1439209_at 2.33777 Transcription factor 12 Tcf12
1439210_at 1.27992 mitochondrial ribosomal protein S9 Mrps9
1439211_at 1.00300 uromodulin-like 1 Umodl1
1439212_at 0.90399 hypothetical protein 6820429M01 6820429M01
1439213_at 1.11848 RIKEN cDNA 4921524J06 gene 4921524J06Rik
1439214_a_at 1.03137 apoptosis inhibitor 5 Api5
1439215_at 1.02557 "ATPase, Ca⁺⁺-sequestering" Atp2c1
1439216_at 1.32237 RIKEN cDNA 2900075B16 gene 2900075B16Rik
1439217_at 0.99502 "NACHT, LRR and PYD containing protein 9c" Nalp9c
1439218_at 1.09904 Uronyl-2-sulfotransferase Ust
1439219_at 0.91776 RIKEN cDNA A730082K24 gene A730082K24Rik
1439220_at 0.94324 "Ankyrin 3, epithelial" Ank3
1439221_s_at 1.00424 CD40 antigen Cd40
1439222_at 0.96929 microrchidia 2B Morc2b
1439223_at 1.07569 tropomodulin 2 Tmod2
1439224_at 1.62288 Ankyrin repeat and IBR domain containing 1 Ankib1
1439225_at 0.89752 "leucyl-tRNA synthetase, mitochondrial" Lars2
1439226_at 0.84897 Dedicator of cytokinesis 8 Dock8
1439227_at 1.73670 "ubiquitin-like, containing PHD and RING finger domains, 1" Uhrf1
1439228_at 1.10653 zinc finger protein 407 Zfp407
1439229_at 0.86924 "pleckstrin homology domain containing, family A member 7" Plekha7
1439230_at 0.88554 RIKEN cDNA 4922505E12 gene 4922505E12Rik
1439231_at 0.96909 --- ---
1439232_at 0.92097 RIKEN cDNA 1500016L03 gene 1500016L03Rik
1439233_at 1.74527 Translocation protein 1 Tloc1
1439234_a_at 0.87853 TM2 domain containing 2 Tm2d2
1439235_x_at 0.85759 TM2 domain containing 2 Tm2d2

1439236_at 0.91586 "DNA segment, Chr 15, ERATO Doi 509, expressed" D15Ert509e
 1439237_a_at 0.88841 slingshot homolog 1 (Drosophila) Ssh1
 1439238_at 1.19739 unc-84 homolog B (C. elegans) Unc84b
 1439239_at 1.04708 lin-7 homolog B (C. elegans) Lin7b
 1439240_x_at 0.96911 lin-7 homolog B (C. elegans) Lin7b
 1439241_x_at 0.88704 steroid 5 alpha-reductase 2-like Srd5a2l
 1439242_at 1.10326 molybdenum cofactor synthesis 3 Mocs3
 1439243_x_at 1.06634 "COP9 (constitutive photomorphogenic) homolog, subunit 5 (Arabidopsis thaliana)" Cops5
 1439244_a_at 0.94724 trinucleotide repeat containing 6a Tnrc6a
 1439245_at 1.05090 trinucleotide repeat containing 6a Tnrc6a
 1439246_x_at 0.86138 trinucleotide repeat containing 6a Tnrc6a
 1439247_at 1.02827 dedicator of cytokinesis 10 /// similar to Dedicator of cytokinesis protein 10 (Protein zizimin 3) Dock10 /// LOC630691
 1439248_at 1.13148 "RMI1, RecQ mediated genome instability 1, homolog (S. cerevisiae)" Rmi1
 1439249_at 0.86930 WW domain containing adaptor with coiled-coil Wac
 1439250_at 0.98513 "SLIT and NTRK-like family, member 3" Slitrk3
 1439251_at 0.66339 Transcribed locus ---
 1439252_at 1.43356 --- ---
 1439253_x_at 1.13562 RIKEN cDNA 2610524G07 gene 2610524G07Rik
 1439254_at 0.93718 A kinase (PRKA) anchor protein 13 Akap13
 1439255_s_at 1.06124 G protein-coupled receptor 137B /// similar to transmembrane 7 superfamily member 1 Gpr137b /// LOC664862
 1439256_x_at 1.08010 G protein-coupled receptor 137B /// similar to transmembrane 7 superfamily member 1 Gpr137b /// LOC664862
 1439257_x_at 0.99433 ribophorin I Rpn1
 1439258_at 1.09825 activating transcription factor 4 Atf4
 1439259_x_at 0.83652 abhydrolase domain containing 4 Abhd4
 1439260_a_at 1.01391 ectonucleotide pyrophosphatase/phosphodiesterase 3 Enpp3
 1439261_x_at 1.08558 "MIT, microtubule interacting and transport, domain containing 1" Mitd1
 1439262_x_at 0.84750 RIKEN cDNA 2310001A20 gene 2310001A20Rik
 1439263_at 0.99172 fibroblast growth factor inducible 15 Fin15
 1439264_x_at 0.99023 LIM and SH3 protein 1 Lasp1
 1439265_at 0.93915 X Kell blood group precursor related family member 6 homolog Xkr6
 1439266_a_at 0.97306 polymerase (RNA) III (DNA directed) polypeptide K Polr3k
 1439267_x_at 0.92634 "cytochrome c oxidase, subunit Va" Cox5a
 1439268_x_at 1.04984 "eukaryotic translation initiation factor 3, subunit 6" Eif3s6
 1439269_x_at 0.92151 minichromosome maintenance deficient 7 (S. cerevisiae) /// similar to minichromosome maintenance protein 7 Mcm7 /// LOC433027
 1439270_x_at 1.04315 "RAN, member RAS oncogene family /// similar to RAN, member RAS oncogene family" Ran /// LOC671251
 1439271_x_at 0.91902 IK cytokine Ik
 1439272_at 0.91747 ligand dependent nuclear receptor corepressor-like Lcorl

1439273_at 1.00219 receptor (TNFRSF)-interacting serine-threonine kinase 1 Ripk1
1439274_at 1.23523 RIKEN cDNA A230083H22 gene A230083H22Rik
1439275_s_at 1.10120 RIKEN cDNA 9530010C24 gene 9530010C24Rik
1439276_at 1.69823 --- ---
1439277_at 0.83189 fibroblast growth factor receptor substrate 3 Frs3
1439278_at 1.01737 zinc finger and BTB domain containing 20 Zbtb20
1439279_at 1.03187 RIKEN cDNA 3110007F17 gene 3110007F17Rik
1439280_at 0.89328 RIKEN cDNA B230396O12 gene B230396O12Rik
1439281_at 0.91651 "PREDICTED: Mus musculus similar to solute carrier family 26, member 8 isoform a (LOC622293), mRNA" ---
1439282_at 1.01558 RIKEN cDNA D930023I05 gene D930023I05Rik
1439283_at 1.70572 Oxysterol binding protein-like 9 Osbpl9
1439284_at 0.98358 RIKEN cDNA E230025E14 gene E230025E14Rik
1439285_at 0.90491 inhibitor of Bruton agammaglobulinemia tyrosine kinase Ibtk
1439286_at 0.96298 "Glutamate receptor, ionotropic, kainate 2 (beta 2)" Grik2
1439287_at 0.77067 --- ---
1439288_at 0.91467 RIKEN cDNA 0710005I19 gene 0710005I19Rik
1439289_s_at 0.98856 RIKEN cDNA 0710005I19 gene 0710005I19Rik
1439290_at 1.06354 "Fragile X mental retardation gene 1, autosomal homolog" Fxr1h
1439291_at 1.01520 "Adult male corpus striatum cDNA, RIKEN full-length enriched library, clone: C030002G09 product:unclassifiable, full insert sequence" ---
1439292_at 1.40604 RIKEN cDNA 2210408F21 gene 2210408F21Rik
1439293_at 0.92136 RIKEN cDNA C130047D21 gene C130047D21Rik
1439294_at 0.96989 hypothetical protein D530008I22 D530008I22
1439295_x_at 0.95478 hypothetical LOC434179 LOC434179
1439296_at 1.01204 --- ---
1439297_at 1.12114 expressed sequence AI449441 AI449441
1439298_at 1.37843 --- ---
1439299_at 0.99858 Spectrin beta 2 Spnb2
1439300_at 1.19172 --- ---
1439301_at 2.20176 RAD51-like 1 (S. cerevisiae) Rad5111
1439302_at 0.90144 ubiquitin-activating enzyme E1-like 2 Ube112
1439303_at 0.80716 Neuronal growth regulator 1 Negr1
1439304_at 1.27073 RIKEN cDNA B230216N24 gene B230216N24Rik
1439305_at 1.42538 --- ---
1439306_at 0.98999 Small optic lobes homolog (Drosophila) Solh
1439307_at 1.98945 Cadherin 2 Cdh2
1439308_at 1.00447 glutamic-oxaloacetic transaminase 1-like 1 Got111
1439309_at 1.17719 RIKEN cDNA 9630044O09 gene 9630044O09Rik
1439310_at 1.12778 RIKEN cDNA A630012P03 gene A630012P03Rik
1439311_at 1.05679 RIKEN cDNA B830012L14 gene B830012L14Rik
1439312_at 0.88038 "Adult male corpora quadrigemina cDNA, RIKEN full-length enriched library, clone:B230337G05 product:unclassifiable, full insert sequence" ---

1439313_at 1.01254 latrophilin 3 Lphn3
1439314_at 1.29300 circadian locomoter output cycles kaput Clock
1439315_at 0.98546 Similar to shroom LOC628919
1439316_at 2.15019 RNA binding motif protein 39 Rbm39
1439317_at 0.98868 "KRIT1, ankyrin repeat containing" Krit1
1439318_at 1.05287 microtubule associated serine/threonine kinase family member 4 Mast4
1439319_at 1.17780 E74-like factor 1 Elf1
1439320_at 1.05756 cDNA sequence BC030477 BC030477
1439321_at 1.95027 AT rich interactive domain 1B (Swi1 like) Arid1b
1439322_at 0.88029 Protocadherin 9 Pcdh9
1439323_a_at 1.01209 mitogen activated protein kinase kinase kinase 1 /// similar to Mitogen-activated protein kinase kinase kinase 1 (MAPK/ERK kinase kinase 1) (MEK kinase kinase 1) (MEKKK 1) (Hematopoietic progenitor kinase) (HPK) Map4k1 /// LOC672367
1439324_at 0.91388 RIKEN cDNA D230004J03 gene /// RIKEN cDNA B430006D22 gene D230004J03Rik /// B430006D22Rik
1439325_at 1.55150 Bleomycin hydrolase Blmh
1439326_at 1.11739 leucyl/cystinyl aminopeptidase Lnpep
1439327_at 0.99447 collagen and calcium binding EGF domains 1 Ccbe1
1439328_at 1.90308 nuclear factor of activated T-cells 5 Nfat5
1439329_a_at 0.85087 BR serine/threonine kinase 2 Brsk2
1439330_at 2.56516 RIKEN cDNA D230040J21 gene D230040J21Rik
1439331_at 1.13925 RIKEN cDNA 4932439E07 gene 4932439E07Rik
1439332_at 0.86117 DNA-damage-inducible transcript 4-like Ddit4l
1439333_at 0.98296 "potassium channel, subfamily V, member 1" Kcnv1
1439334_at 0.86100 Transcribed locus ---
1439335_at 1.05066 --- ---
1439336_at 0.93313 Transcription factor 4 Tcf4
1439337_at 0.97468 target of myb1-like 1 (chicken) Tom111
1439338_at 1.01753 hypothetical protein LOC620416 LOC620417
1439339_at 0.94213 "3 days neonate thymus cDNA, RIKEN full-length enriched library, clone: A630050C01 product:unclassifiable, full insert sequence" ---
1439340_at 1.03958 RIKEN cDNA D630036G22 gene D630036G22Rik
1439341_at 1.08553 cDNA sequence AK220484 AK220484
1439342_at 1.53257 Caseinolytic peptidase X (E.coli) Clpx
1439343_at 0.96689 --- ---
1439344_at 1.25844 Transcribed locus ---
1439345_at 0.97990 glycoprotein (transmembrane) nmb Gpnmb
1439346_at 1.04736 --- ---
1439347_at 1.00596 tensin 4 Tns4
1439348_at 2.05824 S100 calcium binding protein A10 (calpactin) S100a10
1439349_at 1.53606 strawberry notch homolog (Drosophila) Stno
1439350_s_at 0.95173 CDC91 cell division cycle 91-like 1 (S. cerevisiae) Cdc9111
1439351_at 0.94886 RIKEN cDNA A930031D07 gene A930031D07Rik

1439352_at 1.07238 tripartite motif protein 7 Trim7
1439353_x_at 1.10438 transmembrane protein 56 Tmem56
1439354_at 1.01798 region containing RIKEN cDNA 2310008H09 gene; RIKEN cDNA
A230094G09 gene /// region containing RIKEN cDNA 2310008H09 gene; RIKEN cDNA A230094G09
gene LOC668373 /// LOC670145
1439355_at 0.93232 "Pleckstrin homology domain containing, family A member 5" Plekha5
1439356_at 1.14239 flightless I homolog (Drosophila) Fliih
1439357_at 0.97711 interleukin 17 receptor E Il17re
1439358_a_at 0.95053 neurexin I Nrnx1
1439359_x_at 0.92322 neurexin I Nrnx1
1439360_x_at 0.90419 "DNA segment, Chr 15, ERATO Doi 785, expressed /// similar to kleisin beta
isoform 2 /// similar to kleisin beta isoform 2" D15Ertd785e /// LOC665187 /// LOC670255
1439361_at 1.01544 Myosin IXa Myo9a
1439362_at 1.01747 "Adult male urinary bladder cDNA, RIKEN full-length enriched library,
clone:9530098H01 product:unclassifiable, full insert sequence" ---
1439363_at 1.31765 RIKEN cDNA 1200014J11 gene 1200014J11Rik
1439364_a_at 0.95480 matrix metalloproteinase 2 Mmp2
1439365_at 1.07618 Myelin transcription factor 1 Myt1
1439366_at 0.96331 G-rich RNA sequence binding factor 1 Grsf1
1439367_x_at 0.84344 ADP-ribosylation factor 4 Arf4
1439368_a_at 0.81441 "solute carrier family 9 (sodium/hydrogen exchanger), isoform 3 regulator 2"
Slc9a3r2
1439369_x_at 0.81259 "solute carrier family 9 (sodium/hydrogen exchanger), isoform 3 regulator 2"
Slc9a3r2
1439370_x_at 1.28272 "splicing factor 3b, subunit 5" Sf3b5
1439371_x_at 0.92904 translocase of inner mitochondrial membrane 44 Timm44
1439372_at 0.94094 SMT3 suppressor of mif two 3 homolog 3 (yeast) Sumo3
1439373_x_at 0.90107 wingless-related MMTV integration site 5B Wnt5b
1439374_x_at 1.02447 ribosomal protein S10 /// similar to ribosomal protein S10 /// similar to
ribosomal protein S10 /// similar to ribosomal protein S10 /// similar to ribosomal protein S10 Rps10 ///
LOC621772 /// LOC636539 /// LOC666274 /// LOC668457
1439375_x_at 1.16033 "aldolase 1, A isoform /// similar to Fructose-bisphosphate aldolase A (Muscle-
type aldolase) (Aldolase 1) /// similar to Fructose-bisphosphate aldolase A (Muscle-type aldolase)
(Aldolase 1)" Aldoa /// LOC667483 /// LOC671023
1439376_x_at 0.93024 --- ---
1439377_x_at 0.98676 cell division cycle 20 homolog (S. cerevisiae) Cdc20
1439378_at 0.95551 cDNA sequence BC048671 BC048671
1439379_x_at 0.94790 protamine 1 Prm1
1439380_x_at 1.04911 "GTL2, imprinted maternally expressed untranslated mRNA" Gtl2
1439381_x_at 0.88941 MARVEL (membrane-associating) domain containing 1 Marveld1
1439382_x_at 0.96134 "discoidin domain receptor family, member 1" Ddr1
1439383_x_at 0.95514 "protein phosphatase 2A, regulatory subunit B (PR 53)" Ppp2r4
1439384_at 1.04288 "solute carrier family 13 (sodium-dependent dicarboxylate transporter), member

3" Slc13a3
1439385_x_at 1.03442 "Solute carrier family 13 (sodium-dependent dicarboxylate transporter), member 3" Slc13a3
1439386_x_at 0.90739 "methionine adenosyltransferase II, alpha" Mat2a
1439387_x_at 1.00521 RIKEN cDNA 2310061F22 gene 2310061F22Rik
1439388_s_at 1.02064 breast cancer anti-estrogen resistance 1 Bcar1
1439389_s_at 1.05332 myeloid-associated differentiation marker Myadm
1439390_at 0.93440 RIKEN cDNA 1300018I17 gene 1300018I17Rik
1439391_at 0.85157 RIKEN cDNA 2810002I04 gene 2810002I04Rik
1439392_x_at 1.03627 "protease (prosome, macropain) 26S subunit, ATPase 1 /// hypothetical gene supported by NM_008947" Psmc1 /// LOC386042
1439393_x_at 0.94386 "protein phosphatase 2A, regulatory subunit B (PR 53)" Ppp2r4
1439394_x_at 0.87031 cell division cycle 20 homolog (S. cerevisiae) Cdc20
1439395_at 0.94915 RIKEN cDNA 9630025I21 gene 9630025I21Rik
1439396_x_at 0.77236 glycerol-3-phosphate dehydrogenase 1 (soluble) Gpd1
1439397_at 0.89213 formin 1 Fmn1
1439398_x_at 0.87427 nasal embryonic LHRH factor Nelf
1439399_a_at 0.96188 "small nucleolar RNA, C/D box 22" Snord22
1439400_x_at 0.95705 RIKEN cDNA 5430433E21 gene 5430433E21Rik
1439401_x_at 0.97585 "Protein phosphatase 2, regulatory subunit B (B56), epsilon isoform" Ppp2r5e
1439402_at 1.04724 --- ---
1439403_x_at 1.00026 ring finger protein 12 Rnf12
1439404_x_at 0.90884 "zinc finger protein, autosomal /// zinc finger protein X-linked" Zfa /// Zfx
1439405_x_at 0.90277 "Beclin 1 (coiled-coil, myosin-like BCL2-interacting protein)" Becn1
1439406_x_at 1.04462 phenylalanine-tRNA synthetase 2 (mitochondrial) Fars2
1439407_x_at 1.06909 transgelin 2 /// similar to transgelin 2 Tagln2 /// LOC672466
1439408_a_at 1.07133 peroxisome proliferator activated receptor binding protein Pparbp
1439409_x_at 1.13686 tyrosinase-related protein 1 Tyrp1
1439410_x_at 0.87546 "DNA segment, Chr 11, ERATO Doi 333, expressed" D11Ertd333e
1439411_a_at 0.93955 exportin 7 Xpo7
1439412_at 0.96891 "Cyclic AMP-regulated phosphoprotein, 21" Arpp21
1439413_x_at 1.02016 mortality factor 4 like 2 Morf4l2
1439414_x_at 1.34630 "Sno, strawberry notch homolog 1 (Drosophila)" Sbnol
1439415_x_at 0.93041 similar to 40S ribosomal protein S21 LOC622373
1439416_x_at 1.01334 "tubulin, beta 2c" Tubb2c
1439417_at 0.91168 Quiescin Q6 Qscn6
1439418_x_at 0.88892 transmembrane protein 43 Tmem43
1439419_at 1.26302 RIKEN cDNA 2310057M21 gene 2310057M21Rik
1439420_x_at 0.93688 gamma-glutamyltransferase-like activity 1 Ggtla1
1439421_x_at 0.97630 chromobox homolog 3 (Drosophila HP1 gamma) /// similar to chromobox homolog 3 Cbx3 /// LOC633016
1439422_a_at 1.17519 C1q domain containing 2 C1qdc2
1439423_x_at 1.05304 cDNA sequence U46068 U46068

1439424_x_at 0.94074 HERPUD family member 2 Herpud2
1439425_x_at 1.08778 cDNA sequence BC024814 BC024814
1439426_x_at 1.00781 P lysozyme structural Lzp-s
1439427_at 0.83997 claudin 9 Cldn9
1439428_x_at 0.95646 "GDP-mannose 4, 6-dehydratase" Gmds
1439429_x_at 0.83808 deltex 2 homolog (Drosophila) Dtx2
1439430_x_at 1.11091 RIKEN cDNA 4930403C10 gene 4930403C10Rik
1439431_x_at 1.04397 --- ---
1439432_x_at 0.92461 mortality factor 4 like 2 /// similar to Mortality factor 4-like protein 2 (MORF-related gene X protein) (Transcription factor-like protein MRGX) (Sid 393) /// similar to Mortality factor 4-like protein 2 (MORF-related gene X protein) (Transcription factor-like protein MRGX) (Sid 393) Morf4l2 /// LOC433261 /// LOC677515
1439433_a_at 0.88933 "solute carrier family 35 (UDP-galactose transporter), member 2" Slc35a2
1439434_x_at 0.93772 SH2 domain containing 5 Sh2d5
1439435_x_at 1.28160 phosphoglycerate kinase 1 /// phosphoglycerate kinase 1-like /// phosphoglycerate kinase 1 pseudogene /// phosphoglycerate kinase 1-like Pgk1 /// LOC433594 /// LOC665053 /// LOC668435
1439436_x_at 1.00076 inner centromere protein Incenp
1439437_x_at 1.10512 carboxypeptidase E /// similar to carboxypeptidase E Cpe /// LOC677374
1439438_a_at 0.96087 RIKEN cDNA 1110005A23 gene /// similar to cytokine induced protein 29 kDa /// similar to cytokine induced protein 29 kDa 1110005A23Rik /// LOC625193 /// LOC636633
1439439_x_at 0.99776 eukaryotic translation elongation factor 1 delta (guanine nucleotide exchange protein) Eef1d
1439440_x_at 0.97579 protein tyrosine kinase 9-like (A6-related protein) Ptk9l
1439441_x_at 0.91557 large tumor suppressor 2 Lats2
1439442_x_at 1.09549 tyrosyl-tRNA synthetase 2 (mitochondrial) Yars2
1439443_x_at 0.98891 transketolase Tkt
1439444_x_at 1.06724 transmembrane emp24-like trafficking protein 10 (yeast) /// similar to transmembrane trafficking protein Tmed10 /// LOC634748
1439445_x_at 1.04781 ATP citrate lyase Acly
1439446_at 1.05868 cDNA sequence BC048507 BC048507
1439447_x_at 1.03696 ribosomal protein L37a /// similar to 60S ribosomal protein L37a /// similar to 60S ribosomal protein L37a /// similar to 60S ribosomal protein L37a /// similar to 60S ribosomal protein L37a /// similar to 60S ribosomal protein L37a Rpl37a /// LOC629949 /// LOC632394 /// LOC632599 /// LOC640991 /// LOC668935 /// LOC668941 /// LOC669803 /// LOC674834
1439448_x_at 0.97775 transmembrane emp24 protein transport domain containing 9 Tmed9
1439449_at 0.90628 special AT-rich sequence binding protein 1 Satb1
1439450_x_at 1.21804 RIKEN cDNA A230046K03 gene A230046K03Rik
1439451_x_at 0.90285 G protein-coupled receptor 172B Gpr172b
1439452_x_at 0.89649 aspartyl aminopeptidase Dnpep
1439453_x_at 0.96960 RIKEN cDNA 1500026D16 gene 1500026D16Rik

1439454_x_at 0.87773 TM2 domain containing 2 Tm2d2
1439455_x_at 0.95063 "capping protein (actin filament) muscle Z-line, alpha 1" Capza1
1439456_x_at 0.98884 "ATPase, H+ transporting, lysosomal accessory protein 2" Atp6ap2
1439457_x_at 1.02191 autophagy-related 12 (yeast) Atg12
1439458_x_at 1.02759 CEA-related cell adhesion molecule 11 Ceacam11
1439459_x_at 1.23013 ATP citrate lyase Acly
1439460_a_at 0.96936 zinc finger protein 289 Zfp289
1439461_x_at 0.89991 RIKEN cDNA 2410003A14 gene 2410003A14Rik
1439462_x_at 1.01555 transmembrane emp24-like trafficking protein 10 (yeast) /// similar to transmembrane trafficking protein Tmed10 /// LOC634748
1439463_x_at 0.94119 high mobility group box 1 /// similar to High mobility group protein 1 (HMG-1) (High mobility group protein B1) (Amphoterin) (Heparin-binding protein p30) /// similar to High mobility group protein 1 (HMG-1) (High mobility group protein B1) (Amphoterin) (Heparin-binding protein p30) /// similar to High mobility group protein 1 (HMG-1) (High mobility group protein B1) (Amphoterin) (Heparin-binding protein p30) /// similar to High mobility group protein 1 (HMG-1) (High mobility group protein B1) (Amphoterin) (Heparin-binding protein p30) /// similar to High mobility group protein 1 (HMG-1) (High mobility group protein B1) (Amphoterin) (Heparin-binding protein p30) /// similar to High mobility group protein 1 (HMG-1) (High mobility group protein B1) (Amphoterin) (Heparin-binding protein p30) /// similar to High mobility group protein 1 (HMG-1) (High mobility group protein B1) (Amphoterin) (Heparin-binding protein p30) /// similar to High mobility group protein 1 (HMG-1) (High mobility group protein B1) (Amphoterin) (Heparin-binding protein p30) /// similar to High mobility group protein 1 (HMG-1) (High mobility group protein B1) (Amphoterin) (Heparin-binding protein p30) Hmgb1 /// LOC434174 /// LOC545555 /// LOC623634 /// LOC637733 /// LOC640060 /// LOC666836 /// LOC669546 /// LOC669692 /// LOC671507 /// LOC677578
1439464_s_at 1.04857 testis expressed gene 10 Tex10
1439465_x_at 0.83251 RIKEN cDNA 9430057O19 gene 9430057O19Rik
1439466_s_at 1.01034 BUD31 homolog (yeast) Bud31
1439467_at 1.15146 Microtubule-associated protein 4 Mtap4
1439468_at 1.01340 BTB and CNC homology 2 Bach2
1439469_at 0.99550 "Leucine rich repeat containing 8 family, member C" Lrrc8c
1439470_at 1.40252 RIKEN cDNA 2410025L10 gene 2410025L10Rik
1439471_at 1.09147 RIKEN cDNA 2610305J24 gene 2610305J24Rik
1439472_at 0.96196 GCN1 general control of amino-acid synthesis 1-like 1 (yeast) Gcn111
1439473_at 1.11763 Pseudouridylate synthase 7 homolog (S. cerevisiae) Pus7
1439474_x_at 0.93593 RIKEN cDNA B230343H07 gene Nell1
1439475_at 1.22572 expressed sequence AI929863 AI929863
1439476_at 0.84368 desmoglein 2 Dsg2
1439477_at 1.64079 "ubiquitin-conjugating enzyme E2B, RAD6 homology (S. cerevisiae)" Ube2b
1439478_at 0.80036 acyl-CoA thioesterase 2 Acot2
1439479_at 1.03003 lactase Lct

1439480_at 1.11833 Expressed sequence AU015850 AU015850
1439481_at 1.09088 importin 9 Ipo9
1439482_at 1.03251 RIKEN cDNA F830021D11 gene F830021D11Rik
1439483_at 1.02222 expressed sequence AI506816 AI506816
1439484_at 1.05670 RIKEN cDNA 4930431H11 gene 4930431H11Rik
1439485_at 0.81223 zinc finger protein 608 /// RIKEN cDNA D430007A19 gene Zfp608 ///
D430007A19Rik
1439486_at 1.00844 "potassium channel, subfamily T, member 1" Kcnt1
1439487_at 0.92029 "ligase IV, DNA, ATP-dependent" Lig4
1439488_at 0.94230 "7 days embryo whole body cDNA, RIKEN full-length enriched library, clone:
C430040N23 product:unclassifiable, full insert sequence" ---
1439489_at 0.93748 G protein-coupled receptor 120 Gpr120
1439490_at 0.77266 "nuclear receptor subfamily 2, group C, member 1" Nr2c1
1439491_at 0.94549 leucine rich repeat containing 38 Lrrc38
1439492_at 1.23568 RIKEN cDNA C130053K05 gene C130053K05Rik
1439493_at 0.91794 RIKEN cDNA D630040G17 gene /// similar to zinc finger protein 64
(predicted) D630040G17Rik /// LOC622675
1439494_at 0.99804 "solute carrier family 5 (sodium/glucose cotransporter), member 9" Slc5a9
1439495_at 0.79133 RIKEN cDNA 4933407H18 gene 4933407H18Rik
1439496_at 1.08388 RIKEN cDNA 4921524J06 gene 4921524J06Rik
1439497_at 1.29687 RIKEN cDNA 4933415E08 gene 4933415E08Rik
1439498_at 1.02845 putative neuronal cell adhesion molecule Punc
1439499_at 0.95349 expressed sequence AA415398 AA415398
1439500_at 1.01459 --- ---
1439501_at 1.03035 calcium binding atopy-related autoantigen 1 Cbaral
1439502_at 1.02928 RIKEN cDNA 4930573I19 gene 4930573I19Rik
1439503_at 1.15475 zinc finger protein 28 Zfp28
1439504_s_at 0.95582 zinc finger protein 28 Zfp28
1439505_at 0.89412 chloride intracellular channel 5 Clic5
1439506_at 0.78644 "gene model 98, (NCBI)" Gm98
1439507_at 0.98159 TRAF type zinc finger domain containing 1 Traf1
1439508_at 0.92646 RIKEN cDNA A730055L17 gene A730055L17Rik
1439509_at 0.94115 RIKEN cDNA 2900008C10 gene 2900008C10Rik
1439510_at 0.98668 shugoshin-like 1 (S. pombe) Sgol1
1439511_at 0.80298 Cyclin-dependent kinase 7 (homolog of Xenopus MO15 cdk-activating
kinase) Cdk7
1439512_at 0.93279 RIKEN cDNA 5830444B04 gene 5830444B04Rik
1439513_at 1.00432 proline-rich nuclear receptor coactivator 1 Pnrc1
1439514_at 0.89018 CDC28 protein kinase 1b Cks1b
1439515_at 1.17243 SET domain containing 5 Setd5
1439516_at 1.33050 RIKEN cDNA 2610201A13 gene 2610201A13Rik
1439517_at 0.87505 "myb-like, SWIRM and MPN domains 1" Mysm1
1439518_at 1.01065 multimerin 2 Mmrn2

1439519_at 1.00938 "solute carrier family 34 (sodium phosphate), member 3" Slc34a3
1439520_at 1.86622 --- ---
1439521_at 1.03148 "Adult male aorta and vein cDNA, RIKEN full-length enriched library, clone:
A530001O10 product:unclassifiable, full insert sequence" ---
1439522_at 0.90399 --- ---
1439523_at 0.98895 RIKEN cDNA D330027G24 gene D330027G24Rik
1439524_at 0.93873 "DnaJ (Hsp40) homolog, subfamily B, member 4" Dnajb4
1439525_at 0.91086 RIKEN cDNA 1700066D14 gene 1700066D14Rik
1439526_at 1.02820 "Adult male cecum cDNA, RIKEN full-length enriched library,
clone:9130022D06 product:unclassifiable, full insert sequence" ---
1439527_at 0.83939 Expressed sequence BB114106 BB114106
1439528_at 0.97730 RIKEN cDNA 4833423E24 gene 4833423E24Rik
1439529_at 0.91946 RIKEN cDNA A430110N23 gene A430110N23Rik
1439530_a_at 0.98131 similar to NFkB interacting protein 1 LOC669005
1439531_at 1.00022 RIKEN cDNA E130311K13 gene E130311K13Rik
1439532_s_at 0.77317 kinesin family member 13A Kif13a
1439533_at 1.03818 RIKEN cDNA 1700125D06 gene 1700125D06Rik
1439534_at 0.96306 zinc finger protein 248 Zfp248
1439535_at 1.12178 --- ---
1439536_at 0.89297 Expressed sequence AI846133 AI846133
1439537_at 2.67432 --- ---
1439538_at 0.95085 RIKEN cDNA 0610011N22 gene 0610011N22Rik
1439539_at 1.00534 translocating chain-associating membrane protein 2 Tram2
1439540_at 1.03125 membrane-associated ring finger (C3HC4) 2 2-Mar
1439541_at 0.83707 RIKEN cDNA 4930414L22 gene 4930414L22Rik
1439542_at 0.88856 zinc finger protein 651 Zfp651
1439543_at 0.90233 RIKEN cDNA 1110064A23 gene 1110064A23Rik
1439544_at 0.91924 "TBC1 domain family, member 5" Tbc1d5
1439545_at 1.09546 Nuclear respiratory factor 1 Nrf1
1439546_at 0.97668 RIKEN cDNA 4933417O08 gene 4933417O08Rik
1439547_at 0.95313 RIKEN cDNA A930007A09 gene A930007A09Rik
1439548_at 1.09096 "RAP2B, member of RAS oncogene family" Rap2b
1439549_at 1.04596 proline rich Gla (G-carboxyglutamic acid) 3 (transmembrane) Prrg3
1439550_at 1.13249 --- ---
1439551_at 0.88243 cDNA sequence BC006965 BC006965
1439552_at 0.96772 triple functional domain (PTPRF interacting) Trio
1439553_s_at 1.02986 nuclear transport factor 2 /// similar to Nuclear transport factor 2 (NTF-2) ///
similar to Nuclear transport factor 2 (NTF-2) /// similar to Nuclear transport factor 2 (NTF-2) /// similar
to Nuclear transport factor 2 (NTF-2) Nutf2 /// LOC621832 /// LOC668830 /// LOC670887 ///
LOC675580
1439554_at 1.10979 sex comb on midleg homolog 1 Scmh1
1439555_at 0.94979 rearranged L-myc fusion sequence Rlf
1439556_at 0.98961 neural cell adhesion molecule 1 Ncam1

1439557_s_at 0.93048 LIM domain binding 2 Ldb2
1439558_at 0.93508 zinc finger protein 75 Zfp75
1439559_at 0.93194 RIKEN cDNA 1700040D17 gene 1700040D17Rik
1439560_x_at 0.92448 hypothetical gene supported by BC047216 LOC432995
1439561_at 0.95131 RIKEN cDNA 2010012O05 gene 2010012O05Rik
1439562_at 0.94168 RIKEN cDNA F730047E07 gene F730047E07Rik
1439563_at 0.91528 zinc finger protein 3 Zfp3
1439564_at 0.94723 RIKEN cDNA 1700090G07 gene 1700090G07Rik
1439565_at 0.87159 Expressed sequence AW492303 AW492303
1439566_at 0.91304 RIKEN cDNA C030038J10 gene C030038J10Rik
1439567_at 0.97365 T-box 3 Tbx3
1439568_at 1.06239 gene regulated by estrogen in breast cancer protein Greb1
1439569_at 0.82870 G protein-coupled receptor 83 Gpr83
1439570_at 1.02301 "gene model 444, (NCBI)" Gm444
1439571_at 1.08122 RIKEN cDNA E230008J23 gene E230008J23Rik
1439572_at 2.24631 --- ---
1439573_at 1.03775 reticulon 4 receptor-like 2 Rtn4rl2
1439574_at 0.88490 RIKEN cDNA 2600015P10 gene 2600015P10Rik
1439575_at 1.10673 RIKEN cDNA E130009J12 gene E130009J12Rik
1439576_at 1.00295 Transcribed locus ---
1439577_at 0.76919 --- ---
1439578_at 1.00795 U7 snRNP-specific Sm-like protein LSM11 Lsm11
1439579_at 0.97505 hypothetical LOC433408 LOC433408
1439580_at 0.88827 RIKEN cDNA B930053N05 gene B930053N05Rik
1439581_at 0.99227 RIKEN cDNA 6330403L08 gene 6330403L08Rik
1439582_at 1.54258 Microtubule-actin crosslinking factor 1 Macf1
1439583_x_at 0.83920 RIKEN cDNA D130029J02 gene D130029J02Rik
1439584_at 1.02514 zinc finger protein 526 Zfp526
1439585_at 1.04118 midasin homolog (yeast) Mdn1
1439586_at 1.42154 hypothetical LOC548102 LOC548102
1439587_at 1.06035 LEM domain containing 3 Lemd3
1439588_at 0.98029 "solute carrier organic anion transporter family, member 5A1" Slco5a1
1439589_at 0.88946 CD53 antigen Cd53
1439590_at 0.95863 X Kell blood group precursor related family member 8 homolog Xkr8
1439591_at 1.20027 Unc-51 like kinase 2 (C. elegans) Ulk2
1439592_at 1.03045 RIKEN cDNA C030010B13 gene C030010B13Rik
1439593_s_at 0.89000 RIKEN cDNA C030010B13 gene C030010B13Rik
1439594_at 1.09793 RIKEN cDNA D130062J10 gene D130062J10Rik
1439595_at 1.04824 RIKEN cDNA A430107P09 gene A430107P09Rik
1439596_at 1.08588 --- ---
1439597_at 1.11808 RIKEN cDNA 4932417H02 gene 4932417H02Rik
1439598_at 1.25533 Forkhead box O3a Foxo3a
1439599_at 0.95862 galactose-3-O-sulfotransferase 2 Gal3st2

1439600_at 1.09615 CASP8 and FADD-like apoptosis regulator Cflar
1439601_at 1.02890 RIKEN cDNA 2700008B19 gene 2700008B19Rik
1439602_at 1.03806 "10 days neonate cerebellum cDNA, RIKEN full-length enriched library, clone: B930049N01 product:unclassifiable, full insert sequence" ---
1439603_at 1.12343 "Adult male hypothalamus cDNA, RIKEN full-length enriched library, clone: A230032A12 product:unclassifiable, full insert sequence" ---
1439604_at 0.97727 "a disintegrin-like and metalloproteinase (reprolysin type) with thrombospondin type 1 motif, 16" Adamts16
1439605_at 1.02469 --- ---
1439606_at 0.93987 Transcribed locus ---
1439607_at 0.90747 --- ---
1439608_at 0.81462 Transcribed locus ---
1439609_at 1.01071 Transcribed locus ---
1439610_at 0.81600 "RAB27b, member RAS oncogene family" Rab27b
1439611_at 0.92326 "cholinergic receptor, muscarinic 1, CNS" Chrm1
1439612_at 0.90049 "calcium channel, voltage-dependent, N type, alpha 1B subunit" Cacna1b
1439613_at 1.01329 soluble adenylyl cyclase Sacy
1439614_at 1.08069 5 hydroxytryptamine (serotonin) receptor 4 Htr4
1439615_at 1.03114 giant axonal neuropathy Gan
1439616_at 1.14529 Transcribed locus ---
1439617_s_at 1.02139 "phosphoenolpyruvate carboxykinase 1, cytosolic" Pck1
1439618_at 1.07782 phosphodiesterase 10A Pde10a
1439619_at 1.12121 transcription factor 12 Tcf12
1439620_at 1.02354 carbonic anhydrase 13 Car13
1439621_at 0.86987 "17 days embryo kidney cDNA, RIKEN full-length enriched library, clone: I920178P20 product:unclassifiable, full insert sequence" ---
1439622_at 1.24929 Ras association (RalGDS/AF-6) domain family 4 Rassf4
1439623_at 1.03795 OTU domain containing 1 Otud1
1439624_at 0.62714 "UDP glucuronosyltransferase 2 family, polypeptide B35" Ugt2b35
1439625_at 0.98575 Keratin associated protein 5-5 Krtap5-5
1439626_at 0.94470 tropomodulin 3 Tmod3
1439627_at 0.90001 zinc finger protein of the cerebellum 1 Zic1
1439628_x_at 0.93371 "Rab38, member of RAS oncogene family" Rab38
1439629_at 1.01697 neurexin III Nrnx3
1439630_x_at 0.90461 suprabasin Sbsn
1439631_at 1.33838 --- ---
1439632_at 0.88819 "12 days embryo spinal ganglion cDNA, RIKEN full-length enriched library, clone: D130046A02 product:unclassifiable, full insert sequence" ---
1439633_at 1.04103 RIKEN cDNA B230112P13 gene B230112P13Rik
1439634_at 0.94545 RIKEN cDNA 4930505D03 gene 4930505D03Rik
1439635_at 0.97801 regulator of G-protein signaling 9 Rgs9
1439636_at 1.07379 "NACHT, LRR and PYD containing protein 9c" Nalp9c
1439637_at 0.88943 kinesin family member 7 Kif7

1439638_at 3.09105 Erbb2 interacting protein Erbb2ip
1439639_at 1.50287 Polyhomeotic-like 2 (Drosophila) Phc2
1439640_at 1.01346 "16 days neonate thymus cDNA, RIKEN full-length enriched library, clone:
A130046K21 product:hypothetical protein, full insert sequence" ---
1439641_at 0.96445 hypothetical LOC553090 LOC553090
1439642_at 0.90473 RIKEN cDNA 5930403N24 gene 5930403N24Rik
1439643_at 2.86074 Activin receptor IIA Acvr2a
1439644_at 0.89385 Early B-cell factor 1 Ebf1
1439645_at 1.08045 "adrenergic receptor, alpha 2b" Adra2b
1439646_at 0.89302 "Adult male pituitary gland cDNA, RIKEN full-length enriched library,
clone:5330431D23 product:unclassifiable, full insert sequence" ---
1439647_at 0.99861 hypothetical LOC545611 LOC545611
1439648_at 0.91789 "anillin, actin binding protein (scraps homolog, Drosophila)" Anln
1439649_at 0.86800 arginine decarboxylase Adc
1439650_at 2.00418 reticulon 4 Rtn4
1439651_at 1.04002 LIM domain only 4 Lmo4
1439652_at 2.12963 --- ---
1439653_at 1.13672 a disintegrin and metallopeptidase domain 6 Adam6
1439654_at 1.02483 hypothetical LOC319225 /// hypothetical protein LOC633385
LOC319225 /// LOC633385
1439655_at 1.22106 Ubiquitin-conjugating enzyme E2D 2 Ube2d2
1439656_at 1.52367 "Platelet-activating factor acetylhydrolase, isoform 1b, beta1 subunit" Pafah1b1
1439657_at 1.12150 hypothetical LOC319980 LOC319980
1439658_at 1.00399 leiomodulin 3 (fetal) Lmod3
1439659_at 1.01584 Trf (TATA binding protein-related factor)-proximal protein homolog
(Drosophila) /// ubiquitin specific peptidase 49 Trfp /// Usp49
1439660_at 0.87484 human immunodeficiency virus type I enhancer binding protein 3 Hivep3
1439661_at 1.04593 "solute carrier family 16 (monocarboxylic acid transporters), member 14"
Slc16a14
1439662_at 1.16154 homer homolog 1 (Drosophila) Homer1
1439663_at 1.06429 Patched homolog 1 Ptch1
1439664_at 1.43506 WD repeat domain 68 Wdr68
1439665_at 0.97781 G protein-coupled receptor 23 Gpr23
1439666_at 0.96486 RIKEN cDNA 2010001J22 gene 2010001J22Rik
1439667_at 1.02529 --- ---
1439668_at 1.02175 "Adult male hypothalamus cDNA, RIKEN full-length enriched library, clone:
A230035D19 product:hypothetical protein, full insert sequence" ---
1439669_at 0.97746 RIKEN cDNA 6430571L13 gene 6430571L13Rik
1439670_at 1.13500 RIKEN cDNA B020006M18 gene B020006M18Rik
1439671_at 0.76427 RIKEN cDNA 4930466K18 gene 4930466K18Rik
1439672_at 1.19330 Synaptojanin 1 Synj1
1439673_at 1.73454 RIKEN cDNA E230008O15 gene E230008O15Rik
1439674_at 0.91261 "solute carrier family 4 (anion exchanger), member 8" Slc4a8

1439675_at 1.02234 RIKEN cDNA 4933429D07 gene 4933429D07Rik
1439676_at 1.19289 apoptosis antagonizing transcription factor Aatf
1439677_at 0.96409 "epilepsy, progressive myoclonic epilepsy, type 2 gene alpha" Epm2a
1439678_at 2.54301 Glutaminase Gls
1439679_at 1.07733 chaperonin subunit 3 (gamma) Cct3
1439680_at 1.00442 "tumor necrosis factor (ligand) superfamily, member 10" Tnfsf10
1439681_at 1.03995 "FERM, RhoGEF and pleckstrin domain protein 2" Farp2
1439682_at 0.99582 --- ---
1439683_at 0.88096 "In vitro fertilized eggs cDNA, RIKEN full-length enriched library,
clone:7420434E04 product:unclassifiable, full insert sequence" ---
1439684_at 0.92762 RIKEN cDNA 4930570G19 gene 4930570G19Rik
1439685_at 1.19982 Ring finger protein 11 Rnf11
1439686_at 1.09335 Muscleblind-like 1 (Drosophila) Mbnl1
1439687_at 2.06667 "RAB14, member RAS oncogene family" Rab14
1439688_at 1.04472 fibulin 1 Fbln1
1439689_at 0.96655 RIKEN cDNA 9030224M15 gene 9030224M15Rik
1439690_at 1.08014 RIKEN cDNA 5133401N09 gene 5133401N09Rik
1439691_at 1.77355 "DNA segment, Chr 5, ERATO Doi 579, expressed" D5Ert579e
1439692_at 1.03274 "16 days neonate cerebellum cDNA, RIKEN full-length enriched library,
clone:9630042A05 product:unclassifiable, full insert sequence" ---
1439693_a_at 0.94982 F-box and WD-40 domain protein 16 Fbxw16
1439694_at 1.16766 F-box and WD-40 domain protein 16 Fbxw16
1439695_a_at 1.16016 M-phase phosphoprotein 1 Mphosph1
1439696_at 1.10197 "nuclear receptor subfamily 2, group C, member 2" Nr2c2
1439697_at 1.05068 interleukin 1 receptor accessory protein Il1rap
1439698_at 0.94194 Zinc finger protein (C2H2 type) 276 Zfp276
1439699_at 1.05464 Progesterone receptor Pgr
1439700_at 0.99992 A disintegrin and metallopeptidase domain 1b Adam1b
1439701_at 1.09901 RIKEN cDNA 9330121K16 gene 9330121K16Rik
1439702_at 0.92898 --- ---
1439703_at 1.03368 CD200 receptor 1 Cd200r1
1439704_at 1.24160 histone deacetylase 2 Hdac2
1439705_at 1.40209 "Protein tyrosine phosphatase, non-receptor type 12" Ptpn12
1439706_at 1.05446 RIKEN cDNA A330106F07 gene A330106F07Rik
1439707_at 0.90520 RIKEN cDNA C230034O21 gene C230034O21Rik
1439708_at 1.09864 "myomesin family, member 3" Myom3
1439709_at 0.99853 --- ---
1439710_at 1.00141 nuclear receptor coactivator 6 interacting protein Ncoa6ip
1439711_at 1.07873 --- ---
1439712_at 1.26685 integrator complex subunit 10 Ints10
1439713_at 0.96395 integrin alpha 1 Itga1
1439714_at 0.91668 --- ---
1439715_at 1.15470 O-sialoglycoprotein endopeptidase-like 1 Osgepl1

1439716_at	0.96367 piwi-like homolog 4 (Drosophila)	Piwil4	
1439717_at	0.97020 RIKEN cDNA B230362M20 gene	B230362M20Rik	
1439718_at	0.93312 "Protein kinase inhibitor, gamma"	Pkig	
1439719_at	0.97197 RIKEN cDNA E430004N04 gene	E430004N04Rik	
1439720_at	1.05032 Ral GEF with PH domain and SH3 binding motif 1	Ralgps1	
1439721_at	0.86170 expressed sequence C85627	C85627	
1439722_at	1.40771 --- ---		
1439723_at	0.95568 RIKEN cDNA 5730590G19 gene	5730590G19Rik	
1439724_at	0.92815 resistance to inhibitors of cholinesterase 8 homolog B (C. elegans)	Ric8b	
1439725_at	1.04699 "protein tyrosine phosphatase, receptor type, T"	Ptptrt	
1439726_at	1.25037 RIKEN cDNA 4432406C05 gene	4432406C05Rik	
1439727_at	0.99738 chloride channel calcium activated 6 /// expressed sequence AI747448		
	Clca6 /// AI747448		
1439728_at	0.92319 RIKEN cDNA D330027H18 gene	D330027H18Rik	
1439729_at	0.98531 RIKEN cDNA A930038B10 gene	A930038B10Rik	
1439730_at	0.95060 Neurocalcin delta	Ncald	
1439731_at	0.93274 RIKEN cDNA E130309F12 gene	E130309F12Rik	
1439732_at	0.96286 "16 days neonate cerebellum cDNA, RIKEN full-length enriched library, clone:9630041I08 product:unclassifiable, full insert sequence"	---	
1439733_at	0.98128 "13 days embryo stomach cDNA, RIKEN full-length enriched library, clone: D530023N09 product:unclassifiable, full insert sequence"	---	
1439734_at	0.73580 Matrix metalloproteinase 15	Mmp15	
1439735_at	0.92092 "Adult male testis cDNA, RIKEN full-length enriched library, clone:4933414F24 product:unclassifiable, full insert sequence"	---	
1439736_at	0.94571 RIKEN cDNA 5830453J16 gene	5830453J16Rik	
1439737_x_at	0.99965 "Mediator of RNA polymerase II transcription, subunit 6 homolog (yeast)"	Med6	
1439738_at	0.99203 --- ---		
1439739_at	0.99877 RIKEN cDNA 1700112C13 gene /// testes-specific protease 50	1700112C13Rik /// Tsp50	
1439740_s_at	1.06116 uridine-cytidine kinase 2	Uck2	
1439741_x_at	1.01775 uridine-cytidine kinase 2	Uck2	
1439742_at	0.88732 Signal transducing adaptor molecule (SH3 domain and ITAM motif) 2	Stam2	
1439743_at	0.96276 expressed sequence AW048948	AW048948	
1439744_at	0.81154 C-src tyrosine kinase	Csk	
1439745_at	0.90047 "calcium channel, voltage-dependent, gamma subunit 7"	Cacng7	
1439746_at	1.00289 dual specificity phosphatase 27 (putative)	Dusp27	
1439747_at	0.94309 prostaglandin E synthase	Ptges	
1439748_at	0.97421 dipeptidylpeptidase 6	Dpp6	
1439749_at	1.04745 zeta-chain (TCR) associated protein kinase	Zap70	
1439750_at	0.95899 Transcribed locus	---	
1439751_at	1.03427 zinc finger protein 708	Zfp708	
1439752_at	0.97793 "PREDICTED: Mus musculus similar to O-acyltransferase (membrane bound)		

domain containing 1 (LOC234155), mRNA" ---

1439753_x_at 1.07829 sine oculis-related homeobox 4 homolog (Drosophila) Six4
 1439754_at 0.91373 SRY-box containing gene 12 Sox12
 1439755_at 0.84379 --- ---
 1439756_at 0.86501 RIKEN cDNA 4933406J04 gene 4933406J04Rik
 1439757_s_at 1.01271 Eph receptor A4 Epha4
 1439758_at 0.97362 "amyotrophic lateral sclerosis 2 (juvenile) chromosome region, candidate 12 (human)" Als2cr12
 1439759_x_at 0.98972 RIKEN cDNA 2410078J06 gene 2410078J06Rik
 1439760_x_at 0.87493 uroplakin 1B Upk1b
 1439761_x_at 0.96204 RIKEN cDNA D830026I12 gene D830026I12Rik
 1439762_x_at 0.97093 "adrenergic receptor, alpha 2c" Adra2c
 1439763_at 0.88603 "DnaJ (Hsp40) homolog, subfamily C, member 11" Dnajc11
 1439764_s_at 1.23973 insulin-like growth factor 2 mRNA binding protein 2 Igf2bp2
 1439765_x_at 1.00134 keratin 42 Krt42
 1439766_x_at 0.76335 vascular endothelial growth factor C Vegfc
 1439767_at 0.96796 "discs, large (Drosophila) homolog-associated protein 2" Dlgap2
 1439768_x_at 0.85288 "sema domain, immunoglobulin domain (Ig), TM domain, and short cytoplasmic domain" Sema4f
 1439769_at 1.02994 Zinc finger protein 644 Zfp644
 1439770_at 0.86182 RIKEN cDNA 6430598A04 gene 6430598A04Rik
 1439771_s_at 1.01435 "DNA segment, Chr 13, ERATO Doi 608, expressed" D13Ertd608e
 1439772_at 1.10554 "DNA segment, Chr 1, Brigham & Women's Genetics 0212 expressed" D1Bwg0212e
 1439773_at 0.94159 "lymphocyte antigen 6 complex, locus E" Ly6e
 1439774_at 1.10020 paired related homeobox 1 Prrx1
 1439775_at 0.88511 bromodomain and WD repeat domain containing 3 Brwd3
 1439776_at 0.99749 2-oxoglutarate and iron-dependent oxygenase domain containing 1 Ogfod1
 1439777_at 0.89091 hypothetical protein B230218O03 B230218O03
 1439778_at 1.03074 Cdk5 and Abl enzyme substrate 1 Cables1
 1439779_at 0.97398 "16 days neonate cerebellum cDNA, RIKEN full-length enriched library, clone:9630050J22 product:unclassifiable, full insert sequence" ---
 1439780_at 1.01365 ribosomal protein L7-like 1 Rpl7l1
 1439781_at 0.98365 RIKEN cDNA D930048N14 gene D930048N14Rik
 1439782_s_at 1.08841 RIKEN cDNA D930048N14 gene D930048N14Rik
 1439783_at 0.84306 RIKEN cDNA C330018D20 gene C330018D20Rik
 1439784_at 0.87036 RIKEN cDNA 2310046O06 gene 2310046O06Rik
 1439785_at 0.99072 RIKEN cDNA 9630013A20 gene 9630013A20Rik
 1439786_at 1.05650 growth factor receptor bound protein 2-associated protein 2 Gab2
 1439787_at 0.95967 "purinergic receptor P2X, ligand-gated ion channel, 7" P2rx7
 1439788_at 0.68701 coiled-coil domain containing 111 Ccdc111
 1439789_at 1.02310 --- ---
 1439790_at 1.07424 "serine (or cysteine) peptidase inhibitor, clade B, member 9" Serpinb9

1439791_at 0.92176 "gene model 527, (NCBI)" Gm527
 1439792_at 1.01518 "Transcribed locus, strongly similar to XP_579813.1 PREDICTED:
 hypothetical protein XP_579813 [Rattus norvegicus]" ---
 1439793_at 1.08020 gap junction membrane channel protein alpha 3 Gja3
 1439794_at 0.76498 --- ---
 1439795_at 1.04161 G protein-coupled receptor 64 Gpr64
 1439796_at 1.02294 expressed sequence AI747448 AI747448
 1439797_at 1.09236 peroxisome proliferator activator receptor delta Ppard
 1439798_at 0.95424 homeo box C10 Hoxc10
 1439799_at 0.82052 Transcribed locus ---
 1439800_at 2.22805 restin (Reed-Steinberg cell-expressed intermediate filament-associated protein)
 Rsn
 1439801_at 0.94509 Sad1 and UNC84 domain containing 1 Sunc1
 1439802_at 0.92956 serine/threonine kinase 35 Stk35
 1439803_at 0.94240 RIKEN cDNA 6430526N21 gene 6430526N21Rik
 1439804_at 0.91950 RIKEN cDNA 2310015A10 gene 2310015A10Rik
 1439805_at 2.08641 Nuclear factor of activated T-cells 5 Nfat5
 1439806_at 1.08829 "Transcriptional adaptor 1 (HFI1 homolog, yeast) like" Tada11
 1439807_at 1.02312 transmembrane protein 74 Tmem74
 1439808_at 1.03561 RIKEN cDNA A130090K04 gene A130090K04Rik
 1439809_at 1.00876 preferentially expressed antigen in melanoma like 7 Pramel7
 1439810_s_at 1.00256 preferentially expressed antigen in melanoma like 7 Pramel7
 1439811_at 1.09237 5-methyltetrahydrofolate-homocysteine methyltransferase Mtr
 1439812_at 1.10614 RIKEN cDNA 4930402H24 gene 4930402H24Rik
 1439813_at 1.53851 Signal peptide peptidase 3 Sppl3
 1439814_at 0.94295 "ATPase, class I, type 8B, member 4" Atp8b4
 1439815_at 0.95442 "RIKEN cDNA A230048G03 gene /// similar to CG2747-PB, isoform B"
 A230048G03Rik /// LOC670614
 1439816_at 0.95703 Hypothetical gene supported by AK087915 LOC433270
 1439817_at 1.17320 RIKEN cDNA 2900064A13 gene 2900064A13Rik
 1439818_at 1.00970 expressed sequence AI931714 AI931714
 1439819_at 0.96002 expressed sequence AU015263 AU015263
 1439820_at 1.06616 Early B-cell factor 1 Ebf1
 1439821_at 0.95016 Lrp2 binding protein Lrp2bp
 1439822_at 1.04254 --- ---
 1439823_at 1.28844 C-terminal binding protein 2 Ctbp2
 1439824_at 0.93904 choroideremia Chm
 1439825_at 1.19356 deltex 3-like (Drosophila) Dtx3l
 1439826_at 1.26065 heat shock protein 14 Hspa14
 1439827_at 0.75609 "a disintegrin-like and metallopeptidase (reprolysin type) with thrombospondin
 type 1 motif, 12" Adamts12
 1439828_at 1.13346 MAP kinase-activated protein kinase 5 Mapkapk5
 1439829_at 1.06599 Adenylate cyclase 5 Adcy5

1439830_at 0.79012 similar to mitogen activated protein kinase kinase kinase 5 LOC675366
1439831_at 1.08975 Hypothetical LOC240327 LOC240327
1439832_at 1.70974 Wiskott-Aldrich syndrome-like (human) Wasl
1439833_at 0.92608 septin 3 3-Sep
1439834_at 1.14751 RIKEN cDNA 2400009B08 gene 2400009B08Rik
1439835_x_at 1.04215 "ATP-binding cassette, sub-family D (ALD), member 2" Abcd2
1439836_at 1.00511 ankyrin repeat and SOCS box-containing protein 15 Asb15
1439837_at 0.75508 trinucleotide repeat containing 15 Tnrc15
1439838_a_at 1.01934 transmembrane inner ear Tmie
1439839_at 1.79666 RIKEN cDNA D130051D11 gene D130051D11Rik
1439840_at 1.19656 RIKEN cDNA A430088C08 gene A430088C08Rik
1439841_at 0.93172 "zinc finger, FYVE domain containing 27" Zfyve27
1439842_at 1.10541 Zinc finger protein 287 Zfp287
1439843_at 0.96516 calcium/calmodulin-dependent protein kinase IV Camk4
1439844_at 0.94754 RIKEN cDNA 8430426J06 gene 8430426J06Rik
1439845_at 1.12087 Transcribed locus ---
1439846_at 0.95047 Kruppel-like factor 12 Klf12
1439847_s_at 0.99388 Kruppel-like factor 12 Klf12
1439848_at 1.04044 blood vessel epicardial substance Bves
1439849_at 1.00215 "Mus musculus, clone IMAGE:4222153, mRNA" ---
1439850_at 1.00653 cDNA sequence BC066028 BC066028
1439851_at 1.00737 Transcribed locus ---
1439852_at 0.92509 --- ---
1439853_at 0.92272 "beta-1,4-N-acetyl-galactosaminyl transferase 2" B4galnt2
1439854_at 0.86109 "harakiri, BCL2 interacting protein (contains only BH3 domain)" Hrk
1439855_at 0.98309 cDNA sequence BC023818 BC023818
1439856_at 0.89651 activin receptor IIB Acvr2b
1439857_at 1.69167 Ubiquitin specific peptidase 32 Usp32
1439858_at 0.84105 "12 days embryo spinal ganglion cDNA, RIKEN full-length enriched library, clone:D130031D15 product:unclassifiable, full insert sequence" ---
1439859_at 1.02680 RIKEN cDNA 9630033F20 gene 9630033F20Rik
1439860_at 0.97806 eukaryotic elongation factor-2 kinase Eef2k
1439861_at 0.98189 zinc finger protein 583 Zfp583
1439862_at 0.77326 "Visual cortex cDNA, RIKEN full-length enriched library, clone:K530032J01 product:unclassifiable, full insert sequence" ---
1439863_at 1.12108 UDP-glucose ceramide glucosyltransferase Ugcg
1439864_at 0.89980 coiled-coil domain containing 40 Ccdc40
1439865_at 1.09516 kelch repeat and BTB (POZ) domain containing 7 Kbtbd7
1439866_at 0.87183 p53-associated parkin-like cytoplasmic protein Parc
1439867_at 0.85953 U box domain containing 5 Ubox5
1439868_at 0.94520 "Adult male spinal cord cDNA, RIKEN full-length enriched library, clone:A330108B13 product:unclassifiable, full insert sequence" ---
1439869_at 0.92098 RIKEN cDNA 2900054J07 gene 2900054J07Rik

1439870_at	1.01382	RIKEN cDNA A330008L17 gene	A330008L17Rik
1439871_at	1.74729	Lemur tyrosine kinase 2 Lmtk2	
1439872_at	0.93502	Transcribed locus	---
1439873_at	1.06054	"diacylglycerol kinase, delta" Dgkd	
1439874_at	0.98591	RIKEN cDNA 9330102E08 gene	9330102E08Rik
1439875_at	0.88771	zinc finger protein 128 Zfp128	
1439876_at	1.02406	vesicle transport through interaction with t-SNAREs homolog 1A (yeast) Vti1a	
1439877_at	3.21269	Exostoses (multiple) 1 Ext1	
1439878_at	0.98131	involucrin Iv1	
1439879_at	1.16960	Zinc finger protein 568 Zfp568	
1439880_at	0.84684	RIKEN cDNA D630023F18 gene	D630023F18Rik
1439881_at	1.21897	RIKEN cDNA C030013E06 gene	C030013E06Rik
1439882_at	0.91618	Sec23 interacting protein Sec23ip	
1439883_at	1.02983	Nuclear receptor coactivator 1 Ncoa1	
1439884_at	0.96150	nudix (nucleoside diphosphate linked moiety X)-type motif 16 Nudt16	
1439885_at	1.05295	homeo box C5 Hoxc5	
1439886_at	0.88653	RIKEN cDNA 4632415L05 gene	4632415L05Rik
1439887_at	1.19193	"16 days embryo head cDNA, RIKEN full-length enriched library, clone: C130025B04 product:unclassifiable, full insert sequence"	---
1439888_at	1.10320	"Mammary gland RCB-0527 Jyg-MC(B) cDNA, RIKEN full-length enriched library, clone:G930041P03 product:unclassifiable, full insert sequence"	---
1439889_at	0.85822	"sodium channel, voltage-gated, type VIII, alpha"	Scn8a
1439890_at	0.91989	F-box and leucine-rich repeat protein 7 Fbx17	
1439891_at	0.99882	Zinc finger protein 703 Zfp703	
1439892_at	0.99102	BR serine/threonine kinase 1 Brsk1	
1439893_at	1.01413	CDNA clone IMAGE:3471071	---
1439894_at	1.01809	RIKEN cDNA A730056I06 gene	A730056I06Rik
1439895_at	1.90541	expressed sequence AU021025	AU021025
1439896_at	1.03744	LIM motif-containing protein kinase 2 Limk2	
1439897_at	0.91233	Nebulette Nebl	
1439898_at	0.96924	EBNA1 binding protein 2 Ebna1bp2	
1439899_at	0.98337	UDP-N-acetyl-alpha-D-galactosamine:polypeptide N-acetylgalactosaminyltransferase 13 Galnt13	
1439900_at	1.09011	Thiamine triphosphatase Thtpa	
1439901_at	1.19366	RIKEN cDNA 2610208M17 gene	2610208M17Rik
1439902_at	1.01426	complement component 5a receptor 1 C5ar1	
1439903_at	0.92716	cDNA sequence BC028663	BC028663
1439904_at	0.96790	follistatin-like 5 Fstl5	
1439905_at	0.99835	RIKEN cDNA A330062J17 gene	A330062J17Rik
1439906_at	0.92003	"0 day neonate eyeball cDNA, RIKEN full-length enriched library, clone: E130004J04 product:unclassifiable, full insert sequence"	---
1439907_at	0.91089	Radixin Rdx	
1439908_at	0.84291	zinc finger with KRAB and SCAN domains 1	Zkscan1

1439909_at 1.14461 zinc finger protein 521 Zfp521
1439910_a_at 0.79376 --- ---
1439911_at 1.08865 "Adult male urinary bladder cDNA, RIKEN full-length enriched library, clone:9530055O04 product:unclassifiable, full insert sequence" ---
1439912_at 1.18402 RIKEN cDNA 9430098F02 gene 9430098F02Rik
1439913_at 0.92315 RIKEN cDNA 1810053B01 gene 1810053B01Rik
1439914_at 1.21437 RIKEN cDNA 4932416H05 gene 4932416H05Rik
1439915_at 1.30276 Myeloid ecotropic viral integration site-related gene 1 Mrg1
1439916_at 0.93490 hypothetical protein LOC545182 LOC545182
1439917_at 1.94002 PDZ domain containing 8 Pdzd8
1439918_at 1.10071 outer dense fiber of sperm tails 2 Odf2
1439919_at 0.87576 Transcribed locus ---
1439920_at 1.81061 Transcribed locus ---
1439921_at 0.96258 --- ---
1439922_at 1.11713 RIKEN cDNA 1190002C06 gene 1190002C06Rik
1439923_at 1.01048 "tubulin, gamma complex associated protein 5" Tubgcp5
1439924_x_at 1.05605 "tubulin, gamma complex associated protein 5" Tubgcp5
1439925_at 0.81313 transmembrane 4 superfamily member 1 Tm4sf1
1439926_at 1.08113 RIKEN cDNA 4632417D23 gene 4632417D23Rik
1439927_at 1.18726 Paralemmin Palm
1439928_at 1.13009 Spectrin beta 2 Spnb2
1439929_at 1.47690 Malignant fibrous histiocytoma amplified sequence 1 Mfhas1
1439930_at 1.01545 "SWI/SNF related, matrix associated, actin dependent regulator of chromatin, subfamily a, member 2" Smarca2
1439931_at 1.61861 glycogen synthase kinase 3 beta Gsk3b
1439932_at 1.06463 RIKEN cDNA 3110003A22 gene 3110003A22Rik
1439933_at 1.60963 RIKEN cDNA B430316J06 gene B430316J06Rik
1439934_at 0.98314 "solute carrier family 30, member 10" Slc30a10
1439935_at 0.94928 transmembrane protein 122 Tmem122
1439936_at 0.96354 --- ---
1439937_at 1.17587 --- ---
1439938_at 1.28534 serine/threonine kinase 38 Stk38
1439939_at 0.90328 RIKEN cDNA E030042N06 gene E030042N06Rik
1439940_at 1.20741 RIKEN cDNA 2900019G14 gene 2900019G14Rik
1439941_at 0.89223 --- ---
1439942_at 1.14594 prolyl endopeptidase Prep
1439943_at 1.10133 vacuolar protein sorting 54 (yeast) Vps54
1439944_at 0.97700 --- ---
1439945_at 0.88241 zinc finger protein 449 Zfp449
1439946_at 0.86724 Myocyte enhancer factor 2C Mef2c
1439947_at 1.00780 "cytochrome P450, family 11, subfamily a, polypeptide 1" Cyp11a1
1439948_at 1.31383 cDNA sequence BC046401 BC046401
1439949_at 2.40634 Glycogen synthase kinase 3 beta Gsk3b

1439950_at 1.15956 dynein cytoplasmic 1 heavy chain 1 Dync1h1
1439951_at 1.01671 --- ---
1439952_at 1.54704 RIKEN cDNA 3110082M05 gene 3110082M05Rik
1439953_at 0.99873 phosphomannomutase 2 Pmm2
1439954_at 0.94547 RIKEN cDNA 6430514M23 gene 6430514M23Rik
1439955_at 0.85782 RAD9 homolog B (S. cerevisiae) Rad9b
1439956_at 1.12376 "membrane-spanning 4-domains, subfamily A, member 6B" Ms4a6b
1439957_at 0.99165 expressed sequence AI843190 AI843190
1439958_at 1.01324 similar to transcription elongation factor B polypeptide 3 binding protein 1 isoform 1 /// similar to transcription elongation factor B polypeptide 3 binding protein 1 isoform 1 LOC546257 /// LOC619649
1439959_at 1.01171 fibroblast growth factor 11 Fgf11
1439960_at 1.01675 RNA pseudouridylate synthase domain containing 2 Rpsud2
1439961_x_at 0.94507 Transcribed locus ---
1439962_at 0.92096 RIKEN cDNA 2310010J17 gene 2310010J17Rik
1439963_x_at 0.91576 RIKEN cDNA 1700129I04 gene 1700129I04Rik
1439964_at 0.90801 "DNA segment, Chr 8, Brigham & Women's Genetics 1414 expressed" D8Bwg1414e
1439965_at 1.13634 Transcribed locus ---
1439966_x_at 0.91647 sideroflexin 2 Sfxn2
1439967_at 1.02162 RIKEN cDNA 1700071A11 gene 1700071A11Rik
1439968_x_at 0.91319 "Adult male corpora quadrigemina cDNA, RIKEN full-length enriched library, clone:B230215D24 product:unclassifiable, full insert sequence" ---
1439969_at 0.86266 RIKEN cDNA 4930500O05 gene 4930500O05Rik
1439970_at 2.01654 WNK lysine deficient protein kinase 1 Wnk1
1439971_at 0.90564 RIKEN cDNA 6330439K17 gene 6330439K17Rik
1439972_at 1.34463 ethanolamine kinase 1 Etnk1
1439973_at 0.95459 Crx opposite strand transcript 1 Crxos1
1439974_at 1.09151 RIKEN cDNA C430014M02 gene C430014M02Rik
1439975_at 0.93491 cDNA sequence BC062109 BC062109
1439976_at 1.08399 "asparagine-linked glycosylation 11 homolog (yeast, alpha-1,2-mannosyltransferase)" Alg11
1439977_at 0.90186 hypothetical protein 4930515I15 4930515I15
1439978_at 0.99141 RIKEN cDNA D030022P06 gene D030022P06Rik
1439979_at 0.95708 zinc finger protein 398 Zfp398
1439980_at 2.18163 "DNA segment, Chr 14, ERATO Doi 725, expressed" D14Ertd725e
1439981_at 0.93780 RIKEN cDNA A430001F24 gene A430001F24Rik
1439982_at 0.90074 "Adult male liver tumor cDNA, RIKEN full-length enriched library, clone: C730006I10 product:unclassifiable, full insert sequence" ---
1439983_a_at 0.91218 amiloride-sensitive cation channel 3 Accn3
1439984_at 0.95525 phosphatidylserine synthase 2 Ptdss2
1439985_at 1.05452 "ATP-binding cassette, sub-family C (CFTR/MRP), member 12" Abcc12
1439986_at 0.96859 "diacylglycerol kinase, iota" Dgki

1439987_at 0.99819 "glutamate receptor, ionotropic, kainate 1" Grik1
1439988_at 1.11508 --- ---
1439989_at 2.03452 tuberous sclerosis 1 Tsc1
1439990_at 1.03403 "12 days embryo spinal ganglion cDNA, RIKEN full-length enriched library, clone:D130065P14 product:unclassifiable, full insert sequence" ---
1439991_a_at 1.05608 poly(A)binding protein nuclear-like 1 Pabpn1
1439992_at 1.05376 RIKEN cDNA 2610206G21 gene 2610206G21Rik
1439993_at 1.11823 Arginine/serine-rich coiled-coil 1 Rsrc1
1439994_at 0.70014 "Adult male corpora quadrigemina cDNA, RIKEN full-length enriched library, clone:B230306E21 product:unclassifiable, full insert sequence" ---
1439995_at 1.08301 expressed sequence C80638 C80638
1439996_at 1.00113 TatD DNase domain containing 1 Tatdn1
1439997_at 0.86792 synaptojanin 1 Synj1
1439998_at 1.86542 jumonji domain containing 1C Jmjd1c
1439999_at 1.53362 --- ---
1440000_at 0.99060 RIKEN cDNA E330013P04 gene E330013P04Rik
1440001_at 0.97554 RNA imprinted and accumulated in nucleus Rian
1440002_at 1.24599 RNA (guanine-7-) methyltransferase Rnmt
1440003_at 0.91262 Transcribed locus ---
1440004_at 1.07113 Suppressor of fused homolog (Drosophila) Sufu
1440005_at 1.09259 hypothetical protein C730009D12 C730009D12
1440006_at 1.01475 cDNA sequence BC026600 BC026600
1440007_at 1.07458 "solute carrier family 43, member 1" Slc43a1
1440008_at 0.99799 RIKEN cDNA 2310043L19 gene 2310043L19Rik
1440009_at 1.08733 olfactory receptor 78 Olfr78
1440010_at 1.16478 RIKEN cDNA C230066K19 gene C230066K19Rik
1440011_at 2.55834 Exostoses (multiple) 1 Ext1
1440012_at 1.29897 --- ---
1440013_at 2.16964 Tripartite motif-containing 44 Trim44
1440014_at 1.46349 Phosphofurin acidic cluster sorting protein 1 Pacs1
1440015_at 0.88393 Transcribed locus ---
1440016_at 1.06820 Dihydrolipoamide branched chain transacylase E2 Dbt
1440017_at 0.96270 "pleckstrin homology domain containing, family G (with RhoGef domain) member 3" Plekhg3
1440018_at 0.90986 RIKEN cDNA A330043J11 gene A330043J11Rik
1440019_at 1.07611 G protein-coupled receptor 22 Gpr22
1440020_at 2.26044 Rearranged L-myc fusion sequence Rlf
1440021_at 1.00614 G protein-coupled receptor 20 Gpr20
1440022_at 1.77622 "polymerase (DNA-directed), delta interacting protein 3" Poldip3
1440023_at 1.01890 protocadherin 12 Pcdh12
1440024_at 0.81793 "inositol 1,4,5-trisphosphate 3-kinase C" Itpkc
1440025_at 1.20221 kaptin Kptn
1440026_at 1.54447 Trichorhinophalangeal syndrome I (human) Trps1

1440027_at 0.97180 "10 days neonate cerebellum cDNA, RIKEN full-length enriched library, clone: B930082K12 product:unclassifiable, full insert sequence" ---

1440028_at 2.10734 Autophagy-related 7 (yeast) Atg7

1440029_at 1.04935 "ST8 alpha-N-acetyl-neuraminide alpha-2,8-sialyltransferase 3" St8sia3

1440030_at 1.08321 --- ---

1440031_at 1.04069 dact 3 Dact3

1440032_at 0.95097 RIKEN cDNA E130008O17 gene E130008O17Rik

1440033_at 1.09603 Phosphoprotein associated with glycosphingolipid microdomains 1 Pag1

1440034_at 1.17277 signal transducing adaptor molecule (SH3 domain and ITAM motif) 2 Stam2

1440035_at 1.11671 --- ---

1440036_x_at 0.99967 "similar to poly(A) binding protein, cytoplasmic 4 isoform 1" LOC230010

1440037_at 3.11741 pre B-cell leukemia transcription factor 1 Pbx1

1440038_at 1.06237 RIKEN cDNA B430007K19 gene B430007K19Rik

1440039_at 1.13415 Bromodomain and WD repeat domain containing 3 Brwd3

1440040_at 3.01534 Cyclin-dependent kinase 6 Cdk6

1440041_at 1.06233 "Transforming growth factor, beta receptor III" Tgfbr3

1440042_at 1.09643 "ankyrin 2, brain" Ank2

1440043_at 0.97828 "ankyrin 2, brain" Ank2

1440044_at 0.93967 Transcription termination factor 1 Ttf1

1440045_at 1.22443 "Crystallin, zeta (quinone reductase)-like 1" Cryz11

1440046_at 1.10576 cDNA sequence BC031748 BC031748

1440047_at 0.97715 suppressor of cytokine signaling 1 Socs1

1440048_at 1.39403 "SWI/SNF related, matrix associated, actin dependent regulator of chromatin, subfamily a, member 5" Smarca5

1440049_at 0.93313 Ribosomal protein 10 Rpl10

1440050_at 0.97117 Hbs1-like (S. cerevisiae) Hbs11

1440051_at 1.40984 "Protein phosphatase 3, catalytic subunit, alpha isoform" Ppp3ca

1440052_at 2.76216 Ephrin A5 Efna5

1440053_at 1.06395 Collagen and calcium binding EGF domains 1 Ccbe1

1440054_at 1.06487 "phosphatidylinositol 3-kinase, catalytic, alpha polypeptide" Pik3ca

1440055_at 1.04380 Expressed sequence AI848100 AI848100

1440056_at 1.00733 "Adult male hippocampus cDNA, RIKEN full-length enriched library, clone: C630001C05 product:unclassifiable, full insert sequence" ---

1440057_at 1.03041 hydroxysteroid (17-beta) dehydrogenase 7 Hsd17b7

1440058_at 1.04841 "solute carrier family 22 (organic cation transporter), member 14" Slc22a14

1440059_at 1.19071 Mitogen-activated protein kinase kinase kinase 5 Map4k5

1440060_at 1.04917 hypothetical LOC331480 LOC331480

1440061_at 1.34370 ring-box 1 Rbx1

1440062_at 0.97623 RIKEN cDNA 2510009E07 gene 2510009E07Rik

1440063_at 1.02512 "phenylalanine-tRNA synthetase-like, alpha subunit" Farsla

1440064_at 0.98170 cDNA sequence BC026657 BC026657

1440065_at 1.18427 Polymerase (RNA) II (DNA directed) polypeptide J Polr2j

1440066_at 1.39101 "SWI/SNF-related, matrix-associated actin-dependent regulator of chromatin,

subfamily a, containing DEAD/H box 1`" Smarcd1

1440067_at 1.03453 Neural cell adhesion molecule 1 Ncam1

1440068_at 1.16612 Regulatory factor X domain containing 2 homolog (human) Rfxdc2

1440069_at 1.08675 Ephrin B2 Efnb2

1440070_at 0.83018 choline acetyltransferase Chat

1440071_at 1.00408 "membrane associated guanylate kinase, WW and PDZ domain containing 1" Magi1

1440072_at 0.84987 RIKEN cDNA E430022K19 gene E430022K19Rik

1440073_at 1.26631 ribonucleotide reductase M1 Rrm1

1440074_at 0.94946 hypothetical gene supported by AK082832 LOC432988

1440075_at 1.08776 Transcribed locus ---

1440076_at 1.42070 Sequestosome 1 Sqstm1

1440077_at 1.06130 RIKEN cDNA 6330549D23 gene 6330549D23Rik

1440078_at 2.48942 Bromodomain containing 4 Brd4

1440079_at 1.00108 RIKEN cDNA 3632454L22 gene 3632454L22Rik

1440080_at 0.95824 Serine/arginine-rich protein specific kinase 2 Srpk2

1440081_at 1.44764 RIKEN cDNA 4631422C13 gene 4631422C13Rik

1440082_at 1.80372 PTK2 protein tyrosine kinase 2 Ptk2

1440083_at 2.15093 RIKEN cDNA A430061O12 gene A430061O12Rik

1440084_at 0.86489 Expressed sequence AI663975 AI663975

1440085_at 0.87971 ectodysplasin A2 isoform receptor Eda2r

1440086_at 0.82726 ring finger protein 182 Rnf182

1440087_at 1.04077 --- ---

1440088_at 1.04205 Transcribed locus ---

1440089_at 0.99023 cAMP responsive element binding protein 5 Creb5

1440090_at 1.14674 "Solute carrier family 25, member 27" Slc25a27

1440091_at 1.56568 myeloid ecotropic viral integration site-related gene 1 Mrg1

1440092_at 4.42931 Exostoses (multiple) 1 Ext1

1440093_at 0.91555 endothelin receptor type A Ednra

1440094_at 0.94367 cDNA sequence BC030870 BC030870

1440095_at 1.72215 --- ---

1440096_at 1.00619 "extracellular matrix protein 2, female organ and adipocyte specific" Ecm2

1440097_at 1.13938 miRNA containing gene Mirg

1440098_at 1.13390 RIKEN cDNA D030018L15 gene D030018L15Rik

1440099_at 0.95533 "0 day neonate cerebellum cDNA, RIKEN full-length enriched library, clone: C230026N07 product:unclassifiable, full insert sequence" ---

1440100_at 0.82461 RIKEN cDNA 4930527D15 gene 4930527D15Rik

1440101_at 1.10950 --- ---

1440102_at 0.95393 "solute carrier family 36 (proton/amino acid symporter), member 3" Slc36a3

1440103_at 0.94763 Transcribed locus ---

1440104_at 1.71037 RAN binding protein 2 Ranbp2

1440105_at 0.95381 jumonji domain containing 2D Jmjd2d

1440106_at 0.96645 Transcription factor 4 Tcf4

1440107_at 1.09828 "Smg-7 homolog, nonsense mediated mRNA decay factor (C. elegans)"
Smg7

1440108_at 0.56310 forkhead box P2 Foxp2

1440109_at 0.97121 "DNA segment, Chr 7, ERATO Doi 413, expressed" D7Ertd413e

1440110_at 1.17686 RIKEN cDNA B130019D13 gene B130019D13Rik

1440111_at 1.45660 Formin-like 2 Fmnl2

1440112_at 0.92819 leucine rich repeat transmembrane neuronal 3 Lrrtm3

1440113_at 0.99923 metastasis associated 1 Mta1

1440114_x_at 0.94739 ankyrin repeat domain 47 Ankrd47

1440115_at 0.94463 Olfactomedin 1 Olfm1

1440116_at 0.88663 RIKEN cDNA D630045J12 gene D630045J12Rik

1440117_at 1.00384 leucine rich repeat containing 48 Lrrc48

1440118_at 0.94265 "DNA segment, Chr 11, ERATO Doi 333, expressed" D11Ertd333e

1440119_at 1.13305 signal-induced proliferation-associated 1 like 1 Sipal11

1440120_at 1.53785 "Guanine nucleotide binding protein (G protein), beta polypeptide 2 like 1"
Gnb211

1440121_at 0.86679 "ST6 (alpha-N-acetyl-neuraminyl-2,3-beta-galactosyl-1,3)-N-
acetylgalactosaminide alpha-2,6-sialyltransferase 3" St6galnac3

1440122_at 0.90601 hypothetical protein LOC666504 /// hypothetical protein LOC672638 ///
hypothetical protein LOC675536 LOC666504 /// LOC672638 /// LOC675536

1440123_at 1.59540 --- ---

1440124_at 1.03277 RIKEN cDNA B230334C09 gene B230334C09Rik

1440125_at 0.88553 RIKEN cDNA A530054K11 gene A530054K11Rik

1440126_at 1.11536 cDNA sequence BC037704 BC037704

1440127_a_at 1.06835 Transcribed locus ---

1440128_s_at 0.88649 seizure related 6 homolog like Sez6l

1440129_at 0.99661 Aurora kinase A Aurka

1440130_at 0.92458 expressed sequence BB116930 BB116930

1440131_at 0.92145 Poliovirus receptor-related 1 Pvr1l

1440132_s_at 0.96305 "protein kinase, cAMP dependent regulatory, type I beta" Prkar1b

1440133_x_at 0.85107 "protein kinase, cAMP dependent regulatory, type I beta" Prkar1b

1440134_at 1.06006 "cytochrome P450, family 4, subfamily a, polypeptide 10 /// cDNA sequence
BC013476" Cyp4a10 /// BC013476

1440135_at 1.13692 "Amyloid beta (A4) precursor protein-binding, family B, member 2" Apbb2

1440136_at 0.77104 Zinc finger protein 509 Zfp509

1440137_at 0.84418 B-cell leukemia/lymphoma 11B Bcl11b

1440138_at 0.99740 "AT hook, DNA binding motif, containing 1" Ahdc1

1440139_at 1.04793 "Neural precursor cell expressed, developmentally down-regulated gene 4-
like" Nedd4l

1440140_at 1.03862 Nuclear factor I/C Nfic

1440141_at 0.89490 "potassium voltage-gated channel, subfamily Q, member 2" Kcnq2

1440142_s_at 1.01964 glial fibrillary acidic protein Gfap

1440143_at 1.00406 "phosphatidylinositol glycan anchor biosynthesis, class Z" Pigz

1440144_x_at 1.09331 hypothetical protein C330046E03 C330046E03
 1440145_at 1.01803 Histocompatibility 60 H60
 1440146_at 0.88418 vacuolar protein sorting 13A (yeast) Vps13a
 1440147_at 1.01218 "leucine-rich repeat LGI family, member 2" Lgi2
 1440148_at 1.03657 G protein-coupled receptor 6 Gpr6
 1440149_at 0.93886 PAN3 polyA specific ribonuclease subunit homolog (S. cerevisiae) Pan3
 1440150_at 1.02587 "Transglutaminase 3, E polypeptide" Tgm3
 1440151_s_at 0.96133 endothelial differentiation-related factor 1 Edf1
 1440152_x_at 0.96070 endothelial differentiation-related factor 1 /// similar to Endothelial differentiation-related factor 1 (EDF-1) (Multiprotein bridging factor 1) (MBF1) (Calmodulin-associated peptide 19) (CAP-19) /// similar to Endothelial differentiation-related factor 1 (EDF-1) (Multiprotein bridging factor 1) (MBF1) (Calmodulin-associated peptide 19) (CAP-19) /// similar to Endothelial differentiation-related factor 1 (EDF-1) (Multiprotein bridging factor 1) (MBF1) (Calmodulin-associated peptide 19) (CAP-19) Edf1 /// LOC665181 /// LOC671865 /// LOC674653
 1440153_at 0.95419 "Adult male brain cDNA, RIKEN full-length enriched library, clone:3526403L10 product:unclassifiable, full insert sequence" ---
 1440154_at 0.94196 Pre B-cell leukemia transcription factor 3 Pbx3
 1440155_at 1.01566 follistatin-like 4 Fstl4
 1440156_s_at 0.99528 expressed sequence AI851523 AI851523
 1440157_at 0.91610 Sex comb on midleg-like 4 (Drosophila) Scml4
 1440158_x_at 1.13338 Coiled-coil-helix-coiled-coil-helix domain containing 7 Chchd7
 1440159_at 0.86362 RIKEN cDNA 1700010B09 gene 1700010B09Rik
 1440160_x_at 1.00696 RIKEN cDNA 2010321M09 gene 2010321M09Rik
 1440161_at 0.95025 Matrix metalloproteinase 16 Mmp16
 1440162_x_at 0.88980 hypothetical protein A630043P06 A630043P06
 1440163_at 1.91820 RIKEN cDNA 6030490B17 gene 6030490B17Rik
 1440164_x_at 0.87809 "CD8 antigen, alpha chain" Cd8a
 1440165_at 1.06719 "Protein tyrosine phosphatase, receptor type, C" Ptpcr
 1440166_x_at 0.98931 5-hydroxytryptamine (serotonin) receptor 1D Htr1d
 1440167_s_at 1.01078 LIM domain containing preferred translocation partner in lipoma Lpp
 1440168_x_at 0.96656 potassium channel tetramerisation domain containing 7 Kctd7
 1440169_x_at 0.98565 interferon (alpha and beta) receptor 2 Ifnar2
 1440170_at 0.98425 ankyrin repeat domain 10 Ankrd10
 1440171_x_at 1.10492 RIKEN cDNA 1700049G17 gene 1700049G17Rik
 1440172_s_at 0.92188 RIKEN cDNA 1700113H08 gene 1700113H08Rik
 1440173_x_at 1.06328 "selectin, platelet" Selp
 1440174_at 0.96689 RIKEN cDNA E130307A14 gene E130307A14Rik
 1440175_at 1.05312 Epsin 2 Epn2
 1440176_x_at 0.90158 RIKEN cDNA 1700003F12 gene 1700003F12Rik
 1440177_at 1.03319 hypothetical protein 9630027E11 9.63E+17
 1440178_x_at 0.91681 zeta-chain (TCR) associated protein kinase Zap70
 1440179_x_at 1.05994 IBR domain containing 1 Ibrdc1
 1440180_x_at 1.01743 zinc finger and BTB domain containing 3 Zbtb3

1440181_at	1.01176	"gene model 1568, (NCBI)"	Gm1568
1440182_at	1.01475	frizzled homolog 10 (Drosophila)	Fzd10
1440183_x_at	1.11881	Coenzyme Q9 homolog (yeast)	Coq9
1440184_at	1.04103	RIKEN cDNA 2610005L07 gene	2610005L07Rik
1440185_x_at	1.05427	Ubiquitin-like 1 (sentrin) activating enzyme E1A	Uble1a
1440186_s_at	0.90612	RIKEN cDNA 2310020A21 gene	2310020A21Rik
1440187_at	0.78275	"TAF3 RNA polymerase II, TATA box binding protein (TBP)-associated factor" Taf3	
1440188_at	1.05354	---	---
1440189_at	0.93144	"protein phosphatase 2, regulatory subunit B (B56), alpha isoform"	Ppp2r5a
1440190_at	0.92812	leukocyte receptor cluster (LRC) member 9	Leng9
1440191_s_at	1.04217	leukocyte receptor cluster (LRC) member 9	Leng9
1440192_at	1.03918	RIKEN cDNA 1810054D07 gene	1810054D07Rik
1440193_at	0.99148	ankyrin repeat domain 12	Ankrd12
1440194_at	0.87050	RIKEN cDNA 2210011C24 gene	2210011C24Rik
1440195_at	0.77990	Serpine1 mRNA binding protein 1	Serbp1
1440196_at	1.02180	---	---
1440197_at	0.91769	cDNA sequence BC032203	BC032203
1440198_at	1.13774	Enhancer of polycomb homolog 1 (Drosophila)	Epc1
1440199_at	0.88728	expressed sequence AU042651	AU042651
1440200_at	0.84436	RIKEN cDNA 9630031F12 gene	9630031F12Rik
1440201_at	1.09984	mitogen activated protein kinase 10	Mapk10
1440202_at	0.81637	similar to gonadotropin inducible ovarian transcription factor 2	LOC622175
1440203_at	1.00067	Polycomb group ring finger 3	Pcgf3
1440204_at	0.99908	RIKEN cDNA 3110039M20 gene	3110039M20Rik
1440205_at	1.20590	---	---
1440206_at	1.06097	RIKEN cDNA A930024E05 gene	A930024E05Rik
1440207_at	1.35519	"Nacylglycerol-3-phosphate O-acyltransferase 7 (lysophosphatidic acid acyltransferase, eta)"	Agpat7
1440208_at	0.98860	RAS-related C3 botulinum substrate 2	Rac2
1440209_at	1.05111	membrane-associated ring finger (C3HC4) 1	1-Mar
1440210_at	0.95614	"calcium channel, voltage-dependent, gamma subunit 2"	Cacng2
1440211_at	1.00694	"cytochrome P450, family 2, subfamily j, polypeptide 11"	Cyp2j11
1440212_at	1.02618	---	---
1440213_a_at	0.89983	RIKEN cDNA 2010001M06 gene	2010001M06Rik
1440214_at	0.97929	RIKEN cDNA A630001G21 gene	A630001G21Rik
1440215_at	0.72580	hypothetical protein C130086A10	C130086A10
1440216_at	0.96840	interferon induced transmembrane protein 5	Ifitm5
1440217_at	1.00511	similar to hypothetical protein FLJ39743	LOC434197
1440218_at	0.99823	cDNA sequence BC040758	BC040758
1440219_at	1.68394	Centrosome and spindle pole associated protein 1	Cspp1
1440220_at	0.96169	expressed sequence AI503564	AI503564
1440221_at	0.94908	expressed sequence AA408650	AA408650

1440222_at	1.03408	---	---	
1440223_at	2.04038	RNA binding motif protein 6	Rbm6	
1440224_at	1.03526	RIKEN cDNA 9330133O14 gene	9330133O14Rik	
1440225_at	1.04127	G protein-coupled receptor 116	Gpr116	
1440226_at	0.91959	zinc finger protein 760	Zfp760	
1440227_at	1.14173	expressed sequence BF642829	BF642829	
1440228_at	0.81496	RAN binding protein 6	Ranbp6	
1440229_at	0.93139	RIKEN cDNA 2310034G01 gene	2310034G01Rik	
1440230_at	1.06399	leucine rich repeat containing 54	Lrrc54	
1440231_at	0.99372	microtubule-associated protein 9	Mtap9	
1440232_at	0.95014	similar to KIAA1913 /// similar to KIAA1913	LOC622645 /// LOC674857	
1440233_at	0.94068	RIKEN cDNA A930013B10 gene	A930013B10Rik	
1440234_at	0.90280	RIKEN cDNA 1810012P15 gene	1810012P15Rik	
1440235_at	0.85853	"integrin, alpha 10"	Itga10	
1440236_at	1.10297	RIKEN cDNA A830039H05 gene	A830039H05Rik	
1440237_at	0.75229	"excision repair cross-complementing rodent repair deficiency, complementation group 4"	Ercc4	
1440238_at	0.94526	glioma tumor suppressor candidate region gene 1	Gltscr1	
1440239_at	0.91522	RIKEN cDNA 2310066E14 gene	2310066E14Rik	
1440240_at	1.02893	neuropeptide B	Npb	
1440241_at	1.14341	"Transient receptor potential cation channel, subfamily M, member 1"	Trpm1	
1440242_at	1.15145	RIKEN cDNA C030014K22 gene	C030014K22Rik	
1440243_at	1.07896	copine VII	Cpne7	
1440244_at	1.07476	avian erythroblastosis virus E-26 (v-ets) oncogene related	Erg	
1440245_at	0.88850	coiled-coil domain containing 88	Ccdc88	
1440246_at	1.04055	longevity assurance homolog 6 (<i>S. cerevisiae</i>)	Lass6	
1440247_at	1.20146	PHD finger protein 14	Phf14	
1440248_at	0.93794	cancer susceptibility candidate 4	Casc4	
1440249_at	0.98146	RIKEN cDNA 4921525H12 gene	4921525H12Rik	
1440250_at	0.50592	"procollagen, type IV, alpha 4"	Col4a4	
1440251_s_at	1.03135	zinc finger protein 64	Zfp64	
1440252_at	0.94287	"H2A histone family, member J"	H2afj	
1440253_at	1.06549	---	---	
1440254_at	1.00545	"Transcribed locus, weakly similar to XP_417295.1 PREDICTED: similar to TAF3 protein [<i>Gallus gallus</i>]"	---	
1440255_at	1.07831	MBD2-interacting zinc finger	Mizf	
1440256_at	0.91791	regulator of G-protein signalling 9 binding protein	Rgs9bp	
1440257_at	0.86263	oocyte specific homeobox 6	Obox6	
1440258_at	1.02299	"Potassium voltage-gated channel, subfamily Q, member 2"	Kcnq2	
1440259_at	1.50403	"Mannosidase, alpha, class 1A, member 2"	Man1a2	
1440260_at	2.87528	RIKEN cDNA A930007B11 gene	A930007B11Rik	
1440261_at	1.10311	"adaptor-related protein complex AP-4, epsilon 1"	Ap4e1	
1440262_at	0.85839	---	---	

1440263_at	1.53053	neural regeneration protein	Nrp
1440264_at	0.91689	---	---
1440265_at	0.94326	Jun proto-oncogene related gene d1	Jund1
1440266_at	1.02037	RIKEN cDNA 5830428M24 gene	5830428M24Rik
1440267_at	1.13271	peroxisome biogenesis factor 1	Pex1
1440268_at	1.81218	tripartite motif-containing 41	Trim41
1440269_at	0.96654	SCY1-like 2 (<i>S. cerevisiae</i>)	Scyl2
1440270_at	0.94422	fibroblast growth factor 12	Fgf12
1440271_at	1.00545	Transcribed locus	---
1440272_at	0.93363	"Eukaryotic translation initiation factor 2, subunit 1 alpha"	Eif2s1
1440273_at	0.95039	"10 days neonate cerebellum cDNA, RIKEN full-length enriched library, clone: B930073D03 product: hypothetical protein, full insert sequence"	---
1440274_at	1.04303	Rap guanine nucleotide exchange factor (GEF) 4	Rapgef4
1440275_at	0.94060	runt related transcription factor 3	Runx3
1440276_at	0.94500	"gene model 62, (NCBI)"	Gm62
1440277_at	2.09901	expressed sequence AU041474	AU041474
1440278_at	0.84969	---	---
1440279_at	0.96700	---	---
1440280_at	0.95397	coiled-coil domain containing 45	Ccdc45
1440281_at	1.20342	RIKEN cDNA 0610042C05 gene	0610042C05Rik
1440282_at	0.85557	tubby like protein 4 /// RIKEN cDNA 2210038L17 gene	Tulp4 /// 2210038L17Rik
1440283_at	1.18582	RIKEN cDNA 1810059H22 gene	1810059H22Rik
1440284_at	0.99295	"Transcribed locus, weakly similar to XP_576460.1 PREDICTED: similar to hypothetical protein PB402898.00.0 [<i>Rattus norvegicus</i>]"	---
1440285_at	0.99178	"protein phosphatase 1, regulatory (inhibitor) subunit 9A"	Ppp1r9a
1440286_at	0.99823	F-box and leucine-rich repeat protein 16	Fbxl16
1440287_at	3.30812	RIKEN cDNA 1110014D18 gene	1110014D18Rik
1440288_at	1.10640	patched domain containing 2	Ptchd2
1440289_at	0.95822	"GA repeat binding protein, beta 2"	Gabpb2
1440290_at	0.93458	hypothetical gene supported by AK039686 LOC434089	
1440291_at	1.12160	Transcribed locus	---
1440292_at	1.49177	"DNA segment, Chr 13, ERATO Doi 666, expressed"	D13Ertd666e
1440293_at	0.94028	RIKEN cDNA C230081A13 gene	C230081A13Rik
1440294_at	1.38194	SET domain containing 2	Setd2
1440295_at	1.56762	RIKEN cDNA 1110057K04 gene	1110057K04Rik
1440296_at	1.09270	hypothetical protein LOC666250 /// hypothetical protein LOC669419	LOC666250 /// LOC669419
1440297_at	1.01278	RIKEN cDNA 4931406I20 gene	4931406I20Rik
1440298_at	0.90521	triggering receptor expressed on myeloid cells-like 2	Trem12
1440299_at	1.09725	RIKEN cDNA E330016A19 gene	E330016A19Rik
1440300_at	0.99073	Transcribed locus	---
1440301_at	1.11174	---	---

1440302_at 0.98031 "solute carrier family 25 (mitochondrial carrier, phosphate carrier), member 3" Slc25a3

1440303_at 1.07996 "Solute carrier family 7, member 6 opposite strand" Slc7a6os

1440304_at 1.39634 BTB and CNC homology 2 Bach2

1440305_at 1.12920 --- ---

1440306_at 1.01117 "Nuclear receptor subfamily 3, group C, member 2" Nr3c2

1440307_at 1.62882 "general transcription factor IIIC, polypeptide 2, beta" Gtf3c2

1440308_at 1.06639 cytidylate kinase Cmpk

1440309_at 1.04117 RIKEN cDNA 4833430A08 gene 4833430A08Rik

1440310_at 0.98607 "Runt-related transcription factor 1; translocated to, 1 (cyclin D-related)" Runx1t1

1440311_at 1.14467 sorbin and SH3 domain containing 1 Sorbs1

1440312_at 0.90528 "ELOVL family member 7, elongation of long chain fatty acids (yeast)" Elovl7

1440313_at 0.99419 "fucosidase, alpha-L- 2, plasma" Fuca2

1440314_at 1.72932 Thyroid hormone receptor interactor 12 Trip12

1440315_at 1.01623 muscleblind-like 1 (Drosophila) Mbnl1

1440316_at 1.09007 RIKEN cDNA 6430526N21 gene 6430526N21Rik

1440317_at 2.64126 RIKEN cDNA C130068B02 gene C130068B02Rik

1440318_at 0.83972 WD repeat domain 35 Wdr35

1440319_at 0.92258 "gene model 817, (NCBI)" Gm817

1440320_at 1.19843 --- ---

1440321_at 0.95554 Microtubule associated serine/threonine kinase 2 Mast2

1440322_at 1.06895 "CUG triplet repeat, RNA binding protein 2" Cugbp2

1440323_at 1.08277 synaptotagmin II Syt2

1440324_at 0.85631 mitochondrial ribosomal protein L19 Mrpl19

1440325_at 0.98026 "Adult male medulla oblongata cDNA, RIKEN full-length enriched library, clone:6330531L09 product:unclassifiable, full insert sequence" ---

1440326_at 0.96917 Src family associated phosphoprotein 1 Scap1

1440327_at 1.00903 expressed sequence AI195470 AI195470

1440328_at 1.15198 nuclear autoantigenic sperm protein (histone-binding) Nasp

1440329_s_at 1.00866 G patch domain containing 2 Gpatc2

1440330_at 0.97971 "Histone 1, H1e" Hist1h1e

1440331_at 0.98663 RIKEN cDNA 9430079B08 gene 9430079B08Rik

1440332_at 1.38285 carnitine deficiency-associated gene expressed in ventricle 3 Cdv3

1440333_at 1.01276 MAM domain containing 4 Mamdc4

1440334_at 1.33673 Zinc finger protein 90 Zfp90

1440335_at 0.95314 --- ---

1440336_at 0.99584 protein arginine N-methyltransferase 6 Prmt6

1440337_at 1.08578 Transcribed locus ---

1440338_at 1.03077 RIKEN cDNA E430028B21 gene E430028B21Rik

1440339_at 0.61613 RIKEN cDNA 4833416E15 gene 4833416E15Rik

1440340_at 0.92580 "UDP-Gal:betaGal beta 1,3-galactosyltransferase, polypeptide 6" B3galt6

1440341_at 1.32105 similar to SMF protein LOC629345

1440342_at 1.04716 RIKEN cDNA G530011O06 gene G530011O06Rik
1440343_at 0.98328 "ribosomal protein S6 kinase, polypeptide 5" Rps6ka5
1440344_at 0.95993 Transcribed locus ---
1440345_at 1.17666 expressed sequence AI450353 AI450353
1440346_at 0.97748 Jumonji domain containing 3 Jmjd3
1440347_at 1.42935 Rho GTPase activating protein 10 Arhgap10
1440348_at 1.08740 "zinc finger, FYVE domain containing 9" Zfyve9
1440349_at 1.19930 Cyclin D binding myb-like transcription factor 1 Dmtf1
1440350_at 1.01215 --- ---
1440351_at 1.50132 Baculoviral IAP repeat-containing 4 Birc4
1440352_at 0.96814 region containing RIKEN cDNA 1700028E10 gene; hypothetical protein
A630054D14 LOC625175
1440353_at 0.84763 neurotrophin 5 Ntf5
1440354_at 0.95460 "ELOVL family member 7, elongation of long chain fatty acids (yeast)" Elovl7
1440355_at 1.01103 potassium channel tetramerisation domain containing 12b Kctd12b
1440356_at 1.06763 cytokine induced apoptosis inhibitor 1 Ciapin1
1440357_at 1.03521 Transcribed locus ---
1440358_at 0.85249 Rho guanine nucleotide exchange factor (GEF) 15 Arhgef15
1440359_at 0.96573 RIKEN cDNA 1700012H17 gene 1700012H17Rik
1440360_at 1.59167 ADP-ribosylation factor-like 6 interacting protein 5 Arl6ip5
1440361_at 1.11261 "16 days embryo head cDNA, RIKEN full-length enriched library, clone:
C130014E09 product:unclassifiable, full insert sequence" ---
1440362_at 0.92802 RIKEN cDNA A330068P14 gene A330068P14Rik
1440363_at 1.01500 Transcribed locus ---
1440364_a_at 0.82290 RIKEN cDNA A230062G08 gene A230062G08Rik
1440365_at 1.59660 "Actin, beta, cytoplasmic" Actb
1440366_at 1.19276 "Solute carrier family 24 (sodium/potassium/calcium exchanger), member
2" Slc24a2
1440367_at 1.04211 Phosphatidylglycerophosphate synthase 1 Pgs1
1440368_at 1.12991 Jumonji domain containing 2B Jmjd2b
1440369_at 1.05535 Prostaglandin E receptor 3 (subtype EP3) Ptger3
1440370_at 1.02952 "ATP-binding cassette, sub-family A (ABC1), member 13 /// similar to ATP
binding cassette, sub-family A (ABC1), member 13" Abca13 /// LOC672028
1440371_at 0.95217 "2 days neonate sympathetic ganglion cDNA, RIKEN full-length enriched
library, clone:7120457A19 product:unclassifiable, full insert sequence" ---
1440372_at 2.04957 ADP-ribosylation factor-like 15 Arl15
1440373_at 2.14789 "ELAV (embryonic lethal, abnormal vision, Drosophila)-like 1 (Hu antigen
R)" Elavl1
1440374_at 0.91395 Phosphodiesterase 1C Pde1c
1440375_at 1.35200 RIKEN cDNA 5730419I09 gene 5730419I09Rik
1440376_at 0.95634 F-box protein 41 Fbxo41
1440377_at 0.92139 tigger transposable element derived 5 Tigd5
1440378_at 0.96265 --- ---

1440379_at	1.12340 "solute carrier family 1 (neutral amino acid transporter), member 5"	Slc1a5
1440380_at	1.13538 "Capping protein (actin filament) muscle Z-line, alpha 1"	Capza1
1440381_at	1.16733 RIKEN cDNA 2410085M17 gene	2410085M17Rik
1440382_at	0.94778 cDNA sequence BC051628	BC051628
1440383_at	0.85595 "DNA cross-link repair 1B, PSO2 homolog (S. cerevisiae)"	Dclre1b
1440384_at	0.96752 transmembrane and coiled coil domains 1	Tmcc1
1440385_at	0.80030 RIKEN cDNA 9530051E23 gene	9530051E23Rik
1440386_at	1.04218 glucuronyl C5-epimerase	Glce
1440387_at	0.94099 RIKEN cDNA 2610005L07 gene	2610005L07Rik
1440388_at	1.06160 Transcribed locus	---
1440389_at	0.94273 ---	---
1440390_at	0.91927 cDNA sequence BC025575	BC025575
1440391_at	1.49955 hypothetical protein LOC638038	LOC638038
1440392_at	1.29281 A kinase (PRKA) anchor protein 13	Akap13
1440393_at	0.87953 Odd Oz/ten-m homolog 2 (Drosophila)	Odz2
1440394_at	1.10164 ---	---
1440395_at	1.09625 RNA binding motif protein 21	Rbm21
1440396_at	1.19333 "Membrane protein, palmitoylated 6 (MAGUK p55 subfamily member 6)"	Mpp6
1440397_at	0.92007 ---	---
1440398_at	1.05509 ---	---
1440399_at	0.94885 Ubiquitin specific peptidase 32	Usp32
1440400_at	1.49395 ---	---
1440401_at	0.95921 ---	---
1440402_at	0.96444 RIKEN cDNA 4831426I19 gene	4831426I19Rik
1440403_at	1.24589 Rho guanine nucleotide exchange factor (GEF) 1	Arhgef1
1440404_at	1.61869 ---	---
1440405_at	1.08208 Adenylate cyclase 3	Adcy3
1440406_at	0.94641 G patch domain containing 3	Gpatc3
1440407_at	0.86331 ---	---
1440408_at	1.12810 RIKEN cDNA B830008J18 gene	B830008J18Rik
1440409_at	1.11822 RIKEN cDNA 2210401J11 gene	2210401J11Rik
1440410_at	0.96425 RIKEN cDNA A930011G23 gene	A930011G23Rik
1440411_at	0.89847 Transcribed locus	---
1440412_at	1.03841 "Immunoglobulin superfamily, member 4"	Igsf4d
1440413_at	0.99231 RIKEN cDNA A830006F12 gene	A830006F12Rik
1440414_at	1.04628 metal response element binding transcription factor 1	Mtf1
1440415_at	0.89050 "0 day neonate thymus cDNA, RIKEN full-length enriched library, clone:	
A430095E15	product:unclassifiable, full insert sequence"	---
1440416_at	1.06993 Ubiquitin specific peptidase 46	Usp46
1440417_at	2.54722 "DNA segment, Chr 19, ERATO Doi 409, expressed"	D19Ert409e
1440418_at	0.97417 ---	---
1440419_at	1.04683 "gene model 1679, (NCBI)"	Gm1679

1440420_at	0.91498	"Serine/threonine kinase 24 (STE20 homolog, yeast)"	Stk24
1440421_at	1.24849	RIKEN cDNA 2700085M18 gene	2700085M18Rik
1440422_at	0.95050	5-hydroxytryptamine (serotonin) receptor 1F	Htr1f
1440423_at	0.95865	zinc finger and BTB domain containing 37	Zbtb37
1440424_at	0.86899	"Troponin T2, cardiac"	Tnnt2
1440425_at	0.90666	---	---
1440426_at	0.94557	"nuclear factor of activated T-cells, cytoplasmic, calcineurin-dependent 2"	Nfatc2
1440427_at	1.02678	expressed sequence BB124205	BB124205
1440428_at	0.99122	sideroflexin 5	Sfxn5
1440429_at	1.16044	"cyclic AMP-regulated phosphoprotein, 21"	Arpp21
1440430_at	1.06966	RIKEN cDNA A130004G07 gene	A130004G07Rik
1440431_at	1.36636	Myeloid ecotropic viral integration site 1	Meis1
1440432_at	0.98878	Leucine zipper transcription factor-like 1	Lztfl1
1440433_at	1.04839	N-myristoyltransferase 2	Nmt2
1440434_at	1.18418	---	---
1440435_at	0.92155	kyphoscoliosis peptidase	Ky
1440436_at	0.93468	hypothetical protein A730030A06	A730030A06
1440437_at	1.02440	hect (homologous to the E6-AP (UBE3A) carboxyl terminus) domain and RCC1 (CHC1)-like domain (RLD) 1	Herc1
1440438_at	0.89073	WASP family 1	Wasf1
1440439_at	1.42308	Expressed sequence AI591476	AI591476
1440440_at	2.24390	---	---
1440441_at	1.48349	F-box and leucine-rich repeat protein 7	Fbxl7
1440442_at	0.96809	mitogen activated protein kinase kinase 7	Map2k7
1440443_at	1.18324	RIKEN cDNA E030016H06 gene	E030016H06Rik
1440444_at	1.42972	fatty acid desaturase 1	Fads1
1440445_at	1.00080	Pax6 opposite strand transcript 1	Pax6os1
1440446_at	0.97111	similar to naked cuticle 1 homolog	/// similar to naked cuticle 1 homolog
LOC634379	///	LOC671838	
1440447_at	0.89983	abhydrolase domain containing 2	Abhd2
1440448_at	1.25869	Transcribed locus	---
1440449_at	1.00411	leucine rich repeat containing 40	Lrrc40
1440450_at	1.01448	Heat shock protein 1-like	Hspa11
1440451_at	0.95263	RIKEN cDNA D730048J04 gene	D730048J04Rik
1440452_at	0.94188	dystrophin related protein 2	Drp2
1440453_at	1.02426	RIKEN cDNA 1700012C15 gene	1700012C15Rik
1440454_at	0.99632	RIKEN cDNA A530088I07 gene	A530088I07Rik
1440455_at	0.90338	expressed sequence AI848599	AI848599
1440456_at	0.95820	similar to KIAA0944 protein	LOC381255
1440457_at	1.18749	"Clathrin, heavy polypeptide (Hc)"	Cltc
1440458_at	1.06707	cytidine monophospho-N-acetylneuraminic acid hydroxylase	Cmah
1440459_at	1.42278	senataxin	Setx

1440460_at 0.92529 RIKEN cDNA 4930504O13 gene 4930504O13Rik
1440461_at 4.62722 RIKEN cDNA 0910001A06 gene 0910001A06Rik
1440462_at 1.61564 RIKEN cDNA B930068K11 gene B930068K11Rik
1440463_at 0.94147 N-glycanase 1 Ngly1
1440464_at 1.46557 "ELAV (embryonic lethal, abnormal vision, Drosophila)-like 1 (Hu antigen R)" Elavl1
1440465_at 0.91149 Cut-like 1 (Drosophila) Cutl1
1440466_at 1.08236 cDNA sequence BC003965 BC003965
1440467_at 0.91046 cDNA sequence BC028799 /// RIKEN cDNA 4922501C03 gene BC028799 /// 4922501C03Rik
1440468_at 0.96985 RIKEN cDNA D630023F18 gene D630023F18Rik
1440469_at 0.87690 hypothetical protein LOC629141 LOC629141
1440470_at 0.82802 Transcribed locus ---
1440471_x_at 0.92320 --- ---
1440472_at 0.94346 sperm associated antigen 17 Spag17
1440473_at 1.07927 RAS related protein 2a Rap2a
1440474_at 0.95671 --- ---
1440475_at 0.94870 expressed sequence AW011738 AW011738
1440476_at 0.94055 "DNA segment, Chr 6, ERATO Doi 474, expressed" D6Ertd474e
1440477_at 1.08207 --- ---
1440478_at 1.04339 "DNA segment, Chr 10, ERATO Doi 438, expressed" D10Ertd438e
1440479_at 0.96309 "Chromobox homolog 4 (Drosophila Pc class) /// 10 days lactation, adult female mammary gland cDNA, RIKEN full-length enriched library, clone:D730014L03 product:unclassifiable, full insert sequence" Cbx4
1440480_at 1.06670 cDNA sequence AB182283 AB182283
1440481_at 1.19721 signal transducer and activator of transcription 1 Stat1
1440482_at 1.15802 vacuolar protein sorting 13A (yeast) Vps13a
1440483_at 1.12924 "Pleckstrin homology domain containing, family M (with RUN domain) member 2" Plekhm2
1440484_at 0.90516 unc-5 homolog D (C. elegans) Unc5d
1440485_at 0.98938 --- ---
1440486_at 0.93815 RIKEN cDNA 2310037I24 gene 2310037I24Rik
1440487_at 1.10066 "0 day neonate cerebellum cDNA, RIKEN full-length enriched library, clone: C230056E17 product:unclassifiable, full insert sequence" ---
1440488_at 1.20283 --- ---
1440489_at 0.76386 BTB (POZ) domain containing 4 Btbd4
1440490_at 1.29365 "Membrane protein, palmitoylated 6 (MAGUK p55 subfamily member 6)" Mpp6
1440491_at 1.00865 "solute carrier family 1 (glial high affinity glutamate transporter), member 3" Slc1a3
1440492_at 1.16118 Exportin 6 Xpo6
1440493_at 1.03773 UDP-N-acetyl-alpha-D-galactosamine:polypeptide N-acetylgalactosaminyltransferase 10 Galnt10

1440494_at 1.01345 MIF4G domain containing Mif4gd
 1440495_at 0.93382 dorso-medial telencephalon gene 2 Dmt2
 1440496_at 1.06174 sorting and assembly machinery component 50 homolog (S. cerevisiae)
 Samm50
 1440497_at 0.95853 CDNA clone IMAGE:5690974 ---
 1440498_at 1.18761 RAB interacting factor Rabif
 1440499_at 0.88850 "DNA segment, Chr 9, ERATO Doi 26, expressed" D9Ertd26e
 1440500_at 1.05276 Mitogen activated protein kinase kinase kinase 10 Map3k10
 1440501_at 0.96591 expressed sequence AU041480 AU041480
 1440502_at 0.95993 RIKEN cDNA 4930485B16 gene 4930485B16Rik
 1440503_at 1.01806 "3 days neonate thymus cDNA, RIKEN full-length enriched library, clone:
 A630095O16 product:unclassifiable, full insert sequence" ---
 1440504_at 1.05682 Coiled-coil domain containing 39 Ccdc39
 1440505_at 0.83414 "phosphatidylinositol-specific phospholipase C, X domain containing 1" Plcxd1
 1440506_at 1.12205 "Solute carrier family 7 (cationic amino acid transporter, y+ system), member
 2" Slc7a2
 1440507_at 0.96569 CD151 antigen Cd151
 1440508_at 1.01978 "Adult male hippocampus cDNA, RIKEN full-length enriched library, clone:
 C630033O16 product:unclassifiable, full insert sequence" ---
 1440509_at 1.04273 SRY-box containing gene 30 Sox30
 1440510_at 0.86188 RIKEN cDNA C430002N11 gene C430002N11Rik
 1440511_at 1.31159 "Proteasome (prosome, macropain) activator subunit 4" Psme4
 1440512_at 1.16180 expressed sequence AW048066 AW048066
 1440513_at 2.05183 expressed sequence C80258 C80258
 1440514_at 1.00630 "DNA segment, Chr 3, ERATO Doi 508, expressed" D3Ertd508e
 1440515_at 1.02366 similar to Doublecortin domain-containing protein 2 /// similar to Doublecortin
 domain-containing protein 2 /// similar to Doublecortin domain-containing protein 2 LOC635529 ///
 LOC667452 /// LOC672313
 1440516_at 0.97277 "13 days embryo heart cDNA, RIKEN full-length enriched library, clone:
 D330017K01 product:unclassifiable, full insert sequence" ---
 1440517_x_at 0.97525 cytidine monophospho-N-acetylneuraminic acid hydroxylase Cmah
 1440518_at 0.97861 RIKEN cDNA 1700029I01 gene 1700029I01Rik
 1440519_at 1.00657 trans-acting transcription factor 8 Sp8
 1440520_a_at 1.02492 RIKEN cDNA 1700051A21 gene 1700051A21Rik
 1440521_x_at 0.88514 RIKEN cDNA 1700051A21 gene 1700051A21Rik
 1440522_at 0.88379 hypothetical protein B930075F07 B930075F07
 1440523_at 0.90481 retinal short chain dehydrogenase reductase 2 Rdhe2
 1440524_at 0.98187 "ubiquitously transcribed tetratricopeptide repeat gene, X chromosome" Utx
 1440525_at 0.79717 "In vitro fertilized eggs cDNA, RIKEN full-length enriched library,
 clone:7420418N15 product:unclassifiable, full insert sequence" ---
 1440526_at 1.02867 RIKEN cDNA D130029J02 gene D130029J02Rik
 1440527_at 0.81640 "Transcribed locus, strongly similar to NP_035188.1 platelet derived growth
 factor receptor, alpha polypeptide [Mus musculus]" ---

1440528_at 1.08273 Transcribed locus ---
1440529_at 1.09582 Transcribed locus ---
1440530_at 1.02586 expressed sequence AU019752 AU019752
1440531_at 0.93126 RNA binding motif protein 11 Rbm11
1440532_a_at 1.01714 "Adult male diencephalon cDNA, RIKEN full-length enriched library, clone:9330176F03 product:unclassifiable, full insert sequence" ---
1440533_at 1.06741 Bifunctional apoptosis regulator Bfar
1440534_at 1.04236 "Adult male hypothalamus cDNA, RIKEN full-length enriched library, clone:A230069K02 product:hypothetical protein, full insert sequence" ---
1440535_at 0.96372 uroporphyrinogen III synthase Uros
1440536_at 1.08932 "Solute carrier family 22 (organic cation transporter), member 5" Slc22a5
1440537_at 0.97191 "potassium channel, subfamily V, member 2" Kcnv2
1440538_at 0.99737 --- ---
1440539_at 1.16427 --- ---
1440540_at 1.50364 GLE1 RNA export mediator-like (yeast Gle11
1440541_at 1.11001 RIKEN cDNA 2810442I21 gene 2810442I21Rik
1440542_at 0.99110 RIKEN cDNA 7420416P09 gene 7420416P09Rik
1440543_at 1.13805 RIKEN cDNA D930036F22 gene D930036F22Rik
1440544_at 1.01878 "Solute carrier family 14 (urea transporter), member 2" Slc14a2
1440545_at 0.89276 "Adult male corpora quadrigemina cDNA, RIKEN full-length enriched library, clone:B230378A03 product:hypothetical protein, full insert sequence" ---
1440546_at 1.12331 RIKEN cDNA 9630002D21 gene 9630002D21Rik
1440547_at 1.07272 --- ---
1440548_at 1.19605 BTB (POZ) domain containing 9 Btbd9
1440549_at 1.31542 RIKEN cDNA B230334L07 gene B230334L07Rik
1440550_at 1.03427 "Adult male corpora quadrigemina cDNA, RIKEN full-length enriched library, clone:B230313K13 product:unclassifiable, full insert sequence" ---
1440551_at 2.04308 "DnaJ (Hsp40) homolog, subfamily C, member 1" Dnajc1
1440552_at 1.02347 Ubiquitin protein ligase E3 component n-recognin 1 Ubr1
1440553_at 1.02014 mitochondrial trans-2-enoyl-CoA reductase Mecr
1440554_at 1.08535 --- ---
1440555_at 0.99411 Ras-related GTP binding D Rragd
1440556_at 1.00940 --- ---
1440557_at 0.95838 imprinted gene in the Prader-Willi syndrome region Ipw
1440558_at 1.00868 --- ---
1440559_at 1.17749 "high mobility group AT-hook 2, pseudogene 1" Hmga2-ps1
1440560_at 1.12757 Yippee-like 2 (Drosophila) Ypel2
1440561_at 1.03892 expressed sequence C87259 C87259
1440562_at 0.97202 "Adult male corpora quadrigemina cDNA, RIKEN full-length enriched library, clone:B230304I02 product:hypothetical protein, full insert sequence" ---
1440563_at 0.92456 RIKEN cDNA 2010111I01 gene 2010111I01Rik
1440564_at 1.19076 prokineticin receptor 2 Prokr2
1440565_at 1.29988 Zinc finger and BTB domain containing 20 Zbtb20

1440566_at	0.87665	RIKEN cDNA D330005C11 gene	D330005C11Rik
1440567_at	1.43001	RIKEN cDNA D030051N19 gene	D030051N19Rik
1440568_at	1.00786	protection of telomeres 1	Pot1
1440569_at	1.02736	"solute carrier family 16 (monocarboxylic acid transporters), member 10"	
Slc16a10			
1440570_at	0.98678	Lipoma HMGIC fusion partner-like 3	Lhfp13
1440571_at	0.94512	"Eukaryotic translation initiation factor 4 gamma, 3"	Eif4g3
1440572_at	0.93576	dual specificity phosphatase 19	Dusp19
1440573_at	2.94189	ErbB2 interacting protein	ErbB2ip
1440574_at	0.93050	Recombination activating gene 1	Rag1
1440575_at	1.59884	heat shock protein 4	Hspa4
1440576_at	0.88318	"carboxypeptidase N, polypeptide 1"	Cpn1
1440577_at	1.10106	"Muskelin 1, intracellular mediator containing kelch motifs"	Mkln1
1440578_at	1.35553	Kinesin family member 13A	Kif13a
1440579_at	1.68560	---	---
1440580_at	1.21658	RIKEN cDNA C030014M07 gene	/// similar to RIKEN cDNA C920006C10
C030014M07Rik		/// LOC668212	
1440581_at	0.95683	RIKEN cDNA A530088H08 gene	A530088H08Rik
1440582_at	1.37406	itchy	/// RIKEN cDNA 8030492O04 gene Itch
1440583_at	0.94781	"DNA segment, Chr 6, Wayne State University 116, expressed"	D6Wsu116e
1440584_at	1.06744	RIKEN cDNA 9130221L21 gene	9130221L21Rik
1440585_at	1.01220	Syntaxin 6	Stx6
1440586_at	1.40725	RIKEN cDNA B430203I24 gene	B430203I24Rik
1440587_at	1.14036	---	---
1440588_at	0.99099	---	---
1440589_at	0.89684	Sterile alpha motif domain containing 12	Samd12
1440590_at	1.35796	---	---
1440591_at	0.95560	---	---
1440592_at	1.02789	CD80 antigen	Cd80
1440593_at	1.01989	Basonuclin 2	Bnc2
1440594_at	1.19252	"Protein phosphatase 1, catalytic subunit, gamma isoform"	Ppp1cc
1440595_at	1.04520	erythrocyte protein band 4.1-like 3	Epb4.113
1440596_at	1.03948	calcineurin binding protein 1	Cabin1
1440597_at	1.07118	---	---
1440598_at	0.95638	"syntrophin, gamma 1"	Sntg1
1440599_at	1.08539	olfactory receptor 658	OlfR658
1440600_at	0.95091	RIKEN cDNA 4930407I10 gene	4930407I10Rik
1440601_at	1.00584	Transcribed locus	---
1440602_at	1.08933	"Glutamate receptor, ionotropic, kainate 2 (beta 2)"	Grik2
1440603_at	1.02134	"Adult male corpus striatum cDNA, RIKEN full-length enriched library, clone: C030004I15 product:unclassifiable, full insert sequence"	---
1440604_at	1.08380	Riken cDNA 8030494B02 gene	8030494B02Rik
1440605_at	0.97461	"fascin homolog 2, actin-bundling protein, retinal (Strongylocentrotus	

purpuratus)" Fscn2
1440606_at 1.08326 Transmembrane protein 97 Tmem97
1440607_at 1.11705 Checkpoint suppressor 1 Ches1
1440608_at 0.96985 "10 days neonate skin cDNA, RIKEN full-length enriched library,
clone:4732488M12 product:unclassifiable, full insert sequence" ---
1440609_at 1.63761 mitogen-activated protein kinase kinase kinase 4 Map4k4
1440610_at 0.97447 CAS1 domain containing 1 Casd1
1440611_at 0.94331 RIKEN cDNA 6230409E13 gene 6230409E13Rik
1440612_at 0.99717 --- ---
1440613_at 0.89852 Per1-like domain containing 1 Perld1
1440614_at 1.05327 Rho GTPase activating protein 23 Arhgap23
1440615_at 1.42950 dual specificity phosphatase 16 Dusp16
1440616_at 1.00396 expressed sequence AU042410 AU042410
1440617_at 0.96269 carboxypeptidase A6 Cpa6
1440618_at 1.24519 hypothetical protein 9130416B15 9130416B15
1440619_at 0.88703 mitogen activated protein kinase 8 interacting protein 1 Mapk8ip1
1440620_at 0.95564 "RAB8A, member RAS oncogene family" Rab8a
1440621_at 1.25766 Kit ligand Kitl
1440622_at 1.22560 SEC15-like 2 (S. cerevisiae) Sec15l2
1440623_at 0.92397 G protein-coupled receptor 26 Gpr26
1440624_at 0.97363 "Protein tyrosine phosphatase, receptor type, D" Ptpnd
1440625_at 0.98720 RIKEN cDNA D930015E06 gene D930015E06Rik
1440626_at 1.00927 Homeo box D13 Hoxd13
1440627_at 1.07238 "ATPase, aminophospholipid transporter-like, class I, type 8A, member 2"
Atp8a2
1440628_at 1.08790 --- ---
1440629_at 1.03818 eukaryotic translation termination factor 1 Etf1
1440630_at 0.94717 "Adult male olfactory brain cDNA, RIKEN full-length enriched library,
clone:6430514A06 product:unclassifiable, full insert sequence" ---
1440631_at 1.08241 --- ---
1440632_at 0.97184 protocadherin beta 4 Pcdh4
1440633_at 1.23720 nei like 3 (E. coli) Neil3
1440634_at 1.04197 Glypican 6 Gpc6
1440635_at 0.91635 "palladin, cytoskeletal associated protein" Pall1
1440636_at 0.86051 Mitochondrial ribosomal protein L3 Mrpl3
1440637_at 2.22743 Intersectin 1 (SH3 domain protein 1A) Itsn1
1440638_at 0.96117 RIKEN cDNA A230107N01 gene A230107N01Rik
1440639_at 1.00129 "discs, large (Drosophila) homolog-associated protein 1" Dlgap1
1440640_at 0.91983 RIKEN cDNA 2010111I01 gene 2010111I01Rik
1440641_at 1.65187 vacuolar protein sorting 29 (S. pombe) Vps29
1440642_at 0.88897 RIKEN cDNA D630042P16 gene D630042P16Rik
1440643_at 1.47774 --- ---
1440644_at 1.02632 "poly (ADP-ribose) polymerase family, member 6" Parp6

1440645_at 1.03697 similar to deleted in malignant brain tumors 1 isoform c precursor
 LOC546006
 1440646_at 1.23291 CDNA sequence BC052040 BC052040
 1440647_at 1.14865 signal-induced proliferation-associated 1 like 1 Sipal11
 1440648_at 0.95970 Coenzyme Q10 homolog A (yeast) Coq10a
 1440649_at 1.10655 Testis derived transcript Tes
 1440650_at 1.02981 Slit homolog 2 (Drosophila) Slit2
 1440651_at 1.56735 Dual specificity phosphatase 16 Dusp16
 1440652_at 1.08427 ankyrin repeat domain 44 Ankrd44
 1440653_at 1.47720 Pleckstrin homology domain interacting protein Phip
 1440654_at 0.92468 Hypothetical protein LOC574403 MGC99845
 1440655_at 1.00052 Transcribed locus ---
 1440656_at 1.05228 "HECT, C2 and WW domain containing E3 ubiquitin protein ligase 1"
 Hecw1
 1440657_at 1.02136 Contactin 3 Cntn3
 1440658_at 1.04821 RIKEN cDNA E230022H04 gene E230022H04Rik
 1440659_at 1.19347 RIKEN cDNA 2610027L16 gene 2610027L16Rik
 1440660_at 1.13880 Nuclear factor I/A Nfia
 1440661_at 0.85726 RIKEN cDNA A430010J10 gene A430010J10Rik
 1440662_at 1.16525 "Ral guanine nucleotide dissociation stimulator,-like 1" Rgl1
 1440663_at 0.95566 Exportin 4 Xpo4
 1440664_at 0.95148 "cat eye syndrome chromosome region, candidate 5 homolog (human)"
 Cocr5
 1440665_at 0.84305 --- ---
 1440666_at 0.99131 RIKEN cDNA 4933430I17 gene 4933430I17Rik
 1440667_at 0.89376 "DNA cross-link repair 1B, PSO2 homolog (S. cerevisiae)" Dclre1b
 1440668_at 0.95989 ADAMTS-like 3 Adamts13
 1440669_at 0.96792 RIKEN cDNA B230217O12 gene B230217O12Rik
 1440670_at 1.10182 Transcribed locus ---
 1440671_at 2.00065 RIKEN cDNA A130012E19 gene A130012E19Rik
 1440672_at 0.84615 similar to zinc finger protein 541 /// similar to zinc finger protein 541 /// similar
 to zinc finger protein 541 LOC638394 /// LOC666528 /// LOC670763
 1440673_at 0.87791 Exonuclease domain containing 1 Exod1
 1440674_at 0.89958 EGF-like repeats and discoidin I-like domains 3 Edil3
 1440675_at 1.04282 Glucocorticoid receptor DNA binding factor 1 Grlf1
 1440676_at 0.77971 RIKEN cDNA A830043J08 gene A830043J08Rik
 1440677_at 1.03726 similar to hypothetical protein FLJ38281 LOC245305
 1440678_at 1.12246 Latent transforming growth factor beta binding protein 1 Ltbp1
 1440679_at 1.09541 Transcribed locus ---
 1440680_at 0.97840 Transcribed locus ---
 1440681_at 0.87571 "cholinergic receptor, nicotinic, alpha polypeptide 7" Chrna7
 1440682_at 0.91534 Odd Oz/ten-m homolog 4 (Drosophila) Odz4
 1440683_at 1.18235 RIKEN cDNA A930004D18 gene A930004D18Rik

1440684_at 0.98240 acyltransferase like 1 Ayt1l
1440685_at 0.91677 Calcitonin receptor-like Calcr1
1440686_at 1.10847 "protease, serine 27" Prss27
1440687_at 0.86750 "6 days neonate head cDNA, RIKEN full-length enriched library, clone:5430439N10 product:unclassifiable, full insert sequence" ---
1440688_at 1.06111 Rho GTPase activating protein 26 Arhgap26
1440689_at 1.05830 forkhead box K2 Foxk2
1440690_at 3.72533 Nipped-B homolog (Drosophila) Nipbl
1440691_at 0.90477 "cytochrome P450, family 2, subfamily j, polypeptide 6" Cyp2j6
1440692_at 0.98371 "gene model 364, (NCBI)" Gm364
1440693_at 1.31374 "palladin, cytoskeletal associated protein" Palld
1440694_at 1.05301 --- ---
1440695_at 1.15932 Zinc finger protein 40 Zfp40
1440696_at 0.90118 RIKEN cDNA 2300002D11 gene 2300002D11Rik
1440697_at 0.93829 expressed sequence AI746446 AI746446
1440698_at 0.91669 Unc-5 homolog C (C. elegans) Unc5c
1440699_at 0.88783 microtubule-associated protein 2 Mtap2
1440700_a_at 0.99851 rho/rac guanine nucleotide exchange factor (GEF) 18 Arhgef18
1440701_at 0.96966 RIKEN cDNA C530008M17 gene C530008M17Rik
1440702_at 0.95013 expressed sequence AU022297 AU022297
1440703_at 0.96185 RIKEN cDNA 2810429I04 gene 2810429I04Rik
1440704_at 1.00725 relaxin/insulin-like family peptide receptor 2 Rxfp2
1440705_at 0.92057 expressed sequence AU021720 AU021720
1440706_at 1.00960 bone morphogenetic protein 8b Bmp8b
1440707_at 1.01889 doublesex and mab-3 related transcription factor 3 Dmrt3
1440708_at 1.36964 "myosin, heavy polypeptide 9, non-muscle" Myh9
1440709_at 0.73449 RIKEN cDNA C130081A10 gene C130081A10Rik
1440710_at 0.94049 zinc finger protein 706 Zfp706
1440711_at 0.93139 RIKEN cDNA C630001G18 gene C630001G18Rik
1440712_at 1.10711 Poly(rC) binding protein 3 Pcbp3
1440713_at 0.93876 RIKEN cDNA 4933407H18 gene 4933407H18Rik
1440714_at 1.01854 Transcribed locus ---
1440715_s_at 1.01068 "DNA segment, Chr 11, ERATO Doi 497, expressed" D11Ertd497e
1440716_at 0.98829 RIKEN cDNA 6430604M11 gene 6430604M11Rik
1440717_at 1.16168 expressed sequence AA407881 AA407881
1440718_at 1.14709 cDNA sequence BC052328 BC052328
1440719_at 1.29104 GLE1 RNA export mediator-like (yeast Gle1l
1440720_s_at 1.04882 GLE1 RNA export mediator-like (yeast Gle1l
1440721_at 1.75311 RIKEN cDNA 5930433N17 gene 5930433N17Rik
1440722_at 1.09057 "DNA segment, Chr 19, ERATO Doi 386, expressed" D19Ertd386e
1440723_at 0.96592 --- ---
1440724_at 1.07072 expressed sequence AI854703 AI854703
1440725_at 1.01354 G protein-coupled receptor 63 Gpr63

1440726_at 1.07001 RIKEN cDNA A230062I15 gene A230062I15Rik
1440727_at 1.01171 Transcribed locus ---
1440728_at 0.95670 "Potassium large conductance calcium-activated channel, subfamily M, alpha member 1" Kcnma1
1440729_at 1.11865 epidermal growth factor receptor pathway substrate 15 Eps15
1440730_at 1.25815 --- ---
1440731_at 0.97454 expressed sequence AI413759 AI413759
1440732_at 0.88933 RIKEN cDNA D030018L15 gene D030018L15Rik
1440733_at 0.87282 RIKEN cDNA 1200009O22 gene 1200009O22Rik
1440734_at 1.05716 tau tubulin kinase 2 Ttbk2
1440735_at 1.06168 polymerase (RNA) III (DNA directed) polypeptide K Polr3k
1440736_at 1.07015 expressed sequence AI131651 AI131651
1440737_at 1.00610 InaD-like (Drosophila) Inadl
1440738_at 1.02367 RIKEN cDNA B020011L13 gene B020011L13Rik
1440739_at 0.71926 vascular endothelial growth factor C Vegfc
1440740_at 1.66776 "CTD (carboxy-terminal domain, RNA polymerase II, polypeptide A) small phosphatase like 2" Ctdspl2
1440741_at 1.04880 5-hydroxytryptamine (serotonin) receptor 1D Htr1d
1440742_at 1.04562 Enhancer of polycomb homolog 1 (Drosophila) Epc1
1440743_at 0.87304 RIKEN cDNA 4931428F04 gene 4931428F04Rik
1440744_at 0.95016 Peptidase inhibitor 16 Pi16
1440745_at 0.96355 PR domain containing 16 Prdm16
1440746_at 1.02138 MYST histone acetyltransferase 2 Myst2
1440747_at 1.01965 coproporphyrinogen oxidase Cpox
1440748_at 0.90392 --- ---
1440749_at 0.93525 Mitogen-activated protein kinase kinase kinase 13 Map3k13
1440750_at 0.99093 Transcribed locus ---
1440751_at 1.13479 E1A binding protein p400 Ep400
1440752_at 0.94117 Sal-like 4 (Drosophila) Sall4
1440753_at 0.89581 Transcribed locus ---
1440754_at 1.18692 Ankyrin repeat and KH domain containing 1 Ankhd1
1440755_at 1.14636 Neighbor of Brca1 gene 1 Nbr1
1440756_at 1.10321 --- ---
1440757_at 1.01240 "10 days neonate cerebellum cDNA, RIKEN full-length enriched library, clone: B930073N06 product:unclassifiable, full insert sequence" ---
1440758_at 1.01499 --- ---
1440759_at 0.97320 "Adult male corpora quadrigemina cDNA, RIKEN full-length enriched library, clone:B230398E24 product:hypothetical protein, full insert sequence" ---
1440760_at 0.96327 Lymphoid enhancer binding factor 1 Lef1
1440761_at 0.99515 RIKEN cDNA 4833422C13 gene 4833422C13Rik
1440762_at 0.97931 synapsin II Syn2
1440763_at 0.90759 Transcribed locus ---
1440764_at 1.11552 v-raf murine sarcoma 3611 viral oncogene homolog Araf

1440765_at	1.12170	Fraser syndrome 1 homolog (human)	Fras1	
1440766_at	0.97872	Guanine nucleotide binding protein-like 3 (nucleolar)	Gnl3	
1440767_at	1.03572	defensin beta 41	Defb41	
1440768_x_at	1.09606	RIKEN cDNA 1700052I22 gene	1700052I22Rik	
1440769_at	0.88926	RIKEN cDNA 2010305A19 gene	2010305A19Rik	
1440770_at	1.25012	B-cell leukemia/lymphoma 2	Bcl2	
1440771_at	1.58814	zinc finger with KRAB and SCAN domains 1	Zkscan1	
1440772_x_at	0.99607	RIKEN cDNA 9530077C05 gene	9530077C05Rik	
1440773_at	0.95187	RIKEN cDNA 2410004P22 gene	2410004P22Rik	
1440774_x_at	1.05095	cDNA sequence BC052883	BC052883	
1440775_at	0.92995	RIKEN cDNA 1700054K19 gene	1700054K19Rik	
1440776_at	0.88759	RIKEN cDNA 3732412D22 gene	3732412D22Rik	
1440777_x_at	1.03823	---	---	
1440778_x_at	0.93304	Zinc finger protein 712	Zfp712	
1440779_s_at	1.10108	"solute carrier family 5 (sodium/glucose cotransporter), member 9"	Slc5a9	
1440780_x_at	0.83708	RIKEN cDNA 1500015O10 gene	1500015O10Rik	
1440781_at	2.10805	RIKEN cDNA B830007D08 gene	B830007D08Rik	
1440782_at	1.04522	src family associated phosphoprotein 1	Scap1	
1440783_at	0.91892	RIKEN cDNA 4930529M08 gene	4930529M08Rik	
1440784_at	0.87576	Eukaryotic translation elongation factor 2	Eef2	
1440785_at	0.85488	relaxin/insulin-like family peptide receptor 1	Rxfp1	
1440786_x_at	0.88323	Sideroflexin 2	Sfxn2	
1440787_s_at	0.87080	Bardet-Biedl syndrome 10 (human)	Bbs10	
1440788_at	0.98904	Growth arrest specific 8	Gas8	
1440789_at	1.57314	neogenin	Neo1	
1440790_x_at	0.98022	RIKEN cDNA 5033414K04 gene	5033414K04Rik	
1440791_x_at	0.99689	"transcription elongation factor A (SII), 2"	Tcea2	
1440792_x_at	1.07254	"opioid receptor, sigma 1"	Oprs1	
1440793_at	1.04824	RIKEN cDNA 1700029M03 gene	1700029M03Rik	
1440794_x_at	1.00734	RIKEN cDNA 1700029M03 gene	1700029M03Rik	
1440795_x_at	1.03498	"rabaptin, RAB GTPase binding effector protein 2"	Rabep2	
1440796_at	1.39723	RIKEN cDNA A330021E22 gene	A330021E22Rik	
1440797_at	0.94223	Dlx6 opposite strand transcript 2	Dlx6os2	
1440798_x_at	1.00391	Leucine rich repeat containing 6 (testis)	Lrrc6	
1440799_s_at	0.81166	"FERM, RhoGEF and pleckstrin domain protein 2"	Farp2	
1440800_at	0.85874	hypothetical protein LOC665193	LOC665193	
1440801_s_at	1.05421	"adrenergic receptor kinase, beta 2"	Adrbk2	
1440802_at	1.07835	CLIP associating protein 2	Clasp2	
1440803_x_at	0.93787	tachykinin receptor 3	Tacr3	
1440804_at	1.12256	natural killer tumor recognition sequence	Nktr	
1440805_at	0.99861	---	---	
1440806_x_at	1.04156	RIKEN cDNA 5730526G10 gene	5730526G10Rik	
1440807_at	1.16917	"Membrane associated guanylate kinase, WW and PDZ domain containing 2"		

Magi2

1440808_x_at 1.09885 RIKEN cDNA E430002D04 gene E430002D04Rik
 1440809_at 0.96446 SWA-70 protein Swap70
 1440810_x_at 1.06732 "PREDICTED: Mus musculus similar to ribosomal protein L27a
 (LOC628200), mRNA" ---
 1440811_x_at 1.12731 "CD8 antigen, alpha chain" Cd8a
 1440812_at 0.97098 Transcribed locus ---
 1440813_s_at 1.05842 plexin B3 Plxnb3
 1440814_x_at 0.96779 heparan sulfate (glucosamine) 3-O-sulfotransferase 2 Hs3st2
 1440815_x_at 1.09445 hypothetical protein LOC624112 LOC624112
 1440816_x_at 0.96997 DEAD (Asp-Glu-Ala-Asp) box polypeptide 1 Ddx1
 1440817_x_at 0.92891 RIKEN cDNA G630024C07 gene G630024C07Rik
 1440818_s_at 0.98982 "splicing factor 3b, subunit 1" Sf3b1
 1440819_s_at 1.14074 cDNA sequence BC059842 BC059842
 1440820_x_at 0.93925 transmembrane and coiled-coil domains 2 Tmco2
 1440821_x_at 0.95034 outer dense fiber of sperm tails 1 Odf1
 1440822_x_at 0.96770 RalBP1 associated Eps domain containing protein Repl1
 1440823_x_at 0.99656 RIKEN cDNA D130058I21 gene D130058I21Rik
 1440824_at 1.24834 F-box protein 45 Fbxo45
 1440825_s_at 0.85762 coiled-coil domain containing 28A Ccdc28a
 1440826_s_at 1.01095 transmembrane protein 103 Tmem103
 1440827_x_at 0.93074 SRY-box containing gene 5 Sox5
 1440828_x_at 1.04891 PHD finger protein 7 Phf7
 1440829_x_at 0.91376 "tubulin tyrosine ligase-like family, member 11" Ttll11
 1440830_at 0.95403 G protein-coupled receptor 116 Gpr116
 1440831_at 1.02773 BTB and CNC homology 1 Bach1
 1440832_at 0.97221 "angiogenin, ribonuclease A family, member 4" Ang4
 1440833_at 0.83789 cell division cycle 2-like 5 (cholinesterase-related cell division controller)
 Cdc215
 1440834_at 1.03371 "solute carrier family 5 (sodium/glucose cotransporter), member 10" Slc5a10
 1440835_at 1.00800 zinc finger protein 27 Zfp27
 1440836_at 0.96132 SET domain containing 1B Setd1b
 1440837_at 1.02327 "histocompatibility 2, O region beta locus" H2-Ob
 1440838_at 1.08843 Expressed sequence AI852064 AI852064
 1440839_x_at 1.02890 RIKEN cDNA 1700065I16 gene 1700065I16Rik
 1440840_at 0.94655 RIKEN cDNA D630004K10 gene D630004K10Rik
 1440841_at 3.04281 "Tyrosine 3-monooxygenase/tryptophan 5-monooxygenase activation protein,
 epsilon polypeptide" Ywhae
 1440842_at 0.92819 RIKEN cDNA C230085N15 gene C230085N15Rik
 1440843_at 1.00932 RIKEN cDNA A230051N06 gene A230051N06Rik
 1440844_at 0.89623 transducer of ErbB-2.1 Tob1
 1440845_at 0.99648 RIKEN cDNA E030031F02 gene E030031F02Rik
 1440846_at 0.86285 RIKEN cDNA A930041I02 gene A930041I02Rik

1440847_at 2.03535 metastasis suppressor 1 Mtss1
1440848_at 1.17893 RIKEN cDNA 1110028C15 gene 1110028C15Rik
1440849_at 1.04092 RIKEN cDNA 6330417G04 gene 6330417G04Rik
1440850_at 1.21959 ring finger protein 12 Rnf12
1440851_at 1.15258 RIKEN cDNA 4933407N01 gene 4933407N01Rik
1440852_at 0.99060 RIKEN cDNA 4833405L16 gene 4833405L16Rik
1440853_at 1.13054 "ras homolog gene family, member T1" Rhot1
1440854_at 1.45811 RIKEN cDNA 2810403A07 gene 2810403A07Rik
1440855_at 0.96387 "Machado-Joseph disease (spinocerebellar ataxia 3, olivopontocerebellar ataxia 3, autosomal dominant, ataxin 3) homolog (human)" Mjd
1440856_at 0.81821 mitogen activated protein kinase 8 Mapk8
1440857_at 0.86985 "carbamoyl-phosphate synthetase 2, aspartate transcarbamylase, and dihydroorotase" Cad
1440858_at 1.00664 "Cdc2-related kinase, arginine/serine-rich" Crkrs
1440859_at 1.01155 A kinase (PRKA) anchor protein 6 Akap6
1440860_at 1.09333 mab-21-like 1 (C. elegans) Mab21l1
1440861_a_at 1.01184 --- ---
1440862_at 0.97682 G protein-coupled receptor 178 Gpr178
1440863_at 0.98406 --- ---
1440864_at 1.04994 expressed sequence AI836737 AI836737
1440865_at 1.06101 interferon induced transmembrane protein 6 Ifitm6
1440866_at 1.22137 eukaryotic translation initiation factor 2-alpha kinase 2 Eif2ak2
1440867_at 0.98768 sprouty homolog 4 (Drosophila) Spry4
1440868_at 0.80892 "GA repeat binding protein, beta 2" Gabpb2
1440869_x_at 0.83153 RIKEN cDNA 6720489N17 gene 6720489N17Rik
1440870_at 1.04429 PR domain containing 16 Prdm16
1440871_at 1.02036 "membrane associated guanylate kinase, WW and PDZ domain containing 1" Magi1
1440872_at 1.09230 CDNA sequence BC010304 BC010304
1440873_at 0.85840 --- ---
1440874_at 1.00302 "solute carrier organic anion transporter family, member 5A1" Slco5a1
1440875_a_at 1.28325 radical S-adenosyl methionine domain containing 1 Rsad1
1440876_at 1.01088 Retinitis pigmentosa 9 homolog (human) Rp9h
1440877_at 0.97327 Fibroblast growth factor receptor substrate 2 Frs2
1440878_at 0.97885 runt related transcription factor 1 Runx1
1440879_at 0.97811 ATP-binding cassette transporter sub-family A member 9 Abca9
1440880_at 0.85454 metallophosphoesterase 1 Mppe1
1440881_at 1.06063 bromodomain and WD repeat domain containing 3 Brwd3
1440882_at 1.34806 "low density lipoprotein receptor-related protein 8, apolipoprotein e receptor" Lrp8
1440883_at 1.32320 Transcribed locus ---
1440884_s_at 1.28282 RIKEN cDNA A530047J11 gene A530047J11Rik
1440885_at 1.29516 Ena-vasodilator stimulated phosphoprotein Evl

1440886_at 0.90878 cell division cycle 37 homolog (S. cerevisiae)-like 1 Cdc3711
1440887_at 0.96345 RIKEN cDNA C330005M16 gene C330005M16Rik
1440888_at 0.92944 oxytocin receptor Oxt
1440889_at 0.97159 cDNA sequence BC006965 BC006965
1440890_a_at 1.01932 expressed sequence BB114266 BB114266
1440891_at 0.98766 "glutamate receptor, ionotropic, AMPA4 (alpha 4)" Gria4
1440892_at 1.49503 CDNA sequence BC017647 BC017647
1440893_at 1.09595 --- ---
1440894_at 1.06877 "similar to CG4050-PA, isoform A" LOC670793
1440895_at 0.95670 --- ---
1440896_at 1.16129 "Superoxide dismutase 1, soluble" Sod1
1440897_at 1.01391 "Mus musculus, clone IMAGE:4458738, mRNA" ---
1440898_at 0.91402 RIKEN cDNA 4632412N22 gene 4632412N22Rik
1440899_at 1.06396 flavin containing monooxygenase 5 Fmo5
1440900_at 0.87480 G protein-coupled receptor 174 Gpr174
1440901_at 1.04703 "diacylglycerol kinase, beta" Dgkb
1440902_at 0.93096 RIKEN cDNA A330104H05 gene A330104H05Rik
1440903_at 1.26111 Hect (homologous to the E6-AP (UBE3A) carboxyl terminus) domain and
RCC1 (CHC1)-like domain (RLD) 1 Herc1
1440904_at 0.98415 SUMO/sentrin specific peptidase 5 Senp5
1440905_at 2.09451 Heparan sulfate 2-O-sulfotransferase 1 Hs2st1
1440906_at 1.11106 B-cell CLL/lymphoma 7C Bcl7c
1440907_at 0.99718 RIKEN cDNA A130038H09 gene A130038H09Rik
1440908_at 0.97859 hypothetical protein D030063E12 D030063E12
1440909_at 0.88720 "immunity-related GTPase family, cinema 1" Irgc1
1440910_at 0.92593 CDNA clone IMAGE:5252333 ---
1440911_at 0.93250 "procollagen, type XXIII, alpha 1" Col23a1
1440912_at 0.93688 ring finger protein 39 Rnf39
1440913_at 1.12424 cDNA sequence BC037704 BC037704
1440914_s_at 1.09285 aftiphilin Aftph
1440915_at 1.11591 M-phase phosphoprotein 9 Mphosph9
1440916_at 0.98832 RIKEN cDNA 2510049J12 gene 2510049J12Rik
1440917_at 1.09235 RIKEN cDNA 2900093K20 gene 2900093K20Rik
1440918_at 0.95275 --- ---
1440919_at 0.91175 "MOB1, Mps One Binder kinase activator-like 2C (yeast)" Mobk12c
1440920_at 0.85602 Transcribed locus ---
1440921_at 1.01948 "NACHT, LRR and PYD containing protein 12" Nalp12
1440922_at 0.93691 RIKEN cDNA 9130208D14 gene 9130208D14Rik
1440923_at 0.82992 --- ---
1440924_at 1.04554 M-phase phosphoprotein 1 Mphosph1
1440925_at 1.63343 --- ---
1440926_at 1.16294 FMS-like tyrosine kinase 1 Flt1
1440927_x_at 1.08512 RIKEN cDNA A330102K04 gene A330102K04Rik

1440928_at 0.88099 RIKEN cDNA D630037F22 gene D630037F22Rik
1440929_at 1.03044 --- ---
1440930_a_at 1.02844 "scratch homolog 2, zinc finger protein (Drosophila)" Scrt2
1440931_at 0.87666 "scratch homolog 2, zinc finger protein (Drosophila)" Scrt2
1440932_at 0.99213 --- ---
1440933_at 1.00144 RIKEN cDNA 4732460I02 gene 4732460I02Rik
1440934_at 0.82498 expressed sequence AW742931 AW742931
1440935_at 1.02298 Growth factor receptor bound protein 10 Grb10
1440936_at 1.04366 serine active site containing 1 Serac1
1440937_at 0.90906 RNA binding motif protein 41 Rbm41
1440938_at 0.99625 triggering receptor expressed on myeloid cells-like 2 Trem12
1440939_at 0.96984 Septin 6 6-Sep
1440940_at 1.01284 "Calcium channel, voltage-dependent, beta 1 subunit" Cacnb1
1440941_at 0.97293 hypothetical LOC552912 LOC552912
1440942_at 0.96581 "dapper homolog 2, antagonist of beta-catenin (xenopus)" Dact2
1440943_at 1.13946 RIKEN cDNA B230208H17 gene B230208H17Rik
1440944_at 1.01993 ankyrin repeat and sterile alpha motif domain containing 1B /// similar to AIDA-1b Anks1b /// LOC544718
1440945_at 0.89926 --- ---
1440946_at 1.02164 Zinc finger protein 483 Zfp483
1440947_at 0.93236 BAT2 domain containing 1 Bat2d
1440948_at 1.83560 Fyn-related kinase Frk
1440949_at 0.97573 --- ---
1440950_at 1.05504 Thymoma viral proto-oncogene 1 Akt1
1440951_x_at 0.95355 Wings apart-like homolog (Drosophila) Wapal
1440952_at 1.06208 MAD homolog 7 (Drosophila) Smad7
1440953_at 0.95079 RIKEN cDNA 1700096K18 gene 1700096K18Rik
1440954_at 3.32849 Pre B-cell leukemia transcription factor 1 Pbx1
1440955_at 1.00944 cysteine rich BMP regulator 2 (chordin like) Crim2
1440956_at 1.17220 coiled-coil domain containing 94 Ccdc94
1440957_at 1.00994 defensin beta 29 Defb29
1440958_at 1.00699 --- ---
1440959_s_at 0.91992 myoneurin Mynn
1440960_at 0.89984 "3 days neonate thymus cDNA, RIKEN full-length enriched library, clone: A630060L04 product:unclassifiable, full insert sequence" ---
1440961_at 0.86751 RIKEN cDNA 9130604C24 gene 9130604C24Rik
1440962_at 1.02239 "solute carrier family 8 (sodium/calcium exchanger), member 3" Slc8a3
1440963_at 0.64084 expressed sequence AI465270 AI465270
1440964_s_at 0.71864 expressed sequence AI465270 AI465270
1440965_at 0.96314 "Phosphatidylinositol glycan, class L" Pigl
1440966_at 1.63558 membrane-associated ring finger (C3HC4) 7 7-Mar
1440967_at 1.17592 Brain protein I3 /// BAI1-associated protein 2-like 1 Bri3 /// Baiap211
1440968_at 0.86610 --- ---

1440969_at 0.87856 cDNA sequence BC030308 BC030308
 1440970_at 0.92219 Transcribed locus ---
 1440971_x_at 0.92948 RIKEN cDNA G630024C07 gene G630024C07Rik
 1440972_at 1.01206 nuclear receptor-binding SET-domain protein 1 Nsd1
 1440973_at 0.94154 hypothetical LOC552874 LOC552874
 1440974_at 1.38087 RIKEN cDNA 9030203C11 gene 9030203C11Rik
 1440975_at 0.95312 RIKEN cDNA 1810057P16 gene 1810057P16Rik
 1440976_at 0.91741 RIKEN cDNA 1110035M17 gene 1110035M17Rik
 1440977_at 1.14860 A kinase (PRKA) anchor protein 13 Akap13
 1440978_at 0.86266 RIKEN cDNA C920011G20 gene C920011G20Rik
 1440979_at 0.98537 Insulin-like growth factor 2 receptor Igf2r
 1440980_at 1.10891 "potassium voltage-gated channel, subfamily H (eag-related), member 8" Kcnh8
 1440981_at 1.03834 p21 (CDKN1A)-activated kinase 7 Pak7
 1440982_at 1.56210 hypothetical LOC552909 LOC552909
 1440983_at 1.07075 Tetratricopeptide repeat domain 15 Ttc15
 1440984_at 0.71452 "bromodomain adjacent to zinc finger domain, 2B" Baz2b
 1440985_at 2.09516 RIKEN cDNA D130023J23 gene D130023J23Rik
 1440986_at 1.14319 RNA polymerase II associated protein 1 Rpap1
 1440987_at 0.81187 --- ---
 1440988_at 1.05032 "9.5 days embryo parthenogenote cDNA, RIKEN full-length enriched library, clone:B130035E07 product:unclassifiable, full insert sequence" ---
 1440989_at 0.91957 --- ---
 1440990_at 0.90983 cDNA sequence BC056349 BC056349
 1440991_at 1.02666 "pleckstrin homology domain containing, family G (with RhoGef domain) member 1" Plekhg1
 1440992_at 0.85367 RIKEN cDNA 3110052M02 gene 3110052M02Rik
 1440993_at 1.02047 Expressed sequence AW111846 AW111846
 1440994_at 0.96123 SNF2 histone linker PHD RING helicase Shprh
 1440995_at 0.97041 RIKEN cDNA 6430531B16 gene 6430531B16Rik
 1440996_at 1.14989 "Neural precursor cell expressed, developmentally down-regulated gene 8" Nedd8
 1440997_at 1.08694 trinucleotide repeat containing 6C Tnrc6c
 1440998_at 1.22814 RIKEN cDNA 2310021P13 gene 2310021P13Rik
 1440999_at 1.18611 zinc finger protein 697 Zfp697
 1441000_at 1.04608 "PREDICTED: Mus musculus similar to T-cell immunoglobulin and mucin domain containing 2 (LOC237749), mRNA" ---
 1441001_at 1.47376 expressed sequence AI225934 AI225934
 1441002_at 1.00388 "processing of precursor 1, ribonuclease P/MRP family, (S. cerevisiae)" Pop1
 1441003_at 0.86368 "excision repair cross-complementing rodent repair deficiency, complementation group 4" Ercc4
 1441004_at 0.84255 Exosome component 4 Exosc4
 1441005_at 0.97974 RIKEN cDNA 2610304G08 gene 2610304G08Rik
 1441006_at 0.95743 dihydrouridine synthase 4-like (S. cerevisiae) Dus4l

1441007_at 1.17949 RIKEN cDNA A330102I10 gene A330102I10Rik
 1441008_at 0.96874 dymeclin Dym
 1441009_at 0.94760 RIKEN cDNA 4732491K20 gene 4732491K20Rik
 1441010_at 0.83694 "Adult male vesicular gland cDNA, RIKEN full-length enriched library, clone:
 G630068J03 product:unclassifiable, full insert sequence" ---
 1441011_at 1.08093 Transcribed locus ---
 1441012_at 1.02617 Huntingtin interacting protein 2 Hip2
 1441013_at 1.27924 expressed sequence C81521 C81521
 1441014_at 0.97035 Alpha thalassemia/mental retardation syndrome X-linked homolog (human)
 Atrx
 1441015_at 1.04075 Fucosyltransferase 8 Fut8
 1441016_at 0.93874 "Adult male corpora quadrigemina cDNA, RIKEN full-length enriched library,
 clone:B230340D10 product:unclassifiable, full insert sequence" ---
 1441017_at 1.31410 "zinc finger, CCHC domain containing 14" Zcchc14
 1441018_at 1.07365 ubiquitin specific peptidase 24 Usp24
 1441019_at 1.10797 F-box only protein 3 Fbxo3
 1441020_at 1.27682 RIKEN cDNA 2900056M20 gene 2900056M20Rik
 1441021_at 1.03918 LETM1 domain containing 1 Letmd1
 1441022_at 0.96025 ariadne ubiquitin-conjugating enzyme E2 binding protein homolog 1
 (Drosophila) Arih1
 1441023_at 1.39265 "eukaryotic translation initiation factor 2, subunit 2 (beta)" Eif2s2
 1441024_at 0.97917 stromal interaction molecule 2 Stim2
 1441025_at 1.11526 RIKEN cDNA 3110047P20 gene 3110047P20Rik
 1441026_at 1.02864 "poly (ADP-ribose) polymerase family, member 4" Parp4
 1441027_at 0.88263 Transcribed locus ---
 1441028_at 1.01157 "DNA segment, Chr 2, ERATO Doi 210, expressed" D2Ertd210e
 1441029_at 1.21704 Axin 1 Axin1
 1441030_at 2.75005 retinoic acid induced 14 Rai14
 1441031_at 1.20080 "peptidyl arginine deiminase, type I" Padi1
 1441032_at 0.97555 "Adult male liver tumor cDNA, RIKEN full-length enriched library, clone:
 C730040K02 product:hypothetical protein, full insert sequence" ---
 1441033_at 0.98792 transmembrane and tetratricopeptide repeat containing 2 Tmtc2
 1441034_at 1.00153 Intersectin 1 (SH3 domain protein 1A) Itsn1
 1441035_at 1.02743 "potassium voltage-gated channel, Isk-related subfamily, member 1" Kcne1
 1441036_at 0.93139 --- ---
 1441037_at 1.02243 "Synuclein, alpha interacting protein (synphilin)" Sncap
 1441038_at 1.89278 Utrophin Utrn
 1441039_at 0.95760 RIKEN cDNA 4930532J02 gene 4930532J02Rik
 1441040_at 1.05289 Kruppel-like factor 12 Klf12
 1441041_at 1.08461 Zinc finger protein 746 Zfp746
 1441042_at 0.98966 fibroblast growth factor 1 Fgf1
 1441043_at 0.87985 "Adult male testis cDNA, RIKEN full-length enriched library,
 clone:4930579G14 product:unclassifiable, full insert sequence" ---

1441044_at	1.04595	RIKEN cDNA C130002M15 gene	C130002M15Rik
1441045_at	0.91905	---	---
1441046_at	1.21825	---	---
1441047_at	0.93819	"solute carrier family 45, member 4"	Slc45a4
1441048_at	0.92337	RIKEN cDNA 5930430L01 gene	5930430L01Rik
1441049_at	0.86770	"potassium voltage-gated channel, shaker-related, subfamily, member 6"	Kcna6
1441050_at	2.24202	"Striatin, calmodulin binding protein 3"	Strn3
1441051_at	0.92591	tripartite motif protein 27	Trim27
1441052_at	1.06495	RIKEN cDNA 2310035C23 gene	2310035C23Rik
1441053_at	0.96067	Transcribed locus	---
1441054_at	1.12446	"apolipoprotein L, 2"	Apol2
1441055_at	0.98097	paralemmin 2	Palm2
1441056_at	1.34720	ubiquitin specific peptidase 3	Usp3
1441057_at	1.34840	"Myosin, heavy polypeptide 10, non-muscle"	Myh10
1441058_at	1.22221	"inositol 1,4,5-trisphosphate 3-kinase B"	Itpkb
1441059_at	0.97498	RIKEN cDNA 1700049G17 gene	1700049G17Rik
1441060_at	0.93783	RIKEN cDNA 2900052L18 gene	2900052L18Rik
1441061_at	1.08615	"Visual cortex cDNA, RIKEN full-length enriched library, clone:K530018N01 product:unclassifiable, full insert sequence"	---
1441062_at	1.03791	"DEAD/H (Asp-Glu-Ala-Asp/His) box polypeptide 11 (CHL1-like helicase homolog, <i>S. cerevisiae</i>)"	Ddx11
1441063_at	1.14021	"eukaryotic translation initiation factor 2C, 3"	Eif2c3
1441064_at	0.94696	RIKEN cDNA E330039K12 gene	E330039K12Rik
1441065_at	0.99535	---	---
1441066_at	0.93481	"16 days embryo head cDNA, RIKEN full-length enriched library, clone: C130061K19 product:unclassifiable, full insert sequence"	---
1441067_at	0.90850	Expressed sequence AL033314	AL033314
1441068_at	1.02297	RIKEN cDNA A130001G05 gene	A130001G05Rik
1441069_at	0.82499	"zinc finger, DHHC domain containing 23"	Zdhhc23
1441070_at	0.96268	Homeo box A3	Hoxa3
1441071_at	0.93806	"potassium voltage-gated channel, subfamily Q, member 5"	Kcnq5
1441072_at	2.51876	"CUG triplet repeat, RNA binding protein 2"	Cugbp2
1441073_at	0.94834	---	---
1441074_at	0.99300	Ewing sarcoma breakpoint region 1	Ewsr1
1441075_at	1.10201	nitric oxide synthase trafficker	Nostrin
1441076_at	1.10743	Transcribed locus	---
1441077_at	1.15278	expressed sequence AW124847	AW124847
1441078_at	0.90602	RIKEN cDNA 1700023F02 gene	1700023F02Rik
1441079_at	1.02102	Transcribed locus	---
1441080_at	0.85457	Transcribed locus	---
1441081_a_at	1.08711	RIKEN cDNA 1110038B12 gene	1110038B12Rik
1441082_at	0.88956	Down syndrome cell adhesion molecule	Dscam
1441083_at	1.05805	cleft lip and palate associated transmembrane protein 1	Clptm1

1441084_at 0.99023 expressed sequence BE686333 BE686333
1441085_at 0.93122 RAR-related orphan receptor alpha Rora
1441086_at 0.88156 RIKEN cDNA 4930451I11 gene 4930451I11Rik
1441087_at 0.91416 RIKEN cDNA 2810011L19 gene 2810011L19Rik
1441088_at 0.97915 "zinc finger protein, subfamily 1A, 3 (Aiolos)" Zfpn1a3
1441089_at 1.30090 "eukaryotic translation initiation factor 2C, 3" Eif2c3
1441090_at 1.01975 Nuclear mitotic apparatus protein 1 Numal
1441091_at 1.00168 Transcribed locus ---
1441092_at 1.09066 RIKEN cDNA 9330159M07 gene 9330159M07Rik
1441093_at 1.16357 --- ---
1441094_at 0.95501 alcohol dehydrogenase 6B (class V) Adh6b
1441095_at 0.95777 RIKEN cDNA A430041B07 gene A430041B07Rik
1441096_at 1.03283 tumor suppressor candidate 5 Tusc5
1441097_at 0.94363 Transcribed locus ---
1441098_at 1.01760 poly(A)-specific ribonuclease (PARN)-like domain containing 1 Pnlcd1
1441099_at 1.03695 "Pleckstrin homology domain containing, family G (with RhoGef domain)
member 3" Plekhg3
1441100_at 2.34890 mbt domain containing 1 Mbtd1
1441101_at 0.97720 "HECT, C2 and WW domain containing E3 ubiquitin protein ligase 1"
Hecw1
1441102_at 1.01818 prolactin receptor Prlr
1441103_at 1.07871 "DNA segment, Chr 12, ERATO Doi 551, expressed" D12Ertd551e
1441104_at 0.93624 myosin Vb Myo5b
1441105_at 1.02671 "NOD-derived CD11c +ve dendritic cells cDNA, RIKEN full-length enriched
library, clone:F630006H17 product:hypothetical protein, full insert sequence" ---
1441106_at 1.15380 "zinc finger, AN1-type domain 5" Zfand5
1441107_at 0.98796 doublesex and mab-3 related transcription factor like family A2 Dmrta2
1441108_at 1.30955 "12 days embryo eyeball cDNA, RIKEN full-length enriched library, clone:
D230035N06 product:unclassifiable, full insert sequence" ---
1441109_at 0.98878 RIKEN cDNA 5430414B12 gene 5430414B12Rik
1441110_at 0.91003 leucine rich repeat containing 21 Lrrc21
1441111_at 0.94468 "15 days embryo embryonic body below diaphragm cDNA, RIKEN full-length
enriched library, clone:8230401F05 product:unclassifiable, full insert sequence" ---
1441112_at 2.37095 similar to Zinc finger BED domain containing protein 4 /// similar to Zinc finger
BED domain containing protein 4 /// similar to Zinc finger BED domain containing protein 4
LOC667118 /// LOC669797 /// LOC675760
1441113_at 0.91744 RIKEN cDNA A430071A18 gene A430071A18Rik
1441114_at 1.84154 RIKEN cDNA 9330156P08 gene 9330156P08Rik
1441115_at 0.97022 "DNA segment, Chr 18, ERATO Doi 232, expressed" D18Ertd232e
1441116_at 0.80150 --- ---
1441117_at 3.29931 "Ash1 (absent, small, or homeotic)-like (Drosophila)" Ash11
1441118_at 0.96549 cDNA sequence BC017647 BC017647
1441119_at 1.35477 GTPase activating RANGAP domain-like 1 Garnl1

1441120_at 1.08743 CEA-related cell adhesion molecule 16 Ceacam16
1441121_at 1.30671 PCI domain containing 2 Pcid2
1441122_at 0.92414 expressed sequence AU024342 AU024342
1441123_at 1.08751 open reading frame 34 ORF34
1441124_at 1.09632 "16 days embryo head cDNA, RIKEN full-length enriched library, clone:
C130032O22 product:unclassifiable, full insert sequence" ---
1441125_at 1.24234 SET domain containing 5 Setd5
1441126_at 1.12576 "Phenylalanine-tRNA synthetase-like, beta subunit" Farslb
1441127_at 0.93451 "12 days embryo spinal ganglion cDNA, RIKEN full-length enriched library,
clone:D130065D02 product:unclassifiable, full insert sequence" ---
1441128_at 1.00941 Renal tumor antigen Rage
1441129_at 1.31447 Septin 11 11-Sep
1441130_at 1.02698 Transcribed locus ---
1441131_at 1.17268 --- ---
1441132_at 0.90382 "Adult male liver tumor cDNA, RIKEN full-length enriched library, clone:
C730010N16 product:unclassifiable, full insert sequence" ---
1441133_at 0.83314 Syntaxin 6 Stx6
1441134_at 0.91901 RIKEN cDNA 6820424L24 gene 6820424L24Rik
1441135_at 1.04333 Zinc finger protein 672 Zfp672
1441136_at 0.97337 "7 days neonate cerebellum cDNA, RIKEN full-length enriched library, clone:
A730055K08 product:unclassifiable, full insert sequence" ---
1441137_at 1.00286 bicaudal C homolog 1 (Drosophila) Biccl
1441138_at 1.19388 forkhead box N2 Foxn2
1441139_at 0.83760 hypothetical protein B330003H21 B330003H21
1441140_at 2.10585 --- ---
1441141_at 1.36737 RIKEN cDNA C730024G19 gene C730024G19Rik
1441142_at 1.02614 RIKEN cDNA 2700081L22 gene 2700081L22Rik
1441143_at 0.96862 RIKEN cDNA E130309D14 gene E130309D14Rik
1441144_at 1.00600 "arrestin 3, retinal" Arr3
1441145_at 3.29511 RIKEN cDNA D030065N23 gene D030065N23Rik
1441146_at 0.99997 expressed sequence C76132 C76132
1441147_at 1.10736 "DNA segment, Chr 3, ERATO Doi 229, expressed" D3Ertd229e
1441148_at 1.13954 Fukuyama type congenital muscular dystrophy homolog (human) Fcmd
1441149_at 0.99571 --- ---
1441150_x_at 1.16966 Transcribed locus ---
1441151_at 1.16294 expressed sequence AU044157 AU044157
1441152_at 2.43917 --- ---
1441153_at 1.35048 Utrophin Utn
1441154_at 0.87144 Chromodomain helicase DNA binding protein 7 Chd7
1441155_at 1.38168 RIKEN cDNA 2810446P07 gene 2810446P07Rik
1441156_at 1.14488 "COP9 (constitutive photomorphogenic) homolog, subunit 3 (Arabidopsis
thaliana)" Cops3
1441157_at 1.02863 hypothetical LOC403344 C230006B20

1441158_at 1.08525 Chimerin (chimaerin) 2 Chn2
1441159_at 1.06521 "mediator of RNA polymerase II transcription, subunit 12 homolog (yeast)-like" Med12l
1441160_at 0.99597 RIKEN cDNA D330050G23 gene D330050G23Rik
1441161_at 1.04095 RIKEN cDNA B230216G23 gene B230216G23Rik
1441162_at 1.05277 Rho-associated coiled-coil forming kinase 1 Rock1
1441163_at 0.95650 "mediator of RNA polymerase II transcription, subunit 12 homolog (yeast)-like" Med12l
1441164_at 0.91124 "Phosphatidylinositol-4-phosphate 5-kinase, type II, beta" Pip5k2b
1441165_s_at 0.93620 calyntenin 2 Clstn2
1441166_at 0.94674 RIKEN cDNA A330050F15 gene A330050F15Rik
1441167_at 0.93397 Transcribed locus ---
1441168_at 0.95929 Interleukin 1 receptor accessory protein-like 2 Il1rapl2
1441169_at 0.97931 "UDP-Gal:betaGlcNAc beta 1,4-galactosyltransferase, polypeptide 6" B4galt6
1441170_a_at 0.92050 disabled homolog 2 (Drosophila) interacting protein Dab2ip
1441171_at 0.90175 "gene model 1614, (NCBI)" Gm1614
1441172_at 1.11097 "AF4/FMR2 family, member 3" Aff3
1441173_at 1.02921 RIKEN cDNA 4930451G09 gene 4930451G09Rik
1441174_a_at 1.16326 leishmanolysin-like (metallopeptidase M8 family) Lmln
1441175_at 0.96982 Aristaless related homeobox gene (Drosophila) Arx
1441176_at 0.96369 --- ---
1441177_at 1.78933 --- ---
1441178_at 0.84723 DTW domain containing 2 Dtwd2
1441179_at 1.00137 RIKEN cDNA 9530020O07 gene 9530020O07Rik
1441180_at 0.82697 RIKEN cDNA C330024D21 gene C330024D21Rik
1441181_at 0.96352 Transcribed locus ---
1441182_at 0.90618 bromodomain and WD repeat domain containing 3 Brwd3
1441183_at 1.01153 Jumonji domain containing 2B Jmjd2b
1441184_at 0.89074 "Transient receptor potential cation channel, subfamily C, member 4 associated protein" Trpc4ap
1441185_at 1.69194 Musashi homolog 2 (Drosophila) Msi2
1441186_at 0.91052 sal-like 3 (Drosophila) Sall3
1441187_at 1.04252 RIKEN cDNA E330037M01 gene E330037M01Rik
1441188_at 1.01686 --- ---
1441189_at 0.89626 Transcribed locus ---
1441190_at 0.91192 "actin related protein 2/3 complex, subunit 5-like" Arpc5l
1441191_at 1.16982 RIKEN cDNA 2810454H06 gene 2810454H06Rik
1441192_at 1.48019 selenocysteine lyase Scly
1441193_at 1.01671 --- ---
1441194_at 1.10060 Neuregulin 3 Nrg3
1441195_at 1.26195 RIKEN cDNA 6030487A22 gene 6030487A22Rik
1441196_at 1.12105 RIKEN cDNA C230076A16 gene C230076A16Rik

1441197_at	1.11705	RIKEN cDNA 9530059O14 gene	9530059O14Rik
1441198_at	0.92598	zinc finger protein 39	Zfp39
1441199_at	0.97416	"Protein tyrosine phosphatase, receptor type, M"	Ptprm
1441200_at	1.08521	Kruppel-like factor 3 (basic)	Klf3
1441201_at	0.95600	"Ubiquitously transcribed tetratricopeptide repeat gene, X chromosome"	Utx
1441202_at	0.92814	RAB GTPase activating protein 1-like	Rabgap11
1441203_at	1.01067	CDNA clone IMAGE:30543871	---
1441204_at	1.12962	RIKEN cDNA A130019P10 gene	A130019P10Rik
1441205_at	0.97172	---	---
1441206_at	1.16969	synaptopodin 2	Synpo2
1441207_at	0.97101	RIKEN cDNA 9630041N07 gene	9630041N07Rik
1441208_at	0.98356	Haloacid dehalogenase-like hydrolase domain containing 2	Hdhd2
1441209_at	1.06715	Ras and Rab interactor 2	Rin2
1441210_at	0.95301	Myosin IB	Myo1b
1441211_at	0.92725	"K+ voltage-gated channel, subfamily S, 2"	Kcns2
1441212_at	1.13212	RIKEN cDNA 6720467C03 gene	6720467C03Rik
1441213_at	0.95909	---	---
1441214_at	0.91397	exophilin 5	Exph5
1441215_at	0.96048	TraB domain containing	Trabd
1441216_at	0.97990	RIKEN cDNA 5330418N22 gene	5330418N22Rik
1441217_at	1.04973	RAB GTPase activating protein 1	Rabgap1
1441218_at	0.96695	Tetratricopeptide repeat domain 21A	ttc21a
1441219_at	1.09470	"11 days pregnant adult female ovary and uterus cDNA, RIKEN full-length enriched library, clone:5031405O07 product:unclassifiable, full insert sequence"	---
1441220_at	0.98759	---	---
1441221_at	1.24673	RIKEN cDNA 1700008J07 gene	1700008J07Rik
1441222_x_at	1.04352	---	---
1441223_at	0.92910	membrane-associated ring finger (C3HC4) 4	4-Mar
1441224_at	0.94474	Dystrophin related protein 2	Drp2
1441225_at	0.93234	Gelsolin	Gsn
1441226_at	0.98674	"spondin 1, (f-spondin) extracellular matrix protein"	Spon1
1441227_at	0.95482	CDNA clone IMAGE:6815954	---
1441228_at	1.08049	apolipoprotein L domain containing 1	Apold1
1441229_at	1.70293	RIKEN cDNA D230019N24 gene	D230019N24Rik
1441230_at	2.07786	Fibronectin type III domain containing 3B	Fndc3b
1441231_at	2.12481	"PREDICTED: Mus musculus hypothetical protein LOC626097 (LOC626097), mRNA"	---
1441232_at	0.96917	Transcribed locus	---
1441233_at	1.64009	Signal recognition particle 54	Srp54
1441234_at	0.83782	---	---
1441235_at	0.86916	nucleoporin 133	Nup133
1441236_at	1.07797	"solute carrier family 9 (sodium/hydrogen exchanger), member 3"	Slc9a3
1441237_at	1.08157	"DNA segment, Chr 5, ERATO Doi 163, expressed"	D5Erttd163e

1441238_at	1.66703	RIKEN cDNA 9030416H16 gene	9030416H16Rik
1441239_at	1.11540	potassium channel tetramerisation domain containing 18	Kctd18
1441240_at	1.08455	Replication protein A1	Rpa1
1441241_at	0.53763	RIKEN cDNA 9630013D21 gene	9630013D21Rik
1441242_at	0.99798	Dipeptidylpeptidase 4	Dpp4
1441243_at	0.96565	Zinc finger protein 532	Zfp532
1441244_at	0.93865	Transcribed locus	---
1441245_a_at	1.01740	"gene model 632, (NCBI)"	Gm632
1441246_s_at	1.14584	dihydropyrimidinase-like 5	Dpysl5
1441247_at	0.91709	"Phospholipase A2, group III"	Pla2g3
1441248_at	1.95364	---	---
1441249_at	1.13585	RIKEN cDNA 4930403D09 gene	4930403D09Rik
1441250_at	1.01759	---	---
1441251_a_at	1.05422	RIKEN cDNA 2010001A14 gene	2010001A14Rik
1441252_at	0.85800	RIKEN cDNA 2010001A14 gene	2010001A14Rik
1441253_at	2.47363	"Regulatory factor X, 3 (influences HLA class II expression)"	Rfx3
1441254_at	0.74450	RIKEN cDNA 1810008K04 gene	1810008K04Rik
1441255_at	0.99218	cadherin 3	Cdh3
1441256_at	1.02810	Wdr45 like	Wdr45l
1441257_x_at	1.09794	Wdr45 like	Wdr45l
1441258_at	0.89880	cDNA sequence AF529169	AF529169
1441259_s_at	0.69503	intraflagellar transport 122 homolog (Chlamydomonas)	Ift122
1441260_a_at	1.02900	---	---
1441261_at	0.89348	---	---
1441262_at	2.18488	STE20-like kinase (yeast)	Slk
1441263_a_at	1.07125	RIKEN cDNA A930005H10 gene	A930005H10Rik
1441264_x_at	1.10971	RIKEN cDNA A930005H10 gene	A930005H10Rik
1441265_at	0.87973	"Pleckstrin homology domain containing, family M (with RUN domain) member 2"	Plekhm2
1441266_at	1.60832	"Striatin, calmodulin binding protein 3"	Strn3
1441267_at	1.11832	Arginine/serine-rich coiled-coil 1	Rsrc1
1441268_at	0.88140	RIKEN cDNA 1500011B03 gene	1500011B03Rik
1441269_at	1.07824	Hydrocephalus inducing	Hydin
1441270_at	1.26070	A kinase (PRKA) anchor protein 2	Akap2
1441271_at	1.05875	Isocitrate dehydrogenase 3 (NAD+) beta	Idh3b
1441272_at	1.75585	---	---
1441273_at	1.02142	Leucine-rich repeats and calponin homology (CH) domain containing 2	Lrch2
1441274_at	1.06309	Cancer susceptibility candidate 3	Casc3
1441275_at	0.95820	kelch repeat and BTB (POZ) domain containing 8	Kbtbd8
1441276_at	0.99009	"protein tyrosine phosphatase, receptor type, K"	Ptprk
1441277_s_at	0.94462	"protein tyrosine phosphatase, receptor type, K"	Ptprk
1441278_at	0.92726	RIKEN cDNA 4930513N10 gene	4930513N10Rik
1441279_at	0.94621	RIKEN cDNA C430002E04 gene	C430002E04Rik

1441280_at 1.01000 "potassium channel, subfamily K, member 12" Kcnk12
1441281_s_at 0.97383 ninjurin 1 Ninj1
1441282_at 0.94821 Ssu72 RNA polymerase II CTD phosphatase homolog (yeast) Ssu72
1441283_at 0.95695 RIKEN cDNA 5730405I09 gene 5730405I09Rik
1441284_at 1.14503 Transcribed locus ---
1441285_at 1.08681 RIKEN cDNA 4833426J09 gene 4833426J09Rik
1441286_at 1.00022 similar to T-cell receptor alpha chain V region CTL-F3 precursor
LOC328483
1441287_at 1.20992 --- ---
1441288_at 1.09533 --- ---
1441289_at 0.83847 Transcribed locus ---
1441290_at 1.14544 RIKEN cDNA 4930506M07 gene 4930506M07Rik
1441291_at 0.92683 v-abl Abelson murine leukemia oncogene 1 Abl1
1441292_at 0.96283 Hypothetical gene Rp23-14f5.7 RP23-14F5.7
1441293_at 0.95350 "Adult male accessory axillary lymph node cDNA, RIKEN full-length enriched library, clone:G630004G16 product:unclassifiable, full insert sequence" ---
1441294_at 1.02496 --- ---
1441295_at 0.87997 "lectin, mannose-binding 2-like" Lman2l
1441296_at 0.97825 --- ---
1441297_at 1.53867 "mucin 6, gastric" Muc6
1441298_at 1.05759 --- ---
1441299_at 0.99311 Transcribed locus ---
1441300_at 0.92689 "potassium voltage-gated channel, subfamily F, member 1" Kcnf1
1441301_at 1.04069 Janus kinase and microtubule interacting protein 1 Jakmip1
1441302_at 1.14416 --- ---
1441303_at 0.94599 --- ---
1441304_at 1.06595 ribosomal protein L31 /// similar to ribosomal protein L31 /// similar to ribosomal protein L31 /// similar to ribosomal protein L31 /// similar to ribosomal protein L31 /// similar to ribosomal protein L31 /// similar to ribosomal protein L31 /// similar to ribosomal protein L31 /// similar to ribosomal protein L31 /// similar to ribosomal protein L31 /// similar to ribosomal protein L31 /// similar to ribosomal protein L31 /// similar to ribosomal protein L31 /// similar to ribosomal protein L31 /// similar to ribosomal protein L31 /// similar to ribosomal protein L31 /// similar to ribosomal protein L31 /// similar to ribosomal protein L31 /// similar to ribosomal protein L31
Rpl31 /// LOC435486 /// LOC544868 /// LOC625917 /// LOC626367 /// LOC631731 /// LOC637860 /// LOC638399 /// LOC667682 /// LOC668409 /// LOC668850 /// LOC671205 /// LOC672957 /// LOC673582 /// LOC676592
1441305_at 0.94484 "Adult male corpus striatum cDNA, RIKEN full-length enriched library, clone: C030044L04 product:unclassifiable, full insert sequence" ---
1441306_at 0.92493 RIKEN cDNA 6820408C15 gene 6820408C15Rik
1441307_at 0.98437 hypothetical protein LOC665891 /// hypothetical protein LOC670335
LOC665891 /// LOC670335
1441308_at 1.00742 myelin-associated oligodendrocytic basic protein Mobp
1441309_at 1.07586 "a disintegrin-like and metallopeptidase (reprolysin type) with thrombospondin type 1 motif, 10" Adamts10
1441310_at 0.87656 "A disintegrin-like and metallopeptidase (reprolysin type) with thrombospondin

type 1 motif, 15" Adamts15
 1441311_at 0.96232 "ribosomal protein S6 kinase, polypeptide 2" Rps6ka2
 1441312_at 1.00065 Cyclin M1 Cnnm1
 1441313_x_at 0.93011 LIM homeobox protein 9 Lhx9
 1441314_at 1.02651 inner centromere protein Incenp
 1441315_s_at 1.42136 "solute carrier family 19 (thiamine transporter), member 2" Slc19a2
 1441316_at 1.00909 similar to wingless related MMTV integration site 8b LOC677060
 1441317_x_at 0.96901 janus kinase and microtubule interacting protein 1 Jakmip1
 1441318_at 0.82854 "jumonji, AT rich interactive domain 1A (Rbp2 like)" Jarid1a
 1441319_at 1.27461 RNA binding motif protein 5 Rbm5
 1441320_a_at 0.85227 expressed sequence AI413194 AI413194
 1441321_at 1.23194 hypothetical LOC552911 LOC552911
 1441322_at 1.01266 src family associated phosphoprotein 1 Scap1
 1441323_at 1.05308 Transcribed locus ---
 1441324_at 1.19373 Zinc finger protein 395 Zfp395
 1441325_at 1.07222 NF-kappaB repressing factor Nkrf
 1441326_at 0.90212 ceruloplasmin Cp
 1441327_a_at 1.02682 "signal sequence receptor, alpha" Ssr1
 1441328_at 0.97735 Phosphatidylinositol binding clathrin assembly protein Picalm
 1441329_at 1.13380 galanin receptor 1 Galr1
 1441330_at 0.96571 crumbs homolog 1 (Drosophila) Crb1
 1441331_at 1.37736 RIKEN cDNA A230061C15 gene A230061C15Rik
 1441332_at 1.07627 --- ---
 1441333_at 0.94844 Trichorhinophalangeal syndrome I (human) Trps1
 1441334_at 1.01737 Transcribed locus ---
 1441335_at 0.98629 RIKEN cDNA 4930474N05 gene 4930474N05Rik
 1441336_at 0.96502 nuclear protein in testis Nut
 1441337_at 1.03918 Transcribed locus ---
 1441338_at 0.86250 RIKEN cDNA 5930412G12 gene 5930412G12Rik
 1441339_at 0.94970 chromodomain helicase DNA binding protein 9 Chd9
 1441340_at 1.10193 centrosomal protein 68 Cep68
 1441341_at 0.91889 Transcribed locus ---
 1441342_at 0.98590 Dipeptidylpeptidase 4 Dpp4
 1441343_at 0.92178 --- ---
 1441344_at 1.03992 expressed sequence C80197 C80197
 1441345_at 1.12059 expressed sequence C80143 C80143
 1441346_at 1.22740 RIKEN cDNA 2810421E14 gene 2810421E14Rik
 1441347_at 1.37617 HIV-1 Rev binding protein-like Hrbl
 1441348_at 1.56841 "Zinc finger protein 422, related sequence 1" Zfp422-rs1
 1441349_at 1.25304 hypothetical LOC552904 LOC552904
 1441350_at 1.15388 --- ---
 1441351_at 1.56791 Ankyrin repeat and IBR domain containing 1 Ankib1
 1441352_at 1.34850 RIKEN cDNA 9430034N14 gene 9430034N14Rik

1441353_at 0.91895 RIKEN cDNA 5330421F07 gene 5330421F07Rik
1441354_at 1.35950 expressed sequence AU018740 AU018740
1441355_at 1.32001 RIKEN cDNA E030047P09 gene E030047P09Rik
1441356_at 1.01544 src homology 2 domain-containing transforming protein C2 Shc2
1441357_at 0.94118 kin of IRRE like 3 (Drosophila) Kirrel3
1441358_at 0.79682 protocadherin beta 16 Pcdhb16
1441359_at 2.31409 RIKEN cDNA 9230115F04 gene 9230115F04Rik
1441360_at 1.40431 Ribosomal protein S6 kinase polypeptide 3 Rps6ka3
1441361_at 1.12013 multiple PDZ domain protein Mpdz
1441362_at 1.32393 "Protein tyrosine phosphatase, receptor type, G" Ptpnrg
1441363_at 0.95325 --- ---
1441364_at 0.92247 GATA binding protein 4 Gata4
1441365_at 0.97806 forkhead box P2 Foxp2
1441366_at 1.02790 --- ---
1441367_a_at 1.06488 Transcribed locus ---
1441368_at 0.93094 "12 days embryo embryonic body between diaphragm region and neck cDNA,
RIKEN full-length enriched library, clone:9430084D06 product:unclassifiable, full insert sequence"

1441369_at 0.95482 RIKEN cDNA C030017B01 gene C030017B01Rik
1441370_at 1.43465 "H2A histone family, member Z" H2afz
1441371_at 0.99376 plexin A4 Plxna4
1441372_at 0.90228 RIKEN cDNA 5930405F01 gene 5930405F01Rik
1441373_at 1.03921 Musashi homolog 2 (Drosophila) Msi2
1441374_at 0.98992 expressed sequence C79798 C79798
1441375_at 1.36277 Leucine-rich repeats and immunoglobulin-like domains 1 Lrig1
1441376_at 1.15450 gamma-aminobutyric acid (GABA-A) receptor-associated protein-like 2
Gabarapl2
1441377_at 1.07682 kinesin family member 6 Kif6
1441378_at 0.97170 --- ---
1441379_at 0.99100 expressed sequence AU045094 AU045094
1441380_at 0.97221 RIKEN cDNA 2810439F02 gene 2810439F02Rik
1441381_at 0.99276 RIKEN cDNA 9430029E18 gene 9430029E18Rik
1441382_at 1.00251 G protein-coupled receptor 101 Gpr101
1441383_at 1.16110 RIKEN cDNA 1500002O10 gene 1500002O10Rik
1441384_at 1.01826 Transcribed locus ---
1441385_at 0.88399 "ras homolog gene family, member H" Rhoh
1441386_at 1.86338 Rap guanine nucleotide exchange factor (GEF) 1 Rapgef1
1441387_at 1.14811 cDNA sequence BC030343 BC030343
1441388_at 1.11759 "Adult male hippocampus cDNA, RIKEN full-length enriched library, clone:
C630020O05 product:unclassifiable, full insert sequence" ---
1441389_at 0.93517 Transcribed locus ---
1441390_at 0.87157 speedy homolog A (Drosophila) Spdya
1441391_at 0.89985 "Guanine nucleotide binding protein (G protein), beta polypeptide 1-like"

Gnb11

1441392_at	0.94608	Transcribed locus	---
1441393_at	0.94679	---	---
1441394_at	1.06477	expressed sequence C76554	C76554
1441395_at	1.00156	---	---
1441396_at	0.98114	"UDP-Gal:betaGlcNAc beta 1,3-galactosyltransferase, polypeptide 1"	
B3galt1			
1441397_at	1.44223	Par-3 (partitioning defective 3) homolog (C. elegans)	Pard3
1441398_at	1.04413	---	---
1441399_at	0.96955	---	---
1441400_at	1.40589	RIKEN cDNA 1810029B16 gene	1810029B16Rik
1441401_at	0.92640	expressed sequence C79329	C79329
1441402_at	1.17333	similar to cysteine and glycine-rich protein 2 binding protein	LOC674229
1441403_at	1.45120	RIKEN cDNA 6430501K19 gene	6430501K19Rik
1441404_at	1.26007	"Platelet-activating factor acetylhydrolase, isoform 1b, beta1 subunit"	Pafah1b1
1441405_at	0.91554	RIKEN cDNA 9630014M24 gene	9630014M24Rik
1441406_at	1.00589	---	---
1441407_at	1.30218	ROD1 regulator of differentiation 1 (S. pombe)	Rod1
1441408_at	1.10312	expressed sequence AU044856	AU044856
1441409_at	0.84376	Grp94 neighboring nucleotidase variant 4	Gnn
1441410_at	1.53095	"DNA segment, Chr 2, ERATO Doi 112, expressed"	D2Erttd112e
1441411_at	1.47836	LIM and senescent cell antigen-like domains 1	Lims1
1441412_s_at	0.94334	tripartite motif-containing 45	Trim45
1441413_at	1.04878	Catalase	Cat
1441414_at	1.13682	Eukaryotic translation elongation factor 2	Eef2
1441415_at	1.71300	"Sprouty-related, EVH1 domain containing 2"	Spred2
1441416_at	1.11227	"inositol polyphosphate-4-phosphatase, type II"	Inpp4b
1441417_at	1.17198	"STT3, subunit of the oligosaccharyltransferase complex, homolog A (S. cerevisiae)"	Stt3a
1441418_at	1.00950	Src family associated phosphoprotein 1	Scap1
1441419_at	1.08768	RIKEN cDNA 1700030G06 gene	1700030G06Rik
1441420_at	1.04806	"immunoglobulin superfamily, member 9"	Igsf9
1441421_at	1.14562	RIKEN cDNA C530030P08 gene	C530030P08Rik
1441422_at	1.00264	growth arrest-specific 2 like 1	Gas2l1
1441423_at	0.93136	endothelin converting enzyme 1	Ece1
1441424_at	1.09790	Mitofusin 2	Mfn2
1441425_at	1.01618	Transcribed locus	---
1441426_at	1.40849	CDNA sequence BC002230	BC002230
1441427_at	1.01139	von Willebrand factor C domain containing 2	Vwc2
1441428_at	1.05751	"Adult male liver tumor cDNA, RIKEN full-length enriched library, clone: C730003N19 product:unclassifiable, full insert sequence"	---
1441429_at	1.06731	insulin receptor substrate 4	Irs4
1441430_at	1.00137	Transcribed locus	---

1441431_at	0.95986	RIKEN cDNA 1700041C02 gene	1700041C02Rik
1441432_at	1.06985	RIKEN cDNA C820005J03 gene	C820005J03Rik
1441433_at	1.01347	Transcribed locus	---
1441434_at	0.94913	RIKEN cDNA 2810426N06 gene	2810426N06Rik
1441435_at	1.97608	Transducin (beta)-like 1 X-linked	Tbl1x
1441436_at	1.05158	dual specificity phosphatase 12	Dusp12
1441437_at	2.25442	"RNA binding motif, single stranded interacting protein 1"	Rbms1
1441438_at	1.30377	glypican 6	Gpc6
1441439_at	0.95973	urocortin 3	Ucn3
1441440_at	1.38398	autophagy-related 4C (yeast)	Atg4c
1441441_at	1.80985	RIKEN cDNA 2010007L08 gene	2010007L08Rik
1441442_at	0.95947	---	---
1441443_at	0.84644	Testis expressed gene 2	Tex2
1441444_at	1.80317	Neurobeachin like 1	Nbeal1
1441445_at	1.16438	period homolog 3 (Drosophila)	Per3
1441446_at	1.02144	---	---
1441447_at	1.11661	nucleolar protein 4	Nol4
1441448_at	1.19802	Pumilio 2 (Drosophila)	Pum2
1441449_at	0.95248	"jumonji, AT rich interactive domain 1C (Rbp2 like)"	Jarid1c
1441450_s_at	1.40117	"jumonji, AT rich interactive domain 1C (Rbp2 like)"	Jarid1c
1441451_at	0.97701	bromodomain and WD repeat domain containing 3	Brwd3
1441452_at	1.90202	"LSM1 homolog, U6 small nuclear RNA associated (S. cerevisiae)"	Lsm1
1441453_at	0.97907	RIKEN cDNA 8030453O22 gene	8030453O22Rik
1441454_at	0.89189	zinc fingers and homeoboxes 3	Zhx3
1441455_at	1.10852	CAMP-regulated phosphoprotein 19	Arpp19
1441456_at	0.97462	matrix metalloproteinase 24	Mmp24
1441457_at	1.46392	---	---
1441458_at	0.97472	RAS p21 protein activator 4	Rasa4
1441459_at	0.99742	"16 days embryo head cDNA, RIKEN full-length enriched library, clone: C130020A17 product:unclassifiable, full insert sequence"	---
1441460_at	1.16945	FGFR1 oncogene partner 2	Fgfr1op2
1441461_at	1.02662	FERM domain containing 4A	Frmd4a
1441462_at	0.90176	dedicator of cytokinesis 4	Dock4
1441463_at	1.02805	---	---
1441464_at	1.05676	RIKEN cDNA A630035D09 gene	A630035D09Rik
1441465_at	1.27894	"CCR4-NOT transcription complex, subunit 3"	Cnot3
1441466_at	1.27375	"splicing factor, arginine/serine-rich 10 (transformer 2 homolog, Drosophila)"	Sfrs10
1441467_at	1.75018	Tetraspanin 5	Tspan5
1441468_at	0.93872	"PREDICTED: Mus musculus similar to dachsous 2 isoform 2 (LOC627094), mRNA"	---
1441469_at	1.04569	dedicator of cytokinesis 4	Dock4
1441470_at	1.11597	expressed sequence AV249152	AV249152

1441471_at 0.88987 RIKEN cDNA 2810002D19 gene 2810002D19Rik
1441472_at 1.18324 CDC-like kinase 3 Clk3
1441473_at 1.15487 hypothetical protein LOC666775 /// hypothetical protein LOC671556
LOC666775 /// LOC671556
1441474_at 0.94384 RIKEN cDNA C130070B15 gene C130070B15Rik
1441475_at 1.51071 PTK2 protein tyrosine kinase 2 Ptk2
1441476_at 0.90846 suppressor of cytokine signaling 2 Socs2
1441477_at 1.40442 calumenin Calu
1441478_at 0.95130 RIKEN cDNA 9930013L23 gene 9930013L23Rik
1441479_at 1.83176 "Protein phosphatase 1B, magnesium dependent, beta isoform" Ppm1b
1441480_at 1.33689 Signal recognition particle 54 Srp54
1441481_at 1.08024 microfibrillar-associated protein 3-like Mfap3l
1441482_at 1.09410 --- ---
1441483_at 0.85445 "SLIT and NTRK-like family, member 2" Slitrk2
1441484_at 1.25522 Transcription factor 2 Tcf2
1441485_at 1.02113 "Yip1 domain family, member 6" Yipf6
1441486_at 1.20201 RIKEN cDNA C430014M02 gene C430014M02Rik
1441487_at 1.03991 tripartite motif protein 2 Trim2
1441488_at 0.90089 Transcribed locus ---
1441489_at 1.00137 hypothetical LOC552878 LOC552878
1441490_at 1.02709 transmembrane protein 39a Tmem39a
1441491_at 1.01650 RIKEN cDNA A330068G13 gene A330068G13Rik
1441492_at 1.22197 Zinc finger protein 64 Zfp64
1441493_at 0.92207 ELKS/RAB6-interacting/CAST family member 1 Erc1
1441494_at 1.02207 Retinoblastoma 1 Rb1
1441495_at 0.84588 potassium channel tetramerisation domain containing 8 Kctd8
1441496_at 0.84672 tudor domain containing 3 Tdrd3
1441497_at 0.97712 expressed sequence AA388235 AA388235
1441498_at 1.05790 "Protein tyrosine phosphatase, receptor type, D" Ptpnd
1441499_at 1.27782 "Adult male hypothalamus cDNA, RIKEN full-length enriched library, clone:
A230096K23 product:39 KDA ANTIGEN homolog [Leishmania donovani], full insert sequence" ---
1441500_at 0.99167 --- ---
1441501_at 1.07875 Serpine1 mRNA binding protein 1 Serbp1
1441502_at 1.16199 Leucine rich repeat and fibronectin type III domain containing 5 Lrfn5
1441503_at 1.01099 doublecortin domain containing 2a Dcdc2a
1441504_at 1.03374 "gene model 817, (NCBI)" Gm817
1441505_at 2.35778 --- ---
1441506_at 1.03156 decorin Dcn
1441507_at 1.05655 spectrin beta 2 Spnb2
1441508_at 0.92514 "Adult retina cDNA, RIKEN full-length enriched library, clone:A930026H04
product:hypothetical protein, full insert sequence" ---
1441509_at 0.86253 RIKEN cDNA A130009I22 gene A130009I22Rik
1441510_at 1.02672 "Ras homolog gene family, member f" Rhof

1441511_at 1.03958 "potassium channel, subfamily T, member 1" Kcnt1
1441512_at 0.86169 RIKEN cDNA 6030445D17 gene 6030445D17Rik
1441513_at 0.95953 TRAF family member-associated Nf-kappa B activator Tank
1441514_at 0.86228 "3-oxoacyl-ACP synthase, mitochondrial" Oxsm
1441515_at 0.94753 --- ---
1441516_a_at 0.94249 RIKEN cDNA C130050O18 gene C130050O18Rik
1441517_at 1.08234 "Sprouty protein with EVH-1 domain 1, related sequence" Spred1
1441518_at 1.10818 "Adult retina cDNA, RIKEN full-length enriched library, clone:A930003C13
product:unclassifiable, full insert sequence" ---
1441519_at 1.05582 nidogen 1 Nid1
1441520_at 0.82263 "asp (abnormal spindle)-like, microcephaly associated (Drosophila)" Aspm
1441521_at 1.08109 "Calcium channel, voltage-dependent, beta 2 subunit" Cacnb2
1441522_at 1.22810 expressed sequence AW495713 AW495713
1441523_at 0.98745 neurofibromatosis 1 Nf1
1441524_at 0.96543 "cytoplasmic tyrosine kinase, Dscr28C related (Drosophila)" Tec
1441525_at 1.09591 NIMA (never in mitosis gene a)-related expressed kinase 11 Nek11
1441526_at 1.32109 Mbt domain containing 1 Mbtd1
1441527_at 0.93686 RIKEN cDNA 4930558J18 gene 4930558J18Rik
1441528_at 1.24686 tetratricopeptide repeat domain 15 Ttc15
1441529_at 1.11164 RIKEN cDNA 9130230N09 gene 9130230N09Rik
1441530_at 1.05718 Muscleblind-like 3 (Drosophila) Mbnl3
1441531_at 0.98233 "phospholipase C, beta 4" Plcb4
1441532_at 0.98631 Transcribed locus ---
1441533_at 1.04360 "Adult male diencephalon cDNA, RIKEN full-length enriched library,
clone:9330162K03 product:unclassifiable, full insert sequence" ---
1441534_at 0.86327 expressed sequence C86753 C86753
1441535_at 1.87787 Myeloid/lymphoid or mixed lineage-leukemia translocation to 3 homolog
(Drosophila) Mllt3
1441536_at 1.89533 3-hydroxy-3-methylglutaryl-Coenzyme A synthase 1 Hmgcs1
1441537_at 1.03182 RecQ protein-like 5 Recql5
1441538_at 0.92460 Dipeptidylpeptidase 6 Dpp6
1441539_at 3.48177 --- ---
1441540_at 1.03996 Nuclear receptor coactivator 7 Ncoa7
1441541_at 0.90690 Expressed sequence AW011752 AW011752
1441542_at 0.93392 dystrophin related protein 2 Drp2
1441543_at 1.05298 Eyes absent 3 homolog (Drosophila) Eya3
1441544_at 1.16713 Cyclin M3 Cnm3
1441545_at 1.72850 RIKEN cDNA 9230115F04 gene 9230115F04Rik
1441546_at 1.48973 hypothetical LOC624524 LOC624524
1441547_at 1.49462 Nuclear factor I/A Nfia
1441548_at 0.93004 FERM domain containing 4B Frmd4b
1441549_at 1.68797 --- ---
1441550_at 1.07502 RIKEN cDNA 9330184L24 gene 9330184L24Rik

1441551_at 1.05178 myopalladin Mypn
1441552_at 0.91232 RIKEN cDNA A430107P09 gene A430107P09Rik
1441553_at 0.99361 androgen-induced proliferation inhibitor Aprin
1441554_at 0.93863 RIKEN cDNA 4833446K15 gene 4833446K15Rik
1441555_at 1.06659 --- ---
1441556_at 1.22523 "Bromodomain adjacent to zinc finger domain, 2B" Baz2b
1441557_at 1.17950 SET domain containing (lysine methyltransferase) 8 Setd8
1441558_at 1.86988 RIKEN cDNA D230044B12 gene D230044B12Rik
1441559_at 0.98662 similar to CG11212-PA LOC627626
1441560_at 0.95545 "catenin (cadherin associated protein), alpha 3" Ctnna3
1441561_at 1.29611 F-box and leucine-rich repeat protein 3 Fbxl3
1441562_at 1.05095 RIKEN cDNA A230048O21 gene A230048O21Rik
1441563_at 0.95905 RIKEN cDNA 2410017P07 gene 2410017P07Rik
1441564_at 1.01760 Transcribed locus ---
1441565_at 0.98148 --- ---
1441566_at 0.84578 Transcribed locus ---
1441567_at 1.71986 Myosin IXa Myo9a
1441568_at 1.09633 cDNA sequence BC010311 BC010311
1441569_at 1.09511 RIKEN cDNA E330009P21 gene E330009P21Rik
1441570_at 1.03327 RIKEN cDNA 2810021G02 gene 2810021G02Rik
1441571_at 1.11586 RIKEN cDNA 2410089E03 gene 2410089E03Rik
1441572_at 1.03069 RIKEN cDNA C030036D22 gene C030036D22Rik
1441573_at 0.98884 Sex comb on midleg homolog 1 Scmh1
1441574_at 1.02180 "Phosphatidylinositol transfer protein, cytoplasmic 1" Pitpnc1
1441575_at 0.94510 --- ---
1441576_at 1.18350 RIKEN cDNA 2410002O22 gene 2410002O22Rik
1441577_at 2.08447 RIKEN cDNA C530014P21 gene C530014P21Rik
1441578_at 1.04269 --- ---
1441579_at 0.98346 doublesex and mab-3 related transcription factor like family A1 Dmrta1
1441580_at 1.00221 Sphingosine-1-phosphate phosphatase 2 Sgpp2
1441581_at 1.08223 ankyrin repeat and SOCS box-containing protein 10 Asb10
1441582_at 1.04863 --- ---
1441583_at 1.01170 --- ---
1441584_at 0.91373 Friend leukemia integration 1 Fli1
1441585_at 0.97607 Spastic paraplegia 7 homolog (human) Spg7
1441586_at 0.99086 Target of myb1-like 2 (chicken) Tom1l2
1441587_at 0.95669 phospholipase C-like 4 Plcl4
1441588_at 1.28126 KCNQ1 overlapping transcript 1 Kcnq1ot1
1441589_at 1.02303 Src family associated phosphoprotein 1 Scap1
1441590_at 1.07806 "potassium inwardly-rectifying channel, subfamily J, member 5" Kcnj5
1441591_at 0.97088 --- ---
1441592_at 1.33294 Jumonji domain containing 1C Jmjd1c
1441593_at 1.09436 Phosphatase and tensin homolog Pten

1441594_at 1.15514 "AF4/FMR2 family, member 1" Aff1
1441595_at 1.00495 "Excision repair cross-complementing rodent repair deficiency, complementation group 6" Ercc6
1441596_at 0.92573 RIKEN cDNA 5730601F06 gene 5730601F06Rik
1441597_at 1.31132 Ankyrin repeat and KH domain containing 1 Ankhd1
1441598_at 0.92731 transmembrane protein with EGF-like and two follistatin-like domains 2 Tmeff2
1441599_at 0.87295 RIKEN cDNA 2900075B16 gene 2900075B16Rik
1441600_at 0.93780 hypothetical protein C920021A13 C920021A13
1441601_at 1.10789 "Origin recognition complex, subunit 5-like (*S. cerevisiae*)" Orc5l
1441602_at 1.32085 RIKEN cDNA 1110005A03 gene 1110005A03Rik
1441603_at 1.00805 somatostatin receptor 3 Sstr3
1441604_at 1.70830 Esterase D/formylglutathione hydrolase Esd
1441605_at 1.08781 --- ---
1441606_at 1.00307 Neuregulin 3 Nrg3
1441607_at 1.23835 expressed sequence AU015603 AU015603
1441608_at 1.10485 "Calcium channel, voltage-dependent, alpha2/delta subunit 1" Cacna2d1
1441609_at 1.07631 Transcribed locus ---
1441610_at 0.89342 expressed sequence N28178 N28178
1441611_at 1.21299 "zinc finger, DHHC domain containing 6 /// similar to Probable palmitoyltransferase ZDHHC6 (Zinc finger DHHC domain-containing protein 6) (DHHC-6) (H4 homolog)" Zdhhc6 /// LOC669170
1441612_at 1.57039 Zinc finger protein 148 Zfp148
1441613_at 1.00127 RIKEN cDNA 1110033F14 gene 1110033F14Rik
1441614_at 1.00587 --- ---
1441615_at 1.38036 "core-binding factor, runt domain, alpha subunit 2, translocated to, 2 homolog (human)" Cbfa2t2h
1441616_at 0.95589 --- ---
1441617_at 2.07524 "positive cofactor 2, multiprotein complex, glutamine/Q-rich-associated protein" Pcqap
1441618_at 1.05380 Rho GTPase activating protein 29 Arhgap29
1441619_at 0.91639 zinc finger protein 14 Zfp14
1441620_at 1.24933 Growth factor receptor bound protein 10 Grb10
1441621_at 1.03358 Transcribed locus ---
1441622_at 0.86886 hypothetical gene supported by AK078896 LOC433023
1441623_at 1.03642 --- ---
1441624_at 1.32186 sorbin and SH3 domain containing 2 Sorbs2
1441625_at 0.91776 RIMS binding protein 2 /// similar to RIM binding protein 2 (RIM-BP2) Rimpb2 /// LOC637661
1441626_at 0.99338 --- ---
1441627_at 1.05299 Early B-cell factor 1 Ebf1
1441628_at 1.07524 Diaphanous homolog 3 (*Drosophila*) Diap3
1441629_at 1.97684 Glutamate receptor interacting protein 1 Grip1
1441630_at 1.48050 E1A binding protein p400 Ep400

1441631_at 1.26289 DEAD (Asp-Glu-Ala-Asp) box polypeptide 24 Ddx24
1441632_at 1.37054 RIKEN cDNA C130079B09 gene C130079B09Rik
1441633_at 0.95835 Nudix (nucleoside diphosphate linked moiety X)-type motif 9 Nudt9
1441634_at 1.15510 netrin G1 Ntnng1
1441635_at 1.20931 Olfactomedin-like 2A Olfml2a
1441636_at 0.98624 Transcribed locus ---
1441637_at 2.28940 RIKEN cDNA F730047E07 gene F730047E07Rik
1441638_at 0.98402 RIKEN cDNA A630014C17 gene A630014C17Rik
1441639_at 0.93446 "zinc finger, CCHC domain containing 8" Zcchc8
1441640_at 1.01159 Suppressor of cytokine signaling 5 Socs5
1441641_at 0.98427 RIKEN cDNA D630032N06 gene D630032N06Rik
1441642_at 1.29421 Like-glycosyltransferase Large
1441643_at 0.90726 membrane-associated ring finger (C3HC4) 3 3-Mar
1441644_at 1.09530 "Protein tyrosine phosphatase, receptor type, K" Ptpkr
1441645_s_at 0.95116 importin 9 Ipo9
1441646_at 0.99802 RIKEN cDNA 1110059P08 gene 1110059P08Rik
1441647_at 0.89465 Fyn proto-oncogene Fyn
1441648_at 1.07310 C1q and tumor necrosis factor related protein 4 C1qtnf4
1441649_at 0.98103 Sestrin 1 Sesn1
1441650_at 1.09568 Rho GTPase activating protein 15 Arhgap15
1441651_at 0.90574 RIKEN cDNA 4930414N06 gene 4930414N06Rik
1441652_at 1.06717 "bone morphogenic protein receptor, type II (serine/threonine kinase)" Bmpr2
1441653_at 1.38422 RIKEN cDNA D030022P06 gene D030022P06Rik
1441654_at 1.07574 "Guanine nucleotide binding protein, alpha stimulating, olfactory type" Gnal
1441655_at 0.87575 "gene model 884, (NCBI)" Gm884
1441656_at 1.96017 RIKEN cDNA B930068K11 gene B930068K11Rik
1441657_at 0.98191 BTB and CNC homology 2 Bach2
1441658_at 0.89388 Transcribed locus ---
1441659_at 0.98148 "D4, zinc and double PHD fingers, family 3" Dpf3
1441660_at 1.26051 ARP2 actin-related protein 2 homolog (yeast) Actr2
1441661_at 0.98865 --- ---
1441662_at 1.13867 "Cytochrome P450, family 4, subfamily x, polypeptide 1" Cyp4x1
1441663_at 1.29329 Prickle like 1 (Drosophila) Prickle1
1441664_at 1.06736 hypothetical gene supported by AK081809 LOC433351
1441665_at 1.02723 "isoleucine-tRNA synthetase 2, mitochondrial" Iars2
1441666_at 0.93872 "Visual cortex cDNA, RIKEN full-length enriched library, clone:K530010N24
product:unclassifiable, full insert sequence" ---
1441667_s_at 1.04056 SET and MYND domain containing 1 Smyd1
1441668_at 1.13028 RIKEN cDNA E130107B13 gene E130107B13Rik
1441669_at 1.16086 "centaurin, beta 2" Centb2
1441670_at 1.07783 T-cell lymphoma invasion and metastasis 2 Tiam2
1441671_at 1.16588 Glucocorticoid induced transcript 1 Glcci1
1441672_at 1.04530 "DNA segment, Chr 3, ERATO Doi 547, expressed" D3Ert547e

1441673_at	1.19049	plasma membrane proteolipid	Pllp
1441674_at	1.04529	---	---
1441675_at	0.99347	---	---
1441676_at	0.76214	zinc finger protein 406	Zfp406
1441677_at	1.41615	SMC4 structural maintenance of chromosomes 4-like 1 (yeast)	Smc4l1
1441678_at	1.04319	---	---
1441679_at	1.03206	"Calcium channel, voltage-dependent, L type, alpha 1C subunit"	Cacna1c
1441680_at	0.91794	"RNA binding motif, single stranded interacting protein"	Rbms3
1441681_at	1.03778	"exportin, tRNA (nuclear export receptor for tRNAs)"	Xpot
1441682_s_at	1.15280	"exportin, tRNA (nuclear export receptor for tRNAs)"	Xpot
1441683_at	1.76037	RIKEN cDNA 1110033M05 gene	1110033M05Rik
1441684_at	1.19851	tetratricopeptide repeat domain 3	Ttc3
1441685_at	0.85477	proline synthetase co-transcribed	Prosc
1441686_at	0.89059	---	---
1441687_at	0.91095	---	---
1441688_at	1.62272	RIKEN cDNA 6430537I21 gene	6430537I21Rik
1441689_at	1.24300	nucleoporin 153	Nup153
1441690_at	0.97842	cadherin 8	Cdh8
1441691_at	0.94041	Transcribed locus	---
1441692_at	1.00117	PFTAIRE protein kinase 1	Pftk1
1441693_at	0.64497	"ADAM metalloproteinase with thrombospondin type 1 motif, 3"	Adamts3
1441694_at	0.96659	Transcribed locus	---
1441695_at	1.02973	Similar to Potassium voltage-gated channel subfamily G member 2 (Voltage-gated potassium channel subunit Kv6.2) (Cardiac potassium channel subunit)	LOC240444
1441696_at	1.07922	SEC22 vesicle trafficking protein-like A (S. cerevisiae)	Sec22a
1441697_at	0.94807	Mastermind like 1 (Drosophila)	Maml1
1441698_at	1.24001	RIKEN cDNA 2410025L10 gene	2410025L10Rik
1441699_at	0.94230	"Atpase, class I, type 8B, member 2"	Atp8b2
1441700_at	0.86303	"solute carrier family 4 (anion exchanger), member 8"	Slc4a8
1441701_at	0.89147	Zinc finger protein 148	Zfp148
1441702_at	1.12940	"Regulatory factor X, 3 (influences HLA class II expression)"	Rfx3
1441703_at	1.07711	RIKEN cDNA 2210018M03 gene	2210018M03Rik
1441704_at	0.97388	"Pleckstrin homology domain containing, family A member 5"	Plekha5
1441705_at	0.99978	expressed sequence AU015680	AU015680
1441706_at	1.12556	Down syndrome cell adhesion molecule-like 1	Dscam1l
1441707_at	1.58575	---	---
1441708_at	0.96612	sperm associated antigen 16	Spag16
1441709_at	1.29714	"solute carrier family 11 (proton-coupled divalent metal ion transporters), member 2"	Slc11a2
1441710_at	0.93750	Transcribed locus	---
1441711_at	1.01009	unc-45 homolog B (C. elegans)	Unc45b
1441712_at	0.91434	Trf (TATA binding protein-related factor)-proximal protein homolog (Drosophila)	Trfp

1441713_at 0.94645 anaphase promoting complex subunit 7 Anapc7
 1441714_at 0.98236 Transcribed locus ---
 1441715_at 0.96306 RIKEN cDNA B130055D15 gene B130055D15Rik
 1441716_at 0.99691 "12 days embryo eyeball cDNA, RIKEN full-length enriched library, clone:
 D230024G22 product:unclassifiable, full insert sequence" ---
 1441717_at 1.02907 "DNA segment, Chr 9, ERATO Doi 596, expressed" D9Ertd596e
 1441718_at 1.50587 "E3 ubiquitin protein ligase, HECT domain containing, 1" Edd1
 1441719_at 0.96178 "Transcribed locus, weakly similar to XP_578477.1 PREDICTED: similar to
 hypothetical protein DKFZp434P0316 [Rattus norvegicus]" ---
 1441720_at 1.07358 RIKEN cDNA 4921530L18 gene 4921530L18Rik
 1441721_at 0.95060 --- ---
 1441722_at 1.50391 Apoptosis antagonizing transcription factor Aatf
 1441723_at 0.99834 dual specificity phosphatase 22 Dusp22
 1441724_at 1.18826 --- ---
 1441725_at 0.94061 RIKEN cDNA 6530402F18 gene 6530402F18Rik
 1441726_at 1.06353 CLIP associating protein 2 Clasp2
 1441727_s_at 1.16379 zinc finger protein 467 Zfp467
 1441728_at 0.92797 "sodium channel, voltage-gated, type I, alpha" Scn1a
 1441729_at 0.93894 Transcribed locus ---
 1441730_at 0.94836 "Protein phosphatase 1, regulatory (inhibitor) subunit 12B" Ppp1r12b
 1441731_at 0.95279 Transcribed locus ---
 1441732_at 0.90810 Teashirt zinc finger family member 1 Tshz1
 1441733_s_at 0.95421 nucleoporin 153 Nup153
 1441734_at 0.93912 Calcium/calmodulin-dependent protein kinase II alpha Camk2a
 1441735_at 0.97022 --- ---
 1441736_at 1.05162 Integrin alpha FG-GAP repeat containing 1 Itfg1
 1441737_s_at 1.11219 Ras association (RalGDS/AF-6) domain family 1 Rassf1
 1441738_at 0.84768 --- ---
 1441739_at 0.92317 Early B-cell factor 4 Ebf4
 1441740_at 0.92169 --- ---
 1441741_at 1.02937 RIKEN cDNA 1700049E17 gene /// RIKEN cDNA 4930503E14 gene ///
 RIKEN cDNA 1700024B05 gene /// similar to RIKEN cDNA 4930503E14 /// hypothetical protein
 LOC624112 /// similar to Spetex-2C protein /// similar to Spetex-2C protein /// similar to Spetex-2C
 protein 1700049E17Rik /// 4930503E14Rik /// 1700024B05Rik /// LOC434459 /// LOC624112 ///
 LOC638262 /// LOC666214 /// LOC666264
 1441742_at 0.96425 "potassium voltage-gated channel, subfamily H (eag-related), member 5" Kcnh5
 1441743_at 1.10530 paired box gene 3 Pax3
 1441744_at 0.94126 "Calcium channel, voltage-dependent, L type, alpha 1C subunit" Cacna1c
 1441745_at 1.01066 Fetal Alzheimer antigen Falz
 1441746_at 0.93943 --- ---
 1441747_at 0.89825 "Adult male spinal cord cDNA, RIKEN full-length enriched library, clone:
 A330024P14 product:unclassifiable, full insert sequence" ---
 1441748_at 0.96313 "Zinc finger, ZZ domain containing 3" Zzz3

1441749_at 1.47763 RIKEN cDNA 4930447F24 gene 4930447F24Rik
1441750_x_at 1.23394 RIKEN cDNA 4930447F24 gene 4930447F24Rik
1441751_at 0.85372 expressed sequence AW493563 AW493563
1441752_at 0.99313 ADP-ribosyltransferase 3 Art3
1441753_at 1.85321 RIKEN cDNA 4921505C17 gene 4921505C17Rik
1441754_at 0.90622 --- ---
1441755_at 0.81922 mitogen-activated protein kinase 15 Mapk15
1441756_at 1.34285 "Transcription factor 7-like 2, T-cell specific, HMG-box" Tcf7l2
1441757_at 1.38869 RIKEN cDNA 1190002F15 gene 1190002F15Rik
1441758_at 0.95381 "ras homolog gene family, member T1" Rhot1
1441759_at 0.89749 CD82 antigen Cd82
1441760_at 1.09147 similar to 40S ribosomal protein S25 /// similar to 40S ribosomal protein S25 ///
hypothetical protein LOC669102 /// similar to 40S ribosomal protein S25 LOC243302 /// LOC664840 ///
LOC669102 /// LOC676458
1441761_at 0.94525 Src-like adaptor Sla
1441762_at 0.96067 B-cell leukemia/lymphoma 11B Bcl11b
1441763_at 1.44044 "zinc finger, DHHC domain containing 6" Zdhhc6
1441764_at 0.99780 PR domain containing 10 Prdm10
1441765_at 1.19626 "PREDICTED: Mus musculus similar to Heterogeneous nuclear
ribonucleoprotein A1 (Helix-destabilizing protein) (Single-strand binding protein) (hnRNP core protein
A1) (HDP) (LOC628392), mRNA" ---
1441766_at 1.19708 Transcribed locus ---
1441767_at 0.98311 zinc finger protein 142 Zfp142
1441768_at 1.24023 RIKEN cDNA 9430051O21 gene 9430051O21Rik
1441769_at 1.76161 ADP-ribosylation factor-like 15 Arl15
1441770_at 0.99334 phosphoribosyl pyrophosphate amidotransferase Ppat
1441771_at 1.80642 Alpha thalassemia/mental retardation syndrome X-linked homolog (human)
Atrx
1441772_at 1.01740 "Purinergic receptor P2X, ligand-gated ion channel 4" P2rx4
1441773_at 0.92384 RIKEN cDNA 1700061N14 gene 1700061N14Rik
1441774_at 0.93055 "Rab38, member of RAS oncogene family" Rab38
1441775_at 1.15101 WD repeat domain 7 Wdr7
1441776_at 0.86441 tetraspanin 11 Tspan11
1441777_at 0.88074 empty spiracles homolog 1 (Drosophila) Emx1
1441778_at 1.05710 adenylate cyclase activating polypeptide 1 Adcyap1
1441779_at 1.54620 RIKEN cDNA 9530006C21 gene 9530006C21Rik
1441780_at 1.16200 Transcribed locus ---
1441781_at 0.86426 expressed sequence AI317158 AI317158
1441782_at 1.01863 AP2 associated kinase 1 Aak1
1441783_at 0.91286 RIKEN cDNA 2410022M11 gene 2410022M11Rik
1441784_at 1.07742 "Adult male hypothalamus cDNA, RIKEN full-length enriched library, clone:
A230076B10 product:unclassifiable, full insert sequence" ---
1441785_at 1.04093 "gene model 879, (NCBI)" Gm879

1441786_at 1.07133 "PREDICTED: Mus musculus similar to HYPOTHETICAL PROTEIN ORF-1137 (LOC622016), mRNA" ---

1441787_at 1.10471 Zinc finger CCCH-type containing 12C Zc3h12c

1441788_s_at 1.03393 cDNA sequence BC068171 BC068171

1441789_at 1.13432 "Rosbin, round spermatid basic protein 1" Rsb1

1441790_at 0.94886 Transcribed locus ---

1441791_at 1.01904 "Opioid receptor, mu 1" Oprm1

1441792_at 1.09384 RIKEN cDNA A630033E08 gene A630033E08Rik

1441793_at 1.00286 ring finger protein 39 Rnf39

1441794_at 1.00568 RIKEN cDNA 4933437K13 gene 4933437K13Rik

1441795_at 1.16927 "small nuclear RNA activating complex, polypeptide 4" Snapc4

1441796_at 1.07657 RIKEN cDNA 2010309E21 gene 2010309E21Rik

1441797_at 1.00310 --- ---

1441798_at 1.08321 --- ---

1441799_at 1.81105 RIKEN cDNA 6030422H21 gene 6030422H21Rik

1441800_at 1.06660 --- ---

1441801_at 0.83750 potassium channel tetramerisation domain containing 4 Kctd4

1441802_at 1.21388 "sprouty-related, EVH1 domain containing 2" Spred2

1441803_at 1.04922 RIKEN cDNA A130090K04 gene A130090K04Rik

1441804_at 0.86681 --- ---

1441805_at 0.98931 "Adult male hypothalamus cDNA, RIKEN full-length enriched library, clone: A230056E14 product:hypothetical protein, full insert sequence" ---

1441806_at 0.89174 Chromodomain helicase DNA binding protein 1-like Chd11

1441807_s_at 0.88587 --- ---

1441808_at 0.93003 RIKEN cDNA 2410002I01 gene 2410002I01Rik

1441809_at 0.95299 RIKEN cDNA A930009A15 gene A930009A15Rik

1441810_at 0.94064 Transcribed locus ---

1441811_x_at 1.43933 RIKEN cDNA 0610011I04 gene 0610011I04Rik

1441812_at 0.96862 "cytochrome P450, family 2, subfamily u, polypeptide 1" Cyp2u1

1441813_at 0.81941 hypothetical protein LOC670358 LOC670358

1441814_s_at 1.01842 RPA interacting protein Rpain

1441815_at 1.05359 expressed sequence AI851453 AI851453

1441816_at 1.20404 RIKEN cDNA 2900056M20 gene 2900056M20Rik

1441817_at 1.13406 RIKEN cDNA 1700099I09 gene 1700099I09Rik

1441818_at 0.98882 Transcribed locus ---

1441819_x_at 1.00475 translocase of inner mitochondrial membrane 50 homolog (yeast) Timm50

1441820_at 1.07454 cyclin-dependent kinase-like 2 (CDC2-related kinase) Cdkl2

1441821_at 0.84006 --- ---

1441822_at 0.75813 RIKEN cDNA 1700008F19 gene 1700008F19Rik

1441823_at 2.20712 retinoic acid induced 17 Rai17

1441824_at 1.03241 membrane bound O-acyltransferase domain containing 1 Mboat1

1441825_x_at 0.91505 Immunoglobulin mu binding protein 2 Ighmbp2

1441826_x_at 0.89799 Ubiquitin domain containing 1 Ubtd1

1441827_x_at 1.00104 "heat shock protein 90kDa alpha (cytosolic), class A member 1 /// similar to heat shock protein 1, alpha" Hsp90aa1 /// LOC670148
1441828_at 1.00360 StAR-related lipid transfer (START) domain containing 5 Stard5
1441829_s_at 1.37902 A kinase (PRKA) anchor protein 10 Akap10
1441830_x_at 1.17987 --- ---
1441831_x_at 1.01719 Chloride channel 5 Clcn5
1441832_at 0.91811 histone deacetylase 10 Hdac10
1441833_at 0.91979 N-glycanase 1 Ngly1
1441834_x_at 0.96620 Cytotoxic granule-associated RNA binding protein 1 Tia1
1441835_x_at 0.89408 myotubularin related protein 11 Mtmr11
1441836_x_at 1.20549 RIKEN cDNA 1700006H03 gene 1700006H03Rik
1441837_at 1.00956 "cholinergic receptor, nicotinic, beta polypeptide 2 (neuronal)" Chrnb2
1441838_at 0.93845 "excision repair cross-complementing rodent repair deficiency, complementation group 1" Ercc1
1441839_s_at 0.97081 cDNA sequence BC024814 BC024814
1441840_x_at 1.21302 Strawberry notch homolog (Drosophila) Stno
1441841_at 1.08679 "a disintegrin-like and metalloproteinase (reprolysin type) with thrombospondin type 1 motif, 16" Adamts16
1441842_s_at 0.87249 zinc finger protein 707 Zfp707
1441843_s_at 2.29152 RIKEN cDNA 5230400M03 gene 5230400M03Rik
1441844_at 1.02624 Epidermal growth factor receptor pathway substrate 15-like 1 Eps1511
1441845_at 0.93173 calcyphosphine 2 Caps2
1441846_x_at 1.05027 Transmembrane emp24 domain containing 3 Tmed3
1441847_at 1.12853 Microtubule associated serine/threonine kinase 2 Mast2
1441848_at 1.04504 Phosphomannomutase 2 Pmm2
1441849_at 1.02323 --- ---
1441850_x_at 0.76547 transcobalamin 2 Tcn2
1441851_x_at 0.80127 "lymphocyte antigen 6 complex, locus G6C" Ly6g6c
1441852_x_at 1.05506 autophagy-related 16-like 1 (yeast) Atg1611
1441853_at 1.02345 Dipeptidylpeptidase 10 Dpp10
1441854_at 1.10672 "Polymerase (DNA directed), epsilon" Pole
1441855_x_at 2.66323 chemokine (C-X-C motif) ligand 1 Cxcl1
1441856_x_at 0.84471 trypsin domain containing 1 Tysnd1
1441857_x_at 1.02040 SEH1-like (S. cerevisiae) Seh11
1441858_at 1.91695 Uridine-cytidine kinase 2 Uck2
1441859_x_at 1.02892 Coiled-coil domain containing 64 Ccdc64
1441860_x_at 0.97648 Insulin degrading enzyme Ide
1441861_at 0.93416 RIKEN cDNA 1700018G05 gene 1700018G05Rik
1441862_at 0.91142 protoporphyrinogen oxidase Ppox
1441863_x_at 0.85502 --- ---
1441864_x_at 0.90399 centromere protein A Cenpa
1441865_at 0.92197 --- ---
1441866_s_at 0.94111 phosphatidylserine synthase 1 Ptdss1

1441867_x_at	0.83864	RIKEN cDNA 4930534B04 gene	4930534B04Rik
1441868_x_at	1.02271	DC-STAMP domain containing 2	Dcst2
1441869_x_at	1.14915	Autism susceptibility candidate 2	Auts2
1441870_s_at	0.90403	polycystic kidney disease 2	Pkd2
1441871_at	1.09709	RIKEN cDNA 1810044D09 gene	1810044D09Rik
1441872_x_at	0.96905	PHD finger protein 17	Phf17
1441873_at	1.04223	prolactin-like protein E	Prlpe
1441874_x_at	0.92521	prolactin-like protein E	Prlpe
1441875_at	0.89243	RIKEN cDNA 1700063J08 gene	1700063J08Rik
1441876_x_at	0.96595	Zinc finger protein 93	Zfp93
1441877_x_at	0.86125	tumor suppressing subtransferable candidate 1	Tssc1
1441878_s_at	0.82643	RIKEN cDNA 1810049H13 gene	1810049H13Rik
1441879_x_at	0.93296	"makorin, ring finger protein, 1"	Mkrn1
1441880_x_at	0.97977	hypothetical protein MGC30332	MGC30332
1441881_x_at	0.98838	RIKEN cDNA 3110032G18 gene	3110032G18Rik
1441882_at	0.92563	---	---
1441883_at	0.80902	RIKEN cDNA 0610010O12 gene	0610010O12Rik
1441884_x_at	1.06326	chemokine (C-X3-C motif) ligand 1	Cx3cl1
1441885_s_at	1.06295	"Adult male urinary bladder cDNA, RIKEN full-length enriched library, clone:9530098H01 product:unclassifiable, full insert sequence" ---	
1441886_at	0.95855	RIKEN cDNA 4930532D21 gene	4930532D21Rik
1441887_x_at	1.02068	hypothetical protein LOC622976	LOC622976
1441888_x_at	1.10411	---	---
1441889_x_at	0.98246	Rac GTPase-activating protein 1	Racgap1
1441890_x_at	1.04465	transmembrane protein with EGF-like and two follistatin-like domains 1	Tmeff1
1441891_x_at	0.85090	"ELOVL family member 7, elongation of long chain fatty acids (yeast)" Elov17	
1441892_x_at	1.16987	RIKEN cDNA 4933440H19 gene	4933440H19Rik
1441893_at	0.93451	testis expressed gene 24	Tex24
1441894_s_at	1.06247	GRP1 (general receptor for phosphoinositides 1)-associated scaffold protein Grasp	
1441895_x_at	0.95526	Transcribed locus	---
1441896_x_at	1.32073	Oligonucleotide/oligosaccharide-binding fold containing 1	Obfc1
1441897_at	0.93852	RIKEN cDNA B230120H23 gene	B230120H23Rik
1441898_at	0.93252	RIKEN cDNA E530011F12 gene	E530011F12Rik
1441899_x_at	0.93236	Brevican	Bcan
1441900_x_at	1.04615	Hspb associated protein 1	Hspbap1
1441901_x_at	0.95792	fumarylacetoacetate hydrolase domain containing 2A	Fahd2a
1441902_x_at	1.02937	---	---
1441903_x_at	0.98232	"meteorin, glial cell differentiation regulator-like" Metrnl	
1441904_x_at	0.78303	RIKEN cDNA 9130005N14 gene	9130005N14Rik
1441905_x_at	0.87555	Small nuclear ribonucleoprotein N	Snrpn

1441906_x_at	1.11211	Synapse associated protein 1	Syap1	
1441907_s_at	0.97340	CD93 antigen	Cd93	
1441908_x_at	1.09889	CD93 antigen	Cd93	
1441909_s_at	0.90960	RIKEN cDNA 9530066K23 gene	9530066K23Rik	
1441910_x_at	0.87934	cyclin E1	Ccne1	
1441911_x_at	0.98176	phosphoribosylglycinamide formyltransferase	Gart	
1441912_x_at	0.91762	complement component 2 (within H-2S)	C2	
1441913_at	1.06674	leucine zipper protein 2	Luzp2	
1441914_x_at	0.99733	Fibroblast growth factor 3	Fgf3	
1441915_s_at	1.05605	RIKEN cDNA 2310076L09 gene	2310076L09Rik	
1441916_s_at	0.94365	outer dense fiber of sperm tails 3	Odf3	
1441917_s_at	0.95653	transmembrane protein 40	Tmem40	
1441918_x_at	0.95898	similar to NK13 RP23-262J21.5		
1441919_x_at	0.88110	carnitine acetyltransferase	Crat	
1441920_x_at	1.01551	HpaII tiny fragments locus 9c	Htf9c	
1441921_x_at	0.87349	"estrogen related receptor, beta"	Esrrb	
1441922_x_at	1.04980	vacuolar protein sorting 11 (yeast)	Vps11	
1441923_s_at	1.00866	endothelin 3	Edn3	
1441924_x_at	1.27092	endothelin 3	Edn3	
1441925_at	1.05354	"solute carrier family 15, member 4"	Slc15a4	
1441926_x_at	0.97238	transmembrane inner ear	Tmie	
1441927_at	0.79748	Synaptotagmin VII	Syt7	
1441928_x_at	1.02217	elongation factor RNA polymerase II	Ell	
1441929_at	0.90201	Midline 1	Mid1	
1441930_x_at	0.98342	Vesicle amine transport protein 1 homolog (T californica)	Vat1	
1441931_x_at	0.85224	glutathione synthetase	Gss	
1441932_at	1.06220	"Protein tyrosine phosphatase, receptor type, D"	Ptprd	
1441933_x_at	1.03639	"phosphatidylinositol glycan anchor biosynthesis, class Z"	Pigz	
1441934_at	0.98326	---	---	
1441935_at	1.23751	"ankyrin repeat, family A (RFXANK-like), 2"	Ankra2	
1441936_x_at	0.92401	RIKEN cDNA 4930447M23 gene	4930447M23Rik	
1441937_s_at	0.88525	PTEN induced putative kinase 1	Pink1	
1441938_x_at	1.06204	Cdk5 and Abl enzyme substrate 1 /// similar to Cdk5 and Abl enzyme substrate 1	Cables1 /// LOC635753	
1441939_x_at	1.04059	RIKEN cDNA 2410003I16 gene	2410003I16Rik	
1441940_x_at	1.00008	kelch domain containing 8A	Klhdc8a	
1441941_x_at	0.96191	"serine (or cysteine) peptidase inhibitor, clade B, member 5"	Serpib5	
1441942_x_at	1.08445	snurportin 1	Snupn	
1441943_x_at	0.97833	Cytokine induced apoptosis inhibitor 1	Ciapin1	
1441944_s_at	0.96184	G protein-coupled receptor 135	Gpr135	
1441945_s_at	1.06019	abhydrolase domain containing 14A	Abhd14a	
1441946_at	0.96070	inter-alpha (globulin) inhibitor H5	Itih5	
1441947_x_at	1.12408	cDNA sequence BC033915	BC033915	

1441948_x_at	0.93063	"zinc finger, AN1-type domain 3"	Zfand3
1441949_x_at	1.06536	"solute carrier family 39 (metal ion transporter), member 6"	Slc39a6
1441950_s_at	1.12166	RIKEN cDNA 2010109A12 gene	2010109A12Rik
1441951_x_at	0.88065	RIKEN cDNA 2010109A12 gene	2010109A12Rik
1441952_x_at	0.69024	Ly6/neurotoxin 1	Lynx1
1441953_at	1.11801	Storkhead box 2	Stox2
1441954_s_at	0.96484	"H1 histone family, member N, testis-specific"	H1fnt
1441955_s_at	1.00042	polyadenylate binding protein-interacting protein 1 /// similar to poly(A) binding protein interacting protein 1	Paip1 /// LOC676674
1441956_s_at	2.36598	---	---
1441957_x_at	0.95130	RIKEN cDNA 2410076I21 gene	2410076I21Rik
1441958_s_at	0.68858	advanced glycosylation end product-specific receptor	Ager
1441959_s_at	0.90932	RIKEN cDNA 1200003C05 gene	1200003C05Rik
1441960_x_at	0.86827	RIKEN cDNA 5730494M16 gene	5730494M16Rik
1441961_at	0.91967	microtubule-associated protein 9	Mtap9
1441962_at	0.99171	arachidonate 5-lipoxygenase	Alox5
1441963_at	0.95533	ProSAPiP1 protein	Prosapip1
1441964_at	0.98214	RIKEN cDNA 1110003F05 gene	1110003F05Rik
1441965_at	0.91820	coiled-coil domain containing 38	Ccdc38
1441966_at	0.90172	"transient receptor potential cation channel, subfamily M, member 3"	Trpm3
1441967_at	0.83798	Parkinson disease 7 domain containing 1	Pddc1
1441968_at	0.89010	tetraspanin 9	Tspan9
1441969_at	0.92070	tripartite motif-containing 36	Trim36
1441970_at	1.15607	RIKEN cDNA E430010N07 gene	E430010N07Rik
1441971_at	0.85439	Transcribed locus	---
1441972_at	0.64591	RIKEN cDNA 6230424C14 gene	6230424C14Rik
1441973_at	0.88087	zinc finger protein 295	Zfp295
1441974_at	0.97516	Transcribed locus	---
1441975_at	0.96029	"acid phosphatase, prostate"	Acpp
1441976_at	1.95503	RIKEN cDNA 5430434G16 gene	5430434G16Rik
1441977_at	1.05216	RIKEN cDNA 9630023C09 gene	9630023C09Rik
1441978_at	1.17757	aquaporin 6	Aqp6
1441979_at	1.05120	cDNA sequence BC060267	BC060267
1441980_at	0.78686	RIKEN cDNA C030007I09 gene	C030007I09Rik
1441981_at	0.95847	"transmembrane protease, serine 11f"	Tmprss11f
1441982_at	0.87141	"TAF7 RNA polymerase II, TATA box binding protein (TBP)-associated factor"	Taf7
1441983_at	1.03018	"0 day neonate cerebellum cDNA, RIKEN full-length enriched library, clone: C230096I08 product:hypothetical protein, full insert sequence"	---
1441984_at	1.14788	claspin homolog (Xenopus laevis)	Clspn
1441985_at	1.02224	RIKEN cDNA 4933416C03 gene	4933416C03Rik
1441986_at	0.82603	"zinc finger, CCHC domain containing 6"	Zcchc6
1441987_at	1.15394	methyl-CpG binding domain protein 5	Mbd5

1441988_at	0.99110	protein phosphatase 1K (PP2C domain containing)	Ppm1k
1441989_at	0.91476	"BCL2/adenovirus E1B interacting protein 1, NIP2"	Bnip2
1441990_at	1.13203	meiotic recombination 11 homolog A (<i>S. cerevisiae</i>)	Mre11a
1441991_at	0.91794	cDNA sequence BC039632	BC039632
1441992_at	1.41621	"RAB14, member RAS oncogene family"	Rab14
1441993_at	0.92045	"adaptor-related protein complex 3, sigma 2 subunit"	Ap3s2
1441994_at	0.79449	protocadherin beta 16	Pcdhb16
1441995_at	1.11181	Neural cell adhesion molecule 1	Ncam1
1441996_at	1.09609	RIKEN cDNA 9530082P21 gene	9530082P21Rik
1441997_at	0.92130	zinc finger protein 184 (Kruppel-like)	Zfp184
1441998_at	0.83270	Transcribed locus	---
1441999_at	1.06330	Transcribed locus	---
1442000_at	1.05698	"gene model 1964, (NCBI)"	Gm1964
1442001_at	0.98730	"protein kinase, AMP-activated, beta 2 non-catalytic subunit"	Prkab2
1442002_at	0.92511	RIKEN cDNA 7030402D04 gene	7030402D04Rik
1442003_at	1.05391	"Mus musculus, clone IMAGE:4219318, mRNA"	---
1442004_at	1.14293	tripartite motif-containing 65	Trim65
1442005_at	1.15645	expressed sequence AW987390	AW987390
1442006_at	0.89010	Transcribed locus	---
1442007_at	0.94524	RIKEN cDNA 9830124H08 gene	9830124H08Rik
1442008_at	0.76502	"Adult male hypothalamus cDNA, RIKEN full-length enriched library, clone: A230090L04 product:unclassifiable, full insert sequence"	---
1442009_at	0.92619	"eukaryotic translation initiation factor 2C, 3"	Eif2c3
1442010_at	0.93196	"castor homolog 1, zinc finger (<i>Drosophila</i>) /// similar to castor homolog 1, zinc finger" Casz1 /// LOC547380	
1442011_at	1.07103	RIKEN cDNA A230065H16 gene	A230065H16Rik
1442012_at	0.96658	expressed sequence AU015791	AU015791
1442013_at	1.02353	RIKEN cDNA 4930532D21 gene	4930532D21Rik
1442014_at	1.83279	Interferon-related developmental regulator 1	Ifrd1
1442015_at	1.33978	RIKEN cDNA 2500002B13 gene	2500002B13Rik
1442016_at	1.00720	RIKEN cDNA D630004N19 gene	D630004N19Rik
1442017_at	1.02069	nitrogen fixation gene 1 (<i>S. cerevisiae</i>)	Nfs1
1442018_at	1.32105	expressed sequence AI426953	AI426953
1442019_at	0.97957	RIKEN cDNA B230343A10 gene	B230343A10Rik
1442020_at	0.93744	Olfactory receptor 1329	Olf1329
1442021_at	0.96208	"guanine nucleotide binding protein, alpha stimulating, olfactory type"	Gnal
1442022_at	0.93403	RIKEN cDNA E430004N04 gene	E430004N04Rik
1442023_at	0.95431	RIKEN cDNA A530030E21 gene	A530030E21Rik
1442024_at	1.09636	"Protein phosphatase 1, regulatory (inhibitor) subunit 3E"	Ppp1r3e
1442025_a_at	0.85226	---	---
1442026_at	0.83420	---	---
1442027_at	1.24521	neurobeachin like 1	Nbeal1
1442028_at	0.92947	expressed sequence AI593864	AI593864

1442029_at 0.98441 "Potassium voltage-gated channel, subfamily Q, member 1" Kcnq1
1442030_at 0.92497 --- ---
1442031_at 1.03187 RIKEN cDNA D130073L02 gene D130073L02Rik
1442032_at 0.89686 cDNA sequence BC030500 BC030500
1442033_at 0.87904 C1q and tumor necrosis factor related protein 9 C1qtnf9
1442034_at 0.97011 FAST kinase domains 1 Fastkd1
1442035_at 1.07034 "cholinergic receptor, nicotinic, alpha polypeptide 5" Chrna5
1442036_a_at 0.94533 hypothetical protein LOC668580 /// hypothetical protein LOC670695
LOC668580 /// LOC670695
1442037_at 0.98715 hypothetical protein LOC668580 LOC668580
1442038_at 1.22497 RNA binding motif protein 26 Rbm26
1442039_at 1.06398 thymocyte selection-associated HMG box gene Tox
1442040_at 0.92606 RIKEN cDNA 6330530A05 gene 6330530A05Rik
1442041_at 1.29583 hypothetical LOC552876 LOC552876
1442042_at 1.06011 Transcription factor 20 Tcf20
1442043_at 0.85159 lipoma HMGIC fusion partner-like protein 4 Lhfpl4
1442044_at 1.14346 Transcribed locus ---
1442045_at 0.96180 --- ---
1442046_at 0.94052 shroom family member 2 /// similar to Apical-like protein (APXL protein)
Shroom2 /// LOC670546
1442047_at 1.02401 ras responsive element binding protein 1 Rreb1
1442048_at 0.96306 ring finger protein 11 Rnf11
1442049_at 0.97406 Transferrin Trf
1442050_at 0.73845 zinc finger protein 608 /// RIKEN cDNA D430007A19 gene Zfp608 ///
D430007A19Rik
1442051_at 1.17810 "histone 2, H3c1" Hist2h3c1
1442052_at 1.54914 RIKEN cDNA C330019G07 gene C330019G07Rik
1442053_at 1.00901 Phosphorylase kinase beta Phkb
1442054_at 1.45355 "PREDICTED: Mus musculus similar to serine-threonine protein kinase pim-2
isoform 1 (LOC622647), mRNA" ---
1442055_at 0.88850 Transcribed locus ---
1442056_at 0.94668 Zinc finger protein 608 Zfp608
1442057_at 0.95908 expressed sequence AA986695 AA986695
1442058_s_at 0.89375 "proteasome (prosome, macropain) 26S subunit, ATPase 3, interacting
protein" Psmc3ip
1442059_at 1.01674 "fragile X mental retardation gene 1, autosomal homolog" Fxr1h
1442060_at 1.01296 prolyl endopeptidase-like Prepl
1442061_at 0.97585 BTB (POZ) domain containing 7 Btbd7
1442062_at 0.90485 RIKEN cDNA 7120426M23 gene 7120426M23Rik
1442063_at 1.01504 RIKEN cDNA 5930437A14 gene 5930437A14Rik
1442064_at 0.98407 expressed sequence AW556556 AW556556
1442065_at 0.94793 Rho GTPase activating protein 17 Arhgap17
1442066_at 0.90252 RIKEN cDNA C530028O21 gene C530028O21Rik

1442067_at 1.10043 Receptor tyrosine kinase-like orphan receptor 1 Ror1
1442068_at 1.04228 "Mus musculus, clone IMAGE:5363897, mRNA" ---
1442069_at 1.32594 "DNA segment, Chr 5, Wayne State University 178, expressed" D5Wsu178e
1442070_at 1.01753 myeloid/lymphoid or mixed lineage-leukemia translocation to 10 homolog
(Drosophila) Mllt10
1442071_at 1.20651 "ATP-binding cassette, sub-family E (OABP), member 1" Abce1
1442072_at 0.90189 RIKEN cDNA C230081A13 gene C230081A13Rik
1442073_at 1.00189 --- ---
1442074_at 0.96897 "C-type lectin domain family 2, member g" Clec2g
1442075_at 0.99075 expressed sequence AI314604 AI314604
1442076_at 1.00311 cordon-bleu Cobl
1442077_at 0.64330 RIKEN cDNA 2310076G05 gene 2310076G05Rik
1442078_at 1.27104 "CTD (carboxy-terminal domain, RNA polymerase II, polypeptide A) small
phosphatase like 2" Ctdspl2
1442079_at 1.15551 transmembrane protein 23 Tmem23
1442080_at 1.13769 CAMP responsive element binding protein 3-like 2 Creb3l2
1442081_at 0.88559 Expressed sequence AU017455 AU017455
1442082_at 0.97543 complement component 3a receptor 1 C3ar1
1442083_at 1.20930 RIKEN cDNA 1500011J06 gene 1500011J06Rik
1442084_at 0.81584 Prkr interacting protein 1 (IL11 inducible) Prkrip1
1442085_at 1.11149 Wolf-Hirschhorn syndrome candidate 1-like 1 (human) Whsc1l1
1442086_at 1.03926 Metastasis associated 3 Mta3
1442087_at 1.52210 "H3 histone, family 3A" H3f3a
1442088_at 1.25025 RIKEN cDNA 9230111E07 gene 9230111E07Rik
1442089_at 0.98225 --- ---
1442090_at 1.03192 RIKEN cDNA 8030463A06 gene 8030463A06Rik
1442091_at 0.89194 Transcribed locus ---
1442092_at 1.04219 "Fanconi anemia, complementation group D2" Fancd2
1442093_at 1.04750 Arginine decarboxylase Adc
1442094_at 1.24223 "inhibitor of growth family, member 5" Ing5
1442095_at 1.03052 additional sex combs like 3 (Drosophila) Asxl3
1442096_at 0.91019 cDNA sequence BC031901 BC031901
1442097_at 0.88413 "0 day neonate kidney cDNA, RIKEN full-length enriched library, clone:
D630011L08 product:hypothetical protein, full insert sequence" ---
1442098_at 0.91697 expressed sequence AU022434 AU022434
1442099_at 1.04299 ubiquitin specific peptidase 31 Usp31
1442100_at 0.98023 inositol polyphosphate-5-phosphatase F Inpp5f
1442101_at 1.06828 RIKEN cDNA A930017N06 gene A930017N06Rik
1442102_at 1.39620 "Aconitase 2, mitochondrial" Aco2
1442103_at 2.74477 Nipped-B homolog (Drosophila) Nipbl
1442104_at 0.97745 Ubiquitin specific peptidase 25 Usp25
1442105_at 0.95619 RIKEN cDNA 4831416G18 gene 4831416G18Rik
1442106_at 0.96752 "Fanconi anemia, complementation group M" Fancm

1442107_at 1.90116 "filamin, beta" Flnb
1442108_at 1.18563 Ring finger protein 25 Rnf25
1442109_at 1.76378 --- ---
1442110_at 0.90131 SET and MYND domain containing 3 Smyd3
1442111_at 1.99389 RIKEN cDNA D430033H22 gene D430033H22Rik
1442112_at 1.32615 vacuolar protein sorting 18 (yeast) Vps18
1442113_at 1.01470 RIKEN cDNA 5330417C22 gene 5330417C22Rik
1442114_at 0.77918 XPA binding protein 1 Xab1
1442115_at 0.93820 RIKEN cDNA 9430028L06 gene 9430028L06Rik
1442116_at 0.94480 G protein-coupled receptor 176 Gpr176
1442117_at 0.90400 "Adult retina cDNA, RIKEN full-length enriched library, clone:A930003C13
product:unclassifiable, full insert sequence" ---
1442118_at 0.97975 Transcribed locus ---
1442119_at 0.88666 expressed sequence AI449212 AI449212
1442120_at 0.89198 "Lung RCB-0558 LLC cDNA, RIKEN full-length enriched library, clone:
G730007D18 product:hypothetical protein, full insert sequence" ---
1442121_at 0.91881 "cyclic AMP-regulated phosphoprotein, 21" Arpp21
1442122_at 1.20772 expressed sequence AI451458 AI451458
1442123_at 1.14660 RIKEN cDNA 2010012P19 gene 2010012P19Rik
1442124_at 0.82084 expressed sequence AI450326 AI450326
1442125_at 1.24064 "KRIT1, ankyrin repeat containing" Krit1
1442126_at 1.22359 RIKEN cDNA 5830417C01 gene 5830417C01Rik
1442127_at 0.94236 Transcribed locus ---
1442128_at 1.34485 Receptor accessory protein 5 Reep5
1442129_at 0.82676 RIKEN cDNA 1810058I24 gene 1810058I24Rik
1442130_at 1.10150 Hematopoietic SH2 domain containing Hsh2d
1442131_at 0.96374 RIKEN cDNA 1700048M11 gene 1700048M11Rik
1442132_at 1.05109 CDNA sequence BC057552 BC057552
1442133_at 1.08049 "gene model 237, (NCBI)" Gm237
1442134_at 1.18972 proline rich 11 Prr11
1442135_at 0.98207 "Gene model 237, (NCBI)" Gm237
1442136_at 0.87408 ankyrin repeat and IBR domain containing 1 Ankib1
1442137_at 1.04838 RIKEN cDNA 2010000I03 gene 2010000I03Rik
1442138_at 0.98642 G protein-coupled receptor 62 Gpr62
1442139_at 1.01216 "Adult male diencephalon cDNA, RIKEN full-length enriched library,
clone:9330140C08 product:unclassifiable, full insert sequence" ---
1442140_at 1.01158 tenascin N Tnn
1442141_at 0.85053 cDNA sequence BC028454 BC028454
1442142_at 1.00370 RIKEN cDNA 2700050L05 gene 2700050L05Rik
1442143_at 1.07763 transmembrane protein 16D (eight membrane-spanning domains) Tmem16d
1442144_at 0.91365 EMI domain containing 1 Emid1
1442145_at 1.05196 ATPase type 13A3 Atp13a3
1442146_at 0.91571 Metastasis associated 3 Mta3

1442147_at 1.03006 Phosphorylase kinase alpha 2 Phka2
1442148_at 0.93786 PC4 and SFRS1 interacting protein 1 Psip1
1442149_at 0.97364 "TBC1 domain family, member 16" Tbc1d16
1442150_at 1.49748 --- ---
1442151_at 2.21932 RIKEN cDNA D230040A04 gene D230040A04Rik
1442152_at 1.05181 RIKEN cDNA 4930513N10 gene 4930513N10Rik
1442153_at 1.12156 "Ubiquitously transcribed tetratricopeptide repeat gene, Y chromosome" Uty
1442154_at 0.94430 "U2 small nuclear ribonucleoprotein auxiliary factor (U2AF) 1, related sequence 2" U2af1-rs2
1442155_at 1.47149 RIKEN cDNA 4632427E13 gene 4632427E13Rik
1442156_at 0.88196 RIKEN cDNA E030030I06 gene E030030I06Rik
1442157_at 0.84882 RIKEN cDNA A930041H05 gene A930041H05Rik
1442158_at 1.13094 microtubule associated serine/threonine kinase family member 4 Mast4
1442159_at 0.99980 "DNA segment, Chr 2, ERATO Doi 48, expressed" D2Ertd48e
1442160_at 1.06087 RIKEN cDNA 7530404M11 gene 7530404M11Rik
1442161_at 0.96912 immunoglobulin (CD79A) binding protein 1 Igbp1
1442162_at 0.97933 --- ---
1442163_at 1.51960 "HECT domain and ankyrin repeat containing, E3 ubiquitin protein ligase 1" Hace1
1442164_at 0.97241 basic leucine zipper nuclear factor 1 Blzf1
1442165_at 0.74389 armadillo repeat containing 9 Armc9
1442166_at 0.83777 copine V Cpne5
1442167_at 1.00821 cytoplasmic FMR1 interacting protein 2 Cyfip2
1442168_at 1.54424 RIKEN cDNA A230071A22 gene A230071A22Rik
1442169_at 0.86268 very low density lipoprotein receptor Vldlr
1442170_at 1.17837 "Elastase 3, pancreatic" Ela3
1442171_at 1.03044 PRP40 pre-mRNA processing factor 40 homolog A (yeast) Prpf40a
1442172_at 0.93816 Prickle like 1 (Drosophila) Prickle1
1442173_at 1.06759 PHD finger protein 3 Phf3
1442174_at 1.00141 tetraspanin 18 Tspan18
1442175_at 0.86366 RIKEN cDNA C030027H14 gene C030027H14Rik
1442176_at 1.00148 AT rich interactive domain 5B (Mrf1 like) Arid5b
1442177_at 0.98250 "solute carrier family 6 (neurotransmitter transporter, glycine), member 5 /// similar to Sodium- and chloride-dependent glycine transporter 2 (GlyT2) (GlyT-2) (Solute carrier family 6 member 5)" Slc6a5 /// LOC677366
1442178_at 1.07839 "GDP-mannose 4, 6-dehydratase" Gmds
1442179_at 1.00700 RIKEN cDNA 9430053O09 gene 9430053O09Rik
1442180_at 1.08139 "deleted in lymphocytic leukemia, 7" Dleu7
1442181_at 1.88265 Glucocorticoid receptor DNA binding factor 1 Grlf1
1442182_at 0.92494 "DnaJ (Hsp40) homolog, subfamily C, member 19" Dnajc19
1442183_at 1.16284 similar to carnitine deficiency-associated gene expressed in ventricle 3 isoform CDV3B LOC236069
1442184_at 1.14882 "Solute carrier family 24 (sodium/potassium/calcium exchanger), member

3" Slc24a3

1442185_at 1.08666 --- ---
 1442186_at 1.07897 ataxin 7 Atxn7
 1442187_at 0.96578 "bradykinin receptor, beta 2" Bdkrb2
 1442188_at 1.12657 "0 day neonate cerebellum cDNA, RIKEN full-length enriched library, clone:
 C230031C13 product:unclassifiable, full insert sequence" ---
 1442189_at 1.03470 "Pleckstrin homology, Sec7 and coiled-coil domains 1" Pscd1
 1442190_at 0.98477 --- ---
 1442191_at 0.94056 cDNA sequence AF397014 AF397014
 1442192_at 0.97498 thymidylate synthase Tyms
 1442193_at 0.88456 RIKEN cDNA A430028G04 gene A430028G04Rik
 1442194_at 1.90229 expressed sequence AU018552 AU018552
 1442195_at 0.97662 expressed sequence BB283564 BB283564
 1442196_at 0.90058 RIKEN cDNA A930006K02 gene A930006K02Rik
 1442197_at 1.20831 expressed sequence AI480624 AI480624
 1442198_at 1.63829 NIMA (never in mitosis gene a)-related expressed kinase 7 Nek7
 1442199_at 0.98894 butyrophilin-like 2 Btnl2
 1442200_at 1.06636 RIKEN cDNA 1810063B07 gene 1810063B07Rik
 1442201_at 0.94929 Transcribed locus ---
 1442202_at 0.87570 RIKEN cDNA 6430706D22 gene 6430706D22Rik
 1442203_at 0.86931 hypothetical LOC552880 LOC552880
 1442204_at 1.03468 Pannexin 2 Panx2
 1442205_at 0.99339 "Adult retina cDNA, RIKEN full-length enriched library, clone:A930031I16
 product:unclassifiable, full insert sequence" ---
 1442206_at 0.98590 MAM domain containing 1 Mamdc1
 1442207_at 0.93422 Autophagy related 16 like 2 (S. cerevisiae) Atg16l2
 1442208_at 0.88131 "Sodium channel, voltage-gated, type VIII, alpha" Scn8a
 1442209_at 0.92790 "7 days neonate cerebellum cDNA, RIKEN full-length enriched library, clone:
 A730047K09 product:unclassifiable, full insert sequence" ---
 1442210_at 0.95687 RIKEN cDNA 1700058C01 gene 1700058C01Rik
 1442211_at 0.99903 expressed sequence C78893 C78893
 1442212_at 0.94577 carboxypeptidase D Cpd
 1442213_at 1.02842 hypothetical LOC552908 LOC552908
 1442214_at 1.72942 Nuclear factor I/B Nfib
 1442215_at 1.03639 smoothed homolog (Drosophila) Smo
 1442216_at 1.31363 "ATPase, H+ transporting, lysosomal V0 subunit D1" Atp6v0d1
 1442217_at 1.23718 "DnaJ (Hsp40) homolog, subfamily B, member 6" Dnajb6
 1442218_at 0.96316 mitogen-activated protein kinase kinase kinase 9 Map3k9
 1442219_at 0.94342 "Membrane-spanning 4-domains, subfamily A, member 4B" Ms4a4b
 1442220_at 1.02404 --- ---
 1442221_at 0.97264 Secretion regulating guanine nucleotide exchange factor Sergef
 1442222_at 1.43979 interferon (alpha and beta) receptor 1 Ifnar1
 1442223_at 1.85646 Enabled homolog (Drosophila) Enah

1442224_at	1.16947	"Pleckstrin homology domain containing, family A member 6"	Plekha6
1442225_at	0.88318	Cell division cycle 2-like 6 (CDK8-like)	Cdc216
1442226_at	0.93939	hypothetical protein 6430702L12	6430702L12
1442227_at	1.05272	"Adaptor protein complex AP-2, mu1"	Ap2m1
1442228_at	0.96870	RIKEN cDNA D630036G22 gene	D630036G22Rik
1442229_at	0.91469	Paired box gene 9	Pax9
1442230_at	0.90353	---	---
1442231_at	1.13788	expressed sequence AI616248	AI616248
1442232_at	1.02752	RIKEN cDNA C530025M09 gene	C530025M09Rik
1442233_at	0.94784	Prohibitin	Phb
1442234_at	0.92302	hypothetical protein LOC667085	LOC667085
1442235_at	1.12525	Pleiomorphic adenoma gene-like 2	Plagl2
1442236_at	0.93983	"zinc finger, AN1-type domain 6"	Zfand6
1442237_at	0.86945	Sequestosome 1	Sqstm1
1442238_a_at	1.03892	kinesin family member 6	Kif6
1442239_at	1.12070	"Tumor necrosis factor, alpha-induced protein 8"	Tnfaip8
1442240_at	1.09936	"Ctr9, Paf1/RNA polymerase II complex component, homolog (S. cerevisiae)"	Ctr9
1442241_at	1.26996	Serine/arginine-rich protein specific kinase 2	Srpk2
1442242_at	0.98881	expressed sequence C79127	C79127
1442243_at	1.28540	period homolog 3 (Drosophila)	Per3
1442244_at	0.93593	InaD-like (Drosophila)	Inadl
1442245_at	0.96863	Protein arginine N-methyltransferase 4	Prmt4
1442246_at	1.06269	hypothetical protein LOC668224	/// hypothetical protein LOC669738
hypothetical protein LOC676027		LOC668224	/// LOC669738
1442247_at	1.08295	---	---
1442248_at	1.84304	Triple functional domain (PTPRF interacting)	Trio
1442249_at	0.80039	Transcribed locus	---
1442250_at	1.29255	X-ray repair complementing defective repair in Chinese hamster cells 6	Xrcc6
1442251_at	1.02251	valosin containing protein (p97)/p47 complex interacting protein 1	Vcpi1
1442252_at	1.13544	kinesin family member 13A	Kif13a
1442253_at	0.97520	---	---
1442254_at	1.14856	"Diacylglycerol kinase, delta"	Dgkd
1442255_at	1.32983	RIKEN cDNA 5033414D02 gene	5033414D02Rik
1442256_at	1.55780	"protein kinase C, delta"	Prkcd
1442257_at	0.50478	RIKEN cDNA B130021B11 gene	B130021B11Rik
1442258_at	1.20335	---	---
1442259_at	1.05714	---	---
1442260_at	0.92410	Cdc42 binding protein kinase alpha	Cdc42bpa
1442261_at	1.00070	G protein-coupled receptor 150	Gpr150
1442262_at	0.94643	---	---
1442263_at	0.94845	regulator of G-protein signaling 13	Rgs13
1442264_at	0.82412	"RAS, guanyl releasing protein 2"	Rasgrp2

1442265_at 0.88128 "Fanconi anemia, complementation group C" FancC
1442266_at 0.97523 expressed sequence AI662175 AI662175
1442267_at 1.08347 syntaxin binding protein 4 Stxbp4
1442268_a_at 1.06542 "Adult male epididymis cDNA, RIKEN full-length enriched library, clone:9230001K01 product:unclassifiable, full insert sequence" ---
1442269_at 1.07216 "Adult male epididymis cDNA, RIKEN full-length enriched library, clone:9230001K01 product:unclassifiable, full insert sequence" ---
1442270_at 1.91402 Pre B-cell leukemia transcription factor 3 Pbx3
1442271_at 1.15263 Cytoskeleton associated protein 5 Ckap5
1442272_at 0.91374 Transcribed locus ---
1442273_at 1.04068 RIKEN cDNA 4932413O14 gene 4932413O14Rik
1442274_at 0.87772 "zinc finger, DHHC domain containing 15" Zdhhc15
1442275_at 0.97766 RIKEN cDNA 4933402J07 gene 4933402J07Rik
1442276_at 1.00055 Latrophilin 3 Lphn3
1442277_at 2.01496 choline kinase alpha Chka
1442278_at 1.02478 "Jumonji, AT rich interactive domain 1B (Rbp2 like)" Jarid1b
1442279_at 1.60905 Enhancer of polycomb homolog 1 (Drosophila) Epc1
1442280_at 1.09387 "DNA segment, Chr 2, ERATO Doi 750, expressed" D2Ertd750e
1442281_at 0.91632 "DNA segment, Chr 4, ERATO Doi 796, expressed" D4Ertd796e
1442282_at 1.05506 "Similar to vomeronasal 2, receptor, 2" LOC625068
1442283_at 1.04274 Phosphoglycerate dehydrogenase like 1 Phgdh1
1442284_at 0.92455 --- ---
1442285_at 1.27163 synaptic nuclear envelope 2 Syne2
1442286_at 1.18914 WD repeat domain 67 Wdr67
1442287_at 0.99050 Transcribed locus ---
1442288_at 1.12040 Annexin A6 Anxa6
1442289_at 1.12492 Josephin domain containing 3 Jsd3
1442290_at 0.89697 --- ---
1442291_at 0.97070 "endothelial differentiation, lysophosphatidic acid G-protein-coupled receptor 4" Edg4
1442292_at 0.99966 Rac/Cdc42 guanine nucleotide exchange factor (GEF) 6 Arhgef6
1442293_at 0.94187 Transcribed locus ---
1442294_at 0.99912 RIKEN cDNA A530053G22 gene A530053G22Rik
1442295_at 1.63233 "Actin related protein 2/3 complex, subunit 2" Arpc2
1442296_at 1.08475 WD repeat domain 75 Wdr75
1442297_at 0.90227 expressed sequence BB212172 BB212172
1442298_at 1.56127 Bicaudal C homolog 1 (Drosophila) Bicc1
1442299_at 1.05960 --- ---
1442300_at 1.01079 --- ---
1442301_at 0.92516 expressed sequence AW060742 AW060742
1442302_at 1.48810 N-methylpurine-DNA glycosylase Mpg
1442303_at 1.01132 "10, 11 days embryo whole body cDNA, RIKEN full-length enriched library, clone:2810048J24 product:unclassifiable, full insert sequence" ---

1442304_at 0.99034 "16 days neonate heart cDNA, RIKEN full-length enriched library, clone:
 D830044D09 product:unclassifiable, full insert sequence" ---
 1442305_at 1.12789 GTP binding protein 2 Gtpbp2
 1442306_at 1.05202 Transcribed locus ---
 1442307_at 0.81981 Transcribed locus ---
 1442308_at 1.11880 SET and MYND domain containing 4 Smyd4
 1442309_at 1.08140 DNA methyltransferase 3A Dnmt3a
 1442310_at 1.18800 "Phosphatidylinositol-4-phosphate 5-kinase, type II, alpha" Pip5k2a
 1442311_at 3.24750 Zinc finger RNA binding protein Zfr
 1442312_at 0.93710 transducin (beta)-like 1X-linked receptor 1 Tbl1xr1
 1442313_at 0.92251 RIKEN cDNA 4933400E14 gene 4933400E14Rik
 1442314_at 0.90046 nucleoporin 188 Nup188
 1442315_at 1.03231 expressed sequence AI426778 AI426778
 1442316_x_at 1.03915 transformation related protein 53 binding protein 1 Trp53bp1
 1442317_at 1.54119 T-complex protein 1 Tcp1
 1442318_at 1.13281 Transcribed locus ---
 1442319_at 0.95907 ubiquitin specific peptidase 4 (proto-oncogene) Usp4
 1442320_at 2.97558 hypothetical LOC553096 LOC553096
 1442321_at 0.92627 COMM domain containing 2 Commd2
 1442322_at 1.03252 Slit homolog 2 (Drosophila) Slit2
 1442323_at 0.98011 RIKEN cDNA 2410193C02 gene 2410193C02Rik
 1442324_at 1.01892 "PREDICTED: Mus musculus hypothetical LOC547070 (LOC547070),
 mRNA" ---
 1442325_at 0.97168 "TBC1 domain family, member 24" Tbc1d24
 1442326_at 0.99093 protocadherin 15 Pcdh15
 1442327_at 0.89651 RIKEN cDNA 4930482J15 gene 4930482J15Rik
 1442328_at 0.95060 --- ---
 1442329_at 1.07053 "DNA cross-link repair 1A, PSO2 homolog (S. cerevisiae)" Dclre1a
 1442330_at 1.14110 Serine/arginine repetitive matrix 1 Srrm1
 1442331_at 1.03366 Aminolevulinic acid synthase 1 Alas1
 1442332_at 1.07884 "Transforming growth factor, beta receptor III" Tgfbr3
 1442333_a_at 0.94952 "sodium channel, voltage-gated, type IX, alpha /// similar to sodium channel
 25" Scn9a /// LOC671835
 1442334_at 0.88761 PRAME family member LOC545693 MGC118150
 1442335_at 1.04557 Transcribed locus ---
 1442336_at 1.07284 --- ---
 1442337_at 1.03890 Adenylate cyclase activating polypeptide 1 receptor 1 Adcyap1r1
 1442338_at 1.19445 Transcribed locus ---
 1442339_at 1.00610 stefin A2 like 1 Stfa2l1
 1442340_x_at 0.78496 cysteine rich protein 61 Cyr61
 1442341_at 0.93413 "kalirin, RhoGEF kinase" Kalrn
 1442342_at 0.96462 trans-acting transcription factor 6 Sp6
 1442343_at 0.91784 Thioredoxin domain containing 11 Txndc11

1442344_at 1.07472 AP1 gamma subunit binding protein 1 Ap1gbp1
1442345_at 1.07841 RIKEN cDNA 1700031L13 gene 1700031L13Rik
1442346_at 1.00072 Transcribed locus ---
1442347_at 1.09861 "low density lipoprotein receptor-related protein 8, apolipoprotein e receptor"
Lrp8
1442348_at 1.05068 --- ---
1442349_at 1.21509 RIKEN cDNA C630028N24 gene C630028N24Rik
1442350_at 1.46060 "0 day neonate skin cDNA, RIKEN full-length enriched library,
clone:4632424N07 product:unclassifiable, full insert sequence" ---
1442351_a_at 1.02035 cDNA sequence BC029214 BC029214
1442352_at 0.99441 RIKEN cDNA 9430091N11 gene 9430091N11Rik
1442353_at 0.91465 inosine triphosphatase (nucleoside triphosphate pyrophosphatase) Itpa
1442354_at 0.93614 RIKEN cDNA B230325K18 gene B230325K18Rik
1442355_at 0.99140 hypothetical LOC546100 LOC546100
1442356_at 1.04917 MAX gene associated Mga
1442357_at 1.05882 Rho guanine nucleotide exchange factor (GEF) 17 Arhgef17
1442358_at 1.76116 expressed sequence AA409587 AA409587
1442359_at 1.03663 six transmembrane epithelial antigen of prostate 2 Steap2
1442360_at 1.00973 Autism susceptibility candidate 2 Auts2
1442361_at 1.02103 Integrin alpha 2b Itga2b
1442362_at 0.97559 "gene model 104, (NCBI)" Gm104
1442363_at 1.11549 RIKEN cDNA 1110012J17 gene 1110012J17Rik
1442364_at 1.09496 Mitogen activated protein kinase 14 Mapk14
1442365_at 1.72271 Reticulon 3 Rtn3
1442366_at 1.09499 RIKEN cDNA 6820408C15 gene 6820408C15Rik
1442367_at 0.87583 "Atpase, class VI, type 11C" Atp11c
1442368_at 0.92332 potassium channel tetramerisation domain containing 12b Kctd12b
1442369_at 1.03689 RIKEN cDNA 4832406H04 gene 4832406H04Rik
1442370_at 1.00918 "10 days neonate cortex cDNA, RIKEN full-length enriched library, clone:
A830028E23 product:unclassifiable, full insert sequence" ---
1442371_at 0.84155 RIKEN cDNA 1110048D14 gene 1110048D14Rik
1442372_at 0.93378 Transcribed locus ---
1442373_at 0.98933 RIKEN cDNA 2610203C20 gene 2610203C20Rik
1442374_at 1.07980 "DNA segment, Chr 2, ERATO Doi 127, expressed" D2Ertd127e
1442375_at 1.36612 Ubiquitin-associated protein 2 Ubap2
1442376_at 1.06079 actin-binding LIM protein 1 Ablim1
1442377_at 1.06566 "Adult male corpora quadrigemina cDNA, RIKEN full-length enriched library,
clone:B230366E07 product:unclassifiable, full insert sequence" ---
1442378_x_at 1.19910 "In vitro fertilized eggs cDNA, RIKEN full-length enriched library,
clone:7420405G17 product:unclassifiable, full insert sequence" ---
1442379_at 1.03157 hypothetical protein LOC574403 MGC99845
1442380_at 1.09374 lipocalin 4 Lcn4
1442381_at 2.41477 "Muskelin 1, intracellular mediator containing kelch motifs" Mkl1

1442382_at	1.33692	microtubule associated serine/threonine kinase family member 4	Mast4
1442383_at	1.38649	TRIO and F-actin binding protein	Triobp
1442384_at	1.00788	"G elongation factor, mitochondrial 2"	Gfm2
1442385_at	1.14149	Methionine sulfoxide reductase A	Msra
1442386_at	0.97559	RIKEN cDNA C030010B13 gene	C030010B13Rik
1442387_at	1.00953	Histidyl-tRNA synthetase	Hars
1442388_at	0.91817	RIKEN cDNA 2410137F16 gene	2410137F16Rik
1442389_at	0.97494	2'-5' oligoadenylate synthetase 1F	Oas1f
1442390_at	1.04020	Transcribed locus	---
1442391_at	0.95837	RIKEN cDNA 4930519H02 gene	4930519H02Rik
1442392_at	0.95164	"0 day neonate thymus cDNA, RIKEN full-length enriched library, clone: A430024M08 product:unclassifiable, full insert sequence"	---
1442393_at	1.16547	Zinc finger homeobox 1b	Zfhx1b
1442394_at	0.97333	"0 day neonate thymus cDNA, RIKEN full-length enriched library, clone: A430024H08 product:unclassifiable, full insert sequence"	---
1442395_at	1.45564	Dystonin	Dst
1442396_at	1.05733	RIKEN cDNA A730020M07 gene	A730020M07Rik
1442397_at	0.98159	"nuclear transcription factor, X-box binding 1"	Nfx1
1442398_at	0.98326	ubiquitin-conjugating enzyme E2T (putative)	Ube2t
1442399_at	0.99138	5'-3' exoribonuclease 1	Xrn1
1442400_at	1.15838	Prickle like 1 (Drosophila)	Prickle1
1442401_at	1.11888	Sorbin and SH3 domain containing 1	Sorbs1
1442402_at	0.99898	SH3 domain containing ring finger 1	Sh3rf1
1442403_at	1.05818	RIKEN cDNA B130034C11 gene	B130034C11Rik
1442404_at	0.96571	nucleolin	Ncl
1442405_at	0.94737	Cysteine and glycine-rich protein 2 binding protein	Csrp2bp
1442406_at	0.98159	RIKEN cDNA 9230104K21 gene	9230104K21Rik
1442407_at	0.99822	---	---
1442408_at	0.84270	RIKEN cDNA 2010305B15 gene	2010305B15Rik
1442409_at	1.03465	"DNA segment, Chr 9, Wayne State University 90, expressed"	D9Wsu90e
1442410_at	1.08199	DEP domain containing 5	Depdc5
1442411_at	0.99619	Glucocorticoid induced transcript 1	Glcci1
1442412_at	0.81971	"dynein, axonemal, heavy chain 9"	Dnahc9
1442413_at	1.17436	"potassium voltage-gated channel, shaker-related subfamily, member 1"	Kcna1
1442414_at	1.07991	Ring finger protein 103	Rnf103
1442415_at	0.89530	RIKEN cDNA 5830454E08 gene	5830454E08Rik
1442416_at	1.33092	hypothetical protein D930001B02	D930001B02
1442417_at	1.01353	"mediator of RNA polymerase II transcription, subunit 8 homolog (yeast)"	Med8
1442418_at	1.82480	RIKEN cDNA B930096F20 gene	B930096F20Rik
1442419_at	1.01032	Transcribed locus	---
1442420_at	1.15197	Like-glycosyltransferase	Large
1442421_at	1.07050	---	---

1442422_at 1.60046 Transmembrane protein 55A Tmem55a
1442423_at 0.99216 Neurexin III Nrnx3
1442424_at 0.96969 "CTD (carboxy-terminal domain, RNA polymerase II, polypeptide A) small phosphatase like 2" Ctdspl2
1442425_at 1.09618 RIKEN cDNA A030014E15 gene A030014E15Rik
1442426_at 0.93968 wings apart-like homolog (Drosophila) Wapal
1442427_at 1.27952 RIKEN cDNA 9630026M06 gene 9630026M06Rik
1442428_at 0.94656 "Adult male hypothalamus cDNA, RIKEN full-length enriched library, clone: A230084D06 product:unclassifiable, full insert sequence" ---
1442429_at 1.11325 "Adaptor protein complex AP-2, mu1" Ap2m1
1442430_at 0.92722 cDNA sequence BC048562 BC048562
1442431_at 0.97744 Transcribed locus ---
1442432_x_at 0.88880 ataxin 7-like 1 Atxn7l1
1442433_at 1.07292 --- ---
1442434_at 0.69580 "DNA segment, Chr 8, ERATO Doi 82, expressed" D8Ertd82e
1442435_at 1.87888 "Smg-6 homolog, nonsense mediated mRNA decay factor (C. elegans)" Smg6
1442436_at 0.90472 fructosamine 3 kinase Fn3k
1442437_at 0.98463 PTK2 protein tyrosine kinase 2 beta Ptk2b
1442438_at 0.95805 --- ---
1442439_at 0.95844 expressed sequence W91776 W91776
1442440_at 0.97952 Neuronal d4 domain family member Neud4
1442441_at 1.00890 "DNA segment, Chr 11, Brigham & Women's Genetics 0434 expressed" D11Bwg0434e
1442442_at 1.14678 THO complex 7 homolog (Drosophila) Thoc7
1442443_at 1.00897 Retinoblastoma binding protein 4 Rbbp4
1442444_at 1.07041 DEAD (Asp-Glu-Ala-Asp) box polypeptide 19b Ddx19b
1442445_at 1.12528 RIKEN cDNA 2610027H17 gene 2610027H17Rik
1442446_at 1.03029 Zinc finger protein 523 Zfp523
1442447_at 0.94270 --- ---
1442448_at 1.01206 RIKEN cDNA 1810064F22 gene 1810064F22Rik
1442449_at 1.06850 "solute carrier family 6 (neurotransmitter transporter, noradrenalin), member 2" Slc6a2
1442450_at 0.94468 "DNA segment, Chr 2, ERATO Doi 501, expressed" D2Ertd501e
1442451_at 0.95819 Tetratricopeptide repeat domain 15 Ttc15
1442452_at 0.99942 "amyotrophic lateral sclerosis 2 (juvenile) chromosome region, candidate 2 (human)" Als2cr2
1442453_at 1.73429 FCH domain only 2 Fcho2
1442454_at 1.43981 topoisomerase (DNA) II alpha Top2a
1442455_at 1.18438 expressed sequence C79845 C79845
1442456_at 1.03029 spermatogenesis associated 5 Spata5
1442457_at 0.86720 --- ---
1442458_at 0.91054 Transcribed locus ---

1442459_at 0.93591 "a disintegrin-like and metallopeptidase (reprolysin type) with thrombospondin type 1 motif, 19" Adamts19

1442460_at 0.96565 --- ---

1442461_at 1.03726 glucose-fructose oxidoreductase domain containing 1 Gfod1

1442462_at 1.02131 expressed sequence C87042 C87042

1442463_at 1.05191 "RAB10, member RAS oncogene family" Rab10

1442464_at 1.01519 F-box and leucine-rich repeat protein 20 Fbxl20

1442465_s_at 0.89992 spermatid perinuclear RNA binding protein Strbp

1442466_a_at 0.96513 histidine acid phosphatase domain containing 2A Hisppd2a

1442467_at 1.06131 RIKEN cDNA 5330421F07 gene 5330421F07Rik

1442468_at 1.15849 "0 day neonate thymus cDNA, RIKEN full-length enriched library, clone: A430088L10 product:unclassifiable, full insert sequence" ---

1442469_at 1.00217 --- ---

1442470_at 1.92696 --- ---

1442471_at 1.04504 Eph receptor B2 Ephb2

1442472_at 0.92101 Tetratricopeptide repeat domain 21A ttc21a

1442473_at 2.48532 --- ---

1442474_at 0.93845 --- ---

1442475_at 1.14582 "DNA segment, Chr 14, ERATO Doi 574, expressed" D14Ertd574e

1442476_at 0.98873 Intraflagellar transport 81 homolog (Chlamydomonas) Ift81

1442477_at 1.13537 --- ---

1442478_at 1.05182 --- ---

1442479_at 1.01256 "10 days neonate cortex cDNA, RIKEN full-length enriched library, clone: A830040F21 product:unclassifiable, full insert sequence" ---

1442480_at 0.89281 Transcribed locus ---

1442481_at 0.91854 hypothetical protein LOC675578 LOC675578

1442482_at 0.94637 RIKEN cDNA 1700106N22 gene 1700106N22Rik

1442483_at 1.11551 Fucosyltransferase 8 Fut8

1442484_at 2.08246 "DNA segment, Chr 9, ERATO Doi 306, expressed" D9Ertd306e

1442485_at 0.92068 --- ---

1442486_at 1.15622 --- ---

1442487_at 0.91716 "DNA segment, Chr 14, ERATO Doi 24, expressed" D14Ertd24e

1442488_at 0.95334 Glypican 5 Gpc5

1442489_at 1.15835 "DNA segment, Chr 1, ERATO Doi 564, expressed" D1Ertd564e

1442490_at 1.03729 RIKEN cDNA C130090I23 gene C130090I23Rik

1442491_at 1.10974 Dipeptidylpeptidase 10 Dpp10

1442492_at 1.10252 "DNA segment, Chr X, ERATO Doi 11, expressed" DXErtd11e

1442493_at 1.08456 Transcribed locus ---

1442494_at 1.79748 Ubiquitin protein ligase E3 component n-recognin 2 Ubr2

1442495_at 1.17290 Baculoviral IAP repeat-containing 6 Birc6

1442496_at 1.14117 Cyclin-dependent kinase-like 3 Cdkl3

1442497_at 1.14384 SMC hinge domain containing 1 Smchd1

1442498_at 1.19616 EST C78662 C78662

1442499_at 1.14943 carnitine deficiency-associated gene expressed in ventricle 3 Cdv3
1442500_at 0.88132 expressed sequence C77406 C77406
1442501_at 1.02848 neurexin I Nrnx1
1442502_at 0.99164 leucine-rich repeats and calponin homology (CH) domain containing 1 Lrch1
1442503_at 0.84181 --- ---
1442504_at 0.96431 "DNA segment, Chr 18, ERATO Doi 653, expressed" D18Ertd653e
1442505_at 1.48728 Fibronectin type III domain containing 3B Fndc3b
1442506_at 0.97118 Transcribed locus ---
1442507_at 1.15624 whirlin Whrn
1442508_at 1.10808 RIKEN cDNA 4933404M19 gene /// similar to hypothetical protein
MGC39650 /// similar to proline-rich cyclin A1-interacting protein 4933404M19Rik /// RP23-
185A18.6 /// LOC216974
1442509_at 2.14001 Ecotropic viral integration site 5 Evi5
1442510_at 1.19827 hypothetical protein LOC665792 /// hypothetical protein LOC669957 ///
hypothetical protein LOC674416 LOC665792 /// LOC669957 /// LOC674416
1442511_at 1.53929 importin 7 Ipo7
1442512_at 1.43588 CDC42 effector protein (Rho GTPase binding) 1 Cdc42ep1
1442513_at 1.14384 cDNA sequence BC016423 /// similar to cDNA sequence BC016423
BC016423 /// LOC669774
1442514_a_at 0.90117 elongation factor RNA polymerase II-like 3 Ell3
1442515_at 1.07444 --- ---
1442516_at 0.96273 "DNA segment, Chr 2, ERATO Doi 623, expressed" D2Ertd623e
1442517_a_at 1.04267 RIKEN cDNA 9630013A20 gene 9630013A20Rik
1442518_at 0.93882 RIKEN cDNA C030044O21 gene C030044O21Rik
1442519_at 0.91607 Dendritic cell protein GA17 Ga17
1442520_at 1.02439 RIKEN cDNA 5730509C05 gene 5730509C05Rik
1442521_at 1.15320 "Adult male testis cDNA, RIKEN full-length enriched library,
clone:4930568O12 product:unclassifiable, full insert sequence" ---
1442522_at 0.99683 --- ---
1442523_at 1.00980 similar to Nucleolar phosphoprotein p130 (Nucleolar 130 kDa protein) (140
kDa nucleolar phosphoprotein) (Nopp140) (Nucleolar and coiled-body phosphoprotein 1) LOC331392
1442524_at 1.10052 Transcribed locus ---
1442525_at 0.93305 Transcribed locus ---
1442526_at 0.89674 expressed sequence C77681 C77681
1442527_at 0.97530 expressed sequence C87583 C87583
1442528_at 1.20635 exportin 4 Xpo4
1442529_at 0.92387 Opioid growth factor receptor-like 1 Ogfr1
1442530_at 0.93817 --- ---
1442531_at 1.37756 "DNA segment, Chr 12, ERATO Doi 123, expressed" D12Ertd123e
1442532_at 1.11987 --- ---
1442533_at 0.92884 "Nuclear receptor subfamily 3, group C, member 2" Nr3c2
1442534_at 0.93590 Glycosyltransferase 25 domain containing 2 Glt25d2
1442535_at 1.60347 Zinc finger protein 609 Zfp609

1442536_at 1.01003 hypothetical protein LOC619719 LOC619719
1442537_at 1.01111 --- ---
1442538_at 1.23326 REST corepressor 3 Rcor3
1442539_at 0.93072 "WW, C2 and coiled-coil domain containing 2" Wwc2
1442540_at 1.04736 expressed sequence C77609 C77609
1442541_at 1.13283 Ornithine decarboxylase antizyme 2 Oaz2
1442542_at 0.67018 eyes absent 4 homolog (Drosophila) Eya4
1442543_at 1.08324 Testis expressed gene 264 Tex264
1442544_at 0.92853 immunoglobulin heavy chain 4 (serum IgG1) Igh-4
1442545_at 0.94922 --- ---
1442546_at 0.77294 RIKEN cDNA C730027H18 gene C730027H18Rik
1442547_at 1.16858 Transcribed locus ---
1442548_at 1.43507 Cysteine rich transmembrane BMP regulator 1 (chordin like) Crim1
1442549_at 1.18982 muscleblind-like 3 (Drosophila) Mbnl3
1442550_at 0.97904 RIKEN cDNA 1110033M05 gene 1110033M05Rik
1442551_at 0.93358 --- ---
1442552_at 1.09554 "thrombospondin, type I, domain containing 4" Thsd4
1442553_at 0.81045 "microtubule-associated protein, RP/EB family, member 2" Mapre2
1442554_s_at 0.94154 "kalirin, RhoGEF kinase" Kalrn
1442555_at 1.15772 "potassium voltage-gated channel, Shal-related family, member 3" Kcnd3
1442556_at 1.10390 --- ---
1442557_at 0.87550 Synaptotagmin I Syt1
1442558_at 1.02300 "Adult male hypothalamus cDNA, RIKEN full-length enriched library, clone:
A230083A21 product:hypothetical protein, full insert sequence" ---
1442559_at 1.02483 "10 days neonate cerebellum cDNA, RIKEN full-length enriched library, clone:
B930020D16 product:hypothetical protein, full insert sequence" ---
1442560_at 0.88792 "Adult male testis cDNA, RIKEN full-length enriched library,
clone:4933433O05 product:unclassifiable, full insert sequence" ---
1442561_at 1.05350 MAM domain containing 1 Mamdc1
1442562_at 0.84291 RIKEN cDNA D630040G17 gene /// similar to zinc finger protein 64
(predicted) D630040G17Rik /// LOC622675
1442563_at 1.04437 Transmembrane protein 8 (five membrane-spanning domains) Tmem8
1442564_at 1.01848 --- ---
1442565_at 0.92856 protocadherin 10 Pcdh10
1442566_at 1.09895 expressed sequence C78878 C78878
1442567_at 1.00042 zinc finger and BTB domain containing 24 Zbtb24
1442568_at 0.93658 RIKEN cDNA C330002I19 gene C330002I19Rik
1442569_at 1.19148 Chaperonin subunit 4 (delta) Cct4
1442570_at 1.00879 "Zinc finger, MYM domain containing 1" Zmym1
1442571_at 1.02791 "13 days embryo head cDNA, RIKEN full-length enriched library,
clone:3110033I20 product:hypothetical protein, full insert sequence" ---
1442572_at 0.98388 Gamma-aminobutyric acid (GABA(A)) receptor-associated protein-like 1
Gabarapl1

1442573_at	0.82737	RIKEN cDNA A930028L21 gene	A930028L21Rik
1442574_at	0.90739	"Crystallin, zeta (quinone reductase)-like 1"	Cryz11
1442575_at	1.06071	---	---
1442576_at	0.89340	CAMP responsive element binding protein 5	Creb5
1442577_at	1.19326	KIAA0415 protein	KIAA0415
1442578_at	1.04346	---	---
1442579_at	1.11273	RIKEN cDNA 9330111N05 gene	9330111N05Rik
1442580_at	0.95382	RIKEN cDNA A530020G20 gene	A530020G20Rik
1442581_at	1.64087	kinase suppressor of ras 1	Ksr1
1442582_at	0.93126	germinal center expressed transcript 2	Gcet2
1442583_a_at	1.08482	expressed sequence AI854703	AI854703
1442584_at	1.03612	---	---
1442585_at	0.97954	RIKEN cDNA B230220N19 gene	B230220N19Rik
1442586_at	0.84151	suppressor of cytokine signaling 2	Socs2
1442587_at	0.83671	---	---
1442588_at	1.19547	"Mus musculus, clone IMAGE:1533345, mRNA"	---
1442589_at	0.89061	"Transient receptor potential cation channel, subfamily C, member 5"	Trpc5
1442590_at	1.03970	"tumor necrosis factor receptor superfamily, member 23"	Tnfrsf23
1442591_at	1.01899	Expressed sequence AW060207	AW060207
1442592_at	0.98637	Phosphatidic acid phosphatase type 2 domain containing 3	Ppapdc3
1442593_at	1.37290	---	---
1442594_at	1.21322	Ttk protein kinase	Ttk
1442595_at	0.94422	Expressed sequence C77713	C77713
1442596_at	0.97453	kelch domain containing 2	Klhdc2
1442597_at	1.11349	expressed sequence C79122	C79122
1442598_at	0.86435	Prkr interacting protein 1 (IL11 inducible)	Prkrip1
1442599_at	1.19340	"Solute carrier family 12 (potassium/chloride transporters), member 9"	Slc12a9
1442600_at	0.97585	---	---
1442601_at	1.07279	glucocorticoid modulatory element binding protein 1	Gmeb1
1442602_at	0.86678	---	---
1442603_at	1.54349	Polybromo 1	Pb1
1442604_at	1.03595	"excision repair cross-complementing rodent repair deficiency, complementation group 6"	Ercc6
1442605_at	1.05464	BTB and CNC homology 2	Bach2
1442606_at	1.13548	Dynamin binding protein Dnmbp	
1442607_a_at	0.98496	---	---
1442608_at	0.90301	layilin	Layn
1442609_at	1.66803	"Protein kinase C, epsilon"	Prkce
1442610_at	0.89417	Plexin domain containing 1	Plxdc1
1442611_at	1.12667	RIKEN cDNA 4930592C13 gene	4930592C13Rik
1442612_at	1.09607	RIKEN cDNA C730036E19 gene	C730036E19Rik
1442613_at	1.23695	---	---
1442614_at	0.96621	interleukin 1 receptor accessory protein	Il1rap

1442615_at 0.98967 Adenosine kinase Adk
1442616_at 1.11435 GRP1 (general receptor for phosphoinositides 1)-associated scaffold protein
Grasp
1442617_at 1.14553 Core binding factor beta Cbfb
1442618_at 0.82220 Lactate dehydrogenase B Ldhb
1442619_at 1.08561 Kruppel-like factor 15 Klf15
1442620_at 1.06346 RIKEN cDNA 4931420C21 gene 4931420C21Rik
1442621_at 1.06357 expressed sequence C77798 C77798
1442622_at 1.18678 Spermatid perinuclear RNA binding protein Strbp
1442623_at 1.01225 "LysM, putative peptidoglycan-binding, domain containing 4" Lysmd4
1442624_at 1.80902 RIKEN cDNA C920008N22 gene C920008N22Rik
1442625_at 1.12212 "Pleckstrin homology domain containing, family C (with FERM domain)
member 1" Plekhc1
1442626_at 1.07650 RIKEN cDNA 5730405I09 gene 5730405I09Rik
1442627_at 1.02841 --- ---
1442628_at 1.03204 "15 days embryo head cDNA, RIKEN full-length enriched library, clone:
D930002E06 product:hypothetical protein, full insert sequence" ---
1442629_at 1.71047 "5'-nucleotidase, cytosolic II" Nt5c2
1442630_at 0.97601 Transcribed locus ---
1442631_at 0.99324 RIKEN cDNA 2700070H01 gene 2700070H01Rik
1442632_at 2.35581 "Centaurin, gamma 2" Centg2
1442633_at 1.07846 "Transcribed locus, weakly similar to XP_576460.1 PREDICTED: similar to
hypothetical protein PB402898.00.0 [Rattus norvegicus]" ---
1442634_at 0.90281 Transcribed locus ---
1442635_at 1.05879 "Adult male corpora quadrigemina cDNA, RIKEN full-length enriched library,
clone:B230312L02 product:unclassifiable, full insert sequence" ---
1442636_at 0.98517 CDNA sequence BC004636 BC004636
1442637_at 1.03240 Intersectin 1 (SH3 domain protein 1A) Itsn1
1442638_at 0.95326 --- ---
1442639_at 1.00320 --- ---
1442640_at 0.91638 Schlafen 5 Slfn5
1442641_at 1.37401 Ankyrin repeat and KH domain containing 1 Ankhd1
1442642_at 1.03625 RIKEN cDNA 4930413O22 gene 4930413O22Rik
1442643_at 1.05236 jumonji domain containing 3 Jmjd3
1442644_at 1.02883 "PREDICTED: Mus musculus hypothetical protein LOC622030 (LOC622030),
mRNA" ---
1442645_at 1.03287 "ATPase, Ca++ transporting, plasma membrane 3" Atp2b3
1442646_at 1.00203 B-cell scaffold protein with ankyrin repeats 1 Bank1
1442647_at 1.12011 "Yip1 domain family, member 1" Yipf1
1442648_at 0.86459 hypothetical protein D130054H01 D130054H01
1442649_at 0.96560 RIKEN cDNA 1500009L16 gene 1500009L16Rik
1442650_at 1.05247 Transcribed locus ---
1442651_at 1.53540 WW domain-containing oxidoreductase Wwox

1442652_at	0.93321	"gene model 410, (NCBI)"	Gm410
1442653_at	1.05545	---	---
1442654_at	0.89725	RIKEN cDNA A530054K11 gene	A530054K11Rik
1442655_at	1.07799	DNA methyltransferase 3B	Dnmt3b
1442656_at	1.34084	---	---
1442657_at	0.95717	"DNA segment, Chr 1, ERATO Doi 273, expressed"	D1Ertd273e
1442658_at	0.99229	"NADH dehydrogenase (ubiquinone) 1 beta subcomplex, 2"	Ndufb2
1442659_at	0.93430	protocadherin 9 Pcdh9	
1442660_at	1.02021	Yippee-like 1 (Drosophila)	Ypel1
1442661_at	1.21823	expressed sequence C80113	C80113
1442662_at	0.96273	RIKEN cDNA 0610041E09 gene	0610041E09Rik
1442663_at	0.91447	---	---
1442664_at	1.17795	Muscleblind-like 3 (Drosophila)	Mbnl3
1442665_at	1.06866	expressed sequence C77874	C77874
1442666_at	1.09082	"DNA segment, Chr 9, ERATO Doi 12, expressed"	D9Ertd12e
1442667_at	1.25505	cell division cycle 40 homolog (yeast)	Cdc40
1442668_at	0.87631	---	---
1442669_at	0.97243	---	---
1442670_at	0.99860	RIKEN cDNA 6430517E21 gene	6430517E21Rik
1442671_at	1.28346	Huntingtin interacting protein 2	Hip2
1442672_at	1.00320	RIKEN cDNA 0610040F04 gene	0610040F04Rik
1442673_at	1.39452	Signal transducing adaptor molecule (SH3 domain and ITAM motif) 2	Stam2
1442674_at	1.24324	F-box and WD-40 domain protein 4	Fbxw4
1442675_at	0.94675	RIKEN cDNA 2010111I01 gene	2010111I01Rik
1442676_at	0.89095	monoamine oxidase A	Maoa
1442677_at	1.19277	Transcribed locus	---
1442678_at	0.89854	coiled coil domain containing 28B	Ccdc28b
1442679_at	1.30923	Mitogen activated protein kinase kinase 4	Map2k4
1442680_at	1.07725	Neural cell adhesion molecule 1	Ncam1
1442681_at	1.17366	"TRNA splicing endonuclease 2 homolog (SEN2, S. cerevisiae)"	Tsen2
1442682_at	1.36629	---	---
1442683_at	1.07140	high mobility group 20A	Hmg20a
1442684_x_at	1.04566	high mobility group 20A	Hmg20a
1442685_at	0.96424	---	---
1442686_at	1.00366	"D4, zinc and double PHD fingers, family 3"	Dpf3
1442687_at	1.02467	RIKEN cDNA 1700064N11 gene	1700064N11Rik
1442688_at	0.91477	---	---
1442689_at	1.06163	"DNA segment, Chr 9, ERATO Doi 278, expressed"	D9Ertd278e
1442690_at	0.99161	Transmembrane protein 143	Tmem143
1442691_at	1.03724	"0 day neonate lung cDNA, RIKEN full-length enriched library, clone: E030049N05 product:unclassifiable, full insert sequence"	---
1442692_at	1.32165	Mitogen-activated protein kinase associated protein 1	Mapkap1
1442693_at	0.95306	similar to tripartite motif protein TRIM5	LOC667823

1442694_at	0.93811	---	---	
1442695_at	1.00339	RIKEN cDNA C030007I01 gene	C030007I01Rik	
1442696_at	1.00788	7-dehydrocholesterol reductase	Dhcr7	
1442697_at	1.17427	---	---	
1442698_at	1.15919	"7 days neonate cerebellum cDNA, RIKEN full-length enriched library, clone:		
A730053D01		product:unclassifiable, full insert sequence"	---	
1442699_at	0.91914	---	---	
1442700_at	1.07166	"phosphodiesterase 4B, cAMP specific"	Pde4b	
1442701_at	0.91646	---	---	
1442702_at	0.90446	Dorso-medial telencephalon gene 2	Dmt2	
1442703_at	1.22261	cDNA sequence AK220484	AK220484	
1442704_at	1.80840	"Bromodomain adjacent to zinc finger domain, 2B"	Baz2b	
1442705_at	1.18584	expressed sequence C76938	C76938	
1442706_at	1.04073	nasal embryonic LHRH factor	Nelf	
1442707_at	1.01378	calcium/calmodulin-dependent protein kinase II alpha	Camk2a	
1442708_at	0.98157	"0 day neonate kidney cDNA, RIKEN full-length enriched library, clone:		
D630048D13		product:unclassifiable, full insert sequence"	---	
1442709_at	1.13168	Zinc finger protein 521	Zfp521	
1442710_at	1.26629	PDZ and LIM domain 5	Pdlim5	
1442711_at	1.00606	Transcribed locus	---	
1442712_at	0.81645	"Similar to H3 histone, family 3B"	LOC627894	
1442713_at	1.01177	Oxysterol binding protein-like 10	Osbp110	
1442714_at	0.89010	hypothetical protein LOC667963	LOC667963	
1442715_at	1.05693	---	---	
1442716_at	0.94725	Transcribed locus	---	
1442717_at	1.09092	"TBC1 domain family, member 13"	Tbc1d13	
1442718_at	0.94202	"7 days neonate cerebellum cDNA, RIKEN full-length enriched library, clone:		
A730040F03		product:unclassifiable, full insert sequence"	---	
1442719_at	1.04856	protein arginine N-methyltransferase 3	Prmt3	
1442720_at	1.08959	---	---	
1442721_at	0.90734	Transcribed locus	---	
1442722_at	0.97532	Transcribed locus	---	
1442723_at	0.97831	Transcribed locus	---	
1442724_at	0.98431	expressed sequence AW060763	AW060763	
1442725_at	0.85671	---	---	
1442726_s_at	1.05429	expressed sequence AI845619	AI845619	
1442727_at	0.92270	---	---	
1442728_at	1.06604	Nuclear autoantigenic sperm protein (histone-binding)	Nasp	
1442729_at	0.90707	"DNA segment, Chr 12, ERATO Doi 553, expressed"	D12Ertd553e	
1442730_at	1.02857	---	---	
1442731_at	0.97418	RIKEN cDNA 9030416H16 gene	9030416H16Rik	
1442732_at	1.20841	---	---	
1442733_at	0.86755	---	---	

1442734_at	0.98238 --- ---	
1442735_at	1.02801 Ornithine decarboxylase antizyme 2	Oaz2
1442736_at	0.98947 kelch-like 11 (Drosophila)	Klhl11
1442737_at	0.77592 Hypothetical protein 4933423E17	4.93E+23
1442738_at	1.10246 CAMP responsive element binding protein-like 2	Crebl2
1442739_at	0.87987 cDNA sequence BC031441	BC031441
1442740_at	1.10069 --- ---	
1442741_at	0.96448 mutated in colorectal cancers	Mcc
1442742_at	0.98966 "ATPase, Ca ⁺⁺ -sequestering"	Atp2c1
1442743_at	0.92624 "Adult male olfactory brain cDNA, RIKEN full-length enriched library, clone:6430530F17 product:unclassifiable, full insert sequence" ---	
1442744_at	2.91189 expressed sequence C79248	C79248
1442745_x_at	2.16519 expressed sequence C79248	C79248
1442746_at	0.88381 --- ---	
1442747_at	1.02870 --- ---	
1442748_at	1.03703 BTB (POZ) domain containing 3	Btbd3
1442749_at	1.16609 Braf transforming gene	Braf
1442750_at	1.19230 Transcribed locus	---
1442751_at	0.94621 Triple functional domain (PTPRF interacting)	Trio
1442752_at	1.05986 Opioid binding protein/cell adhesion molecule-like	Opcml
1442753_at	0.94650 "tumor necrosis factor, alpha-induced protein 8"	Tnfaip8
1442754_at	0.90069 RIKEN cDNA C030013G03 gene	C030013G03Rik
1442755_at	0.97519 Transcribed locus	---
1442756_at	0.96635 Linker of T-cell receptor pathways	Lnk
1442757_at	0.96840 leucine-rich repeats and calponin homology (CH) domain containing 1	Lrch1
1442758_at	1.01772 Heterogeneous nuclear ribonucleoprotein A1	Hnrpa1
1442759_at	0.89132 thymoma viral proto-oncogene 1	Akt1
1442760_x_at	2.49097 Reticulon 4	Rtn4
1442761_at	1.49684 Tumor protein D52-like 2	Tpd52l2
1442762_at	1.01978 "tubulin tyrosine ligase-like family, member 10"	Ttl10
1442763_s_at	1.07365 "tubulin tyrosine ligase-like family, member 10"	Ttl10
1442764_at	1.10714 suppressor of variegation 4-20 homolog 1 (Drosophila)	Suv420h1
1442765_at	1.12304 Transcribed locus	---
1442766_at	1.06640 Transcribed locus	---
1442767_s_at	1.07827 "ubiquitin-activating enzyme E1, Chr X"	Ube1x
1442768_at	1.01102 RIKEN cDNA E130112L23 gene	E130112L23Rik
1442769_at	0.98614 RIKEN cDNA 8030451F13 gene	8030451F13Rik
1442770_at	0.88663 RIKEN cDNA 9530033F24 gene	9530033F24Rik
1442771_at	1.15453 RAD51-like 1 (S. cerevisiae)	Rad51l1
1442772_at	0.98890 EH domain binding protein 1	Ehbp1
1442773_at	1.06106 Transcribed locus	---
1442774_x_at	1.05665 Transcribed locus	---
1442775_at	1.53590 Suppression of tumorigenicity 13	St13

1442776_at 0.96093 RIKEN cDNA 2310009E04 gene 2310009E04Rik
1442777_at 1.03247 Transcribed locus ---
1442778_at 1.00360 expressed sequence AA511254 AA511254
1442779_at 1.00459 RIKEN cDNA 1700071K01 gene 1700071K01Rik
1442780_at 1.08617 RIKEN cDNA 2610002J02 gene 2610002J02Rik
1442781_at 1.01923 Ferric-chelate reductase 1 Frrs1
1442782_at 1.10489 Kelch-like 8 (Drosophila) Khlh8
1442783_x_at 1.07814 "3 days neonate thymus cDNA, RIKEN full-length enriched library, clone: A630030O14 product:unclassifiable, full insert sequence" ---
1442784_at 0.87665 Zinc finger protein 608 Zfp608
1442785_at 1.50337 SET and MYND domain containing 3 Smyd3
1442786_s_at 1.04833 RUN and FYVE domain containing 3 Rufy3
1442787_at 0.87811 --- ---
1442788_at 1.03092 RIKEN cDNA 2600003E23 gene 2600003E23Rik
1442789_at 1.14624 Transcribed locus ---
1442790_at 0.96416 --- ---
1442791_x_at 1.00638 RIKEN cDNA 6720407P12 gene 6720407P12Rik
1442792_x_at 0.83921 RIKEN cDNA 4930471M23 gene 4930471M23Rik
1442793_s_at 1.10813 transforming growth factor beta regulated gene 4 Tbrg4
1442794_at 0.98215 "procollagen, type VIII, alpha 1" Col8a1
1442795_x_at 1.00560 RIKEN cDNA 1700016G05 gene 1700016G05Rik
1442796_at 0.91755 activating transcription factor 7 interacting protein 2 Atf7ip2
1442797_x_at 0.87047 CKLF-like MARVEL transmembrane domain containing 2A Cmtm2a
1442798_x_at 1.01460 hexokinase 3 Hk3
1442799_x_at 0.99579 NudC domain containing 3 Nudcd3
1442800_x_at 0.98197 RIKEN cDNA A830059I20 gene A830059I20Rik
1442801_x_at 0.96559 transmembrane 7 superfamily member 3 Tm7sf3
1442802_x_at 0.99374 HLA-B associated transcript 2 Bat2
1442803_at 1.12500 RAD51-like 3 (S. cerevisiae) Rad51l3
1442804_at 0.85249 Gardner-Rasheed feline sarcoma viral (Fgr) oncogene homolog Fgr
1442805_at 1.20971 RIKEN cDNA A130086G11 gene A130086G11Rik
1442806_at 0.86559 RIKEN cDNA 9430030N17 gene 9430030N17Rik
1442807_at 0.92562 --- ---
1442808_at 0.88253 "Transcribed locus, weakly similar to XP_342027.2 PREDICTED: similar to MS4A8B protein [Rattus norvegicus]" ---
1442809_at 1.03054 "sodium channel, voltage-gated, type IX, alpha /// similar to sodium channel 25" Scn9a /// LOC671835
1442810_x_at 0.98794 "sodium channel, voltage-gated, type IX, alpha /// similar to sodium channel 25" Scn9a /// LOC671835
1442811_at 2.03451 "RGM domain family, member B" Rgmb
1442812_at 1.11189 similar to Anapc5 protein LOC433844
1442813_at 1.02629 "Diacylglycerol kinase, iota" Dgki
1442814_at 1.09501 Glucocorticoid induced transcript 1 Glcci1

1442815_at 1.01385 Peroxisome biogenesis factor 19 Pex19
1442816_at 0.78003 heparan sulfate (glucosamine) 3-O-sulfotransferase 3B1 Hs3st3b1
1442817_at 1.09050 RIKEN cDNA A730091E23 gene A730091E23Rik
1442818_at 1.06209 hypothetical protein LOC671796 /// hypothetical protein LOC675117
LOC671796 /// LOC675117
1442819_at 1.09886 "rhomboid, veinlet-like 2 (Drosophila)" Rhbdl2
1442820_at 1.12050 "Protein tyrosine phosphatase, non-receptor type 22 (lymphoid)" Ptpn22
1442821_at 2.07618 Transmembrane 9 superfamily member 3 Tm9sf3
1442822_at 1.03513 Calcium/calmodulin-dependent protein kinase IV Camk4
1442823_at 0.93251 "0 day neonate eyeball cDNA, RIKEN full-length enriched library, clone:
E130001F20 product:hypothetical protein, full insert sequence" ---
1442824_at 1.01051 RIKEN cDNA 8030497I03 gene 8030497I03Rik
1442825_at 1.08850 Golgi reassembly stacking protein 2 Gorasp2
1442826_at 1.09648 Ubiquitin specific peptidase 25 Usp25
1442827_at 1.00018 Toll-like receptor 4 Tlr4
1442828_at 0.93462 Interferon activated gene 203 Ifi203
1442829_at 0.99494 Ras and Rab interactor 2 Rin2
1442830_at 1.17349 Nucleolar and spindle associated protein 1 Nusap1
1442831_at 0.98604 RIKEN cDNA 2900026A02 gene 2900026A02Rik
1442832_at 1.00164 ankyrin repeat domain 52 Ankrd52
1442833_at 1.16514 "DNA segment, Chr 15, ERATO Doi 30, expressed" D15ErtD30e
1442834_at 1.60916 "Protein phosphatase 4, regulatory subunit 2" Ppp4r2
1442835_at 0.96918 "0 day neonate lung cDNA, RIKEN full-length enriched library, clone:
E030050O06 product:unclassifiable, full insert sequence" ---
1442836_at 1.06514 exportin 7 Xpo7
1442837_at 0.92058 "Protein tyrosine phosphatase, receptor type, D" Ptpnd
1442838_at 0.94944 "Adult male testis cDNA, RIKEN full-length enriched library,
clone:4930532C19 product:hypothetical protein, full insert sequence" ---
1442839_at 1.14733 --- ---
1442840_at 1.03983 "DNA segment, Chr 4, ERATO Doi 669, expressed" D4ErtD669e
1442841_at 1.07681 Tetraspanin 3 Tspan3
1442842_at 0.93974 Hspb associated protein 1 Hspbap1
1442843_at 1.67966 RIKEN cDNA 4933411D12 gene 4933411D12Rik
1442844_at 0.98100 RIKEN cDNA A830052D11 gene A830052D11Rik
1442845_at 3.83691 RIKEN cDNA C130075A20 gene C130075A20Rik
1442846_at 1.09050 Pre B-cell leukemia transcription factor 1 Pbx1
1442847_at 1.10037 --- ---
1442848_at 0.94499 3-hydroxyisobutyrate dehydrogenase Hibadh
1442849_at 1.36708 low density lipoprotein receptor-related protein 1 Lrp1
1442850_at 0.99354 --- ---
1442851_at 1.05210 expressed sequence C79818 C79818
1442852_at 0.91472 --- ---
1442853_at 0.95553 Mitogen-activated protein kinase 8 interacting protein 3 Mapk8ip3

1442854_at 0.91040 Transcribed locus ---
1442855_at 1.14716 dopey family member 1 Dopey1
1442856_at 1.12444 adrenomedullin 2 Adm2
1442857_at 1.04040 Astrotactin 2 Astn2
1442858_at 0.95849 "Pam, highwire, rpm 1" Phr1
1442859_at 1.05770 "Sprouty protein with EVH-1 domain 1, related sequence" Spred1
1442860_at 0.95826 "diacylglycerol kinase, beta" Dgkb
1442861_at 0.96522 CREB regulated transcription coactivator 1 Crtc1
1442862_at 1.09850 --- ---
1442863_at 0.98277 "calcium channel, voltage-dependent, alpha 2/delta subunit 4" Cacna2d4
1442864_at 1.15213 "estrogen related receptor, alpha" Esrra
1442865_at 0.97235 diacylglycerol kinase kappa Dgkk
1442866_at 1.08568 RCSD domain containing 1 Rcsd1
1442867_at 0.99580 --- ---
1442868_at 0.86600 RIKEN cDNA C130026L21 gene C130026L21Rik
1442869_at 0.86093 docking protein 7 Dok7
1442870_at 0.90331 "CDNA, clone:Y2G0105B03, strand:unspecified" ---
1442871_at 1.07317 "PREDICTED: Mus musculus similar to Nucleolar transcription factor 1
(Upstream binding factor 1) (UBF-1) (LOC628897), mRNA" ---
1442872_at 1.09441 cDNA sequence BC021438 BC021438
1442873_at 0.64779 "10 days neonate cerebellum cDNA, RIKEN full-length enriched library, clone:
B930049N01 product:unclassifiable, full insert sequence" ---
1442874_at 0.98919 Activating signal cointegrator 1 complex subunit 1 Ascc1
1442875_at 0.92171 --- ---
1442876_at 1.03462 Mitogen activated protein kinase 1 Mapk1
1442877_at 0.96475 --- ---
1442878_at 1.22366 peroxiredoxin 6 Prdx6
1442879_at 1.03822 --- ---
1442880_at 1.60960 --- ---
1442881_at 1.05884 --- ---
1442882_at 1.01684 --- ---
1442883_s_at 0.97590 "DNA segment, Chr 10, Brigham & Women's Genetics 1364 expressed"
D10Bwg1364e
1442884_at 1.10035 hepatocyte growth factor Hgf
1442885_at 0.93152 SREBP cleavage activating protein Scap
1442886_at 1.73199 transformer 2 alpha homolog (Drosophila) Tra2a
1442887_at 1.02487 expressed sequence AI853363 AI853363
1442888_at 1.07644 InaD-like (Drosophila) Inadl
1442889_at 0.95948 Transcribed locus ---
1442890_at 1.29315 Suppressor of cytokine signaling 5 Socs5
1442891_at 0.93404 expressed sequence AU023386 AU023386
1442892_at 1.47299 Mutated in colorectal cancers Mcc
1442893_at 0.99646 "Catenin (cadherin associated protein), alpha 2" Ctnna2

1442894_at 0.98213 "dynein, axonemal, heavy chain 6" Dnahc6
1442895_at 0.99393 "PREDICTED: Mus musculus similar to zinc finger protein 547 (LOC628043), mRNA" ---
1442896_at 0.94410 Heat shock protein 12A Hspa12a
1442897_at 1.30920 RIKEN cDNA 2610024E20 gene 2610024E20Rik
1442898_at 1.01097 hypothetical protein LOC671579 LOC671579
1442899_at 0.93429 Transcribed locus ---
1442900_at 1.01155 Transcribed locus ---
1442901_at 1.34777 Tax1 (human T-cell leukemia virus type I) binding protein 1 Tax1bp1
1442902_at 0.96034 glutamine rich 2 Qrich2
1442903_at 1.15133 RIKEN cDNA 4732423E21 gene 4732423E21Rik
1442904_at 0.91483 similar to RNA (guanine-9-) methyltransferase domain containing 2
LOC622118
1442905_at 0.98454 ADP-ribosylation factor 4 Arf4
1442906_at 0.99419 --- ---
1442907_at 1.07022 "Adult male aorta and vein cDNA, RIKEN full-length enriched library, clone: A530062L09 product:unclassifiable, full insert sequence" ---
1442908_at 0.93324 Transmembrane 9 superfamily member 1 Tm9sf1
1442909_at 0.96089 "Potassium voltage-gated channel, subfamily Q, member 4" Kcnq4
1442910_at 0.95206 Expressed sequence AL033314 AL033314
1442911_at 1.18234 RIO kinase 2 (yeast) Riok2
1442912_at 0.92280 "hydroxy-delta-5-steroid dehydrogenase, 3 beta- and steroid delta-isomerase 7" Hsd3b7
1442913_at 0.97191 "12 days embryo spinal ganglion cDNA, RIKEN full-length enriched library, clone:D130048N24 product:unclassifiable, full insert sequence" ---
1442914_at 0.87112 "Transcribed locus, weakly similar to XP_858458.1 PREDICTED: hypothetical protein XP_853365 [Canis familiaris]" ---
1442915_at 1.02217 "solute carrier family 9 (sodium/hydrogen exchanger), member 4" Slc9a4
1442916_at 2.11065 RIKEN cDNA 4931420C21 gene 4931420C21Rik
1442917_at 0.78085 "Adult male vesicular gland cDNA, RIKEN full-length enriched library, clone: G630023F10 product:unclassifiable, full insert sequence" ---
1442918_at 0.97225 neuron navigator 3 /// RIKEN cDNA 9630020C08 gene /// region containing RIKEN cDNA 9630020C08 gene; neuron navigator 3 Nav3 /// 9630020C08Rik /// LOC676640
1442919_at 1.23009 SH3 and PX domains 2B Sh3pxd2b
1442920_at 1.07004 Kruppel-like factor 3 (basic) Klf3
1442921_at 1.01092 Matrix metalloproteinase 24 Mmp24
1442922_at 0.97207 checkpoint with forkhead and ring finger domains Chfr
1442923_at 1.05620 PTK6 protein tyrosine kinase 6 Ptk6
1442924_at 0.98692 Calsyntenin 2 Clstn2
1442925_at 1.03190 RIKEN cDNA 2610207I05 gene 2610207I05Rik
1442926_at 1.05199 RIKEN cDNA 1700011B04 gene 1700011B04Rik
1442927_at 0.91496 PTK2 protein tyrosine kinase 2 beta Ptk2b
1442928_at 0.98384 Karyopherin (importin) alpha 4 Kpna4

1442929_at 0.89271 "ST8 alpha-N-acetyl-neuraminide alpha-2,8-sialyltransferase 5" St8sia5
1442930_at 0.89644 expressed sequence AW061147 AW061147
1442931_at 1.00687 RIKEN cDNA 2600003E23 gene 2600003E23Rik
1442932_at 0.98251 expressed sequence C79709 C79709
1442933_at 1.20587 forkhead box K2 Foxk2
1442934_at 1.09283 expressed sequence C81608 C81608
1442935_at 0.81034 expressed sequence C81452 C81452
1442936_at 1.01113 Desmoglein 3 Dsg3
1442937_at 1.03015 expressed sequence C77190 C77190
1442938_at 1.01417 "DNA segment, Chr 12, ERATO Doi 247, expressed" D12Ertd247e
1442939_at 1.03313 Rap1 interacting factor 1 homolog (yeast) /// similar to Telomere-associated protein RIF1 (Rap1-interacting factor 1 homolog) (mRif1) Rif1 /// LOC671598
1442940_at 1.02001 "Adult male corpora quadrigemina cDNA, RIKEN full-length enriched library, clone:B230110I22 product:unclassifiable, full insert sequence" ---
1442941_at 1.00019 expressed sequence C77027 C77027
1442942_at 0.93863 Williams-Beuren syndrome chromosome region 1 homolog (human) Wbscr1
1442943_at 0.95215 Ubiquitin specific peptidase 20 Usp20
1442944_at 1.29089 expressed sequence C76555 C76555
1442945_at 1.33728 RIKEN cDNA 2010109K09 gene 2010109K09Rik
1442946_at 1.30019 Ataxin 7 Atxn7
1442947_x_at 0.95130 "13 days embryo heart cDNA, RIKEN full-length enriched library, clone: D330042P15 product:unclassifiable, full insert sequence" ---
1442948_at 1.04776 --- ---
1442949_at 1.28560 "Nuclear factor of kappa light chain gene enhancer in B-cells 1, p105" Nfkb1
1442950_at 0.95427 Neuregulin 3 Nrg3
1442951_at 1.43783 Developmentally regulated repeat element-containing transcript 1 Drr1
1442952_at 1.02895 Shroom Shrm
1442953_at 0.97024 Transcribed locus ---
1442954_at 0.92431 Transcribed locus ---
1442955_at 1.04726 --- ---
1442956_at 1.05155 "Protein phosphatase 1, regulatory (inhibitor) subunit 13B" Ppp1r13b
1442957_at 0.95521 RIKEN cDNA G730013B05 gene G730013B05Rik
1442958_at 1.02722 "Lung RCB-0558 LLC cDNA, RIKEN full-length enriched library, clone: G730002L02 product:unclassifiable, full insert sequence" ---
1442959_at 2.23475 baculoviral IAP repeat-containing 6 Birc6
1442960_at 1.03026 mucin 20 Muc20
1442961_at 0.95425 --- ---
1442962_at 1.03113 expressed sequence AU022052 AU022052
1442963_at 1.08877 zinc fingers and homeoboxes 3 Zhx3
1442964_at 1.26753 Expressed sequence AU020772 AU020772
1442965_at 1.03332 --- ---
1442966_at 0.96172 RIKEN cDNA 5330421F07 gene 5330421F07Rik
1442967_at 1.11615 expressed sequence AU020147 AU020147

1442968_at	0.99270	"DNA segment, Chr 9, ERATO Doi 241, expressed"	D9Ertd241e
1442969_at	0.96428	---	---
1442970_at	0.99476	---	---
1442971_at	1.84839	"bromodomain adjacent to zinc finger domain, 2B"	Baz2b
1442972_at	1.65861	"Coronin, actin binding protein 1C"	Coro1c
1442973_at	0.92386	expressed sequence C80865	C80865
1442974_at	1.00498	"DNA segment, Chr 6, ERATO Doi 160, expressed"	D6Ertd160e
1442975_at	0.90858	Transcribed locus	---
1442976_at	0.86846	expressed sequence C81072	C81072
1442977_at	1.03034	"Procollagen, type III, alpha 1"	Col3a1
1442978_at	0.91884	Multiple PDZ domain protein	Mpdz
1442979_at	0.80599	"PREDICTED: Mus musculus similar to zinc finger protein 709 (LOC629191), mRNA"	---
1442980_at	1.00240	Serine/arginine-rich protein specific kinase 2	Srpk2
1442981_at	0.99812	---	---
1442982_at	0.97898	coiled-coil domain containing 66	Ccdc66
1442983_at	1.36652	"Solute carrier family 23 (nucleobase transporters), member 2"	Slc23a2
1442984_at	0.96026	expressed sequence C80060	C80060
1442985_at	1.04151	Hemicentin 1	Hmcn1
1442986_at	1.00177	"ELK3, member of ETS oncogene family"	Elk3
1442987_at	0.96862	---	---
1442988_at	1.04325	RIKEN cDNA A630001O12 gene	A630001O12Rik
1442989_at	1.10900	Asparaginyl-tRNA synthetase 2 (mitochondrial)(putative)	Nars2
1442990_at	1.00084	---	---
1442991_at	1.00901	"beta-1,3-glucuronyltransferase 2 (glucuronosyltransferase S)"	B3gat2
1442992_at	2.80315	hypothetical LOC403343	130004C03
1442993_at	1.11152	integrin beta 3 binding protein (beta3-endonexin)	Itgb3bp
1442994_at	0.96266	"Superoxide dismutase 2, mitochondrial"	Sod2
1442995_at	1.28338	N-deacetylase/N-sulfotransferase (heparan glucosaminyl) 1	Ndst1
1442996_x_at	1.09438	"0 day neonate kidney cDNA, RIKEN full-length enriched library, clone: D630027M10 product:unclassifiable, full insert sequence"	---
1442997_at	1.37006	Odd Oz/ten-m homolog 3 (Drosophila)	Odz3
1442998_at	0.95373	Cytidine 5'-triphosphate synthase 2	Ctps2
1442999_at	0.88102	RIKEN cDNA B930036G03 gene	B930036G03Rik
1443000_at	1.09569	"DNA segment, Chr 3, ERATO Doi 300, expressed"	D3Ertd300e
1443001_at	1.00585	expressed sequence C78344	C78344
1443002_at	1.69729	zinc finger RNA binding protein Zfr	
1443003_at	1.26660	---	---
1443004_at	1.01643	"Adult male liver tumor cDNA, RIKEN full-length enriched library, clone: C730045N16 product:unclassifiable, full insert sequence"	---
1443005_at	1.07094	zinc finger homeobox 1a	Zfhx1a
1443006_at	1.00169	Synapsin III	Syn3
1443007_at	1.05772	---	---

1443008_at 1.00217 Musashi homolog 2 (Drosophila) Msi2
1443009_at 1.02785 Transcribed locus ---
1443010_at 1.05099 F-box and WD-40 domain protein 11 Fbxw11
1443011_at 0.91714 protein arginine N-methyltransferase 7 Prmt7
1443012_at 1.01496 Transcription factor 12 Tcf12
1443013_at 1.04538 "A disintegrin-like and metalloproteinase (reprolysin type) with thrombospondin type 1 motif, 16" Adamts16
1443014_at 1.01279 RIKEN cDNA 4833403I15 gene 4833403I15Rik
1443015_at 0.88692 "Ras homolog gene family, member H" Rhoh
1443016_at 1.00830 DNA methyltransferase 3B Dnmt3b
1443017_at 0.95378 cytoplasmic polyadenylation element binding protein 2 Cpeb2
1443018_at 1.28519 Neural cell adhesion molecule 1 Ncam1
1443019_at 0.84676 --- ---
1443020_at 1.71927 Homeobox containing 1 Hmbox1
1443021_at 0.93138 "5, 10-methenyltetrahydrofolate synthetase" Mthfs
1443022_at 0.89750 RIKEN cDNA 4930521E07 gene 4930521E07Rik
1443023_at 1.06746 Transcribed locus ---
1443024_at 1.08711 "PREDICTED: Mus musculus hypothetical protein LOC623199 (LOC623199), mRNA" ---
1443025_at 1.07779 "10 days neonate medulla oblongata cDNA, RIKEN full-length enriched library, clone:B830002A04 product:unclassifiable, full insert sequence" ---
1443026_at 1.07190 Glutamate receptor interacting protein 1 Grip1
1443027_at 0.98854 RIKEN cDNA 4930523C07 gene 4930523C07Rik
1443028_at 0.98386 "La ribonucleoprotein domain family, member 6" Larp6
1443029_at 1.42457 "PTPRF interacting protein, binding protein 1 (liprin beta 1)" Ppfibp1
1443030_at 0.97553 l(3)mbt-like 3 (Drosophila) L3mbtl3
1443031_at 1.21335 "15 days embryo head cDNA, RIKEN full-length enriched library, clone: D930037E02 product:unclassifiable, full insert sequence" ---
1443032_at 1.05899 Mitochondrial ribosomal protein S22 Mrps22
1443033_at 1.00616 RNA binding motif protein 14 Rbm14
1443034_at 0.84383 Hermansky-Pudlak syndrome 1 homolog (human) Hps1
1443035_at 1.08310 --- ---
1443036_at 1.08337 RIKEN cDNA C630007C17 gene C630007C17Rik
1443037_at 2.19341 Neuroplastin Nptn
1443038_at 0.98134 --- ---
1443039_at 0.92074 --- ---
1443040_x_at 0.92728 --- ---
1443041_at 0.89212 zinc finger and BTB domain containing 17 Zbtb17
1443042_at 1.13755 Roundabout homolog 1 (Drosophila) Robo1
1443043_at 1.10712 otopetrin 2 Otop2
1443044_at 1.13944 hypothetical A830091E24 A830091E24
1443045_at 1.19197 Integrator complex subunit 7 Ints7
1443046_at 0.99595 "ATP-binding cassette, sub-family D (ALD), member 3" Abcd3

1443047_at 1.05676 PDZ domain containing RING finger 3 Pdzn3
1443048_at 1.10776 --- ---
1443049_at 1.21817 Transmembrane protein 19 Tmem19
1443050_at 0.96327 CDNA sequence BC032265 BC032265
1443051_at 0.92510 "10 days neonate cerebellum cDNA, RIKEN full-length enriched library, clone:
B930095F19 product:unclassifiable, full insert sequence" ---
1443052_at 0.93007 RIKEN cDNA C330019L16 gene C330019L16Rik
1443053_at 0.95787 "Protein tyrosine phosphatase, receptor type, D" Ptprd
1443054_at 1.02759 "potassium inwardly-rectifying channel, subfamily J, member 15" Kcnj15
1443055_at 0.89086 RIKEN cDNA B230369F24 gene B230369F24Rik
1443056_at 1.03000 "Protein tyrosine phosphatase, receptor type, D" Ptprd
1443057_at 0.92468 --- ---
1443058_at 0.86389 RIKEN cDNA 2900006F19 gene 2900006F19Rik
1443059_at 0.94889 dehydrogenase/reductase (SDR family) member 8 Dhrr8
1443060_at 1.02882 RIKEN cDNA 2810003K23 gene 2810003K23Rik
1443061_at 0.99776 Transcribed locus ---
1443062_at 1.00678 autism susceptibility candidate 2 Aut2
1443063_at 0.96670 RIKEN cDNA A330019N05 gene A330019N05Rik
1443064_at 1.02130 RIKEN cDNA 9130004C02 gene 9130004C02Rik
1443065_at 1.17818 Rhomboid domain containing 1 Rhbdd1
1443066_at 0.89291 --- ---
1443067_at 1.01596 Transcribed locus ---
1443068_at 1.58084 RIKEN cDNA D130084N16 gene D130084N16Rik
1443069_at 1.23848 "Protein kinase C, alpha" Prkca
1443070_at 1.02895 Autism susceptibility candidate 2 Aut2
1443071_at 0.98584 Expressed sequence AI839979 AI839979
1443072_at 1.15422 --- ---
1443073_at 1.06932 hypothetical protein LOC545681 LOC545681
1443074_at 1.34946 --- ---
1443075_at 0.89665 "Protein tyrosine phosphatase, receptor type, D" Ptprd
1443076_at 1.07037 RIKEN cDNA D030041N04 gene D030041N04Rik
1443077_at 1.24429 RIKEN cDNA 1700081L11 gene 1700081L11Rik
1443078_at 1.04930 RIKEN cDNA 6030439D06 gene 6030439D06Rik
1443079_at 0.92581 Transcribed locus ---
1443080_at 0.81425 Adenylate cyclase 5 Adcy5
1443081_at 0.96865 GATA binding protein 6 Gata6
1443082_at 1.00430 --- ---
1443083_at 1.01979 Cleavage and polyadenylation specific factor 2 Cpsf2
1443084_at 0.92848 cathepsin O Ctso
1443085_at 0.94106 "Splicing factor, arginine/serine-rich 2, interacting protein" Sfrs2ip
1443086_at 0.91772 activated leukocyte cell adhesion molecule Alcam
1443087_at 0.97662 "CDC23 (cell division cycle 23, yeast, homolog)" Cdc23
1443088_at 3.87249 RIKEN cDNA 9930031P18 gene 9930031P18Rik

1443089_at 0.90082 RIKEN cDNA 4930506C02 gene 4930506C02Rik
1443090_at 1.12733 Choline/ethanolaminephosphotransferase 1 Cept1
1443091_at 1.01006 --- ---
1443092_at 1.29667 myotubularin related protein 3 Mtmr3
1443093_at 1.03190 Adherens junction associated protein 1 Ajap1
1443094_at 0.96193 expressed sequence BB031773 BB031773
1443095_at 1.10661 --- ---
1443096_at 0.98394 RIKEN cDNA 4921526F01 gene 4921526F01Rik
1443097_at 1.24605 Neurofibromatosis 1 Nf1
1443098_at 0.93907 "Adult male corpora quadrigemina cDNA, RIKEN full-length enriched library, clone:B230340D10 product:unclassifiable, full insert sequence" ---
1443099_at 0.70093 CDNA sequence BC087945 BC087945
1443100_at 0.98465 Thyroid hormone receptor beta Thrb
1443101_at 0.92237 "Glutamate receptor, ionotropic, delta 2" Grid2
1443102_at 0.98588 zinc finger protein 597 Zfp597
1443103_at 1.05228 RIKEN cDNA D830046C22 gene D830046C22Rik
1443104_at 0.89964 --- ---
1443105_at 1.02513 Zinc finger protein 398 Zfp398
1443106_at 0.92959 RIKEN cDNA 4930562C15 gene 4930562C15Rik
1443107_at 1.90466 Transmembrane protein 16F Tmem16f
1443108_at 0.98592 "15 days embryo head cDNA, RIKEN full-length enriched library, clone: D930002E06 product:hypothetical protein, full insert sequence" ---
1443109_at 1.20624 RIKEN cDNA 5830417C01 gene 5830417C01Rik
1443110_at 0.86617 "general transcription factor II E, polypeptide 1 (alpha subunit)" Gtf2e1
1443111_at 0.97001 "pleckstrin homology domain containing, family A (phosphoinositide binding specific) member 1" Plekha1
1443112_at 0.98801 apoptosis inhibitor 5 Api5
1443113_at 0.92858 Jumonji domain containing 2B Jmjd2b
1443114_at 1.38240 RIKEN cDNA A730059M13 gene A730059M13Rik
1443115_at 0.89137 "Transforming growth factor, beta receptor II" Tgfbr2
1443116_at 1.35486 --- ---
1443117_at 1.32551 eyes absent 1 homolog (Drosophila) Eya1
1443118_at 0.91254 --- ---
1443119_at 1.07381 "glutamate receptor, metabotropic 7" Grm7
1443120_at 0.96613 PDZ domain containing 4 Pdzd4
1443121_at 0.97062 --- ---
1443122_at 1.02147 "15 days embryo head cDNA, RIKEN full-length enriched library, clone: D930037A07 product:unclassifiable, full insert sequence" ---
1443123_at 1.08030 "Tetratricopeptide repeat, ankyrin repeat and coiled-coil containing 2" Tanc2
1443124_at 1.15176 --- ---
1443125_at 1.21955 thyroid hormone receptor interactor 12 Trip12
1443126_at 1.01881 "Solute carrier family 12, member 1" Slc12a1
1443127_at 1.01066 RIKEN cDNA 9630021D06 gene 9630021D06Rik

1443128_at 1.55467 integrator complex subunit 6 Ints6
1443129_at 0.92052 --- ---
1443130_at 1.00343 hypothetical protein LOC628951 LOC628951
1443131_at 1.01062 "RAN, member RAS oncogene family" Ran
1443132_at 1.07672 hypothetical protein LOC544809 LOC544809
1443133_at 0.89153 RIKEN cDNA A230070E04 gene A230070E04Rik
1443134_at 0.95643 RIKEN cDNA A930005G22 gene A930005G22Rik
1443135_at 1.21235 component of oligomeric golgi complex 4 Cog4
1443136_at 0.96577 hypothetical gene supported by AK082906 LOC432459
1443137_at 0.97733 "0 day neonate lung cDNA, RIKEN full-length enriched library, clone:
E030034L15 product:unclassifiable, full insert sequence" ---
1443138_at 1.07063 "Sulfotransferase family 5A, member 1" Sult5a1
1443139_at 1.10373 --- ---
1443140_at 1.05789 hypothetical protein LOC627777 LOC627777
1443141_at 1.13836 --- ---
1443142_at 1.28688 Beta-transducin repeat containing protein Btrc
1443143_at 0.93233 fibronectin leucine rich transmembrane protein 3 Flrt1
1443144_at 1.21474 "protein kinase C, beta 1" Prkcb1
1443145_at 1.01060 "amyloid beta (A4) precursor protein-binding, family B, member 1 interacting
protein" Apbb1ip
1443146_at 1.97070 "ATPase family, AAA domain containing 2" Atad2
1443147_at 0.93915 Acyl-CoA thioesterase 3 Acot3
1443148_at 1.14245 huntingtin interacting protein 2 Hip2
1443149_at 1.16046 Nicotinamide nucleotide adenylyltransferase 3 Nmnat3
1443150_at 1.07962 Dynamin 2 Dnm2
1443151_at 1.05795 Synaptojanin 2 binding protein Synj2bp
1443152_at 1.11242 Pentatricopeptide repeat domain 3 Ptdc3
1443153_at 1.10442 Thyroid hormone receptor interactor 11 Trip11
1443154_at 1.00549 hypothetical protein 6030438J01 6030438J01
1443155_at 0.95586 --- ---
1443156_at 1.50906 --- ---
1443157_at 2.99367 GLIS family zinc finger 3 Glis3
1443158_at 1.21815 Sex comb on midleg homolog 1 Scmh1
1443159_at 2.00389 RIKEN cDNA 9130221J17 gene 9130221J17Rik
1443160_at 1.51706 SET binding factor 2 Sbf2
1443161_at 2.51331 Trichorhinophalangeal syndrome I (human) Trps1
1443162_at 1.17733 "Protein tyrosine phosphatase, non-receptor type 3" Ptpn3
1443163_at 1.06304 "solute carrier family 39 (zinc transporter), member 2" Slc39a2
1443164_at 1.01179 Vacuolar protein sorting 13D (yeast) Vps13d
1443165_at 1.02752 mitochondrial ribosomal protein S31 Mrps31
1443166_at 1.60311 CDNA sequence BC021395 BC021395
1443167_at 2.38766 Ring finger protein 12 Rnf12
1443168_at 0.98266 "Adult male colon cDNA, RIKEN full-length enriched library,

clone:9030003C19 product:unclassifiable, full insert sequence" ---

1443169_at 1.35779 "CCR4-NOT transcription complex, subunit 4" Cnot4

1443170_at 0.94710 --- ---

1443171_at 0.82938 --- ---

1443172_at 1.24368 "origin recognition complex, subunit 1-like (S.cereviaiae)" Orc11

1443173_at 0.97033 pleiotrophin Ptn

1443174_at 1.02631 Transmembrane and tetratricopeptide repeat containing 2 Tmtc2

1443175_at 0.91588 RIKEN cDNA A830010M09 gene A830010M09Rik

1443176_at 0.86838 Scm-like with four mbt domains 2 Sfmbt2

1443177_at 0.92281 Transcribed locus ---

1443178_at 0.90562 ring finger protein 13 Rnf13

1443179_at 0.94331 CUB and Sushi multiple domains 1 Csm1

1443180_at 1.40850 ERGIC and golgi 2 Ergic2

1443181_at 0.98426 "12 days embryo spinal ganglion cDNA, RIKEN full-length enriched library, clone:D130046G19 product:inferred: RNP particle component {Mus musculus}, full insert sequence" ---

1443182_at 1.09523 Expressed sequence AW060207 AW060207

1443183_at 1.12012 "HECT, UBA and WWE domain containing 1" Huwe1

1443184_at 0.96584 CDC14 cell division cycle 14 homolog A (S. cerevisiae) Cdc14a

1443185_at 1.06730 lipoma HMGIC fusion partner-like 2 Lhfp12

1443186_at 1.10004 RIKEN cDNA 4930425N13 gene 4930425N13Rik

1443187_at 0.87893 --- ---

1443188_at 1.78021 ubiquitin-conjugating enzyme E2W (putative) Ube2w

1443189_at 0.99788 --- ---

1443190_at 0.91554 "15 days embryo head cDNA, RIKEN full-length enriched library, clone: D930036L24 product:unclassifiable, full insert sequence" ---

1443191_at 0.89587 --- ---

1443192_at 0.88614 --- ---

1443193_at 0.94799 tubulin tyrosine ligase-like 1 Ttll1

1443194_at 0.97152 --- ---

1443195_at 0.96262 --- ---

1443196_at 0.79976 --- ---

1443197_at 1.01745 wingless-type MMTV integration site 9B Wnt9b

1443198_at 0.91725 "1 cell embryo 1 cell cDNA, RIKEN full-length enriched library, clone: IOC0049K08 product:unclassifiable, full insert sequence" ---

1443199_at 1.12376 Leucine-rich repeats and calponin homology (CH) domain containing 3 Lrch3

1443200_at 0.97375 hypothetical protein A930011G24 A930011G24

1443201_at 1.18674 Glypican 6 Gpc6

1443202_at 1.40482 Cytochrome b5 type B Cyb5b

1443203_at 1.00981 Zinc finger homeobox 1a Zfhx1a

1443204_at 0.94337 "RNA binding motif, single stranded interacting protein" Rbms3

1443205_at 0.99861 "DNA segment, Chr 5, Bucan 30 expressed" D5Buc30e

1443206_at 1.14420 "ATP-binding cassette, sub-family C (CFTR/MRP), member 3" Abcc3

1443207_at 1.10036 "Adult male diencephalon cDNA, RIKEN full-length enriched library,

clone:9330196A19 product:unclassifiable, full insert sequence" ---
1443208_at 1.57625 G protein-coupled receptor 137B Gpr137b
1443209_at 1.05549 "Histone 1, H1e" Hist1h1e
1443210_at 1.05427 --- ---
1443211_at 0.89690 --- ---
1443212_at 1.90877 Like-glycosyltransferase Large
1443213_at 1.07767 glycosyltransferase-like domain containing 1 Gtdc1
1443214_at 0.98426 G protein-coupled receptor 155 Gpr155
1443215_at 1.06141 regulating synaptic membrane exocytosis 2 Rims2
1443216_at 1.20406 "TBC1 domain family, member 5" Tbc1d5
1443217_at 0.94386 "Adult male epididymis cDNA, RIKEN full-length enriched library,
clone:9230118O19 product:unclassifiable, full insert sequence" ---
1443218_at 0.99007 Phosphatidylinositol binding clathrin assembly protein Picalm
1443219_at 1.03684 --- ---
1443220_at 1.20759 Reticulon 3 Rtn3
1443221_at 1.01191 Wilms tumor homolog Wt1
1443222_at 1.03054 Thymoma viral proto-oncogene 3 Akt3
1443223_at 0.84067 --- ---
1443224_at 0.97465 ectonucleotide pyrophosphatase/phosphodiesterase 2 Enpp2
1443225_at 1.04796 "activin A receptor, type IC" Acvr1c
1443226_at 1.25416 RIKEN cDNA 5730470L24 gene 5730470L24Rik
1443227_at 1.05789 basic leucine zipper and W2 domains 2 Bzw2
1443228_at 1.10942 --- ---
1443229_at 1.22810 "ATPase family, AAA domain containing 2" Atad2
1443230_at 1.27106 RIKEN cDNA 5330421F07 gene 5330421F07Rik
1443231_at 2.96807 "Membrane associated guanylate kinase, WW and PDZ domain containing 1"
Magi1
1443232_at 0.94823 Transcribed locus ---
1443233_at 1.53229 "General transcription factor IIF, polypeptide 2" Gtf2f2
1443234_at 1.03044 CDNA sequence BC088983 BC088983
1443235_at 1.18797 Eukaryotic translation initiation factor 2 alpha kinase 4 Eif2ak4
1443236_at 1.15721 RIKEN cDNA B230118H07 gene B230118H07Rik
1443237_at 1.04497 --- ---
1443238_at 1.07132 PHD finger protein 14 Phf14
1443239_at 1.07357 Microtubule-associated protein 2 Mtap2
1443240_at 0.96908 Glypican 3 Gpc3
1443241_at 1.12121 Pseudouridylate synthase 7 homolog (S. cerevisiae) Pus7
1443242_at 1.28133 "DNA segment, Chr 5, ERATO Doi 121, expressed" D5Ert121e
1443243_at 0.98156 --- ---
1443244_at 0.98567 "zinc binding alcohol dehydrogenase, domain containing 1" Zadh1
1443245_at 1.01673 --- ---
1443246_at 1.03652 FAST kinase domains 2 Fastkd2
1443247_at 1.11956 Glucose-fructose oxidoreductase domain containing 1 Gfod1

1443248_at 1.14265 transmembrane and coiled-coil domains 3 Tmco3
1443249_at 0.97056 Membrane-associated ring finger (C3HC4) 3 3-Mar
1443250_at 1.12198 --- ---
1443251_at 0.89180 PR domain containing 16 Prdm16
1443252_at 0.92793 IBR domain containing 2 Ibrdc2
1443253_at 2.68078 GLIS family zinc finger 3 Glis3
1443254_at 1.07378 RIKEN cDNA D030011O10 gene D030011O10Rik
1443255_s_at 1.00085 Cdc42 GTPase-activating protein Cdgap
1443256_at 0.92390 Chloride channel calcium activated 6 Clca6
1443257_at 1.16889 RIKEN cDNA 9630050E16 gene 9630050E16Rik
1443258_at 1.45312 forkhead box P1 Foxp1
1443259_at 1.16690 Transcribed locus ---
1443260_at 1.04565 myeloid ecotropic viral integration site 1 Meis1
1443261_at 0.92657 RIKEN cDNA E130307A14 gene E130307A14Rik
1443262_at 1.02813 mitochondrial ribosomal protein S14 Mrps14
1443263_at 1.03210 BTB and CNC homology 2 Bach2
1443264_at 0.89920 "membrane-spanning 4-domains, subfamily A, member 2" Ms4a2
1443265_at 0.98831 CD44 antigen Cd44
1443266_at 1.25141 Niemann Pick type C2 Npc2
1443267_at 1.04722 "Tetratricopeptide repeat, ankyrin repeat and coiled-coil containing 2" Tanc2
1443268_at 0.94389 Transcribed locus ---
1443269_at 0.95302 RIKEN cDNA D130009B15 gene D130009B15Rik
1443270_at 1.12617 "Adult male hypothalamus cDNA, RIKEN full-length enriched library, clone:
A230020K01 product:unclassifiable, full insert sequence" ---
1443271_at 0.94710 Transmembrane protein 28 Tmem28
1443272_at 0.91784 neighbor of Punc E11 Nope
1443273_at 0.90574 Eph receptor A3 Epha3
1443274_at 0.97261 RIKEN cDNA 4932435O22 gene 4932435O22Rik
1443275_at 1.13942 Cdc42 GTPase-activating protein Cdgap
1443276_at 0.99031 5-hydroxytryptamine (serotonin) receptor 5A Htr5a
1443277_at 1.03679 N-deacetylase/N-sulfotransferase (heparan glucosaminyl) 3 Ndst3
1443278_at 1.03769 Deleted in colorectal carcinoma Dcc
1443279_at 1.14754 Nemo like kinase Nlk
1443280_at 0.99088 --- ---
1443281_at 0.98518 RIKEN cDNA C130076O07 gene C130076O07Rik
1443282_at 1.00559 PRP38 pre-mRNA processing factor 38 (yeast) domain containing A Prpf38a
1443283_at 1.10041 RIKEN cDNA C330002D13 gene C330002D13Rik
1443284_at 0.85644 Special AT-rich sequence binding protein 1 Satb1
1443285_at 1.09037 "Glutamate receptor, ionotropic, AMPA4 (alpha 4)" Gria4
1443286_at 1.05292 expressed sequence AU019754 AU019754
1443287_at 0.94485 "gene model 1337, (NCBI)" Gm1337
1443288_at 1.01240 Tnf receptor-associated factor 6 Traf6
1443289_at 1.02254 "Phosphodiesterase 4D, cAMP specific" Pde4d

1443290_at 0.92641 "Adult male medulla oblongata cDNA, RIKEN full-length enriched library, clone:6330436D20 product:unclassifiable, full insert sequence" ---

1443291_at 1.01433 "Ankyrin 1, erythroid" Ank1

1443292_at 0.99124 Transcribed locus ---

1443293_at 1.68391 --- ---

1443294_at 1.20493 "Cdc2-related kinase, arginine/serine-rich" Crkrs

1443295_at 0.94530 Transcribed locus ---

1443296_at 1.06433 PCTAIRE-motif protein kinase 1 Pctk1

1443297_at 1.23992 "TBC1D12: TBC1 domain family, member 12" Tbc1d12

1443298_at 1.04047 F-box and WD-40 domain protein 13 Fbxw13

1443299_at 1.01367 PDZ and LIM domain 3 Pdlim3

1443300_at 0.89688 Notch gene homolog 3 (Drosophila) Notch3

1443301_at 1.09366 RIKEN cDNA C230086A09 gene C230086A09Rik

1443302_at 1.16422 RIKEN cDNA 6720403M19 gene 6720403M19Rik

1443303_at 1.16957 RIKEN cDNA 4930521E07 gene 4930521E07Rik

1443304_at 1.03837 "metallothionein-like 5, testis-specific (tesmin)" Mtl5

1443305_at 1.01436 "13 days embryo heart cDNA, RIKEN full-length enriched library, clone: D330014J04 product:unclassifiable, full insert sequence" ---

1443306_at 1.08909 "0 day neonate cerebellum cDNA, RIKEN full-length enriched library, clone: C230004E12 product:hypothetical protein, full insert sequence" ---

1443307_at 1.41492 "Proteasome (prosome, macropain) 26S subunit, ATPase 2" Psmc2

1443308_at 0.99693 unc-13 homolog B (C. elegans) Unc13b

1443309_at 1.20330 PDZ domain containing RING finger 3 Pdzn3

1443310_at 0.92841 nucleoporin 210 Nup210

1443311_at 0.78745 THAP domain containing 4 Thap4

1443312_at 0.99664 "Regulatory factor X, 4 (influences HLA class II expression)" Rfx4

1443313_at 0.93309 RIKEN cDNA A530058N18 gene A530058N18Rik

1443314_at 2.15156 RIKEN cDNA 2410042D21 gene 2410042D21Rik

1443315_at 0.88943 --- ---

1443316_at 0.99892 Transcribed locus ---

1443317_at 1.09760 "DNA segment, Chr 6, ERATO Doi 47, expressed" D6Ertd47e

1443318_at 1.11158 RIKEN cDNA E130016E03 gene E130016E03Rik

1443319_at 0.95915 phosphoprotein associated with glycosphingolipid microdomains 1 Pag1

1443320_at 0.88363 CLIP associating protein 2 Clasp2

1443321_at 1.03857 Transmembrane and tetratricopeptide repeat containing 2 Tmtc2

1443322_at 0.99471 Transcribed locus ---

1443323_at 1.01738 RIKEN cDNA A930028L21 gene A930028L21Rik

1443324_at 1.02641 "Transcribed locus, moderately similar to XP_515274.1 PREDICTED: similar to FLJ00254 protein [Pan troglodytes]" ---

1443325_at 1.35804 Transcribed locus ---

1443326_at 1.06163 --- ---

1443327_at 1.14526 RIKEN cDNA D130043K22 gene D130043K22Rik

1443328_at 0.98908 RIKEN cDNA 4921505C17 gene 4921505C17Rik

1443329_at 1.09781 membrane-associated ring finger (C3HC4) 7 7-Mar
1443330_at 1.01123 "DNA segment, Chr 1, ERATO Doi 646, expressed" D1Ert646e
1443331_at 0.95862 "Pleckstrin homology domain containing, family G (with RhoGef domain)
member 1" Plekhg1
1443332_at 1.22795 "Solute carrier family 12, member 2" Slc12a2
1443333_at 1.02226 "Adult male spinal cord cDNA, RIKEN full-length enriched library, clone:
A330092A19 product:unclassifiable, full insert sequence" ---
1443334_at 1.08907 RIKEN cDNA D430042O09 gene D430042O09Rik
1443335_at 1.30210 PDZ domain containing RING finger 3 Pdzn3
1443336_at 1.01020 F-box and leucine-rich repeat protein 11 Fbxl11
1443337_at 3.08022 Glutamate receptor interacting protein 1 Grip1
1443338_at 1.60460 "Adaptor protein complex AP-2, mu1" Ap2m1
1443339_at 1.33318 RIKEN cDNA 2310056P07 gene 2310056P07Rik
1443340_at 1.09789 calcium binding atopy-related autoantigen 1 Cbara1
1443341_at 2.22127 Annexin A4 Anxa4
1443342_at 1.04768 Transcribed locus ---
1443343_at 1.00893 --- ---
1443344_at 0.98244 RIKEN cDNA 4932409I22 gene 4932409I22Rik
1443345_at 0.95275 RIKEN cDNA 1700047E16 gene 1700047E16Rik
1443346_at 1.06688 --- ---
1443347_at 1.08688 --- ---
1443348_at 1.07263 RIKEN cDNA D030028M11 gene D030028M11Rik
1443349_at 0.97575 Deducator of cytokinesis 10 Dock10
1443350_at 0.93291 metastasis associated 3 Mta3
1443351_at 0.93058 "Glycine receptor, alpha 2 subunit" Glra2
1443352_at 1.09517 EST C77614 C77614
1443353_at 1.26545 Zinc finger RNA binding protein Zfr
1443354_at 1.41318 Tripartite motif-containing 59 Trim59
1443355_at 1.84830 --- ---
1443356_at 0.94724 "Cyclic AMP-regulated phosphoprotein, 21" Arpp21
1443357_at 0.96007 Zinc finger RNA binding protein Zfr
1443358_at 0.86641 Ribonuclease P 14 subunit (human) Rpp14
1443359_at 0.97862 Odd Oz/ten-m homolog 4 (Drosophila) Odz4
1443360_x_at 0.91996 Rho GTPase activating protein 4 Arhgap4
1443361_at 1.80912 Tetraspanin 9 Tspan9
1443362_at 1.27184 zinc finger protein 277 Zfp277
1443363_at 0.89442 RIKEN cDNA A430107O13 gene A430107O13Rik
1443364_at 1.05008 Transcribed locus ---
1443365_at 1.06848 5 hydroxytryptamine (serotonin) receptor 4 Htr4
1443366_at 1.23387 N-ethylmaleimide sensitive fusion protein attachment protein gamma Napg
1443367_at 0.94834 --- ---
1443368_at 0.99493 "2 cells egg cDNA, RIKEN full-length enriched library, clone:B020017L01
product:unclassifiable, full insert sequence" ---

1443369_at 1.07202 cDNA sequence BC052055 BC052055
1443370_at 1.02389 Activating signal cointegrator 1 complex subunit 3 Ascc3
1443371_at 0.95881 RIKEN cDNA 4930473A06 gene 4930473A06Rik
1443372_at 0.97177 "12 days embryo spinal cord cDNA, RIKEN full-length enriched library, clone: C530005L06 product:unclassifiable, full insert sequence" ---
1443373_at 1.03390 Mitogen-activated protein kinase kinase kinase kinase 5 Map4k5
1443374_at 0.89362 Transcribed locus ---
1443375_at 0.99189 "GNAS (guanine nucleotide binding protein, alpha stimulating) complex locus" Gnas
1443376_at 0.83008 Transcribed locus ---
1443377_at 0.95834 MAP kinase-activated protein kinase 5 /// a disintegrin and metallopeptidase domain 1a Mapkapk5 /// Adam1a
1443378_s_at 1.03544 MAP kinase-activated protein kinase 5 /// a disintegrin and metallopeptidase domain 1a Mapkapk5 /// Adam1a
1443379_at 0.84292 RIKEN cDNA A430033K04 gene A430033K04Rik
1443380_at 0.95451 "alcohol dehydrogenase 5 (class III), chi polypeptide" Adh5
1443381_at 1.31673 "ets variant gene 4 (E1A enhancer binding protein, E1AF)" Etv4
1443382_s_at 1.30729 BTB (POZ) domain containing 14B Btbd14b
1443383_at 1.03738 Muscleblind-like 3 (Drosophila) Mbnl3
1443384_at 1.64959 PTK2 protein tyrosine kinase 2 Ptk2
1443385_at 1.01477 RIKEN cDNA A930012N16 gene A930012N16Rik
1443386_at 1.03600 expressed sequence C86727 C86727
1443387_at 0.98354 expressed sequence BB045044 BB045044
1443388_at 1.33853 RIKEN cDNA 1300018I05 gene 1300018I05Rik
1443389_at 0.97249 EGL nine homolog 1 (C. elegans) Egl1
1443390_at 2.34275 Dedicator of cyto-kinesis 1 Dock1
1443391_at 0.85915 "5'-nucleotidase, cytosolic II" Nt5c2
1443392_at 1.02731 "Transient receptor potential cation channel, subfamily V, member 1" Trpv1
1443393_at 2.10114 SMAD specific E3 ubiquitin protein ligase 1 Smurf1
1443394_at 1.64906 High mobility group AT-hook 2 Hmga2
1443395_at 1.00733 "DNA segment, Chr 6, ERATO Doi 490, expressed" D6Ertd490e
1443396_at 1.16974 survival motor neuron domain containing 1 /// similar to survival motor neuron domain containing 1 Smndc1 /// LOC545792
1443397_at 1.01859 nuclear receptor coactivator 1 Ncoa1
1443398_at 0.97491 Synaptosomal-associated protein 25 Snap25
1443399_at 1.09782 expressed sequence AU021001 AU021001
1443400_at 1.37792 Mesoderm induction early response 1 homolog (Xenopus laevis) Mier1
1443401_at 1.01601 Astrotactin 2 Astn2
1443402_at 1.21055 Phosphorylase kinase alpha 2 Phka2
1443403_at 1.05956 RIKEN cDNA 4933428G09 gene 4933428G09Rik
1443404_s_at 0.93822 "DNA segment, Chr 13, ERATO Doi 94, expressed" D13Ertd94e
1443405_at 0.99849 "9 days embryo whole body cDNA, RIKEN full-length enriched library, clone: D030059B04 product:unclassifiable, full insert sequence" ---

1443406_at 1.04368 phospholipid scramblase 4 Plscr4
1443407_at 1.26698 ring finger protein 13 Rnf13
1443408_at 0.91094 Polo-like kinase 1 (Drosophila) Plk1
1443409_at 0.92239 expressed sequence C76434 C76434
1443410_at 1.07237 "Glutamate receptor, metabotropic 5" Grm5
1443411_at 0.93162 Spermatogenesis associated 9 Spata9
1443412_s_at 1.00119 matrix metalloproteinase 16 Mmp16
1443413_s_at 1.00046 "transient receptor potential cation channel, subfamily M, member 5" Trpm5
1443414_at 0.89740 EST C78513 C78513
1443415_at 0.96736 RIKEN cDNA 4931432P07 gene 4931432P07Rik
1443416_at 0.82020 expressed sequence C79741 C79741
1443417_at 1.00080 expressed sequence C87122 C87122
1443418_at 0.98270 Phosphoribosyl pyrophosphate synthetase 2 Prps2
1443419_at 0.93539 "DNA segment, Chr 5, ERATO Doi 525, expressed" D5Ertd525e
1443420_at 0.96158 Ubiquitin specific peptidase 8 Usp8
1443421_s_at 0.96428 protocadherin beta 15 Pcdhb15
1443422_at 1.25313 RIKEN cDNA 2410089E03 gene /// hypothetical LOC546032 /// hypothetical protein LOC669465 2410089E03Rik /// LOC546032 /// LOC669465
1443423_at 0.88490 WNK lysine deficient protein kinase 2 /// hypothetical LOC624678 Wnk2 /// LOC624678
1443424_at 1.08120 RIKEN cDNA 2210018M11 gene 2210018M11Rik
1443425_at 0.94315 "A disintegrin-like and metalloproteinase (reprolysin type) with thrombospondin type 1 motif, 6" Adamts6
1443426_at 1.31975 "SAFB-like, transcription modulator" Sltm
1443427_at 0.93698 --- ---
1443428_at 1.02594 --- ---
1443429_at 1.04909 RIKEN cDNA E130310K16 gene E130310K16Rik
1443430_at 1.41299 CASP2 and RIPK1 domain containing adaptor with death domain Cradd
1443431_at 1.04800 --- ---
1443432_at 0.90328 expressed sequence AU022680 AU022680
1443433_at 0.92089 plexin C1 Plxnc1
1443434_s_at 0.95275 plexin C1 Plxnc1
1443435_at 1.03039 RIKEN cDNA 3732413I11 gene 3732413I11Rik
1443436_at 1.08612 Mitogen activated protein kinase kinase 2 Map2k2
1443437_at 0.96063 similar to apobec-1 complementation factor LOC633285
1443438_at 1.09836 Nuclear receptor coactivator 1 Ncoa1
1443439_at 1.07452 Acyl-CoA synthetase short-chain family member 1 Acss1
1443440_at 1.07859 --- ---
1443441_x_at 0.82892 similar to hypothetical protein FLJ14345 LOC380850
1443442_at 0.95414 expressed sequence AU022084 AU022084
1443443_at 0.94879 --- ---
1443444_at 0.97562 Mannoside acetylglucosaminyltransferase 5 Mgat5
1443445_at 1.43009 Diaphanous homolog 3 (Drosophila) Diap3

1443446_at 0.98141 --- ---
1443447_at 1.01369 RIKEN cDNA 2010110K16 gene 2010110K16Rik
1443448_x_at 0.97576 RIKEN cDNA 2010110K16 gene 2010110K16Rik
1443449_at 0.98115 --- ---
1443450_at 1.03028 "7 days neonate cerebellum cDNA, RIKEN full-length enriched library, clone:
A730086M21 product:unclassifiable, full insert sequence" ---
1443451_at 1.08029 --- ---
1443452_at 1.03002 Elongation protein 3 homolog (S. cerevisiae) Elp3
1443453_at 0.89944 zinc finger protein 651 Zfp651
1443454_at 0.89262 secretin receptor Sctr
1443455_at 1.36831 CUE domain containing 1 Cuedc1
1443456_at 1.11067 RIKEN cDNA 6330416L07 gene 6330416L07Rik
1443457_at 1.10322 RIKEN cDNA A230055J12 gene A230055J12Rik
1443458_at 1.11514 RIKEN cDNA D630033O11 gene D630033O11Rik
1443459_at 1.06236 expressed sequence C86415 C86415
1443460_at 1.12658 --- ---
1443461_at 1.08234 expressed sequence C79130 C79130
1443462_at 1.03598 "DNA segment, Chr 8, ERATO Doi 572, expressed" D8Ertd572e
1443463_at 1.07328 Wingless-related MMTV integration site 5B Wnt5b
1443464_at 0.96034 "Syntrophin, basic 1" Sntb1
1443465_at 0.91063 --- ---
1443466_s_at 1.07537 polymerase (RNA) III (DNA directed) polypeptide B Polr3b
1443467_at 1.57510 dedicator of cytokinesis 11 Dock11
1443468_at 0.90963 --- ---
1443469_at 0.93022 Transcribed locus ---
1443470_at 0.98266 RIKEN cDNA A530058N18 gene A530058N18Rik
1443471_at 1.20635 zinc finger and BTB domain containing 20 Zbtb20
1443472_at 1.20052 expressed sequence C77691 C77691
1443473_at 0.93785 expressed sequence C79562 C79562
1443474_at 0.97887 "DNA segment, Chr 7, ERATO Doi 495, expressed" D7Ertd495e
1443475_at 0.93425 "Histone 1, H3i" Hist1h3i
1443476_at 0.75448 "3 days neonate thymus cDNA, RIKEN full-length enriched library, clone:
A630019K16 product:paired box gene 1, full insert sequence" ---
1443477_at 1.00009 "proteasome (prosome, macropain) 26S subunit, non-ATPase, 2 /// similar to
proteasome 26S non-ATPase subunit 2 /// similar to proteasome 26S non-ATPase subunit 2" Psmd2 ///
LOC432448 /// LOC672350
1443478_at 0.97843 expressed sequence C86695 C86695
1443479_at 0.88581 "7 days neonate cerebellum cDNA, RIKEN full-length enriched library, clone:
A730096N24 product:unclassifiable, full insert sequence" ---
1443480_at 1.06115 --- ---
1443481_at 1.29383 epsilon-tubulin 1 Tube1
1443482_at 0.94926 RIKEN cDNA 1700124P09 gene 1700124P09Rik
1443483_at 1.05429 X-linked lymphocyte-regulated 5D Xlr5d

1443484_at 0.97820 RNA polymerase II associated protein 1 Rpap1
1443485_at 0.85856 Eph receptor A7 Epha7
1443486_at 0.96679 Component of oligomeric golgi complex 7 Cog7
1443487_at 0.95688 expressed sequence AU022238 AU022238
1443488_at 1.06455 --- ---
1443489_at 1.53694 RIKEN cDNA C330002D13 gene C330002D13Rik
1443490_at 1.56825 "DNA segment, Chr 3, ERATO Doi 34, expressed" D3Ertd34e
1443491_at 1.39348 "Protein tyrosine phosphatase, receptor type, K" Ptpkr
1443492_at 1.02305 "Inositol 1,4,5-triphosphate receptor 1" Itpr1
1443493_at 1.02924 DEAH (Asp-Glu-Ala-His) box polypeptide 37 Dhx37
1443494_at 0.93964 polymerase (RNA) III (DNA directed) polypeptide B Polr3b
1443495_at 1.11578 Zinc finger protein 99 Zfp99
1443496_at 1.05573 ELMO domain containing 1 Elmod1
1443497_at 0.97539 Ankyrin repeat and BTB (POZ) domain containing 2 Abtb2
1443498_at 1.27541 methylmalonic aciduria (cobalamin deficiency) type B homolog (human)
Mmab
1443499_at 1.05905 hypothetical protein A430075N02 A430075N02
1443500_at 1.14543 Myeloid/lymphoid or mixed lineage-leukemia translocation to 10 homolog
(Drosophila) Mllt10
1443501_at 1.05837 "Transcribed locus, moderately similar to XP_345375.2 PREDICTED: similar
to Homeobox protein Hox-D8 (Hox-4.3) (Hox-5.4) [Rattus norvegicus]" ---
1443502_at 0.96734 baculoviral IAP repeat-containing 7 (livin) Birc7
1443503_at 1.08443 inositol hexaphosphate kinase 3 Ihpk3
1443504_at 1.06425 hypothetical protein E330022O07 E330022O07
1443505_at 0.96626 Prokineticin 1 Prok1
1443506_at 1.01680 potassium channel tetramerisation domain containing 2 Kctd2
1443507_at 1.06696 "Solute carrier family 4 (anion exchanger), member 1, adaptor protein"
Slc4a1ap
1443508_at 1.03736 "discs, large (Drosophila) homolog-associated protein 1" Dlgap1
1443509_at 0.97601 Cadherin 1 Cdh1
1443510_at 1.00515 expressed sequence AW822216 AW822216
1443511_at 0.80408 RAR-related orphan receptor alpha Rora
1443512_at 1.59812 Spermatogenesis associated 5 Spata5
1443513_at 1.53210 Transcribed locus ---
1443514_at 0.96265 --- ---
1443515_at 1.03431 RIKEN cDNA 1110032F04 gene 1110032F04Rik
1443516_at 2.45120 Ataxin 2 Atxn2
1443517_at 1.33417 RIKEN cDNA 6030443O07 gene 6030443O07Rik
1443518_at 0.89914 "sarcoglycan, epsilon" Sgce
1443519_at 1.07547 --- ---
1443520_at 0.97661 Expressed sequence AU022870 AU022870
1443521_at 2.27607 Gametogenetin binding protein 2 Ggnbp2
1443522_s_at 1.11361 pleckstrin homology domain interacting protein Phip

1443523_at 0.96745 --- ---
1443524_x_at 1.01510 B-cell leukemia/lymphoma 10 Bcl10
1443525_at 1.00039 Developing brain homeobox 2 Dbx2
1443526_at 2.94218 --- ---
1443527_at 1.40789 telomeric repeat binding factor 1 Terf1
1443528_at 1.01092 tau tubulin kinase 1 Ttbk1
1443529_at 0.99921 RUN and TBC1 domain containing 2 Rutbc2
1443530_at 0.92955 Musashi homolog 2 (Drosophila) Msi2
1443531_at 1.08866 RIKEN cDNA A930005I04 gene A930005I04Rik
1443532_at 1.13530 --- ---
1443533_at 1.99156 "Protein phosphatase 2, regulatory subunit B (B56), epsilon isoform" Ppp2r5e
1443534_at 1.41138 --- ---
1443535_at 2.61502 RAB GTPase activating protein 1 Rabgap1
1443536_at 1.01998 "solute carrier family 7 (cationic amino acid transporter, y+ system), member 11" Slc7a11
1443537_at 1.12206 RIKEN cDNA C130009A20 gene C130009A20Rik
1443538_at 0.94318 --- ---
1443539_at 0.83799 Rho guanine nucleotide exchange factor (GEF) 11 Arhgef11
1443540_at 1.27426 mitogen activated protein kinase kinase kinase 1 /// similar to Mitogen-activated protein kinase kinase kinase 1 (MAPK/ERK kinase kinase 1) (MEK kinase 1) (MEKK 1) /// similar to Mitogen-activated protein kinase kinase kinase 1 (MAPK/ERK kinase kinase 1) (MEK kinase 1) (MEKK 1) Map3k1 /// LOC670493 /// LOC674573
1443541_at 0.83617 Hypothetical gene supported by AK039231; AK039519; AK039710; AK045832 LOC381438
1443542_at 0.97963 --- ---
1443543_at 0.96172 "Adult male olfactory brain cDNA, RIKEN full-length enriched library, clone:6430501N10 product:unclassifiable, full insert sequence" ---
1443544_at 1.11912 "DNA segment, Chr 18, ERATO Doi 653, expressed" D18Ertd653e
1443545_at 1.07043 "CUG triplet repeat, RNA binding protein 2" Cugbp2
1443546_at 1.36311 High density lipoprotein (HDL) binding protein Hdlbp
1443547_at 1.12567 --- ---
1443548_at 0.96184 "Ankyrin 2, brain" Ank2
1443549_at 1.10552 RIKEN cDNA 4930579E17 gene 4930579E17Rik
1443550_at 0.93181 ClpB caseinolytic peptidase B homolog (E. coli) Clpb
1443551_at 1.23704 "ATPase, Ca++ transporting, cardiac muscle, slow twitch 2" Atp2a2
1443552_at 0.88682 RIKEN cDNA E230008N13 gene E230008N13Rik
1443553_at 1.03553 --- ---
1443554_at 1.04438 Single-stranded DNA binding protein 3 Ssbp3
1443555_at 1.09025 --- ---
1443556_at 0.96574 RIKEN cDNA 1110007F12 gene 1110007F12Rik
1443557_at 1.45095 --- ---
1443558_s_at 1.03323 RIKEN cDNA C630002B14 gene C630002B14Rik
1443559_at 1.06253 RIKEN cDNA 1500011J06 gene 1500011J06Rik

1443560_at 0.94036 "Adult male testis cDNA, RIKEN full-length enriched library, clone:4930506M03 product:hypothetical protein, full insert sequence" ---
1443561_at 1.01066 Myeloid ecotropic viral integration site-related gene 1 Mrg1
1443562_at 1.03277 RIKEN cDNA 2610017I09 gene 2610017I09Rik
1443563_at 1.03688 --- ---
1443564_at 0.86062 --- ---
1443565_at 1.01566 expressed sequence AI849538 AI849538
1443566_at 1.15881 Insulin-like growth factor I receptor Igf1r
1443567_at 1.06137 cDNA sequence BC032203 BC032203
1443568_x_at 0.92966 --- ---
1443569_at 1.01839 RIKEN cDNA 4930430E16 gene 4930430E16Rik
1443570_at 1.03224 "COP9 (constitutive photomorphogenic) homolog, subunit 3 (Arabidopsis thaliana)" Cops3
1443571_at 1.00577 Transcribed locus ---
1443572_at 0.92705 --- ---
1443573_at 1.05129 "poly (ADP-ribose) polymerase family, member 1" Parp1
1443574_at 0.93617 hypothetical protein 6530439I21 6530439I21
1443575_at 0.97268 RIKEN cDNA 2310040G24 gene 2310040G24Rik
1443576_at 1.03498 Transcribed locus ---
1443577_at 0.93445 "Adult male testis cDNA, RIKEN full-length enriched library, clone:4930422D11 product:unclassifiable, full insert sequence" ---
1443578_at 0.90291 Transcribed locus ---
1443579_s_at 0.78797 DEP domain containing 6 Depdc6
1443580_at 0.98667 Syntaxin binding protein 6 (amisyn) Stxbp6
1443581_at 1.07717 "Beta galactoside alpha 2,6 sialyltransferase 1" St6gal1
1443582_at 0.92823 similar to growth arrest-specific 2 like 2; GAS2-related protein RP23-249K18.2
1443583_at 1.04014 RIKEN cDNA C030010B13 gene C030010B13Rik
1443584_at 0.84992 RIKEN cDNA 1110028C15 gene 1110028C15Rik
1443585_at 0.98699 zinc finger protein 526 Zfp526
1443586_at 1.56982 FIP1 like 1 (S. cerevisiae) Fip111
1443587_at 0.93239 cDNA sequence AK129128 AK129128
1443588_at 1.05484 Transcribed locus ---
1443589_at 1.01322 "DNA segment, Chr X, ERATO Doi 242, expressed" DXErtd242e
1443590_at 1.10777 --- ---
1443591_at 1.23480 distal-less homeobox 4 Dlx4
1443592_at 1.26195 expressed sequence AA617406 AA617406
1443593_at 1.01320 --- ---
1443594_at 1.07328 expressed sequence AI043046 AI043046
1443595_at 0.93950 "CCR4-NOT transcription complex, subunit 6-like" Cnot6l
1443596_at 0.98459 "PREDICTED: Mus musculus similar to SRY-box 13, transcript variant 2 (LOC621627), mRNA" ---
1443597_at 1.10323 "cofactor required for Sp1 transcriptional activation, subunit 2" Crsp2

1443598_at	0.88341	Similar to zinc finger protein 709	LOC626854
1443599_at	1.00037	DC-STAMP domain containing 1	Dcst1
1443600_at	0.94443	expressed sequence AA414992	AA414992
1443601_at	1.10407	Hypothetical protein LOC545677	LOC545677
1443602_at	0.99882	Interferon gamma receptor 1	Ifngr1
1443603_at	1.48197	Protein-L-isoaspartate (D-aspartate) O-methyltransferase domain containing 1	Pcmt1
1443604_at	1.03422	---	---
1443605_at	1.07125	expressed sequence C81272	C81272
1443606_at	1.06270	Transcribed locus	---
1443607_at	1.13379	---	---
1443608_at	0.84389	---	---
1443609_s_at	0.88423	"synovial apoptosis inhibitor 1, synoviolin"	Syvn1
1443610_at	0.90386	Transcribed locus	---
1443611_at	0.92845	"DnaJ (Hsp40) homolog, subfamily B, member 4"	Dnajb4
1443612_at	0.84587	Expressed sequence AI838058	AI838058
1443613_x_at	1.12491	Activating signal cointegrator 1 complex subunit 2	Ascc2
1443614_at	1.03837	Transcribed locus	---
1443615_at	1.04081	RIKEN cDNA 1110057K04 gene	1110057K04Rik
1443616_at	0.99213	Synaptophysin-like protein	Sypl
1443617_at	1.02790	SEC15-like 2 (S. cerevisiae)	Sec15l2
1443618_at	0.92328	PDZ domain containing 2	Pdzd2
1443619_at	1.03035	transmembrane protein 30A	Tmem30a
1443620_at	0.84634	Glypican 4	Gpc4
1443621_at	0.92633	RIKEN cDNA 1700010H23 gene	1700010H23Rik
1443622_at	0.98518	Down syndrome critical region gene 3	Dscr3
1443623_a_at	0.95059	"21 days neonate cerebellum cDNA, RIKEN full-length enriched library, clone: G630010A17 product:unclassifiable, full insert sequence"	---
1443624_at	1.11027	CDNA sequence BC013481	BC013481
1443625_at	1.49804	N-acetyltransferase 11	Nat11
1443626_at	1.27800	Chimerin (chimaerin) 2	Chn2
1443627_at	0.81031	---	---
1443628_at	1.24011	---	---
1443629_at	1.09161	neuron navigator 1	Nav1
1443630_at	0.98780	---	---
1443631_at	1.04393	"Diacylglycerol kinase, beta"	Dgkb
1443632_at	0.96237	"obscurin, cytoskeletal calmodulin and titin-interacting RhoGEF"	Obscn
1443633_at	0.96474	RIKEN cDNA 1700086O06 gene	1700086O06Rik
1443634_at	1.01821	"7 days neonate cerebellum cDNA, RIKEN full-length enriched library, clone: A730097O22 product:unclassifiable, full insert sequence"	---
1443635_at	0.92213	"gene model 128, (NCBI)"	Gm128
1443636_at	0.94307	"ATP-binding cassette, sub-family B (MDR/TAP), member 9"	Abcb9
1443637_at	0.95603	RIKEN cDNA 2900084M01 gene	2900084M01Rik

1443638_at	1.06049	"Catenin (cadherin associated protein), delta 2"	Ctnnd2
1443639_at	1.02544	Adenomatosis polyposis coli down-regulated 1	Apccd1
1443640_at	0.87755	Zinc finger protein 617	Zfp617
1443641_at	1.05724	---	---
1443642_at	0.90028	TAO kinase 2	Taok2
1443643_at	1.04464	---	---
1443644_at	1.08903	Brain and reproductive organ-expressed protein	Bre
1443645_at	0.96678	RIKEN cDNA 1700026B20 gene	1700026B20Rik
1443646_at	1.02884	"immunoglobulin superfamily, member 10"	Igsf10
1443647_at	1.03690	RAR-related orphan receptor alpha	Rora
1443648_at	1.07583	RIKEN cDNA 9530003O04 gene	9530003O04Rik
1443649_at	2.78378	"HECT, UBA and WWE domain containing 1"	Huwe1
1443650_at	1.00490	Ankyrin repeat and SAM domain containing 1	Anks1
1443651_at	1.00539	Polyhomeotic-like 3 (Drosophila)	Phc3
1443652_x_at	0.99359	"sprouty protein with EVH-1 domain 1, related sequence"	Spred1
1443653_at	0.96896	---	---
1443654_at	1.14228	Ankyrin repeat and FYVE domain containing 1	Ankfy1
1443655_s_at	1.02224	"phosphatidylinositol 3-kinase, C2 domain containing, alpha polypeptide"	Pik3c2a
1443656_at	0.89676	fucosyltransferase 8	Fut8
1443657_at	1.01243	RIKEN cDNA 6330505N24 gene	6330505N24Rik
1443658_at	1.09324	"DNA segment, Chr 8, ERATO Doi 457, expressed"	D8Ert457e
1443659_at	0.85460	Zinc finger protein 292	Zfp292
1443660_at	0.94164	Neuregulin 3	Nrg3
1443661_at	1.11212	---	---
1443662_at	1.02329	"Catenin (cadherin associated protein), alpha 1"	Ctnna1
1443663_at	0.97515	"Protein kinase, cGMP-dependent, type I"	Prkg1
1443664_s_at	0.91608	D164 sialomucin-like 2	Cd164l2
1443665_at	1.09790	Expressed sequence AI852444	AI852444
1443666_at	1.05547	Midline 1	Mid1
1443667_at	1.03193	expressed sequence C79407	C79407
1443668_x_at	0.91975	expressed sequence C79407	C79407
1443669_at	1.04386	"Zinc finger, FYVE domain containing 28"	Zfyve28
1443670_at	0.82118	RIKEN cDNA 2010001J22 gene	2010001J22Rik
1443671_x_at	1.10170	RIKEN cDNA 2010001J22 gene	2010001J22Rik
1443672_at	0.96463	"leucyl-tRNA synthetase, mitochondrial"	Lars2
1443673_x_at	2.15301	---	---
1443674_at	0.94334	"11 days embryo spinal cord cDNA, RIKEN full-length enriched library, clone: G630090G13 product:unclassifiable, full insert sequence"	---
1443675_at	1.33235	Enhancer of polycomb homolog 1 (Drosophila)	Epc1
1443676_at	1.06948	"Solute carrier family 4 (anion exchanger), member 4"	Slc4a4
1443677_at	0.95539	RIKEN cDNA E030011O05 gene	E030011O05Rik
1443678_at	1.31275	"10 days neonate cerebellum cDNA, RIKEN full-length enriched library, clone:	

B930065L22 product:unclassifiable, full insert sequence" ---

1443679_at 1.05386 RIKEN cDNA 2510003B16 gene 2510003B16Rik

1443680_at 1.11701 Otx2 opposite strand transcript 1 Otx2os1

1443681_at 1.11706 Expressed sequence AI595406 AI595406

1443682_at 0.92363 expressed sequence AI662476 AI662476

1443683_at 1.05121 "Protein tyrosine phosphatase, receptor type, J" Ptpnj

1443684_at 0.88717 --- ---

1443685_at 0.99818 RIKEN cDNA 6430526N21 gene 6430526N21Rik

1443686_at 0.98431 "Histocompatibility 2, class II, locus Mb1" H2-DMb1

1443687_x_at 1.03591 "Histocompatibility 2, class II, locus Mb1" H2-DMb1

1443688_at 0.98415 cDNA sequence BC050092 BC050092

1443689_at 0.96626 Ubiquitin specific peptidase 32 Usp32

1443690_at 1.09400 Transcribed locus ---

1443691_at 0.94022 RIKEN cDNA 2410089E03 gene 2410089E03Rik

1443692_at 1.05088 cDNA sequence BC057022 BC057022

1443693_at 0.97991 expressed sequence AU022332 AU022332

1443694_at 0.93815 regulator of G-protein signaling 20 Rgs20

1443695_at 0.89258 hyaluronic acid binding protein 2 Habp2

1443696_s_at 0.80222 hyaluronic acid binding protein 2 Habp2

1443697_at 1.21633 --- ---

1443698_at 1.31480 F-box protein 39 Fbxo39

1443699_at 0.84843 "protein tyrosine phosphatase, receptor type, K" Ptprk

1443700_at 1.04073 B-cell CLL/lymphoma 9-like Bcl9l

1443701_at 0.85218 cDNA sequence BC025575 BC025575

1443702_at 1.11242 microtubule-associated protein 4 Mtap4

1443703_at 0.97013 CD28 antigen Cd28

1443704_at 0.83370 Adenylate cyclase 8 Adcy8

1443705_at 0.88676 "DNA segment, Chr 14, ERATO Doi 171, expressed" D14Ertd171e

1443706_at 1.01338 C1GALT1-specific chaperone 1 C1galt1c1

1443707_at 1.00436 RIKEN cDNA 2900046B09 gene 2900046B09Rik

1443708_at 1.05073 cDNA sequence BC068281 BC068281

1443709_at 1.13143 Yippee-like 2 (Drosophila) Ypel2

1443710_s_at 0.92260 "spermatogenesis associated, serine-rich 1" Spats1

1443711_at 0.99877 Chromodomain helicase DNA binding protein 8 Chd8

1443712_x_at 0.94858 Coiled-coil domain containing 82 Ccdc82

1443713_at 0.93114 cell adhesion molecule with homology to L1CAM Chl1

1443714_at 1.04493 RIKEN cDNA C530044N13 gene C530044N13Rik

1443715_at 1.02033 similar to RNA binding motif protein 24 /// similar to RNA binding motif protein 24 /// similar to RNA binding motif protein 24 LOC380843 /// LOC638266 /// LOC666794

1443716_at 0.98472 --- ---

1443717_at 1.05126 dihydropyrimidine dehydrogenase Dpyd

1443718_at 1.07165 Transcribed locus ---

1443719_x_at 0.99965 DEAD (Asp-Glu-Ala-Asp) box polypeptide 42 Ddx42

1443720_s_at 1.01315 "bone morphogenetic protein receptor, type 1B" Bmpr1b
1443721_x_at 1.93459 strawberry notch homolog (Drosophila) Stno
1443722_at 0.93784 RIKEN cDNA 6030419C18 gene 6030419C18Rik
1443723_at 1.03080 "transient receptor potential cation channel, subfamily M, member 3" Trpm3
1443724_at 0.98882 junctophilin 3 Jph3
1443725_at 0.96512 --- ---
1443726_at 0.91723 SET and MYND domain containing 1 Smyd1
1443727_x_at 1.00479 SET and MYND domain containing 1 Smyd1
1443728_at 0.96427 Transcribed locus ---
1443729_at 1.01958 metastasis suppressor 1 Mtss1
1443730_at 0.91538 RIKEN cDNA 9130404D14 gene 9130404D14Rik
1443731_at 0.97496 Transcribed locus ---
1443732_at 0.82712 "ATPase, H⁺ transporting, lysosomal V0 subunit A4" Atp6v0a4
1443733_x_at 1.02367 "polymerase (DNA-directed), delta 3, accessory subunit" Pold3
1443734_at 0.99578 "DNA segment, Chr 5, Wayne State University 148, expressed" D5Wsu148e
1443735_at 1.11680 CDNA sequence BC052688 BC052688
1443736_at 0.92930 "Tubulin tyrosine ligase-like family, member 10" Ttl10
1443737_at 0.89983 Exportin 6 Xpo6
1443738_at 0.94747 Phosphatidylserine synthase 2 Ptdss2
1443739_at 1.13833 LPS-responsive beige-like anchor Lrba
1443740_at 1.04094 "ST3 beta-galactoside alpha-2,3-sialyltransferase 6" St3gal6
1443741_x_at 0.97128 Wolf-Hirschhorn syndrome candidate 1 (human) /// similar to Wolf-
Hirschhorn syndrome candidate 1 protein isoform 1 Whsc1 /// LOC622397
1443742_x_at 1.12834 Ubiquitin-conjugating enzyme E2R 2 Ube2r2
1443743_at 1.01294 RIKEN cDNA 1700054O13 gene 1700054O13Rik
1443744_at 0.94502 Deleted in colorectal carcinoma Dcc
1443745_s_at 1.05506 dentin matrix protein 1 Dmp1
1443746_x_at 0.82044 dentin matrix protein 1 Dmp1
1443747_at 1.16641 "DNA segment, Chr 3, ERATO Doi 300, expressed" D3Ertd300e
1443748_x_at 0.94060 ankyrin repeat domain 44 Ankrd44
1443749_x_at 1.11445 "solute carrier family 1 (glial high affinity glutamate transporter), member 3"
Slc1a3
1443750_s_at 1.11396 ribonuclease P 40 subunit (human) Rpp40
1443751_at 1.08629 "Calcium channel, voltage-dependent, beta 4 subunit" Cacnb4
1443752_at 0.97170 RIKEN cDNA E230008N13 gene E230008N13Rik
1443753_at 0.94410 Galactosamine (N-acetyl)-6-sulfate sulfatase Galns
1443754_x_at 0.94131 Limbic system-associated membrane protein Lsamp
1443755_at 0.81973 --- ---
1443756_at 0.96029 "Serine (or cysteine) peptidase inhibitor, clade I, member 1" Serpini1
1443757_x_at 0.88885 "DNA segment, Chr 1, ERATO Doi 161, expressed" D1Ertd161e
1443758_at 0.97291 WD repeat domain 18 Wdr18
1443759_at 1.06175 --- ---
1443760_at 0.94468 Deducator of cyto-kinesis 1 Dock1

1443761_at 0.90455 nurim (nuclear envelope membrane protein) Nrm
1443762_s_at 1.01854 SET binding factor 2 Sbf2
1443763_at 1.06982 Breast cancer 1 Brca1
1443764_x_at 1.01448 "RAB27b, member RAS oncogene family" Rab27b
1443765_at 0.90974 RIKEN cDNA 1810058N05 gene 1810058N05Rik
1443766_x_at 1.02147 RAB11 family interacting protein 4 (class II) Rab11fip4
1443767_at 1.01182 Protein-L-isoaspartate (D-aspartate) O-methyltransferase domain containing 1
Pcmd1
1443768_at 1.09939 --- ---
1443769_at 0.94207 Transcribed locus ---
1443770_x_at 0.92433 Autism susceptibility candidate 2 Aut2
1443771_x_at 0.77658 MAD homolog 7 (Drosophila) Smad7
1443772_at 1.00498 --- ---
1443773_at 1.06719 YLP motif containing 1 Ylpm1
1443774_at 0.88907 zinc finger protein 406 Zfp406
1443775_x_at 0.91111 zinc finger protein 406 Zfp406
1443776_at 0.97350 --- ---
1443777_at 1.04302 --- ---
1443778_at 0.88053 similar to Transcription factor SOX-4 LOC672274
1443779_s_at 1.01232 ligand dependent nuclear receptor corepressor Lcor
1443780_at 0.97928 Transcribed locus ---
1443781_at 1.04957 --- ---
1443782_x_at 0.93612 "cytochrome P450, family 20, subfamily A, polypeptide 1" Cyp20a1
1443783_x_at 1.00376 "Histocompatibility 2, class II antigen A, alpha" H2-Aa
1443784_at 1.01988 RIKEN cDNA A230046K03 gene A230046K03Rik
1443785_x_at 0.77355 PDZ and LIM domain 7 Pdlim7
1443786_at 0.87268 undifferentiated embryonic cell transcription factor 1 Utf1
1443787_x_at 0.86659 Caspase 14 Casp14
1443788_at 0.95476 "Runt-related transcription factor 1; translocated to, 1 (cyclin D-related)"
Runx1t1
1443789_x_at 0.89446 "cytochrome c oxidase, subunit VIIIc" Cox8c
1443790_x_at 0.81471 RIKEN cDNA 4930414L22 gene 4930414L22Rik
1443791_at 1.01576 RIKEN cDNA 4931406I20 gene 4931406I20Rik
1443792_at 1.09781 testis specific gene A14 Tsga14
1443793_x_at 0.96361 integrator complex subunit 1 Ints1
1443794_x_at 1.02807 nucleolar complex associated 4 homolog (S. cerevisiae) Noc4l
1443795_at 0.84651 transcription factor 19 Tcf19
1443796_at 0.93070 interleukin 16 Il16
1443797_at 0.97752 Alkylglycerone phosphate synthase Agps
1443798_at 0.93024 phosphatidylinositol 3-kinase catalytic delta polypeptide Pik3cd
1443799_at 0.97477 "Protein phosphatase 1, regulatory (inhibitor) subunit 14c" Ppp1r14c
1443800_at 0.98998 Ngg1 interacting factor 3-like 1 (S. pombe) Nif3l1
1443801_at 0.82843 Expressed sequence AW822216 AW822216

1443802_at 1.05331 TRAF2 and NCK interacting kinase Tnik
1443803_x_at 0.88240 homeo box A5 Hoxa5
1443804_at 0.88564 "Protein phosphatase 2 (formerly 2A), regulatory subunit B (PR 52), gamma isoform" Ppp2r2c
1443805_at 0.93684 dact 3 Dact3
1443806_x_at 1.06487 Erythrocyte protein band 4.1-like 1 Epb4.111
1443807_x_at 1.28995 cyclin F Ccnf
1443808_at 0.97286 "DNA segment, Chr 2, Brigham & Women's Genetics 1423 expressed" D2Bwg1423e
1443809_at 0.97905 Pleiotrophin Ptn
1443810_at 1.12969 testis-specific serine kinase 4 Tssk4
1443811_at 1.06643 "Protein kinase, cGMP-dependent, type I" Prkg1
1443812_x_at 0.95656 "Protein kinase, cGMP-dependent, type I" Prkg1
1443813_x_at 0.97688 branched chain ketoacid dehydrogenase kinase Bckdk
1443814_x_at 0.84494 cathepsin H Ctsh
1443815_x_at 0.97535 STE20-like kinase (yeast) Slk
1443816_s_at 0.91055 expressed sequence BB220380 BB220380
1443817_x_at 0.86157 "NK2 transcription factor related, locus 6 (Drosophila)" Nkx2-6
1443818_at 0.98291 --- ---
1443819_x_at 1.01033 fission 1 (mitochondrial outer membrane) homolog (yeast) Fis1
1443820_x_at 0.97541 "elongation of very long chain fatty acids (FEN1/Elo2, SUR4/Elo3, yeast)-like 1" Elovl1
1443821_at 0.96806 leukocyte cell-derived chemotaxin 2 Lect2
1443822_s_at 0.97874 --- ---
1443823_s_at 1.01513 "ATPase, Na+/K+ transporting, alpha 2 polypeptide" Atp1a2
1443824_s_at 1.09992 carbonic anhydrase 7 Car7
1443825_x_at 0.88367 sperm acrosome associated 3 Spaca3
1443826_x_at 0.94402 multiple endocrine neoplasia 1 Men1
1443827_x_at 0.89362 cDNA sequence BC004044 BC004044
1443828_x_at 0.71876 HERPUD family member 2 Herpud2
1443829_x_at 0.95444 Coenzyme A synthase Coasy
1443830_x_at 0.93506 ring finger protein 103 Rnf103
1443831_s_at 0.97517 "metallothionein-like 5, testis-specific (tesmin)" Mtl5
1443832_s_at 0.82934 serum deprivation response Sdpr
1443833_at 0.91432 Transcribed locus ---
1443834_at 0.98149 Transcribed locus ---
1443835_x_at 1.02255 RIKEN cDNA 1110014K08 gene /// hypothetical protein LOC664786 /// hypothetical protein LOC664849 /// hypothetical protein LOC669054 /// hypothetical protein LOC672175 1110014K08Rik /// LOC664786 /// LOC664849 /// LOC669054 /// LOC672175
1443836_x_at 0.95316 WD repeat domain 48 Wdr48
1443837_x_at 1.13089 B-cell leukemia/lymphoma 2 Bcl2
1443838_x_at 1.15737 fatty acid desaturase 2 Fads2
1443839_at 1.01357 RIKEN cDNA 1700010H22 gene 1700010H22Rik

1443840_x_at 0.94989 RIKEN cDNA 1700010H22 gene 1700010H22Rik
1443841_x_at 0.97724 --- ---
1443842_at 1.01762 ADP-ribosylation factor guanine nucleotide-exchange factor 2 (brefeldin A-inhibited) /// similar to ADP-ribosylation factor guanine nucleotide-exchange factor 2 (brefeldin A-inhibited) Arfgef2 /// LOC670796
1443843_x_at 0.99844 similar to ribosomal protein L9 LOC433546
1443844_at 1.06346 RIKEN cDNA 1700123J19 gene 1700123J19Rik
1443845_x_at 1.09825 RIKEN cDNA 1700012P22 gene 1700012P22Rik
1443846_x_at 1.19606 Pantothenate kinase 2 (Hallervorden-Spatz syndrome) Pank2
1443847_x_at 0.78710 Transcribed locus ---
1443848_at 1.69803 RNA binding motif protein 16 Rbm16
1443849_x_at 1.06320 uroporphyrinogen decarboxylase Urod
1443850_at 0.95125 Transcribed locus ---
1443851_at 0.88673 RIKEN cDNA 8430415E04 gene 8430415E04Rik
1443852_at 1.03178 Transcribed locus ---
1443853_x_at 0.91364 Scm-like with four mbt domains 1 Sfmbt1
1443854_at 1.01483 --- ---
1443855_at 1.07289 "potassium voltage gated channel, Shaw-related subfamily, member 1" Kcnc1
1443856_at 0.98163 --- ---
1443857_at 0.88045 hook homolog 3 (Drosophila) Hook3
1443858_at 0.87917 Tripartite motif protein 34 Trim34
1443859_at 1.13461 "rosbin, round spermatid basic protein 1" Rsbn1
1443860_at 0.99835 "protein tyrosine phosphatase, receptor type, D" Ptpkd
1443861_at 0.84165 cingulin-like 1 /// similar to cingulin-like 1 isoform 1 Cgnl1 /// LOC677485
1443862_at 2.05305 --- ---
1443863_at 0.96024 fibronectin type III domain containing 3a Fndc3a
1443864_at 1.01419 denticleless homolog (Drosophila) Dtl
1443865_at 1.06791 "gamma-aminobutyric acid (GABA-A) receptor, subunit alpha 2" Gabra2
1443866_at 1.00173 leucine-rich repeats and transmembrane domains 1 Lrtm1
1443867_at 1.21946 ankyrin repeat domain 12 Ankrd12
1443868_at 0.98101 ATPase type 13A3 Atp13a3
1443869_at 0.91765 RIKEN cDNA E430028B21 gene E430028B21Rik
1443870_at 1.28569 "ATP-binding cassette, sub-family C (CFTR/MRP), member 4" Abcc4
1443871_at 0.86568 "Mannosidase, alpha, class 1A, member 2" Man1a2
1443872_at 0.99316 Membrane-associated ring finger (C3HC4) 2 2-Mar
1443873_at 1.16259 --- ---
1443874_at 1.02345 RIKEN cDNA 2810002I04 gene 2810002I04Rik
1443875_at 1.12465 "gene model 603, (NCBI)" Gm603
1443876_at 0.92652 Calcium/calmodulin-dependent protein kinase II alpha Camk2a
1443877_a_at 1.11612 Rap guanine nucleotide exchange factor (GEF) 6 Rapgef6
1443878_at 1.05427 Rap guanine nucleotide exchange factor (GEF) 6 Rapgef6
1443879_at 1.01834 --- ---
1443880_at 1.03633 zinc finger and BTB domain containing 39 Zbtb39

1443881_at 1.01313 protein O-fucosyltransferase 1 Pofut1
1443882_at 1.23400 "Solute carrier family 26 (sulfate transporter), member 2" Slc26a2
1443883_at 1.08579 RIKEN cDNA 2610042O14 gene 2610042O14Rik
1443884_at 1.03779 thyroid adenoma associated Thada
1443885_at 0.96967 "7 days neonate cerebellum cDNA, RIKEN full-length enriched library, clone:
A730060K18 product:unclassifiable, full insert sequence" ---
1443886_at 0.95683 "protein kinase C, zeta 2" Prkcz2
1443887_at 0.86017 ankyrin repeat domain 13c Ankrd13c
1443888_at 0.85420 Expressed sequence AU023762 AU023762
1443889_at 0.88487 RIKEN cDNA 9030619P08 gene 9030619P08Rik
1443890_at 0.97724 Transcribed locus ---
1443891_at 0.87540 RAD51-like 3 (S. cerevisiae) Rad51l3
1443892_at 0.97799 hypothetical LOC435970 /// similar to gonadotropin inducible ovarian
transcription factor 2 LOC435970 /// LOC668620
1443893_at 1.06648 "Tyrosine 3-monooxygenase/tryptophan 5-monooxygenase activation protein,
zeta polypeptide" Ywhaz
1443894_at 1.26013 Ecotropic viral integration site 2a Evi2a
1443895_at 1.12957 hypothetical protein LOC621440 /// hypothetical protein LOC633947 /// region
containing hypothetical protein LOC621440; hypothetical protein LOC633947 LOC621440 ///
LOC633947 /// LOC669138
1443896_at 0.89232 "TBC1 domain family, member 5" Tbc1d5
1443897_at 1.75793 DNA-damage inducible transcript 3 Ddit3
1443898_at 0.96043 B-cell CLL/lymphoma 7C Bcl7c
1443899_at 0.95901 cDNA sequence BC028799 /// RIKEN cDNA 4922501C03 gene BC028799 ///
4922501C03Rik
1443900_at 0.87479 chromodomain helicase DNA binding protein 8 Chd8
1443901_at 0.92068 --- ---
1443902_at 0.91129 RIKEN cDNA 6430573F11 gene 6430573F11Rik
1443903_at 0.91746 "Protein phosphatase 1A, magnesium dependent, alpha isoform" Ppm1a
1443904_at 0.98070 "fatty acid desaturase domain family, member 6" Fads6
1443905_at 2.10711 "Splicing factor, arginine/serine-rich 8" Sfrs8
1443906_at 1.06167 CD55 antigen Cd55
1443907_at 0.87479 leucyl/cystinyl aminopeptidase Lnpep
1443908_at 0.91921 "0 day neonate thymus cDNA, RIKEN full-length enriched library, clone:
A430069K07 product:unclassifiable, full insert sequence" ---
1443909_at 1.27444 "Cleavage stimulation factor, 3' pre-RNA, subunit 3" Cstf3
1443910_at 1.10990 RIKEN cDNA 6330408A02 gene 6330408A02Rik
1443911_at 1.16616 --- ---
1443912_at 0.98490 --- ---
1443913_at 0.96751 Vacuolar protein sorting 33A (yeast) Vps33a
1443914_at 0.79985 "Lipase-like, ab-hydrolase domain containing 2" Lip12
1443915_at 0.76511 mitochondrial ribosomal protein L47 Mrpl47
1443916_at 0.93098 RIKEN cDNA 2900026A02 gene 2900026A02Rik

1443917_at 1.20944 transmembrane BAX inhibitor motif containing 4 Tmbim4
1443918_at 1.04995 RIKEN cDNA 2700050L05 gene 2700050L05Rik
1443919_at 0.84460 proline-rich transmembrane protein 3 Prrt3
1443920_at 0.82332 coiled-coil domain containing 66 Ccdc66
1443921_at 1.00882 RIKEN cDNA C130037N17 gene C130037N17Rik
1443922_at 1.11752 REST corepressor 3 Rcor3
1443923_at 0.82928 A kinase (PRKA) anchor protein 13 Akap13
1443924_at 1.09524 WNK lysine deficient protein kinase 3 Wnk3
1443925_at 0.96507 opposite strand transcription unit to Stag3 Gats
1443926_at 1.16243 Myeloid ecotropic viral integration site-related gene 1 Mrg1
1443927_at 0.95368 ubiquitin-activating enzyme E1-like 2 Ube1l2
1443928_at 0.83642 expressed sequence AI462064 AI462064
1443929_at 0.94886 "In vitro fertilized eggs cDNA, RIKEN full-length enriched library,
clone:7420432F05 product:hypothetical protein, full insert sequence /// In vitro fertilized eggs cDNA,
RIKEN full-length enriched library, clone:7420438K14 product:unclassifiable, full insert sequence"

1443930_at 0.98581 cDNA sequence BC052046 BC052046
1443931_at 0.96905 "Cyclin B2, pseudogene" Ccnb2-ps
1443932_at 0.80154 kelch domain containing 1 Klhdc1
1443933_at 1.08713 membrane targeting (tandem) C2 domain containing 1 Mtac2d1
1443934_at 0.83613 RIKEN cDNA 9230110C19 gene 9230110C19Rik
1443935_at 0.91966 cDNA sequence BC032203 BC032203
1443936_at 1.05261 Olfactory receptor 976 Olfr976
1443937_at 1.20253 Expressed sequence AI790276 AI790276
1443938_at 0.91169 Transcribed locus ---
1443939_at 1.02128 similar to C05G5.5 /// similar to C05G5.5 /// similar to C05G5.5
LOC230628 /// LOC666048 /// LOC670598
1443940_at 0.93676 leucine rich repeat containing 22 Lrrc22
1443941_at 0.94174 hypothetical gene supported by AK049058; BC025881 LOC433886
1443942_at 1.16570 "GA repeat binding protein, beta 2" Gabpb2
1443943_at 1.09469 pleiomorphic adenoma gene 1 Plag1
1443944_at 0.87460 RIKEN cDNA C530005A16 gene C530005A16Rik
1443945_at 0.89609 RIKEN cDNA C130073F10 gene C130073F10Rik
1443946_s_at 0.92141 --- ---
1443947_at 0.96631 LIM and senescent cell antigen-like domains 1 Lims1
1443948_at 1.00130 tetratricopeptide repeat domain 9C Ttc9c
1443949_at 2.25999 "Protein phosphatase 2, regulatory subunit B (B56), epsilon isoform" Ppp2r5e
1443950_at 1.87609 RIKEN cDNA A630042L21 gene A630042L21Rik
1443951_at 1.21069 RIKEN cDNA C230076A16 gene C230076A16Rik
1443952_at 0.90157 thyroid hormone receptor alpha Thra
1443953_at 0.94446 testis expressed gene 2 Tex2
1443954_at 2.16046 RAD18 homolog (S. cerevisiae) Rad18
1443955_at 1.10517 --- ---

1443988_at	1.40620	RIKEN cDNA B330012G18 gene	B330012G18Rik
1443989_at	0.99762	tripartite motif protein 9	Trim9
1443990_at	0.95202	"neurotrophic tyrosine kinase, receptor, type 1"	Ntrk1
1443991_at	0.82035	dedicator of cyto-kinesis 1	Dock1
1443992_at	1.05853	Nipped-B homolog (Drosophila)	Nipbl
1443993_at	0.97231	zinc finger CCCH type containing 12A	Zc3h12a
1443994_at	1.06172	RIKEN cDNA 6030407O03 gene	6030407O03Rik
1443995_at	0.95506	"gene model 9, (NCBI)"	Gm9
1443996_at	1.07005	Fibroblast growth factor receptor 2	Fgfr2
1443997_at	0.97960	RIKEN cDNA C130040D06 gene	C130040D06Rik
1443998_at	0.99619	Transcribed locus	---
1443999_at	1.03108	NIMA (never in mitosis gene a)-related expressed kinase 2	Nek2
1444000_at	0.92462	RIKEN cDNA B930059L03 gene	B930059L03Rik
1444001_at	0.87262	Spermatid perinuclear RNA binding protein	Strbp
1444002_at	0.88653	forkhead box L2	Foxl2
1444003_at	1.39775	lung-inducible neuralized-related C3HC4 RING domain protein	Lincr
1444004_at	1.07138	THO complex 2	Thoc2
1444005_at	0.89391	---	---
1444006_at	0.95946	SET domain containing 2	Setd2
1444007_at	0.81220	RIKEN cDNA A830082K12 gene	A830082K12Rik
1444008_at	0.82520	"H2A histone family, member Y2"	H2afy2
1444009_at	0.91220	Ras association (RalGDS/AF-6) domain family 4	Rassf4
1444010_at	1.06112	hypothetical protein LOC668879 /// hypothetical protein LOC669500	
LOC668879 /// LOC669500			
1444011_at	0.93312	---	---
1444012_at	0.99595	"Yip1 domain family, member 3"	Yipf3
1444013_at	1.02650	RIKEN cDNA 5930412G12 gene	5930412G12Rik
1444014_at	1.02867	hypothetical protein LOC670853	LOC670853
1444015_at	1.03918	Poly (A) polymerase alpha	Papola
1444016_at	0.89434	RIKEN cDNA C430014K04 gene	C430014K04Rik
1444017_at	1.68220	Pellino 1	Peli1
1444018_at	0.87849	RIKEN cDNA B930098A02 gene	B930098A02Rik
1444019_at	1.00426	leucine rich repeat containing 19	Lrrc19
1444020_at	0.97089	chondroitin sulfate proteoglycan 3	Cspg3
1444021_at	0.98450	---	---
1444022_at	0.90149	---	---
1444023_at	1.06203	"ankyrin 2, brain"	Ank2
1444024_at	0.98129	excision repair cross-complementing rodent repair deficiency complementation group 6 - like	Ercc6l
1444025_at	1.09970	---	---
1444026_at	1.08472	expressed sequence AI593442	AI593442
1444027_at	0.96036	"solute carrier family 30 (zinc transporter), member 8"	Slc30a8
1444028_s_at	0.98207	dedicator of cytokinesis 9 /// similar to Dedicator of cytokinesis protein 9	

(Cdc42 guanine nucleotide exchange factor zizimin 1) Dock9 /// LOC670309

1444029_at 0.97914 "poly (ADP-ribose) polymerase family, member 11" Parp11

1444030_at 1.03180 DCP1 decapping enzyme homolog b (S. cerevisiae) Dcp1b

1444031_at 1.02600 --- ---

1444032_at 0.95792 Kidney expressed gene 1 Keg1

1444033_at 1.06173 Anaphase promoting complex subunit 10 Anapc10

1444034_at 0.91851 Rap guanine nucleotide exchange factor (GEF) 4 Rapgef4

1444035_at 0.86609 mesoderm development candiate 2 Mesdc2

1444036_at 1.00664 "Protein phosphatase 2, regulatory subunit B (B56), alpha isoform" Ppp2r5a

1444037_at 0.92023 "lectin, mannose-binding, 1" Lman1

1444038_at 1.03244 expressed sequence AU015836 AU015836

1444039_at 0.90103 dispatched homolog 2 (Drosophila) Disp2

1444040_at 1.11504 leukocyte-associated Ig-like receptor 1 Lair1

1444041_at 0.88614 expressed sequence AU041133 AU041133

1444042_at 1.04845 thyroid hormone receptor interactor 11 Trip11

1444043_at 1.69529 GLI-Kruppel family member GLI3 Gli3

1444044_at 1.03210 "Neural precursor cell expressed, developmentally down-regulted gene 4" Nedd4

1444045_at 0.91967 Transcribed locus ---

1444046_at 1.09199 RIKEN cDNA D430041B17 gene D430041B17Rik

1444047_at 0.98364 Potassium channel tetramerisation domain containing 8 Kctd8

1444048_at 1.26527 Retinitis pigmentosa 9 homolog (human) Rp9h

1444049_at 1.00252 "adenosine deaminase, tRNA-specific 1" Adat1

1444050_at 0.97998 "PREDICTED: Mus musculus WNT1 inducible signaling pathway protein 3 (Wisp3), mRNA" ---

1444051_at 1.03830 RIKEN cDNA 5830405N20 gene 5830405N20Rik

1444052_at 1.22920 Transcribed locus ---

1444053_at 0.94979 single-stranded DNA binding protein 1 Ssbp1

1444054_at 1.01825 --- ---

1444055_at 0.99962 Heparan sulfate (glucosamine) 3-O-sulfotransferase 3B1 Hs3st3b1

1444056_at 1.42531 RIKEN cDNA 2900006B13 gene 2900006B13Rik

1444057_at 1.22846 Transcribed locus ---

1444058_at 1.05473 RIKEN cDNA 2310047C04 gene 2310047C04Rik

1444059_at 0.96452 RIKEN cDNA 6330415G19 gene 6330415G19Rik

1444060_at 1.02785 RIKEN cDNA 2610027K06 gene 2610027K06Rik

1444061_at 1.08081 RIKEN cDNA A030004J04 gene A030004J04Rik

1444062_at 1.12791 RIKEN cDNA 2900056L01 gene 2900056L01Rik

1444063_at 0.87324 RIKEN cDNA 5430435G22 gene 5430435G22Rik

1444064_at 1.24126 "SAM domain and HD domain, 1" Samhd1

1444065_at 1.24580 Cytochrome b5 domain containing 2 Cyb5d2

1444066_at 1.06607 GTPase activating protein and VPS9 domains 1 Gapvd1

1444067_at 0.96363 phosphodiesterase 1C Pde1c

1444068_at 1.11331 thyroid hormone receptor associated protein 1 Thrapp1

1444069_at 0.99927 Autism susceptibility candidate 2 Aut2

1444070_at 0.86628 --- ---
 1444071_at 0.79239 RIKEN cDNA 9630013A20 gene 9630013A20Rik
 1444072_at 0.82951 Transcribed locus ---
 1444073_at 0.82612 avian musculoaponeurotic fibrosarcoma (v-maf) AS42 oncogene homolog
 Maf
 1444074_at 0.95785 synaptic nuclear envelope 2 /// similar to Nesprin-2 (Nuclear envelope spectrin
 repeat protein 2) (Syne-2) (Synaptic nuclear envelope protein 2) (Nucleus and actin connecting element
 protein) (NUANCE protein) Syne2 /// LOC677266
 1444075_at 1.02558 filamin A interacting protein 1 Filip1
 1444076_at 0.94017 zinc finger protein 81 Zfp81
 1444077_at 0.99431 Transcribed locus ---
 1444078_at 0.98566 "CD8 antigen, alpha chain /// similar to T-cell surface glycoprotein CD8 alpha
 chain precursor (T-cell surface glycoprotein Lyt-2) /// similar to T-cell surface glycoprotein CD8 alpha
 chain precursor (T-cell surface glycoprotein Lyt-2) (CD8a antigen)" Cd8a /// LOC636147 ///
 LOC669166
 1444079_at 1.02693 defensin beta 8 Defb8
 1444080_at 0.90409 --- ---
 1444081_at 0.87094 RIKEN cDNA 5033428A16 gene 5033428A16Rik
 1444082_at 0.86327 RIKEN cDNA A730017C20 gene A730017C20Rik
 1444083_at 1.04425 titin Ttn
 1444084_at 0.96560 proline rich Gla (G-carboxyglutamic acid) 4 (transmembrane) /// similar to
 Transmembrane gamma-carboxyglutamic acid protein 4 precursor (Proline-rich Gla protein 4) (Proline-
 rich gamma-carboxyglutamic acid protein 4) Prrg4 /// LOC676960
 1444085_at 1.02773 "prenyl (solanesyl) diphosphate synthase, subunit 2" Pdss2
 1444086_at 0.67971 RIKEN cDNA E030049G20 gene E030049G20Rik
 1444087_at 1.30362 PRP38 pre-mRNA processing factor 38 (yeast) domain containing A Prpf38a
 1444088_at 0.97329 --- ---
 1444089_at 1.27118 spectrin beta 2 Spnb2
 1444090_at 0.96111 PML-RAR alpha-regulated adaptor molecule 1 Pram1
 1444091_a_at 0.94170 RIKEN cDNA 6430571L13 gene 6430571L13Rik
 1444092_at 1.06831 RIKEN cDNA 9430025M13 gene 9430025M13Rik
 1444093_at 0.96671 RIKEN cDNA 4930557M11 gene 4930557M11Rik
 1444094_at 1.06934 --- ---
 1444095_a_at 0.84594 --- ---
 1444096_at 3.08438 Zinc finger RNA binding protein Zfr
 1444097_at 1.05078 "meteorin, glial cell differentiation regulator-like" Metrnl
 1444098_at 1.13195 RIKEN cDNA 1700010H15 gene 1700010H15Rik
 1444099_at 0.93725 "PREDICTED: Mus musculus similar to apolipoprotein L, 3 (predicted),
 transcript variant 2 (LOC278677), mRNA" ---
 1444100_at 1.05993 podoplanin Pdpn
 1444101_at 1.18434 outer dense fiber of sperm tails 2 Odf2
 1444102_at 0.91827 --- ---
 1444103_at 0.98501 Epithelial stromal interaction 1 (breast) Epsti1

1444104_at 1.02781 --- ---
1444105_at 0.91004 "actin, alpha 2, smooth muscle, aorta" Acta2
1444106_at 1.04487 RIKEN cDNA D330012F22 gene D330012F22Rik
1444107_at 1.09423 RIKEN cDNA 9430029N19 gene 9430029N19Rik
1444108_at 1.07903 "Adult male liver tumor cDNA, RIKEN full-length enriched library, clone:
C730040G13 product:unclassifiable, full insert sequence" ---
1444109_at 1.76619 RIKEN cDNA C130009A20 gene C130009A20Rik
1444110_at 1.03293 transcobalamin 2 Tcn2
1444111_at 1.06323 --- ---
1444112_at 0.80021 "6 days neonate head cDNA, RIKEN full-length enriched library,
clone:5430409O07 product:unclassifiable, full insert sequence" ---
1444113_at 1.06528 Transcribed locus ---
1444114_at 1.49930 "Muskelin 1, intracellular mediator containing kelch motifs" Mkln1
1444115_at 0.97722 "PREDICTED: Mus musculus similar to membrane-associated ring finger
(C3HC4) 2 (LOC623771), mRNA" ---
1444116_at 1.15510 "MAK10 homolog, amino-acid N-acetyltransferase subunit, (S. cerevisiae)"
Mak10
1444117_at 1.02024 adhesion molecule with Ig like domain 1 Amigo1
1444118_at 1.00723 --- ---
1444119_at 1.02087 RIKEN cDNA B930006L02 gene B930006L02Rik
1444120_at 0.99327 bridging integrator 1 Bin1
1444121_at 0.93551 Transcribed locus ---
1444122_at 1.01141 RIKEN cDNA 4930518F03 gene 4930518F03Rik
1444123_at 1.01432 "PR domain containing 2, with ZNF domain" Prdm2
1444124_a_at 1.19368 RIKEN cDNA E330017L17 gene E330017L17Rik
1444125_at 0.94749 "Platelet-activating factor acetylhydrolase, isoform 1b, beta1 subunit" Pafah1b1
1444126_at 1.58539 "Praja 2, RING-H2 motif containing" Pja2
1444127_at 0.96164 ankyrin repeat and SOCS box-containing protein 1 Asb1
1444128_at 1.07531 "Phosphatidylinositol-4-phosphate 5-kinase, type II, beta" Pip5k2b
1444129_at 1.03467 --- ---
1444130_at 1.74102 RIKEN cDNA 1700081L11 gene 1700081L11Rik
1444131_at 0.90166 RIKEN cDNA 2610101N10 gene 2610101N10Rik
1444132_at 1.13282 RIKEN cDNA 2310003H01 gene 2310003H01Rik
1444133_at 0.92911 WNK lysine deficient protein kinase 4 Wnk4
1444134_at 0.97642 v-abl Abelson murine leukemia oncogene 1 Abl1
1444135_at 0.91319 RIKEN cDNA 6332401O19 gene 6332401O19Rik
1444136_at 1.16387 Retinoic acid induced 12 Rai12
1444137_at 1.00803 RIKEN cDNA A430108G06 gene A430108G06Rik
1444138_at 1.06123 "cytochrome P450, family 2, subfamily r, polypeptide 1" Cyp2r1
1444139_at 0.92412 DNA-damage-inducible transcript 4-like Ddit4l
1444140_at 1.64672 --- ---
1444141_at 0.88249 sorting nexin 13 Snx13
1444142_at 0.95276 hypothetical LOC432591 /// hypothetical protein LOC639155

LOC432591 /// LOC639155

1444143_at 0.94789 RIKEN cDNA B230342M21 gene B230342M21Rik
 1444144_at 1.48388 RIKEN cDNA 5530400K22 gene 5530400K22Rik
 1444145_at 0.88034 Methyltransferase like 6 Mettl6
 1444146_at 0.91008 ring finger protein 20 Rnf20
 1444147_at 1.00783 proprotein convertase subtilisin/kexin type 2 Pcsk2
 1444148_at 0.97573 hypothetical LOC497253 LOC497253
 1444149_at 0.91047 zinc finger protein 96 Zfp96
 1444150_at 1.29806 erythrocyte protein band 4.1 Epb4.1
 1444151_at 1.00602 RIKEN cDNA 2600009P04 gene 2600009P04Rik
 1444152_at 1.01315 "CUG triplet repeat, RNA binding protein 2" Cugbp2
 1444153_at 0.90493 "16 days neonate heart cDNA, RIKEN full-length enriched library, clone:
 D830033B01 product:unclassifiable, full insert sequence" ---
 1444154_at 1.11903 NADH dehydrogenase (ubiquinone) 1 alpha subcomplex 11 Ndufa11
 1444155_at 0.85319 Ninein Nin
 1444156_at 0.89976 RIKEN cDNA 9230112E08 gene 9230112E08Rik
 1444157_a_at 0.95726 "jumonji, AT rich interactive domain 1C (Rbp2 like)" Jarid1c
 1444158_at 0.89277 "jumonji, AT rich interactive domain 1C (Rbp2 like)" Jarid1c
 1444159_at 1.36631 "Phosphatidylinositol-4-phosphate 5-kinase, type II, alpha" Pip5k2a
 1444160_at 1.02166 Polycystic kidney disease 1 like 3 Pkd113
 1444161_at 0.93165 RIKEN cDNA A530020G20 gene A530020G20Rik
 1444162_at 1.15983 fibroblast growth factor receptor substrate 2 Frs2
 1444163_at 1.09818 NIMA (never in mitosis gene a)-related expressed kinase 5 Nek5
 1444164_at 0.88532 FK506 binding protein 3 Fkbp3
 1444165_at 0.84811 layilin Layn
 1444166_at 1.02883 thyroid hormone responsive SPOT14 homolog (Rattus) Thrsp
 1444167_at 0.89548 Transcribed locus ---
 1444168_at 1.09006 xenotropic and polytropic retrovirus receptor 1 Xpr1
 1444169_at 1.16729 RIKEN cDNA 3110052M02 gene 3110052M02Rik
 1444170_at 0.87562 RIKEN cDNA 6030446N20 gene 6030446N20Rik
 1444171_at 1.08543 Programmed cell death 4 Pcd4
 1444172_at 0.95259 Transcribed locus ---
 1444173_at 1.02787 "8 days neonate male testis cDNA, RIKEN full-length enriched library, clone:
 G630012C09 product:unclassifiable, full insert sequence" ---
 1444174_at 0.99541 "A disintegrin-like and metallopeptidase (reprolysin type) with thrombospondin
 type 1 motif, 8" Adamts8
 1444175_at 1.41458 ADP-ribosylation factor guanine nucleotide-exchange factor 1(brefeldin A-
 inhibited) Arfgef1
 1444176_at 1.17595 "ATPase, H+ transporting, lysosomal V0 subunit D2" Atp6v0d2
 1444177_at 0.92166 "PREDICTED: Mus musculus hypothetical protein LOC626058 (LOC626058),
 mRNA" ---
 1444178_at 1.05748 hypothetical gene supported by AK028012 LOC433237
 1444179_at 1.01546 RIKEN cDNA A530023O14 gene A530023O14Rik

1444180_at 1.31702 "12 days embryo spinal ganglion cDNA, RIKEN full-length enriched library, clone:D130055E11 product:unclassifiable, full insert sequence" ---
1444181_at 0.94348 "GTPase, IMAP family member 5" Gimap5
1444182_at 0.95801 RIKEN cDNA 2210010C17 gene 2210010C17Rik
1444183_at 1.26020 RIKEN cDNA C030013C21 gene C030013C21Rik
1444184_at 1.30807 "alkB, alkylation repair homolog 3 (E. coli)" Alkbh3
1444185_at 1.09099 Stromal interaction molecule 2 Stim2
1444186_at 1.14255 RIKEN cDNA 2900006B13 gene 2900006B13Rik
1444187_at 1.09545 similar to spermatogenesis associated glutamate (E)-rich protein 4d /// similar to cyclin-dependent kinase 6 LOC544917 /// LOC667335
1444188_at 0.98439 --- ---
1444189_at 0.96029 Transcribed locus ---
1444190_at 0.91652 RIKEN cDNA E030049G20 gene E030049G20Rik
1444191_at 1.06340 --- ---
1444192_at 1.12614 RIKEN cDNA 9630020C08 gene 9630020C08Rik
1444193_at 0.99949 "alcohol dehydrogenase, iron containing, 1" Adhfe1
1444194_at 3.69498 --- ---
1444195_at 1.82125 Required for meiotic nuclear division 5 homolog A (S. cerevisiae) Rmnd5a
1444196_at 0.93902 expressed sequence AI662501 AI662501
1444197_at 0.94986 RIKEN cDNA 9930116N10 gene 9930116N10Rik
1444198_at 1.00266 Dorso-medial telencephalon gene 2 Dmt2
1444199_at 0.72568 --- ---
1444200_at 0.88356 "DNA segment, Chr 19, ERATO Doi 386, expressed" D19Erttd386e
1444201_at 0.96852 centrosomal protein 135 Cep135
1444202_at 0.91456 zinc finger protein 703 Zfp703
1444203_at 0.98806 "PREDICTED: Mus musculus similar to HYPOTHETICAL PROTEIN ORF-1137 (LOC626244), mRNA" ---
1444204_at 0.93816 Homeo box B2 Hoxb2
1444205_at 1.24541 MAD homolog 4 (Drosophila) Smad4
1444206_at 0.87655 RIKEN cDNA A030001O10 gene A230042K10Rik
1444207_at 1.06879 Camello-like 3 Cml3
1444208_at 1.03307 "Dynein, axonemal, heavy chain 1" Dnahc1
1444209_at 0.85498 "DNA segment, Chr 19, ERATO Doi 737, expressed" D19Erttd737e
1444210_at 0.96019 --- ---
1444211_at 1.06739 Endonuclease G-like 1 Endogl1
1444212_at 0.97053 Huntingtin interacting protein 2 Hip2
1444213_at 1.06347 RIKEN cDNA 1700012D14 gene 1700012D14Rik
1444214_at 0.99857 RIKEN cDNA 2810484G07 gene 2810484G07Rik
1444215_at 1.02582 Protocadherin 9 Pcdh9
1444216_at 1.00569 empty spiracles homolog 1 (Drosophila) Emx1
1444217_at 1.22058 mitochondrial ribosomal protein L38 Mrpl38
1444218_at 1.51417 "DNA segment, Chr 19, ERATO Doi 737, expressed" D19Erttd737e
1444219_at 1.05478 cDNA sequence AY702103 AY702103

1444220_at 1.37686 RIKEN cDNA 2410089E03 gene 2410089E03Rik
1444221_at 0.99514 "TBC1D12: TBC1 domain family, member 12" Tbc1d12
1444222_x_at 0.97596 Transcribed locus ---
1444223_at 0.83709 hypocretin (orexin) receptor 2 Hcrtr2
1444224_at 1.11700 "Tumor necrosis factor, alpha-induced protein 8" Tnfaip8
1444225_at 0.95683 SH3-domain GRB2-like (endophilin) interacting protein 1 Sgip1
1444226_at 0.95527 forkhead box O3a Foxo3a
1444227_at 1.02697 Transcribed locus ---
1444228_s_at 0.93645 hect (homologous to the E6-AP (UBE3A) carboxyl terminus) domain and
RCC1 (CHC1)-like domain (RLD) 2 Herc2
1444229_at 1.04261 "nuclear receptor subfamily 2, group F, member 2" Nr2f2
1444230_at 1.05256 RIKEN cDNA 2900046D03 gene 2900046D03Rik
1444231_at 1.37808 "Zinc finger, DHHC domain containing 6" Zdhhc6
1444232_at 0.79513 "protein kinase, cGMP-dependent, type I" Prkg1
1444233_at 0.92437 G protein-coupled receptor 132 Gpr132
1444234_at 0.93136 --- ---
1444235_at 0.93099 RIKEN cDNA 1700025G04 gene 1700025G04Rik
1444236_at 0.95231 expressed sequence AW552393 AW552393
1444237_at 1.11917 RIKEN cDNA C330019G07 gene C330019G07Rik
1444238_at 1.00012 "7 days neonate cerebellum cDNA, RIKEN full-length enriched library, clone:
A730093O14 product:unclassifiable, full insert sequence" ---
1444239_at 0.91397 CDNA sequence BC023488 BC023488
1444240_at 0.95408 SH3/ankyrin domain gene 1 Shank1
1444241_at 1.09928 WW domain containing adaptor with coiled-coil Wac
1444242_at 1.01424 --- ---
1444243_at 1.01923 "WAS protein family, member 3" Wasf3
1444244_at 0.82055 Transcribed locus ---
1444245_at 1.13978 RIKEN cDNA B230118H07 gene B230118H07Rik
1444246_at 1.37033 Chromodomain helicase DNA binding protein 2 Chd2
1444247_at 1.05986 expressed sequence C81557 C81557
1444248_at 0.98148 reticulocalbin 2 Rcn2
1444249_at 1.00384 Transcribed locus ---
1444250_at 1.01110 --- ---
1444251_x_at 0.97850 "DNA segment, Chr 4, ERATO Doi 111, expressed" D4Ertd111e
1444252_at 1.01489 "0 day neonate eyeball cDNA, RIKEN full-length enriched library, clone:
E130003J08 product:unclassifiable, full insert sequence" ---
1444253_at 0.85666 "a disintegrin-like and metallopeptidase (reprolysin type) with thrombospondin
type 1 motif, 18" Adamts18
1444254_at 0.91764 tensin 4 Tns4
1444255_at 1.01064 RIKEN cDNA D630045J12 gene D630045J12Rik
1444256_at 1.10173 RAE1 RNA export 1 homolog (S. pombe) Rae1
1444257_at 1.25328 proline rich 11 Prr11
1444258_at 1.53520 Actinin alpha 4 Actn4

1444259_at 0.78961 Dimethylglycine dehydrogenase precursor Dmgdh
 1444260_at 0.87800 "Transcribed locus, weakly similar to XP_573711.1 PREDICTED: similar to ORF2 consensus sequence encoding endonuclease and reverse transcriptase minus RNaseH [Rattus norvegicus]" ---
 1444261_at 1.08545 --- ---
 1444262_at 0.82518 RIKEN cDNA 1110017F19 gene 1110017F19Rik
 1444263_at 0.96918 Enhancer of zeste homolog 2 (Drosophila) Ezh2
 1444264_at 1.16470 "Tumor necrosis factor superfamily, member 5-induced protein 1" Tnfsf5ip1
 1444265_at 1.13265 --- ---
 1444266_at 1.16386 Homeobox only domain Hod
 1444267_at 0.89765 Xenotropic and polytropic retrovirus receptor 1 Xpr1
 1444268_at 1.34946 cDNA sequence BC004004 BC004004
 1444269_at 1.12906 --- ---
 1444270_at 1.05746 radial spokehead-like 3 Rshl3
 1444271_at 0.98201 anaphase promoting complex subunit 1 Anapc1
 1444272_at 0.93595 PDZ domain containing 6 Pdzd6
 1444273_at 2.40158 expressed sequence AW555355 AW555355
 1444274_at 0.99565 --- ---
 1444275_at 1.04001 Uronyl-2-sulfotransferase Ust
 1444276_at 0.98750 immature colon carcinoma transcript 1 Ict1
 1444277_at 1.03436 "Adult male liver tumor cDNA, RIKEN full-length enriched library, clone: C730026F15 product:unclassifiable, full insert sequence" ---
 1444278_at 0.98718 --- ---
 1444279_at 1.61931 "HECT, UBA and WWE domain containing 1" Huwe1
 1444280_at 0.95324 RIKEN cDNA A930001M12 gene A930001M12Rik
 1444281_at 0.94894 "Calcium/calmodulin-dependent protein kinase II, delta" Camk2d
 1444282_at 1.01059 Sterile alpha motif domain containing 10 Samd10
 1444283_at 1.02715 "GTPase, IMAP family member 7" Gimap7
 1444284_at 0.99121 RIKEN cDNA A130050O07 gene A130050O07Rik
 1444285_at 1.12870 Mortality factor 4 like 1 Morf411
 1444286_at 0.93319 "Transcribed locus, weakly similar to XP_530798.1 PREDICTED: hypothetical protein XP_530798 [Pan troglodytes]" ---
 1444287_at 0.92484 expressed sequence AI853106 AI853106
 1444288_at 0.90387 polyribonucleotide nucleotidyltransferase 1 Pnpt1
 1444289_at 0.86458 "Yip1 domain family, member 5" Yipf5
 1444290_at 1.15607 six transmembrane epithelial antigen of prostate 2 Steap2
 1444291_at 1.02670 "LysM, putative peptidoglycan-binding, domain containing 4" Lysmd4
 1444292_at 0.99030 "DNA segment, Chr 7, ERATO Doi 143, expressed" D7Ert143e
 1444293_at 0.90126 cDNA sequence BC028471 BC028471
 1444294_at 0.91741 "Adult male testis cDNA, RIKEN full-length enriched library, clone:4930551A22 product:unclassifiable, full insert sequence" ---
 1444295_at 1.66944 neogenin Neo1
 1444296_a_at 0.98673 "serine (or cysteine) peptidase inhibitor, clade A, member 4, pseudogene 1"

Serpina4-ps1

1444297_at 0.91607 "serine (or cysteine) peptidase inhibitor, clade A, member 4, pseudogene 1"

Serpina4-ps1

1444298_at 1.10071 RIKEN cDNA A130090K04 gene A130090K04Rik

1444299_at 1.09707 RIKEN cDNA A430093F15 gene A430093F15Rik

1444300_at 0.96116 G protein-coupled receptor 125 Gpr125

1444301_at 0.97710 protocadherin 10 Pcdh10

1444302_at 0.87887 cDNA sequence BC049816 BC049816

1444303_at 1.01190 MAP/microtubule affinity-regulating kinase 2 Mark2

1444304_at 0.96156 RIKEN cDNA A330102K23 gene A330102K23Rik

1444305_at 0.90545 hypothetical protein LOC668940 /// hypothetical protein LOC669705 ///
hypothetical protein LOC674786 LOC668940 /// LOC669705 /// LOC674786

1444306_at 1.05181 Musashi homolog 1(Drosophila) Msi1

1444307_at 1.20266 Transcribed locus ---

1444308_at 0.93170 --- ---

1444309_at 0.99072 "Succinate-Coenzyme A ligase, GDP-forming, beta subunit" Suclg2

1444310_at 1.01806 polymerase (RNA) III (DNA directed) polypeptide A Polr3a

1444311_at 0.86632 --- ---

1444312_at 0.88096 "solute carrier family 22 (organic anion/cation transporter), member 15"

Slc22a15

1444313_at 1.02221 Dymeclin Dym

1444314_at 1.15289 RIKEN cDNA 4921506I22 gene 4921506I22Rik

1444315_at 1.15760 RIKEN cDNA D630024D03 gene D630024D03Rik

1444316_at 1.01112 --- ---

1444317_at 1.05713 protocadherin 15 Pcdh15

1444318_at 0.94867 Coiled-coil-helix-coiled-coil-helix domain containing 7 Chchd7

1444319_at 0.84657 E2F transcription factor 8 E2f8

1444320_at 1.32623 DDHD domain containing 2 Ddhd2

1444321_at 0.95978 "Proteasome (prosome, macropain) 26S subunit, non-ATPase, 9" Psmc9

1444322_at 0.81940 lipocalin 11 Lcn11

1444323_at 1.04051 cyclin D3 Ccnd3

1444324_at 0.91772 RIKEN cDNA E530001K10 gene E530001K10Rik

1444325_at 0.99317 Phosphatase and tensin homolog Pten

1444326_at 1.31644 "Prolyl 4-hydroxylase, beta polypeptide" P4hb

1444327_at 0.78976 "E3 ubiquitin protein ligase, HECT domain containing, 1" Edd1

1444328_at 1.35998 "clathrin, light polypeptide (Lca)" Clta

1444329_at 1.01380 glutamine fructose-6-phosphate transaminase 1 Gfpt1

1444330_at 1.23904 "DNA segment, Chr 2, ERATO Doi 173, expressed" D2Ert173e

1444331_at 1.02939 Synaptoporin Synpr

1444332_at 0.87857 RIKEN cDNA 9130019P16 gene 9130019P16Rik

1444333_at 1.80441 --- ---

1444334_at 0.94562 --- ---

1444335_at 0.99578 "ER degradation enhancer, mannosidase alpha-like 2" Edem2

1444336_at 1.00338 "Potassium inwardly-rectifying channel, subfamily J, member 3" Kcnj3
1444337_at 1.24635 Numb gene homolog (Drosophila) Numb
1444338_at 1.04035 --- ---
1444339_at 0.91487 "0 day neonate kidney cDNA, RIKEN full-length enriched library, clone:
D630003P16 product:unclassifiable, full insert sequence" ---
1444340_at 1.13591 RIKEN cDNA C230066G23 gene C230066G23Rik
1444341_at 0.92448 RIKEN cDNA 8030451F13 gene 8030451F13Rik
1444342_at 1.06548 Transcribed locus ---
1444343_at 1.67019 RIKEN cDNA A130064L14 gene A130064L14Rik
1444344_at 1.00673 --- ---
1444345_at 1.03332 Transcribed locus ---
1444346_at 1.05231 SH3-domain binding protein 5 (BTK-associated) Sh3bp5
1444347_at 1.03955 "mannoside acetylglucosaminyltransferase 4, isoenzyme A" Mgat4a
1444348_at 0.86692 --- ---
1444349_at 1.24280 --- ---
1444350_at 1.14852 similar to schlafen 10 LOC672855
1444351_at 1.11489 Cytoplasmic FMR1 interacting protein 2 Cyfip2
1444352_at 0.93726 zinc finger protein 287 Zfp287
1444353_at 1.01798 RIKEN cDNA C030040A22 gene C030040A22Rik
1444354_at 1.07357 "AF4/FMR2 family, member 3" Aff3
1444355_at 1.01651 "ATPase, aminophospholipid transporter (APLT), class I, type 8A, member
1" Atp8a1
1444356_at 0.91928 Rab6 interacting protein 2 Rab6ip2
1444357_at 0.95105 A kinase (PRKA) anchor protein 6 Akap6
1444358_at 1.38844 F-box and leucine-rich repeat protein 11 Fbxl11
1444359_at 0.99639 RIKEN cDNA 2500002G23 gene 2500002G23Rik
1444360_at 0.94002 CDNA sequence BC038822 BC038822
1444361_at 0.99966 --- ---
1444362_at 1.08069 neurogenic differentiation 2 Neurod2
1444363_at 0.92425 "10 days neonate cortex cDNA, RIKEN full-length enriched library, clone:
A830040F21 product:unclassifiable, full insert sequence" ---
1444364_at 1.20826 REST corepressor 3 Rcor3
1444365_at 1.05231 TSC22 domain family 3 Tsc22d3
1444366_at 1.09499 TAO kinase 3 Taok3
1444367_at 1.02264 fibronectin type III domain containing 7 Fndc7
1444368_at 1.02184 --- ---
1444369_at 1.00176 RIKEN cDNA 2310016D23 gene 2310016D23Rik
1444370_at 0.83953 expressed sequence C77058 C77058
1444371_at 1.15572 RIKEN cDNA A630007B06 gene A630007B06Rik
1444372_at 1.31082 Archain 1 Arcn1
1444373_at 1.00212 T-cell lymphoma invasion and metastasis 1 Tiam1
1444374_at 1.05120 "PREDICTED: Mus musculus hypothetical LOC381451 (LOC381451),
mRNA" ---

1444375_at 0.96359 Transcribed locus ---
1444376_at 1.16693 Sestrin 1 Sesn1
1444377_at 1.39855 "Proteasome (prosome, macropain) subunit, beta type 2" Psmb2
1444378_at 1.73923 "Casein kinase 1, delta" Csnk1d
1444379_at 0.87178 PWWP domain containing 2 Pwwp2
1444380_at 0.90276 expressed sequence AI844869 AI844869
1444381_at 1.13463 RIKEN cDNA A830005F24 gene A830005F24Rik
1444382_at 1.03081 RIKEN cDNA 4930579G22 gene 4930579G22Rik
1444383_at 0.92961 RIKEN cDNA 1110014J01 gene 1110014J01Rik
1444384_at 1.43561 Expressed sequence AI591476 AI591476
1444385_at 1.27595 TAO kinase 1 Taok1
1444386_at 0.86441 phosphodiesterase 1C Pde1c
1444387_at 1.28591 N-myristoyltransferase 2 Nmt2
1444388_at 1.03361 Transcribed locus ---
1444389_at 0.94355 "DNA segment, Chr 14, ERATO Doi 426, expressed" D14Ertd426e
1444390_at 0.87405 PR domain containing 14 /// similar to PR domain containing 14 Prdm14 ///
LOC637693
1444391_at 0.94116 Transcribed locus ---
1444392_at 1.15411 Leprecan-like 1 Leprel1
1444393_at 1.08746 Regulating synaptic membrane exocytosis 1 Rims1
1444394_at 1.00591 "acyl-Coenzyme A dehydrogenase family, member 9" Acad9
1444395_at 0.85511 DIX domain containing 1 Dixdc1
1444396_at 1.13577 Tumor protein p53 inducible nuclear protein 2 Trp53inp2
1444397_at 0.92451 sperm flagellar 1 Spef1
1444398_at 1.10291 "fascin homolog 1, actin bundling protein (Strongylocentrotus purpuratus)"
Fscn1
1444399_at 1.02725 RIKEN cDNA 5830411E10 gene 5830411E10Rik
1444400_at 1.09386 retinoblastoma 1 Rb1
1444401_at 1.36931 Expressed sequence C80913 C80913
1444402_at 1.24631 zinc finger CCCH type containing 12C Zc3h12c
1444403_at 0.98500 "Core-binding factor, runt domain, alpha subunit 2, translocated to, 2 homolog
(human)" Cbfa2t2h
1444404_at 0.88261 RIKEN cDNA 330010C22 gene D330010C22Rik
1444405_at 1.01015 RIKEN cDNA E430029J22 gene E430029J22Rik
1444406_at 2.12310 Mitogen-activated protein kinase kinase kinase kinase 3 Map4k3
1444407_at 0.99675 Transcribed locus ---
1444408_at 1.14527 RIKEN cDNA B130040O20 gene B130040O20Rik
1444409_at 0.80788 Kelch repeat and BTB (POZ) domain containing 8 Kbtbd8
1444410_at 0.88671 "acid phosphatase 6, lysophosphatidic" Acp6
1444411_at 1.24995 --- ---
1444412_at 1.03248 Absent in melanoma 1 Aim1
1444413_at 1.18381 "adaptor-related protein complex 3, sigma 2 subunit" Ap3s2
1444414_at 1.04766 Apolipoprotein D Apod

1444415_at 1.02893 "SWI/SNF related, matrix associated, actin dependent regulator of chromatin, subfamily d, member 3" Smarcd3
1444416_at 1.29019 Centromere autoantigen A Cenpa
1444417_at 1.00074 RIKEN cDNA 9330101J02 gene /// similar to Protein C10orf93 /// similar to Protein C10orf93 9330101J02Rik /// LOC664857 /// LOC672120
1444418_at 0.86038 "Inositol 1,4,5-triphosphate receptor 2" Itpr2
1444419_at 1.04047 prolylcarboxypeptidase (angiotensinase C) Prcp
1444420_at 0.97007 RIKEN cDNA E230032D23 gene E230032D23Rik
1444421_at 0.87179 ethanol decreased 2 Etohd2
1444422_at 1.10063 protocadherin 19 Pcdh19
1444423_at 1.03934 Transcribed locus ---
1444424_at 1.12880 Ubiquitin-like 1 (sentrin) activating enzyme E1A Uble1a
1444425_at 0.84728 Zinc finger protein 53 Zfp53
1444426_at 1.13780 RIKEN cDNA F730031O20 gene F730031O20Rik
1444427_at 0.97103 RIKEN cDNA D930038D03 gene D930038D03Rik
1444428_at 0.93133 --- ---
1444429_at 0.85990 leucine-rich repeats and transmembrane domains 1 Lrtm1
1444430_at 1.37367 Armadillo repeat containing 8 Armc8
1444431_at 0.95767 RCSD domain containing 1 Rcsd1
1444432_at 0.93289 RIKEN cDNA D330040H18 gene D330040H18Rik
1444433_at 1.14312 AU RNA binding protein/enoyl-coenzyme A hydratase Auh
1444434_at 0.96302 WD repeat domain 33 Wdr33
1444435_at 0.93708 vacuolar protein sorting 13B (yeast) /// similar to vacuolar protein sorting 13B isoform 4 /// region containing RIKEN cDNA C330002D13 gene; RIKEN cDNA 2310042E16 gene; vacuolar protein sorting 13B (yeast) Vps13b /// LOC666173 /// LOC669978
1444436_at 0.88006 heat shock factor 2 /// RIKEN cDNA 9630030I15 gene Hsf2 /// 9630030I15Rik
1444437_at 0.99808 ubiquitin specific peptidase 34 Usp34
1444438_at 1.08657 calcium and integrin binding family member 3 Cib3
1444439_at 1.02117 Bri3 binding protein Bri3bp
1444440_at 0.86435 RIKEN cDNA 4121402D02 gene 4121402D02Rik
1444441_at 1.11925 "PREDICTED: Mus musculus hypothetical protein LOC629847 (LOC629847), mRNA" ---
1444442_at 0.85629 oral cancer overexpressed 1 Oraov1
1444443_at 0.87364 RIKEN cDNA A830039H10 gene A830039H10Rik
1444444_at 0.99053 "Adult male medulla oblongata cDNA, RIKEN full-length enriched library, clone:6330524O11 product:unclassifiable, full insert sequence" ---
1444445_at 1.46207 expressed sequence C77648 C77648
1444446_at 1.54960 --- ---
1444447_at 1.50283 NIMA (never in mitosis gene a)-related expressed kinase 7 Nek7
1444448_at 0.92780 RIKEN cDNA 1300007F04 gene 1300007F04Rik
1444449_at 1.12934 Ubiquitin specific peptidase 18 Usp18
1444450_at 1.03982 RIKEN cDNA 9530096D07 gene 9530096D07Rik

1444451_at	0.89870	pappalysin 2	Pappa2	
1444452_at	1.07455	RIKEN cDNA D930002L09	gene	D930002L09Rik
1444453_at	0.95649	Trp53 regulating kinase	Trp53rk	
1444454_at	1.02184	Retinoblastoma 1	Rb1	
1444455_at	0.97445	"calcium channel, voltage-dependent, P/Q type, alpha 1A subunit"		Cacna1a
1444456_at	1.20914	RIKEN cDNA 9030425P06	gene	9030425P06Rik
1444457_at	1.02389	---	---	
1444458_at	1.39477	Meningioma expressed antigen 6 (coiled-coil proline-rich)		Mgea6
1444459_at	1.19240	---	---	
1444460_at	0.96674	ubiquitin-conjugating enzyme E2R 2	Ube2r2	
1444461_at	0.90901	"Zinc finger, MYND domain containing 11"		Zmynd11
1444462_at	1.01233	"0 day neonate thymus cDNA, RIKEN full-length enriched library, clone: A430098J12 product:unclassifiable, full insert sequence"	---	
1444463_at	1.03185	Rho GTPase activating protein 12	Arhgap12	
1444464_at	0.97825	NADH dehydrogenase (ubiquinone) Fe-S protein 4	Ndufs4	
1444465_at	1.12799	"Ubiquitination factor E4B, UFD2 homolog (S. cerevisiae)"		Ube4b
1444466_at	1.16966	Neurocalcin delta	Ncald	
1444467_at	1.01210	expressed sequence C77534	C77534	
1444468_at	0.85196	progesterin and adipoQ receptor family member VIII		Paqr8
1444469_at	0.90451	RIKEN cDNA 1110059H15	gene	1110059H15Rik
1444470_x_at	0.86663	Bone morphogenetic protein 5	Bmp5	
1444471_at	0.91038	RIKEN cDNA 4432416J03	gene	4432416J03Rik
1444472_at	1.24014	SNF1-like kinase 2	Snf1lk2	
1444473_at	1.98550	P300/CBP-associated factor	Pcaf	
1444474_at	0.95353	RIKEN cDNA C630028L02	gene	C630028L02Rik
1444475_at	0.96224	---	---	
1444476_at	1.03042	RIKEN cDNA D030007L05	gene	D030007L05Rik
1444477_at	1.02738	Leucine rich repeat transmembrane neuronal 3		Lrrtm3
1444478_at	0.82957	RIKEN cDNA 2900057D21	gene	2900057D21Rik
1444479_at	0.74907	---	---	
1444480_at	0.87843	---	---	
1444481_at	1.11900	---	---	
1444482_at	1.27655	RIKEN cDNA A130078K24	gene	A130078K24Rik
1444483_at	1.16065	RIKEN cDNA 4931428D14	gene	4931428D14Rik
1444484_at	1.00880	RIKEN cDNA A130009E19	gene	A130009E19Rik
1444485_at	1.57703	SHQ1 homolog (S. cerevisiae)	Shq1	
1444486_at	0.92763	kelch-like 5 (Drosophila)	Klh15	
1444487_at	1.18750	lecithin-retinol acyltransferase (phosphatidylcholine-retinol-O-acyltransferase)	Lrat	
1444488_at	0.86548	Ca ²⁺ dependent activator protein for secretion	Cadps	
1444489_at	0.97730	"Solute carrier family 25 (mitochondrial carrier, Aralar), member 12"		Slc25a12
1444490_at	1.33980	---	---	
1444491_at	0.93608	"Transcribed locus, strongly similar to XP_344663.2 PREDICTED: neuregulin		

2 [Rattus norvegicus]" ---

1444492_at 0.98206 "protein tyrosine phosphatase, receptor type, D" Ptprd
1444493_at 1.10602 "TBC1 domain family, member 2B" Tbc1d2b
1444494_at 0.99611 kelch repeat and BTB (POZ) domain containing 10 Kbtbd10
1444495_at 1.03262 --- ---
1444496_at 0.91374 RIKEN cDNA 2600011C06 gene 2600011C06Rik
1444497_at 1.06884 cDNA sequence BC059050 BC059050
1444498_at 1.10715 --- ---
1444499_at 0.95295 Transcribed locus ---
1444500_at 1.09229 "AHA1, activator of heat shock protein ATPase homolog 1 (yeast)" Ahsa1
1444501_at 1.31865 expressed sequence AI462521 AI462521
1444502_at 0.87606 PET112-like (yeast) Pet112l
1444503_at 1.18717 Glioblastoma amplified sequence Gbas
1444504_at 1.01882 dehydrogenase/reductase (SDR family) member 7C Dhrr7c
1444505_at 0.73590 retinoic acid induced 17 Rai17
1444506_at 1.54435 Translocating chain-associating membrane protein 1 Tram1
1444507_at 1.34085 ubiquitin specific peptidase 53 Usp53
1444508_s_at 0.98130 arrestin domain containing 1 Arrdc1
1444509_at 1.15090 Transcribed locus ---
1444510_at 0.93946 Transcribed locus ---
1444511_at 0.96692 "solute carrier family 41, member 1" Slc41a1
1444512_at 0.85861 Rho GTPase activating protein 29 Arhgap29
1444513_at 0.94666 connector enhancer of kinase suppressor of Ras 2 Cnksr2
1444514_at 0.96507 RIKEN cDNA B930096F20 gene B930096F20Rik
1444515_at 1.76624 Myosin ID Myo1d
1444516_at 1.11871 "Protein tyrosine phosphatase, receptor-type, F interacting protein, binding protein 2" Ppfibp2
1444517_at 0.96924 --- ---
1444518_at 1.21085 "Acyl-Coenzyme A oxidase 1, palmitoyl" Acox1
1444519_at 0.86778 leucine rich repeat containing G protein coupled receptor 5 Lgr5
1444520_at 1.19882 furry homolog-like (Drosophila) Fryl
1444521_at 0.96544 Transcribed locus ---
1444522_at 0.89209 Transcribed locus ---
1444523_s_at 1.01499 ubiquitin-conjugating enzyme E2 variant 1 /// similar to ubiquitin-conjugating enzyme E2 variant 1 /// similar to ubiquitin-conjugating enzyme E2 variant 1 Ube2v1 /// LOC545332 /// LOC666853
1444524_at 1.07205 "Activated spleen cDNA, RIKEN full-length enriched library, clone: F830206P12 product:hypothetical Cytochrome c heme-binding site containing protein, full insert sequence" ---
1444525_at 0.96381 "EGF, latrophilin seven transmembrane domain containing 1" Eltd1
1444526_at 0.83609 "signal sequence receptor, alpha" Ssr1
1444527_at 1.02426 Runt related transcription factor 1 Runx1
1444528_at 0.91445 --- ---

1444529_at 0.91820 similar to X-linked eukaryotic translation initiation factor 1A LOC666806
1444530_at 1.48434 Transcribed locus ---
1444531_at 0.86578 "Superoxide dismutase 2, mitochondrial" Sod2
1444532_at 1.27377 --- ---
1444533_at 1.07641 "CUG triplet repeat, RNA binding protein 2" Cugbp2
1444534_at 0.94865 similar to CG3950-PA /// similar to CG3950-PA LOC666801 /// LOC669012
1444535_at 0.94329 "Adult male urinary bladder cDNA, RIKEN full-length enriched library, clone:9530001M05 product:unclassifiable, full insert sequence /// Osteoclast-like cell cDNA, RIKEN full-length enriched library, clone:I420017N03 product:unclassifiable, full insert sequence" ---
1444536_at 0.88339 expressed sequence AI462171 AI462171
1444537_at 0.90079 expressed sequence AI429363 AI429363
1444538_at 1.89582 Expressed sequence AL033314 AL033314
1444539_at 0.92209 RIKEN cDNA D530039A21 gene D530039A21Rik
1444540_at 1.06451 --- ---
1444541_at 1.46578 Myosin IXa Myo9a
1444542_at 0.90863 "Adult male aorta and vein cDNA, RIKEN full-length enriched library, clone:A530017F16 product:unclassifiable, full insert sequence" ---
1444543_at 1.01500 Transcribed locus ---
1444544_at 1.02113 "DNA primase, p49 subunit" Prim1
1444545_at 1.02233 Erythrocyte protein band 4.1 Epb4.1
1444546_at 1.04456 cDNA sequence BC027057 BC027057
1444547_at 1.03113 --- ---
1444548_at 1.04176 RIKEN cDNA 1600025M17 gene 1600025M17Rik
1444549_at 0.98499 "Adult male adrenal gland cDNA, RIKEN full-length enriched library, clone:7330409L06 product:unclassifiable, full insert sequence" ---
1444550_at 0.75842 RIKEN cDNA 1110021J02 gene 1110021J02Rik
1444551_at 1.15211 "Inositol 1,4,5-triphosphate receptor 2" Itpr2
1444552_at 1.40314 --- ---
1444553_at 0.97835 Postmeiotic segregation increased 1 (S. cerevisiae) Pms1
1444554_at 1.08673 Trinucleotide repeat containing 6b Tnrc6b
1444555_at 1.06031 Checkpoint suppressor 1 Ches1
1444556_at 1.14896 Schlafen 1 Slfn1
1444557_at 1.86361 SMT3 suppressor of mif two 3 homolog 1 (yeast) Sumo1
1444558_at 0.92134 WD repeat domain 33 Wdr33
1444559_at 0.89832 Putative homeodomain transcription factor 2 Phtf2
1444560_at 0.77909 Transcribed locus ---
1444561_at 1.06335 Transmembrane protein 39a Tmem39a
1444562_at 1.50194 Ubiquitin protein ligase E3C Ube3c
1444563_at 1.01834 "UDP-Gal:betaGlcNAc beta 1,4-galactosyltransferase, polypeptide 5" B4galt5
1444564_at 1.06495 apolipoprotein D Apod
1444565_at 1.01899 expressed sequence BB166591 BB166591
1444566_at 0.96612 "DnaJ (Hsp40) related, subfamily B, member 13" Dnajb13

1444567_at 1.02604 RIKEN cDNA 1700066J03 gene 1700066J03Rik
1444568_at 0.96935 "Actin related protein 2/3 complex, subunit 5" Arpc5
1444569_at 1.12787 hypothetical protein LOC666752 /// hypothetical protein LOC669447 ///
hypothetical protein LOC677552 LOC666752 /// LOC669447 /// LOC677552
1444570_at 1.17076 Craniofacial development protein 1 Cfdp1
1444571_at 0.94475 "Zinc finger, SWIM domain containing 3" Zswim3
1444572_at 0.95176 monocyte to macrophage differentiation-associated 2 Mmd2
1444573_at 0.99755 Lemur tyrosine kinase 3 Lmtk3
1444574_at 0.95156 E74-like factor 2 Elf2
1444575_at 1.21601 RIKEN cDNA 1500002O20 gene 1500002O20Rik
1444576_at 1.07745 Nuclear mitotic apparatus protein 1 Numa1
1444577_x_at 1.31085 Nuclear mitotic apparatus protein 1 Numa1
1444578_at 1.12450 Speckle-type POZ protein Spop
1444579_at 1.04124 GATA zinc finger domain containing 2B Gatad2b
1444580_at 0.96854 Transcribed locus ---
1444581_at 0.93403 RAN GTPase activating protein 1 Rangap1
1444582_at 1.05350 "DCN1, defective in cullin neddylation 1, domain containing 5 (S. cerevisiae)"
Dcun1d5
1444583_at 2.11756 "DNA segment, Chr 4, ERATO Doi 767, expressed" D4Ertd767e
1444584_at 1.02611 piggyBac transposable element derived 1 (Pgbd1) pseudogene /// RIKEN cDNA
4921509E05 gene LOC238535 /// 4921509E05Rik
1444585_at 0.89826 arginine decarboxylase Adc
1444586_at 0.99332 Ras association (RalGDS/AF-6) and pleckstrin homology domains 1 Raph1
1444587_at 1.00971 expressed sequence AI481207 AI481207
1444588_at 1.08722 --- ---
1444589_at 1.01752 similar to reduced expression 2 LOC240038
1444590_at 0.99851 DDHD domain containing 2 Ddhd2
1444591_at 0.94338 "Phosphatidylinositol 3-kinase, regulatory subunit, polypeptide 1 (p85 alpha)"
Pik3r1
1444592_at 1.04921 O-sialoglycoprotein endopeptidase Osgep
1444593_at 1.02593 PH domain and leucine rich repeat protein phosphatase-like Phlpp1
1444594_at 1.08659 T-box 3 Tbx3
1444595_at 0.99775 "zinc finger, X-linked, duplicated A /// similar to Zinc finger X-linked protein
ZXDA" Zxda /// LOC668171
1444596_at 1.09106 paired box gene 7 Pax7
1444597_at 1.02389 Transcribed locus ---
1444598_at 1.30143 Ets variant gene 6 (TEL oncogene) Etv6
1444599_at 1.12483 --- ---
1444600_at 0.99764 nuclear receptor binding protein Nrpb
1444601_at 1.12681 --- ---
1444602_at 0.92975 "Transcribed locus, weakly similar to XP_343428.2 PREDICTED: neural
precursor cell expressed, developmentally down-regulated gene 4A [Rattus norvegicus]" ---
1444603_at 0.93535 RIKEN cDNA 2810030D12 gene 2810030D12Rik

1444604_at 0.88348 Enhancer of zeste homolog 1 (Drosophila) Ezh1
1444605_at 0.95379 RIKEN cDNA 1700061F12 gene 1700061F12Rik
1444606_at 0.89233 ephrin A2 Efna2
1444607_at 0.82348 Corneodesmosin Cdsn
1444608_at 1.01832 Ankyrin repeat domain 17 Ankrd17
1444609_at 1.57652 Exocyst complex component 4 Exoc4
1444610_at 0.95622 Transcribed locus ---
1444611_at 0.98305 "U2 small nuclear ribonucleoprotein auxiliary factor (U2AF) 1, related sequence 2" U2af1-rs2
1444612_at 0.88741 RIKEN cDNA 3222402P14 gene 3222402P14Rik
1444613_at 1.07793 Transcribed locus ---
1444614_x_at 0.94178 RIKEN cDNA 6430537H07 gene 6430537H07Rik
1444615_x_at 1.02345 "runt-related transcription factor 1; translocated to, 1 (cyclin D-related)" Runx1t1
1444616_x_at 1.06051 Transcribed locus ---
1444617_at 0.97325 Deleted in colorectal carcinoma Dcc
1444618_at 1.05857 RNA binding motif protein 5 Rbm5
1444619_x_at 1.00070 "proteasome (prosome, macropain) subunit, beta type 8 (large multifunctional peptidase 7)" Psmb8
1444620_at 1.01439 Transcription factor 12 Tcf12
1444621_at 0.86743 IQ motif and Sec7 domain 2 Iqsec2
1444622_at 0.98033 "Protein tyrosine phosphatase, receptor type, D" Ptprd
1444623_at 1.01637 RIKEN cDNA E530011L22 gene E530011L22Rik
1444624_at 1.06824 ubiquitin-associated protein 1 Ubap1
1444625_at 1.10643 Hypothetical LOC545822 LOC545822
1444626_at 1.10581 hypothetical protein D930001B02 D930001B02
1444627_at 1.02068 "Purinergic receptor P2Y, G-protein coupled, 5" P2ry5
1444628_at 0.96017 a disintegrin and metallopeptidase domain 33 Adam33
1444629_at 0.94738 "amyotrophic lateral sclerosis 2 (juvenile) chromosome region, candidate 13 (human)" Als2cr13
1444630_at 0.94922 "DNA segment, Chr 2, ERATO Doi 127, expressed" D2Ertd127e
1444631_at 0.99205 Zinc finger protein 410 Zfp410
1444632_at 1.02151 cDNA sequence BC064078 BC064078
1444633_at 0.97424 --- ---
1444634_at 0.92443 NOL1/NOP2/Sun domain family 3 Nsun3
1444635_at 1.06623 Transcribed locus ---
1444636_at 0.98106 RIKEN cDNA 4930592I03 gene 4930592I03Rik
1444637_at 0.83558 --- ---
1444638_at 0.97782 titin Ttn
1444639_at 1.14794 Nephronophthisis 4 (juvenile) homolog (human) Nphp4
1444640_at 0.95122 "phosphatidylinositol glycan anchor biosynthesis, class G" Pigg
1444641_at 0.93070 Adenylate cyclase 3 Adcy3
1444642_at 0.96886 Transcribed locus ---

1444643_at 1.00600 Transcribed locus ---
1444644_x_at 1.13878 TatD DNase domain containing 1 Tatdn1
1444645_at 2.55292 SERTA domain containing 2 Sertad2
1444646_at 0.98499 RIKEN cDNA 8430420F16 gene 8430420F16Rik
1444647_at 1.15773 "phospholipase A2, activating protein" Plaa
1444648_at 0.93913 cDNA sequence BC050210 BC050210
1444649_at 1.07275 "Protein kinase C, epsilon" Prkce
1444650_at 1.33362 rho/rac guanine nucleotide exchange factor (GEF) 18 Arhgef18
1444651_at 1.52182 hypothetical LOC553089 LOC553089
1444652_at 0.91105 RIKEN cDNA B830004H01 gene B830004H01Rik
1444653_at 0.94813 "10 days neonate skin cDNA, RIKEN full-length enriched library,
clone:4732412F15 product:unclassifiable, full insert sequence" ---
1444654_at 1.23134 Transcriptional regulating factor 1 Trefl
1444655_at 1.02232 --- ---
1444656_at 1.06976 cDNA sequence BC030499 BC030499
1444657_at 0.76147 Bcl3 binding protein B3bp
1444658_at 0.99959 --- ---
1444659_at 1.04503 Thyroid hormone receptor interactor 11 Trip11
1444660_at 1.45884 RIKEN cDNA 4932438A13 gene 4932438A13Rik
1444661_at 1.00541 G protein-coupled receptor 26 Gpr26
1444662_at 1.04243 "phospholipase A2, group IVC (cytosolic, calcium-independent)" Pla2g4c
1444663_at 1.40386 Vascular endothelial zinc finger 1 Vezf1
1444664_at 0.88050 "Immunoglobulin superfamily, member 4" Igsf4d
1444665_at 0.97477 RIKEN cDNA 4930402E16 gene 4930402E16Rik
1444666_at 1.05549 brix domain containing 1 Bxdc1
1444667_at 1.05093 "bromodomain, testis-specific" Brdt
1444668_at 1.02723 "Adult male testis cDNA, RIKEN full-length enriched library,
clone:4930534N17 product:unclassifiable, full insert sequence" ---
1444669_at 0.97492 leucine rich repeat and fibronectin type III domain containing 1 Lrfn1
1444670_at 2.19602 RIKEN cDNA A130004G11 gene A130004G11Rik
1444671_at 0.92686 RAS protein activator like 2 Rasal2
1444672_at 0.90030 Transcribed locus ---
1444673_at 0.99229 RIKEN cDNA E030031F02 gene E030031F02Rik
1444674_at 1.08422 expressed sequence AU015680 AU015680
1444675_at 0.96060 expressed sequence AL023051 AL023051
1444676_at 1.21598 --- ---
1444677_at 1.96184 expressed sequence C77673 C77673
1444678_at 1.09453 --- ---
1444679_at 1.31919 PHD finger protein 21A Phf21a
1444680_at 0.93100 "Positive cofactor 2, multiprotein complex, glutamine/Q-rich-associated
protein" Pcqap
1444681_at 0.86605 "DNA segment, Chr 14, ERATO Doi 171, expressed" D14Ertd171e
1444682_at 0.87916 cDNA Sequence BC037032 BC037032

1444683_at 0.99755 --- ---
1444684_at 0.95518 RIKEN cDNA 8030475D13 gene 8030475D13Rik
1444685_at 0.87650 --- ---
1444686_at 0.93159 RIKEN cDNA A930016P21 gene A930016P21Rik
1444687_at 0.97547 "complement component 1, q subcomponent-like 2" C1ql2
1444688_at 1.10618 "positive cofactor 2, multiprotein complex, glutamine/Q-rich-associated protein" Pcqap
1444689_at 1.16695 WD repeat domain 67 Wdr67
1444690_at 0.82461 Eph receptor A5 Epha5
1444691_at 1.02616 "protein kinase C, nu" Prkc
1444692_at 0.97322 EST AI316844 AI316844
1444693_at 2.20405 "Calcium channel, voltage-dependent, beta 2 subunit" Cacnb2
1444694_at 1.39126 Tuftelin 1 Tuft1
1444695_at 1.02970 "Adult male colon cDNA, RIKEN full-length enriched library, clone:9030003C19 product:unclassifiable, full insert sequence" ---
1444696_at 0.92838 CDNA sequence BC059841 BC059841
1444697_at 1.12583 RIKEN cDNA 4732490B19 gene 4732490B19Rik
1444698_at 0.99708 Quaking Qk
1444699_at 1.14354 splicing factor 4 Sf4
1444700_at 0.88313 Neurexin III Nrxn3
1444701_at 1.64682 RIKEN cDNA A130001D14 gene A130001D14Rik
1444702_at 1.02652 "Adult male epididymis cDNA, RIKEN full-length enriched library, clone:9230116A06 product:unclassifiable, full insert sequence" ---
1444703_at 0.92319 RIKEN cDNA 2810403D21 gene 2810403D21Rik
1444704_at 0.92453 "7 days neonate cerebellum cDNA, RIKEN full-length enriched library, clone:A730050L22 product:unclassifiable, full insert sequence" ---
1444705_at 1.65599 Amyloid beta (A4) precursor protein App
1444706_at 1.10272 RIKEN cDNA E430014L09 gene E430014L09Rik
1444707_at 0.97950 Transcribed locus ---
1444708_at 0.98464 transmembrane protein 29 Tmem29
1444709_at 1.22965 inversin Invs
1444710_at 1.03061 "Adult male epididymis cDNA, RIKEN full-length enriched library, clone:9230020K05 product:unclassifiable, full insert sequence" ---
1444711_at 0.82799 ankyrin repeat domain 15 Ankrd15
1444712_at 1.05443 "gene model 739, (NCBI) /// hypothetical protein LOC672930" Gm739 /// LOC672930
1444713_at 1.11159 --- ---
1444714_at 1.01066 RIKEN cDNA 2510006D16 gene 2510006D16Rik
1444715_at 1.23018 Transcribed locus ---
1444716_at 1.76886 --- ---
1444717_at 1.10385 ZW10 interactor Zwint
1444718_at 1.08937 triggering receptor expressed on myeloid cells-like 2 Trem12
1444719_at 0.89467 "DNA Segment, Chr 7, Mouse Genome Informatics 29" D7Mgi29

1444720_at	1.25608	expressed sequence AA408456	AA408456
1444721_at	0.89479	---	---
1444722_at	1.70662	---	---
1444723_at	0.92152	RIKEN cDNA 6530418L21 gene	6530418L21Rik
1444724_at	1.02069	Protocadherin 9 Pcdh9	
1444725_at	0.95277	microtubule associated serine/threonine kinase-like	Mastl
1444726_at	0.89565	"KRR1, small subunit (SSU) processome component, homolog (yeast)"	Krr1
1444727_at	0.90267	---	---
1444728_at	2.81317	Exocyst complex component 4	Exoc4
1444729_at	1.07760	RIKEN cDNA E130317F20 gene	E130317F20Rik
1444730_at	0.86548	tousled-like kinase 1	Tlk1
1444731_at	1.03380	"phosphodiesterase 4B, cAMP specific"	Pde4b
1444732_at	1.03040	---	---
1444733_at	1.02086	Transcribed locus	---
1444734_at	0.92511	RIKEN cDNA A330001L22 gene	A330001L22Rik
1444735_at	1.21642	---	---
1444736_at	1.09180	"cadherin 7, type 2"	Cdh7
1444737_at	1.03015	"DNA segment, Chr 2, Wayne State University 107, expressed"	D2Wsu107e
1444738_at	1.20726	RIKEN cDNA 5730419I09 gene	5730419I09Rik
1444739_at	1.04916	"Polymerase (DNA-directed), epsilon 4 (p12 subunit)"	Pole4
1444740_at	1.00526	RIKEN cDNA 4930550L11 gene	4930550L11Rik
1444741_at	0.93374	---	---
1444742_at	0.94790	Ankyrin repeat and SOCS box-containing protein 14	Asb14
1444743_at	1.02461	"0 day neonate cerebellum cDNA, RIKEN full-length enriched library, clone: C230024F22 product:unclassifiable, full insert sequence"	---
1444744_at	0.97531	Death inducer-obliterator 1	Dido1
1444745_at	1.00884	Diaphanous homolog 2 (Drosophila)	Diap2
1444746_at	1.48165	Polypyrimidine tract binding protein 2	Ptbp2
1444747_at	1.13296	Protein-L-isoaspartate (D-aspartate) O-methyltransferase domain containing 1	Pcmt1
1444748_at	1.01958	Paraspeckle protein 1	Pspc1
1444749_at	1.04763	Apolipoprotein D	Apod
1444750_at	1.40836	Restin (Reed-Steinberg cell-expressed intermediate filament-associated protein)	Rsn
1444751_at	0.96369	"POU domain, class 6, transcription factor 1"	Pou6f1
1444752_at	0.85672	RIKEN cDNA C230073G13 gene	C230073G13Rik
1444753_at	1.58350	NIMA (never in mitosis gene a)-related expressed kinase 7	Nek7
1444754_at	1.00931	additional sex combs like 2 (Drosophila)	Asxl2
1444755_at	1.01441	mitotic arrest deficient 1-like 1	Mad111
1444756_at	0.92481	Transcribed locus	---
1444757_at	1.09668	Eukaryotic translation elongation factor 1 epsilon 1	Eef1e1
1444758_at	1.25496	Tnf receptor-associated factor 3	Traf3
1444759_at	0.91553	prickle like 1 (Drosophila)	Prickle1

1444760_at	1.03552	nuclear receptor co-repressor 1	Ncor1
1444761_at	0.98807	Coiled-coil domain containing 8	Ccdc8
1444762_at	1.76525	"Protein phosphatase 1, regulatory (inhibitor) subunit 12A"	Ppp1r12a
1444763_at	1.14962	"protein tyrosine phosphatase, receptor type, K"	Ptprk
1444764_at	1.45680	RIKEN cDNA A130022J21 gene	A130022J21Rik
1444765_at	1.52073	RNA binding protein gene with multiple splicing	Rbpms
1444766_at	1.14600	ataxin 7-like 1	Atxn7l1
1444767_at	1.05769	"GNAS (guanine nucleotide binding protein, alpha stimulating) complex locus"	Gnas
1444768_at	0.95656	Transcribed locus	---
1444769_at	1.06716	Testis expressed gene 9	Tex9
1444770_at	1.04416	RIKEN cDNA 1700095A13 gene	1700095A13Rik
1444771_at	1.03792	ankyrin repeat and BTB (POZ) domain containing 2	Abtb2
1444772_at	1.07270	SET and MYND domain containing 3	Smyd3
1444773_at	1.01027	RIKEN cDNA D430040D24 gene	D430040D24Rik
1444774_at	1.02992	de-etiolated homolog 1 (Arabidopsis)	Det1
1444775_at	1.45447	RIKEN cDNA 9930033D15 gene	9930033D15Rik
1444776_at	0.95356	RIKEN cDNA 2700005E23 gene	2700005E23Rik
1444777_at	2.13902	Retinoic acid induced 14	Rai14
1444778_at	0.98569	septin 3	3-Sep
1444779_s_at	0.95353	zinc finger protein 59	Zfp59
1444780_at	0.96489	RIKEN cDNA 5330421F07 gene	5330421F07Rik
1444781_at	0.95674	XPA binding protein 1	Xab1
1444782_at	1.08533	RIKEN cDNA 4930417H01 gene	4930417H01Rik
1444783_at	1.04618	Abl-interactor 2	Abi2
1444784_at	0.85387	RIKEN cDNA 4930564K09 gene	4930564K09Rik
1444785_at	1.93930	Sloan-Kettering viral oncogene homolog	Ski
1444786_at	0.95881	nucleolar protein 3 (apoptosis repressor with CARD domain)	Nol3
1444787_at	1.24018	Chromodomain helicase DNA binding protein 1	Chd1
1444788_x_at	1.03613	Chromodomain helicase DNA binding protein 1	Chd1
1444789_at	0.84870	RIKEN cDNA 1700120C14 gene	1700120C14Rik
1444790_at	1.00357	RIKEN cDNA 1810005K13 gene	1810005K13Rik
1444791_at	0.93151	Purkinje cell protein 4-like 1	Pcp4l1
1444792_at	0.98038	---	---
1444793_at	0.83414	---	---
1444794_at	1.03902	---	---
1444795_at	0.97280	expressed sequence AU015723	AU015723
1444796_at	0.98530	expressed sequence C76876	C76876
1444797_at	0.95621	RIKEN cDNA 8030474K03 gene	8030474K03Rik
1444798_at	0.80126	---	---
1444799_at	1.12745	expressed sequence AU022054	AU022054
1444800_at	1.04941	Nuclear factor I/B	Nfib
1444801_at	0.96219	RIKEN cDNA 2900041M22 gene	2900041M22Rik

1444802_at	1.04542	SEC22 vesicle trafficking protein-like A (<i>S. cerevisiae</i>)	Sec22a
1444803_at	1.13064	"Zinc finger protein, multitype 2"	Zfpm2
1444804_at	1.02940	"DNA segment, Chr 17, ERATO Doi 96, expressed"	D17Ertd96e
1444805_at	0.95029	RIKEN cDNA 4930473A06 gene	4930473A06Rik
1444806_at	1.74801	RIKEN cDNA 6720456B07 gene	6720456B07Rik
1444807_at	1.00948	---	---
1444808_at	0.99352	"PRKC, apoptosis, WT1, regulator"	Pawr
1444809_at	0.99690	MARVEL (membrane-associating) domain containing 3	Marveld3
1444810_at	2.43483	Acetyl-Coenzyme A carboxylase alpha	Acaca
1444811_at	1.67817	translocation protein 1	Tloc1
1444812_at	0.87560	PR domain containing 10	Prdm10
1444813_at	1.07009	Microtubule associated serine/threonine kinase 2	Mast2
1444814_at	1.41692	"DNA segment, Chr 15, ERATO Doi 180, expressed"	D15Ertd180e
1444815_at	0.94603	expressed sequence C85328	C85328
1444816_at	0.96386	---	---
1444817_at	1.13951	"Pleckstrin homology domain containing, family H (with MyTH4 domain) member 2"	Plekhh2
1444818_at	1.09665	---	---
1444819_at	0.89945	expressed sequence C86090	C86090
1444820_at	0.95556	Transmembrane protein 49	Tmem49
1444821_at	0.88895	"DNA segment, Chr 8, ERATO Doi 82, expressed"	D8Ertd82e
1444822_at	1.00506	"Sarcoglycan, delta (dystrophin-associated glycoprotein)"	Sgcd
1444823_at	1.03506	---	---
1444824_at	1.12063	expressed sequence C76793	C76793
1444825_at	1.01098	---	---
1444826_at	0.96668	---	---
1444827_at	2.20500	BTB (POZ) domain containing 9	Btbd9
1444828_at	1.30615	"Protein phosphatase 2, regulatory subunit B (B56), gamma isoform"	Ppp2r5c
1444829_at	0.86393	expressed sequence C78532	C78532
1444830_at	1.00146	Transcribed locus	---
1444831_at	0.91786	Serologically defined colon cancer antigen 8	Sdccag8
1444832_at	0.97538	Huntingtin interacting protein 1 related	Hip1r
1444833_at	0.85962	RIKEN cDNA 2410012H22 gene	2410012H22Rik
1444834_at	0.96959	"Protein phosphatase 1, regulatory (inhibitor) subunit 12A"	Ppp1r12a
1444835_at	0.99850	cDNA sequence BC030499	BC030499
1444836_at	0.89299	DEK oncogene (DNA binding)	Dek
1444837_at	0.93222	---	---
1444838_at	0.91278	hypothetical protein C630010N09	C630010N09
1444839_at	1.00460	---	---
1444840_at	0.99890	---	---
1444841_at	1.01290	---	---
1444842_at	0.97209	"jumonji, AT rich interactive domain 1A (Rbp2 like)"	Jarid1a
1444843_at	0.97993	---	---

1444844_at	1.00777	"fatty acid binding protein 3, muscle and heart"	Fabp3
1444845_at	1.92274	---	---
1444846_at	1.00492	"DNA segment, Chr 15, ERATO Doi 529, expressed"	D15Ert529e
1444847_at	0.84938	---	---
1444848_at	1.13610	R3H domain 1 (binds single-stranded nucleic acids)	R3hdm1
1444849_at	0.94878	expressed sequence C76872	C76872
1444850_at	0.91456	hepatitis A virus cellular receptor 1-like	RP23-248K2.4
1444851_at	1.00184	Zinc finger protein 532	Zfp532
1444852_at	0.93288	Flavin containing monooxygenase 4	Fmo4
1444853_at	0.99894	---	---
1444854_at	0.94699	"PREDICTED: Mus musculus similar to gonadotropin inducible ovarian transcription factor 1 (LOC625996), mRNA"	---
1444855_at	1.29705	B-cell CLL/lymphoma 9-like	Bcl9l
1444856_at	1.59140	CREB binding protein	Crebbp
1444857_at	1.02049	expressed sequence C87312	C87312
1444858_at	0.98090	INO80 complex homolog 1 (S. cerevisiae)	Inoc1
1444859_at	0.92118	---	---
1444860_at	0.84512	"Laminin, alpha 3"	Lama3
1444861_at	0.91256	"DNA segment, Chr 13, ERATO Doi 297, expressed"	D13Ert297e
1444862_at	0.91696	RIKEN cDNA 1700112E06 gene	1700112E06Rik
1444863_at	0.87549	NHL repeat containing 2	Nhlrc2
1444864_at	0.97058	"DNA segment, Chr 12, ERATO Doi 125, expressed"	D12Ert125e
1444865_at	0.94368	---	---
1444866_at	0.95349	RIKEN cDNA E230006M18 gene	E230006M18Rik
1444867_at	0.95796	expressed sequence C76751	C76751
1444868_at	1.04375	RIKEN cDNA 4921529O18 gene	4921529O18Rik
1444869_at	1.05116	Gephyrin	Gphn
1444870_at	0.99537	expressed sequence AU015438	AU015438
1444871_at	0.99153	expressed sequence AU022697	AU022697
1444872_at	1.89835	"DNA segment, Chr 3, ERATO Doi 740, expressed"	D3Ert740e
1444873_at	1.01373	Huntingtin interacting protein 1	Hip1
1444874_at	0.95542	"ATP synthase, H ⁺ transporting, mitochondrial F0 complex, subunit c (subunit 9), isoform 1"	Atp5g1
1444875_at	1.17354	"Protein phosphatase 2 (formerly 2A), catalytic subunit, alpha isoform"	Ppp2ca
1444876_at	0.96959	RIKEN cDNA 2700049A03 gene	2700049A03Rik
1444877_at	0.99920	---	---
1444878_at	0.91880	Dedicator of cytokinesis 10	Dock10
1444879_at	0.95551	"PREDICTED: Mus musculus similar to long interspersed repetitive DNA sequence LINE4 (L1Rn) (LOC621812), mRNA"	---
1444880_at	0.95519	Transcribed locus	---
1444881_at	1.02411	TAO kinase 3	Taok3
1444882_at	1.09887	---	---
1444883_at	0.98192	transmembrane protein 19	Tmem19

1444884_at	1.02320	palmitoyl-protein thioesterase 1	Ppt1
1444885_at	0.82874	Transcribed locus	---
1444886_at	1.28051	Amyloid beta precursor protein (cytoplasmic tail) binding protein 2	Appbp2
1444887_at	0.94626	expressed sequence C79015	C79015
1444888_at	0.91404	expressed sequence AU022852	AU022852
1444889_at	0.96719	Ras association (RalGDS/AF-6) domain family 2	Rassf2
1444890_at	1.62356	Expressed sequence AA536749	AA536749
1444891_at	0.94284	---	---
1444892_at	0.89402	nucleoporin 160	Nup160
1444893_at	1.05777	---	---
1444894_at	0.93853	"Protein tyrosine phosphatase, non-receptor type 18"	Ptpn18
1444895_at	0.85201	Junction adhesion molecule 3	Jam3
1444896_at	0.96045	---	---
1444897_at	1.03082	expressed sequence AU017263	AU017263
1444898_at	0.98339	---	---
1444899_at	1.03170	RIKEN cDNA D630037F22 gene	D630037F22Rik
1444900_at	1.00279	"DNA segment, Chr 18, ERATO Doi 289, expressed"	D18Ertd289e
1444901_at	0.90980	---	---
1444902_at	1.01575	F-box and leucine-rich repeat protein 10	Fbxl10
1444903_at	1.03520	Membrane-associated ring finger (C3HC4) 6	6-Mar
1444904_at	0.94872	"Runt-related transcription factor 1; translocated to, 1 (cyclin D-related)"	Runx1t1
1444905_at	0.99169	"DNA segment, Chr 1, ERATO Doi 705, expressed"	D1Ertd705e
1444906_at	0.90726	latrophilin 2	Lphn2
1444907_at	1.07643	expressed sequence AU022138	AU022138
1444908_at	0.96203	hyaluronic acid binding protein 4 /// similar to hyaluronic acid binding protein 4	Habp4 /// LOC671769
1444909_at	1.09882	Transcribed locus	---
1444910_at	1.01097	---	---
1444911_at	1.01812	RIKEN cDNA 9030605H24 gene	9030605H24Rik
1444912_at	1.00764	---	---
1444913_at	1.14442	Hedgehog acyltransferase	Hhat
1444914_at	1.00137	---	---
1444915_at	0.99991	"DNA segment, Chr 1, ERATO Doi 399, expressed"	D1Ertd399e
1444916_at	1.18417	"PREDICTED: Mus musculus similar to Kinesin-Like Protein family member (klp-6) (LOC383592), mRNA"	---
1444917_at	1.01331	Ubiquitin-conjugating enzyme E2W (putative)	Ube2w
1444918_at	0.94462	---	---
1444919_at	1.07749	---	---
1444920_at	1.05057	"DNA segment, Chr 19, ERATO Doi 386, expressed"	D19Ertd386e
1444921_at	0.99279	"DNA segment, Chr 15, ERATO Doi 466, expressed"	D15Ertd466e
1444922_at	0.96272	"PREDICTED: Mus musculus similar to microtubule-associated protein, RP/EB family, member 3 (LOC620153), mRNA"	---

1444923_at 0.91056 "Adult male diencephalon cDNA, RIKEN full-length enriched library, clone:9330152C01 product:unclassifiable, full insert sequence" ---
1444924_at 0.93829 --- ---
1444925_at 0.89799 --- ---
1444926_at 0.93422 --- ---
1444927_at 1.00145 "DNA segment, Chr 11, ERATO Doi 729, expressed" D11Ertd729e
1444928_at 1.03502 "Solute carrier family 35, member F1" Slc35f1
1444929_at 1.03242 --- ---
1444930_at 1.00139 "Rabaptin, RAB GTPase binding effector protein 1" Rabep1
1444931_at 1.05357 Endothelial-specific receptor tyrosine kinase Tek
1444932_at 0.92669 --- ---
1444933_at 0.90425 "Transcribed locus, moderately similar to XP_231006.3 PREDICTED: similar to formin [Rattus norvegicus]" ---
1444934_at 1.18694 zinc finger protein (C2H2 type) 276 Zfp276
1444935_at 0.82759 --- ---
1444936_at 0.97877 "DNA segment, Chr 5, ERATO Doi 566, expressed" D5Ertd566e
1444937_at 1.02381 "AF4/FMR2 family, member 1" Aff1
1444938_at 0.78935 Transcribed locus ---
1444939_at 1.00521 --- ---
1444940_at 0.92762 expressed sequence C76411 C76411
1444941_at 0.97684 --- ---
1444942_at 0.94873 helicase with zinc finger domain Helz
1444943_at 1.38213 SH3 domain containing ring finger 1 Sh3rf1
1444944_at 0.89091 --- ---
1444945_at 0.91846 expressed sequence AU024605 AU024605
1444946_at 1.07051 Transcribed locus ---
1444947_at 1.09788 LIM and SH3 protein 1 Lasp1
1444948_at 1.05304 --- ---
1444949_at 0.97994 --- ---
1444950_at 1.03211 zinc finger protein 509 Zfp509
1444951_at 0.90881 RIKEN cDNA 4930467M19 gene 4930467M19Rik
1444952_a_at 1.13351 nuclear casein kinase and cyclin-dependent kinase substrate 1 Nucks1
1444953_at 1.00997 Nuclear casein kinase and cyclin-dependent kinase substrate 1 Nucks1
1444954_at 1.04235 Cyclin-dependent kinase-like 4 Cdk14
1444955_at 0.88024 "DNA segment, Chr 6, ERATO Doi 469, expressed" D6Ertd469e
1444956_at 0.98262 Gamma-aminobutyric acid (GABA-A) receptor-associated protein-like 2 Gabarapl2
1444957_at 1.01603 Paraspeckle protein 1 Pspc1
1444958_at 1.12132 expressed sequence C79017 C79017
1444959_at 1.07158 "DNA segment, Chr 1, ERATO Doi 83, expressed" D1Ertd83e
1444960_at 1.05097 "cytochrome P450, family 2, subfamily u, polypeptide 1" Cyp2u1
1444961_at 0.93873 --- ---
1444962_at 1.07891 similar to component of Sp100-rs LOC665317

1444963_at	1.06908	---	---	
1444964_at	0.99894	"casein kinase 1, gamma 1"	Csnk1g1	
1444965_at	0.95168	RIKEN cDNA E130002L11 gene	E130002L11Rik	
1444966_at	0.94746	"Ribonucleoprotein, PTB-binding 1"	Raver1	
1444967_at	1.08414	Dystroglycan 1	Dag1	
1444968_at	0.96900	---	---	
1444969_at	1.09135	RIKEN cDNA 5033414K04 gene	5033414K04Rik	
1444970_at	0.91699	expressed sequence C79870	C79870	
1444971_at	1.10758	RNA binding motif protein 5	Rbm5	
1444972_at	1.10912	expressed sequence C77626	C77626	
1444973_at	0.99408	"Potassium large conductance calcium-activated channel, subfamily M, alpha member 1"	Kcnma1	
1444974_at	0.92119	expressed sequence AU023617	AU023617	
1444975_at	0.76097	Coiled-coil-helix-coiled-coil-helix domain containing 6	Chchd6	
1444976_at	1.01243	---	---	
1444977_at	1.00366	---	---	
1444978_at	0.92675	---	---	
1444979_at	1.06482	protein C	Proc	
1444980_at	1.09778	Hypothetical protein C730009D12	C730009D12	
1444981_at	0.90717	Putative neuronal cell adhesion molecule	Punc	
1444982_at	0.84805	"Ras homolog gene family, member J"	Rhoj	
1444983_at	0.86506	RIKEN cDNA 4933437K13 gene	4933437K13Rik	
1444984_at	0.83636	expressed sequence C77649	C77649	
1444985_at	1.04128	"Solute carrier family 8 (sodium/calcium exchanger), member 1"	Slc8a1	
1444986_at	0.95252	---	---	
1444987_at	1.12296	Cathepsin B	Ctsb	
1444988_at	1.01992	---	---	
1444989_at	1.02632	hypothetical protein LOC665914	/// hypothetical protein LOC671445	///
hypothetical protein LOC677108		LOC665914	/// LOC671445	/// LOC677108
1444990_at	1.00558	Human immunodeficiency virus type I enhancer binding protein 2	Hivep2	
1444991_at	0.89410	---	---	
1444992_at	1.25091	expressed sequence AI120166	AI120166	
1444993_at	0.98379	similar to mortality factor 4 like 1 (predicted)	/// similar to mortality factor 4 like 1 (predicted)	/// similar to mortality factor 4 like 1 (predicted) LOC664949
LOC675080		/// LOC671072	///	
1444994_at	0.97886	RIKEN cDNA A230038B01 gene	A230038B01Rik	
1444995_at	0.85318	"Chromodomain protein, Y chromosome-like 2"	Cdyl2	
1444996_at	1.13046	DEP domain containing 5	Depdc5	
1444997_at	0.96923	"PR domain containing 1, with ZNF domain"	Prdm1	
1444998_at	0.81735	Transcribed locus	---	
1444999_at	0.98347	expressed sequence C77651	C77651	
1445000_at	0.99047	---	---	
1445001_at	1.13278	---	---	

1445002_at	0.92904	---	---	
1445003_at	1.07713	"phenylalanine-tRNA synthetase-like, alpha subunit"	Farsla	
1445004_a_at	0.87589	RIKEN cDNA A330068P14 gene	A330068P14Rik	
1445005_at	0.92437	expressed sequence C79491	C79491	
1445006_at	1.02784	---	---	
1445007_at	1.02093	Leucine rich repeat containing 50	Lrrc50	
1445008_at	1.90574	Fibronectin type III domain containing 3B	Fndc3b	
1445009_at	1.12171	"Solute carrier family 9 (sodium/hydrogen exchanger), member 8"	Slc9a8	
1445010_at	0.95031	Transcribed locus	---	
1445011_at	0.96428	---	---	
1445012_at	0.96807	NIMA (never in mitosis gene a)-related expressed kinase 11	Nek11	
1445013_at	1.01924	expressed sequence C86098	C86098	
1445014_at	0.98467	expressed sequence AU019157	AU019157	
1445015_at	0.99452	---	---	
1445016_at	0.91292	---	---	
1445017_at	1.02882	"CCR4-NOT transcription complex, subunit 2"	Cnot2	
1445018_at	0.99648	Protein serine kinase H1	Pskh1	
1445019_at	1.03066	---	---	
1445020_at	0.90403	---	---	
1445021_at	0.95796	RIKEN cDNA 6030465E24 gene	6030465E24Rik	
1445022_at	1.07364	Transcribed locus	---	
1445023_at	0.89828	Transcribed locus	---	
1445024_at	1.03458	START domain containing 7	Stard7	
1445025_at	1.05376	expressed sequence AU015536	AU015536	
1445026_at	0.98254	Activin receptor IIA	Acvr2a	
1445027_at	1.02826	cerebellar degeneration-related protein 2-like	Cdr2l	
1445028_at	1.07819	"Protein kinase C, alpha"	Prkca	
1445029_at	1.08506	RIKEN cDNA A330019N05 gene	A330019N05Rik	
1445030_at	1.04276	"PREDICTED: Mus musculus similar to Glyceraldehyde-3-phosphate dehydrogenase (GAPDH) (38 kDa BFA-dependent ADP-ribosylation substrate) (BARS-38) (LOC623178), mRNA"	---	
1445031_at	1.08825	"TBC1 domain family, member 8"	Tbc1d8	
1445032_at	1.05961	Death associated protein kinase 1	Dapk1	
1445033_at	1.02209	"ATP-binding cassette, sub-family A (ABC1), member 16"	Abca16	
1445034_at	0.98503	"Coronin, actin binding protein 2A"	Coro2a	
1445035_at	1.01980	immunoglobulin kappa chain complex variable region	Igk-V	
1445036_at	1.04893	"DNA segment, Chr 11, ERATO Doi 518, expressed"	D11Ertd518e	
1445037_at	1.84121	RIKEN cDNA 6430510B20 gene	6430510B20Rik	
1445038_at	0.86590	---	---	
1445039_at	0.90435	---	---	
1445040_at	0.87814	"DNA segment, Chr 7, ERATO Doi 783, expressed"	D7Ertd783e	
1445041_at	1.00981	"DNA segment, Chr 16, ERATO Doi 6, expressed"	D16Ertd6e	
1445042_at	0.90380	Amino adipate-semialdehyde synthase	Aass	

1445043_at 0.99396 --- ---
1445044_at 1.04626 Prostaglandin I2 (prostacyclin) synthase Ptgis
1445045_at 0.94991 Mastermind like 1 (Drosophila) Maml1
1445046_at 1.06734 "DNA segment, Chr 8, ERATO Doi 317, expressed" D8Ertd317e
1445047_at 0.95052 expressed sequence C79246 C79246
1445048_at 1.10729 "bromodomain adjacent to zinc finger domain, 2B" Baz2b
1445049_at 0.94253 expressed sequence C80893 C80893
1445050_at 0.92661 expressed sequence AU015247 AU015247
1445051_at 1.01271 --- ---
1445052_at 1.07717 Protocadherin 9 Pcdh9
1445053_at 1.00069 expressed sequence AU022537 AU022537
1445054_at 0.95168 Forkhead box N2 Foxn2
1445055_at 0.93763 "Adult male hypothalamus cDNA, RIKEN full-length enriched library, clone:
A230013J07 product:unclassifiable, full insert sequence" ---
1445056_at 1.01728 RIKEN cDNA 9230102K24 gene 9230102K24Rik
1445057_at 1.05887 expressed sequence C81285 C81285
1445058_at 1.09187 Fraser syndrome 1 homolog (human) Fras1
1445059_at 1.01850 expressed sequence C80435 C80435
1445060_at 0.91131 "DNA segment, Chr 5, ERATO Doi 470, expressed" D5Ertd470e
1445061_at 1.02630 Protocadherin 9 Pcdh9
1445062_at 1.79659 "ELOVL family member 6, elongation of long chain fatty acids (yeast)" Elov16
1445063_at 0.95060 "DNA segment, Chr 1, ERATO Doi 228, expressed" D1Ertd228e
1445064_at 0.84096 developmentally regulated repeat element-containing transcript 1 Drr1
1445065_at 0.85379 "Atpase, class VI, type 11C" Atp11c
1445066_at 0.91857 Tetratricopeptide repeat domain 7B Ttc7b
1445067_at 1.07772 ADP-ribosylation factor GTPase activating protein 1 Arfgap1
1445068_at 1.15360 mucosa associated lymphoid tissue lymphoma translocation gene 1 Malt1
1445069_at 1.02996 --- ---
1445070_at 0.86328 Transcribed locus ---
1445071_at 0.93797 --- ---
1445072_at 1.05268 --- ---
1445073_at 0.98137 --- ---
1445074_at 1.09071 "16 days neonate heart cDNA, RIKEN full-length enriched library, clone:
D830036I24 product:weakly similar to CYTOCHROME C OXIDASE POLYPEPTIDE VA,
MITOCHONDRIAL PRECURSOR (EC 1.9.3.1) [Rattus norvegicus], full insert sequence" ---
1445075_at 1.06435 --- ---
1445076_at 0.95493 --- ---
1445077_at 0.85880 --- ---
1445078_at 1.07870 "DNA segment, Chr 7, ERATO Doi 526, expressed" D7Ertd526e
1445079_at 0.99723 "DNA segment, Chr 15, ERATO Doi 320, expressed" D15Ertd320e
1445080_at 1.27876 "WW, C2 and coiled-coil domain containing 1" Wwc1
1445081_at 0.89332 RIKEN cDNA A930041I02 gene A930041I02Rik
1445082_at 1.18397 Kinesin family member 16B Kif16b

1445083_at	1.08770	Transcribed locus	---
1445084_at	0.95874	---	---
1445085_at	0.98639	Calsyntenin 1	Clstn1
1445086_at	1.02768	"Cytochrome P450, family 2, subfamily j, polypeptide 5"	Cyp2j5
1445087_at	1.07539	---	---
1445088_at	1.09272	"Membrane associated guanylate kinase, WW and PDZ domain containing 1"	Magi1
1445089_at	0.95550	"DNA segment, Chr 16, ERATO Doi 778, expressed"	D16Ertd778e
1445090_at	0.98969	"Glutamate receptor, ionotropic, kainate 2 (beta 2)"	Grik2
1445091_at	1.56975	ataxia telangiectasia and Rad3 related	Atr
1445092_at	0.98184	---	---
1445093_at	1.07131	Transcription factor 12	Tcf12
1445094_at	0.99595	---	---
1445095_at	0.97621	Interleukin 31 receptor A	Il31ra
1445096_at	0.94900	Coiled-coil domain containing 51	Ccdc51
1445097_at	1.02334	"Phosphatidylinositol 3 kinase, regulatory subunit, polypeptide 3 (p55)"	Pik3r3
1445098_at	0.89392	expressed sequence AU022706	AU022706
1445099_at	0.93397	Transcribed locus	---
1445100_at	0.98763	---	---
1445101_at	1.13150	---	---
1445102_at	1.05978	RIKEN cDNA B930096M23 gene	B930096M23Rik
1445103_at	0.99641	Transmembrane protein 23	Tmem23
1445104_at	1.04471	RIKEN cDNA E230029C05 gene	E230029C05Rik
1445105_at	1.20438	P300/CBP-associated factor	Pcaf
1445106_at	1.25279	Muscleblind-like 1 (Drosophila)	Mbnl1
1445107_at	1.03546	RIKEN cDNA A530064N14 gene	A530064N14Rik
1445108_at	0.97165	Latrophilin 3	Lphn3
1445109_at	1.01827	WD repeat domain 4	Wdr4
1445110_at	0.96150	"KH domain containing, RNA binding, signal transduction associated 2"	Khdrbs2
1445111_at	1.76603	Exocyst complex component 4	Exoc4
1445112_at	1.01216	Adenylate cyclase 1	Adcy1
1445113_at	1.13287	---	---
1445114_at	0.97501	"Catenin (cadherin associated protein), delta 2"	Ctnnd2
1445115_at	0.97226	RIKEN cDNA D130067I03 gene	D130067I03Rik
1445116_at	0.94540	Ubiquitin specific peptidase 25	Usp25
1445117_at	0.96528	Glycosyltransferase 28 domain containing 1	Glt28d1
1445118_at	1.04096	---	---
1445119_at	1.04670	"DNA segment, Chr 7, ERATO Doi 661, expressed"	D7Ertd661e
1445120_at	0.99050	expressed sequence AU021977	AU021977
1445121_at	1.02608	RIKEN cDNA 1110007J02 gene	1110007J02Rik
1445122_at	1.12469	RIKEN cDNA A830082N09 gene	A830082N09Rik

1445123_at 0.93083 "PREDICTED: Mus musculus similar to ubiquitin A-52 residue ribosomal protein fusion product 1 (LOC629750), mRNA" ---
1445124_at 1.04751 RIKEN cDNA A930009L07 gene A930009L07Rik
1445125_at 0.95269 "DNA segment, Chr 8, ERATO Doi 107, expressed" D8Ertd107e
1445126_at 1.03986 LIM domain binding 2 Ldb2
1445127_at 0.90275 collagen and calcium binding EGF domains 1 Ccbe1
1445128_at 0.98355 CDNA clone IMAGE:30463708 ---
1445129_at 1.08371 --- ---
1445130_at 1.06825 expressed sequence AA407111 AA407111
1445131_at 0.97305 F-box and leucine-rich repeat protein 13 Fbxl13
1445132_at 1.06104 Arginyl aminopeptidase (aminopeptidase B) Rnpep
1445133_at 1.04170 Transcription factor 12 Tcf12
1445134_at 0.97221 MKL/myocardin-like 2 Mkl2
1445135_at 1.06893 --- ---
1445136_at 1.03241 --- ---
1445137_at 2.26036 PTK2 protein tyrosine kinase 2 Ptk2
1445138_at 0.93604 RIKEN cDNA A530083I20 gene A530083I20Rik
1445139_at 0.96522 Expressed sequence AV312086 AV312086
1445140_at 1.05987 expressed sequence AU022804 AU022804
1445141_at 1.15824 inhibitor of kappaB kinase beta Ikbkb
1445142_at 0.96665 "Adaptor-related protein complex 2, beta 1 subunit" Ap2b1
1445143_at 1.05821 vasohibin 1 Vash1
1445144_at 1.00804 "Syntrophin, basic 1" Sntb1
1445145_at 0.94890 Stromal antigen 1 Stag1
1445146_at 1.07713 RIKEN cDNA B230219N05 gene B230219N05Rik
1445147_at 0.92948 "proteasome (prosome, macropain) activator subunit 4" Psme4
1445148_at 0.98322 Phospholipase C-like 1 Plcl1
1445149_at 1.03696 Poliovirus receptor Pvr
1445150_at 1.00120 cDNA sequence BC023744 BC023744
1445151_at 1.16063 --- ---
1445152_at 1.03562 Calcium/calmodulin-dependent serine protein kinase (MAGUK family) Cask
1445153_at 1.13397 V-abl Abelson murine leukemia oncogene 1 Abl1
1445154_at 0.84573 RIKEN cDNA 3300002I08 gene 3300002I08Rik
1445155_at 0.99213 Transcribed locus ---
1445156_at 1.08046 expressed sequence C81001 C81001
1445157_at 1.08589 --- ---
1445158_at 0.94093 synaptophysin-like protein Sypl
1445159_at 1.03984 Transcribed locus ---
1445160_at 1.06238 RIKEN cDNA 9630020C08 gene /// region containing RIKEN cDNA 9630020C08 gene; neuron navigator 3 9630020C08Rik /// LOC676640
1445161_at 1.24313 --- ---
1445162_at 1.02363 Transcribed locus ---
1445163_at 1.02532 --- ---

1445164_at 1.06775 RIKEN cDNA 2210418O10 gene 2210418O10Rik
1445165_at 0.95259 RIKEN cDNA 1200003I07 gene 1200003I07Rik
1445166_at 0.98414 Polycomb group ring finger 3 Pcgf3
1445167_at 0.99403 RIKEN cDNA 9630001P10 gene 9630001P10Rik
1445168_at 1.05665 --- ---
1445169_at 1.27662 --- ---
1445170_at 0.97925 RIKEN cDNA 2700012I20 gene 2700012I20Rik
1445171_at 1.01280 RIKEN cDNA 1700067C01 gene /// RIKEN cDNA A230107C01 gene
1700067C01Rik /// A230107C01Rik
1445172_at 1.02978 ligatin Lgtn
1445173_at 0.90817 cDNA sequence AK129341 AK129341
1445174_at 1.06227 --- ---
1445175_at 0.84969 "Adult male aorta and vein cDNA, RIKEN full-length enriched library, clone:
A530052F04 product:unclassifiable, full insert sequence" ---
1445176_at 0.97014 --- ---
1445177_at 1.02026 "4 days neonate male adipose cDNA, RIKEN full-length enriched library, clone:
B430206K05 product:unclassifiable, full insert sequence" ---
1445178_at 2.22360 SH3 domain containing ring finger 1 Sh3rf1
1445179_at 0.96126 "DNA segment, Chr 10, ERATO Doi 494, expressed" D10Ertd494e
1445180_at 1.29916 steroid 5 alpha-reductase 2-like Srd5a2l
1445181_at 0.96783 echinoderm microtubule associated protein like 5 Eml5
1445182_at 0.98483 Zinc finger protein 672 Zfp672
1445183_s_at 0.98847 "DNA segment, Chr 7, ERATO Doi 523, expressed" D7Ertd523e
1445184_at 0.92295 RIKEN cDNA 9130019O22 gene 9130019O22Rik
1445185_at 0.99881 "tubulin tyrosine ligase-like family, member 10" Ttl10
1445186_at 0.98709 stanniocalcin 2 Stc2
1445187_at 1.00073 RIKEN cDNA 9430070O13 gene 9430070O13Rik
1445188_at 0.95037 Gephyrin Gphn
1445189_at 0.88000 G patch domain containing 2 Gpatc2
1445190_at 0.95460 golgi phosphoprotein 4 Golph4
1445191_at 1.02199 Vexonuclease 3'-5' domain-like 1 Exdl1
1445192_at 0.98601 "DNA segment, Chr 8, ERATO Doi 575, expressed" D8Ertd575e
1445193_x_at 1.00852 "DNA segment, Chr 8, ERATO Doi 575, expressed" D8Ertd575e
1445194_at 0.97548 connector enhancer of kinase suppressor of Ras 2 Cnksr2
1445195_at 1.17536 expressed sequence C77631 C77631
1445196_at 1.09570 expressed sequence C79595 C79595
1445197_x_at 1.03359 expressed sequence C79595 C79595
1445198_at 2.44614 "ubiquitously transcribed tetratricopeptide repeat gene, X chromosome" Utx
1445199_at 0.93410 Transcribed locus ---
1445200_at 1.09330 "FERM, RhoGEF (Arhgef) and pleckstrin domain protein 1 (chondrocyte-
derived)" Farp1
1445201_at 1.26154 zinc finger protein 53 Zfp53
1445202_at 1.01791 "0 day neonate cerebellum cDNA, RIKEN full-length enriched library, clone:

C230022P04 product:unclassifiable, full insert sequence" ---

1445203_at 0.99682 PDZ domain containing 2 Pdzd2

1445204_at 0.99821 Coiled-coil domain containing 85A Ccdc85a

1445205_at 1.04930 GTP binding protein 3 Gtpbp3

1445206_at 0.94299 "ZW10 homolog (Drosophila), centromere/kinetochore protein" Zw10

1445207_at 1.04015 Hypothetical protein LOC208426 D230050A05

1445208_at 0.95791 exonuclease 3"-5" domain-like 2 Exdl2

1445209_at 0.93610 expressed sequence AU021745 AU021745

1445210_at 2.19703 Transcribed locus ---

1445211_at 1.20603 "PREDICTED: Mus musculus hypothetical protein LOC626411 (LOC626411), mRNA" ---

1445212_at 1.35391 "3 days neonate thymus cDNA, RIKEN full-length enriched library, clone: A630021H04 product:unclassifiable, full insert sequence" ---

1445213_at 0.99901 cylindromatosis (turban tumor syndrome) Cyld

1445214_at 1.04315 Similar to reduced expression 2 LOC195531

1445215_at 1.09626 RIKEN cDNA 2600003E23 gene 2600003E23Rik

1445216_at 0.98842 --- ---

1445217_at 1.06028 RIKEN cDNA A730073F16 gene A730073F16Rik

1445218_at 1.12782 Ventral anterior homeobox containing gene 2 Vax2

1445219_at 0.94179 olfactory receptor 315 Olfr315

1445220_at 1.04873 BTB (POZ) domain containing 9 Btbd9

1445221_at 0.98214 Transcribed locus ---

1445222_at 1.01405 CD22 antigen Cd22

1445223_at 1.10958 --- ---

1445224_at 0.93648 "Adaptor-related protein complex 3, beta 2 subunit" Ap3b2

1445225_at 1.34210 RIKEN cDNA 2410001C21 gene 2410001C21Rik

1445226_at 1.92883 cDNA sequence BC023969 BC023969

1445227_at 1.09376 "Tumor necrosis factor (ligand) superfamily, member 13" Tnfsf13

1445228_at 0.92737 "Sprouty protein with EVH-1 domain 1, related sequence" Spred1

1445229_at 1.18284 diacylglycerol O-acyltransferase 1 Dgat1

1445230_at 1.04725 Nuclear receptor co-repressor 1 Ncor1

1445231_at 0.90116 RIKEN cDNA B230112C05 gene B230112C05Rik

1445232_at 1.07461 Deducator of cytokinesis 9 Dock9

1445233_at 1.07893 hypothetical protein 9130416B15 9130416B15

1445234_at 1.00588 RIKEN cDNA C130030K03 gene C130030K03Rik

1445235_at 1.48658 YTH domain family 3 Ythdf3

1445236_at 1.00307 Hypothetical protein LOC626410 LOC626410

1445237_at 1.07552 --- ---

1445238_at 1.02801 chemokine (C-C motif) ligand 21b Ccl21b

1445239_at 1.13817 GATA zinc finger domain containing 2A Gatad2a

1445240_at 1.05446 expressed sequence AU019796 AU019796

1445241_at 0.94591 RAB11 family interacting protein 4 (class II) Rab11fip4

1445242_at 0.70381 glycerol kinase Gyk

1445243_at 1.07013 "DNA segment, Chr 10, ERATO Doi 533, expressed" D10ErtD533e
1445244_at 1.15552 TM2 domain containing 1 Tm2d1
1445245_at 1.33805 RIKEN cDNA B130065D12 gene B130065D12Rik
1445246_at 0.84768 "G elongation factor, mitochondrial 2" Gfm2
1445247_at 0.96183 RIKEN cDNA C530044C16 gene C530044C16Rik
1445248_at 0.88765 Tetratricopeptide repeat domain 18 Ttc18
1445249_at 1.02687 --- ---
1445250_at 0.94073 Hephaestin-like 1 Heph11
1445251_at 1.15065 "tumor necrosis factor (ligand) superfamily, member 13b" Tnfsf13b
1445252_at 0.97073 kringle containing transmembrane protein 1 Kremen1
1445253_at 1.01919 Olfactory receptor 1417 Olfr1417
1445254_at 1.03933 --- ---
1445255_at 0.94322 Rap guanine nucleotide exchange factor (GEF) 5 Rapgef5
1445256_at 1.09885 Vinculin Vcl
1445257_at 1.03666 expressed sequence AU022531 AU022531
1445258_at 0.88111 Fibroblast growth factor 9 Fgf9
1445259_at 0.94731 "sodium channel, voltage-gated, type II, alpha 1" Scn2a1
1445260_at 1.40226 "DCUN1D1 DCN1, defective in cullin neddylation 1, domain containing 1 (S. cerevisiae)" Dcun1d1
1445261_at 0.97421 "DNA segment, Chr 17, ERATO Doi 763, expressed" D17ErtD763e
1445262_at 0.97340 --- ---
1445263_at 1.49653 Cystatin F (leukocystatin) Cst7
1445264_at 0.93047 "cation channel, sperm associated 2" Catsper2
1445265_at 1.03277 "Ubiquitously transcribed tetratricopeptide repeat gene, Y chromosome" Uty
1445266_at 0.92516 "excision repair cross-complementing rodent repair deficiency, complementation group 1" Ercc1
1445267_at 1.22833 Similar to zinc finger protein 709 LOC626854
1445268_at 0.96242 "Transcribed locus, weakly similar to NP_059506.1 coatomer protein complex, subunit gamma 2 [Mus musculus]" ---
1445269_at 1.11173 Coiled-coil domain containing 98 Ccdc98
1445270_at 1.04366 Transcribed locus ---
1445271_at 1.07995 RIKEN cDNA 9230105E10 gene 9230105E10Rik
1445272_at 1.07206 Ring finger protein 38 Rnf38
1445273_at 1.52967 "DNA segment, Chr 10, ERATO Doi 638, expressed" D10ErtD638e
1445274_at 0.79979 cDNA sequence BC024063 BC024063
1445275_at 0.92743 RIKEN cDNA C130092O11 gene C130092O11Rik
1445276_at 0.96575 expressed sequence C80154 C80154
1445277_at 1.78639 "16 days neonate thymus cDNA, RIKEN full-length enriched library, clone: A130006J10 product:unclassifiable, full insert sequence" ---
1445278_at 0.91670 gametogenetin Ggn
1445279_at 0.95791 Transcribed locus ---
1445280_at 0.95798 "adaptor-related protein complex 3, delta 1 subunit" Ap3d1
1445281_a_at 1.11658 RIKEN cDNA B230311B06 gene B230311B06Rik

1445282_at 1.08017 "Muskelin 1, intracellular mediator containing kelch motifs" Mkl1
1445283_at 1.09605 --- ---
1445284_at 0.90432 --- ---
1445285_at 0.90398 Histamine receptor H 3 Hrh3
1445286_at 1.06437 RIKEN cDNA 6430701C03 gene 6430701C03Rik
1445287_at 0.92288 "Adult male spinal cord cDNA, RIKEN full-length enriched library, clone:
A330010P20 product:unclassifiable, full insert sequence" ---
1445288_at 0.98192 --- ---
1445289_at 1.03358 --- ---
1445290_at 1.03792 Hormonally upregulated Neu-associated kinase Hunk
1445291_at 1.00895 arylsulfatase A Arsa
1445292_at 1.05233 CD300A antigen Cd300a
1445293_at 1.02447 RIKEN cDNA 2610002D18 gene 2610002D18Rik
1445294_at 1.03360 "Adult male corpora quadrigemina cDNA, RIKEN full-length enriched library,
clone:B230354C23 product:unclassifiable, full insert sequence" ---
1445295_at 1.09453 Erythrocyte protein band 4.1 Epb4.1
1445296_at 0.89854 "DNA segment, Chr 7, ERATO Doi 715, expressed" D7Ertd715e
1445297_at 0.95622 Transcribed locus ---
1445298_at 0.92445 --- ---
1445299_at 1.07741 Protein phosphatase 1H (PP2C domain containing) Ppm1h
1445300_at 1.13069 --- ---
1445301_at 0.95553 CDNA sequence BC028528 BC028528
1445302_at 1.38836 "immunoglobulin superfamily, member 6" Igsf6
1445303_at 0.89401 --- ---
1445304_at 1.38366 "mahogunin, ring finger 1" Mgrn1
1445305_at 1.07181 --- ---
1445306_at 0.96875 "CUG triplet repeat, RNA binding protein 2" Cugbp2
1445307_at 1.00984 Autism susceptibility candidate 2 Aut2
1445308_at 1.03124 tumor suppressing subtransferable candidate 1 Tssc1
1445309_at 0.83237 --- ---
1445310_at 0.88080 synaptotagmin-like 3 Sytl3
1445311_at 0.98002 A kinase (PRKA) anchor protein 13 Akap13
1445312_at 1.19371 RIKEN cDNA C530043G21 gene C530043G21Rik
1445313_at 1.10796 coenzyme Q9 homolog (yeast) Coq9
1445314_at 1.04477 ets variant gene 1 Etv1
1445315_at 0.98928 --- ---
1445316_at 0.96826 Armadillo repeat containing 9 Armc9
1445317_at 1.10742 "Succinate dehydrogenase complex, subunit A, flavoprotein (Fp)" Sdha
1445318_at 0.91476 Kelch-like 4 (Drosophila) Klhl4
1445319_at 0.87698 "Adult male testis cDNA, RIKEN full-length enriched library,
clone:4930509G02 product:unclassifiable, full insert sequence" ---
1445320_at 0.83441 "12 days embryo spinal ganglion cDNA, RIKEN full-length enriched library,
clone:D130033J21 product:unclassifiable, full insert sequence" ---

1445321_at	1.38641	RIKEN cDNA E430025E21 gene	E430025E21Rik	
1445322_x_at	1.54534	RIKEN cDNA E430025E21 gene	E430025E21Rik	
1445323_at	1.07471	gene regulated by estrogen in breast cancer protein	Greb1	
1445324_s_at	0.86801	gene regulated by estrogen in breast cancer protein	Greb1	
1445325_at	1.14640	"Protein kinase, cAMP dependent regulatory, type II alpha"	Prkar2a	
1445326_at	0.94391	sidekick homolog 1 (chicken)	Sdk1	
1445327_at	1.10520	Cytoplasmic FMR1 interacting protein 1	Cyfip1	
1445328_at	0.48694	"procollagen, type IV, alpha 4"	Col4a4	
1445329_at	1.16117	"dystrobrevin, beta"	Dtnb	
1445330_at	0.94215	protein disulfide isomerase associated 3	Pdia3	
1445331_at	1.00919	Transcribed locus	---	
1445332_at	1.04783	Transcribed locus	---	
1445333_at	0.96251	---	---	
1445334_at	1.12185	similar to signal recognition particle 54	LOC665155	
1445335_at	0.97358	expressed sequence C78888	C78888	
1445336_at	0.94637	RIKEN cDNA 1110032A03 gene	1110032A03Rik	
1445337_at	1.12130	"DnaJ (Hsp40) homolog, subfamily C, member 13"	Dnajc13	
1445338_at	1.11296	Ring finger protein 144	Rnf144	
1445339_at	1.08626	cyclic nucleotide gated channel alpha 4	Cnga4	
1445340_at	1.48557	"pam, highwire, rpm 1"	Phr1	
1445341_at	1.66719	---	---	
1445342_at	1.03967	---	---	
1445343_at	0.91510	transmembrane emp24 protein transport domain containing 5	Tmed5	
1445344_at	0.99008	Src homology three (SH3) and cysteine rich domain	Stac	
1445345_at	1.20068	"Pleckstrin homology domain containing, family A member 5"	Plekha5	
1445346_at	1.07592	"Ankyrin 2, brain"	Ank2	
1445347_at	1.09562	---	---	
1445348_at	0.98949	cell division cycle 40 homolog (yeast)	Cdc40	
1445349_at	0.93156	"0 day neonate lung cDNA, RIKEN full-length enriched library, clone: E030049A20 product:unclassifiable, full insert sequence"	---	
1445350_at	0.96289	"protein phosphatase 4, catalytic subunit"	Ppp4c	
1445351_at	1.04990	---	---	
1445352_at	1.14310	Transcribed locus	---	
1445353_at	0.85489	---	---	
1445354_at	1.21452	Transcribed locus	---	
1445355_at	0.99218	spermatogenesis associated 6	Spata6	
1445356_at	0.94497	---	---	
1445357_at	1.16018	expressed sequence AI851613	AI851613	
1445358_at	0.90240	"tankyrase, TRF1-interacting ankyrin-related ADP-ribose polymerase"	Tnks	
1445359_at	1.07963	adenylate cyclase 1	Adcy1	
1445360_at	0.99032	"Transforming growth factor, beta 1"	Tgfb1	
1445361_at	1.30828	---	---	
1445362_at	0.95878	expressed sequence AW456874	AW456874	

1445363_at	1.12032	RIKEN cDNA 2810055G20 gene	2810055G20Rik
1445364_at	0.84264	---	---
1445365_at	0.97128	"membrane-spanning 4-domains, subfamily A, member 7"	Ms4a7
1445366_at	1.00584	---	---
1445367_at	0.81880	Transcribed locus	---
1445368_at	0.85309	---	---
1445369_at	0.94827	RIKEN cDNA B230359F08 gene	B230359F08Rik
1445370_at	1.05605	"Vesicle-associated membrane protein, associated protein B and C"	Vapb
1445371_at	1.05333	"gene model 540, (NCBI)"	Gm540
1445372_at	0.94927	MAM domain containing 1	Mamdc1
1445373_at	1.06975	"Phosphodiesterase 3A, cGMP inhibited"	Pde3a
1445374_at	1.56355	Vesicle transport through interaction with t-SNAREs homolog 1A (yeast)	Vti1a
1445375_at	1.13516	intersectin 2	Itsn2
1445376_at	1.05861	Spastin	Spast
1445377_at	1.66869	"Amyotrophic lateral sclerosis 2 (juvenile) chromosome region, candidate 19 (human)"	Als2cr19
1445378_at	1.01472	---	---
1445379_at	1.01037	"Atpase, class VI, type 11C"	Atp11c
1445380_at	0.90055	RIKEN cDNA A230072C01 gene	A230072C01Rik
1445381_at	1.05653	---	---
1445382_at	1.02578	Transcribed locus	---
1445383_at	0.98676	transmembrane protein 44	Tmem44
1445384_at	1.03530	RIO kinase 2 (yeast)	Riok2
1445385_at	0.93235	"Adult inner ear cDNA, RIKEN full-length enriched library, clone: F930025M14 product:unclassifiable, full insert sequence"	---
1445386_at	0.95972	Transcribed locus	---
1445387_at	1.05190	SUMO/sentrin specific peptidase 6	Senp6
1445388_at	0.89773	CD226 antigen	Cd226
1445389_at	0.95045	Zona pellucida 3 receptor	Zp3r
1445390_at	1.14664	"PRKC, apoptosis, WT1, regulator"	Pawr
1445391_at	1.24644	diaphanous homolog 1 (Drosophila) /// similar to Protein diaphanous homolog 1 (Diaphanous-related formin-1) (DRF1) (mDIA1) (p140mDIA) Diap1 /// LOC671545	
1445392_at	0.86556	Transmembrane 7 superfamily member 3	Tm7sf3
1445393_at	0.96888	"Solute carrier family 24 (sodium/potassium/calcium exchanger), member 2"	Slc24a2
1445394_at	0.90008	RIKEN cDNA 4933409K07 gene	4933409K07Rik
1445395_at	1.28650	"Protein kinase C, alpha"	Prkca
1445396_at	0.99295	leucine-rich repeats and calponin homology (CH) domain containing 1	Lrch1
1445397_at	1.22815	---	---
1445398_at	0.99730	RIKEN cDNA 3732412D22 gene	3732412D22Rik
1445399_at	0.99814	killer cell lectin-like receptor subfamily B member 1D	Klrk1d
1445400_at	1.50082	Hepatitis B virus x interacting protein	Hbxip
1445401_at	0.97518	Shroom	Shrm

1445402_at	1.67760	Adenosine kinase	Adk
1445403_at	0.96031	Transcribed locus	---
1445404_at	0.84082	kinesin family member 27	Kif27
1445405_at	1.04417	---	---
1445406_at	0.84294	expressed sequence C77494	C77494
1445407_at	0.94691	---	---
1445408_at	1.06488	"similar to dynein, axonemal, intermediate polypeptide 2"	LOC432611
1445409_at	0.99260	LPS-responsive beige-like anchor	Lrba
1445410_at	1.24667	"MAK10 homolog, amino-acid N-acetyltransferase subunit, (<i>S. cerevisiae</i>)"	
Mak10			
1445411_at	1.16038	---	---
1445412_at	1.07523	coiled-coil domain containing 73	Ccdc73
1445413_at	0.88308	"Bone morphogenetic protein receptor, type 1A"	Bmpr1a
1445414_at	1.08769	Transcribed locus	---
1445415_at	0.92801	Hypothetical protein B230303A05	B230303A05
1445416_at	1.00575	Testis specific gene A14	Tsga14
1445417_at	1.16588	Eph receptor B2	Ephb2
1445418_at	1.13342	"Smg-6 homolog, nonsense mediated mRNA decay factor (<i>C. elegans</i>)"	
Smg6			
1445419_at	1.00261	GTP-binding protein 8 (putative)	Gtpbp8
1445420_at	1.14596	Myocyte enhancer factor 2C	Mef2c
1445421_at	0.93444	Transcribed locus	---
1445422_at	1.03015	hypothetical protein LOC621549	LOC621549
1445423_at	1.06266	AT rich interactive domain 5B (<i>Mrf1</i> like)	Arid5b
1445424_at	1.05252	---	---
1445425_at	0.94017	hypothetical gene supported by AK040104	LOC432958
1445426_at	1.38544	Mitogen-activated protein kinase kinase kinase kinase 5	Map4k5
1445427_at	1.27641	Nemo like kinase	Nlk
1445428_at	0.95459	---	---
1445429_at	0.88832	Prune homolog (<i>Drosophila</i>)	Prune
1445430_at	0.98750	Rho GTPase activating protein 28	Arhgap28
1445431_at	0.96708	"Serine/threonine kinase 39, STE20/SPS1 homolog (yeast)"	Stk39
1445432_at	1.03825	---	---
1445433_at	0.92323	glycosyltransferase 8 domain containing 2	Glt8d2
1445434_at	1.05347	snurportin 1	Snupn
1445435_at	1.04635	"gene model 732, (NCBI)"	Gm732
1445436_at	1.02741	Transcribed locus	---
1445437_at	0.99295	RIKEN cDNA 2310015A05 gene	2310015A05Rik
1445438_at	1.09380	DDHD domain containing 1	Ddhd1
1445439_at	1.05445	erythrocyte protein band 4.9	Epb4.9
1445440_at	1.77944	RIKEN cDNA A430106J12 gene	A430106J12Rik
1445441_at	0.93198	similar to RIKEN cDNA 1700129C05	LOC380907
1445442_at	1.04242	microtubule-associated protein 9	Mtap9

1445443_at 1.49647 --- ---
1445444_at 1.08406 "0 day neonate cerebellum cDNA, RIKEN full-length enriched library, clone: C230092G11 product:unclassifiable, full insert sequence" ---
1445445_s_at 1.05769 prostaglandin E receptor 1 (subtype EP1) Ptger1
1445446_at 0.86804 Transcribed locus ---
1445447_at 0.93410 --- ---
1445448_at 1.10297 --- ---
1445449_at 0.86557 "Phosphodiesterase 4B, cAMP specific" Pde4b
1445450_x_at 1.01278 hypothetical protein A530021J07 A530021J07
1445451_at 0.96194 --- ---
1445452_at 1.09015 Tnf receptor-associated factor 1 Traf1
1445453_at 1.06401 --- ---
1445454_at 1.03758 "DNA segment, Chr 2, ERATO Doi 282, expressed" D2Ertd282e
1445455_at 1.02962 "DNA segment, Chr 3, ERATO Doi 270, expressed" D3Ertd270e
1445456_at 1.17373 --- ---
1445457_at 1.03938 "Adult male cecum cDNA, RIKEN full-length enriched library, clone:9130219A07 product:hypothetical protein, full insert sequence" ---
1445458_at 1.03024 "MAS-related GPR, member G" Mrgprg
1445459_at 1.00203 somatostatin receptor 5 Sstr5
1445460_at 1.02884 Transcribed locus ---
1445461_at 0.91222 Transcribed locus ---
1445462_at 0.91399 --- ---
1445463_at 1.14287 aminopeptidase puromycin sensitive Npepps
1445464_at 0.93857 --- ---
1445465_at 0.84020 expressed sequence N28178 N28178
1445466_at 1.19643 MAP6 domain containing 1 Map6d1
1445467_at 0.87701 Transcribed locus ---
1445468_at 1.14806 "PREDICTED: Mus musculus hypothetical protein LOC625847 (LOC625847), mRNA" ---
1445469_at 1.00777 Autism susceptibility candidate 2 Aut2
1445470_at 0.87928 similar to alpha 3 type VI collagen isoform 4 precursor /// similar to alpha 3 type VI collagen isoform 4 precursor LOC235580 /// LOC669304
1445471_at 0.92030 "Transcriptional adaptor 2 (ADA2 homolog, yeast)-like" Tada2l
1445472_at 0.87274 RIKEN cDNA 2210412E05 gene /// similar to Carcinoembryonic antigen-related cell adhesion molecule 1 precursor (Biliary glycoprotein 1) (BGP-1) (Murine hepatitis virus receptor) (MHV-R) (Biliary glycoprotein D) 2210412E05Rik /// LOC672817
1445473_at 1.01413 "5-azacytidine induced gene 2 /// DIRAS family, GTP-binding RAS-like 2" Azi2 /// Diras2
1445474_at 0.86267 "DNA primase, p58 subunit" Prim2
1445475_at 0.90536 p21 (CDKN1A)-activated kinase 6 Pak6
1445476_at 1.04421 "Mus musculus, clone IMAGE:1313576, mRNA" ---
1445477_at 1.04976 Transcribed locus ---
1445478_at 0.86416 --- ---

1445479_at	1.11442 --- ---	
1445480_at	0.96859 G protein-coupled receptor 19	Gpr19
1445481_at	0.82265 expressed sequence AI317158	AI317158
1445482_at	1.00014 Transcribed locus	---
1445483_at	0.96363 "v-abl Abelson murine leukemia viral oncogene 2 (arg, Abelson-related gene)"	Abl2
1445484_at	0.84611 "cytochrome c, somatic"	Cycc
1445485_at	0.96633 "DNA segment, Chr 7, ERATO Doi 187, expressed"	D7Erttd187e
1445486_at	1.02400 Leucine rich repeat containing 49	Lrrc49
1445487_at	0.96482 Rap1 interacting factor 1 homolog (yeast)	Rif1
1445488_at	1.10368 Transcribed locus	---
1445489_at	1.13518 RIKEN cDNA 5830472M02 gene	5830472M02Rik
1445490_at	1.20125 expressed sequence C77805	C77805
1445491_at	1.02846 expressed sequence C79685	C79685
1445492_at	1.20488 "Microtubule-associated protein, RP/EB family, member 2"	Mapre2
1445493_at	0.98671 expressed sequence AU022255	AU022255
1445494_at	1.08669 "Achalasia, adrenocortical insufficiency, alacrimia"	Aaas
1445495_at	0.98214 --- ---	
1445496_at	0.90550 AU RNA binding protein/enoyl-coenzyme A hydratase	Auh
1445497_at	0.95608 Carbonic anhydrase 10	Car10
1445498_at	0.95465 "Ryanodine receptor 2, cardiac"	Ryr2
1445499_at	1.03869 zinc finger CCCH type containing 13	Zc3h13
1445500_at	1.07470 A disintegrin and metallopeptidase domain 17	Adam17
1445501_at	1.04043 "12 days embryo spinal cord cDNA, RIKEN full-length enriched library, clone: C530045D16 product:unclassifiable, full insert sequence"	---
1445502_at	1.00058 "15 days embryo head cDNA, RIKEN full-length enriched library, clone: D930047H10 product:unclassifiable, full insert sequence"	---
1445503_at	1.01168 "gene model 715, (NCBI) /// similar to CG4768-PA"	Gm715 /// LOC671594
1445504_at	0.85489 similar to Nucleolar phosphoprotein p130 (Nucleolar 130 kDa protein) (140 kDa nucleolar phosphoprotein) (Nopp140) (Nucleolar and coiled-body phosphoprotein 1)	LOC331392
1445505_at	1.01303 N-deacetylase/N-sulfotransferase (heparan glucosaminyl) 1	Ndst1
1445506_at	1.11188 RIKEN cDNA 1300007F04 gene	1300007F04Rik
1445507_at	1.15655 RIKEN cDNA 1110054O05 gene	1110054O05Rik
1445508_at	0.95396 --- ---	
1445509_at	0.95332 Activating transcription factor 7	Atf7
1445510_at	0.92935 expressed sequence C79601	C79601
1445511_at	1.04833 centrin 4 /// similar to centrin 4	Cetn4 /// LOC632802
1445512_at	1.39189 --- ---	
1445513_at	0.94358 "Vezatin, adherens junctions transmembrane protein"	Vezt
1445514_at	1.20358 von Hippel-Lindau syndrome homolog	Vhlh
1445515_at	1.00306 --- ---	
1445516_at	1.04342 --- ---	
1445517_at	1.24755 --- ---	

1445518_at 1.12253 Zinc finger homeobox 1b Zfhx1b
1445519_at 0.90633 "potassium voltage-gated channel, delayed-rectifier, subfamily S, member 3"
Kcns3
1445520_at 0.94478 "RAB4A, member RAS oncogene family" Rab4a
1445521_at 1.05033 "ELAV (embryonic lethal, abnormal vision, Drosophila)-like 1 (Hu antigen
R)" Elavl1
1445522_at 1.02381 "ATPase, class I, type 8B, member 4" Atp8b4
1445523_at 1.05024 Gap junction membrane channel protein epsilon 1 Gje1
1445524_at 1.03914 Dopa decarboxylase Ddc
1445525_at 0.95359 kinesin family member 14 Kif14
1445526_at 0.97466 cDNA sequence AY512915 AY512915
1445527_at 1.12307 Karyopherin (importin) alpha 3 Kpna3
1445528_at 0.97212 "Solute carrier family 28 (sodium-coupled nucleoside transporter), member 3"
Slc28a3
1445529_at 1.04092 --- ---
1445530_at 1.06337 RIKEN cDNA 5330401P04 gene 5330401P04Rik
1445531_at 0.84502 CUB and Sushi multiple domains 1 Csm1
1445532_at 0.82844 RIKEN cDNA C230040D10 gene C230040D10Rik
1445533_at 1.18324 RIKEN cDNA A630047E20 gene A630047E20Rik
1445534_at 3.60083 "Filamin, beta" Flnb
1445535_at 2.41781 expressed sequence AA407107 AA407107
1445536_at 1.08948 expressed sequence AW061096 AW061096
1445537_at 0.94648 SLIT-ROBO Rho GTPase activating protein 3 Srgap3
1445538_at 1.04329 Autism susceptibility candidate 2 Auts2
1445539_at 0.93872 "PREDICTED: Mus musculus similar to 60S ribosomal protein L12
(LOC382344), mRNA" ---
1445540_at 0.85467 Dynamin 3 Dnm3
1445541_at 0.91426 "jumonji, AT rich interactive domain 1A (Rbp2 like)" Jarid1a
1445542_at 0.99984 RIKEN cDNA 4930579C12 gene 4930579C12Rik
1445543_at 1.18564 RIKEN cDNA 9830124H08 gene 9830124H08Rik
1445544_x_at 1.01446 oligonucleotide/oligosaccharide-binding fold containing 1 Obfc1
1445545_at 1.10151 Transmembrane protein 77 Tmem77
1445546_at 1.03688 expressed sequence AI844685 AI844685
1445547_at 1.44347 Transcribed locus ---
1445548_at 0.93215 Radixin Rdx
1445549_at 0.84346 --- ---
1445550_at 0.93687 --- ---
1445551_at 0.90558 nuclear factor related to kappa B binding protein Nfrkb
1445552_at 1.02344 RIKEN cDNA A830039H10 gene A830039H10Rik
1445553_at 1.11101 resistance to inhibitors of cholinesterase 3 homolog (C. elegans) Ric3
1445554_at 0.98099 --- ---
1445555_at 1.02798 "Transient receptor potential cation channel, subfamily M, member 3" Trpm3
1445556_at 0.85630 Transcribed locus ---

1445557_at 0.98094 RIKEN cDNA 0610040B10 gene 0610040B10Rik
1445558_at 0.99046 RIKEN cDNA 5930430L01 gene 5930430L01Rik
1445559_at 1.13398 Retinitis pigmentosa 9 homolog (human) Rp9h
1445560_at 0.96561 Transcribed locus ---
1445561_at 0.83194 Glutamate receptor interacting protein 1 Grip1
1445562_at 0.97283 Zinc finger and BTB domain containing 16 Zbtb16
1445563_at 1.03181 --- ---
1445564_at 0.95115 Transcribed locus ---
1445565_at 1.01883 "Histone 1, H1e" Hist1h1e
1445566_at 1.01622 "Centaurin, beta 2" Centb2
1445567_at 1.07857 Growth associated protein 43 Gap43
1445568_at 1.05678 Expressed sequence AI842396 AI842396
1445569_at 1.09307 similar to kelch-like 9 LOC546611
1445570_at 1.25261 "NOL1/NOP2/Sun domain family, member 4" Nsun4
1445571_at 1.02541 TANK-binding kinase 1 Tbk1
1445572_at 0.83235 forkhead box Q1 Foxq1
1445573_at 1.06143 "Protein phosphatase 1, regulatory (inhibitor) subunit 14B" Ppp1r14b
1445574_at 1.00782 Transcribed locus ---
1445575_at 0.98086 F-box and leucine-rich repeat protein 20 Fbxl20
1445576_at 0.90624 RIKEN cDNA 4930526H21 gene 4930526H21Rik
1445577_at 0.86442 "PTPRF interacting protein, binding protein 1 (liprin beta 1)" Ppfibp1
1445578_at 0.99137 "ELOVL family member 6, elongation of long chain fatty acids (yeast)" Elov16
1445579_at 0.97732 "arachidonate lipoxygenase, epidermal" Alox12e
1445580_at 0.91450 Matrilin 2 Matn2
1445581_at 1.02513 --- ---
1445582_at 0.92488 "7 days neonate cerebellum cDNA, RIKEN full-length enriched library, clone:
A730006J15 product:hypothetical protein, full insert sequence" ---
1445583_x_at 0.98998 "PREDICTED: Mus musculus hypothetical protein LOC621058
(LOC621058), mRNA" ---
1445584_at 0.96227 --- ---
1445585_at 0.95848 zinc finger protein 313 Zfp313
1445586_at 0.95312 Ca<2+>dependent activator protein for secretion Cadps
1445587_at 0.97104 --- ---
1445588_at 0.99289 ankyrin repeat domain 44 Ankrd44
1445589_at 1.12874 "Solute carrier family 23 (nucleobase transporters), member 2" Slc23a2
1445590_at 0.89148 Transcribed locus ---
1445591_at 0.92911 Sarcosine dehydrogenase Sardh
1445592_at 1.04522 --- ---
1445593_at 1.19547 "Adult male testis cDNA, RIKEN full-length enriched library,
clone:4930414J05 product:unclassifiable, full insert sequence" ---
1445594_at 1.10072 Heterogeneous nuclear ribonucleoprotein L Hnrpl
1445595_at 1.10114 Transcribed locus ---
1445596_at 1.00104 "Adenosine deaminase, RNA-specific" Adar

1445597_s_at	0.90638	HRAS like suppressor 3	Hrasls3
1445598_at	1.17520	UDP-N-acetyl-alpha-D-galactosamine: polypeptide N-acetylgalactosaminyltransferase 7	Galnt7
1445599_at	1.00944	fibronectin type III domain containing 7	Fndc7
1445600_at	1.89417	Insulin-like growth factor I receptor	Igf1r
1445601_at	1.09519	"6 days neonate spleen cDNA, RIKEN full-length enriched library, clone: F430214H02 product:unclassifiable, full insert sequence"	---
1445602_at	0.98007	RIKEN cDNA A630055G03 gene	A630055G03Rik
1445603_at	1.01549	Sorcin	Sri
1445604_at	1.06624	Transcribed locus	---
1445605_s_at	1.04002	RIKEN cDNA 4921533L14 gene	4921533L14Rik
1445606_a_at	0.85298	RIKEN cDNA 2900009J06 gene	2900009J06Rik
1445607_at	1.01481	RIKEN cDNA 4833418A01 gene	4833418A01Rik
1445608_at	0.96142	RIKEN cDNA 9130204L05 gene	9130204L05Rik
1445609_at	0.88126	Transcribed locus	---
1445610_at	0.94323	CDNA sequence BC039093	BC039093
1445611_at	1.08779	RIKEN cDNA 1810044A24 gene	1810044A24Rik
1445612_at	1.05628	---	---
1445613_at	1.01370	"Nuclear receptor subfamily 2, group C, member 2"	Nr2c2
1445614_at	0.86099	---	---
1445615_at	1.01743	---	---
1445616_at	1.14584	"Eukaryotic translation initiation factor 4 gamma, 3"	Eif4g3
1445617_at	0.93640	Transcribed locus	---
1445618_at	0.95128	---	---
1445619_at	1.06737	---	---
1445620_at	0.88765	Transcribed locus	---
1445621_at	1.02222	---	---
1445622_at	0.87654	---	---
1445623_at	0.97888	---	---
1445624_at	0.99880	"DNA segment, Chr 8, ERATO Doi 587, expressed"	D8Ertd587e
1445625_at	0.86866	Transcribed locus	---
1445626_at	1.67058	"Lectin, galactose binding, soluble 3"	Lgals3
1445627_at	1.26743	Oxysterol binding protein-like 9	Osbpl9
1445628_at	0.93189	Rho GTPase activating protein 21	Arhgap21
1445629_at	1.69181	"Solute carrier family 35 (UDP-glucuronic acid/UDP-N-acetylgalactosamine dual transporter), member D1"	Slc35d1
1445630_at	1.05682	---	---
1445631_at	0.98373	---	---
1445632_at	1.02884	oxoglutarate dehydrogenase (lipoamide)	Ogdh
1445633_at	1.11573	Reduced expression 2	Rex2
1445634_at	1.17423	Microtubule-associated protein tau	Mapt
1445635_at	0.86871	Eph receptor A7	Epha7
1445636_at	0.99097	Formin-like 1	Fmn11

1445637_at 0.88149 DNA methyltransferase (cytosine-5) 1 Dnmt1
1445638_at 1.07116 Myeloid ecotropic viral integration site-related gene 1 Mrg1
1445639_at 0.92060 RIKEN cDNA 9130014G24 gene 9130014G24Rik
1445640_at 1.06695 Holocarboxylase synthetase (biotin- [propionyl-Coenzyme A-carboxylase (ATP-hydrolysing)] ligase) Hlcs
1445641_at 1.95357 "ELOVL family member 6, elongation of long chain fatty acids (yeast)" Elovl6
1445642_at 0.87641 LEM domain containing 1 Lemd1
1445643_at 1.01183 "PREDICTED: Mus musculus similar to spindle assembly associated Sfi1 homolog isoform b (LOC621860), mRNA" ---
1445644_at 0.91168 RIKEN cDNA A830091I15 gene A830091I15Rik
1445645_at 1.01691 A disintegrin and metallopeptidase domain 22 Adam22
1445646_at 1.42849 BTB (POZ) domain containing 1 Btbd1
1445647_at 0.92796 expressed sequence AI447881 AI447881
1445648_at 0.94984 Transcribed locus ---
1445649_x_at 0.97263 zinc finger protein 142 Zfp142
1445650_at 1.01123 --- ---
1445651_at 1.29318 "ATPase, Class VI, type 11B" Atp11b
1445652_at 1.00329 Stromal antigen 1 Stag1
1445653_at 0.91700 cDNA sequence BC031361 BC031361
1445654_at 1.04589 RIKEN cDNA 4930422I07 gene 4930422I07Rik
1445655_at 0.97376 Transcribed locus ---
1445656_at 0.93756 Myelin-associated oligodendrocytic basic protein Mobp
1445657_at 1.02436 expressed sequence AW111846 AW111846
1445658_at 0.93478 Transcribed locus ---
1445659_at 1.00002 Protective protein for beta-galactosidase Ppqb
1445660_at 1.29015 RIKEN cDNA 2700094F01 gene 2700094F01Rik
1445661_at 0.90873 RIKEN cDNA 4930406D18 gene 4930406D18Rik
1445662_x_at 0.93335 Transcribed locus ---
1445663_at 0.94248 --- ---
1445664_at 1.84285 Insulin-like growth factor I receptor Igf1r
1445665_at 1.28683 Septin 6 6-Sep
1445666_at 0.96660 "DNA segment, Chr 15, Wayne State University 169, expressed" D15Wsu169e
1445667_at 0.94619 --- ---
1445668_at 1.19372 Tubulin-specific chaperone e Tbce
1445669_at 1.16033 sprouty homolog 4 (Drosophila) Spry4
1445670_at 0.76135 --- ---
1445671_at 0.95593 Transcribed locus ---
1445672_at 1.07367 Transcribed locus ---
1445673_at 0.98743 RIKEN cDNA 2900052N01 gene 2900052N01Rik
1445674_at 1.05016 Nitric oxide synthase 3 antisense Nos3as
1445675_at 0.98178 Transcribed locus ---
1445676_at 1.10094 "Potassium intermediate/small conductance calcium-activated channel, subfamily N, member 2" Kcnn2

1445677_x_at 0.96841 "solute carrier family 35, member F2" Slc35f2
1445678_at 1.08611 --- ---
1445679_at 0.94454 "ATPase, Ca++ transporting, plasma membrane 3" Atp2b3
1445680_x_at 1.00382 coagulation factor II (thrombin) receptor-like 2 F2rl2
1445681_at 1.32284 Cell division cycle associated 7 Cdca7
1445682_at 0.88183 RIKEN cDNA 2700099C18 gene 2700099C18Rik
1445683_at 1.13469 --- ---
1445684_s_at 0.92470 histone deacetylase 2 Hdac2
1445685_at 1.00086 growth arrest specific 7 Gas7
1445686_at 1.07336 "12 days embryo spinal ganglion cDNA, RIKEN full-length enriched library, clone:D130056G12 product:unclassifiable, full insert sequence" ---
1445687_at 1.04060 "gene model 885, (NCBI)" Gm885
1445688_at 0.77689 host cell factor C1 Hcfc1
1445689_at 2.64233 Transcribed locus ---
1445690_at 0.95576 --- ---
1445691_at 0.97336 chimerin (chimaerin) 1 Chn1
1445692_x_at 0.82798 --- ---
1445693_at 1.05034 v-raf murine sarcoma 3611 viral oncogene homolog Araf
1445694_at 0.99006 Transcribed locus ---
1445695_at 1.34414 Ataxin 1 Atxn1
1445696_x_at 0.93178 Transcribed locus ---
1445697_at 1.23448 Integrin beta 1 binding protein 1 Itgb1bp1
1445698_at 1.05019 calpain 11 Capn11
1445699_at 0.99130 --- ---
1445700_at 1.01552 RIKEN cDNA 2310047M15 gene 2310047M15Rik
1445701_at 0.98621 "ATPase, Ca++ transporting, plasma membrane 4" Atp2b4
1445702_x_at 1.05025 phosphatidic acid phosphatase type 2 domain containing 2 Ppapdc2
1445703_at 0.91171 expressed sequence AI848149 AI848149
1445704_x_at 1.03386 RIKEN cDNA 2900093B09 gene 2900093B09Rik
1445705_x_at 0.92055 dipeptidylpeptidase 8 Dpp8
1445706_x_at 0.96744 --- ---
1445707_at 0.92995 "zinc finger, DHHC domain containing 6 /// similar to Probable palmitoyltransferase ZDHHC6 (Zinc finger DHHC domain-containing protein 6) (DHHC-6) (H4 homolog)" Zdhhc6 /// LOC669170
1445708_x_at 0.92071 RIKEN cDNA 3110021A11 gene 3110021A11Rik
1445709_at 1.10780 transformed mouse 3T3 cell double minute 1 Mdm1
1445710_x_at 1.05347 RIKEN cDNA 1110051B16 gene 1110051B16Rik
1445711_at 1.02623 expressed sequence BB163080 BB163080
1445712_at 0.97236 Transcribed locus ---
1445713_at 0.92448 RIKEN cDNA 9630041G16 gene 9630041G16Rik
1445714_at 1.01871 RIKEN cDNA B930042K01 gene B930042K01Rik
1445715_at 1.06746 RIKEN cDNA A230001M10 gene A230001M10Rik
1445716_at 1.00151 --- ---

1445717_at 1.48133 RIKEN cDNA E130108L08 gene E130108L08Rik
1445718_at 1.53989 WD repeat and FYVE domain containing 3 Wdfy3
1445719_at 1.02839 "Splicing factor 3a, subunit 1" Sf3a1
1445720_at 1.03891 "SAFB-like, transcription modulator" Sltm
1445721_at 0.93951 hypothetical protein A830021M18 A830021M18
1445722_at 0.97768 "mannosidase, beta A, lysosomal-like" Manbal
1445723_at 0.92945 phospholipase C-like 1 Plcl1
1445724_at 2.00478 IQ motif containing GTPase activating protein 1 Iqgap1
1445725_at 0.85054 --- ---
1445726_at 1.06759 RIKEN cDNA C330011K17 gene C330011K17Rik
1445727_at 1.37102 ubiquitin protein ligase E3A Ube3a
1445728_at 0.85112 --- ---
1445729_at 0.97840 "DnaJ (Hsp40) homolog, subfamily A, member 1" Dnaja1
1445730_at 0.92481 RAB11 family interacting protein 2 (class I) Rab11fip2
1445731_at 0.96069 "Phosphatidylinositol 3-kinase, catalytic, beta polypeptide" Pik3cb
1445732_at 1.00383 --- ---
1445733_at 0.94665 RIKEN cDNA 2610206G21 gene 2610206G21Rik
1445734_at 0.92677 L-threonine dehydrogenase Tdh
1445735_at 1.09537 Hyaluronan and proteoglycan link protein 1 Hapln1
1445736_at 1.12677 --- ---
1445737_at 1.01434 RIKEN cDNA 1700057H21 gene 1700057H21Rik
1445738_at 0.98200 --- ---
1445739_at 0.95570 RIKEN cDNA 9430080K19 gene 9430080K19Rik
1445740_at 0.95585 Transcribed locus ---
1445741_at 0.91435 protein phosphatase 1E (PP2C domain containing) Ppm1e
1445742_at 0.96874 "Smg-6 homolog, nonsense mediated mRNA decay factor (C. elegans)"
Smg6
1445743_at 1.05710 RIKEN cDNA E330024J20 gene E330024J20Rik
1445744_at 1.05073 Transmembrane protein 56 Tmem56
1445745_at 0.98366 RNA binding motif protein 17 /// similar to RNA binding motif protein 17
Rbm17 /// LOC545866
1445746_at 1.12416 hypothetical protein LOC630474 /// hypothetical protein LOC630475 ///
hypothetical LOC638081 LOC630474 /// LOC630475 /// LOC638081
1445747_at 1.10263 "15 days embryo male testis cDNA, RIKEN full-length enriched library,
clone:8030444H21 product:unclassifiable, full insert sequence" ---
1445748_at 1.08850 "CD3 antigen, epsilon polypeptide" Cd3e
1445749_at 1.03357 Ring finger protein 4 Rnf4
1445750_at 0.93027 Transcribed locus ---
1445751_at 1.07703 "Procollagen, type XII, alpha 1" Col12a1
1445752_at 1.08871 --- ---
1445753_at 1.07562 Aldehyde oxidase 3-like 1 Aox3l1
1445754_at 0.97377 Acetyl-Coenzyme A carboxylase beta Acacb
1445755_at 0.94788 Eyes absent 2 homolog (Drosophila) Eya2

1445756_at 0.96647 Transcribed locus ---
 1445757_at 0.93550 T-box 3 Tbx3
 1445758_at 1.07992 CDNA sequence BC049816 BC049816
 1445759_at 1.27608 "Sphingomyelin phosphodiesterase 3, neutral" Smpd3
 1445760_at 1.02423 antimicrobial peptide RYA3 Rya3
 1445761_at 0.92101 "DNA segment, Chr 4, ERATO Doi 628, expressed" D4Ertd628e
 1445762_at 0.96633 "Serine protease inhibitor, Kunitz type 1" Spint1
 1445763_at 1.09449 RIKEN cDNA 1700013F07 gene 1700013F07Rik
 1445764_at 0.82721 "PREDICTED: Mus musculus hypothetical protein LOC628151 (LOC628151), mRNA" ---
 1445765_at 0.99744 RIKEN cDNA 9430011C21 gene 9430011C21Rik
 1445766_at 1.05158 "Protein tyrosine phosphatase, receptor type, f polypeptide (PTPRF), interacting protein (liprin), alpha 2" Ppfia2
 1445767_at 0.92934 "protein tyrosine phosphatase, receptor type, D" Ptprd
 1445768_at 0.97926 similar to Ankyrin repeat domain protein 11 (Ankyrin repeat-containing cofactor 1) LOC625662
 1445769_at 0.92764 Hypothetical LOC546862 LOC546862
 1445770_at 1.03523 --- ---
 1445771_at 0.87082 --- ---
 1445772_at 0.91806 "DNA segment, Chr 16, ERATO Doi 550, expressed" D16Ertd550e
 1445773_at 1.46340 Myeloid ecotropic viral integration site 1 Meis1
 1445774_at 0.91765 "Potassium large conductance calcium-activated channel, subfamily M, alpha member 1" Kcnma1
 1445775_at 0.97354 --- ---
 1445776_at 1.16975 --- ---
 1445777_at 1.01305 UPF2 regulator of nonsense transcripts homolog (yeast) Upf2
 1445778_at 1.26996 E74-like factor 4 (ets domain transcription factor) Elf4
 1445779_at 0.79755 "Glutamate receptor, ionotropic, delta 1" Grid1
 1445780_at 0.87690 --- ---
 1445781_at 2.16882 SH3 domain protein D19 Sh3d19
 1445782_at 0.95772 --- ---
 1445783_at 1.16953 --- ---
 1445784_at 0.99165 RIKEN cDNA 4933435A13 gene 4933435A13Rik
 1445785_at 0.93021 --- ---
 1445786_at 1.56724 Braf transforming gene Braf
 1445787_at 1.06368 RIKEN cDNA 5033413D22 gene 5033413D22Rik
 1445788_at 0.99304 "Insulin-like growth factor 2, binding protein 1" Igf2bp1
 1445789_at 0.91610 "0 day neonate kidney cDNA, RIKEN full-length enriched library, clone: D630046H12 product:unclassifiable, full insert sequence" ---
 1445790_at 1.02486 gastric inhibitory polypeptide receptor Gipr
 1445791_at 1.08299 --- ---
 1445792_at 1.08582 Protein phosphatase methylesterase 1 Ppme1
 1445793_at 1.01100 RIKEN cDNA 1700011I03 gene 1700011I03Rik

1445794_at 1.00599 Sperm associated antigen 16 Spag16
1445795_at 1.16241 choline kinase alpha Chka
1445796_at 1.03791 Expressed sequence AI315068 AI315068
1445797_at 0.94699 Peptidylglycine alpha-amidating monooxygenase Pam
1445798_at 1.78680 "discs, large homolog 1 (Drosophila)" Dlgh1
1445799_at 0.94231 --- ---
1445800_at 0.98508 AU RNA binding protein/enoyl-coenzyme A hydratase Auh
1445801_at 0.90709 CDNA sequence AK129341 AK129341
1445802_at 0.90843 Expressed sequence AU017455 AU017455
1445803_at 1.11845 Diaphanous homolog 3 (Drosophila) Diap3
1445804_at 0.96663 RIKEN cDNA 9630020C08 gene 9630020C08Rik
1445805_x_at 0.95845 "potassium voltage-gated channel, subfamily H (eag-related), member 3"
Kcnh3
1445806_at 0.99749 component of oligomeric golgi complex 5 Cog5
1445807_at 1.04404 "Protein kinase, cGMP-dependent, type I" Prkg1
1445808_at 1.05056 --- ---
1445809_at 0.96841 Eyes absent 4 homolog (Drosophila) Eya4
1445810_at 0.95629 expressed sequence C85163 C85163
1445811_at 1.04036 "Neural precursor cell expressed, developmentally down-regulated gene 8"
Nedd8
1445812_at 0.95024 --- ---
1445813_at 1.02868 WD repeat domain 27 Wdr27
1445814_at 0.90594 "Adult male testis cDNA, RIKEN full-length enriched library,
clone:4930402N10 product:unclassifiable, full insert sequence" ---
1445815_at 0.90450 frizzled homolog 8 (Drosophila) Fzd8
1445816_at 1.02729 --- ---
1445817_at 0.95035 expressed sequence BB168181 BB168181
1445818_at 1.02572 RIKEN cDNA 4930544M13 gene 4930544M13Rik
1445819_at 0.93366 RIKEN cDNA A830018L16 gene A830018L16Rik
1445820_at 0.97562 --- ---
1445821_at 1.04409 Interleukin 1 receptor accessory protein-like 1 Il1rap1l
1445822_at 1.08809 similar to CG18631-PA /// similar to CG18631-PA /// similar to CG18631-
PA LOC639112 /// LOC668310 /// LOC676937
1445823_at 0.99283 Coiled-coil domain containing 91 Ccdc91
1445824_at 0.81384 zinc finger protein 458 Zfp458
1445825_at 0.94632 --- ---
1445826_at 1.13084 ankyrin repeat domain 17 Ankrd17
1445827_at 1.26360 Protein kinase C binding protein 1 Prkcbp1
1445828_at 1.08531 --- ---
1445829_at 0.94840 RIKEN cDNA 8030491N06 gene 8030491N06Rik
1445830_at 1.60225 "catenin (cadherin associated protein), delta 1" Ctnnd1
1445831_at 1.40852 --- ---
1445832_at 0.97285 --- ---

1445833_at 0.93156 RIKEN cDNA 2400010G15 gene 2400010G15Rik
1445834_at 0.87839 Transcribed locus ---
1445835_at 1.00422 Transcribed locus ---
1445836_at 1.11568 Zinc finger protein 516 Zfp516
1445837_at 0.91376 Dendritic cell protein GA17 Ga17
1445838_at 0.95002 UDP-N-acetyl-alpha-D-galactosamine:polypeptide N-acetylgalactosaminyltransferase 5 Galnt5
1445839_at 0.94657 --- ---
1445840_at 1.02848 Transcribed locus ---
1445841_at 1.04609 leucine rich repeat containing 39 Lrrc39
1445842_at 0.94804 --- ---
1445843_at 0.92316 chromodomain helicase DNA binding protein 2 Chd2
1445844_at 1.01440 "Adult male testis cDNA, RIKEN full-length enriched library, clone:4930547H19 product:unclassifiable, full insert sequence" ---
1445845_at 0.95009 Transcribed locus ---
1445846_at 0.92605 "Adult male adrenal gland cDNA, RIKEN full-length enriched library, clone:7330409L06 product:unclassifiable, full insert sequence" ---
1445847_at 0.97805 --- ---
1445848_at 1.10582 hypothetical protein 9630009C16 9630009C16
1445849_at 0.93843 similar to PRAME family member 8 /// similar to PRAME family member 8 /// similar to PRAME family member 8 /// similar to PRAME family member 8 LOC665521 /// LOC665546 /// LOC665755 /// LOC666203
1445850_at 1.02996 RIKEN cDNA 9530009M10 gene 9530009M10Rik
1445851_at 0.98183 --- ---
1445852_at 0.97127 "DNA segment, Chr 19, ERATO Doi 200, expressed" D19Ertd200e
1445853_at 1.02265 --- ---
1445854_at 0.86387 RIKEN cDNA C230004F18 gene C230004F18Rik
1445855_at 1.03385 RIKEN cDNA A030007L17 gene A030007L17Rik
1445856_at 0.98542 RIKEN cDNA 6030443O07 gene 6030443O07Rik
1445857_at 1.04292 cDNA sequence BC026585 BC026585
1445858_at 1.09866 RIKEN cDNA 4933439G19 gene 4933439G19Rik
1445859_at 1.01527 B-cell CLL/lymphoma 9 Bcl9
1445860_at 0.95245 tetraspanin 11 Tspan11
1445861_at 1.22368 Ubiquitin specific peptidase 25 Usp25
1445862_at 1.87424 RIKEN cDNA 4921513D23 gene 4921513D23Rik
1445863_at 0.89692 RIKEN cDNA E030013G06 gene E030013G06Rik
1445864_at 0.94716 Transcribed locus ---
1445865_at 0.93390 --- ---
1445866_at 1.71653 microtubule associated serine/threonine kinase family member 4 Mast4
1445867_at 1.37488 expressed sequence AL023008 AL023008
1445868_at 0.98529 cytoplasmic polyadenylation element binding protein 3 Cpeb3
1445869_at 0.91502 Transmembrane protein 28 Tmem28
1445870_at 1.40481 CDC42 effector protein (Rho GTPase binding) 2 Cdc42ep2

1445871_at	0.98870	expressed sequence	AI315376	AI315376
1445872_at	0.93063	"nuclear receptor subfamily 6, group A, member 1"		Nr6a1
1445873_at	1.28439	Transcription factor Dp 2	Tfdp2	
1445874_at	1.04444	RAR-related orphan receptor beta		Rorb
1445875_at	0.95342	RIKEN cDNA A230057D06	gene	A230057D06Rik
1445876_at	0.90279	---	---	
1445877_at	1.01424	expressed sequence	AI256775	AI256775
1445878_at	0.93364	RIKEN cDNA C920006O11	gene	C920006O11Rik
1445879_at	0.87894	zinc finger protein 507	zfp507	
1445880_at	1.06870	Deleted in colorectal carcinoma	Dcc	
1445881_at	0.91087	RIKEN cDNA 2310035P21	gene	2310035P21Rik
1445882_at	0.84262	CD300 antigen like family member B		Cd300lb
1445883_at	0.96427	RAN binding protein 2	Ranbp2	
1445884_at	1.49066	RIKEN cDNA 1110007A13	gene	1110007A13Rik
1445885_at	1.71118	ubiquitin-conjugating enzyme E2D 2		Ube2d2
1445886_at	1.17450	"ELK3, member of ETS oncogene family"		Elk3
1445887_at	0.99211	---	---	
1445888_x_at	1.16034	"poly (ADP-ribose) polymerase family, member 3"		Parp3
1445889_at	1.13768	"Sortilin-related receptor, LDLR class A repeats-containing"		Sorl1
1445890_at	0.94400	RIKEN cDNA C230079E05	gene	C230079E05Rik
1445891_at	0.89053	cDNA sequence AF529169	AF529169	
1445892_at	1.01066	Period homolog 2 (Drosophila)		Per2
1445893_at	1.15136	"procollagen-lysine, 2-oxoglutarate 5-dioxygenase 1"		Plod1
1445894_at	1.01774	---	---	
1445895_at	0.92463	"T-cell receptor beta, variable 8.2"		Tcrb-V8.2
1445896_at	1.01104	RIKEN cDNA 2810433D01	gene	2810433D01Rik
1445897_s_at	1.12850	interferon-induced protein 35		Ifi35
1445898_at	0.87394	gamma-glutamyl carboxylase		Ggcx
1445899_at	1.09675	expressed sequence AA409289	AA409289	
1445900_at	1.17104	"Kalirin, RhoGEF kinase"		Kalrn
1445901_at	1.00217	low density lipoprotein receptor-related protein 2		Lrp2
1445902_at	0.96642	"Tetratricopeptide repeat, ankyrin repeat and coiled-coil containing 2"		Tanc2
1445903_at	1.12805	---	---	
1445904_at	1.03274	---	---	
1445905_at	0.85156	expressed sequence C79206	C79206	
1445906_at	1.04601	Zinc finger protein 35	Zfp35	
1445907_at	1.02136	---	---	
1445908_at	0.86415	"Eukaryotic translation initiation factor 2B, subunit 5 epsilon"		Eif2b5
1445909_at	0.91843	---	---	
1445910_at	0.92476	expressed sequence C79999	C79999	
1445911_at	1.00634	Cysteine-rich hydrophobic domain 2		Chic2
1445912_at	1.02022	coiled-coil domain containing 60		Ccdc60
1445913_at	0.86493	RIKEN cDNA A730090H04	gene	A730090H04Rik

1445914_at 1.61355 Nuclear respiratory factor 1 Nrf1
1445915_at 0.91348 "Histocompatibility 2, K1, K region" H2-K1
1445916_at 1.04355 "ATPase, H+ transporting, lysosomal V1 subunit A" Atp6v1a
1445917_at 1.09124 expressed sequence C81269 C81269
1445918_at 1.49464 Transmembrane protein 2 Tmem2
1445919_at 1.13474 expressed sequence AA409261 AA409261
1445920_at 0.92279 --- ---
1445921_at 0.96259 "DNA segment, Chr 2, ERATO Doi 397, expressed" D2Ertd397e
1445922_at 1.21507 X-ray repair complementing defective repair in Chinese hamster cells 4 Xrcc4
1445923_at 0.84862 RIKEN cDNA 1700055N04 gene 1700055N04Rik
1445924_at 0.89733 similar to RIKEN cDNA 1700029I01 LOC385211
1445925_at 0.87294 "DNA segment, Chr 15, ERATO Doi 492, expressed" D15Ertd492e
1445926_at 1.05658 --- ---
1445927_at 1.03533 "DNA segment, Chr 13, ERATO Doi 212, expressed" D13Ertd212e
1445928_at 1.59292 membrane-associated ring finger (C3HC4) 6 6-Mar
1445929_at 1.00566 "Potassium channel, subfamily K, member 2" Kcnk2
1445930_at 1.00344 fibronectin type III domain containing 7 Fndc7
1445931_at 0.98300 --- ---
1445932_at 0.89270 Cadherin EGF LAG seven-pass G-type receptor 1 Celsr1
1445933_at 1.10543 "solute carrier family 12, member 5" Slc12a5
1445934_at 1.14390 ariadne homolog 2 (Drosophila) /// similar to ariadne homolog 2 Arih2 ///
LOC432565
1445935_at 0.99430 --- ---
1445936_at 1.09831 Superkiller viralicidic activity 2-like 2 (S. cerevisiae) Skiv2l2
1445937_at 0.99086 RIKEN cDNA A630001G21 gene A630001G21Rik
1445938_at 1.00315 RIKEN cDNA 5930427L02 gene 5930427L02Rik
1445939_at 0.99654 Mitochondrial tumor suppressor 1 Mtus1
1445940_at 1.39358 "DNA segment, Chr 4, ERATO Doi 298, expressed" D4Ertd298e
1445941_at 1.35164 AT motif binding factor 1 Atbf1
1445942_at 0.94866 expressed sequence AU015858 AU015858
1445943_at 1.07752 --- ---
1445944_at 0.89336 "solute carrier family 25 (mitochondrial carrier, adenine nucleotide translocator), member 5" Slc25a5
1445945_at 0.97133 expressed sequence AU021880 AU021880
1445946_at 0.88666 "Phosphodiesterase 4A, cAMP specific" Pde4a
1445947_at 1.18576 Kinesin 2 Kns2
1445948_at 0.96024 Transcribed locus ---
1445949_at 0.88934 "Tumor necrosis factor receptor superfamily, member 19" Tnfrsf19
1445950_at 0.90587 "cDNA sequence, C87926" C87926
1445951_at 1.00597 expressed sequence C86345 C86345
1445952_at 2.48664 "Membrane associated guanylate kinase, WW and PDZ domain containing 1" Magi1
1445953_at 0.86596 "DNA segment, Chr 17, ERATO Doi 191, expressed" D17Ertd191e

1445954_at	0.97343	expressed sequence C81354	C81354
1445955_at	0.88072	RIKEN cDNA 2400010G15 gene	2400010G15Rik
1445956_at	0.97824	---	---
1445957_at	1.55931	Ena-vasodilator stimulated phosphoprotein	Evl
1445958_at	0.94418	Transcribed locus	---
1445959_at	1.12659	---	---
1445960_at	0.97724	---	---
1445961_at	1.24649	---	---
1445962_at	0.97814	"Glutamate receptor, ionotropic, AMPA2 (alpha 2)"	Gria2
1445963_at	1.00098	"Phosphodiesterase 5A, cGMP-specific"	Pde5a
1445964_at	1.01589	---	---
1445965_at	2.14048	---	---
1445966_at	2.04428	Insulin-like growth factor 2 receptor	Igf2r
1445967_at	0.95460	expressed sequence AU022331	AU022331
1445968_at	0.96354	Transmembrane and tetratricopeptide repeat containing 2	Tmtc2
1445969_at	1.04029	Transcribed locus	---
1445970_at	0.94630	expressed sequence AU022526	AU022526
1445971_at	0.92119	RIKEN cDNA 4933439F18 gene	4933439F18Rik
1445972_at	0.95158	"Usher syndrome 2A (autosomal recessive, mild) homolog (human)"	Ush2a
1445973_at	0.94362	expressed sequence C79461	C79461
1445974_at	0.80943	Transcribed locus	---
1445975_at	1.01858	---	---
1445976_at	0.91362	expressed sequence C80279	C80279
1445977_at	0.97109	expressed sequence AU022793	AU022793
1445978_at	0.98259	---	---
1445979_at	0.95483	Cyclin M1	Cnm1
1445980_at	0.88405	"Aldehyde dehydrogenase family 1, subfamily A1"	Aldh1a1
1445981_at	0.97913	---	---
1445982_at	1.03050	Glypican 6	Gpc6
1445983_at	0.91856	NF-kappaB repressing factor	Nkrf
1445984_at	1.82730	PFTAIRE protein kinase 1	Pftk1
1445985_at	1.40863	Traf3 interacting protein 2	Traf3ip2
1445986_at	1.07440	expressed sequence C86195	C86195
1445987_at	1.13741	Transcribed locus	---
1445988_at	1.02543	Interleukin 17D Il17d	
1445989_at	0.95400	Transcribed locus	---
1445990_at	1.06942	expressed sequence AU020096	AU020096
1445991_at	0.83897	Lipoma HMGIC fusion partner	Lhfp
1445992_at	0.97350	Transcribed locus	---
1445993_at	0.92536	---	---
1445994_at	0.90207	---	---
1445995_at	1.16070	cell division cycle 25 homolog A (S. cerevisiae)	Cdc25a
1445996_at	0.92185	Placenta specific 9	Plac9

1445997_at 0.99131 ets variant gene 3 Etv3
1445998_at 0.97082 "Protein kinase, cGMP-dependent, type I" Prkg1
1445999_at 1.19092 "DNA segment, Chr 19, ERATO Doi 132, expressed" D19Ertd132e
1446000_at 0.94873 Zinc finger protein 291 Zfp291
1446001_at 2.49587 RalBP1 associated Eps domain containing protein Repl
1446002_at 0.90967 "ATP-binding cassette, sub-family C (CFTR/MRP), member 1" Abcc1
1446003_at 0.83501 --- ---
1446004_at 0.96655 "ATPase, Cu++ transporting, beta polypeptide" Atp7b
1446005_at 0.82888 "DNA segment, Chr 2, ERATO Doi 337, expressed" D2Ertd337e
1446006_at 0.99509 IMP1 inner mitochondrial membrane peptidase-like (S. cerevisiae) Immp11
1446007_at 1.33846 Bone morphogenetic protein 1 Bmp1
1446008_at 1.07292 "DNA segment, Chr 3, ERATO Doi 246, expressed" D3Ertd246e
1446009_at 1.04034 Similar to Potassium voltage-gated channel subfamily G member 2 (Voltage-gated potassium channel subunit Kv6.2) (Cardiac potassium channel subunit) LOC240444
1446010_at 0.92391 similar to F-box only protein 34 /// similar to F-box only protein 34 /// similar to F-box only protein 34 LOC668019 /// LOC670109 /// LOC674275
1446011_at 0.96779 "Core 1 UDP-galactose:N-acetylgalactosamine-alpha-R beta 1,3-galactosyltransferase" C1galt1
1446012_at 1.15297 "Dystrobrevin, beta" Dtnb
1446013_at 1.04666 "solute carrier family 38, member 3" Slc38a3
1446014_at 1.22240 --- ---
1446015_at 1.03383 "Trafficking protein, kinesin binding 1" Trak1
1446016_at 1.01281 --- ---
1446017_at 1.03368 Transcribed locus ---
1446018_at 0.99406 expressed sequence C79240 C79240
1446019_at 1.08806 Transcribed locus ---
1446020_at 0.88753 --- ---
1446021_at 0.96399 expressed sequence AU018466 AU018466
1446022_at 1.04398 Musashi homolog 2 (Drosophila) Msi2
1446023_at 0.99572 Osteoclast stimulating factor 1 Ostf1
1446024_at 0.92838 --- ---
1446025_at 1.05115 Expressed sequence AI462493 AI462493
1446026_at 1.21909 Kinesin family member 13A Kif13a
1446027_at 0.91346 --- ---
1446028_at 0.97471 --- ---
1446029_at 1.57853 "Tetratricopeptide repeat, ankyrin repeat and coiled-coil containing 1" Tanc1
1446030_at 0.87027 Transcribed locus ---
1446031_at 0.93633 Thyroid hormone receptor associated protein 3 Thrsp3
1446032_at 1.02040 --- ---
1446033_at 0.93780 expressed sequence C87882 C87882
1446034_at 0.93990 "Adult male cortex cDNA, RIKEN full-length enriched library, clone: B530045B12 product:unclassifiable, full insert sequence" ---
1446035_at 0.94848 "PREDICTED: Mus musculus similar to Protein CXorf22 (LOC629131),

mRNA" ---

1446036_at	0.97669 --- ---	
1446037_at	1.08356	RIKEN cDNA C230081A13 gene C230081A13Rik
1446038_at	1.06051	Heterogeneous nuclear ribonucleoprotein A1 Hnrpa1
1446039_at	0.86509	Transcribed locus ---
1446040_at	0.84747	RIKEN cDNA 2700049A03 gene 2700049A03Rik
1446041_at	0.99753	Transcribed locus ---
1446042_at	0.94304	"PREDICTED: Mus musculus similar to preferentially expressed antigen in melanoma-like 3 (LOC621352), mRNA" ---
1446043_at	1.10313	expressed sequence C81086 C81086
1446044_at	0.92064	expressed sequence AU015581 AU015581
1446045_at	0.90813	expressed sequence AU024581 AU024581
1446046_at	0.92646	CDNA sequence BC023818 BC023818
1446047_at	0.95195	"RNA binding motif, single stranded interacting protein" Rbms3
1446048_at	0.96158 --- ---	
1446049_at	1.01212 --- ---	
1446050_at	1.16557	"Membrane associated guanylate kinase, WW and PDZ domain containing 1" Magi1
1446051_at	1.06398	"gene model 1140, (NCBI)" Gm1140
1446052_at	1.02176 --- ---	
1446053_at	1.11131	G patch domain and KOW motifs Gpkow
1446054_at	0.99525	"Zinc metallopeptidase, STE24 homolog (S. cerevisiae)" Zmpste24
1446055_at	1.09241 --- ---	
1446056_at	1.14132 --- ---	
1446057_at	1.93592	rearranged L-myc fusion sequence Rlf
1446058_at	0.84937	"PREDICTED: Mus musculus hypothetical protein LOC628925 (LOC628925), mRNA" ---
1446059_at	0.88814	"ST6 (alpha-N-acetyl-neuraminy1-2,3-beta-galactosyl-1,3)-N-acetylgalactosaminide alpha-2,6-sialyltransferase 5" St6galnac5
1446060_at	0.97738	RIKEN cDNA 9230109A22 gene 9230109A22Rik
1446061_at	1.04165	CDNA sequence BC042698 BC042698
1446062_at	0.97491	RIKEN cDNA B830028B13 gene B830028B13Rik
1446063_at	0.87815	RIKEN cDNA C730034F03 gene C730034F03Rik
1446064_at	0.99490 --- ---	
1446065_at	0.96779	"Potassium voltage-gated channel, Shal-related family, member 2" Kcnd2
1446066_at	0.87083	"zinc finger, CCHC domain containing 17" Zcchc17
1446067_at	1.06919	Piwi-like homolog 2 (Drosophila) Piwil2
1446068_at	1.23052	adenosine kinase Adk
1446069_at	1.03363	"Actin binding LIM protein family, member 3" Ablim3
1446070_at	1.01540	tRNA-yW synthesizing protein 1 homolog (S. cerevisiae) Tyw1
1446071_at	1.23041	six transmembrane epithelial antigen of prostate 2 Steap2
1446072_at	1.08655 --- ---	
1446073_at	0.99233	YY1 associated factor 2 Yaf2

1446074_at 0.96782 MOCO sulphurase C-terminal domain containing 1 Mosc1
1446075_at 1.14249 --- ---
1446076_at 0.93732 Transcribed locus ---
1446077_at 0.87251 RIKEN cDNA 4833446K15 gene 4833446K15Rik
1446078_at 0.91717 FK506 binding protein 7 Fkbp7
1446079_at 1.11336 "zinc finger, matrin type 2" Zmat2
1446080_at 1.04947 Cut-like 2 (Drosophila) Cutl2
1446081_at 1.06609 "Zinc finger, ZZ-type with EF hand domain 1" Zzef1
1446082_at 0.97540 "Jumonji, AT rich interactive domain 2" Jarid2
1446083_at 1.02576 Heterogeneous nuclear ribonucleoprotein D Hnrpd
1446084_at 1.03449 Nexilin Nexn
1446085_at 1.02724 Suppressor of cytokine signaling 2 Socs2
1446086_s_at 0.88084 GLI-Kruppel family member GLI2 Gli2
1446087_at 0.96644 "Golgi autoantigen, golgin subfamily a, 1" Golga1
1446088_at 1.64009 RIKEN cDNA 9430081I23 gene 9430081I23Rik
1446089_at 1.07512 expressed sequence C86400 C86400
1446090_at 1.18543 "15 days embryo head cDNA, RIKEN full-length enriched library, clone:
D930002E06 product:hypothetical protein, full insert sequence" ---
1446091_at 1.10300 "Acidic nuclear phosphoprotein 32 family, member B" Anp32b
1446092_at 0.88652 Transcribed locus ---
1446093_at 0.92600 --- ---
1446094_at 2.23695 AT motif binding factor 1 Atbf1
1446095_at 1.12991 antisense Igf2r RNA Air
1446096_at 1.17189 RIKEN cDNA 2310001H17 gene 2310001H17Rik
1446097_at 0.97898 Retinoblastoma binding protein 8 Rbbp8
1446098_at 1.07331 Zinc finger protein 407 Zfp407
1446099_at 1.11929 Methylenetetrahydrofolate dehydrogenase (NADP+ dependent) 1-like
Mthfd11
1446100_at 0.98101 Transformation related protein 53 binding protein 1 Trp53bp1
1446101_at 1.15416 --- ---
1446102_at 1.21957 "DNA segment, Chr 9, ERATO Doi 292, expressed" D9Ertd292e
1446103_at 1.09199 "RAB8B, member RAS oncogene family" Rab8b
1446104_at 1.00566 SLIT-ROBO Rho GTPase activating protein 3 Srgap3
1446105_at 1.02722 exocyst complex component 1 Exoc1
1446106_at 0.99716 RIKEN cDNA 1700109G14 gene 1700109G14Rik
1446107_at 1.09627 "Serine/threonine kinase 24 (STE20 homolog, yeast)" Stk24
1446108_at 0.97408 "Ankyrin 2, brain" Ank2
1446109_at 1.00454 WD repeat and FYVE domain containing 3 Wdfy3
1446110_at 0.98113 "DNA segment, Chr 3, ERATO Doi 711, expressed" D3Ertd711e
1446111_at 1.01355 "DNA segment, Chr 14, ERATO Doi 611, expressed" D14Ertd611e
1446112_at 1.07846 Ubiquitin specific peptidase 34 Usp34
1446113_at 0.98927 --- ---
1446114_at 1.02751 Inositol polyphosphate-5-phosphatase A Inpp5a

1446115_at 0.97074 --- ---
1446116_at 0.99998 RIKEN cDNA B230378P21 gene B230378P21Rik
1446117_at 1.13973 spire homolog 1 (Drosophila) Spire1
1446118_at 1.30223 "DNA segment, Chr 17, ERATO Doi 165, expressed" D17Ertd165e
1446119_at 1.01491 "Adult male hypothalamus cDNA, RIKEN full-length enriched library, clone:
A230059N19 product:unclassifiable, full insert sequence" ---
1446120_at 1.01174 U2AF homology motif (UHM) kinase 1 Uhmk1
1446121_at 1.00297 Dorso-medial telencephalon gene 2 Dmt2
1446122_at 0.85021 RIKEN cDNA 4732416N19 gene 4732416N19Rik
1446123_at 1.01391 RIKEN cDNA B230396O12 gene B230396O12Rik
1446124_at 1.03774 --- ---
1446125_at 0.92646 --- ---
1446126_at 0.89881 Transcribed locus ---
1446127_at 1.53481 Zinc finger homeobox 1a Zfhx1a
1446128_at 0.93764 Early B-cell factor 2 Ebf2
1446129_at 1.24644 "RAB5A, member RAS oncogene family" Rab5a
1446130_at 0.92727 PCTAIRE-motif protein kinase 2 Pctk2
1446131_at 1.08490 --- ---
1446132_at 1.01368 --- ---
1446133_at 1.19005 "DNA segment, Chr 2, ERATO Doi 295, expressed" D2Ertd295e
1446134_at 0.94881 expressed sequence C78115 C78115
1446135_at 0.98385 RIKEN cDNA D030051J21 gene D030051J21Rik
1446136_at 0.92019 --- ---
1446137_at 0.93843 Glioblastoma amplified sequence Gbas
1446138_at 1.00638 Kinesin-associated protein 3 Kifap3
1446139_at 1.09484 Expressed sequence AI661453 AI661453
1446140_at 1.95936 Pericentriolar material 1 Pcm1
1446141_at 1.16327 "Transforming growth factor, beta 2" Tgfb2
1446142_at 0.92912 --- ---
1446143_at 1.26399 MYST histone acetyltransferase (monocytic leukemia) 3 Myst3
1446144_at 0.88271 peroxin 2 Pex2
1446145_at 0.95290 "eukaryotic translation initiation factor 4 gamma, 3" Eif4g3
1446146_at 1.57646 Gap junction membrane channel protein alpha 7 Gja7
1446147_at 2.47632 expressed sequence C79248 C79248
1446148_x_at 2.47955 expressed sequence C79248 C79248
1446149_at 1.08436 "protein phosphatase 3, catalytic subunit, beta isoform" Ppp3cb
1446150_at 0.89989 "16 days embryo head cDNA, RIKEN full-length enriched library, clone:
C130070I23 product:unclassifiable, full insert sequence" ---
1446151_at 0.91818 --- ---
1446152_at 1.20381 Longevity assurance homolog 6 (S. cerevisiae) Lass6
1446153_at 0.94967 Zinc finger protein 608 Zfp608
1446154_at 0.94975 "16 days neonate cerebellum cDNA, RIKEN full-length enriched library,
clone:9630050E20 product:unclassifiable, full insert sequence" ---

1446155_at 0.88149 RIKEN cDNA 2700089E24 gene 2700089E24Rik
1446156_at 0.89275 "Dystrophin, muscular dystrophy" Dmd
1446157_at 1.04347 RIKEN cDNA A330019N05 gene A330019N05Rik
1446158_at 1.25254 SEC15-like 2 (*S. cerevisiae*) Sec15l2
1446159_at 0.90170 p21 (CDKN1A)-activated kinase 7 Pak7
1446160_x_at 1.06336 transmembrane protein 69 Tmem69
1446161_at 0.96276 "TAF11 RNA polymerase II, TATA box binding protein (TBP)-associated factor" Taf11
1446162_at 1.01463 "poly A binding protein, cytoplasmic 5" Pabpc5
1446163_at 1.05303 --- ---
1446164_at 1.24799 Nuclear receptor coactivator 6 interacting protein Ncoa6ip
1446165_at 0.92574 similar to fibrillarin LOC621312
1446166_at 1.08941 --- ---
1446167_at 1.89040 Homeodomain interacting protein kinase 2 Hipk2
1446168_at 1.10477 --- ---
1446169_at 1.12593 DiGeorge syndrome critical region gene 8 Dgcr8
1446170_at 0.80269 "ATP-binding cassette, sub-family C (CFTR/MRP), member 12" Abcc12
1446171_at 1.02197 RIKEN cDNA A230107N01 gene A230107N01Rik
1446172_at 1.43286 "Adult male urinary bladder cDNA, RIKEN full-length enriched library, clone:9530082M13 product:unclassifiable, full insert sequence" ---
1446173_at 0.97699 --- ---
1446174_at 0.97289 "similar to melanoma antigen family A, 10" LOC245440
1446175_at 0.81611 RIKEN cDNA 1700095A21 gene 1700095A21Rik
1446176_at 1.02991 Myeloid ecotropic viral integration site-related gene 1 Mrg1
1446177_at 0.95657 --- ---
1446178_at 2.14809 Bicaudal C homolog 1 (*Drosophila*) Bicc1
1446179_at 0.91765 RIKEN cDNA C030014K22 gene C030014K22Rik
1446180_at 0.92255 laminin B1 subunit 1 Lamb1-1
1446181_at 0.99839 expressed sequence C85699 C85699
1446182_at 1.53735 "Zinc finger, MYND domain containing 11" Zmynd11
1446183_at 1.20523 Amyotrophic lateral sclerosis 2 (juvenile) homolog (human) Als2
1446184_at 1.03908 --- ---
1446185_at 0.97344 FK506 binding protein 12-rapamycin associated protein 1 Frap1
1446186_at 1.11924 ubiquitin specific peptidase 15 Usp15
1446187_at 1.08279 RIKEN cDNA 4831416G18 gene 4831416G18Rik
1446188_at 1.07359 leucine-rich repeats and IQ motif containing 2 Lrriq2
1446189_at 0.91123 MYST histone acetyltransferase monocytic leukemia 4 Myst4
1446190_at 1.65303 doublecortin and calcium/calmodulin-dependent protein kinase-like 1 Dcamk1l
1446191_at 1.06833 "ABI gene family, member 3" Abi3
1446192_at 0.93522 Amphiphysin Amph
1446193_at 1.61704 --- ---
1446194_at 1.19910 "nuclear receptor subfamily 2, group C, member 1" Nr2c1

1446195_at 0.99218 similar to WD repeat and SOCS box containing protein 2 (WSB-2) (SOCS box-containing WD protein SWiP-2) LOC632689
 1446196_at 1.17798 high mobility group AT-hook 2 Hmga2
 1446197_at 0.93853 Latrophilin 3 Lphn3
 1446198_at 0.82818 "Mesoderm induction early response 1, family member 3" Mier3
 1446199_at 0.98263 regulator of G-protein signaling 20 Rgs20
 1446200_at 1.04797 Deleted in colorectal carcinoma Dcc
 1446201_at 0.94477 RIKEN cDNA 9030420J04 gene 9030420J04Rik
 1446202_at 1.04976 Transcribed locus ---
 1446203_at 0.99099 "muskelin 1, intracellular mediator containing kelch motifs" Mkl1
 1446204_at 1.23542 RIKEN cDNA 8030451K01 gene 8030451K01Rik
 1446205_at 1.00916 "0 day neonate thymus cDNA, RIKEN full-length enriched library, clone: A430057L20 product:unclassifiable, full insert sequence" ---
 1446206_at 1.05966 RIKEN cDNA B230209C24 gene B230209C24Rik
 1446207_at 1.02517 "0 day neonate cerebellum cDNA, RIKEN full-length enriched library, clone: C230011O19 product:unclassifiable, full insert sequence" ---
 1446208_at 1.05698 Tudor domain containing 5 Tdrd5
 1446209_at 1.28840 ubiquitin specific peptidase 40 Usp40
 1446210_at 1.22491 --- ---
 1446211_at 1.00726 V-erb-a erythroblastic leukemia viral oncogene homolog 4 (avian) ErbB4
 1446212_at 0.83786 Dendritic cell protein GA17 Ga17
 1446213_at 1.03607 Myotubularin related protein 2 Mtmr2
 1446214_at 0.95575 RIKEN cDNA D430018E03 gene D430018E03Rik
 1446215_at 0.93684 "zinc finger homeobox 2, antisense" Zfx2as
 1446216_at 0.91289 "Adult male cecum cDNA, RIKEN full-length enriched library, clone:9130202G13 product:unclassifiable, full insert sequence" ---
 1446217_at 1.09101 hypothetical gene supported by AK036329 LOC433202
 1446218_at 0.97227 "protein tyrosine phosphatase, receptor type, f polypeptide (PTPRF), interacting protein (liprin), alpha 2" Ppfia2
 1446219_at 1.03934 RIKEN cDNA D930015E06 gene D930015E06Rik
 1446220_at 1.03593 "gene model 484, (NCBI)" Gm484
 1446221_at 0.97132 Elastin Eln
 1446222_at 1.23237 --- ---
 1446223_at 0.99698 "COX10 homolog, cytochrome c oxidase assembly protein, heme A: farnesyltransferase (yeast)" Cox10
 1446224_at 1.29253 HECT domain containing 2 Hectd2
 1446225_at 0.98367 Ssu72 RNA polymerase II CTD phosphatase homolog (yeast) Ssu72
 1446226_at 1.09392 "Fanconi anemia, complementation group B" Fancb
 1446227_at 0.95776 G protein-coupled receptor 180 Gpr180
 1446228_at 0.97821 "DNA segment, Chr 19, Wayne State University 12, expressed" D19Wsu12e
 1446229_at 1.08174 "16 days embryo head cDNA, RIKEN full-length enriched library, clone: C130045L17 product:unclassifiable, full insert sequence" ---
 1446230_at 0.97398 "Gene model 944, (NCBI)" Gm944

1446231_at	1.00022	Transcribed locus	---	
1446232_at	0.99892	Latent transforming growth factor beta binding protein 1		Ltbp1
1446233_at	0.90750	"Adenosine deaminase, RNA-specific, B2"		Adarb2
1446234_at	3.56919	"Ubiquitously transcribed tetratricopeptide repeat gene, X chromosome"		Utx
1446235_at	1.94270	cDNA sequence BC002199	BC002199	
1446236_at	1.04053	RIKEN cDNA C230072F16 gene	C230072F16Rik	
1446237_at	0.74842	A kinase (PRKA) anchor protein (yotiao) 9		Akap9
1446238_at	0.97006	RIKEN cDNA 4930431P03 gene	4930431P03Rik	
1446239_at	1.12732	RIKEN cDNA 4930444F02 gene	4930444F02Rik	
1446240_at	1.20593	Kinesin 2		Kns2
1446241_at	0.94813	"Adult male thymus cDNA, RIKEN full-length enriched library, clone:5830471B11 product:unclassifiable, full insert sequence"	---	
1446242_at	1.10434	RIKEN cDNA 4930535B03 gene	4930535B03Rik	
1446243_at	1.02882	RIKEN cDNA 5830404H04 gene	5830404H04Rik	
1446244_at	1.04167	RIKEN cDNA 2810482G21 gene	2810482G21Rik	
1446245_at	2.82500	Formin-like 2		Fmnl2
1446246_at	0.89197	"gene model 438, (NCBI)"		Gm438
1446247_at	0.95343	"a disintegrin-like and metallopeptidase (reprolysin type) with thrombospondin type 1 motif, 18"		Adamts18
1446248_at	0.97378	"solute carrier family 18 (vesicular monoamine), member 1"		Slc18a1
1446249_at	1.08916	Anthrax toxin receptor 2		Antxr2
1446250_at	0.93991	peroxisome proliferator activated receptor binding protein		Pparbp
1446251_at	0.93276	hypothetical protein 4930565A17	4930565A17	
1446252_at	0.95958	"Adult retina cDNA, RIKEN full-length enriched library, clone:A930029I05 product:unclassifiable, full insert sequence"	---	
1446253_at	1.06923	"Adult male medulla oblongata cDNA, RIKEN full-length enriched library, clone:6330445G12 product:unclassifiable, full insert sequence"	---	
1446254_at	1.02980	DEP domain containing 6		Depdc6
1446255_at	1.04424	"Potassium voltage-gated channel, subfamily Q, member 1"		Kcnq1
1446256_at	0.96905	---	---	
1446257_at	1.08842	heparan sulfate 6-O-sulfotransferase 2		Hs6st2
1446258_at	1.17720	"Syntrophin, basic 1"		Sntb1
1446259_at	1.18446	---	---	
1446260_at	1.01814	Six6 opposite strand transcript 1		Six6os1
1446261_at	0.90100	"DNA segment, Chr 1, ERATO Doi 507, expressed"		D1Ertd507e
1446262_at	1.20540	"Acyl-Coenzyme A dehydrogenase family, member 10"		Acad10
1446263_at	0.98470	expressed sequence AU015619	AU015619	
1446264_at	1.04725	---	---	
1446265_at	1.02755	dynamamin 3		Dnm3
1446266_at	1.02737	cathepsin 8		Cts8
1446267_at	0.96347	"DNA segment, Chr 17, ERATO Doi 657, expressed"		D17Ertd657e
1446268_at	0.84710	"Pleckstrin homology, Sec7 and coiled-coil domains 3"		Pscd3
1446269_at	1.05224	high mobility group box transcription factor 1		Hbp1

1446270_at	0.96234	RIKEN cDNA E030042O20 gene	E030042O20Rik
1446271_at	1.03722	---	---
1446272_at	1.13633	PCTAIRE-motif protein kinase 2	Pctk2
1446273_at	0.97664	CUB and Sushi multiple domains 1	Csmd1
1446274_at	0.86427	"solute carrier family 16 (monocarboxylic acid transporters), member 1"	Slc16a1
1446275_at	0.83272	Trinucleotide repeat containing 15	Tnrc15
1446276_at	1.45885	RIKEN cDNA 2310008H04 gene	2310008H04Rik
1446277_at	0.87107	---	---
1446278_at	1.95483	"Ubiquitously transcribed tetratricopeptide repeat gene, X chromosome"	Utx
1446279_at	0.92733	Neuronal growth regulator 1	Negr1
1446280_at	1.30293	Forkhead box P1	Foxp1
1446281_at	0.94107	RIKEN cDNA C530008M07 gene	C530008M07Rik
1446282_at	1.08101	hypothetical protein LOC626082	LOC626082
1446283_at	0.84103	Dedicator of cytokinesis 4	Dock4
1446284_at	2.45921	metastasis suppressor 1	Mtss1
1446285_at	0.95276	RIKEN cDNA A130034M23 gene	A130034M23Rik
1446286_at	1.03216	"16 days neonate cerebellum cDNA, RIKEN full-length enriched library, clone:9630014E16 product:unclassifiable, full insert sequence"	---
1446287_at	1.01238	RIKEN cDNA 4931440B09 gene	4931440B09Rik
1446288_at	1.32127	expressed sequence C78692	C78692
1446289_at	1.05749	"Serine/threonine kinase 3 (Ste20, yeast homolog)"	Stk3
1446290_at	1.15983	RIKEN cDNA 2410129H14 gene	2410129H14Rik
1446291_at	0.82593	RIKEN cDNA 9330175H22 gene	9330175H22Rik
1446292_at	1.06880	Spermatogenesis associated 5	Spata5
1446293_at	0.89791	B-cell CLL/lymphoma 11A (zinc finger protein)	Bcl11a
1446294_at	1.13105	Transcribed locus	---
1446295_at	1.14035	tripartite motif protein 24	Trim24
1446296_at	0.88315	RIKEN cDNA 4631422O05 gene	4631422O05Rik
1446297_at	0.99695	3-hydroxyisobutyrate dehydrogenase	Hibadh
1446298_at	1.01337	RIKEN cDNA C630016I17 gene	C630016I17Rik
1446299_at	1.34240	RIKEN cDNA 6720456B07 gene	6720456B07Rik
1446300_at	0.95756	"Adult male epididymis cDNA, RIKEN full-length enriched library, clone:9230007L17 product:unclassifiable, full insert sequence"	---
1446301_at	1.00703	"Phenylalanine-tRNA synthetase-like, alpha subunit"	Farsla
1446302_at	0.90597	Cysteine conjugate-beta lyase 1	Ccbl1
1446303_at	1.64229	insulin-like growth factor I receptor	Igf1r
1446304_at	1.04485	---	---
1446305_at	0.92207	"Adult male corpora quadrigemina cDNA, RIKEN full-length enriched library, clone:B230216H12 product:unclassifiable, full insert sequence"	---
1446306_at	0.89090	---	---
1446307_at	1.07252	Poly(A) polymerase gamma	Papolg
1446308_at	0.98476	RIKEN cDNA 1700106J16 gene	1700106J16Rik
1446309_at	1.02235	RIKEN cDNA 5830483C08 gene	5830483C08Rik

1446310_at	0.92080	RIKEN cDNA C130053K05 gene	C130053K05Rik
1446311_at	0.87380	membrane bound C2 domain containing protein	Mbc2
1446312_at	1.10779	---	---
1446313_at	0.96625	zinc finger protein 30	Zfp30
1446314_at	1.13822	RIKEN cDNA A330041C17 gene	A330041C17Rik
1446315_at	0.94485	ATP citrate lyase	Acly
1446316_at	1.00900	Lipin 2	Lpin2
1446317_at	0.97501	AT rich interactive domain 4B (Rbp1 like)	Arid4b
1446318_at	1.78541	"Solute carrier family 44, member 1"	Slc44a1
1446319_at	1.14389	ankyrin repeat and SOCS box-containing protein 7	Asb7
1446320_at	1.02506	RIKEN cDNA 2310010I16 gene	2310010I16Rik
1446321_at	1.04953	RIKEN cDNA B230208B08 gene	B230208B08Rik
1446322_at	1.07150	expressed sequence C78568	C78568
1446323_at	1.05542	---	---
1446324_at	0.98798	"Calcium channel, voltage-dependent, alpha2/delta subunit 1"	Cacna2d1
1446325_at	0.88032	Prenylcysteine oxidase 1	Pcyox1
1446326_at	1.01713	"procollagen, type I, alpha 2"	Colla2
1446327_at	0.87328	---	---
1446328_at	1.08041	Transcribed locus	---
1446329_at	1.20704	---	---
1446330_at	1.28803	---	---
1446331_at	0.93564	prostaglandin F receptor	Ptgfr
1446332_at	0.96146	"Protocadherin gamma subfamily C, 3"	Pcdhgc3
1446333_at	1.09830	---	---
1446334_at	1.01146	---	---
1446335_at	1.15267	CDNA sequence BC013481	BC013481
1446336_at	1.15674	---	---
1446337_at	0.98441	Huntington disease gene homolog	Hdh
1446338_at	1.06116	---	---
1446339_at	0.95417	PDZ and LIM domain 5	Pdlim5
1446340_at	1.01874	Androgen receptor	Ar
1446341_at	0.87078	RIKEN cDNA 6030403N03 gene	6030403N03Rik
1446342_at	1.00550	RIKEN cDNA 2310001H17 gene	2310001H17Rik
1446343_at	0.92927	Transcribed locus	---
1446344_at	0.87245	nyctalopin	Nyx
1446345_at	1.73107	---	---
1446346_at	1.22068	---	---
1446347_at	1.01348	RIKEN cDNA 4631405J19 gene	4631405J19Rik
1446348_at	1.03866	Mitogen-activated protein kinase kinase kinase 5	Map4k5
1446349_at	0.92370	zinc finger protein 78	Zfp78
1446350_at	0.89002	Integrin alpha 4	Itga4
1446351_at	1.09821	hypothetical protein D430034N21	D430034N21
1446352_at	0.87743	RIKEN cDNA 2410016O06 gene	2410016O06Rik

1446353_at	0.94945	"Tubulin, beta 6"	Tubb6
1446354_at	0.91907	RIKEN cDNA C130098B18 gene	C130098B18Rik
1446355_at	1.06144	RIKEN cDNA 9530085L11 gene	9530085L11Rik
1446356_at	1.03444	"Protein phosphatase 2, regulatory subunit B (B56), epsilon isoform"	Ppp2r5e
1446357_at	1.04785	cDNA sequence BC020402	BC020402
1446358_at	0.99347	Brain-specific angiogenesis inhibitor 3	Bai3
1446359_at	0.94833	"guanine nucleotide binding protein, alpha stimulating, olfactory type"	Gnal
1446360_at	0.99423	RAR-related orphan receptor alpha	Rora
1446361_at	0.90943	G protein-coupled receptor kinase 5	Gprk5
1446362_at	0.97354	---	---
1446363_at	0.99367	---	---
1446364_at	1.07690	WW domain containing E3 ubiquitin protein ligase 1	Wwp1
1446365_at	2.20958	vesicle transport through interaction with t-SNAREs homolog 1A (yeast)	Vti1a
1446366_at	0.92098	"transient receptor potential cation channel, subfamily C, member 4"	Trpc4
1446367_at	1.09074	"0 day neonate cerebellum cDNA, RIKEN full-length enriched library, clone: C230077G16 product:unclassifiable, full insert sequence"	---
1446368_at	1.03662	RIKEN cDNA 9130221J18 gene	9130221J18Rik
1446369_at	0.89624	armadillo repeat containing 1	Armc1
1446370_at	0.97437	RAB11 family interacting protein 4 (class II)	Rab11fip4
1446371_at	1.07013	---	---
1446372_at	1.17332	"CD8 antigen, beta chain 1"	Cd8b1
1446373_at	0.90265	LPS-responsive beige-like anchor	Lrba
1446374_at	1.17311	Rho guanine nucleotide exchange factor (GEF) 10	Arhgef10
1446375_at	0.93665	Exocyst complex component 6	Exoc6
1446376_at	0.96007	"ribosomal protein S6 kinase, polypeptide 1"	Rps6kb1
1446377_at	0.97322	fibulin 5	Fbln5
1446378_at	1.01580	"DNA segment, Chr 2, ERATO Doi 357, expressed"	D2Ertd357e
1446379_at	1.06742	expressed sequence C79296	C79296
1446380_at	1.02446	RIKEN cDNA 9430076C15 gene	9430076C15Rik
1446381_at	1.01107	CDNA sequence BC030499	BC030499
1446382_at	1.05414	"0 day neonate cerebellum cDNA, RIKEN full-length enriched library, clone: C230094E06 product:unclassifiable, full insert sequence"	---
1446383_at	2.34244	Ephrin A5	Efna5
1446384_at	0.94472	RIKEN cDNA B230339M05 gene	B230339M05Rik
1446385_at	0.97585	RIKEN cDNA 9430083A17 gene	9430083A17Rik
1446386_at	0.95725	Transcription factor 4	Tcf4
1446387_at	1.11137	---	---
1446388_at	0.96309	Testis expressed gene 9	Tex9
1446389_at	1.00057	Nuclear receptor interacting protein 1	Nrip1
1446390_at	0.97601	"ELK1, member of ETS oncogene family"	Elk1
1446391_at	0.98904	"Synuclein, alpha"	Snca
1446392_at	0.85311	Slit homolog 2 (Drosophila)	Slit2
1446393_at	1.07594	"DNA segment, Chr 5, ERATO Doi 159, expressed"	D5Ertd159e

1446394_at 1.05988 "opioid receptor, kappa 1" Oprk1
1446395_at 1.03022 Roundabout homolog 2 (Drosophila) Robo2
1446396_at 0.93279 Angiopoietin-like 1 Angptl1
1446397_at 1.82175 Nucleoporin 98 Nup98
1446398_at 0.81324 hypothetical gene supported by AK028463 LOC433197
1446399_at 0.97206 cadherin 10 Cdh10
1446400_at 1.10649 olfactory receptor 558 Olfr558
1446401_at 1.04416 Attractin like 1 Atrnl1
1446402_at 0.90006 --- ---
1446403_at 1.10248 expressed sequence AU022899 AU022899
1446404_at 1.04691 --- ---
1446405_at 1.15941 MYST histone acetyltransferase (monocytic leukemia) 3 Myst3
1446406_at 1.00030 Progesterone and adipoQ receptor family member VIII Paqr8
1446407_at 1.19936 "Myb-like, SWIRM and MPN domains 1" Mysm1
1446408_at 1.01452 Homeo box A10 Hoxa10
1446409_at 1.08425 glycosyltransferase-like domain containing 1 Gtdc1
1446410_at 1.16406 Choline/ethanolaminephosphotransferase 1 Cept1
1446411_at 0.86546 RIKEN cDNA 9630020C08 gene 9630020C08Rik
1446412_at 2.23365 WW domain-containing oxidoreductase Wwox
1446413_at 0.85640 RIKEN cDNA 4930555G07 gene 4930555G07Rik
1446414_at 1.13990 RIKEN cDNA 0910001A06 gene 0910001A06Rik
1446415_at 0.83789 heme binding protein 2 Hebp2
1446416_at 1.00873 sulfatase 1 Sulfl
1446417_at 1.62356 "Serine (or cysteine) peptidase inhibitor, clade B, member 6a" Serpinb6a
1446418_at 1.00107 Autophagy-related 7 (yeast) Atg7
1446419_at 1.09672 --- ---
1446420_at 0.91232 Plexin A2 Plxna2
1446421_at 2.43550 Schwannomin interacting protein 1 Schip1
1446422_at 1.02627 Membrane-associated ring finger (C3HC4) 3 3-Mar
1446423_at 1.17830 RAD51-like 1 (S. cerevisiae) Rad511
1446424_at 0.90000 "dynein, axonemal, heavy chain 12" Dnahc12
1446425_at 1.03252 RIKEN cDNA 4732418C07 gene 4732418C07Rik
1446426_at 2.47736 Enabled homolog (Drosophila) Enah
1446427_at 1.31986 RIKEN cDNA B230118G17 gene B230118G17Rik
1446428_at 1.17249 RIKEN cDNA D530005L17 gene D530005L17Rik
1446429_at 0.93275 "purinergic receptor P2X, ligand-gated ion channel 4" P2rx4
1446430_at 0.99980 HECT domain containing 2 Hectd2
1446431_at 0.93656 Dynamin 3 Dnm3
1446432_at 1.02085 RIKEN cDNA B530045E10 gene B530045E10Rik
1446433_at 1.18216 acyl-Coenzyme A binding domain containing 5 Acbd5
1446434_at 1.26270 Chondroitin sulfate GalNAcT-2 Galnact2
1446435_at 1.05680 "0 day neonate eyeball cDNA, RIKEN full-length enriched library, clone:
E130016A22 product:hypothetical protein, full insert sequence" ---

1446436_at 0.97254 Recombining binding protein suppressor of hairless (Drosophila) Rbpsuh
1446437_at 1.00555 RIKEN cDNA 9630028H03 gene 9630028H03Rik
1446438_at 0.96895 ubiquitin protein ligase E3 component n-recognin 1 Ubr1
1446439_at 1.08384 RNA polymerase 1-4 Rpo1-4
1446440_at 0.96431 CDNA clone IMAGE:40051559 ---
1446441_at 0.90809 "solute carrier family 20, member 1" Slc20a1
1446442_at 0.85031 RNA (guanine-7-) methyltransferase Rnmt
1446443_at 1.17303 RIKEN cDNA B230213L16 gene B230213L16Rik
1446444_at 1.56524 membrane associated DNA binding protein Mnab
1446445_at 0.93493 "12 days embryo spinal ganglion cDNA, RIKEN full-length enriched library,
clone:D130018F08 product:unclassifiable, full insert sequence" ---
1446446_at 1.02977 RAN binding protein 10 Ranbp10
1446447_at 1.07324 RIKEN cDNA 4921537D05 gene 4921537D05Rik
1446448_at 1.16970 protein inhibitor of activated STAT 1 Pias1
1446449_at 0.91707 kelch-like 21 (Drosophila) Khl21
1446450_at 1.06789 Titin Ttn
1446451_at 1.17112 --- ---
1446452_at 2.00271 "Transcription factor 7-like 2, T-cell specific, HMG-box" Tcf7l2
1446453_at 1.13328 --- ---
1446454_at 0.95706 --- ---
1446455_at 0.89410 Transcribed locus ---
1446456_at 1.02297 "0 day neonate eyeball cDNA, RIKEN full-length enriched library, clone:
E130003J08 product:unclassifiable, full insert sequence" ---
1446457_at 1.06006 DEAD (Asp-Glu-Ala-Asp) box polypeptide 58 Ddx58
1446458_at 0.84844 "ribonuclease L (2', 5'-oligoadenylate synthetase-dependent)" Rnasel
1446459_at 0.95224 --- ---
1446460_at 0.86283 transmembrane protein 61 Tmem61
1446461_at 1.20390 SRY-box containing gene 5 Sox5
1446462_at 0.91880 Histidyl tRNA synthetase 2 Hars2
1446463_at 0.87366 CDNA sequence BC024969 BC024969
1446464_at 1.09687 "Proteasome (prosome, macropain) activator subunit 4" Psme4
1446465_at 0.94397 "ATPase, H+ transporting, lysosomal V1 subunit B2" Atp6v1b2
1446466_at 0.94844 "Mannose receptor, C type 2" Mrc2
1446467_at 0.89447 "ATPas, class II, type 9B" Atp9b
1446468_at 0.95588 CUB and Sushi multiple domains 1 /// similar to CUB and Sushi multiple
domains 1 Csm1 /// LOC670210
1446469_at 0.99776 hypothetical protein A030008J09 A030008J09
1446470_at 0.88108 --- ---
1446471_at 0.84101 --- ---
1446472_at 1.55949 AT hook containing transcription factor 1 Ahctf1
1446473_at 1.07873 --- ---
1446474_at 1.36251 --- ---
1446475_at 1.43065 Muscleblind-like 2 Mbnl2

1446476_at 0.88596 RIKEN cDNA A430107P09 gene A430107P09Rik
1446477_at 1.09053 zinc finger protein 622 Zfp622
1446478_at 0.97124 RIKEN cDNA D130067I03 gene D130067I03Rik
1446479_at 1.07321 TNNI3 interacting kinase Tnni3k
1446480_at 0.95969 "DNA segment, Chr 8, ERATO Doi 82, expressed" D8Ertd82e
1446481_at 2.65941 "Amyloid beta (A4) precursor protein-binding, family B, member 2" Apbb2
1446482_at 1.00241 centrosomal protein 63 Cep63
1446483_at 1.13968 "aldo-keto reductase family 1, member E1" Akr1e1
1446484_at 0.91384 Myocyte enhancer factor 2C Mef2c
1446485_at 0.94680 RIKEN cDNA 4930553C11 gene 4930553C11Rik
1446486_at 1.22806 DTW domain containing 2 Dtw2
1446487_at 0.91514 --- ---
1446488_at 0.87262 Poliovirus receptor-related 1 Pvr1
1446489_at 1.35811 Suppressor of fused homolog (Drosophila) Sufu
1446490_at 1.10399 Polypyrimidine tract binding protein 2 Ptbp2
1446491_at 0.96245 "Guanine nucleotide binding protein, alpha stimulating, olfactory type" Gnal
1446492_at 1.08844 transmembrane protein 132D Tmem132d
1446493_at 0.92756 CDC14 cell division cycle 14 homolog A (S. cerevisiae) Cdc14a
1446494_at 1.12091 "Non-coding RNA Kis2-T1 mRNA, complete sequence, alternatively spliced"

1446495_at 1.08439 --- ---
1446496_at 1.09665 "splicing factor, arginine/serine-rich 10 (transformer 2 homolog, Drosophila)"
Sfrs10
1446497_at 1.23771 Neuron navigator 1 Nav1
1446498_at 0.83324 "interleukin 20 receptor, alpha" Il20ra
1446499_at 1.05931 Fas (TNFRSF6)-associated via death domain Fadd
1446500_at 1.09492 "LysM, putative peptidoglycan-binding, domain containing 4" Lysmd4
1446501_at 0.97171 RIKEN cDNA A830053O21 gene A830053O21Rik
1446502_at 1.10020 --- ---
1446503_at 0.88364 --- ---
1446504_at 1.11720 --- ---
1446505_at 1.02965 CD84 antigen Cd84
1446506_at 1.14080 hypothetical protein B930041G04 B930041G04
1446507_at 1.32920 Prion protein Prnp
1446508_at 1.85667 --- ---
1446509_at 0.97705 spermine oxidase Smox
1446510_at 1.71733 Pleckstrin homology domain interacting protein Phip
1446511_at 1.34069 mutS homolog 3 (E. coli) Msh3
1446512_at 1.30428 RIKEN cDNA 2610312B22 gene 2610312B22Rik
1446513_at 0.95953 "A disintegrin-like and metallopeptidase (reprolysin type) with thrombospondin
type 1 motif, 19" Adamts19
1446514_at 0.94531 Dipeptidylpeptidase 10 Dpp10
1446515_at 0.94334 Janus kinase 3 Jak3

1446516_at 0.90561 B-cell CLL/lymphoma 7C Bcl7c
1446517_at 1.03912 Carbonic anhydrase 12 Car12
1446518_at 1.04200 Rho-associated coiled-coil forming kinase 1 Rock1
1446519_at 1.02321 RIKEN cDNA A730091E23 gene A730091E23Rik
1446520_at 0.85826 interleukin 1 receptor accessory protein-like 2 Il1rapl2
1446521_at 2.28892 "Proteasome (prosome, macropain) 26S subunit, non-ATPase, 14" Psmd14
1446522_at 0.99765 --- ---
1446523_at 1.05504 --- ---
1446524_at 1.17476 "PREDICTED: Mus musculus similar to HYPOTHETICAL PROTEIN ORF-1137 (LOC626631), mRNA" ---
1446525_at 1.03180 Glypican 3 Gpc3
1446526_at 1.00135 RIKEN cDNA 9330199F22 gene 9330199F22Rik
1446527_at 0.92825 "angiotensin II receptor, type 1b" Agtr1b
1446528_at 1.00331 SCY1-like 2 (S. cerevisiae) Scyl2
1446529_at 1.60295 RIKEN cDNA 1200003I07 gene 1200003I07Rik
1446530_at 1.10895 --- ---
1446531_at 0.96605 --- ---
1446532_at 1.02416 --- ---
1446533_at 0.74024 CREB regulated transcription coactivator 1 Crtc1
1446534_at 0.95429 --- ---
1446535_at 1.02628 Collectin sub-family member 12 Colec12
1446536_at 0.99100 "sema domain, transmembrane domain (TM), and cytoplasmic domain, (semaphorin) 6D" Sema6d
1446537_at 1.09371 Dendritic cell protein GA17 Ga17
1446538_at 1.11120 Karyopherin (importin) alpha 3 Kpna3
1446539_at 0.84489 "Inositol polyphosphate-4-phosphatase, type II" Inpp4b
1446540_at 1.10357 "15 days embryo head cDNA, RIKEN full-length enriched library, clone: D930047B11 product:unclassifiable, full insert sequence" ---
1446541_at 0.90363 RIKEN cDNA 4930434E21 gene 4930434E21Rik
1446542_at 0.93751 acyl-CoA synthetase short-chain family member 2 Acss2
1446543_at 1.27918 GCN1 general control of amino-acid synthesis 1-like 1 (yeast) Gcn111
1446544_at 0.87654 Aryl hydrocarbon receptor nuclear translocator-like Arntl
1446545_at 1.01097 Spectrin beta 2 Spnb2
1446546_at 0.93247 "trimethyllysine hydroxylase, epsilon" Tmlhe
1446547_at 0.84597 RIKEN cDNA 9330175E14 gene 9330175E14Rik
1446548_at 1.11319 Glioma tumor suppressor candidate region gene 1 Gltscr1
1446549_at 0.89609 Kelch repeat and BTB (POZ) domain containing 11 Kbtbd11
1446550_at 1.24924 G1 to S phase transition 1 Gspt1
1446551_at 1.10127 "Mammary gland RCB-0527 Jyg-MC(B) cDNA, RIKEN full-length enriched library, clone:G930025P03 product:unclassifiable, full insert sequence" ---
1446552_at 0.86509 "solute carrier family 12, member 3" Slc12a3
1446553_at 0.94159 Hemopoietic cell kinase Hck
1446554_at 0.94282 glycoprotein 6 (platelet) /// similar to glycoprotein VI (platelet) Gp6 ///

LOC636242

1446555_at 0.86880 "12 days embryo spinal ganglion cDNA, RIKEN full-length enriched library, clone:D130076B20 product:unclassifiable, full insert sequence" ---

1446556_at 1.07591 "potassium voltage-gated channel, Shal-related family, member 1" Kcnd1

1446557_at 0.96045 RIKEN cDNA E130309F12 gene E130309F12Rik

1446558_at 1.17210 RIKEN cDNA F730015K02 gene F730015K02Rik

1446559_at 0.83690 RIKEN cDNA D830016K09 gene D830016K09Rik

1446560_at 0.92690 "protease, serine, 23" Prss23

1446561_at 0.92606 paired box gene 8 Pax8

1446562_at 0.97914 SH3-domain kinase binding protein 1 Sh3kbp1

1446563_at 1.13946 --- ---

1446564_at 0.85489 "DNA segment, Chr 18, ERATO Doi 169, expressed" D18Ertd169e

1446565_at 1.96545 "Mannosidase, alpha, class 1A, member 2" Man1a2

1446566_at 1.59821 "Adaptor-related protein complex 2, beta 1 subunit" Ap2b1

1446567_at 1.09820 Cyclin D3 Ccnd3

1446568_at 0.93842 SREBP cleavage activating protein Scap

1446569_at 0.92227 Integrin alpha 1 Itga1

1446570_at 1.14586 Mastermind like 2 (Drosophila) Maml2

1446571_at 1.92540 RIKEN cDNA A830039H10 gene A830039H10Rik

1446572_at 0.94588 Ubiquitin specific peptidase 53 Usp53

1446573_at 0.87329 EGF-like repeats and discoidin I-like domains 3 Edil3

1446574_at 0.92348 Nuclear receptor co-repressor 1 Ncor1

1446575_at 1.02366 expressed sequence C77583 C77583

1446576_at 0.98817 RIKEN cDNA 4931419K03 gene 4931419K03Rik

1446577_at 1.10465 "Phosphodiesterase 4B, cAMP specific" Pde4b

1446578_at 1.01399 --- ---

1446579_at 0.99715 --- ---

1446580_at 1.08737 RIKEN cDNA 9430083G14 gene 9430083G14Rik

1446581_at 1.00327 Protocadherin 11 X-linked Pcdh11x

1446582_at 1.09721 DNA polymerase N Poln

1446583_at 1.03938 Transcribed locus ---

1446584_at 0.93090 Thioredoxin domain containing 4 (endoplasmic reticulum) Txndc4

1446585_at 0.92277 "Discs, large homolog 2 (Drosophila)" Dlgh2

1446586_at 0.97648 RIKEN cDNA 8030463A06 gene 8030463A06Rik

1446587_at 1.05836 RIKEN cDNA 2310042E16 gene 2310042E16Rik

1446588_at 1.04845 --- ---

1446589_at 0.97106 Rhomboid domain containing 1 Rhbdd1

1446590_at 0.96661 --- ---

1446591_at 1.25605 "gene model 1574, (NCBI)" Gm1574

1446592_at 1.71566 --- ---

1446593_at 1.33842 Triple functional domain (PTPRF interacting) Trio

1446594_at 1.00422 --- ---

1446595_at 1.21970 intersectin 2 Itsn2

1446596_at	1.00566	RIKEN cDNA C130009A20 gene	C130009A20Rik
1446597_at	0.89718	Hypothetical LOC67660	LOC67660
1446598_at	2.26200	"Protein kinase C, alpha"	Prkca
1446599_at	1.00216	---	---
1446600_at	1.05576	---	---
1446601_at	0.96045	---	---
1446602_at	1.03579	FERM domain containing 3	Frmd3
1446603_at	1.00274	---	---
1446604_at	1.13127	---	---
1446605_at	1.33463	SLIT-ROBO Rho GTPase activating protein 3	Srgap3
1446606_at	1.10511	region containing RIKEN cDNA 1700028E10 gene; hypothetical protein	
A630054D14		LOC625175	
1446607_at	1.18445	Microtubule-associated protein 7	Mtap7
1446608_at	1.05350	Casitas B-lineage lymphoma	Cbl
1446609_at	0.76269	---	---
1446610_at	1.01296	"engulfment and cell motility 1, ced-12 homolog (C. elegans)"	Elmo1
1446611_at	1.07932	Arginine-tRNA-protein transferase 1	Ate1
1446612_at	0.92487	RIKEN cDNA 9330118A15 gene	9330118A15Rik
1446613_at	0.89411	RIKEN cDNA 1190005F20 gene	1190005F20Rik
1446614_at	1.07378	---	---
1446615_at	0.99767	---	---
1446616_at	1.00992	Glioma tumor suppressor candidate region gene 1	Gltscr1
1446617_at	0.86104	"ST8 alpha-N-acetyl-neuraminide alpha-2,8-sialyltransferase 4"	St8sia4
1446618_at	1.32833	hypothetical gene supported by AK038224	LOC432971
1446619_at	1.03794	RIKEN cDNA A130038J17 gene	A130038J17Rik
1446620_at	0.97535	Myosin ID	Myo1d
1446621_at	1.16877	Chromodomain helicase DNA binding protein 7	Chd7
1446622_at	1.02927	RIKEN cDNA A330068G13 gene	A330068G13Rik
1446623_at	0.96635	---	---
1446624_at	0.90824	Fukuyama type congenital muscular dystrophy homolog (human)	Fcmd
1446625_at	0.88336	transmembrane protein 100	Tmem100
1446626_at	1.18829	"DNA segment, Chr 16, human D22S680E, expressed"	D16H22S680E
1446627_at	2.00161	"Serine/threonine kinase 3 (Ste20, yeast homolog)"	Stk3
1446628_at	1.04474	zinc finger and BTB domain containing 24	Zbtb24
1446629_at	1.06523	expressed sequence C85938	C85938
1446630_at	1.17188	---	---
1446631_at	0.91758	hypothetical protein 9430083I22	9430083I22
1446632_at	1.44432	"Calcium channel, voltage-dependent, beta 2 subunit"	Cacnb2
1446633_at	1.22246	autophagy-related 7 (yeast)	Atg7
1446634_at	1.04297	RIKEN cDNA D830036C21 gene	D830036C21Rik
1446635_at	1.09571	"DNA segment, Chr 13, ERATO Doi 476, expressed"	D13Ertd476e
1446636_at	0.86059	tensin 3	Tns3
1446637_at	0.92904	hypothetical protein LOC666539	LOC666539

1446638_at	0.94599	"protein tyrosine phosphatase, receptor type, U"	Ptpru
1446639_at	1.03705	---	---
1446640_at	0.90991	Transcribed locus	---
1446641_at	1.05755	Synaptotagmin VII	Syt7
1446642_at	1.02808	PHD finger protein 14	Phf14
1446643_at	0.93244	RIKEN cDNA 5330409N07 gene	5330409N07Rik
1446644_at	1.25791	Bromodomain containing 3	Brd3
1446645_at	0.91443	RIKEN cDNA A730037C10 gene	A730037C10Rik
1446646_at	1.00217	"TAF3 RNA polymerase II, TATA box binding protein (TBP)-associated factor" Taf3	
1446647_at	1.09627	---	---
1446648_at	1.15703	Syntaxin binding protein 4	Stxbp4
1446649_at	1.07845	"Centaurin, delta 1"	Centd1
1446650_at	0.99786	---	---
1446651_at	1.01287	---	---
1446652_at	1.13872	RALBP1 associated Eps domain containing protein 2	Reps2
1446653_at	1.03494	---	---
1446654_at	1.11163	---	---
1446655_at	1.06903	SCY1-like 3 (<i>S. cerevisiae</i>)	Scyl3
1446656_at	1.00304	Angiopoietin 1	Angpt1
1446657_at	0.97541	---	---
1446658_at	1.14236	Transcribed locus	---
1446659_at	0.99694	"15 days embryo head cDNA, RIKEN full-length enriched library, clone: D930044F01 product:unclassifiable, full insert sequence"	---
1446660_at	1.10829	Coiled-coil domain containing 46	Ccdc46
1446661_at	1.02780	---	---
1446662_at	1.06760	RIKEN cDNA 4930562C15 gene	4930562C15Rik
1446663_at	0.98631	"Calcium/calmodulin-dependent protein kinase II, delta"	Camk2d
1446664_at	0.97715	"DNA segment, Chr 15, ERATO Doi 154, expressed"	D15Ertd154e
1446665_at	1.06251	expressed sequence C80918	C80918
1446666_at	0.91239	"a disintegrin-like and metallopeptidase (reprolysin type) with thrombospondin type 1 motif, 12"	Adamts12
1446667_at	1.10073	---	---
1446668_at	0.87729	"PREDICTED: Mus musculus similar to sidekick homolog 1 (LOC620969), mRNA"	---
1446669_at	0.99760	Early B-cell factor 1	Ebf1
1446670_at	1.01843	"CUG triplet repeat, RNA binding protein 2"	Cugbp2
1446671_at	0.97416	"Protein phosphatase 1, regulatory (inhibitor) subunit 14c"	Ppp1r14c
1446672_at	1.02521	---	---
1446673_at	0.85369	RIKEN cDNA D330013E07 gene	D330013E07Rik
1446674_at	1.06470	Kinesin 2	Kns2
1446675_at	0.97927	adenosine kinase	Adk
1446676_at	0.90662	hypothetical protein A130033P14	A130033P14

1446677_at	1.03313	RIKEN cDNA A830029E22 gene	A830029E22Rik
1446678_at	0.93662	"DNA segment, Chr 14, ERATO Doi 16, expressed"	D14Ertld16e
1446679_at	1.10331	"Glutamate receptor, metabotropic 5"	Grm5
1446680_at	1.05122	---	---
1446681_at	1.01313	expressed sequence BB086117	BB086117
1446682_at	0.91803	Transcribed locus	---
1446683_at	1.08665	epidermal growth factor receptor pathway substrate 15-like 1	Eps1511
1446684_at	0.95263	GTPase activating RANGAP domain-like 1	Garnl1
1446685_at	0.99527	"Similar to matrilin 1, cartilage matrix protein"	LOC436089
1446686_at	1.04010	"DnaJ (Hsp40) homolog, subfamily C, member 11"	Dnajc11
1446687_at	0.94807	"splicing factor 3a, subunit 1"	Sf3a1
1446688_at	2.05188	"Guanine nucleotide binding protein, alpha q polypeptide"	Gnaq
1446689_at	1.00323	RIKEN cDNA D230007K08 gene	D230007K08Rik
1446690_at	0.96862	---	---
1446691_at	1.16124	"Platelet-derived growth factor, D polypeptide"	Pdgfd
1446692_at	1.08891	Nuclear fragile X mental retardation protein interacting protein 2	Nufip2
1446693_at	1.21427	---	---
1446694_at	0.96510	Dynamin 3	Dnm3
1446695_at	0.94505	5'-3' exoribonuclease 1	Xrn1
1446696_at	0.97585	intraflagellar transport 57 homolog (Chlamydomonas)	Ift57
1446697_at	0.96950	---	---
1446698_at	1.31590	Mortality factor 4 like 1	Morf411
1446699_at	0.86888	jumonji domain containing 2A	Jmjd2a
1446700_at	1.08906	---	---
1446701_at	0.90199	"ATP-binding cassette, sub-family B (MDR/TAP), member 1B"	Abcb1b
1446702_at	0.94391	Kinectin 1	Ktn1
1446703_at	0.99863	Ring finger and WD repeat domain 2	Rfwd2
1446704_at	1.49769	Thiamin pyrophosphokinase	Tpk1
1446705_at	1.00275	---	---
1446706_at	0.89216	expressed sequence C79123	C79123
1446707_at	1.09721	---	---
1446708_at	0.87918	"hypoxia inducible factor 3, alpha subunit"	Hif3a
1446709_at	0.90809	---	---
1446710_at	1.10243	expressed sequence C76669	C76669
1446711_at	1.09240	RIKEN cDNA A130040G06 gene	A130040G06Rik
1446712_at	1.12369	"neurotrophic tyrosine kinase, receptor, type 2"	Ntrk2
1446713_at	1.03124	Tial1 cytotoxic granule-associated RNA binding protein-like 1	Tial1
1446714_x_at	1.02665	Transcribed locus	---
1446715_at	1.02123	"ATP-binding cassette, sub-family A (ABC1), member 13 /// similar to ATP binding cassette, sub-family A (ABC1), member 13"	Abca13 /// LOC672028
1446716_at	1.02255	"Transcribed locus, moderately similar to XP_576460.1 PREDICTED: similar to hypothetical protein PB402898.00.0 [Rattus norvegicus]"	---
1446717_at	1.02709	---	---

1446718_at 1.10719 "nuclear factor of kappa light chain gene enhancer in B-cells inhibitor, beta"
Nfkbib
1446719_at 1.14284 RIKEN cDNA D530031C13 gene D530031C13Rik
1446720_at 0.88538 Activated leukocyte cell adhesion molecule Alcam
1446721_at 0.93533 --- ---
1446722_at 1.52687 "Guanine nucleotide binding protein, alpha 13" Gna13
1446723_at 1.04604 --- ---
1446724_at 1.07499 FCH domain only 2 Fcho2
1446725_at 0.99792 RIKEN cDNA A130094D17 gene A130094D17Rik
1446726_at 0.94220 hypothetical protein LOC665490 /// hypothetical protein LOC669883 ///
hypothetical protein LOC676857 LOC665490 /// LOC669883 /// LOC676857
1446727_at 0.97040 --- ---
1446728_at 0.98001 "DNA segment, Chr 3, ERATO Doi 797, expressed" D3ErtD797e
1446729_at 0.94636 dispatched homolog 2 (Drosophila) Disp2
1446730_at 1.06326 Expressed sequence C79267 C79267
1446731_at 1.05089 hypothetical protein A730016A17 A730016A17
1446732_at 1.09523 --- ---
1446733_at 1.16359 --- ---
1446734_at 0.97986 expressed sequence C77438 C77438
1446735_at 1.06059 intersectin 2 Itsn2
1446736_at 0.94399 Transducin (beta)-like 1 X-linked Tbl1x
1446737_a_at 0.92595 hook homolog 3 (Drosophila) Hook3
1446738_at 0.95324 "DNA segment, Chr 7, Wayne State University 130, expressed" D7Wsu130e
1446739_at 1.45057 "Beta galactoside alpha 2,6 sialyltransferase 1" St6gal1
1446740_at 1.04330 Transcribed locus ---
1446741_at 0.96050 hypothetical gene supported by AK052160 LOC432868
1446742_at 0.92434 Nuclear factor I/A Nfia
1446743_at 0.95892 LAS1-like (S. cerevisiae) Las11
1446744_at 0.87156 sorbin and SH3 domain containing 2 Sorbs2
1446745_at 1.02982 hypothetical LOC433637 LOC433637
1446746_at 0.89651 Oxysterol binding protein-like 3 Osbpl3
1446747_at 0.86769 RIKEN cDNA C230086A09 gene C230086A09Rik
1446748_at 1.03001 RIKEN cDNA 2010007H06 gene 2010007H06Rik
1446749_at 1.02044 kinectin 1 Ktn1
1446750_at 1.24923 imprinted and ancient Impact
1446751_s_at 1.48041 imprinted and ancient Impact
1446752_at 1.00333 expressed sequence C78200 C78200
1446753_at 1.10760 --- ---
1446754_a_at 0.98970 Vax2 opposite strand transcript 1 Vax2os1
1446755_at 0.99369 Secretogranin V Scg5
1446756_at 1.11584 growth hormone secretagogue receptor Ghhr
1446757_at 1.10700 "SGT1, suppressor of G2 allele of SKP1 (S. cerevisiae)" Sugt1
1446758_at 1.02254 brain and reproductive organ-expressed protein Bre

1446759_at 1.31945 RIKEN cDNA 5830457H20 gene 5830457H20Rik
1446760_at 1.03276 thyroid hormone receptor interactor 13 Trip13
1446761_at 1.00181 "DNA segment, Chr 8, ERATO Doi 56, expressed" D8Ertd56e
1446762_at 1.00776 Transcribed locus ---
1446763_at 0.90838 "Potassium voltage-gated channel, subfamily H (eag-related), member 1" Kcnh1
1446764_at 0.93661 Wingless-related MMTV integration site 3A Wnt3a
1446765_at 0.88528 expressed sequence C77973 C77973
1446766_at 1.10287 expressed sequence AU022746 AU022746
1446767_at 0.96144 B-cell CLL/lymphoma 9 Bcl9
1446768_at 0.98547 Transcribed locus ---
1446769_at 1.10531 RIKEN cDNA 2810439F02 gene 2810439F02Rik
1446770_at 0.95640 RIKEN cDNA 2610208K16 gene 2610208K16Rik
1446771_at 1.01904 "Tubulin, alpha 8" Tuba8
1446772_at 0.95849 "Sema domain, seven thrombospondin repeats (type 1 and type 1-like),
transmembrane domain (TM) and short cytoplasmic domain, (semaphorin) 5A" Sema5a
1446773_at 1.00095 Hypothetical protein E230025N22 E230025N22
1446774_at 1.11711 "synovial apoptosis inhibitor 1, synoviolin" Syvn1
1446775_at 0.95752 --- ---
1446776_at 1.03715 expressed sequence C77581 C77581
1446777_at 0.94004 Secreted phosphoprotein 2 Spp2
1446778_at 1.06871 Hormonally upregulated Neu-associated kinase Hunk
1446779_at 1.20617 Nuclear protein localization 4 homolog (S. cerevisiae) Nploc4
1446780_at 1.05665 Receptor tyrosine kinase-like orphan receptor 2 Ror2
1446781_at 0.93463 "DNA segment, Chr 8, ERATO Doi 54, expressed" D8Ertd54e
1446782_at 0.92841 lines homolog 2 (Drosophila) Lins2
1446783_at 1.04725 "Growth arrest and DNA-damage-inducible, gamma interacting protein 1"
Gadd45gip1
1446784_at 1.00367 "A disintegrin-like and metallopeptidase (reprolysin type) with thrombospondin
type 1 motif, 6" Adamts6
1446785_at 0.86845 "Transcribed locus, weakly similar to XP_220328.3 PREDICTED: similar to
metalloprotease-disintegrin meltrin beta [Rattus norvegicus]" ---
1446786_at 0.94146 Amyloid beta (A4) precursor-like protein 2 Aplp2
1446787_at 1.22459 COMM domain containing 1 Commd1
1446788_at 1.04336 RIKEN cDNA A430091L06 gene A430091L06Rik
1446789_at 0.96097 Zinc finger CCHC-type and RNA binding motif 1 Zcrb1
1446790_at 0.91082 expressed sequence C78948 C78948
1446791_at 0.98813 Peptidase inhibitor 15 Pi15
1446792_at 0.95110 "Transcribed locus, moderately similar to XP_579119.1 PREDICTED: ATP-
binding cassette, sub-family G (WHITE), member 3 [Rattus norvegicus]" ---
1446793_at 0.97873 Transcribed locus ---
1446794_at 1.12453 G protein-coupled receptor 176 Gpr176
1446795_at 0.87169 Vav 3 oncogene Vav3
1446796_at 1.02400 --- ---

1446797_at	0.90066	RIKEN cDNA A330075M08 gene	A330075M08Rik
1446798_at	1.06119	---	---
1446799_at	0.98254	---	---
1446800_at	0.93711	---	---
1446801_at	0.97260	---	---
1446802_at	1.33146	ATX1 (antioxidant protein 1) homolog 1 (yeast)	Atox1
1446803_at	0.88946	Dedicator of cytokinesis 9	Dock9
1446804_at	0.96214	nucleoredoxin	Nxn
1446805_at	1.04937	"Gamma-aminobutyric acid (GABA-A) receptor, subunit alpha 2"	Gabra2
1446806_at	1.02801	---	---
1446807_at	1.15803	ubiquitin specific peptidase 8	Usp8
1446808_at	0.96079	---	---
1446809_at	0.89857	RIKEN cDNA B830045N13 gene	B830045N13Rik
1446810_at	1.27835	RIKEN cDNA A730062M13 gene	A730062M13Rik
1446811_at	1.02003	Mohawk	Mkx
1446812_at	0.92274	Heart and neural crest derivatives expressed transcript 2	Hand2
1446813_s_at	1.01320	"DNA segment, Chr 5, ERATO Doi 560, expressed"	D5ErtD560e
1446814_at	0.96491	---	---
1446815_at	1.25573	"Zinc finger, CSL-type containing 3"	Zcsl3
1446816_at	1.06149	Transcribed locus	---
1446817_at	1.09455	"DNA segment, Chr 10, ERATO Doi 43, expressed"	D10ErtD43e
1446818_at	1.29748	Mindbomb homolog 1 (Drosophila)	Mib1
1446819_at	1.01179	"RAN, member RAS oncogene family"	Ran
1446820_at	1.07135	zinc finger protein 182	Zfp182
1446821_at	1.06445	Protocadherin 17	Pcdh17
1446822_at	0.95215	---	---
1446823_at	0.93651	"DNA segment, Chr 4, ERATO Doi 103, expressed"	D4ErtD103e
1446824_at	0.91990	Phosphoglucomutase 1	Pgm1
1446825_at	0.95063	---	---
1446826_at	1.00427	exportin 7	Xpo7
1446827_at	0.94843	nuclear receptor coactivator 2	Ncoa2
1446828_at	0.90667	---	---
1446829_at	0.90878	"DNA segment, Chr 12, ERATO Doi 216, expressed"	D12ErtD216e
1446830_at	1.01106	expressed sequence AU015892	AU015892
1446831_at	1.05211	"DNA segment, Chr 5, ERATO Doi 521, expressed"	D5ErtD521e
1446832_at	0.89956	---	---
1446833_at	0.98608	Dedicator of cytokinesis 9	Dock9
1446834_at	1.01062	Cathepsin C	Ctsc
1446835_at	2.25637	Ferritin light chain 1	Ftl1
1446836_at	0.87776	---	---
1446837_at	0.99775	Methyltransferase 5 domain containing 1	Mett5d1
1446838_at	0.90037	"ATPase family, AAA domain containing 1"	Atad1
1446839_at	0.94795	Odd Oz/ten-m homolog 4 (Drosophila)	Odz4

1446840_at 1.13752 Thyroid hormone receptor associated protein 1 Thrap1
1446841_at 1.01192 "12 days embryo spinal ganglion cDNA, RIKEN full-length enriched library, clone:D130065L21 product:unclassifiable, full insert sequence" ---
1446842_at 0.99395 "DNA segment, Chr 4, ERATO Doi 571, expressed" D4Ertd571e
1446843_at 0.94369 --- ---
1446844_at 0.93348 coiled-coil-helix-coiled-coil-helix domain containing 7 Chchd7
1446845_at 1.05369 similar to RIKEN cDNA 1110067D22 LOC272350
1446846_at 0.87698 "16 days neonate thymus cDNA, RIKEN full-length enriched library, clone:A130082C12 product:unclassifiable, full insert sequence" ---
1446847_at 0.99220 RIKEN cDNA 0610010D24 gene 0610010D24Rik
1446848_at 1.09605 --- ---
1446849_at 0.98683 Autism susceptibility candidate 2 Auts2
1446850_at 0.90001 Phosphatidic acid phosphatase type 2B Ppap2b
1446851_at 0.86817 "collagen, type XXII, alpha 1" Col22a1
1446852_at 1.11670 "Serine (or cysteine) peptidase inhibitor, clade B, member 9b" Serpinb9b
1446853_at 0.98261 WW domain containing E3 ubiquitin protein ligase 1 Wwp1
1446854_at 1.03376 "Glutamate receptor, metabotropic 7" Grm7
1446855_at 1.09799 vacuolar protein sorting 13 D (yeast) Vps13d
1446856_at 0.98948 "DNA segment, Chr 3, Wayne State University 167, expressed" D3Wsu167e
1446857_at 1.09028 expressed sequence C87115 C87115
1446858_at 1.00658 expressed sequence AU022245 AU022245
1446859_at 0.99493 --- ---
1446860_at 1.04740 Growth factor receptor bound protein 2-associated protein 2 Gab2
1446861_at 1.38773 glucosamine (N-acetyl)-6-sulfatase Gns
1446862_at 1.09229 --- ---
1446863_at 0.92654 KCNQ1 overlapping transcript 1 Kcnq1ot1
1446864_at 1.03598 S100P binding protein S100pbp
1446865_at 0.95889 "Sema domain, seven thrombospondin repeats (type 1 and type 1-like), transmembrane domain (TM) and short cytoplasmic domain, (semaphorin) 5A" Sema5a
1446866_at 0.85319 RIKEN cDNA A530089I17 gene A530089I17Rik
1446867_at 1.04870 --- ---
1446868_at 1.08488 host cell factor C1 Hcfc1
1446869_at 0.89551 "In vitro fertilized eggs cDNA, RIKEN full-length enriched library, clone:7420455A11 product:unclassifiable, full insert sequence" ---
1446870_at 1.12406 "DNA segment, Chr 4, ERATO Doi 510, expressed" D4Ertd510e
1446871_at 0.94193 "DNA segment, Chr 7, ERATO Doi 481, expressed" D7Ertd481e
1446872_at 0.98900 expressed sequence AU021877 AU021877
1446873_at 0.95440 --- ---
1446874_at 0.96040 --- ---
1446875_at 0.96446 Interleukin 1 receptor accessory protein-like 2 Il1rapl2
1446876_at 1.03584 WD repeat domain 22 Wdr22
1446877_at 0.92932 Expressed sequence C87490 C87490
1446878_at 0.93512 LON peptidase N-terminal domain and ring finger 2 Lonrf2

1446879_at	0.90497	"Glutamate receptor, ionotropic, delta 2"	Grid2
1446880_at	0.91784	SAM and SH3 domain containing 1	Sash1
1446881_at	1.06573	expressed sequence AU022749	AU022749
1446882_at	0.95515	---	---
1446883_at	0.87274	RIKEN cDNA 4933427D06 gene	4933427D06Rik
1446884_at	1.04418	expressed sequence AU022554	AU022554
1446885_at	0.97082	---	---
1446886_at	1.09059	Ubiquitin specific peptidase 3	Usp3
1446887_at	1.07591	Transcribed locus	---
1446888_at	0.84657	---	---
1446889_at	0.93886	"DNA segment, Chr 2, ERATO Doi 105, expressed"	D2Ertd105e
1446890_at	0.97484	---	---
1446891_at	0.95543	"Amyotrophic lateral sclerosis 2 (juvenile) chromosome region, candidate 13 (human)"	Als2cr13
1446892_at	1.39122	Leucine rich repeat containing 16	Lrrc16
1446893_at	0.96385	hypothetical gene supported by AK035124	LOC434228
1446894_at	1.23961	Aryl hydrocarbon receptor nuclear translocator 2	Arnt2
1446895_at	0.83015	---	---
1446896_at	0.97105	expressed sequence AU022186	AU022186
1446897_at	0.95535	---	---
1446898_at	1.07889	Ubiquitin associated domain containing 1	Ubadc1
1446899_at	1.40191	---	---
1446900_at	1.01902	"DNA segment, Chr 1, ERATO Doi 218, expressed"	D1Ertd218e
1446901_at	0.94772	expressed sequence AU022077	AU022077
1446902_at	0.92961	expressed sequence AU022538	AU022538
1446903_at	1.08315	Angiopoietin 2	Angpt2
1446904_at	1.15341	Rho guanine nucleotide exchange factor (GEF) 11	Arhgef11
1446905_at	0.98474	Transcribed locus	---
1446906_at	1.10617	expressed sequence C81615	C81615
1446907_at	1.00930	expressed sequence C87679	C87679
1446908_at	0.99182	Interleukin 28 receptor alpha	Il28ra
1446909_at	0.91548	Angiomotin-like 1	Amotl1
1446910_at	1.04263	"Sema domain, seven thrombospondin repeats (type 1 and type 1-like), transmembrane domain (TM) and short cytoplasmic domain, (semaphorin) 5A"	Sema5a
1446911_at	1.11250	Intersectin 2	Itsn2
1446912_at	1.07626	Transmembrane protein 28	Tmem28
1446913_at	0.94147	---	---
1446914_at	1.12560	expressed sequence C80425	C80425
1446915_at	1.05326	"DNA segment, Chr 7, ERATO Doi 316, expressed"	D7Ertd316e
1446916_at	0.91433	---	---
1446917_at	1.06355	---	---
1446918_at	0.97885	Latrophilin 3	Lphn3
1446919_at	1.05231	CDNA sequence BC062115	BC062115

1446920_at	1.25683	RIKEN cDNA E130308A19 gene	E130308A19Rik
1446921_at	0.96146	Growth factor receptor bound protein 10	Grb10
1446922_at	1.03014	---	---
1446923_at	0.95429	"Protein tyrosine phosphatase-like (proline instead of catalytic arginine), member b"	Ptplb
1446924_at	0.91362	Transcribed locus	---
1446925_at	0.91109	beta-transducin repeat containing protein	Btrc
1446926_at	0.95581	PYD and CARD domain containing	Pycard
1446927_at	0.98020	Dedicator of cytokinesis 9	Dock9
1446928_at	1.05398	"DnaJ (Hsp40) homolog, subfamily C, member 17 /// similar to DnaJ (Hsp40) homolog, subfamily C, member 17"	Dnajc17 /// LOC638909
1446929_at	1.13300	BTB and CNC homology 2	Bach2
1446930_at	1.10771	Runt related transcription factor 1	Runx1
1446931_at	0.92727	Transcribed locus	---
1446932_at	0.98108	---	---
1446933_at	0.99031	---	---
1446934_at	0.90938	---	---
1446935_at	0.97066	RIKEN cDNA 9130011E15 gene	9130011E15Rik
1446936_at	0.97463	---	---
1446937_at	1.00128	fetal Alzheimer antigen Falz	
1446938_at	1.20035	expressed sequence AA408213	AA408213
1446939_at	0.98542	Tripartite motif protein 34	Trim34
1446940_at	0.73334	Tnf receptor-associated factor 6	Traf6
1446941_at	1.22153	expressed sequence C78505	C78505
1446942_at	1.02970	---	---
1446943_at	1.83358	Dystonin	Dst
1446944_at	1.01863	"DNA segment, Chr 1, ERATO Doi 84, expressed"	D1Ertd84e
1446945_at	1.00690	Cdc42 binding protein kinase alpha	Cdc42bpa
1446946_at	1.04878	"Transforming growth factor, beta receptor I"	Tgfr1
1446947_at	1.36516	---	---
1446948_at	1.05953	Density-regulated protein	Denr
1446949_at	1.10712	RIKEN cDNA 4930431P19 gene	4930431P19Rik
1446950_at	1.01833	Thymocyte selection-associated HMG box gene	Tox
1446951_at	0.85703	---	---
1446952_at	1.11722	"DNA segment, Chr 11, ERATO Doi 506, expressed"	D11Ertd506e
1446953_at	1.04230	Transcription factor 4	Tcf4
1446954_at	1.01275	Pyruvate dehydrogenase E1 alpha 1	Pdha1
1446955_at	1.37845	"Nuclear receptor subfamily 2, group C, member 2"	Nr2c2
1446956_at	0.89309	cDNA sequence BC004022	BC004022
1446957_s_at	1.05477	cDNA sequence BC004022	BC004022
1446958_at	1.17872	"DNA segment, Chr 13, ERATO Doi 150, expressed"	D13Ertd150e
1446959_at	0.94977	expressed sequence AU021871	AU021871
1446960_at	0.86015	Hypothetical protein LOC626082	LOC626082

1446961_at	0.98949 --- ---	
1446962_at	1.07411 "ATP-binding cassette, sub-family D (ALD), member 2"	Abcd2
1446963_at	1.76888 --- ---	
1446964_at	1.04050 --- ---	
1446965_at	1.11497 Rho guanine nucleotide exchange factor (GEF) 12	Arhgef12
1446966_at	1.13712 Expressed sequence AU042651	AU042651
1446967_at	1.04061 F-box and leucine-rich repeat protein 7	Fbxl7
1446968_at	1.31341 phosphatidylinositol binding clathrin assembly protein	Picalm
1446969_at	1.10247 RIKEN cDNA 9630037P07 gene	9630037P07Rik
1446970_at	0.86724 "CCR4-NOT transcription complex, subunit 8"	Cnot8
1446971_at	1.02133 "DNA segment, Chr 10, Wayne State University 159, expressed"	D10Wsu159e
1446972_at	4.79519 "DNA segment, Chr 15, Wayne State University 126, expressed"	D15Wsu126e
1446973_at	0.98785 RIKEN cDNA 5730405I09 gene	5730405I09Rik
1446974_at	1.09483 "phosphatidylinositol-3-phosphate/phosphatidylinositol 5-kinase, type III"	
Pip5k3		
1446975_at	1.00350 "Castor homolog 1, zinc finger (Drosophila)"	CasZ1
1446976_at	1.08591 "DNA segment, Chr 7, ERATO Doi 758, expressed"	D7Ertd758e
1446977_at	1.02922 Tnf receptor-associated factor 6	Traf6
1446978_at	0.98905 --- ---	
1446979_at	0.95900 "Decapping enzyme, scavenger"	Dcps
1446980_at	0.78057 Transcribed locus	---
1446981_at	0.99043 RIKEN cDNA A830010M20 gene	A830010M20Rik
1446982_at	1.80336 Par-3 (partitioning defective 3) homolog (C. elegans)	Pard3
1446983_at	0.97035 "Vcell division cycle 73, Paf1/RNA polymerase II complex component, homolog (S. cerevisiae)"	Cdc73
1446984_at	0.97162 Protogenin homolog (Gallus gallus)	Prtg
1446985_at	1.07012 "PREDICTED: Mus musculus similar to serine/threonine kinase (LOC626078), mRNA"	---
1446986_at	0.96949 "scratch homolog 2, zinc finger protein (Drosophila)"	Scr2
1446987_at	0.96653 --- ---	
1446988_at	1.03016 Nucleoporin 98	Nup98
1446989_at	0.94609 Transcribed locus	---
1446990_at	0.75991 nuclear factor I/A	Nfia
1446991_at	1.26105 GCN1 general control of amino-acid synthesis 1-like 1 (yeast)	Gcn111
1446992_at	1.00422 Coiled-coil domain containing 64	Ccdc64
1446993_at	0.86615 Perilipin	Plin
1446994_at	0.86549 Exosome component 2	Exosc2
1446995_at	0.93474 "cytochrome P450, family 7, subfamily b, polypeptide 1"	Cyp7b1
1446996_at	0.96127 --- ---	
1446997_at	1.02000 "eukaryotic translation initiation factor 3, subunit 7 (zeta)"	Eif3s7
1446998_at	0.90957 G protein-coupled receptor kinase 5	Gprk5
1446999_at	1.08067 --- ---	
1447000_at	1.06965 --- ---	

1447001_at	1.03096	expressed sequence C87114	C87114
1447002_at	0.97240	---	---
1447003_at	1.00946	Cell cycle progression 1	Ccpg1
1447004_at	1.01525	"Phospholipase C, beta 1"	Plcb1
1447005_at	0.95622	Transcribed locus	---
1447006_at	1.07523	Myosin ID	Myo1d
1447007_at	1.12364	RIKEN cDNA 1700008I05 gene	1700008I05Rik
1447008_at	1.11218	---	---
1447009_at	1.03412	expressed sequence C79657	C79657
1447010_at	1.05474	Zinc finger protein 609	Zfp609
1447011_at	0.90640	ERGIC and golgi 2	Ergic2
1447012_at	0.96570	---	---
1447013_at	1.05108	CDNA sequence AK220484	AK220484
1447014_at	1.05878	---	---
1447015_at	0.97010	---	---
1447016_at	1.34699	"TBC1 domain family, member 1"	Tbc1d1
1447017_at	1.04594	Transcribed locus	---
1447018_at	0.99900	---	---
1447019_at	1.07672	cytidine monophospho-N-acetylneuraminic acid hydroxylase	Cmah
1447020_at	1.80362	"Zinc finger protein, multitype 2"	Zfpm2
1447021_at	0.92942	CDNA clone IMAGE:5686051	---
1447022_at	0.85164	"Adult male testis cDNA, RIKEN full-length enriched library, clone:4930402K03 product:unclassifiable, full insert sequence"	---
1447023_at	1.02424	---	---
1447024_at	1.07208	---	---
1447025_at	1.47424	"Ubiquitin-conjugating enzyme E2E 2 (UBC4/5 homolog, yeast)"	Ube2e2
1447026_at	1.28665	MAS1 oncogene	Mas1
1447027_s_at	0.93329	lipoic acid synthetase	Lias
1447028_at	0.98347	RIKEN cDNA 4930563E22 gene	4930563E22Rik
1447029_at	0.88542	---	---
1447030_at	0.86510	---	---
1447031_at	1.19242	---	---
1447032_at	0.83581	Citrate lyase beta like	Clybl
1447033_at	0.97019	"gene model 829, (NCBI)"	Gm829
1447034_at	2.08556	Transportin 1	Tnpol
1447035_at	1.07309	RIKEN cDNA A230091C14 gene	A230091C14Rik
1447036_at	0.98094	cell adhesion molecule with homology to L1CAM	Chl1
1447037_at	0.89467	---	---
1447038_x_at	1.28394	pleckstrin homology domain interacting protein	Phip
1447039_at	0.95745	Patched homolog 1	Ptch1
1447040_at	1.03554	Activated leukocyte cell adhesion molecule	Alcam
1447041_at	0.97832	RIKEN cDNA 9530066K23 gene	9530066K23Rik
1447042_at	0.95108	zinc finger and BTB domain containing 34	Zbtb34

1447043_at	0.95663	V-erb-a erythroblastic leukemia viral oncogene homolog 4 (avian)	ErbB4
1447044_at	0.87684	---	---
1447045_at	1.03060	---	---
1447046_at	0.91821	"DnaJ (Hsp40) related, subfamily B, member 13"	Dnajb13
1447047_at	1.10967	RIKEN cDNA 2610203C20 gene	2610203C20Rik
1447048_at	1.00353	RIKEN cDNA 6720458F09 gene	6720458F09Rik
1447049_at	0.95471	IQ motif and WD repeats 1	Iqwd1
1447050_at	1.08083	---	---
1447051_at	0.85300	Ring finger protein 43	Rnf43
1447052_at	1.05822	---	---
1447053_x_at	1.00296	"signal sequence receptor, gamma"	Ssr3
1447054_at	0.96453	expressed sequence C85395	C85395
1447055_at	0.95849	"DnaJ (Hsp40) homolog, subfamily C, member 11"	Dnajc11
1447056_at	0.84482	Slit homolog 2 (Drosophila)	Slit2
1447057_at	0.99293	---	---
1447058_at	0.94799	RIKEN cDNA A530058N18 gene	A530058N18Rik
1447059_at	0.96851	---	---
1447060_at	1.47793	SET and MYND domain containing 3	Smyd3
1447061_at	1.03685	"Hypoxia inducible factor 3, alpha subunit"	Hif3a
1447062_at	1.53561	Vacuolar protein sorting 24 (yeast)	Vps24
1447063_at	1.45442	RIKEN cDNA 1700017B05 gene	1700017B05Rik
1447064_at	1.26159	---	---
1447065_at	0.87806	hypothetical protein 9630041C05	9630041C05
1447066_at	1.02900	"Adult inner ear cDNA, RIKEN full-length enriched library, clone:F930039H20 product:unclassifiable, full insert sequence"	---
1447067_at	0.92858	---	---
1447068_at	0.95458	"Phosphatidylinositol 3 kinase, regulatory subunit, polypeptide 3 (p55)"	Pik3r3
1447069_at	0.97234	Ring finger protein 43	Rnf43
1447070_at	0.97595	Williams-Beuren syndrome chromosome region 1 homolog (human)	Wbscr1
1447071_at	2.55847	"Transcription factor 7-like 2, T-cell specific, HMG-box"	Tcf712
1447072_at	0.87168	GRIP1 associated protein 1	Gripap1
1447073_at	1.07374	thyroid hormone receptor interactor 12	Trip12
1447074_at	0.86870	zinc finger and BTB domain containing 24	Zbtb24
1447075_at	0.98936	Class II transactivator C2ta	
1447076_at	0.95890	NEL-like 1 (chicken)	Nell1
1447077_at	0.87416	Adducin 3 (gamma)	Add3
1447078_at	1.01983	---	---
1447079_at	1.00266	---	---
1447080_at	1.04266	---	---
1447081_at	0.96896	---	---
1447082_at	0.99054	hypothetical protein C630004L07	C630004L07
1447083_at	1.04972	RIKEN cDNA 4933407I05 gene	4933407I05Rik

1447084_at 1.02466 "nuclear factor of activated T-cells, cytoplasmic, calcineurin-dependent 1"
Nfatc1
1447085_s_at 0.90103 "nuclear factor of activated T-cells, cytoplasmic, calcineurin-dependent 1"
Nfatc1
1447086_at 0.87710 SMC6 structural maintenance of chromosomes 6-like 1 (yeast) Smc6l1
1447087_at 1.02091 Cell growth regulator with ring finger domain 1 Cgrrf1
1447088_at 1.08811 Tweety homolog 2 (Drosophila) Ttyh2
1447089_at 1.00210 ADP-ribosylation factor-like 1 Arl1
1447090_s_at 1.04845 ADP-ribosylation factor-like 1 Arl1
1447091_at 1.10053 Jumonji domain containing 2C Jmjd2c
1447092_at 0.99677 Platelet/endothelial cell adhesion molecule 1 Pecam1
1447093_at 0.93966 "Procollagen, type XXIV, alpha 1" Col24a1
1447094_at 1.10938 --- ---
1447095_at 0.83202 expressed sequence C86942 C86942
1447096_at 1.45040 Transcribed locus ---
1447097_at 0.98934 cDNA sequence BC052688 BC052688
1447098_at 1.01090 --- ---
1447099_at 1.00116 Rho GTPase-activating protein Grit
1447100_s_at 1.03820 RIKEN cDNA 5730508B09 gene 5730508B09Rik
1447101_at 1.30923 ring finger protein 122 Rnf122
1447102_at 1.08080 Ninjurin 2 Ninj2
1447103_at 0.89359 CDK5 regulatory subunit associated protein 2 Cdk5rap2
1447104_at 0.99790 AT rich interactive domain 1A (Swi1 like) Arid1a
1447105_at 1.24888 Dephospho-CoA kinase domain containing Dcakd
1447106_at 1.04027 Leucine-rich repeat kinase 2 Lrrk2
1447107_at 1.08470 DEAD (Asp-Glu-Ala-Asp) box polypeptide 55 Ddx55
1447108_at 1.25151 RAD51-like 1 (S. cerevisiae) Rad51l1
1447109_at 1.61399 "ELAV (embryonic lethal, abnormal vision, Drosophila)-like 1 (Hu antigen
R)" Elavl1
1447110_at 0.95991 RIKEN cDNA 4921513D23 gene 4921513D23Rik
1447111_at 1.03653 --- ---
1447112_s_at 0.85148 "crystallin, lamda 1" Cry11
1447113_at 1.12789 --- ---
1447114_x_at 1.14478 RIKEN cDNA 4930414L22 gene 4930414L22Rik
1447115_at 0.84194 Expressed sequence C78409 C78409
1447116_at 1.13113 "Presenilin associated, rhomboid-like" Parl
1447117_at 0.85167 "Catenin (cadherin associated protein), alpha 2" Ctnna2
1447118_at 1.01093 "SAPS domain family, member 3" Saps3
1447119_at 1.13068 "Transcription termination factor, RNA polymerase II" Ttf2
1447120_at 1.15389 Beta-transducin repeat containing protein Btrc
1447121_at 1.12255 RIKEN cDNA 0610038K03 gene 0610038K03Rik
1447122_at 1.28327 --- ---
1447123_at 0.92314 --- ---

1447124_at 0.93444 Transcribed locus ---
1447125_at 0.93127 Keratin associated protein 5-5 Krtap5-5
1447126_at 0.96103 "Chromodomain protein, Y chromosome-like 2" Cdyl2
1447127_at 1.00634 "V-abl Abelson murine leukemia viral oncogene 2 (arg, Abelson-related gene)" Abl2
1447128_at 0.81805 Hypothetical LOC329575 LOC329575
1447129_at 0.96875 --- ---
1447130_at 1.05786 "DNA segment, Chr 8, Wayne State University 26, expressed" D8Wsu26e
1447131_at 0.94892 hypothetical protein LOC666453 /// hypothetical protein LOC671074 /// hypothetical protein LOC674374 LOC666453 /// LOC671074 /// LOC674374
1447132_at 1.04238 CDK5 regulatory subunit associated protein 1-like 1 Cdkal1
1447133_at 1.05683 --- ---
1447134_at 0.89137 SET domain containing 5 Setd5
1447135_at 1.08485 "Protein tyrosine phosphatase, receptor type, T" Ptprt
1447136_at 1.01140 Microtubule associated serine/threonine kinase-like Mastl
1447137_at 0.92478 RIKEN cDNA 4632417K18 gene 4632417K18Rik
1447138_at 0.95444 expressed sequence AI853502 AI853502
1447139_at 1.03117 B-cell CLL/lymphoma 7C Bcl7c
1447140_at 0.93346 hypothetical protein C130078N14 C130078N14
1447141_at 0.87631 expressed sequence AW107722 AW107722
1447142_at 1.02187 cathepsin 7 Cts7
1447143_at 1.45686 PHD finger protein 14 Phf14
1447144_at 0.91821 RIKEN cDNA 9130011E15 gene 9130011E15Rik
1447145_at 1.04000 --- ---
1447146_s_at 0.92385 "DNA segment, Chr 11, ERATO Doi 730, expressed" D11Ertd730e
1447147_at 1.79008 Autophagy-related 7 (yeast) Atg7
1447148_at 1.05424 Transcribed locus ---
1447149_at 1.00567 --- ---
1447150_at 1.42503 "pam, highwire, rpm 1" Phr1
1447151_at 0.95237 RIKEN cDNA 4930519N13 gene 4930519N13Rik
1447152_at 1.06045 Unc-51 like kinase 2 (C. elegans) Ulk2
1447153_x_at 1.21278 Unc-51 like kinase 2 (C. elegans) Ulk2
1447154_at 0.96385 "Calcium channel, voltage-dependent, beta 4 subunit" Cacnb4
1447155_at 1.14414 RIKEN cDNA D730039F16 gene D730039F16Rik
1447156_at 1.73596 Choroideremia Chm
1447157_at 0.90103 RIKEN cDNA 4930480K23 gene 4930480K23Rik
1447158_at 1.04462 --- ---
1447159_at 1.50956 Synaptic nuclear envelope 2 Syne2
1447160_at 1.07215 "non-POU-domain-containing, octamer binding protein /// hypothetical gene supported by AK004830; NM_023144" Nono /// LOC382354
1447161_at 1.11123 Transcribed locus ---
1447162_at 0.98570 --- ---
1447163_x_at 0.92053 --- ---

1447164_at	1.19275	Rearranged L-myc fusion sequence	Rlf
1447165_at	1.52920	Ankyrin repeat domain 12	Ankrd12
1447166_at	1.05523	RIKEN cDNA E230002P03 gene	E230002P03Rik
1447167_at	1.37986	Expressed sequence AL033314	AL033314
1447168_at	0.94127	Transcribed locus	---
1447169_at	0.93892	---	---
1447170_at	1.08692	---	---
1447171_at	1.00906	"Neural precursor cell expressed, developmentally down-regulated gene 4-like"	Nedd4l
1447172_at	0.93269	---	---
1447173_at	0.94465	"PREDICTED: Mus musculus hypothetical protein LOC624743 (LOC624743), mRNA"	---
1447174_at	1.04099	Dachshund 1 (Drosophila)	Dach1
1447175_at	1.09673	RIKEN cDNA 2610528M18 gene	2610528M18Rik
1447176_at	1.07300	RIKEN cDNA A930008G19 gene	A930008G19Rik
1447177_at	0.89514	---	---
1447178_at	0.92408	---	---
1447179_at	1.12364	RIKEN cDNA 2210412D01 gene	2210412D01Rik
1447180_at	0.90607	Transcribed locus	---
1447181_s_at	0.85193	"solute carrier family 7 (cationic amino acid transporter, y+ system), member 7"	Slc7a7
1447182_at	1.01142	expressed sequence C77815	C77815
1447183_at	0.99762	Odd Oz/ten-m homolog 3 (Drosophila)	Odz3
1447184_at	1.37569	expressed sequence AU021760	AU021760
1447185_at	0.92842	RIKEN cDNA 9330158H04 gene	9330158H04Rik
1447186_at	0.95479	"solute carrier family 6, member 16 /// similar to Orphan sodium- and chloride-dependent neurotransmitter transporter NTT5 (Solute carrier family 6 member 16)"	Slc6a16 /// LOC672880
1447187_at	1.10378	Receptor interacting protein kinase 5	Ripk5
1447188_at	0.87539	Transcribed locus	---
1447189_at	0.99135	Glioma tumor suppressor candidate region gene 1	Gltscr1
1447190_at	1.07885	---	---
1447191_at	1.00581	---	---
1447192_at	1.04722	---	---
1447193_at	0.88843	---	---
1447194_at	0.88729	Transcribed locus	---
1447195_at	1.27816	Elongation protein 4 homolog (S. cerevisiae)	Elp4
1447196_at	0.89444	hypothetical protein LOC622866	MGC58416
1447197_at	1.06069	expressed sequence C86544	C86544
1447198_at	1.05731	RecQ protein-like	Recql
1447199_at	1.05340	expressed sequence C80763	C80763
1447200_at	1.00341	"TBC1 domain family, member 5"	Tbc1d5
1447201_at	0.96684	---	---

1447202_at 1.02177 RIKEN cDNA 1200009F10 gene 1200009F10Rik
1447203_at 0.90331 --- ---
1447204_at 0.88511 Mitogen-activated protein kinase kinase kinase kinase 5 Map4k5
1447205_x_at 1.10850 ribosomal protein S12 /// similar to ribosomal protein S12 /// similar to
ribosomal protein S12 /// similar to ribosomal protein S12 /// similar to ribosomal protein S12 /// similar
to ribosomal protein S12 /// similar to ribosomal protein S12 /// similar to ribosomal protein S12 ///
similar to ribosomal protein S12 /// similar to ribosomal protein S12 /// similar to ribosomal protein
S12 /// similar to ribosomal protein S12 /// similar to ribosomal protein S12 /// similar to ribosomal
protein S12 /// similar to ribosomal protein S12 /// similar to 40S ribosomal protein S12 /// similar to
ribosomal protein S12 /// similar to ribosomal protein S12 /// similar to ribosomal protein S12 /// similar
to ribosomal protein S12 Rps12 /// LOC432865 /// LOC434843 /// LOC546371 /// LOC626048 ///
LOC627835 /// LOC665463 /// LOC665501 /// LOC665522 /// LOC665555 /// LOC665579 ///
LOC665611 /// LOC668354 /// LOC670751 /// LOC670832 /// LOC671641 /// LOC672008 ///
LOC672415 /// LOC673522 /// LOC676277
1447206_at 1.43075 Rho GTPase activating protein 21 Arhgap21
1447207_at 0.95755 RIKEN cDNA 2810482G21 gene 2810482G21Rik
1447208_at 1.05472 Transcribed locus ---
1447209_at 1.36770 Forkhead box P1 Foxp1
1447210_at 1.13876 Transcribed locus ---
1447211_at 0.93673 nuclear receptor interacting protein 1 Nrip1
1447212_at 1.00409 "CCR4-NOT transcription complex, subunit 2" Cnot2
1447213_at 1.02063 Nuclear antigen Sp100 Sp100
1447214_at 0.98197 hypothetical protein LOC666827 /// hypothetical protein LOC670370 ///
hypothetical protein LOC674049 /// hypothetical protein LOC676969 LOC666827 ///
LOC670370 /// LOC674049 /// LOC676969
1447215_at 0.92311 WD repeat domain 66 Wdr66
1447216_at 1.00403 neurexin I Nrnx1
1447217_at 1.18821 "ubiquitin-like, containing PHD and RING finger domains 2" Uhrf2
1447218_at 1.23338 RIKEN cDNA 1110003F02 gene 1110003F02Rik
1447219_at 0.85263 Transcribed locus ---
1447220_at 1.14698 RIKEN cDNA 1700125H20 gene 1700125H20Rik
1447221_x_at 0.96996 Fibroblast growth factor 14 Fgf14
1447222_at 0.87130 heat shock protein 12A Hspa12a
1447223_at 1.00870 Transcribed locus ---
1447224_at 0.97083 Transcribed locus ---
1447225_at 0.83789 "solute carrier family 16 (monocarboxylic acid transporters), member 5" Slc16a5
1447226_at 1.09984 --- ---
1447227_at 0.93106 "Solute carrier family 40 (iron-regulated transporter), member 1" Slc40a1
1447228_at 0.91637 RIKEN cDNA 1110051M20 gene 1110051M20Rik
1447229_x_at 0.99454 RIKEN cDNA 1110051M20 gene 1110051M20Rik
1447230_at 0.91493 Expressed sequence AI553587 AI553587
1447231_at 0.96789 "Solute carrier family 8 (sodium/calcium exchanger), member 1" Slc8a1
1447232_at 1.04369 --- ---

1447233_at 0.98300 RIKEN cDNA D330027H18 gene D330027H18Rik
 1447234_s_at 0.94556 sorting nexin 6 Snx6
 1447235_at 1.13614 SH3 domain containing ring finger 1 Sh3rf1
 1447236_at 1.04425 kinesin family member 21A Kif21a
 1447237_at 1.05947 "Phosphodiesterase 4B, cAMP specific" Pde4b
 1447238_at 0.94966 --- ---
 1447239_at 0.86160 Translocase of outer mitochondrial membrane 70 homolog A (yeast)
 Tomm70a
 1447240_at 3.31804 --- ---
 1447241_at 1.10744 "H2A histone family, member Y2" H2afy2
 1447242_at 1.22562 "Presenilin associated, rhomboid-like" Parl
 1447243_at 1.09037 cDNA sequence BC040756 BC040756
 1447244_at 1.03774 "cytochrome b, ascorbate dependent 3" Cybasc3
 1447245_at 1.06835 Transcribed locus ---
 1447246_at 1.08853 "0 day neonate thymus cDNA, RIKEN full-length enriched library, clone:
 A430092C21 product:hypothetical protein, full insert sequence" ---
 1447247_at 1.00088 "Protein phosphatase 1, regulatory subunit 9B" Ppp1r9b
 1447248_at 1.04898 "solute carrier family 39 (zinc transporter), member 12" Slc39a12
 1447249_at 0.96238 Transcribed locus ---
 1447250_a_at 0.92222 RIKEN cDNA 2610301F02 gene 2610301F02Rik
 1447251_x_at 1.00541 RIKEN cDNA 2610301F02 gene 2610301F02Rik
 1447252_s_at 0.89849 mepirin 1 alpha Mep1a
 1447253_x_at 1.01041 nucleoporin 54 Nup54
 1447254_at 1.06835 --- ---
 1447255_at 0.88538 RIKEN cDNA 2310015A10 gene 2310015A10Rik
 1447256_at 0.98046 "17 days embryo kidney cDNA, RIKEN full-length enriched library, clone:
 I920178P20 product:unclassifiable, full insert sequence" ---
 1447257_at 1.11512 TAO kinase 3 Taok3
 1447258_at 1.34112 MYST histone acetyltransferase monocytic leukemia 4 Myst4
 1447259_at 0.96644 "ankyrin 3, epithelial" Ank3
 1447260_at 1.07107 "histocompatibility 2, blastocyst /// RIKEN cDNA 2410017I17 gene /// similar
 to MHC class I antigen precursor /// MHC class I like protein GS17 /// similar to histocompatibility 2,
 blastocyst /// similar to RT1 class Ia, locus A1 /// similar to RT1 class Ia, locus A1 /// hypothetical
 protein LOC677505" H2-B1 /// 2410017I17Rik /// LOC547349 /// H2-gs17 /// LOC673639 ///
 LOC674137 /// LOC676837 /// LOC677505
 1447261_at 0.92146 "PREDICTED: Mus musculus similar to CG12341-PA (LOC621976),
 mRNA" ---
 1447262_at 0.97973 Transcribed locus ---
 1447263_at 1.57932 Methionine aminopeptidase 2 Metap2
 1447264_at 1.02653 RAB11 family interacting protein 1 (class I) Rab11fip1
 1447265_at 1.01841 expressed sequence AI848615 AI848615
 1447266_at 0.96459 "UTP18, small subunit (SSU) processome component, homolog (yeast)"
 Utp18

1447267_at	0.95867	expressed sequence C85319	C85319
1447268_at	1.16304	nucleolar protein 12	Nol12
1447269_at	0.90611	RIKEN cDNA 9630031F12 gene	9630031F12Rik
1447270_at	1.05969	---	---
1447271_at	0.93357	non-catalytic region of tyrosine kinase adaptor protein 1	Nck1
1447272_s_at	0.96665	"ATPase, class V, type 10A"	Atp10a
1447273_x_at	0.94496	"ATPase, class V, type 10A"	Atp10a
1447274_at	1.18808	Wiskott-Aldrich syndrome protein interacting protein	Waspip
1447275_at	0.96965	"gene model 1805, (NCBI)"	Gm1805
1447276_at	1.00195	"DNA segment, Chr 13, ERATO Doi 37, expressed"	D13Ert37e
1447277_s_at	0.84098	prenylcysteine oxidase 1	Pcyox1
1447278_at	1.04014	centrosomal protein 164	Cep164
1447279_at	0.85284	Transcribed locus	---
1447280_at	0.95944	Dystrobrevin alpha	Dtna
1447281_at	0.86804	---	---
1447282_at	0.98521	---	---
1447283_at	0.98405	testis specific 10	Tsga10
1447284_at	1.01542	triggering receptor expressed on myeloid cells 1	Trem1
1447285_at	1.00773	---	---
1447286_at	1.01461	Embigin	Emb
1447287_at	1.11550	expressed sequence C77137	C77137
1447288_at	1.00612	Transcribed locus	---
1447289_at	0.97128	expressed sequence AA763521	AA763521
1447290_at	1.09927	RIKEN cDNA 2610042O14 gene	2610042O14Rik
1447291_at	1.05135	"nephrosis 1 homolog, nephrin antisense"	Nphs1as
1447292_at	1.08060	ARP1 actin-related protein 1 homolog B (yeast)	Actr1b
1447293_x_at	1.02705	ARP1 actin-related protein 1 homolog B (yeast)	Actr1b
1447294_at	1.22273	F-box and WD-40 domain protein 11	Fbxw11
1447295_at	1.66168	Cut-like 1 (Drosophila)	Cutl1
1447296_at	0.95670	F-box protein 27	Fbxo27
1447297_at	0.97209	"Solute carrier family 25, member 36"	Slc25a36
1447298_at	0.97052	RIKEN cDNA C130071C03 gene	C130071C03Rik
1447299_at	0.92428	---	---
1447300_at	0.91227	expressed sequence AW551984	AW551984
1447301_at	1.04708	A kinase (PRKA) anchor protein 5	Akap5
1447302_at	1.01484	Transcribed locus	---
1447303_at	1.01903	---	---
1447304_at	0.83411	"Mammary gland RCB-0527 Jyg-MC(B) cDNA, RIKEN full-length enriched library, clone:G930010E14 product:unclassifiable, full insert sequence"	---
1447305_at	0.96391	T-cell lymphoma breakpoint associated target 1	Tcba1
1447306_at	1.11665	RIKEN cDNA 4930545H06 gene	4930545H06Rik
1447307_at	1.03350	Transcribed locus	---
1447308_at	0.97228	longevity assurance homolog 5 (S. cerevisiae)	Lass5

1447309_at 1.05523 "Adult male testis cDNA, RIKEN full-length enriched library, clone:4930539H17 product:unclassifiable, full insert sequence" ---
1447310_at 1.06320 RIKEN cDNA C630028N24 gene C630028N24Rik
1447311_at 1.03955 "solute carrier family 6 (neurotransmitter transporter, noradrenalin), member 2" Slc6a2
1447312_at 1.23884 Ankyrin repeat and IBR domain containing 1 Ankib1
1447313_at 1.06334 --- ---
1447314_at 1.02292 "Transforming growth factor, beta receptor III" Tgfb3
1447315_at 1.04699 --- ---
1447316_at 0.98795 Transcribed locus ---
1447317_at 0.85991 Transcribed locus ---
1447318_at 0.88268 UDP-N-acetyl-alpha-D-galactosamine:polypeptide N-acetylgalactosaminyltransferase 4 Galnt4
1447319_at 0.91139 --- ---
1447320_x_at 1.05437 RNA polymerase 1-3 Rpo1-3
1447321_at 0.95361 "Amyotrophic lateral sclerosis 2 (juvenile) chromosome region, candidate 19 (human)" Als2cr19
1447322_at 1.00796 --- ---
1447323_at 1.01368 Hypothetical protein LOC621954 LOC621954
1447324_at 1.06484 --- ---
1447325_at 1.06706 Checkpoint suppressor 1 Ches1
1447326_s_at 0.74285 "zinc finger, MYM-type 3" Zmym3
1447327_at 0.84777 --- ---
1447328_at 0.87099 "Inositol 1,4,5-triphosphate receptor 2" Itpr2
1447329_at 1.04304 --- ---
1447330_at 0.97334 zinc finger and BTB domain containing 32 Zbtb32
1447331_at 0.97282 F-box protein 42 Fbxo42
1447332_at 0.86998 expressed sequence C86896 C86896
1447333_at 0.99668 "Solute carrier family 11 (proton-coupled divalent metal ion transporters), member 2" Slc11a2
1447334_at 1.13863 B-cell CLL/lymphoma 11A (zinc finger protein) Bcl11a
1447335_x_at 1.04044 B-cell CLL/lymphoma 11A (zinc finger protein) Bcl11a
1447336_at 0.79071 "Adult male kidney cDNA, RIKEN full-length enriched library, clone: F530012N16 product:hypothetical protein, full insert sequence" ---
1447337_at 1.13774 Dual adaptor for phosphotyrosine and 3-phosphoinositides 1 Dapp1
1447338_at 0.95609 Transcribed locus ---
1447339_at 0.94844 Transcribed locus ---
1447340_at 0.90270 "Adult male kidney cDNA, RIKEN full-length enriched library, clone: F530214C21 product:unclassifiable, full insert sequence" ---
1447341_at 1.06587 "DNA segment, Chr 12, ERATO Doi 551, expressed" D12Ert551e
1447342_at 1.07578 --- ---
1447343_at 1.11856 --- ---
1447344_at 0.84487 Transcribed locus ---

1447345_at 0.99944 RIKEN cDNA A430107O13 gene A430107O13Rik
1447346_s_at 1.03486 RIKEN cDNA 5530400B01 gene 5530400B01Rik
1447347_at 0.97505 Amyloid beta (A4) precursor protein App
1447348_at 1.09215 Wiskott-Aldrich syndrome protein interacting protein Waspip
1447349_s_at 1.00371 E1A binding protein p400 Ep400
1447350_at 1.05289 --- ---
1447351_x_at 1.09533 --- ---
1447352_at 0.97901 "8 days neonate male testis cDNA, RIKEN full-length enriched library, clone:
G630041I22 product:unclassifiable, full insert sequence" ---
1447353_at 1.09764 G protein-coupled receptor 161 Gpr161
1447354_at 0.95857 Centrosomal protein 164 Cep164
1447355_at 0.99115 acyl-CoA synthetase long-chain family member 1 Acs11
1447356_at 0.93492 UbiE-YGHL1 fusion protein LOC554292
1447357_at 0.88306 --- ---
1447358_at 0.96349 expressed sequence C87198 C87198
1447359_at 0.97755 zinc finger protein 575 Zfp575
1447360_at 1.01805 "TSC22 domain family, member 1" Tsc22d1
1447361_at 0.99940 Metaxin 1 Mtx1
1447362_at 1.02465 "budding uninhibited by benzimidazoles 1 homolog, beta (S. cerevisiae)" Bub1b
1447363_s_at 0.93798 "budding uninhibited by benzimidazoles 1 homolog, beta (S. cerevisiae)"
Bub1b
1447364_x_at 0.93018 myosin IB Myo1b
1447365_at 0.89869 Transcribed locus ---
1447366_at 1.08945 --- ---
1447367_at 0.96296 sec1 family domain containing 2 Scfd2
1447368_at 0.87849 RIKEN cDNA 6430706D22 gene 6430706D22Rik
1447369_at 1.00392 RIKEN cDNA 1190005F20 gene 1190005F20Rik
1447370_at 1.09957 Transmembrane protein 16A Tmem16a
1447371_at 0.88853 --- ---
1447372_at 1.00810 "aldehyde dehydrogenase 16 family, member A1" Aldh16a1
1447373_at 1.02173 RIKEN cDNA 1810015C04 gene 1810015C04Rik
1447374_at 0.95110 Ubiquitin-conjugating enzyme E2H Ube2h
1447375_at 0.97439 Transcribed locus ---
1447376_at 0.92595 Moloney leukemia virus 10 Mov10
1447377_at 0.86108 "gene model 428, (NCBI) /// hypothetical LOC623180 /// hypothetical
LOC623197 /// hypothetical LOC623210 /// hypothetical LOC623219" Gm428 /// LOC623180 ///
LOC623197 /// LOC623210 /// LOC623219
1447378_at 1.06798 "PREDICTED: Mus musculus hypothetical protein LOC628013 (LOC628013),
mRNA" ---
1447379_at 1.05723 SAP30-like Sap30l
1447380_at 0.97322 amidohydrolase domain containing 1 Amdhd1
1447381_at 1.07369 Cleavage and polyadenylation specific factor 6 Cpsf6
1447382_at 1.13344 "phosphatidylinositol glycan anchor biosynthesis, class T" Pigt

1447383_at 0.99292 CDNA sequence BC005471 BC005471
1447384_at 1.02779 "ATP-binding cassette, sub-family C (CFTR/MRP), member 5" Abcc5
1447385_at 0.89596 "Adult male corpora quadrigemina cDNA, RIKEN full-length enriched library, clone:B230036G05 product:unclassifiable, full insert sequence" ---
1447386_at 1.02536 Ubiquitin specific peptidase 8 Usp8
1447387_at 1.00031 --- ---
1447388_at 0.85647 RIKEN cDNA 5830484A20 gene 5830484A20Rik
1447389_at 1.10083 expressed sequence AI662168 AI662168
1447390_at 1.14407 --- ---
1447391_at 1.07023 RUN and FYVE domain containing 1 Rufy1
1447392_s_at 0.81359 carboxypeptidase D Cpd
1447393_at 1.13650 "PREDICTED: Mus musculus hypothetical LOC434436 (LOC434436), mRNA" ---
1447394_at 1.07802 chloride channel calcium activated 6 /// expressed sequence AI747448 /// similar to chloride channel calcium activated 6 Clca6 /// AI747448 /// LOC622139
1447395_at 1.01934 RIKEN cDNA 4930525K10 gene 4930525K10Rik
1447396_at 1.06766 Hypothetical protein LOC545253 LOC545253
1447397_at 0.98850 RIKEN cDNA 1700129I04 gene 1700129I04Rik
1447398_at 0.99672 --- ---
1447399_at 1.12248 RIKEN cDNA A330019N05 gene A330019N05Rik
1447400_at 0.89827 receptor-like tyrosine kinase Ryk
1447401_at 0.92883 WW domain binding protein 11 Wbp11
1447402_at 1.14667 Transcribed locus ---
1447403_a_at 1.04970 "Zinc finger, MYND domain containing 19" Zmynd19
1447404_at 0.95780 "Gene model 96, (NCBI)" Gm96
1447405_at 0.89920 Expressed sequence AI449023 AI449023
1447406_at 1.00610 "PREDICTED: Mus musculus similar to Hemoglobin alpha subunit (Hemoglobin alpha chain) (Alpha-globin) (LOC383229), mRNA" ---
1447407_at 1.06621 "centaurin, delta 3" Centd3
1447408_at 3.04457 Nipped-B homolog (Drosophila) Nipbl
1447409_at 1.03844 --- ---
1447410_at 0.97003 --- ---
1447411_at 1.51251 UDP-glucose dehydrogenase Ugdh
1447412_at 0.88782 "gene model 996, (NCBI)" Gm996
1447413_at 1.01928 RIKEN cDNA 1110001A07 gene 1110001A07Rik
1447414_at 1.02697 --- ---
1447415_at 0.92202 --- ---
1447416_at 0.96117 "gene model 239, (NCBI)" Gm239
1447417_at 0.97870 "TAF1 RNA polymerase II, TATA box binding protein (TBP)-associated factor" Taf1
1447418_at 0.89502 "Potassium voltage-gated channel, subfamily Q, member 1" Kcnq1
1447419_at 0.86871 Transcribed locus ---
1447420_at 0.99416 bromodomain containing 1 Brd1

1447421_at	2.32708	Transcribed locus	---	
1447422_at	1.03109	GTPase activating RANGAP domain-like 1	Garnl1	
1447423_at	1.06210	---	---	
1447424_at	1.05686	---	---	
1447425_at	0.97126	RIKEN cDNA 2900093B09 gene	2900093B09Rik	
1447426_at	1.14498	Dip3 beta	Dip3b	
1447427_at	1.04823	"PREDICTED: Mus musculus similar to developmentally regulated repeat element-containing transcript 1 (LOC622619), mRNA"	---	
1447428_at	0.93198	---	---	
1447429_at	1.11091	Transcribed locus	---	
1447430_at	0.98953	Transcribed locus	---	
1447431_at	0.93045	2-deoxyribose-5-phosphate aldolase homolog (C. elegans)	Dera	
1447432_s_at	1.00677	zinc finger protein 263	Zfp263	
1447433_at	1.02404	WD repeat and FYVE domain containing 3	Wdfy3	
1447434_at	0.93227	Cyclin D3	Ccnd3	
1447435_at	0.89212	---	---	
1447436_at	0.98698	expressed sequence AI451250	AI451250	
1447437_at	0.94380	Acyl-CoA synthetase medium-chain family member 1	Acsm1	
1447438_at	1.09199	Whirlin	Whrn	
1447439_at	1.02305	---	---	
1447440_at	0.92354	Nuclear transport factor 2	Nutf2	
1447441_at	0.88300	Rho guanine nucleotide exchange factor (GEF) 17	Arhgef17	
1447442_at	0.90551	Helicase with zinc finger domain	Helz	
1447443_at	0.98294	"Sparc/osteonectin, cwcv and kazal-like domains proteoglycan 2"	Spock2	
1447444_at	0.98862	UBX domain containing 5	Ubx5	
1447445_at	1.11590	Transcribed locus	---	
1447446_at	0.88973	"RAS-like, estrogen-regulated, growth-inhibitor"	Rerg	
1447447_s_at	1.03693	serine/arginine repetitive matrix 1	Srrm1	
1447448_s_at	1.12453	Kruppel-like factor 6	Klf6	
1447449_at	0.99276	---	---	
1447450_at	1.28660	Peroxisome proliferator activated receptor binding protein	Pparbp	
1447451_at	1.04112	RIKEN cDNA 4930556M19 gene	4930556M19Rik	
1447452_at	1.00661	---	---	
1447453_x_at	1.09524	RIKEN cDNA A230083H22 gene	A230083H22Rik	
1447454_at	1.06017	Kinesin family member 5C	Kif5c	
1447455_at	0.99008	---	---	
1447456_x_at	0.88386	similar to common salivary protein 1	LOC671351	
1447457_at	1.03476	"Protein kinase inhibitor, gamma"	Pkig	
1447458_at	0.92677	"ST3 beta-galactoside alpha-2,3-sialyltransferase 4"	St3gal4	
1447459_at	1.24730	Apoptosis antagonizing transcription factor	Aatf	
1447460_at	0.95565	Ribosomal protein L5	Rpl5	
1447461_at	0.91221	---	---	
1447462_at	1.06298	"DNA segment, Chr 7, Wayne State University 130, expressed"	D7Wsu130e	

1447463_at 1.08549 --- ---
1447464_at 0.93588 ankyrin repeat and sterile alpha motif domain containing 1B Anks1b
1447465_at 0.89867 RIKEN cDNA 4930558N01 gene 4930558N01Rik
1447466_at 0.97714 PRP40 pre-mRNA processing factor 40 homolog A (yeast) Prpf40a
1447467_at 0.97532 "protein tyrosine phosphatase, receptor type, A" Ptpra
1447468_at 1.12969 Transcribed locus ---
1447469_at 0.90700 coiled-coil domain containing 66 Ccdc66
1447470_at 0.99488 haloacid dehalogenase-like hydrolase domain containing 2 Hdhd2
1447471_at 1.02670 "Mammary gland RCB-0527 Jyg-MC(B) cDNA, RIKEN full-length enriched library, clone:G930010E14 product:unclassifiable, full insert sequence" ---
1447472_at 1.01244 Transcribed locus ---
1447473_at 1.08092 Transcribed locus ---
1447474_at 1.03301 Receptor (calcitonin) activity modifying protein 1 Ramp1
1447475_at 0.76726 "gene model 967, (NCBI)" Gm967
1447476_at 1.02373 "ATP-binding cassette, sub-family C (CFTR/MRP), member 10" Abcc10
1447477_at 0.93555 RIKEN cDNA 3110006E14 gene 3110006E14Rik
1447478_at 1.07822 "RAN, member RAS oncogene family" Ran
1447479_at 0.93203 --- ---
1447480_at 1.01569 Fibroblast growth factor 14 Fgf14
1447481_at 1.43975 SET domain containing 5 Setd5
1447482_at 1.03284 RIKEN cDNA 5830457H20 gene 5830457H20Rik
1447483_s_at 0.98082 RIKEN cDNA 2610002F03 gene 2610002F03Rik
1447484_x_at 1.10611 RIKEN cDNA 2610002F03 gene 2610002F03Rik
1447485_at 1.01299 G protein-coupled receptor 150 Gpr150
1447486_at 1.00981 "protein phosphatase 1, regulatory subunit 9B" Ppp1r9b
1447487_at 0.98361 --- ---
1447488_at 1.00778 RIKEN cDNA E130016E03 gene E130016E03Rik
1447489_at 0.94167 RIKEN cDNA 1110039B18 gene 1110039B18Rik
1447490_at 0.95966 Transcribed locus ---
1447491_at 1.31633 Leucine rich repeat containing 8D Lrrc8d
1447492_at 0.96085 --- ---
1447493_at 0.91935 RIKEN cDNA A530088H08 gene A530088H08Rik
1447494_at 0.97423 "DNA segment, Chr 7, Brigham & Women's Genetics 0826 expressed" D7Bwg0826e
1447495_at 0.94640 fibronectin type 3 and ankyrin repeat domains 1 Fank1
1447496_s_at 0.99833 fibronectin type 3 and ankyrin repeat domains 1 Fank1
1447497_at 0.97907 "RAB21, member RAS oncogene family" Rab21
1447498_at 0.98881 ectodysplasin-A receptor /// similar to Tumor necrosis factor receptor superfamily member EDAR precursor (Anhidrotic ectodysplasin receptor 1) (Ectodysplasin-A receptor) (Ectodermal dysplasia receptor) (Downless) Edar /// LOC669001
1447499_s_at 0.97528 "oocyte maturation, alpha /// oocyte maturation, beta /// similar to oocyte maturation, beta" Omt2a /// Omt2b /// LOC640894
1447500_at 0.92925 cut-like 2 (Drosophila) Cutl2

1447501_at	1.03393	Transcribed locus	---	
1447502_at	1.00114	Cytidine monophospho-N-acetylneuraminic acid hydroxylase		Cmah
1447503_at	1.12116	CDNA sequence BC016495	BC016495	
1447504_at	1.16284	Growth hormone inducible transmembrane protein	Ghitm	
1447505_at	0.98230	RIKEN cDNA 1700020G03 gene	1700020G03Rik	
1447506_at	1.00891	---	---	
1447507_at	0.92263	expressed sequence C80406	C80406	
1447508_at	1.12476	Transcribed locus	---	
1447509_at	0.91402	RIKEN cDNA 1810057P16 gene	1810057P16Rik	
1447510_at	1.52881	RIKEN cDNA C530014P21 gene	C530014P21Rik	
1447511_at	0.99519	Transcribed locus	---	
1447512_at	1.00303	Cell adhesion molecule with homology to L1CAM	Chl1	
1447513_at	0.95216	"potassium voltage-gated channel, Shal-related family, member 3"		Kcnd3
1447514_at	0.95119	RIKEN cDNA D030041N04 gene	D030041N04Rik	
1447515_at	1.06456	Transcribed locus	---	
1447516_at	0.93207	Transcribed locus	---	
1447517_at	2.11044	superkiller viralicidic activity 2-like 2 (<i>S. cerevisiae</i>)	Skiv2I2	
1447518_at	0.94024	"TPX2, microtubule-associated protein homolog (<i>Xenopus laevis</i>)"	Tpx2	
1447519_x_at	0.99437	"TPX2, microtubule-associated protein homolog (<i>Xenopus laevis</i>)"	Tpx2	
1447520_at	1.21563	---	---	
1447521_x_at	1.00311	"DNA segment, Chr 15, Wayne State University 169, expressed"		
D15Wsu169e				
1447522_s_at	0.94631	"tankyrase, TRF1-interacting ankyrin-related ADP-ribose polymerase 2"		
Tnks2				
1447523_at	1.07432	Zinc finger protein 294	Zfp294	
1447524_at	1.03948	"adaptor-related protein complex 3, mu 2 subunit"	Ap3m2	
1447525_at	0.95609	Intraflagellar transport 172 homolog (<i>Chlamydomonas</i>)	Ift172	
1447526_at	1.09375	"DNA segment, Chr 5, ERATO Doi 255, expressed"	D5Ert255e	
1447527_at	1.37859	Interferon regulatory factor 2	Irf2	
1447528_at	1.09527	---	---	
1447529_at	0.93084	---	---	
1447530_at	0.99272	Factor 8-associated gene A	F8a	
1447531_x_at	0.99042	---	---	
1447532_at	0.96053	RIKEN cDNA 1700025G04 gene	1700025G04Rik	
1447533_at	1.08668	Transcribed locus	---	
1447534_at	1.12562	---	---	
1447535_at	0.96746	BTB (POZ) domain containing 4	Btbd4	
1447536_at	0.98122	"LysM, putative peptidoglycan-binding, domain containing 2"		Lysmd2
1447537_at	1.26556	RIKEN cDNA 1500032P08 gene	1500032P08Rik	
1447538_at	1.01177	CREB regulated transcription coactivator 1	Crtc1	
1447539_at	0.97420	Transcribed locus	---	
1447540_at	0.94610	tigger transposable element derived 3	Tigd3	
1447541_s_at	0.82180	"integrin, alpha E, epithelial-associated"		Itgae

1447542_at	1.02082	cDNA sequence AY702102	AY702102	
1447543_at	0.91931	WD repeat and FYVE domain containing 1	Wdfy1	
1447544_at	1.00478	Transcribed locus	---	
1447545_at	1.11670	"DNA segment, Chr 7, ERATO Doi 193, expressed"	D7Ertd193e	
1447546_s_at	1.32247	Mastermind like 2 (Drosophila)	Maml2	
1447547_at	0.96647	latent transforming growth factor beta binding protein 1	Ltbp1	
1447548_at	0.92892	abhydrolase domain containing 5	Abhd5	
1447549_x_at	1.04278	ninjurin 1	Ninj1	
1447550_at	1.09593	hypothetical protein LOC666892	LOC666892	LOC671332
1447551_x_at	0.85047	latrophilin 3	Lphn3	
1447552_s_at	1.17620	"Catenin (cadherin associated protein), delta 2"	Ctnnd2	
1447553_x_at	1.03488	Resistance to inhibitors of cholinesterase 3 homolog (C. elegans)	Ric3	
1447554_at	1.00046	cDNA sequence BC031781	BC031781	
1447555_at	1.80876	Utrophin	Utnr1	
1447556_x_at	1.09623	RIKEN cDNA 1700094D03 gene	1700094D03Rik	
1447557_at	0.91254	RAB interacting factor	Rabif	
1447558_at	0.91632	---	---	
1447559_at	1.10810	expressed sequence AI834762	AI834762	
1447560_at	0.83517	---	---	
1447561_at	0.92407	Transcribed locus	---	
1447562_at	0.94578	forkhead box F2	Foxf2	
1447563_at	0.91205	ribosomal protein S3	Rps3	
1447564_x_at	1.02437	piwi-like homolog 4 (Drosophila)	Piwil4	
1447565_at	0.94638	RIKEN cDNA 1700082C02 gene	1700082C02Rik	
1447566_at	0.87366	Histone deacetylase 4	Hdac4	
1447567_at	1.26236	RIKEN cDNA D130007H15 gene	D130007H15Rik	
1447568_at	1.35506	RIKEN cDNA 2010003O18 gene	2010003O18Rik	
1447569_at	0.93130	RIKEN cDNA 1700081L11 gene	1700081L11Rik	
1447570_s_at	1.06807	A kinase (PRKA) anchor protein 10	Akap10	
1447571_at	1.03337	"ELK4, member of ETS oncogene family"	Elk4	
1447572_at	1.03782	RIKEN cDNA 1700052I22 gene	1700052I22Rik	
1447573_at	0.89110	"Solute carrier family 32 (GABA vesicular transporter), member 1"	Slc32a1	
1447574_s_at	1.04745	"Solute carrier family 32 (GABA vesicular transporter), member 1"	Slc32a1	
1447575_at	0.95775	---	---	
1447576_at	0.82775	RIKEN cDNA 2010001K21 gene	2010001K21Rik	
1447577_x_at	0.90458	"propionyl-Coenzyme A carboxylase, alpha polypeptide"	Pcca	
1447578_at	0.95195	RIKEN cDNA 1700084J12 gene	1700084J12Rik	
1447579_at	1.01208	Reticulon 4 interacting protein 1	Rtn4ip1	
1447580_at	1.01448	Transcribed locus	---	
1447581_at	0.92747	RIKEN cDNA 3110050K21 gene	3110050K21Rik	
1447582_x_at	0.93375	"Adiponectin, C1Q and collagen domain containing"	Adipoq	
1447583_x_at	0.83793	G protein-coupled receptor 30	Gpr30	

1447584_s_at 1.01511 myc target 1 Myct1
1447585_s_at 1.06398 mitochondrial ribosomal protein S6 Mrps6
1447586_at 0.91229 chondroitin sulfate proteoglycan 2 Cspg2
1447587_at 0.82137 RIKEN cDNA G630024C07 gene G630024C07Rik
1447588_x_at 0.99691 RIKEN cDNA G630024C07 gene G630024C07Rik
1447589_at 0.95192 "Protein phosphatase 4, regulatory subunit 1" Ppp4r1
1447590_at 0.97056 "Adult male bone cDNA, RIKEN full-length enriched library, clone:9830167D18 product:unclassifiable, full insert sequence" ---
1447591_x_at 0.99852 RIKEN cDNA A830082N09 gene A830082N09Rik
1447592_at 0.94465 dopamine beta hydroxylase Dbh
1447593_x_at 0.96691 "guanine nucleotide binding protein, alpha q polypeptide" Gnaq
1447594_at 0.96706 RIKEN cDNA 1810012K16 gene 1810012K16Rik
1447595_x_at 0.93388 RIKEN cDNA 1810012K16 gene 1810012K16Rik
1447596_at 0.93133 Transcribed locus ---
1447597_at 1.05173 RNA polymerase 1-4 Rpo1-4
1447598_x_at 1.26481 RNA polymerase 1-4 Rpo1-4
1447599_x_at 1.00697 epsin 2 Epn2
1447600_at 1.05584 Tetraspanin 5 Tspan5
1447601_x_at 1.01319 N-acetyltransferase 10 Nat10
1447602_x_at 0.79863 sulfatase 2 Sulf2
1447603_x_at 0.92615 RIKEN cDNA 1700027J05 gene 1700027J05Rik
1447604_at 1.06894 cDNA sequence BC053393 BC053393
1447605_at 0.99948 Zinc finger protein 457 Zfp457
1447606_x_at 1.10036 aquaporin 11 Aqp11
1447607_at 0.94264 --- ---
1447608_x_at 0.92591 BTB (POZ) domain containing 14A Btbd14a
1447609_at 1.08756 RIKEN cDNA 5730410E15 gene 5730410E15Rik
1447610_at 0.91034 transmembrane protein 34 Tmem34
1447611_at 0.98036 RIKEN cDNA 1700073E17 gene 1700073E17Rik
1447612_x_at 0.94083 Jumonji domain containing 3 Jmjd3
1447613_at 0.97507 Guanosine diphosphate (GDP) dissociation inhibitor 1 Gdi1
1447614_at 0.96636 Schlafen 1 Slfn1
1447615_at 1.16073 Formin 1 Fmn1
1447616_at 0.97766 Transcribed locus ---
1447617_at 1.04981 cyclin-dependent kinase 2 Cdk2
1447618_at 1.03921 RIKEN cDNA B630019K06 gene B630019K06Rik
1447619_at 0.94882 RIKEN cDNA A830039H05 gene A830039H05Rik
1447620_at 0.95670 cappuccino Cno
1447621_s_at 1.17192 RIKEN cDNA 2610307O08 gene 2610307O08Rik
1447622_at 0.98344 "Fc receptor, IgG, low affinity IIb" Fcgr2b
1447623_s_at 0.87856 RIKEN cDNA C130009A20 gene C130009A20Rik
1447624_s_at 0.90611 storkhead box 2 Stox2
1447625_at 0.90528 E2F transcription factor 5 E2f5

1447626_x_at 0.96862 AT rich interactive domain 3B (Bright like) Arid3b
1447627_at 0.88845 transmembrane protein 132E Tmem132e
1447628_x_at 0.92497 mitochondrial ribosomal protein S5 Mrps5
1447629_at 1.03837 RIKEN cDNA 1810020G14 gene 1810020G14Rik
1447630_x_at 0.94310 "guanine nucleotide binding protein, beta 5" Gnb5
1447631_at 2.16559 MYST histone acetyltransferase 2 Myst2
1447632_at 0.97536 "Fanconi anemia, complementation group C" Fance
1447633_x_at 1.03813 Ubiquitin-conjugating enzyme E2L 6 Ube2l6
1447634_x_at 0.81609 regulator of chromosome condensation (RCC1) and BTB (POZ) domain
containing protein 2 Rcbtb2
1447635_at 1.03606 "protein kinase, cAMP dependent regulatory, type I, alpha" Prkar1a
1447636_x_at 0.93785 transmembrane and coiled-coil domains 4 Tmco4
1447637_at 1.14274 RIKEN cDNA 2410018C17 gene 2410018C17Rik
1447638_at 1.01891 Arylformamidase Afmid
1447639_x_at 1.07832 --- ---
1447640_s_at 0.98454 pre B-cell leukemia transcription factor 3 Pbx3
1447641_at 0.92372 dystrophia myotonica-containing WD repeat motif Dmwd
1447642_x_at 0.95296 dystrophia myotonica-containing WD repeat motif Dmwd
1447643_x_at 1.07529 snail homolog 2 (Drosophila) Snai2
1447644_at 1.09002 Wingless-related MMTV integration site 5B Wnt5b
1447645_x_at 1.01456 "potassium channel, subfamily K, member 13" Kcnk13
1447646_at 0.99928 HEAT repeat containing 2 Heatr2
1447647_at 1.00686 wingless-related MMTV integration site 7A Wnt7a
1447648_at 1.07931 RIKEN cDNA 1700021N20 gene 1700021N20Rik
1447649_x_at 1.20702 "DnaJ (Hsp40) homolog, subfamily C, member 1" Dnajc1
1447650_at 1.00531 NMD3 homolog (S. cerevisiae) Nmd3
1447651_x_at 1.03702 mitochondrial ribosomal protein L30 /// similar to mitochondrial ribosomal
protein L30 /// similar to mitochondrial ribosomal protein L30 Mrpl30 /// LOC621542 /// LOC632468
1447652_at 1.02006 "solute carrier family 35, member A4 /// similar to solute carrier family 35,
member A4" Slc35a4 /// LOC545750
1447653_x_at 1.08285 ribosomal protein L24 /// similar to ribosomal protein L24 /// similar to
ribosomal protein L24 /// similar to ribosomal protein L24 /// similar to ribosomal protein L24 /// similar
to ribosomal protein L24 /// similar to ribosomal protein L24 /// similar to ribosomal protein L24
Rpl24 /// LOC624993 /// LOC664869 /// LOC667996 /// LOC668829 /// LOC672138 /// LOC675789 ///
LOC677192
1447654_at 0.95643 --- ---
1447655_x_at 0.83669 SRY-box containing gene 6 Sox6
1447656_at 0.93904 "zinc finger, DHHC domain containing 17" Zdhhc17
1447657_s_at 0.87670 synaptopodin 2-like Synpo2l
1447658_x_at 0.98609 synaptopodin 2-like Synpo2l
1447659_x_at 1.04103 "ATPase, H+ transporting, lysosomal V1 subunit H" Atp6v1h
1447660_at 0.83824 transducin (beta)-like 3 Tbl3
1447661_at 1.04485 Transcribed locus ---

1447662_x_at 0.92388 "DNA segment, Chr 18, ERATO Doi 653, expressed" D18Ert653e
 1447663_at 1.00422 A kinase (PRKA) anchor protein 13 Akap13
 1447664_x_at 0.96205 expressed sequence AA986860 AA986860
 1447665_at 1.03789 leucine rich repeat neuronal 6C Lrrn6c
 1447666_x_at 0.94525 similar to Protein C11orf51 /// similar to Protein HSPC020 LOC628119 ///
 LOC638436
 1447667_x_at 0.92421 mitogen activated protein kinase kinase kinase 4 Map3k4
 1447668_x_at 1.04222 epidermal growth factor-containing fibulin-like extracellular matrix protein 2
 Efemp2
 1447669_s_at 1.11205 "guanine nucleotide binding protein (G protein), gamma 4 subunit" Gng4
 1447670_at 1.21070 "proteasome (prosome, macropain) 26S subunit, non-ATPase, 9" Psmd9
 1447671_x_at 1.04685 F-box only protein 6b Fbxo6b
 1447672_x_at 0.95153 protein serine kinase H1 Pskh1
 1447673_x_at 1.07142 RIKEN cDNA 1700015C15 gene 1700015C15Rik
 1447674_at 0.98920 RIKEN cDNA 2900080J11 gene 2900080J11Rik
 1447675_x_at 1.02947 myogenic differentiation 1 Myod1
 1447676_x_at 0.87375 S100 calcium binding protein A16 S100a16
 1447677_x_at 0.91316 histidyl tRNA synthetase 2 Hars2
 1447678_at 0.98015 Dehydrogenase/reductase (SDR family) member 10 Dhrs10
 1447679_s_at 1.08328 "BMS1-like, ribosome assembly protein (yeast)" Bms1l
 1447680_at 1.00259 "Acidic (leucine-rich) nuclear phosphoprotein 32 family, member E" Anp32e
 1447681_x_at 0.91238 Pentatricopeptide repeat domain 2 Ptdc2
 1447682_x_at 0.92585 Tnf receptor-associated factor 5 /// similar to TNF receptor-associated factor
 5 Traf5 /// LOC622602
 1447683_x_at 1.22408 methyltransferase-like 1 Mettl1
 1447684_at 1.02009 Leucine zipper and CTNNBIP1 domain containing Lzic
 1447685_x_at 1.05115 "E26 avian leukemia oncogene 2, 3' domain" Ets2
 1447686_at 1.01132 --- ---
 1447687_x_at 1.00600 --- ---
 1447688_at 0.86417 "TBC1 domain family, member 20" Tbc1d20
 1447689_at 1.04663 G protein-coupled receptor associated sorting protein 1 Gprasp1
 1447690_at 0.99500 lipolysis stimulated lipoprotein receptor Lsr
 1447691_x_at 2.46425 Bicaudal C homolog 1 (Drosophila) Bicc1
 1447692_x_at 0.97667 mitogen-activated protein kinase kinase kinase 7 interacting protein 1
 Map3k7ip1
 1447693_s_at 0.80698 neogenin Neo1
 1447694_x_at 0.99013 neogenin Neo1
 1447695_at 0.92451 RIKEN cDNA 2610035D17 gene 2610035D17Rik
 1447696_x_at 0.95197 adenylate cyclase 5 Adcy5
 1447697_at 0.99888 Coiled-coil domain containing 40 Ccdc40
 1447698_x_at 0.91800 --- ---
 1447699_at 1.01756 Mitogen-activated protein kinase kinase kinase 13 Map3k13
 1447700_x_at 0.91044 synovial sarcoma translocation gene on chromosome 18-like 1 Ss18l1

1447701_x_at 0.83256 isocitrate dehydrogenase 3 (NAD+) alpha Idh3a
1447702_x_at 1.12735 "immunoglobulin superfamily, member 1" Igsf1
1447703_x_at 1.05257 zinc finger protein 593 Zfp593
1447704_s_at 1.11664 RIKEN cDNA D530033C11 gene D530033C11Rik
1447705_at 0.79290 RIKEN cDNA 4833432M17 gene 4833432M17Rik
1447706_at 1.80579 --- ---
1447707_s_at 0.90646 "phosphodiesterase 2A, cGMP-stimulated" Pde2a
1447708_x_at 0.94302 "phosphodiesterase 2A, cGMP-stimulated" Pde2a
1447709_at 0.99155 Erythrocyte protein band 4.1-like 3 Epb4.113
1447710_at 0.91104 RIKEN cDNA 1700019P21 gene 1700019P21Rik
1447711_x_at 0.97946 RIKEN cDNA 4933412E12 gene 4933412E12Rik
1447712_x_at 1.12427 E1A binding protein p400 Ep400
1447713_at 1.08407 "tropomyosin 1, alpha" Tpm1
1447714_x_at 0.93161 Transcribed locus ---
1447715_x_at 0.93251 "peroxiredoxin 6, related sequence 1" Prdx6-rs1
1447716_x_at 1.11850 RIKEN cDNA 6330442E10 gene 6330442E10Rik
1447717_x_at 0.94708 neurensin 2 Nrsn2
1447718_at 1.04098 "phosphodiesterase 4B, cAMP specific" Pde4b
1447719_at 0.94035 Apolipoprotein H Apoh
1447720_x_at 0.98531 "protein kinase, cAMP dependent, catalytic, alpha" Prkaca
1447721_at 1.04853 zinc finger and BTB domain containing 26 Zbtb26
1447722_at 0.97628 Transcribed locus ---
1447723_at 1.11088 "16 days neonate cerebellum cDNA, RIKEN full-length enriched library, clone:9630018J10 product:unclassifiable, full insert sequence" ---
1447724_x_at 1.07395 Optic atrophy 3 (human) Opa3
1447725_at 1.03750 RIKEN cDNA C030034E14 gene C030034E14Rik
1447726_at 0.96131 RIKEN cDNA C030002E08 gene C030002E08Rik
1447727_at 0.87960 Aquaporin 3 Aqp3
1447728_x_at 0.99325 heat shock protein 9A Hspa9a
1447729_s_at 1.27933 RIKEN cDNA 1700036D21 gene 1700036D21Rik
1447730_at 1.05091 Sp2 transcription factor Sp2
1447731_at 0.89524 paternally expressed 3 Peg3
1447732_x_at 1.04905 paternally expressed 3 Peg3
1447733_x_at 0.95072 --- ---
1447734_x_at 1.11799 "aldolase 1, A isoform /// similar to Fructose-bisphosphate aldolase A (Muscle-type aldolase) (Aldolase 1) /// similar to Fructose-bisphosphate aldolase A (Muscle-type aldolase) (Aldolase 1)" Aldoa /// LOC667483 /// LOC671023
1447735_x_at 0.93572 ataxin 2 binding protein 1 A2bp1
1447736_at 0.92131 FK506 binding protein 5 Fkbp5
1447737_at 0.86920 "protein kinase, interferon inducible double stranded RNA dependent activator" Prkra
1447738_s_at 0.85104 "ankyrin repeat domain 13 family, member D" Ankrd13d
1447739_x_at 0.92705 kelch domain containing 4 /// hypothetical gene supported by NM_145605

Klhdc4 /// LOC384360

1447740_at 1.14403 RIKEN cDNA 1700034J05 gene /// hypothetical protein LOC635035
1700034J05Rik /// LOC635035

1447741_x_at 0.98649 "acyl-Coenzyme A dehydrogenase family, member 10" Acad10
1447742_at 1.01071 lysosomal-associated protein transmembrane 5 Laptm5
1447743_x_at 0.80650 exostoses (multiple) 2 Ext2
1447744_s_at 1.05783 --- ---
1447745_at 0.88844 aquaporin 4 Aqp4
1447746_at 0.95381 Leucine-rich repeats and calponin homology (CH) domain containing 1 Lrch1
1447747_x_at 1.00636 Adenosine deaminase-like Adal
1447748_x_at 0.89250 "solute carrier family 29 (nucleoside transporters), member 2" Slc29a2
1447749_at 0.85417 SET and MYND domain containing 5 Smyd5
1447750_x_at 1.00591 NOP17 Nop17
1447751_x_at 0.88066 "dihydrouridine synthase 2-like (SMM1, S. cerevisiae)" Dus21
1447752_x_at 0.85293 developmentally regulated GTP binding protein 1 Drg1
1447753_at 0.92139 Cell division cycle 37 homolog (S. cerevisiae)-like 1 Cdc3711
1447754_x_at 1.04482 THAP domain containing 4 Thap4
1447755_at 1.25799 Serine/arginine-rich protein specific kinase 2 Srpk2
1447756_x_at 0.89865 "minichromosome maintenance deficient 6 (MIS5 homolog, S. pombe) (S. cerevisiae)" Mcm6
1447757_x_at 0.84398 inositol polyphosphate-5-phosphatase F Inpp5f
1447758_x_at 1.05418 MYST histone acetyltransferase monocytic leukemia 4 Myst4
1447759_x_at 0.91068 coiled-coil domain containing 22 Ccdc22
1447760_x_at 0.91183 ets homologous factor Ehf
1447761_x_at 0.95540 "superoxide dismutase 1, soluble /// similar to Superoxide dismutase /// similar to Superoxide dismutase" Sod1 /// LOC545845 /// LOC667310
1447762_x_at 1.02313 carbonic anhydrase 12 Car12
1447763_at 1.04924 --- ---
1447764_at 0.97454 "potassium voltage-gated channel, Shal-related family, member 2" Kcnd2
1447765_at 0.84960 WD repeat domain 48 Wdr48
1447766_x_at 0.84267 LIM domain containing 2 /// similar to epithelial protein lost in neoplasm
Limd2 /// LOC632329

1447767_at 0.96145 Cyclin-dependent kinase-like 4 Cdk14
1447768_at 0.95609 "glutamate oxaloacetate transaminase 2, mitochondrial /// similar to Aspartate aminotransferase, mitochondrial precursor (Transaminase A) (Glutamate oxaloacetate transaminase 2)" Got2 /// LOC640847
1447769_x_at 1.07185 adhesion molecule with Ig like domain 2 Amigo2
1447770_at 0.98974 calcium and integrin binding family member 2 Cib2
1447771_at 0.92860 --- ---
1447772_at 0.96266 "pleckstrin homology domain-containing, family A (phosphoinositide binding specific) member 3" Plekha3
1447773_x_at 0.87812 RIKEN cDNA 6330409D20 gene 6330409D20Rik
1447774_x_at 0.91980 RIKEN cDNA 5730469M10 gene 5730469M10Rik

1447775_x_at 1.06599 zinc finger protein 622 Zfp622
1447776_x_at 1.09987 "RAB6, member RAS oncogene family" Rab6
1447777_x_at 1.06652 --- ---
1447778_x_at 0.88927 "BRCA1/BRCA2-containing complex, subunit 3" Brc3
1447779_x_at 0.93056 HLA-B-associated transcript 1A Bat1a
1447780_x_at 1.00989 "Tu translation elongation factor, mitochondrial /// similar to Tu translation elongation factor, mitochondrial" Tufm /// LOC669832
1447781_s_at 1.10570 RNA (guanine-9-) methyltransferase domain containing 2 Rg9mtd2
1447782_x_at 1.10615 transducin (beta)-like 3 Tbl3
1447783_x_at 0.80753 "DNA segment, Chr 11, ERATO Doi 333, expressed" D11Ertd333e
1447784_x_at 1.05609 RIKEN cDNA 9130023D20 gene 9130023D20Rik
1447785_x_at 0.93029 "DNA segment, Chr 11, Wayne State University 99, expressed" D11Wsu99e
1447786_at 1.02165 "pleckstrin homology, Sec7 and coiled-coil domains 1" Pscd1
1447787_x_at 0.96830 gap junction membrane channel protein alpha 7 Gja7
1447788_s_at 0.90682 TSPY-like 3 Tspy13
1447789_x_at 0.92958 --- ---
1447790_at 0.95646 --- ---
1447791_s_at 0.93802 "guanine nucleotide binding protein, alpha 14" Gna14
1447792_x_at 1.03971 G protein-coupled receptor 174 Gpr174
1447793_x_at 0.98881 Ring-box 1 Rbx1
1447794_x_at 1.04410 PLAC8-like 1 Plac8l1
1447795_at 0.96852 B-box and SPRY domain containing Bspry
1447796_at 0.93170 NOP17 Nop17
1447797_x_at 0.90767 NOP17 Nop17
1447798_at 1.01997 RIKEN cDNA 1700123O21 gene 1700123O21Rik
1447799_x_at 0.99843 RIKEN cDNA 1700001P01 gene 1700001P01Rik
1447800_x_at 0.68229 Transcobalamin 2 Tcn2
1447801_x_at 0.98989 --- ---
1447802_x_at 1.03733 --- ---
1447803_x_at 1.03482 "Capping protein (actin filament), gelsolin-like" Capg
1447804_x_at 0.82916 CDNA sequence BC031781 BC031781
1447805_s_at 1.03849 "DNA segment, Chr 11, ERATO Doi 730, expressed" D11Ertd730e
1447806_s_at 1.07314 serine/arginine-rich protein specific kinase 3 Srp3
1447807_s_at 1.15205 "pleckstrin homology domain containing, family H (with MyTH4 domain) member 1" Plekhh1
1447808_s_at 0.96891 "solute carrier family 15 (H+/peptide transporter), member 2" Slc15a2
1447809_x_at 0.90396 --- ---
1447810_x_at 1.02114 "pleckstrin homology domain containing, family B (evectins) member 2" Plekhh2
1447811_s_at 0.75705 adhesion molecule with Ig like domain 1 Amigo1
1447812_x_at 1.02418 "filamin C, gamma (actin binding protein 280)" Flnc
1447813_x_at 0.94287 src-like adaptor Sla
1447814_x_at 1.11097 interleukin 16 Il16

1447815_x_at	1.04948	RIKEN cDNA 6430527G18 gene	6430527G18Rik
1447816_x_at	0.90823	oxidoreductase NAD-binding domain containing 1	Oxnad1
1447817_at	1.03909	"hypoxia-inducible factor 1, alpha subunit inhibitor"	Hif1an
1447818_x_at	0.98398	Ras homolog enriched in brain like 1	Rheb11
1447819_x_at	1.05288	"procollagen, type VIII, alpha 1"	Col8a1
1447820_x_at	1.02960	carnitine palmitoyltransferase 2	Cpt2
1447821_at	1.01423	RIKEN cDNA 1700051K22 gene	1700051K22Rik
1447822_x_at	1.07650	RIKEN cDNA 2700038N03 gene	2700038N03Rik
1447823_x_at	1.02275	mitogen-activated protein kinase 12	Mapk12
1447824_x_at	0.92066	heat shock 70kD protein 5 (glucose-regulated protein)	Hspa5
1447825_x_at	0.96110	protocadherin 8	Pcdh8
1447826_x_at	1.00248	RIKEN cDNA A930037G23 gene	A930037G23Rik
1447827_x_at	1.13279	speckle-type POZ protein	Spop
1447828_x_at	0.96278	RIKEN cDNA 2310005P05 gene	2310005P05Rik
1447829_x_at	0.94663	"TAF9B RNA polymerase II, TATA box binding protein (TBP)-associated factor"	Taf9b
1447830_s_at	0.95006	regulator of G-protein signaling 2	Rgs2
1447831_s_at	1.03393	myotubularin related protein 7	Mtmr7
1447832_x_at	1.13622	Ankyrin repeat domain 10	Ankrd10
1447833_x_at	1.05768	microfibrillar-associated protein 2	Mfap2
1447834_at	0.90591	"Coatamer protein complex, subunit zeta 1"	Copz1
1447835_at	0.97415	RIKEN cDNA 2610528K11 gene	2610528K11Rik
1447836_x_at	1.12978	RIKEN cDNA 4921525O09 gene	4921525O09Rik
1447837_x_at	0.93264	---	---
1447838_x_at	0.82745	echinoderm microtubule associated protein like 4	Eml4
1447839_x_at	1.06466	adrenomedullin	Adm
1447840_x_at	1.01027	E2F transcription factor 1	E2f1
1447841_x_at	0.94478	"ST3 beta-galactoside alpha-2,3-sialyltransferase 6"	St3gal6
1447842_x_at	0.94293	Transcobalamin 2	Tcn2
1447843_at	1.05332	"proteasome (prosome, macropain) 26S subunit, non-ATPase, 13"	Psmd13
1447844_at	1.03210	Transcribed locus	---
1447845_s_at	1.17371	vanin 1	Vnn1
1447846_x_at	1.13747	leukocyte receptor cluster (LRC) member 4	Leng4
1447847_x_at	0.86544	---	---
1447848_at	0.98568	epidermal growth factor receptor pathway substrate 15-like 1	Eps15l1
1447849_s_at	0.91489	avian musculoaponeurotic fibrosarcoma (v-maf) AS42 oncogene homolog	Maf
1447850_x_at	0.92065	"zinc finger, AN1-type domain 3"	Zfand3
1447851_x_at	0.85963	---	---
1447852_x_at	0.95176	RIKEN cDNA 2900002H16 gene	2900002H16Rik
1447853_x_at	0.95619	Kinesin family member 13A	Kif13a
1447854_s_at	0.90110	"histone 2, H2be"	Hist2h2be
1447855_x_at	1.01450	amyloid beta precursor protein (cytoplasmic tail) binding protein 2	Appbp2

1447856_x_at 0.96497 "solute carrier family 25 (mitochondrial carrier, phosphate carrier), member 25" Slc25a25

1447857_at 1.01694 --- ---

1447858_x_at 0.91553 "interleukin 4 receptor, alpha" Il4ra

1447859_at 1.03011 aarF domain containing kinase 2 Adck2

1447860_x_at 0.90317 component of oligomeric golgi complex 8 Cog8

1447861_x_at 1.12800 Myeloid ecotropic viral integration site-related gene 1 Mrg1

1447862_x_at 1.12435 thrombospondin 2 Thbs2

1447863_s_at 0.97127 "nuclear receptor subfamily 4, group A, member 2" Nr4a2

1447864_s_at 0.91680 pogo transposable element with KRAB domain Pogk

1447865_x_at 1.03886 PDZ domain containing 11 Pdzd11

1447866_x_at 0.95005 RIKEN cDNA 2300009A05 gene 2300009A05Rik

1447867_x_at 0.99707 RIKEN cDNA 4930444A02 gene 4930444A02Rik

1447868_x_at 0.90568 thioredoxin-like 2 /// similar to thioredoxin-like 2 Txnl2 /// LOC620016

1447869_x_at 0.95142 Rho-related BTB domain containing 3 Rhobtb3

1447870_x_at 1.76541 RIKEN cDNA 1110002E22 gene 1110002E22Rik

1447871_at 1.02439 Transcribed locus ---

1447872_at 1.03711 Transcribed locus ---

1447873_x_at 1.07703 BH3 interacting domain death agonist Bid

1447874_x_at 0.86674 "sphingomyelin phosphodiesterase 1, acid lysosomal" Smpd1

1447875_x_at 1.05791 EST AA407452 AA407452

1447876_x_at 0.75978 RIKEN cDNA 2310073E15 gene 2310073E15Rik

1447877_x_at 1.00092 DNA methyltransferase (cytosine-5) 1 Dnmt1

1447878_s_at 0.84856 fibroblast growth factor receptor-like 1 Fgfr1

1447879_x_at 0.94123 DALR anticodon binding domain containing 3 /// WD repeat domain 6 Dalrd3 /// Wdr6

1447880_x_at 1.03147 "Endothelial differentiation, sphingolipid G-protein-coupled receptor, 5" Edg5

1447881_x_at 0.83054 --- ---

1447882_x_at 0.97287 DEAD (Asp-Glu-Ala-Asp) box polypeptide 54 Ddx54

1447883_x_at 0.80284 microtubule-associated protein 1 light chain 3 alpha Map1lc3a

1447884_x_at 1.07116 RIKEN cDNA 1300010M03 gene 1300010M03Rik

1447885_x_at 0.87076 "neural precursor cell expressed, developmentally down-regulated gene 9" Nedd9

1447886_at 0.99973 RIKEN cDNA 0610040B09 gene 0610040B09Rik

1447887_x_at 1.04060 chondroitin sulfate proteoglycan 2 Cspg2

1447888_x_at 0.83133 mutS homolog 2 (E. coli) Msh2

1447889_x_at 1.09313 "gene model 50, (NCBI)" Gm50

1447890_at 0.97807 reticulocalbin 1 Rcn1

1447891_at 0.91631 "Transforming, acidic coiled-coil containing protein 2" Tacc2

1447892_at 1.08574 --- ---

1447893_x_at 0.90614 "solute carrier family 39 (metal ion transporter), member 6" Slc39a6

1447894_x_at 0.95961 vacuolar protein sorting 52 (yeast) /// similar to vacuolar protein sorting 52

Vps52 /// LOC631721

1447895_x_at 0.90437 PAN3 polyA specific ribonuclease subunit homolog (*S. cerevisiae*) Pan3
 1447896_s_at 0.92662 RIKEN cDNA 2010109N14 gene 2010109N14Rik
 1447897_x_at 0.90143 Anaphase promoting complex subunit 11 homolog (yeast) Anapc11
 1447898_s_at 0.93371 "splicing factor, arginine/serine-rich 6" Sfrs6
 1447899_x_at 0.90787 tumor-associated calcium signal transducer 1 Tacstd1
 1447900_x_at 0.98528 ectonucleoside triphosphate diphosphohydrolase 4 Entpd4
 1447901_x_at 0.78836 "Sfi1 homolog, spindle assembly associated (yeast) /// similar to spindle assembly associated Sfi1 homolog isoform b /// similar to Sfi1 homolog, spindle assembly associated (yeast)" Sfi1 /// LOC673420 /// LOC673566
 1447902_at 0.94457 RIKEN cDNA 1810013A23 gene 1810013A23Rik
 1447903_x_at 0.96675 "adaptor-related protein complex 1, sigma 2 subunit" Ap1s2
 1447904_s_at 0.94504 "farnesyltransferase, CAAX box, alpha" Fnta
 1447905_x_at 0.97592 nucleoporin 62 Nup62
 1447906_at 0.89832 "Protein tyrosine phosphatase, receptor type, G" Ptpg
 1447907_x_at 0.99648 "Amyloid beta (A4) precursor protein-binding, family A, member 1 binding protein" Apba2bp
 1447908_x_at 0.99335 tetratricopeptide repeat domain 3 Ttc3
 1447909_s_at 1.05647 selenoprotein K Selk
 1447910_x_at 1.01030 selenoprotein K Selk
 1447911_at 1.15911 TEA domain family member 1 Tead1
 1447912_x_at 0.91738 A kinase (PRKA) anchor protein (yotiao) 9 Akap9
 1447913_x_at 1.12874 A kinase (PRKA) anchor protein (yotiao) 9 Akap9
 1447914_x_at 1.07087 RIKEN cDNA 4833411O04 gene 4833411O04Rik
 1447915_x_at 0.93363 cDNA sequence BC054438 BC054438
 1447916_at 1.02563 --- ---
 1447917_x_at 0.98945 N-terminal Asn amidase Ntan1
 1447918_x_at 0.96350 hypothetical protein LOC207685 /// similar to Ig lambda-3 chain C region - mouse LOC207685 /// LOC386520
 1447919_x_at 1.02704 "NADH dehydrogenase (ubiquinone) 1, alpha/beta subcomplex, 1" Ndufab1
 1447920_at 0.98447 Ataxin 10 Atxn10
 1447921_at 0.96045 Transcribed locus ---
 1447922_at 0.95974 Syntaxin binding protein 5-like Stxbp5l
 1447923_at 0.88283 RIKEN cDNA 2810013C04 gene 2810013C04Rik
 1447924_at 1.03202 nuclear casein kinase and cyclin-dependent kinase substrate 1 Nucks1
 1447925_at 1.10011 RIKEN cDNA D630033A02 gene D630033A02Rik
 1447926_at 0.80957 ADP-ribosylation factor-like 5A Arl5a
 1447927_at 1.85134 macrophage activation 2 like /// similar to macrophage activation 2 like /// similar to macrophage activation 2 like Mpa2l /// LOC626578 /// LOC673101
 1447928_at 1.07653 "carbonic anhydrase 5b, mitochondrial" Car5b
 1447929_at 1.04147 slingshot homolog 3 (*Drosophila*) Ssh3
 1447930_at 1.36250 bromodomain adjacent to zinc finger domain 1A Baz1a
 1447931_at 1.08920 Wolf-Hirschhorn syndrome candidate 1-like 1 (human) Whsc11l

1447932_at 1.02106 coiled-coil domain containing 16 Ccdc16
1447933_at 0.89763 kinesin family member 26A /// similar to CG14535-PA /// similar to CG14535-PA Kif26a /// LOC668303 /// LOC676226
1447934_at 1.21312 RIKEN cDNA 9630033F20 gene 9630033F20Rik
1447935_at 0.89798 "Fanconi anemia, complementation group M" Fancm
1447936_at 1.13535 RIKEN cDNA 2410006H16 gene 2410006H16Rik
1447937_a_at 0.88992 RIKEN cDNA 4933409K07 gene /// similar to 4933409K07Rik protein /// similar to 4933409K07Rik protein /// hypothetical protein LOC622738 /// similar to similar to 4933409K07Rik protein /// hypothetical protein LOC665845 4933409K07Rik /// LOC545605 /// LOC545614 /// LOC622738 /// LOC665744 /// LOC665845
1447938_at 1.00872 similar to 4933409K07Rik protein LOC545614
1447939_a_at 0.87475 RIKEN cDNA 4933409K07 gene /// similar to 4933409K07Rik protein /// hypothetical protein LOC665845 4933409K07Rik /// LOC545605 /// LOC665845
1447940_a_at 1.16360 Braf transforming gene Braf
1447941_x_at 1.00042 Braf transforming gene Braf
1447942_x_at 1.19182 Neuraminidase 1 Neu1
1447943_x_at 1.08344 YEATS domain containing 2 Yeats2
1447944_at 0.81944 zinc finger with KRAB and SCAN domains 1 Zkscan1
1447945_at 1.00436 avian musculoaponeurotic fibrosarcoma (v-maf) AS42 oncogene homolog Maf
1447946_at 1.06629 a disintegrin and metallopeptidase domain 23 Adam23
1447947_at 0.94211 "zinc finger, FYVE domain containing 16" Zfyve16
1447948_at 0.97480 RIKEN cDNA A430107O13 gene A430107O13Rik
1447949_at 1.09318 --- ---
1447950_at 1.06390 RIKEN cDNA A730011C13 gene A730011C13Rik
1447951_at 1.50347 RIKEN cDNA 5730403B10 gene 5730403B10Rik
1447952_at 0.98244 AU RNA binding protein/enoyl-coenzyme A hydratase Auh
1447953_at 0.84313 EGF-like domain 8 Egfl8
1447954_at 0.89757 leucine rich repeat containing 49 Lrrc49
1447955_at 1.07043 "similar to heterochromatin protein 1, binding protein 3" LOC670127
1447956_at 0.97271 expressed sequence C76614 C76614
1447957_at 1.15239 --- ---
1447958_at 0.99013 "16 days neonate thymus cDNA, RIKEN full-length enriched library, clone: A130082C12 product:unclassifiable, full insert sequence" ---
1447959_at 0.85393 zinc finger protein 398 Zfp398
1447960_at 1.07377 hypothetical protein 9630017O17 9630017O17
1447961_s_at 1.13675 mitochondrial ribosomal protein L38 Mrpl38
1447962_at 1.00610 "PREDICTED: Mus musculus hypothetical protein LOC626765 (LOC626765), mRNA" ---
1447963_at 0.89421 --- ---
1447964_at 1.12306 tubulin tyrosine ligase Ttl
1447965_at 1.09730 expressed sequence C80012 C80012
1447966_a_at 1.08978 transmembrane protein 69 Tmem69

1447967_at 1.04022 transmembrane protein 69 Tmem69
1447968_at 0.93701 RIKEN cDNA 9330158F14 gene 9330158F14Rik
1447969_at 0.91615 minichromosome maintenance deficient domain containing 1 Mcmdc1
1447970_at 1.34983 --- ---
1447971_at 0.88219 RIKEN cDNA A230061C15 gene A230061C15Rik
1447972_at 1.00624 Transcribed locus ---
1447973_at 0.94613 thyroid hormone receptor interactor 11 Trip11
1447974_s_at 0.98260 aquaporin 6 Aqp6
1447975_a_at 0.98048 hypothetical protein LOC545261 LOC545261
1447976_at 0.97535 expressed sequence AI596198 AI596198
1447977_x_at 0.96433 RIKEN cDNA 3100002L24 gene /// RIKEN cDNA 9230108I15 gene ///
similar to gonadotropin inducible ovarian transcription factor 1 /// similar to zinc finger protein 709 ///
similar to zinc finger protein 709 /// similar to zinc finger protein 709 /// similar to zinc finger protein
709 /// similar to gonadotropin inducible ovarian transcription factor 1 /// similar to zinc finger protein
97 /// similar to gonadotropin inducible ovarian transcription factor 1 3100002L24Rik ///
9230108I15Rik /// LOC626832 /// LOC627914 /// LOC628147 /// LOC665043 /// LOC665078 ///
LOC668030 /// LOC668039 /// LOC668066
1447978_at 0.96875 expressed sequence C80719 C80719
1447979_at 0.90414 RUN and TBC1 domain containing 2 Rutbc2
1447980_s_at 0.79538 RUN and TBC1 domain containing 2 Rutbc2
1447981_at 0.97449 expressed sequence C78441 C78441
1447982_at 1.06719 RIKEN cDNA 1110008P14 gene 1110008P14Rik
1447983_at 1.02029 "Zinc finger, matrin-like" Zfml
1447984_at 1.19595 "DNA segment, Chr 1, ERATO Doi 75, expressed" D1Ertd75e
1447985_s_at 1.03817 ankyrin repeat and IBR domain containing 1 Ankib1
1447986_at 0.97897 expressed sequence D17892 D17892
1447987_at 0.80373 RIKEN cDNA D030041N04 gene D030041N04Rik
1447988_at 0.99780 --- ---
1447989_at 1.09240 "Transcribed locus, weakly similar to XP_576435.1 PREDICTED: similar to
ORF4 [Rattus norvegicus]" ---
1447990_at 0.94657 expressed sequence C76332 C76332
1447991_at 1.04564 proprotein convertase subtilisin/kexin type 2 Pcsk2
1447992_s_at 0.89784 proprotein convertase subtilisin/kexin type 2 Pcsk2
1447993_a_at 0.95336 RIKEN cDNA 1700026B20 gene 1700026B20Rik
1447994_at 0.96828 RIKEN cDNA 1700026B20 gene 1700026B20Rik
1447995_x_at 0.98167 RIKEN cDNA 1700026B20 gene 1700026B20Rik
1447996_at 1.18945 expressed sequence AI848149 AI848149
1447997_s_at 1.11384 translocase of inner mitochondrial membrane 8 homolog a2 (yeast)
Timm8a2
1447998_at 1.02835 expressed sequence AI324046 AI324046
1447999_x_at 1.16279 similar to glyceraldehyde-3-phosphate dehydrogenase /// similar to
glyceraldehyde-3-phosphate dehydrogenase /// cDNA sequence AK190093 /// similar to Glyceraldehyde-
3-phosphate dehydrogenase (GAPDH) (38 kDa BFA-dependent ADP-ribosylation substrate) (BARS-

38) /// glyceraldehyde-3-phosphate dehydrogenase /// similar to Glyceraldehyde-3-phosphate dehydrogenase (GAPDH) /// similar to Glyceraldehyde-3-phosphate dehydrogenase (GAPDH) /// similar to Glyceraldehyde-3-phosphate dehydrogenase (GAPDH) /// similar to Glyceraldehyde-3-phosphate dehydrogenase (GAPDH) /// similar to Glyceraldehyde-3-phosphate dehydrogenase (GAPDH) /// similar to Glyceraldehyde-3-phosphate dehydrogenase (GAPDH) /// similar to Glyceraldehyde-3-phosphate dehydrogenase (GAPDH) /// similar to Glyceraldehyde-3-phosphate dehydrogenase (GAPDH) /// similar to Glyceraldehyde-3-phosphate dehydrogenase (GAPDH) /// similar to Glyceraldehyde-3-phosphate dehydrogenase (GAPDH) /// similar to Glyceraldehyde-3-phosphate dehydrogenase (GAPDH) /// similar to Glyceraldehyde-3-phosphate dehydrogenase (GAPDH) /// hypothetical protein LOC671520 /// similar to Glyceraldehyde-3-phosphate dehydrogenase (GAPDH) /// similar to Glyceraldehyde-3-phosphate dehydrogenase (GAPDH) /// similar to Glyceraldehyde-3-phosphate dehydrogenase (GAPDH) LOC14433 /// LOC380687 /// AK190093 /// LOC384808 /// Gapdh /// LOC432919 /// LOC545984 /// LOC622339 /// LOC630967 /// LOC632720 /// LOC639817 /// LOC665036 /// LOC667048 /// LOC667806 /// LOC670367 /// LOC671520 /// LOC671913 /// LOC672238 /// LOC676700

1448000_at	0.94639	cell division cycle associated 3	Cdca3
1448001_x_at	0.91878	cell division cycle associated 3	Cdca3
1448002_x_at	1.03038	RIKEN cDNA 2610001J05 gene	2610001J05Rik
1448003_at	0.93847	---	---
1448005_at	0.95569	SAM and SH3 domain containing 1	Sash1
1448006_at	1.05581	zinc finger CCCH type containing 7B	Zc3h7b
1448007_at	0.85207	"bromodomain adjacent to zinc finger domain, 2B"	Baz2b
1448008_at	0.98133	ankyrin repeat and KH domain containing 1	Ankhd1
1448009_at	1.09224	UDP-glucose ceramide glucosyltransferase-like 1	Ugcgl1
1448010_at	1.06197	RIKEN cDNA A430104N18 gene	A430104N18Rik
1448011_at	0.93967	---	---
1448012_at	1.08372	expressed sequence C76336	C76336
1448013_at	1.00894	ubiquitin specific peptidase 24	Usp24
1448014_s_at	1.01956	ubiquitin specific peptidase 24	Usp24
1448015_at	0.92989	RIKEN cDNA 2310047C04 gene	2310047C04Rik
1448016_at	1.12487	spindle assembly 6 homolog (C. elegans)	Sass6
1448017_at	1.00961	ubiquitin associated protein 2-like	Ubp2l
1448018_at	1.39650	---	---
1448019_at	1.15435	RIKEN cDNA 2900006A08 gene	2900006A08Rik
1448020_at	1.00421	RAS-related protein-1a	Rap1a
1448021_at	0.62448	Transcribed locus	---
1448022_at	1.00016	Transcribed locus	---
1448023_at	1.02980	"kalirin, RhoGEF kinase"	Kalrn
1448024_at	1.00952	natriuretic peptide receptor 3	Npr3
1448025_at	0.92428	similar to SIRP beta 1 isoform 2	LOC668101
1448026_at	0.97983	---	---
1448027_at	1.05438	RIKEN cDNA 2010305B15 gene	2010305B15Rik
1448028_at	1.17089	"TBC1 domain family, member 24"	Tbc1d24
1448029_at	1.03194	T-box 3	Tbx3
1448030_at	0.97294	RIKEN cDNA 1700023D09 gene	1700023D09Rik

1448031_at 0.77714 "Guanine nucleotide binding protein, alpha o" Gnao1
1448032_at 1.02547 5-azacytidine induced gene 2 Azi2
1448033_at 1.04556 TatD DNase domain containing 1 Tatdn1
1448034_at 0.75673 expressed sequence AI842396 AI842396
1448037_at 1.06226 "Zinc finger, CCHC domain containing 6" Zcchc6
1448038_at 1.00860 --- ---
1448039_at 0.89469 Similar to gonadotropin inducible ovarian transcription factor 2 LOC622175
1448040_at 0.95869 TCF3 (E2A) fusion partner Tfpt
1448041_at 0.81611 ring finger protein 2 Rnf2
1448042_s_at 0.78554 ring finger protein 2 Rnf2
1448043_x_at 0.85392 ring finger protein 2 Rnf2
1448044_a_at 0.90064 "Adult male kidney cDNA, RIKEN full-length enriched library, clone:
F530012N16 product:hypothetical protein, full insert sequence" ---
1448045_at 1.03856 "Adult male kidney cDNA, RIKEN full-length enriched library, clone:
F530012N16 product:hypothetical protein, full insert sequence" ---
1448046_at 0.94107 Rab9 effector protein with kelch motifs Rabepk
1448047_at 0.99734 "Adult male medulla oblongata cDNA, RIKEN full-length enriched library,
clone:6330524D21 product:unclassifiable, full insert sequence" ---
1448048_at 1.05203 cDNA sequence BC016495 BC016495
1448049_at 1.51526 jumonji domain containing 1C Jmjd1c
1448050_s_at 1.09013 mitogen-activated protein kinase kinase kinase 4 Map4k4
1448051_at 1.14396 --- ---
1448052_at 0.86316 cingulin-like 1 /// similar to cingulin-like 1 isoform 1 Cgn1 /// LOC677485
1448053_at 0.89343 "UDP-Gal:betaGlcNAc beta 1,4-galactosyltransferase, polypeptide 3"
B4galt3
1448054_at 1.00375 "Inhibitor of growth family, member 4" Ing4
1448055_at 0.96298 fibronectin type III domain containing 7 Fndc7
1448056_at 1.30100 "HECT, UBA and WWE domain containing 1" Huwe1
1448057_at 0.83164 --- ---
1448058_s_at 1.25421 budding uninhibited by benzimidazoles 3 homolog (S. cerevisiae) Bub3
1448059_at 0.89658 mirror-image polydactyly gene 1 homolog (human) Mipol1
1448060_at 1.02885 "sema domain, transmembrane domain (TM), and cytoplasmic domain,
(semaphorin) 6D" Sema6d
1448061_at 0.99484 macrophage scavenger receptor 1 Msr1
1448062_at 1.02750 serine/threonine kinase 11 Stk11
1448063_at 0.83693 IQ motif and Sec7 domain 2 Iqsec2
1448064_at 0.88856 RAN binding protein 9 Ranbp9
1448065_at 0.94124 protoporphyrinogen oxidase Ppox
1448066_at 0.94077 hypothetical LOC552909 LOC552909
1448067_at 0.95350 expressed sequence AW011956 AW011956
1448068_at 1.06134 sterol O-acyltransferase 1 Soat1
1448069_at 0.81690 transmembrane 4 superfamily member 1 Tm4sf1
1448070_at 0.99539 "exportin 1, CRM1 homolog (yeast)" Xpo1

1448071_at 0.99062 --- ---
1448072_at 1.13130 RIKEN cDNA 2010010I01 gene 2010010I01Rik
1448073_at 0.95403 hypothetical LOC541456 LOC541456
1448074_at 1.07532 Relaxin 1 Rln1
1448075_at 1.01467 expressed sequence AA408251 AA408251
1448076_at 1.23756 SH2 domain binding protein 1 (tetratricopeptide repeat containing) Sh2bp1
1448078_at 0.97837 expressed sequence C76533 C76533
1448079_at 1.36570 ring finger protein 166 Rnf166
1448080_at 1.05730 E2F transcription factor 8 E2f8
1448081_at 1.13986 Collagen-like tail subunit (single strand of homotrimer) of asymmetric acetylcholinesterase Colq
1448082_at 0.94277 Chloride channel 1 Clcn1
1448083_at 0.85461 similar to voltage gated channel like 1 LOC675405
1448085_at 1.00143 "DNA segment, Chr 5, ERATO Doi 102, expressed" D5Ertd102e
1448086_at 0.96323 "DNA segment, Chr 1, ERATO Doi 164, expressed" D1Ertd164e
1448088_at 0.88563 expressed sequence C78228 C78228
1448089_at 1.06245 "DNA segment, Chr 4, ERATO Doi 58, expressed" D4Ertd58e
1448090_at 1.07106 Transcribed locus ---
1448091_at 0.93302 "DNA segment, Chr 15, ERATO Doi 50, expressed" D15Ertd50e
1448092_x_at 0.94567 "serine (or cysteine) peptidase inhibitor, clade A, member 4, pseudogene 1" Serpina4-ps1
1448093_s_at 1.11787 expressed sequence C77405 C77405
1448094_at 1.12440 hypothetical LOC546100 LOC546100
1448095_at 0.97388 Transcribed locus ---
1448096_at 1.46052 2-oxoglutarate and iron-dependent oxygenase domain containing 1 Ogfod1
1448097_at 1.04621 Ubiquitin protein ligase E3C Ube3c
1448098_at 0.97364 expressed sequence AA517545 AA517545
1448099_at 0.89654 "PREDICTED: Mus musculus hypothetical LOC432995 (LOC432995), mRNA" ---
1448100_at 0.94746 RIKEN cDNA 4833439L19 gene 4833439L19Rik
1448101_s_at 0.95268 tripartite motif protein 27 Trim27
1448102_a_at 1.01682 WD repeat domain 61 Wdr61
1448103_s_at 0.98800 "non-POU-domain-containing, octamer binding protein" Nono
1448104_at 0.65969 "aldehyde dehydrogenase family 6, subfamily A1" Aldh6a1
1448105_at 0.99342 protamine 2 Prm2
1448106_at 1.05210 NECAP endocytosis associated 1 Necap1
1448107_x_at 0.89492 kallikrein 1 Klk1
1448108_at 0.89260 serine incorporator 1 Serinc1
1448109_a_at 1.09334 ribosomal protein L26 Rpl26
1448110_at 0.94009 "sema domain, immunoglobulin domain (Ig), transmembrane domain (TM) and short cytoplasmic domain, (semaphorin) 4A" Sema4a
1448111_at 0.85148 cytidine 5'-triphosphate synthase 2 Ctps2
1448112_at 0.95171 "cytochrome c oxidase, subunit VIIc" Cox7c

1448113_at 0.88199 stathmin 1 /// similar to Stathmin (Phosphoprotein p19) (pp19) (Oncoprotein 18) (Op18) (Leukemia-associated phosphoprotein p18) (pp17) (Prosolin) (Metablastin) (Pr22 protein) (Leukemia-associated gene protein) Stmn1 /// LOC639458
 1448114_a_at 1.05156 HpaII tiny fragments locus 9c Htf9c
 1448115_at 0.99960 HpaII tiny fragments locus 9c Htf9c
 1448116_at 0.98813 "ubiquitin-activating enzyme E1, Chr X" Ube1x
 1448117_at 0.84742 kit ligand Kitl
 1448118_a_at 0.97416 cathepsin D Ctsd
 1448119_at 0.98967 "2,3-bisphosphoglycerate mutase" Bpgm
 1448120_at 0.86414 growth differentiation factor 9 Gdf9
 1448121_at 1.02860 WW domain binding protein 2 Wbp2
 1448122_at 1.05883 t-complex protein 1 Tcp1
 1448123_s_at 0.97663 "transforming growth factor, beta induced" Tgfb1
 1448124_at 0.88986 "glucuronidase, beta" Gusb
 1448125_at 0.98716 Ras-like without CAAX 2 Rit2
 1448126_at 1.04539 "teratocarcinoma expressed, serine rich" Tera
 1448127_at 1.00513 ribonucleotide reductase M1 Rrm1
 1448128_at 0.92915 protective protein for beta-galactosidase Ppob
 1448129_at 0.94871 "actin related protein 2/3 complex, subunit 5" Arpc5
 1448130_at 1.35770 farnesyl diphosphate farnesyl transferase 1 Fdft1
 1448131_at 1.04426 mitofusin 2 Mfn2
 1448132_at 1.04083 "solute carrier family 19 (sodium/hydrogen exchanger), member 1" Slc19a1
 1448133_at 1.08609 NMD3 homolog (S. cerevisiae) Nmd3
 1448134_at 0.86947 cDNA sequence X99384 X99384
 1448135_at 0.96342 activating transcription factor 4 Atf4
 1448136_at 0.96710 ectonucleotide pyrophosphatase/phosphodiesterase 2 Enpp2
 1448137_at 0.96350 "aldehyde dehydrogenase family 7, member A1" Aldh7a1
 1448138_at 0.97881 "protein phosphatase 2A, regulatory subunit B (PR 53)" Ppp2r4
 1448139_at 0.93129 megalencephalic leukoencephalopathy with subcortical cysts 1 homolog (human) Mlc1
 1448140_at 1.03334 cytokine induced apoptosis inhibitor 1 Ciapin1
 1448141_at 0.99269 RIKEN cDNA 1110014J01 gene 1110014J01Rik
 1448142_x_at 1.01786 ribosomal protein S13 /// similar to ribosomal protein S13 /// similar to ribosomal protein S13 /// similar to ribosomal protein S13 Rps13 /// LOC625298 /// LOC633920 /// LOC671723
 1448143_at 0.90233 "aldehyde dehydrogenase 2, mitochondrial" Aldh2
 1448144_at 0.97609 heterogeneous nuclear ribonucleoprotein A/B Hnrpab
 1448145_at 0.97843 WW domain containing E3 ubiquitin protein ligase 2 Wwp2
 1448146_at 0.88275 WW domain containing E3 ubiquitin protein ligase 2 Wwp2
 1448147_at 0.97704 "tumor necrosis factor receptor superfamily, member 19" Tnfrsf19
 1448148_at 0.92431 granulin Grn
 1448149_at 0.95136 "catenin (cadherin associated protein), alpha 1" Ctnna1
 1448150_at 0.99376 nucleoporin 62 Nup62

1448151_at 0.98312 "ELAV (embryonic lethal, abnormal vision, Drosophila)-like 1 (Hu antigen R)" Elavl1
 1448152_at 0.93585 insulin-like growth factor 2 Igf2
 1448153_at 0.96144 "cytochrome c oxidase, subunit Va" Cox5a
 1448154_at 1.20303 N-myc downstream regulated gene 2 Ndr2
 1448155_at 0.90440 programmed cell death 6 interacting protein Pdcd6ip
 1448156_at 0.97585 trefoil factor 1 Tff1
 1448157_s_at 1.02472 ribosomal protein 10 Rpl10
 1448158_at 0.82702 syndecan 1 Sdc1
 1448159_at 1.07114 "RAB7, member RAS oncogene family" Rab7
 1448160_at 0.86537 lymphocyte cytosolic protein 1 Lcp1
 1448161_a_at 0.83346 chloride channel 4-2 Clcn4-2
 1448162_at 1.20572 vascular cell adhesion molecule 1 Vcam1
 1448163_at 1.04881 glucosamine-6-phosphate deaminase 1 /// similar to Glucosamine-6-phosphate isomerase (Glucosamine-6-phosphate deaminase) (GNPDA) (GlcN6P deaminase) (Oscillin) /// similar to Glucosamine-6-phosphate isomerase (Glucosamine-6-phosphate deaminase) (GNPDA) (GlcN6P deaminase) (Oscillin) Gnpda1 /// LOC231914 /// LOC667410
 1448164_at 0.92880 kelch domain containing 3 Klhdc3
 1448165_at 0.87170 caspase 2 Casp2
 1448166_a_at 1.07367 "proteasome (prosome, macropain) subunit, beta type 1" Psmb1
 1448167_at 1.01170 interferon gamma receptor 1 Ifngr1
 1448168_a_at 0.96610 salivary protein 1 Spt1
 1448169_at 1.05297 keratin 18 Krt18
 1448170_at 1.04869 seven in absentia 2 Siah2
 1448171_at 1.28782 seven in absentia 2 Siah2
 1448172_at 0.93915 "malate dehydrogenase 1, NAD (soluble)" Mdh1
 1448173_a_at 0.85731 topoisomerase (DNA) III beta Top3b
 1448174_at 1.01044 cullin 1 Cull1
 1448175_at 1.07271 EH-domain containing 1 Ehd1
 1448176_a_at 0.99321 heterogeneous nuclear ribonucleoprotein K Hnrpk
 1448177_at 0.96176 "transporter 1, ATP-binding cassette, sub-family B (MDR/TAP)" Tap1
 1448178_a_at 1.04901 chaperonin subunit 3 (gamma) Cct3
 1448179_at 0.88456 upregulated during skeletal muscle growth 5 Usmg5
 1448180_a_at 0.97827 hematological and neurological expressed sequence 1 Hn1
 1448181_at 0.93466 Kruppel-like factor 15 Klf15
 1448182_a_at 0.92080 CD24a antigen Cd24a
 1448183_a_at 1.04230 "hypoxia inducible factor 1, alpha subunit" Hif1a
 1448184_at 1.00453 FK506 binding protein 1a Fkbp1a
 1448185_at 1.03518 "homocysteine-inducible, endoplasmic reticulum stress-inducible, ubiquitin-like domain member 1" Herpud1
 1448186_at 1.04718 pancreatic lipase-related protein 2 Pnliprp2
 1448187_at 1.00663 "polymerase (DNA directed), delta 1, catalytic subunit" Pold1
 1448188_at 0.91342 "uncoupling protein 2 (mitochondrial, proton carrier)" Ucp2

1448189_a_at 0.97858 flightless I homolog (Drosophila) Fliih
1448190_at 1.15004 mitochondrial ribosomal protein L33 Mrpl33
1448191_at 1.05426 polo-like kinase 1 (Drosophila) Plk1
1448192_s_at 1.02851 phosphoribosyl pyrophosphate synthetase 1 /// expressed sequence
AU021838 Prps1 /// AU021838
1448193_at 1.02585 RIKEN cDNA 5730403B10 gene 5730403B10Rik
1448194_a_at 0.99610 H19 fetal liver mRNA H19
1448195_at 1.09423 "TAF5-like RNA polymerase II, p300/CBP-associated factor (PCAF)-
associated factor" Taf5l
1448196_at 0.98296 "methionine adenosyltransferase II, beta" Mat2b
1448197_at 1.01501 CEA-related cell adhesion molecule 11 Ceacam11
1448198_a_at 0.95676 NADH dehydrogenase (ubiquinone) 1 beta subcomplex 8 Ndufb8
1448199_at 1.08427 ankyrin repeat domain 10 Ankrd10
1448200_at 0.66536 transcobalamin 2 Tcn2
1448201_at 1.19309 secreted frizzled-related protein 2 Sfrp2
1448202_x_at 1.13080 RIKEN cDNA 2610524G07 gene 2610524G07Rik
1448203_at 0.96549 "ATP synthase, H+ transporting, mitochondrial F0 complex, subunit g" Atp5l
1448204_at 1.12708 salvador homolog 1 (Drosophila) Sav1
1448205_at 0.97585 "cyclin B1, related sequence 1 /// cyclin B1" Ccnb1-rs1 /// Ccnb1
1448206_at 1.08551 "proteasome (prosome, macropain) subunit, alpha type 2" Psma2
1448207_at 1.04083 LIM and SH3 protein 1 Lasp1
1448208_at 0.98203 MAD homolog 1 (Drosophila) Smad1
1448209_a_at 0.82216 "solute carrier family 22 (organic cation transporter), member 17" Slc22a17
1448210_at 1.01321 "RAB1, member RAS oncogene family" Rab1
1448211_at 0.83594 "ATPase, H+ transporting, lysosomal V0 subunit E2" Atp6v0e2
1448212_at 1.07841 "tumor necrosis factor superfamily, member 5-induced protein 1" Tnfsf5ip1
1448213_at 0.98413 annexin A1 Anxa1
1448214_at 0.97910 pyruvate dehydrogenase (lipoamide) beta /// similar to pyruvate dehydrogenase
(lipoamide) beta Pdhb /// LOC620009
1448215_a_at 1.04032 dipeptidylpeptidase 3 Dpp3
1448216_at 0.87311 synaptogyrin 3 Syngr3
1448217_a_at 1.06525 ribosomal protein L27 /// similar to ribosomal protein L27 /// similar to
ribosomal protein L27 /// similar to ribosomal protein L27 /// similar to ribosomal protein L27 /// similar
to ribosomal protein L27 Rpl27 /// LOC621100 /// LOC622236 /// LOC669376 /// LOC672370 ///
LOC675088
1448218_s_at 1.14330 "tyrosine 3-monooxygenase/tryptophan 5-monooxygenase activation protein,
zeta polypeptide" Ywhaz
1448219_a_at 1.00718 "tyrosine 3-monooxygenase/tryptophan 5-monooxygenase activation protein,
zeta polypeptide" Ywhaz
1448220_at 1.01080 chymotrypsinogen B1 Ctrb1
1448221_at 0.96497 HLA-B-associated transcript 1A Bat1a
1448222_x_at 0.94261 "cytochrome c oxidase, subunit VIIIa" Cox8a
1448223_at 0.88459 fatso Fto

1448224_at 0.99922 "transcription factor A, mitochondrial" Tfam
 1448225_at 1.01591 GPI anchor attachment protein 1 Gpaa1
 1448226_at 0.89969 ribonucleotide reductase M2 Rrm2
 1448227_at 0.99018 growth factor receptor bound protein 7 Grb7
 1448228_at 0.96040 lysyl oxidase Lox
 1448229_s_at 1.10302 cyclin D2 Ccnd2
 1448230_at 1.07934 ubiquitin specific peptidase 10 Usp10
 1448231_at 1.10412 FK506 binding protein 5 Fkbp5
 1448232_x_at 1.00438 "tubulin, alpha 6 /// similar to Tubulin alpha-6 chain (Alpha-tubulin 6)"
 Tuba6 /// LOC626534
 1448233_at 0.83895 prion protein Prnp
 1448234_at 0.97600 "DnaJ (Hsp40) homolog, subfamily B, member 6" Dnajb6
 1448235_s_at 0.96160 high mobility group box 1 /// RIKEN cDNA 4932431P20 gene /// similar to
 High mobility group protein 1 (HMG-1) (High mobility group protein B1) (Amphoterin) (Heparin-
 binding protein p30) /// similar to High mobility group protein 1 (HMG-1) (High mobility group protein
 B1) (Amphoterin) (Heparin-binding protein p30) /// similar to High mobility group protein 1 (HMG-1)
 (High mobility group protein B1) (Amphoterin) (Heparin-binding protein p30) /// similar to High
 mobility group protein 1 (HMG-1) (High mobility group protein B1) (Amphoterin) (Heparin-binding
 protein p30) /// similar to High mobility group protein 1 (HMG-1) (High mobility group protein B1)
 (Amphoterin) (Heparin-binding protein p30) /// similar to High mobility group protein 1 (HMG-1) (High
 mobility group protein B1) (Amphoterin) (Heparin-binding protein p30) /// similar to High mobility
 group protein 1 (HMG-1) (High mobility group protein B1) (Amphoterin) (Heparin-binding protein
 p30) Hmgb1 /// 4932431P20Rik /// LOC432959 /// LOC434174 /// LOC545555 /// LOC637733 ///
 LOC640060 /// LOC667777 /// LOC669546
 1448236_at 0.99239 radixin Rdx
 1448237_x_at 0.86720 lactate dehydrogenase B Ldhb
 1448238_at 1.02415 RIKEN cDNA 2700060E02 gene 2700060E02Rik
 1448239_at 1.21822 heme oxygenase (decycling) 1 Hmox1
 1448240_at 0.89423 "membrane-bound transcription factor peptidase, site 1" Mbtps1
 1448241_at 1.10189 GM2 ganglioside activator protein Gm2a
 1448242_at 0.97025 Sec61 alpha 1 subunit (*S. cerevisiae*) Sec61a1
 1448243_at 1.02062 N-ethylmaleimide sensitive fusion protein attachment protein alpha Napa
 1448244_at 0.99964 lysophospholipase 1 Lypl1
 1448245_at 1.04484 ribosomal protein SA Rpsa
 1448246_at 0.97250 histone deacetylase 1 Hdac1
 1448247_at 0.92607 B-cell CLL/lymphoma 7B Bcl7b
 1448248_at 1.03288 v-crk sarcoma virus CT10 oncogene homolog (avian) Crk
 1448249_at 0.87851 glycerol-3-phosphate dehydrogenase 1 (soluble) Gpd1
 1448250_at 0.93429 RIKEN cDNA 9030425E11 gene 9030425E11Rik
 1448251_at 1.09184 RIKEN cDNA 9030425E11 gene 9030425E11Rik
 1448252_a_at 1.07194 eukaryotic translation elongation factor 1 beta 2 Eef1b2
 1448253_at 1.06705 glutamate dehydrogenase 1 Glud1
 1448254_at 0.97909 pleiotrophin Ptn

1448255_a_at 0.98657 surfeit gene 4 Surf4
1448256_at 1.09245 golgi SNAP receptor complex member 1 Gosr1
1448257_at 1.03432 "solute carrier family 29 (nucleoside transporters), member 2" Slc29a2
1448258_a_at 0.94760 signal peptidase complex subunit 1 homolog (S. cerevisiae) Spcs1
1448259_at 0.73988 follistatin-like 1 Fstl1
1448260_at 0.92374 ubiquitin carboxy-terminal hydrolase L1 Uchl1
1448261_at 0.88964 cadherin 1 Cdh1
1448262_at 1.03860 "proteasome (prosome, macropain) subunit, beta type 2" Psmb2
1448263_a_at 1.05325 CNDP dipeptidase 2 (metallopeptidase M20 family) Cndp2
1448264_a_at 1.07562 "eukaryotic translation initiation factor 3, subunit 2 (beta)" Eif3s2
1448265_x_at 0.84189 epithelial V-like antigen 1 Eval
1448266_at 1.05253 endothelial differentiation-related factor 1 Edf1
1448267_at 0.85978 syntaxin 5A Stx5a
1448268_at 0.95958 transmembrane emp24 protein transport domain containing 9 Tmed9
1448269_a_at 1.06632 kelch-like 13 (Drosophila) Klhl13
1448270_at 1.19708 DEAD (Asp-Glu-Ala-Asp) box polypeptide 21 Ddx21
1448271_a_at 1.34975 DEAD (Asp-Glu-Ala-Asp) box polypeptide 21 Ddx21
1448272_at 0.71782 "B-cell translocation gene 2, anti-proliferative" Btg2
1448273_at 0.87073 glutathione synthetase Gss
1448274_at 1.12815 "complement component 1, q subcomponent binding protein" C1qbp
1448275_at 1.00273 transmembrane protein 19 Tmem19
1448276_at 1.01699 tetraspanin 4 Tspan4
1448277_at 1.03661 "polymerase (DNA directed), delta 2, regulatory subunit" Pold2
1448278_at 0.95380 XPA binding protein 2 Xab2
1448279_at 0.94638 "actin related protein 2/3 complex, subunit 3" Arpc3
1448280_at 0.98433 synaptophysin Syt
1448281_a_at 0.96836 elastase 2A Ela2a
1448282_at 0.99455 "pleiotropic regulator 1, PRL1 homolog (Arabidopsis)" Plrg1
1448283_a_at 0.99706 ubiquitin-like 1 (sentrin) activating enzyme E1B Uble1b
1448284_a_at 0.87439 "NADH dehydrogenase (ubiquinone) 1, subcomplex unknown, 1" Ndufc1
1448285_at 0.96644 regulator of G-protein signaling 4 Rgs4
1448286_at 1.09051 hydroxyacyl-Coenzyme A dehydrogenase type II Hadh2
1448287_at 1.01474 RNA polymerase 1-3 Rpo1-3
1448288_at 1.02077 nuclear factor I/B Nfib
1448289_at 0.98252 collapsin response mediator protein 1 Crmp1
1448290_at 0.91034 pancreatitis-associated protein Pap
1448291_at 1.01590 matrix metalloproteinase 9 Mmp9
1448292_at 1.01571 ubiquinol-cytochrome c reductase (6.4kD) subunit Uqcrc
1448293_at 0.92368 --- ---
1448294_at 0.94119 LPS-induced TN factor /// hypothetical protein LOC545182 Litaf ///
LOC545182
1448295_at 1.03771 "DNA segment, Chr 13, Wayne State University 50, expressed" D13Wsu50e
1448296_x_at 0.96483 "tubulin, alpha 3 /// tubulin, alpha 7 /// similar to tubulin, alpha 1" Tuba3 ///

Tuba7 /// LOC384954

1448297_a_at 0.94438 "tyrosine kinase, non-receptor, 2 /// similar to Activated CDC42 kinase 1 (ACK-1) (Non-receptor protein tyrosine kinase Ack) (Tyrosine kinase non-receptor protein 2)"

Tnk2 /// LOC672286

1448298_at 0.94216 "tyrosine kinase, non-receptor, 2" Tnk2

1448299_at 0.87997 "solute carrier family 1 (neuronal/epithelial high affinity glutamate transporter, system Xag), member 1" Slc1a1

1448300_at 0.80191 microsomal glutathione S-transferase 3 Mgst3

1448301_s_at 0.90521 "serine (or cysteine) peptidase inhibitor, clade B, member 1a" Serpinb1a

1448302_at 1.00197 potassium channel tetramerisation domain containing 20 Kctd20

1448303_at 0.98930 glycoprotein (transmembrane) nmb Gpnmb

1448304_a_at 1.07662 "RAB6, member RAS oncogene family" Rab6

1448305_at 0.90866 "RAB6, member RAS oncogene family" Rab6

1448306_at 1.49009 "nuclear factor of kappa light chain gene enhancer in B-cells inhibitor, alpha" Nfkbia

1448307_at 1.10793 Down syndrome critical region homolog 2 (human) Dscr2

1448308_at 1.01214 "adaptor-related protein complex 3, mu 1 subunit" Ap3m1

1448309_at 0.99994 "adaptor-related protein complex 3, mu 1 subunit" Ap3m1

1448310_at 1.04109 intestinal cell kinase Ick

1448311_at 1.00759 ubiquitin specific peptidase 5 (isopeptidase T) Usp5

1448312_at 0.91844 proprotein convertase subtilisin/kexin type 2 Pcsk2

1448313_at 0.90808 tripeptidyl peptidase I Tpp1

1448314_at 0.99162 cell division cycle 2 homolog A (S. pombe) Cdc2a

1448315_a_at 1.07310 "pyrroline-5-carboxylate reductase family, member 2" Pycr2

1448316_at 0.97821 CKLF-like MARVEL transmembrane domain containing 3 Cmtm3

1448317_at 0.98159 transmembrane protein 128 Tmem128

1448318_at 1.00977 adipose differentiation related protein Adfp

1448319_at 1.05901 "aldo-keto reductase family 1, member B3 (aldose reductase) /// similar to Aldose reductase (AR) (Aldehyde reductase) /// similar to Aldose reductase (AR) (Aldehyde reductase) /// similar to Aldose reductase (AR) (Aldehyde reductase) /// similar to Aldose reductase (AR) (Aldehyde reductase) /// similar to Aldose reductase (AR) (Aldehyde reductase)" Akr1b3 ///

LOC235852 /// LOC382486 /// LOC385448 /// LOC385495 /// LOC672268

1448320_at 0.78004 stromal interaction molecule 1 Stim1

1448321_at 1.02691 SPARC related modular calcium binding 1 Smoc1

1448322_a_at 0.94325 cytochrome c oxidase subunit IV isoform 1 Cox4i1

1448323_a_at 0.88279 biglycan Bgn

1448324_at 1.10874 ribonucleic acid binding protein S1 Rnps1

1448325_at 0.87062 myeloid differentiation primary response gene 116 Myd116

1448326_a_at 0.86448 cellular retinoic acid binding protein I Crabp1

1448327_at 0.99791 actinin alpha 2 Actn2

1448328_at 1.01413 SH3-domain binding protein 2 Sh3bp2

1448329_at 0.98763 a disintegrin and metallopeptidase domain 3 (cyritestin) Adam3

1448330_at 0.84155 "glutathione S-transferase, mu 1" Gstm1

1448331_at 1.05061 "NADH dehydrogenase (ubiquinone) 1 beta subcomplex, 7" Ndufb7
1448332_at 1.05040 peroxisome biogenesis factor 19 Pex19
1448333_at 0.99566 ADP-ribosylarginine hydrolase Adprh
1448334_a_at 0.87636 cyclin I Ccni
1448335_s_at 0.87154 cyclin I Ccni
1448336_at 0.95683 developmentally regulated GTP binding protein 1 Drg1
1448337_at 0.93591 RIKEN cDNA 2410003P15 gene 2410003P15Rik
1448338_at 1.04688 plasma glutamate carboxypeptidase Pgcg
1448339_at 0.93020 transmembrane protein 30A Tmem30a
1448340_at 0.98001 transmembrane protein 30A Tmem30a
1448341_a_at 0.97687 syntaxin binding protein 2 Stxbp2
1448342_at 1.02903 mitogen activated protein kinase 10 Mapk10
1448343_a_at 0.92371 neighbor of Brca1 gene 1 Nbr1
1448344_at 1.07973 ribosomal protein S12 /// similar to ribosomal protein S12 /// similar to
ribosomal protein S12 /// similar to ribosomal protein S12 /// similar to ribosomal protein S12 /// similar
to ribosomal protein S12 /// similar to 40S ribosomal protein S12 /// similar to ribosomal protein S12 ///
similar to ribosomal protein S12 /// similar to ribosomal protein S12 Rps12 /// LOC432865 ///
LOC627835 /// LOC668354 /// LOC670751 /// LOC670832 /// LOC671641 /// LOC672008 ///
LOC672415 /// LOC676277
1448345_at 0.97094 translocase of outer mitochondrial membrane 34 Tomm34
1448346_at 1.12841 "cofilin 1, non-muscle" Cfl1
1448347_a_at 1.00592 GPI-anchored membrane protein 1 Gpiap1
1448348_at 1.02383 GPI-anchored membrane protein 1 Gpiap1
1448349_at 1.05705 "vesicle-associated membrane protein, associated protein A" Vapa
1448350_at 1.16735 argininosuccinate lyase Asl
1448351_at 1.02173 "coronin, actin binding protein 1B" Coro1b
1448352_at 0.90212 leucine zipper protein 1 Luzp1
1448353_x_at 0.98208 ribophorin I Rpn1
1448354_at 1.07515 glucose-6-phosphate dehydrogenase X-linked G6pdx
1448355_at 0.87206 "protease, serine, 16 (thymus)" Prss16
1448356_at 0.88849 ubiquitin-conjugating enzyme E2D 2 Ube2d2
1448357_at 1.10148 small nuclear ribonucleoprotein polypeptide G /// similar to small nuclear
ribonucleoprotein polypeptide G /// similar to small nuclear ribonucleoprotein polypeptide G Snrpg ///
LOC666609 /// LOC671824
1448358_s_at 1.03406 small nuclear ribonucleoprotein polypeptide G /// similar to small nuclear
ribonucleoprotein polypeptide G /// similar to small nuclear ribonucleoprotein polypeptide G Snrpg ///
LOC666609 /// LOC671824
1448359_a_at 1.22872 "HIG1 domain family, member 1A" Higd1a
1448360_s_at 0.94269 angel homolog 2 (Drosophila) Angel2
1448361_at 0.83441 tetratricopeptide repeat domain 3 Ttc3
1448362_at 1.08440 "DnaJ (Hsp40) homolog, subfamily C, member 7" Dnajc7
1448363_at 1.02548 yes-associated protein 1 Yap1
1448364_at 1.28380 cyclin G2 Ccng2

1448365_at 1.12576 exosome component 7 Exosc7
1448366_at 0.88585 syntaxin 1A (brain) Stx1a
1448367_at 1.02566 stromal cell derived factor 4 Sdf4
1448368_at 0.96619 dynactin 6 Dctn6
1448369_at 0.98984 "polymerase (DNA directed), alpha 2" Pola2
1448370_at 0.83759 Unc-51 like kinase 1 (C. elegans) Ulk1
1448371_at 0.98582 "myosin light chain, phosphorylatable, fast skeletal muscle" Mylpf
1448372_a_at 0.98928 transmembrane protein 4 Tmem4
1448373_at 1.07158 mitochondrial ribosomal protein L18 Mrpl18
1448374_at 0.96051 "mediator of RNA polymerase II transcription, subunit 28 homolog (yeast)"
Med28
1448375_at 0.87943 transmembrane 9 superfamily member 3 Tm9sf3
1448376_at 1.05693 Werner helicase interacting protein 1 Wrnip1
1448377_at 0.94466 secretory leukocyte peptidase inhibitor Slpi
1448378_at 0.98054 "fascin homolog 1, actin bundling protein (Strongylocentrotus purpuratus)"
Fscn1
1448379_at 1.03224 protection of telomeres 1 Pot1
1448380_at 1.17765 "lectin, galactoside-binding, soluble, 3 binding protein" Lgals3bp
1448381_at 0.98972 "G elongation factor, mitochondrial 1" Gfm1
1448382_at 0.93326 "enoyl-Coenzyme A, hydratase/3-hydroxyacyl Coenzyme A dehydrogenase"
Ehhadh
1448383_at 0.73786 matrix metalloproteinase 14 (membrane-inserted) Mmp14
1448384_at 0.86005 protein O-fucosyltransferase 2 Pofut2
1448385_at 1.00407 "solute carrier family 15, member 4" Slc15a4
1448386_a_at 0.92209 whey acidic protein Wap
1448387_at 1.08287 ring-box 1 Rbx1
1448388_a_at 0.96532 RIKEN cDNA 1110002B05 gene 1110002B05Rik
1448389_at 1.02308 WD repeat domain 5 Wdr5
1448390_a_at 0.64187 dehydrogenase/reductase (SDR family) member 3 Dhrs3
1448391_at 1.04344 "RAB9, member RAS oncogene family" Rab9
1448392_at 0.91311 secreted acidic cysteine rich glycoprotein Sparc
1448393_at 0.83193 claudin 7 Cldn7
1448394_at 0.99971 "myosin, light polypeptide 2, regulatory, cardiac, slow" Myl2
1448395_at 0.83874 secreted frizzled-related sequence protein 1 Sfrp1
1448396_at 0.99254 transmembrane protein 131 Tmem131
1448397_at 1.02985 gap junction membrane channel protein beta 6 Gjb6
1448398_s_at 0.88037 ribosomal protein L22 Rpl22
1448399_at 1.02826 Tax1 (human T-cell leukemia virus type I) binding protein 1 Tax1bp1
1448400_a_at 1.00656 "SWI/SNF related, matrix associated, actin dependent regulator of chromatin,
subfamily d, member 2" Smarcd2
1448401_at 1.00494 "SWI/SNF related, matrix associated, actin dependent regulator of chromatin,
subfamily d, member 2" Smarcd2
1448402_at 0.97385 talin 1 Tln1

1448403_at 1.12020 leucyl-tRNA synthetase Lars
1448404_at 1.01155 secretory carrier membrane protein 2 Scamp2
1448405_a_at 0.98712 CREBBP/EP300 inhibitory protein 1 Cri1
1448406_at 0.98267 CREBBP/EP300 inhibitory protein 1 Cri1
1448407_at 0.92472 RIKEN cDNA 4632428N05 gene 4632428N05Rik
1448408_at 0.87721 Hermansky-Pudlak syndrome 1 homolog (human) Hps1
1448409_at 1.09555 lymphoid-restricted membrane protein Lrmp
1448410_at 0.99800 "ubiquitination factor E4B, UFD2 homolog (S. cerevisiae)" Ube4b
1448411_at 0.94012 Wolfram syndrome 1 homolog (human) Wfs1
1448412_a_at 0.92883 TSC22 domain family 4 Tsc22d4
1448413_at 1.00247 RIKEN cDNA 2410016O06 gene 2410016O06Rik
1448414_at 1.00263 RAD1 homolog (S. pombe) Rad1
1448415_a_at 0.89583 "sema domain, immunoglobulin domain (Ig), short basic domain, secreted, (semaphorin) 3B" Sema3b
1448416_at 0.62074 matrix Gla protein Mgp
1448417_at 1.00209 ninjurin 1 Ninj1
1448418_s_at 0.99525 WD repeat domain 23 Wdr23
1448419_at 0.97863 "processing of precursor 4, ribonuclease P/MRP family, (S. cerevisiae)" Pop4
1448420_a_at 1.02790 F-box and leucine-rich repeat protein 12 Fbxl12
1448421_s_at 0.89594 asporin Asp1
1448422_at 0.89248 transmembrane emp24 protein transport domain containing 4 Tmed4
1448423_at 0.96261 transmembrane emp24 protein transport domain containing 4 Tmed4
1448424_at 0.88842 frizzled-related protein Frzb
1448425_at 1.05753 "eukaryotic translation initiation factor 3, subunit 10 (theta)" Eif3s10
1448426_at 0.69052 sarcosine dehydrogenase Sardh
1448427_at 0.89029 "NADH dehydrogenase (ubiquinone) 1 alpha subcomplex, 6 (B14)" Ndufa6
1448428_at 0.76937 "neuroblastoma, suppression of tumorigenicity 1" Nbl1
1448429_at 1.02886 glycogenin Gyg
1448430_a_at 1.10584 nascent polypeptide-associated complex alpha polypeptide /// similar to nascent polypeptide-associated complex alpha polypeptide /// similar to nascent polypeptide-associated complex alpha polypeptide /// similar to nascent polypeptide-associated complex alpha polypeptide Naca /// LOC628277 /// LOC633817 /// LOC669373
1448431_at 1.04655 ankyrin repeat and SOCS box-containing protein 6 Asb6
1448432_at 0.89482 "phospholipase C, delta 1" Plcd1
1448433_a_at 0.96074 procollagen C-endopeptidase enhancer protein Pcolce
1448434_at 0.95069 ring finger protein 103 Rnf103
1448435_at 0.99176 "positive cofactor 2, multiprotein complex, glutamine/Q-rich-associated protein" Pcqap
1448436_a_at 1.08115 interferon regulatory factor 1 Irf1
1448437_a_at 0.99428 GTP binding protein 2 Gtpbp2
1448438_at 0.99620 "Der1-like domain family, member 2" Derl2
1448439_at 1.01957 "DNA segment, Chr 17, Wayne State University 104, expressed" D17Wsu104e
1448440_x_at 1.02595 "DNA segment, Chr 17, Wayne State University 104, expressed"

D17Wsu104e

1448441_at 1.10819 CDC28 protein kinase 1b Cks1b
 1448442_a_at 1.08853 "proteasome (prosome, macropain) subunit, alpha type 3" Psma3
 1448443_at 1.05674 "serine (or cysteine) peptidase inhibitor, clade I, member 1" Serpini1
 1448444_at 0.95776 --- ---
 1448445_at 0.97485 "acid phosphatase 6, lysophosphatidic" Acp6
 1448446_at 0.90284 deformed epidermal autoregulatory factor 1 (Drosophila) Deaf1
 1448447_at 1.04882 vacuolar protein sorting 28 (yeast) Vps28
 1448448_a_at 0.99897 choline kinase beta Chkb
 1448449_at 1.06503 receptor-interacting serine-threonine kinase 3 Ripk3
 1448450_at 0.98026 adenylate kinase 2 Ak2
 1448451_at 1.11767 adenylate kinase 2 Ak2
 1448452_at 0.86522 interferon regulatory factor 8 Irf8
 1448453_at 0.91917 "hydroxy-delta-5-steroid dehydrogenase, 3 beta- and steroid delta-isomerase 1" Hsd3b1
 1448454_at 0.99949 "splicing factor, arginine/serine-rich 6" Sfrs6
 1448455_at 1.13371 "ceroid-lipofuscinosis, neuronal 8" Cln8
 1448456_at 1.12921 "ceroid-lipofuscinosis, neuronal 8" Cln8
 1448457_at 1.06638 keratin 71 Krt71
 1448458_at 1.00469 topoisomerase (DNA) II beta Top2b
 1448459_at 0.95960 Kv channel-interacting protein 1 Kcnip1
 1448460_at 0.77354 "activin A receptor, type 1" Acvr1
 1448461_a_at 0.99947 THO complex 7 homolog (Drosophila) Thoc7
 1448462_at 1.10836 thymine DNA glycosylase /// similar to thymine DNA glycosylase isoform 2 /// similar to G/T mismatch-specific thymine DNA glycosylase (C-JUN leucine zipper interactive protein JZA-3) /// similar to thymine DNA glycosylase isoform 2 /// similar to G/T mismatch-specific thymine DNA glycosylase (C-JUN leucine zipper interactive protein JZA-3) Tdg /// LOC545124 /// LOC624784 /// LOC670176 /// LOC672659
 1448463_at 1.09700 RIKEN cDNA 4933434E20 gene 4933434E20Rik
 1448464_at 1.01476 YKT6 homolog (S. Cerevisiae) Ykt6
 1448465_at 0.80894 4-nitrophenylphosphatase domain and non-neuronal SNAP25-like protein homolog 1 (C. elegans) Nipsnap1
 1448466_at 0.96802 cell division cycle associated 5 Cdca5
 1448467_a_at 1.09475 RIKEN cDNA G430002G23 gene G430002G23Rik
 1448468_a_at 0.94983 "potassium voltage-gated channel, shaker-related subfamily, beta member 1" Kcnab1
 1448469_at 1.00321 nidogen 1 /// similar to Nidogen precursor (Entactin) Nid1 /// LOC630776
 1448470_at 0.99077 fructose biphosphatase 1 Fbp1
 1448471_a_at 0.77372 cytotoxic T lymphocyte-associated protein 2 alpha Ctla2a
 1448472_at 1.17794 valyl-tRNA synthetase 2 Vars2
 1448473_at 0.87023 budding uninhibited by benzimidazoles 3 homolog (S. cerevisiae) Bub3
 1448474_at 1.03555 NIMA (never in mitosis gene a)-related expressed kinase 7 Nek7
 1448475_at 0.94211 olfactomedin-like 3 Olfml3

1448476_at 1.03042 nucleosome assembly protein 1-like 4 Nap114
1448477_at 0.91594 carbohydrate sulfotransferase 12 Chst12
1448478_at 1.00589 Trf (TATA binding protein-related factor)-proximal protein homolog
(Drosophila) Trfp
1448479_at 1.06472 "proteasome (prosome, macropain) 26S subunit, non-ATPase, 3" Psm3
1448480_at 1.21474 nuclear import 7 homolog (S. cerevisiae) Nip7
1448481_at 1.09650 neuraminidase 1 Neu1
1448482_at 0.71269 "solute carrier family 39 (metal ion transporter), member 8" Slc39a8
1448483_a_at 0.96291 "NADH dehydrogenase (ubiquinone) 1 beta subcomplex, 2" Ndufb2
1448484_at 1.15135 S-adenosylmethionine decarboxylase 1 Amd1
1448485_at 0.83575 gamma-glutamyltransferase 1 Ggt1
1448486_at 0.96921 methylmalonyl-Coenzyme A mutase Mut
1448487_at 0.93668 leucine rich repeat (in FLII) interacting protein 1 Lrrfp1
1448488_at 1.00589 mitochondrial ribosomal protein S5 Mrps5
1448489_at 0.81478 platelet-activating factor acetylhydrolase 2 Pafah2
1448490_at 0.95675 aarF domain containing kinase 4 Adck4
1448491_at 0.84291 "enoyl coenzyme A hydratase 1, peroxisomal" Ech1
1448492_a_at 1.12622 "proteasome (prosome, macropain) 26S subunit, non-ATPase, 12" Psm12
1448493_at 1.00082 polyadenylate-binding protein-interacting protein 2 Paip2
1448494_at 0.87627 growth arrest specific 1 Gas1
1448495_at 1.06092 tissue specific transplantation antigen P35B Tsta3
1448496_a_at 1.03954 "inhibitor of growth family, member 1" Ing1
1448497_at 0.99542 "excision repair cross-complementing rodent repair deficiency,
complementation group 3" Ercc3
1448498_at 1.12171 "ribosomal protein S6 kinase, polypeptide 4" Rps6ka4
1448499_a_at 1.01143 "epoxide hydrolase 2, cytoplasmic" Ephx2
1448500_a_at 0.95905 Lck interacting transmembrane adaptor 1 Lime1
1448501_at 1.09820 tetraspanin 6 Tspan6
1448502_at 0.92938 "solute carrier family 16 (monocarboxylic acid transporters), member 7" Slc16a7
1448503_at 0.94264 myeloid cell leukemia sequence 1 Mcl1
1448504_a_at 0.95668 chromobox homolog 3 (Drosophila HP1 gamma) /// similar to chromobox
homolog 3 /// similar to chromobox homolog 3 /// similar to chromobox homolog 3 Cbx3 ///
LOC629578 /// LOC633016 /// LOC672750
1448505_at 1.07858 nuclear DNA binding protein C1d
1448506_at 0.91771 "serine (or cysteine) peptidase inhibitor, clade A, member 6" Serpina6
1448507_at 1.01378 EF hand domain containing 1 Efhd1
1448508_at 1.02014 Traf3 interacting protein 2 Traf3ip2
1448509_at 1.11766 RIKEN cDNA 3110001A13 gene 3110001A13Rik
1448510_at 0.92551 ephrin A1 Efna1
1448511_at 0.96811 "protein tyrosine phosphatase, receptor type, C polypeptide-associated protein"
Ptprcap
1448512_at 0.79914 histone H1-like protein in spermatids 1 Hils1
1448513_a_at 1.06402 Niemann Pick type C2 Npc2

1448514_at 0.89036 "cytochrome c oxidase, subunit Vb" Cox5b
1448515_at 0.99225 translin Tsn
1448516_at 1.02177 translin Tsn
1448517_at 1.02520 translocase of inner mitochondrial membrane 22 homolog (yeast) Timm22
1448518_at 0.80636 translocase of inner mitochondrial membrane 22 homolog (yeast) Timm22
1448519_at 0.88850 TEA domain family member 2 Tead2
1448520_at 1.06082 "DNA cross-link repair 1B, PSO2 homolog (S. cerevisiae)" Dclre1b
1448521_at 1.01207 bromodomain containing 7 /// similar to bromodomain containing 7 Brd7 ///
LOC634327
1448522_at 0.98883 seryl-aminoacyl-tRNA synthetase 2 Sars2
1448523_at 0.91222 nephronophthisis 1 (juvenile) homolog (human) Nphp1
1448524_s_at 1.13822 "signal sequence receptor, delta" Ssr4
1448525_a_at 1.15381 BCL2/adenovirus E1B interacting protein 3-like Bnip3l
1448526_at 1.10400 karyopherin (importin) beta 1 Kpnb1
1448527_at 1.00870 programmed cell death 10 Pdcd10
1448528_at 0.95117 programmed cell death 10 Pdcd10
1448529_at 0.69999 thrombomodulin Thbd
1448530_at 0.86620 guanosine monophosphate reductase Gmpr
1448531_at 0.91629 lamin B2 /// similar to Lamin-B2 Lmnb2 /// LOC671511
1448532_at 0.97981 prolactin-like protein C 2 Prlpc2
1448533_at 1.02435 cytoskeleton-associated protein 1 Ckap1
1448534_at 0.92386 signal-regulatory protein alpha Sirpa
1448535_at 0.95877 elongation protein 4 homolog (S. cerevisiae) Elp4
1448536_at 1.01847 "LSM3 homolog, U6 small nuclear RNA associated (S. cerevisiae)" Lsm3
1448537_at 1.05930 tetratricopeptide repeat domain 1 Ttc1
1448538_a_at 1.11423 "DNA segment, Chr 4, Wayne State University 53, expressed" D4Wsu53e
1448539_a_at 0.86543 aspartoacylase (aminoacylase) 3 Acy3
1448540_a_at 0.98035 RIKEN cDNA 0610012G03 gene /// hypothetical protein LOC638521
0610012G03Rik /// LOC638521
1448541_at 0.89067 kinesin 2 Kns2
1448542_at 1.14075 BRCA2 and CDKN1A interacting protein Bccip
1448543_at 0.94543 RIKEN cDNA 2310042G06 gene 2310042G06Rik
1448544_at 0.85989 carnitine acetyltransferase Crat
1448545_at 0.77727 syndecan 2 Sdc2
1448546_at 0.90201 Ras association (RalGDS/AF-6) domain family 3 Rassf3
1448547_at 0.87814 Ras association (RalGDS/AF-6) domain family 3 Rassf3
1448548_at 0.92478 tubby like protein 4 Tulp4
1448549_a_at 1.03422 dolichyl-phosphate (UDP-N-acetylglucosamine)
acetylglucosaminophosphotransferase 1 (GlcNAc-1-P transferase) Dpagt1
1448550_at 0.99686 lipopolysaccharide binding protein Lbp
1448551_a_at 0.81064 tripartite motif protein 2 Trim2
1448552_s_at 0.94476 RIKEN cDNA 2310028N02 gene 2310028N02Rik
1448553_at 1.06219 "similar to myosin, heavy polypeptide 4, skeletal muscle /// similar to myosin,

heavy polypeptide 7, cardiac muscle, beta" LOC671894 /// LOC674761
1448554_s_at 1.02246 "myosin, heavy polypeptide 6, cardiac muscle, alpha /// similar to myosin,
heavy polypeptide 4, skeletal muscle /// similar to myosin, heavy polypeptide 7, cardiac muscle, beta"
Myh6 /// LOC671894 /// LOC674761
1448555_at 1.00863 "DNA segment, Chr 15, ERATO Doi 682, expressed" D15Ertd682e
1448556_at 0.84297 prolactin receptor Prlr
1448557_at 0.98611 RIKEN cDNA 1200015N20 gene 1200015N20Rik
1448558_a_at 0.81855 "phospholipase A2, group IVA (cytosolic, calcium-dependent)" Pla2g4a
1448559_at 0.96761 flotillin 1 Flot1
1448560_at 1.21630 BH3 interacting domain death agonist Bid
1448561_at 0.92661 neutrophil cytosolic factor 2 Ncf2
1448562_at 1.18752 uridine phosphorylase 1 Upp1
1448563_at 0.93275 prohibitin Phb
1448564_at 1.08195 calcium and integrin binding 1 (calmyrin) Cib1
1448565_at 0.99883 "protein phosphatase 1, regulatory (inhibitor) subunit 11" Ppp1r11
1448566_at 0.66616 "solute carrier family 40 (iron-regulated transporter), member 1" Slc40a1
1448567_at 0.96157 transmembrane protein 115 Tmem115
1448568_a_at 1.27784 "solute carrier family 20, member 1" Slc20a1
1448569_at 0.89725 "CD8 antigen, beta chain 1" Cd8b1
1448570_at 1.01001 "glia maturation factor, beta" Gmfb
1448571_a_at 1.12529 "glia maturation factor, beta" Gmfb
1448572_at 0.87707 prolactin-like protein A Prlpa
1448573_a_at 1.10835 CEA-related cell adhesion molecule 10 Ceacam10
1448574_at 1.10809 "expressed in non-metastatic cells 6, protein" Nme6
1448575_at 0.87707 interleukin 7 receptor Il7r
1448576_at 1.06400 interleukin 7 receptor Il7r
1448577_x_at 1.07663 synaptogyrin 2 Syng2
1448578_at 0.91133 "platelet-activating factor acetylhydrolase, isoform 1b, beta1 subunit" Pafah1b1
1448579_at 0.91513 golgi apparatus protein 1 Glg1
1448580_at 0.95933 golgi apparatus protein 1 Glg1
1448581_at 0.99149 zinc finger protein 346 Zfp346
1448582_at 1.01748 "catenin, beta like 1" Ctnnb1
1448583_at 0.99451 expressed sequence AA960436 AA960436
1448584_at 0.88458 arginine/serine-rich coiled-coil 1 Rsrc1
1448585_at 1.02957 "general transcription factor II H, polypeptide 4" Gtf2h4
1448586_at 1.03860 heat shock protein 14 Hspa14
1448587_at 0.90520 "TBC1 domain family, member 10a" Tbc1d10a
1448588_at 0.91165 CD320 antigen Cd320
1448589_at 0.98942 "NADH dehydrogenase (ubiquinone) 1 beta subcomplex, 5" Ndufb5
1448590_at 1.06370 "procollagen, type VI, alpha 1" Col6a1
1448591_at 1.00533 cathepsin S Ctss
1448592_at 0.85980 cartilage associated protein Crtap
1448593_at 0.99193 WNT1 inducible signaling pathway protein 1 Wispl

1448594_at 0.95658 WNT1 inducible signaling pathway protein 1 Wisp1
1448595_a_at 1.01734 brain expressed gene 1 Bex1
1448596_at 0.93279 "solute carrier family 6 (neurotransmitter transporter, creatine), member 8"
Slc6a8
1448597_at 1.04202 "cleavage stimulation factor, 3' pre-RNA, subunit 1" Cstf1
1448598_at 0.95440 matrix metalloproteinase 17 Mmp17
1448599_s_at 1.03073 "DNA segment, Chr 4, Wayne State University 114, expressed" D4Wsu114e
1448600_s_at 0.95666 vav 3 oncogene Vav3
1448601_s_at 0.87005 "homeo box, msh-like 1" Msx1
1448602_at 0.96614 muscle glycogen phosphorylase Pygm
1448603_at 0.91895 serine/arginine-rich protein specific kinase 2 Srpk2
1448604_at 1.14844 uridine-cytidine kinase 2 Uck2
1448605_at 1.00755 "ras homolog gene family, member C" Rhoc
1448606_at 0.87558 "endothelial differentiation, lysophosphatidic acid G-protein-coupled receptor,
2" Edg2
1448607_at 1.12957 --- ---
1448608_at 1.03847 decidual/trophoblast prolactin-related protein Dtprp
1448609_at 0.92176 "thiosulfate sulfurtransferase, mitochondrial" Tst
1448610_a_at 0.97007 "superoxide dismutase 2, mitochondrial" Sod2
1448611_at 0.89762 "WW, C2 and coiled-coil domain containing 2" Wwc2
1448612_at 0.93909 stratifin Sfn
1448613_at 0.92011 extracellular matrix protein 1 Ecm1
1448614_at 0.85086 "5'-nucleotidase, cytosolic II" Nt5c2
1448615_at 1.00761 copper chaperone for superoxide dismutase Ccs
1448616_at 0.96585 "dishevelled 2, dsh homolog (Drosophila)" Dvl2
1448617_at 1.08529 CD53 antigen Cd53
1448618_at 1.23183 major vault protein Mvp
1448619_at 1.13700 7-dehydrocholesterol reductase Dhcr7
1448620_at 1.04921 "Fc receptor, IgG, low affinity III" Fcgr3
1448621_a_at 0.79141 "sphingomyelin phosphodiesterase 1, acid lysosomal" Smpd1
1448622_at 1.09668 "LSM4 homolog, U6 small nuclear RNA associated (S. cerevisiae)" Lsm4
1448623_at 0.96742 transmembrane protein 123 Tmem123
1448624_at 1.07843 CD2 antigen (cytoplasmic tail) binding protein 2 Cd2bp2
1448625_at 0.92290 "golgi autoantigen, golgin subfamily a, 2" Golga2
1448626_at 1.01349 CDK5 regulatory subunit associated protein 1 Cdk5rap1
1448627_s_at 0.96228 PDZ binding kinase Pbk
1448628_at 0.91434 secretogranin III Scg3
1448629_at 0.89600 Hermansky-Pudlak syndrome 4 homolog (human) Hps4
1448630_a_at 0.93986 "succinate dehydrogenase complex, subunit C, integral membrane protein"
Sdhc
1448631_a_at 0.81786 homeodomain interacting protein kinase 2 Hipk2
1448632_at 1.40561 "proteasome (prosome, macropain) subunit, beta type 10" Psmb10
1448633_at 1.10239 PRP31 pre-mRNA processing factor 31 homolog (yeast) Prpf31

1448634_at 0.97499 ralA binding protein 1 Ralbp1
1448635_at 1.02125 structural maintenance of chromosomes 2 Smc2
1448636_at 1.06535 myozenin 1 Myoz1
1448637_at 0.88236 "mediator of RNA polymerase II transcription, subunit 25 homolog (yeast)"
Med25
1448638_at 1.03954 "Mdm2, transformed 3T3 cell double minute p53 binding protein" Mtbp
1448639_a_at 0.95404 spermatogenesis associated 5 Spata5
1448640_at 0.98760 "solute carrier family 14 (urea transporter), member 1" Slc14a1
1448641_at 1.04500 mbt domain containing 1 Mbtd1
1448642_at 1.03594 poly(rC) binding protein 1 Pcbp1
1448643_at 0.98597 Sjogren's syndrome nuclear autoantigen 1 Ssna1
1448644_at 0.98031 penta-EF hand domain containing 1 Pef1
1448645_at 0.93344 male-specific lethal-3 homolog 1 (Drosophila) Msl31
1448646_at 1.24785 WD repeat domain 12 Wdr12
1448647_at 1.06104 "mannosidase 2, alpha 1" Man2a1
1448648_at 0.70654 RIKEN cDNA 9130005N14 gene 9130005N14Rik
1448649_at 0.56464 glutamyl aminopeptidase Enpep
1448650_a_at 0.98904 "polymerase (DNA directed), epsilon" Pole
1448651_at 1.06931 nudix (nucleoside diphosphate linked moiety X)-type motif 5 Nudt5
1448652_at 1.05589 intraflagellar transport 88 homolog (Chlamydomonas) Ift88
1448653_at 1.15468 embryonic ectoderm development Eed
1448654_at 1.07139 mitochondrial carrier homolog 2 (C. elegans) Mtch2
1448655_at 0.78334 low density lipoprotein receptor-related protein 1 Lrp1
1448656_at 0.82449 "calcium channel, voltage-dependent, beta 3 subunit" Cacnb3
1448657_a_at 1.00026 "DnaJ (Hsp40) homolog, subfamily B, member 10" Dnajb10
1448658_at 1.06490 squamous cell carcinoma antigen recognized by T-cells 1 Sart1
1448659_at 1.00346 caspase 7 Casp7
1448660_at 1.04470 Rho GDP dissociation inhibitor (GDI) gamma Arhgdig
1448661_at 0.90424 "phospholipase C, beta 3" Plcb3
1448662_at 0.79108 frizzled homolog 6 (Drosophila) Fzd6
1448663_s_at 1.55021 mevalonate (diphospho) decarboxylase Mvd
1448664_a_at 0.54146 SPEG complex locus Speg
1448665_at 0.96771 "dystrophin, muscular dystrophy" Dmd
1448666_s_at 0.99548 "transducer of ERBB2, 2" Tob2
1448667_x_at 0.99900 "transducer of ERBB2, 2" Tob2
1448668_a_at 1.13526 interleukin-1 receptor-associated kinase 1 Irak1
1448669_at 0.97343 dickkopf homolog 3 (Xenopus laevis) Dkk3
1448670_at 0.93738 "ubiquitin-conjugating enzyme E2E 3, UBC4/5 homolog (yeast)" Ube2e3
1448671_at 0.80158 "ubiquitin-conjugating enzyme E2E 3, UBC4/5 homolog (yeast)" Ube2e3
1448672_a_at 0.89829 zinc finger protein 289 Zfp289
1448673_at 0.80910 poliovirus receptor-related 3 Pvr13
1448674_at 0.95985 ring finger protein 25 Rnf25
1448675_at 0.89510 "DNA segment, Chr 1, ERATO Doi 161, expressed" D1Ert161e

1448676_at	0.69944	"calcium/calmodulin-dependent protein kinase II, beta"	Camk2b
1448677_at	0.91271	COX4 neighbor	Cox4nb
1448678_at	0.92155	RIKEN cDNA 3110048E14 gene	3110048E14Rik
1448679_at	0.95148	hyaluronoglucosaminidase 2	Hyal2
1448680_at	0.89286	"serine (or cysteine) peptidase inhibitor, clade A, member 1c"	Serpina1c
1448681_at	0.87132	"interleukin 15 receptor, alpha chain"	Il15ra
1448682_at	0.94707	dynein light chain LC8-type 1	Dynl11
1448683_at	0.95336	"cytochrome P450, family 2, subfamily d, polypeptide 26"	Cyp2d26
1448684_at	1.00187	"protein phosphatase 1, regulatory (inhibitor) subunit 2"	Ppp1r2
1448685_at	1.06562	RIKEN cDNA 2900010M23 gene	2900010M23Rik
1448686_at	0.95610	interleukin 16	Il16
1448687_at	1.00591	C1q domain containing 2	C1qdc2
1448688_at	1.04260	podocalyxin-like	Podxl
1448689_at	0.89296	related RAS viral (r-ras) oncogene homolog 2	Rras2
1448690_at	0.97211	"potassium channel, subfamily K, member 1"	Kcnk1
1448691_at	1.06671	ubiquilin 4	Ubqln4
1448692_at	1.05344	ubiquilin 4	Ubqln4
1448693_at	0.94964	cDNA sequence BC002216	BC002216
1448694_at	0.66826	Jun oncogene	Jun
1448695_at	0.99340	"protein kinase C, iota"	Prkci
1448696_at	0.99633	hephaestin	Heph
1448697_s_at	1.13230	ribosomal protein L36a-like	Rpl36al
1448698_at	1.46478	cyclin D1	Ccnd1
1448699_at	1.11046	mitochondrial ribosomal protein L54	Mrpl54
1448700_at	0.96911	G0/G1 switch gene 2	G0s2
1448701_a_at	1.07222	"KRIT1, ankyrin repeat containing"	Krit1
1448702_at	0.97541	immediate early response 3 interacting protein 1	Ier3ip1
1448703_at	1.01165	"LSM8 homolog, U6 small nuclear RNA associated (S. cerevisiae)"	Lsm8
1448704_s_at	0.97083	histocompatibility 47	H47
1448705_at	0.90226	zinc finger and BTB domain containing 22	Zbtb22
1448706_at	0.97770	Traf and Tnf receptor associated protein	Ttrap
1448707_at	1.01398	"TAF13 RNA polymerase II, TATA box binding protein (TBP)-associated factor"	Taf13
1448708_at	0.98852	peroxisome proliferator activated receptor binding protein	Pparbp
1448709_at	0.97296	AT rich interactive domain 1A (Swi1 like) /// similar to AT rich interactive domain 1A isoform a /// similar to AT rich interactive domain 1A isoform a	Arid1a /// LOC669477 /// LOC675933
1448710_at	1.00386	chemokine (C-X-C motif) receptor 4	Cxcr4
1448711_at	1.02748	minichromosome maintenance deficient 3 (S. cerevisiae) associated protein	Mcm3ap
1448712_at	0.98071	choroideremia	Chm
1448713_at	1.15866	signal transducer and activator of transcription 4	Stat4
1448714_at	0.97887	RNA guanylyltransferase and 5'-phosphatase	Rngtt

1448715_x_at 1.15067 SH3-domain GRB2-like (endophilin) interacting protein 1 Sgip1
1448716_at 1.02006 "hemoglobin X, alpha-like embryonic chain in Hba complex" Hba-x
1448717_at 0.83089 glutaryl-Coenzyme A dehydrogenase Gcdh
1448718_at 1.03059 RIKEN cDNA 2400001E08 gene 2400001E08Rik
1448719_at 0.89704 "transcriptional adaptor 3 (NGG1 homolog, yeast)-like" Tada3l
1448720_at 1.00301 leucine rich repeat containing 40 Lrrc40
1448721_at 0.90719 "DNA segment, Chr 1, ERATO Doi 622, expressed" D1Ertd622e
1448722_s_at 1.00186 phosphopantothenoylcysteine synthetase Ppcs
1448723_at 0.97948 retinol dehydrogenase 7 Rdh7
1448724_at 0.86006 cytokine inducible SH2-containing protein Cish
1448725_at 0.94919 poly (ADP-ribose) glycohydrolase Parg
1448726_at 0.94257 "small nuclear RNA activating complex, polypeptide 2" Snapc2
1448727_at 0.89488 "transducin-like enhancer of split 6, homolog of Drosophila E(spl)" Tle6
1448728_a_at 1.22680 "nuclear factor of kappa light polypeptide gene enhancer in B-cells inhibitor, zeta" Nfkbiz
1448729_a_at 1.06629 septin 4 4-Sep
1448730_at 1.07285 "carboxypeptidase A3, mast cell" Cpa3
1448731_at 1.02750 "interleukin 10 receptor, alpha" Il10ra
1448732_at 0.94700 cathepsin B Ctsb
1448733_at 0.83121 B lymphoma Mo-MLV insertion region 1 Bmi1
1448734_at 1.04203 ceruloplasmin Cp
1448735_at 0.84127 ceruloplasmin Cp
1448736_a_at 1.10267 hypoxanthine guanine phosphoribosyl transferase 1 Hprt1
1448737_at 0.65006 tetraspanin 7 Tspan7
1448738_at 0.93119 calbindin-28K Calb1
1448739_x_at 1.03419 ribosomal protein S18 Rps18
1448740_at 1.10216 RIKEN cDNA 2400006H24 gene 2400006H24Rik
1448741_at 0.87532 "solute carrier family 3, member 1" Slc3a1
1448742_at 0.90049 snail homolog 1 (Drosophila) Snai1
1448743_at 0.83721 "synovial sarcoma, X breakpoint 2 interacting protein" Ssx2ip
1448744_at 0.95649 galactosamine (N-acetyl)-6-sulfate sulfatase Galns
1448745_s_at 0.91430 lorcin Lor
1448746_at 0.97897 nibrin Nbn
1448747_at 0.79567 F-box only protein 32 Fbxo32
1448748_at 0.99714 pleckstrin Plek
1448749_at 1.01075 pleckstrin Plek
1448750_at 0.92660 heart and neural crest derivatives expressed transcript 1 Hand1
1448751_at 0.76842 "adaptor-related protein complex 3, mu 2 subunit" Ap3m2
1448752_at 0.64838 carbonic anhydrase 2 Car2
1448753_at 1.09113 signal recognition particle 9 Srp9
1448754_at 0.98470 "retinol binding protein 1, cellular" Rbp1
1448755_at 0.80651 "procollagen, type XV" Col15a1
1448756_at 1.03421 S100 calcium binding protein A9 (calgranulin B) S100a9

1448757_at 0.91888 promyelocytic leukemia Pml
 1448758_at 0.86015 similar to nuclear receptor binding factor 2 /// similar to nuclear receptor binding factor 2 /// nuclear receptor binding factor 2 LOC627232 /// LOC627424 /// Nr1h2
 1448759_at 0.86640 "interleukin 2 receptor, beta chain" Il2rb
 1448760_at 0.81464 zinc finger protein 68 Zfp68
 1448761_a_at 0.91714 "coatamer protein complex, subunit gamma 2" Copg2
 1448762_at 0.95591 RAD17 homolog (S. pombe) Rad17
 1448763_at 0.91518 "ATPase family, AAA domain containing 1" Atad1
 1448764_a_at 1.11358 "fatty acid binding protein 1, liver" Fabp1
 1448765_at 0.82284 Fyn proto-oncogene Fyn
 1448766_at 0.95086 gap junction membrane channel protein beta 1 Gjb1
 1448767_s_at 0.96598 gap junction membrane channel protein beta 1 Gjb1
 1448768_at 0.91955 myelin oligodendrocyte glycoprotein Mog
 1448769_at 1.03164 "solute carrier family 35, member B1" Slc35b1
 1448770_a_at 0.90757 ATPase inhibitory factor 1 Atpif1
 1448771_a_at 1.02773 ferritin heavy chain 1 Fth1
 1448772_at 0.99310 "ubiquitin-conjugating enzyme E2A, RAD6 homolog (S. cerevisiae)" Ube2a
 1448773_at 1.17850 ribosomal protein L11 /// similar to 60S ribosomal protein L11 /// similar to 60S ribosomal protein L11 /// hypothetical gene supported by AK011800; BC025077; NM_025919 /// similar to 60S ribosomal protein L11 Rpl11 /// LOC194197 /// LOC235424 /// LOC328825 /// LOC665330
 1448774_at 1.14880 stomatin (Epb7.2)-like 2 Stoml2
 1448775_at 1.36161 interferon activated gene 203 Ifi203
 1448776_at 1.03850 "growth arrest and DNA-damage-inducible, gamma interacting protein 1" Gadd45gip1
 1448777_at 0.97096 minichromosome maintenance deficient 2 mitotin (S. cerevisiae) Mcm2
 1448778_at 0.95290 "splicing factor, arginine/serine-rich 4 (SRp75)" Sfrs4
 1448779_at 1.09168 CDKN1A interacting zinc finger protein 1 Ciz1
 1448780_at 1.13937 "solute carrier family 12, member 2" Slc12a2
 1448781_at 1.06752 Ngfi-A binding protein 1 Nab1
 1448782_at 1.00415 thioredoxin domain containing 11 Txndc11
 1448783_at 0.97602 "solute carrier family 7 (cationic amino acid transporter, y+ system), member 9" Slc7a9
 1448784_at 1.01607 "TAF10 RNA polymerase II, TATA box binding protein (TBP)-associated factor" Taf10
 1448785_at 1.01235 "runt-related transcription factor 1; translocated to, 1 (cyclin D-related)" Runx1t1
 1448786_at 1.01143 RIKEN cDNA 1100001H23 gene 1100001H23Rik
 1448787_at 0.94295 modulator of apoptosis 1 Moap1
 1448788_at 1.08971 Cd200 antigen Cd200
 1448789_at 0.89178 "aldehyde dehydrogenase family 1, subfamily A3" Aldh1a3
 1448790_at 0.88668 "sema domain, transmembrane domain (TM), and cytoplasmic domain, (semaphorin) 6B" Sema6b

1448791_at 1.01124 sorting nexin 5 Snx5
1448792_a_at 0.95788 "cytochrome P450, family 2, subfamily f, polypeptide 2" Cyp2f2
1448793_a_at 1.17304 syndecan 4 Sdc4
1448794_s_at 1.02773 "DnaJ (Hsp40) homolog, subfamily C, member 2" Dnajc2
1448795_a_at 1.05685 transforming growth factor beta regulated gene 4 Tbrg4
1448796_s_at 1.07062 transforming growth factor beta regulated gene 4 Tbrg4
1448797_at 1.08781 "ELK3, member of ETS oncogene family" Elk3
1448798_at 0.90310 ESP8-like 3 Eps8l3
1448799_s_at 1.01514 mitochondrial ribosomal protein S12 Mrps12
1448800_at 0.93163 reticulon 4 interacting protein 1 Rtn4ip1
1448801_a_at 0.92202 translocase of inner mitochondrial membrane 44 Timm44
1448802_at 1.12862 nuclear fragile X mental retardation protein interacting protein 1 Nufip1
1448803_at 1.04493 "golgi autoantigen, golgin subfamily a, 4" Golga4
1448804_at 0.91127 "cytochrome P450, family 11, subfamily a, polypeptide 1" Cyp11a1
1448805_at 0.99633 upstream transcription factor 1 Usf1
1448806_at 0.96707 matrix metalloproteinase 24 Mmp24
1448807_at 0.98875 histamine receptor H 3 Hrh3
1448808_a_at 1.08585 "expressed in non-metastatic cells 2, protein" Nme2
1448809_at 1.05206 chromosome segregation 1-like (S. cerevisiae) Cse11
1448810_at 1.00469 glucosamine Gne
1448811_at 1.06640 mitochondrial ribosomal protein L2 Mrpl2
1448812_at 1.00314 hippocalcin-like 1 Hpcal1
1448813_at 0.97090 arylacetamide deacetylase (esterase) Aadac
1448814_at 0.83534 growth factor receptor bound protein 2-associated protein 1 Gab1
1448815_at 0.95140 8-oxoguanine DNA-glycosylase 1 Ogg1
1448816_at 1.08330 prostaglandin I2 (prostacyclin) synthase Ptgis
1448817_at 0.95279 "OTU domain, ubiquitin aldehyde binding 1" Otub1
1448818_at 1.01367 wingless-related MMTV integration site 5A Wnt5a
1448819_at 1.00404 "eukaryotic translation initiation factor 2, subunit 2 (beta)" Eif2s2
1448820_a_at 1.10442 "eukaryotic translation initiation factor 2, subunit 2 (beta)" Eif2s2
1448821_at 0.90604 tyrosinase Tyr
1448822_at 1.13053 "proteasome (prosome, macropain) subunit, beta type 6" Psmb6
1448823_at 0.62411 chemokine (C-X-C motif) ligand 12 Cxcl12
1448824_at 0.94156 "ubiquitin-conjugating enzyme E2, J1" Ube2j1
1448825_at 0.62847 "pyruvate dehydrogenase kinase, isoenzyme 2" Pdk2
1448826_at 0.92778 "myosin, heavy polypeptide 6, cardiac muscle, alpha" Myh6
1448827_s_at 1.03216 "myosin, heavy polypeptide 6, cardiac muscle, alpha /// myosin, heavy polypeptide 7, cardiac muscle, beta /// similar to myosin, heavy polypeptide 7, cardiac muscle, beta" Myh6 /// Myh7 /// LOC674761
1448828_at 1.03803 structural maintenance of chromosomes 6 Smc6
1448829_at 1.01612 structural maintenance of chromosomes 6 Smc6
1448830_at 0.83382 dual specificity phosphatase 1 Dusp1
1448831_at 1.15528 angiopoietin 2 Angpt2

1448832_at 0.95919 complexin 1 Cplx1
1448833_at 1.01397 forkhead box M1 Foxm1
1448834_at 1.04647 forkhead box M1 Foxm1
1448835_at 1.12083 E2F transcription factor 6 E2f6
1448836_s_at 1.12846 "ATP binding domain 1 family, member B" Atpbd1b
1448837_at 0.92941 villin 1 Vill
1448838_at 0.92797 "topoisomerase I binding, arginine/serine-rich" Topors
1448839_at 0.99749 ankyrin repeat domain 47 Ankrd47
1448840_at 0.95888 transmembrane and ubiquitin-like domain containing 1 Tmub1
1448841_at 0.99352 pore forming protein-like Pfpl
1448842_at 1.01040 "cysteine dioxygenase 1, cytosolic" Cdo1
1448843_at 1.19158 "similar to signal sequence receptor, alpha" LOC636537
1448844_at 1.15027 cytochrome b5 type B Cyb5b
1448845_at 1.10236 ribonuclease P 25 subunit (human) Rpp25
1448846_a_at 1.16879 ribosomal protein L29 /// similar to 60S ribosomal protein L29 /// similar to 60S ribosomal protein L29 /// similar to 60S ribosomal protein L29 /// similar to 60S ribosomal protein L29 /// similar to 60S ribosomal protein L29 /// similar to 60S ribosomal protein L29 /// similar to 60S ribosomal protein L29 /// similar to 60S ribosomal protein L29 /// similar to 60S ribosomal protein L29 /// similar to 60S ribosomal protein L29 (P23) /// similar to 60S ribosomal protein L29 /// similar to 60S ribosomal protein L29 Rpl29 /// LOC433350 /// LOC433706 /// LOC433941 /// LOC622707 /// LOC626772 /// LOC665032 /// LOC666642 /// LOC667466 /// LOC669999 /// LOC670211 /// LOC670334 /// LOC675846
1448847_at 0.95178 serine incorporator 3 Serinc3
1448848_at 1.01142 "torsin family 1, member B" Tor1b
1448849_at 1.04120 mitochondrial ribosomal protein L40 Mrpl40
1448850_a_at 1.13465 "DnaJ (Hsp40) homolog, subfamily C, member 5" Dnajc5
1448851_a_at 1.10297 "DnaJ (Hsp40) homolog, subfamily C, member 5" Dnajc5
1448852_at 0.96889 regucalcin Rgn
1448853_at 1.06530 synaptojanin 2 binding protein Synj2bp
1448854_s_at 0.97786 "murinoglobulin, pseudogene 1 /// murinoglobulin 1 /// murinoglobulin 2 /// similar to murinoglobulin 1 /// similar to murinoglobulin 1" Mug-ps1 /// Mug1 /// Mug2 /// LOC434083 /// LOC669343
1448855_at 1.12297 Ras association (RalGDS/AF-6) domain family 1 Rassf1
1448856_a_at 0.87543 methionine sulfoxide reductase A Msra
1448857_a_at 0.98943 "zinc finger, DHHC domain containing 12" Zdhhc12
1448858_at 0.82440 Unc-51 like kinase 2 (C. elegans) Ulk2
1448859_at 0.98122 chemokine (C-X-C motif) ligand 13 Cxcl13
1448860_at 0.66352 rad and gem related GTP binding protein 2 Rem2
1448861_at 0.98149 Tnf receptor-associated factor 5 /// similar to TNF receptor-associated factor 5 Traf5 /// LOC622602
1448862_at 1.05690 intercellular adhesion molecule 2 Icam2
1448863_a_at 1.08774 "tumor necrosis factor, alpha-induced protein 1 (endothelial)" Tnfaip1
1448864_at 0.96741 SNF related kinase Snrk

1448865_at	1.47380 hydroxysteroid (17-beta) dehydrogenase 7	Hsd17b7
1448866_at	1.05938 SUMO/sentrin specific peptidase 3	Senp3
1448867_at	0.92293 "TMEM9 domain family, member B"	Tmem9b
1448868_at	1.02793 SCAN domain-containing 1	Scand1
1448869_a_at	1.00314 mitochondrial ribosomal protein S16	Mrps16
1448870_at	0.84621 latent transforming growth factor beta binding protein 1	Ltbp1
1448871_at	1.00337 mitogen activated protein kinase 13	Mapk13
1448872_at	1.00387 regenerating islet-derived 3 gamma	Reg3g
1448873_at	0.89068 occludin	Ocln
1448874_a_at	0.98843 SH2B adaptor protein 1	Sh2b1
1448875_at	0.89346 zinc fingers and homeoboxes protein 1	Zhx1
1448876_at	0.96159 Ellis van Creveld gene homolog (human)	Evc
1448877_at	1.05183 distal-less homeobox 2	Dlx2
1448878_at	0.89948 Max dimerization protein 3	Mxd3
1448879_at	1.14779 ubiquitin-conjugating enzyme E2L 3	Ube2l3
1448880_at	1.09156 ubiquitin-conjugating enzyme E2L 3	Ube2l3
1448881_at	1.08273 haptoglobin	Hp
1448882_at	1.00483 transmembrane protein 93	Tmem93
1448883_at	0.87029 legumain	Lgmn
1448884_at	1.10208 "general transcription factor II E, polypeptide 2 (beta subunit)"	Gtf2e2
1448885_at	1.05930 "RAP2B, member of RAS oncogene family"	Rap2b
1448886_at	1.23528 GATA binding protein 3	Gata3
1448887_x_at	1.06232 fractured callus expressed transcript 1	Fxc1
1448888_at	0.95707 "protein phosphatase 1, regulatory (inhibitor) subunit 7"	Ppp1r7
1448889_at	0.94823 "solute carrier family 38, member 4"	Slc38a4
1448890_at	0.67991 Kruppel-like factor 2 (lung)	Klf2
1448891_at	0.88996 macrophage scavenger receptor 2	Msr2
1448892_at	1.04261 dedicator of cytokinesis 7	Dock7
1448893_at	0.82879 nuclear receptor co-repressor 2	Ncor2
1448894_at	1.02621 "aldo-keto reductase family 1, member B8"	Akr1b8
1448895_a_at	0.93535 "catenin (cadherin associated protein), alpha 2"	Ctnna2
1448896_at	0.98630 "phosphatidylinositol glycan anchor biosynthesis, class F"	Pigf
1448897_at	0.82371 "makorin, ring finger protein, 2"	Mkrn2
1448898_at	1.15998 chemokine (C-C motif) ligand 9	Ccl9
1448899_s_at	0.98528 RAD51 associated protein 1	Rad51ap1
1448900_at	1.04110 "DNA segment, Chr 16, human D22S680E, expressed"	D16H22S680E
1448901_at	0.92097 carboxypeptidase X 1 (M14 family)	Cpxm1
1448902_at	0.93873 tetratricopeptide repeat domain 23	Ttc23
1448903_at	0.97756 selenoprotein 15-Sep	
1448904_at	0.75656 "DNA segment, Chr 6, Wayne State University 176, expressed"	D6Wsu176e
1448905_at	1.02848 expressed in non-metastatic cells 3	Nme3
1448906_at	0.58456 cadherin 16	Cdh16
1448907_at	0.86548 thimet oligopeptidase 1	Thop1

1448908_at 0.75860 phosphatidic acid phosphatase type 2B Ppap2b
1448909_a_at 1.06824 mitochondrial ribosomal protein L39 Mrpl39
1448910_at 1.07143 peroxisomal trans-2-enoyl-CoA reductase Pecr
1448911_at 1.19246 "ATPase, H⁺/K⁺ exchanging, beta polypeptide" Atp4b
1448912_at 0.82442 C1q and tumor necrosis factor related protein 1 C1qtnf1
1448913_at 0.91652 "SWI/SNF related, matrix associated, actin dependent regulator of chromatin, subfamily d, member 1" Smarcd1
1448914_a_at 1.38730 colony stimulating factor 1 (macrophage) Csf1
1448915_at 0.90618 zinc finger protein 524 Zfp524
1448916_at 1.18876 "v-maf musculoaponeurotic fibrosarcoma oncogene family, protein G (avian)" Mafg
1448917_at 1.01992 thyroid hormone receptor associated protein 6 Thrap6
1448918_at 1.06283 "solute carrier organic anion transporter family, member 3a1" Slco3a1
1448919_at 0.99902 CD302 antigen Cd302
1448920_at 1.04254 zinc finger protein 535 Zfp535
1448921_a_at 1.03503 mitochondrial ribosomal protein S9 Mrps9
1448922_at 1.06504 dual specificity phosphatase 19 Dusp19
1448923_at 1.04190 "protein kinase, interferon inducible double stranded RNA dependent activator" Prkra
1448924_at 0.99622 RIKEN cDNA 4432406C05 gene 4432406C05Rik
1448925_at 0.81421 twist homolog 2 (Drosophila) Twist2
1448926_at 0.86579 homeo box A5 Hoxa5
1448927_at 0.86925 "potassium intermediate/small conductance calcium-activated channel, subfamily N, member 2" Kcnn2
1448928_at 0.96301 histone deacetylase 6 /// similar to Histone deacetylase 6 (HD6) (Histone deacetylase mHDA2) Hdac6 /// LOC669168
1448929_at 0.94676 "coagulation factor XIII, A1 subunit" F13a1
1448930_at 0.90578 RIKEN cDNA 3010026O09 gene 3010026O09Rik
1448931_at 0.87203 coagulation factor II (thrombin) receptor-like 1 F2rl1
1448932_at 0.92182 keratin 16 Krt16
1448933_at 0.71348 protocadherin beta 17 Pcdhb17
1448934_at 0.97934 NADH dehydrogenase (ubiquinone) 1 alpha subcomplex 10 Ndufa10
1448935_at 0.89832 "chromodomain protein, Y chromosome-like" Cdyl
1448936_at 0.96398 syntaxin 12 Stx12
1448937_at 0.96001 "solute carrier family 35, member B3" Slc35b3
1448938_at 0.96649 replication protein A3 Rpa3
1448939_at 0.85972 ubiquitin specific peptidase 25 Usp25
1448940_at 0.94974 tripartite motif protein 21 Trim21
1448941_at 0.94611 "UDP-Gal:betaGlcNAc beta 1,4- galactosyltransferase, polypeptide 2" B4galt2
1448942_at 0.85567 "guanine nucleotide binding protein (G protein), gamma 11" Gng11
1448943_at 0.56573 neuropilin 1 Nrp1
1448944_at 0.54203 neuropilin 1 Nrp1

1448945_at 1.06424 plasma membrane proteolipid Pllp
1448946_at 0.96583 kinesin family member 3C Kif3c
1448947_at 0.99337 RIKEN cDNA 2810004N23 gene 2810004N23Rik
1448948_at 0.92565 recombination activating gene 1 activating protein 1 Rag1ap1
1448949_at 1.08167 carbonic anhydrase 4 Car4
1448950_at 0.90121 "interleukin 1 receptor, type I" Il1r1
1448951_at 1.07672 "tumor necrosis factor receptor superfamily, member 1b" Tnfrsf1b
1448952_at 0.81979 RIKEN cDNA A030009H04 gene A030009H04Rik
1448953_at 1.17539 Bloom syndrome homolog (human) Blm
1448954_at 0.90349 nuclear receptor interacting protein 3 Nrip3
1448955_s_at 0.95689 Ca²⁺dependent activator protein for secretion Cadps
1448956_at 0.75399 START domain containing 10 Stard10
1448957_at 1.06306 recombining binding protein suppressor of hairless (Drosophila) Rbpsuh
1448958_at 0.96180 interferon alpha responsive gene Ifrg15
1448959_at 1.07532 NADH dehydrogenase (ubiquinone) Fe-S protein 4 Ndufs4
1448960_at 0.59192 CXXC finger 5 Cxxc5
1448961_at 1.12433 phospholipid scramblase 2 Plscr2
1448962_at 0.91906 "myosin, heavy polypeptide 11, smooth muscle" Myh11
1448963_at 0.92431 nuclear transcription factor-Y gamma Nfyc
1448964_at 1.04891 S100 calcium binding protein G S100g
1448965_at 0.80634 INO80 complex homolog 1 (S. cerevisiae) Inoc1
1448966_a_at 0.79249 nuclear factor of activated T-cells 5 Nfat5
1448967_at 0.89283 nipsnap homolog 3A (C. elegans) Nipsnap3a
1448968_at 0.99884 "DNA segment, Chr 7, Wayne State University 128, expressed" D7Wsu128e
1448969_at 0.88869 FtsJ homolog 2 (E. coli) Ftsj2
1448970_at 0.99391 RIKEN cDNA 1200007B05 gene 1200007B05Rik
1448971_at 1.23948 RIKEN cDNA 2410022L05 gene 2410022L05Rik
1448972_at 0.94887 "glutamate receptor, ionotropic, AMPA1 (alpha 1)" Gria1
1448973_at 1.05036 "sulfotransferase family 1D, member 1" Sult1d1
1448974_at 0.93691 doublecortin Dcx
1448975_s_at 1.05292 renin 1 structural /// renin 2 tandem duplication of Ren1 Ren1 /// Ren2
1448976_at 0.99171 tuftelin interacting protein 11 Tfip11
1448977_at 1.02070 "transcription factor AP-2, gamma" Tcfap2c
1448978_at 0.84114 neuronal guanine nucleotide exchange factor Ngef
1448979_at 0.92704 muted Muted
1448980_at 1.09574 ghrelin Ghrl
1448981_x_at 0.94745 "T-cell leukemia/lymphoma 1B, 1" Tcl1b1
1448982_at 0.98698 kallikrein 6 Klk6
1448983_at 1.05331 CMT1A duplicated region transcript 4 Cdrt4
1448984_at 0.94024 "excision repair cross-complementing rodent repair deficiency, complementation group 4" Ercc4
1448985_at 0.99384 dual specificity phosphatase 22 Dusp22
1448986_x_at 0.94206 deoxyribonuclease II alpha Dnase2a

1448987_at 0.96360 "acyl-Coenzyme A dehydrogenase, long-chain" Acadl
1448988_at 0.96622 "acyl-Coenzyme A dehydrogenase, long-chain" Acadl
1448989_a_at 1.02845 myosin IB Myo1b
1448990_a_at 1.12797 myosin IB Myo1b
1448991_a_at 0.92993 "internexin neuronal intermediate filament protein, alpha" Ina
1448992_at 1.02805 "internexin neuronal intermediate filament protein, alpha" Ina
1448993_at 1.03100 autophagy-related 3 (yeast) Atg3
1448994_at 0.94012 trans-acting transcription factor 1 Sp1
1448995_at 1.44155 chemokine (C-X-C motif) ligand 4 Cxcl4
1448996_at 1.02410 rod outer segment membrane protein 1 Rom1
1448997_at 0.96430 "pleckstrin homology, Sec7 and coiled-coil domains 1" Pscd1
1448998_at 1.07254 lactoperoxidase Lpo
1448999_at 0.92576 trafficking protein particle complex 5 Trappc5
1449000_at 0.91982 "DNA segment, Chr 10, Johns Hopkins University 81 expressed" D10Jhu81e
1449001_at 0.84891 isovaleryl coenzyme A dehydrogenase Ivd
1449002_at 0.96060 "pleckstrin homology-like domain, family A, member 3" Phlda3
1449003_a_at 1.04337 vesicle transport through interaction with t-SNAREs 1B homolog Vti1b
1449004_at 1.03538 mitochondrial ribosomal protein L46 Mrpl46
1449005_at 1.52308 "solute carrier family 16 (monocarboxylic acid transporters), member 3" Slc16a3
1449006_at 1.03854 "galactosidase, alpha" Gla
1449007_at 1.08608 B-cell translocation gene 3 /// similar to BTG3 protein (Tob5 protein)
(Abundant in neuroepithelium area protein) /// B-cell translocation gene 3 pseudogene Btg3 ///
LOC640416 /// LOC654432
1449008_at 1.07374 tubby-like protein 3 Tulp3
1449009_at 1.13992 T-cell specific GTPase Tgtp
1449010_at 1.05688 heat shock protein 4 like Hspa4l
1449011_at 0.72663 "solute carrier family 12, member 7" Slc12a7
1449012_s_at 0.82644 fibronectin type III domain containing 4 Fndc4
1449013_at 0.87807 eukaryotic elongation factor-2 kinase Eef2k
1449014_at 0.93871 "lactamase, beta" Lactb
1449015_at 0.91964 resistin like alpha Retnla
1449016_at 0.95455 zona pellucida glycoprotein 2 Zp2
1449017_at 0.86138 nuclear transport factor 2 Nutf2
1449018_at 1.02484 profilin 1 Pfn1
1449019_at 0.98723 A kinase (PRKA) anchor protein 1 Akap1
1449020_at 0.85623 phospholipid scramblase 3 Plscr3
1449021_at 0.94152 ribonuclease P 21 subunit (human) Rpp21
1449022_at 1.07752 nestin Nes
1449023_a_at 1.01589 enhancer of zeste homolog 1 (Drosophila) Ezh1
1449024_a_at 0.92556 hexosaminidase A Hexa
1449025_at 1.43011 interferon-induced protein with tetratricopeptide repeats 3 Ifit3
1449026_at 1.07887 interferon (alpha and beta) receptor 1 Ifnar1
1449027_at 0.98538 "ras homolog gene family, member U" Rhou

1449028_at 0.95319 "ras homolog gene family, member U" Rhou
 1449029_at 0.90160 MAP kinase-interacting serine/threonine kinase 2 Mknk2
 1449030_at 0.90958 synapsin II Syn2
 1449031_at 0.89762 Cbp/p300-interacting transactivator with Glu/Asp-rich carboxy-terminal domain
 1 Cited1
 1449032_at 0.91365 prolactin-like protein M Prlpm
 1449033_at 0.99000 "tumor necrosis factor receptor superfamily, member 11b (osteoprotegerin)"
 Tnfrsf11b
 1449034_at 0.84556 "kallikrein B, plasma 1" Klkb1
 1449035_at 1.00566 RIKEN cDNA 4930544G11 gene 4930544G11Rik
 1449036_at 0.92261 ring finger protein 128 Rnf128
 1449037_at 0.78141 cAMP responsive element modulator Crem
 1449038_at 0.89144 hydroxysteroid 11-beta dehydrogenase 1 Hsd11b1
 1449039_a_at 0.98589 heterogeneous nuclear ribonucleoprotein D-like Hnrpdl
 1449040_a_at 1.06687 selenophosphate synthetase 2 Sephs2
 1449041_a_at 0.91699 thyroid hormone receptor interactor 6 Trip6
 1449042_at 0.92850 CCCTC-binding factor Ctf
 1449043_at 0.87233 "N-acetyl galactosaminidase, alpha" Naga
 1449044_at 1.17771 eukaryotic translation elongation factor 1 epsilon 1 Eef1e1
 1449045_at 1.03408 AFG3(ATPase family gene 3)-like 1 (yeast) Afg3l1
 1449046_a_at 0.93842 Josephin domain containing 2 Josd2
 1449047_at 1.13240 2-hydroxyacyl-CoA lyase 1 Hacl1
 1449048_s_at 0.95138 "RAB4A, member RAS oncogene family" Rab4a
 1449049_at 0.80474 toll-like receptor 1 Tlr1
 1449050_at 0.95762 replication factor C 1 Recc1
 1449051_at 0.92071 peroxisome proliferator activated receptor alpha Ppara
 1449052_a_at 0.91344 DNA methyltransferase 3B Dnmt3b
 1449053_s_at 0.99078 component of oligomeric golgi complex 1 Cog1
 1449054_a_at 0.99762 poly(rC) binding protein 4 Pcbp4
 1449055_x_at 0.92501 poly(rC) binding protein 4 Pcbp4
 1449056_at 0.83915 RIKEN cDNA E330009J07 gene E330009J07Rik
 1449057_at 0.92486 Kell blood group Kel
 1449058_at 0.94270 GLI-Kruppel family member GLI1 Gli1
 1449059_a_at 0.88456 3-oxoacid CoA transferase 1 Oxct1
 1449060_at 1.05857 kinesin family member 2C /// similar to Kinesin-like protein KIF2C (Mitotic
 centromere-associated kinesin) (MCAK) Kif2c /// LOC631653
 1449061_a_at 0.90372 "DNA primase, p49 subunit" Prim1
 1449062_at 0.84747 ketoheksokinase Khk
 1449063_at 0.93037 SEC22 vesicle trafficking protein homolog B (S. cerevisiae) Sec22b
 1449064_at 0.93315 L-threonine dehydrogenase Tdh
 1449065_at 0.89772 acyl-CoA thioesterase 1 Acot1
 1449066_a_at 0.90792 Rho guanine nucleotide exchange factor (GEF7) Arhgef7
 1449067_at 0.93461 "solute carrier family 2 (facilitated glucose transporter), member 2" Slc2a2

1449068_at 1.09335 zinc finger protein 148 Zfp148
1449069_at 1.05837 zinc finger protein 148 Zfp148
1449070_x_at 0.42863 adenomatosis polyposis coli down-regulated 1 Apcdd1
1449071_at 0.90389 "myosin, light polypeptide 7, regulatory" Myl7
1449072_a_at 1.06118 RIKEN cDNA 2510005D08 gene 2510005D08Rik
1449073_at 0.96013 "filamin C, gamma (actin binding protein 280)" Flnc
1449074_at 0.99614 RIKEN cDNA 1700019N12 gene 1700019N12Rik
1449075_at 0.81840 exonuclease 3"-5" domain-like 2 Exdl2
1449076_x_at 0.91293 acireductone dioxygenase 1 Adi1
1449077_at 0.96611 erythroid associated factor Eraf
1449078_at 0.94073 "ST3 beta-galactoside alpha-2,3-sialyltransferase 6" St3gal6
1449079_s_at 0.80268 "ST3 beta-galactoside alpha-2,3-sialyltransferase 6" St3gal6
1449080_at 0.86213 histone deacetylase 2 Hdac2
1449081_at 0.89970 carboxylesterase 3 Ces3
1449082_at 1.07462 microfibrillar associated protein 5 Mfap5
1449083_at 1.08476 coiled-coil domain containing 91 Ccdc91
1449084_s_at 1.01106 SH3 domain protein D19 Sh3d19
1449085_at 1.19137 PHD finger protein 10 Phf10
1449086_at 0.90893 ring finger protein 141 Rnf141
1449087_at 1.02352 ring finger protein 141 Rnf141
1449088_at 0.96947 fructose bisphosphatase 2 Fbp2
1449089_at 0.90973 nuclear receptor interacting protein 1 Nrip1
1449090_a_at 0.99745 Yamaguchi sarcoma viral (v-yes) oncogene homolog 1 Yes1
1449091_at 0.90031 claudin 8 Cldn8
1449092_at 0.88222 "protein phosphatase 1D magnesium-dependent, delta isoform" Ppm1d
1449093_at 0.84654 cardiotrophin 1 Ctf1
1449094_at 0.99928 gap junction membrane channel protein alpha 7 Gja7
1449095_at 1.02753 vacuolar protein sorting 54 (yeast) Vps54
1449096_at 1.01565 RIKEN cDNA 0610011N22 gene 0610011N22Rik
1449097_at 0.91234 thioredoxin reductase 2 Txnrd2
1449098_a_at 1.04577 "polymerase (DNA directed), iota" Poli
1449099_at 0.92209 LPS-responsive beige-like anchor Lrba
1449100_at 0.92656 "par-6 (partitioning defective 6,) homolog alpha (C. elegans)" Pard6a
1449101_at 1.03519 early B-cell factor 2 Ebf2
1449102_at 0.99704 early B-cell factor 2 Ebf2
1449103_at 0.96865 testis expressed gene 101 Tex101
1449104_at 1.03764 uroplakin 3A Upk3a
1449105_at 0.97657 SH2 domain protein 2A Sh2d2a
1449106_at 0.97134 glutathione peroxidase 3 Gpx3
1449107_at 0.90081 nudix (nucleoside diphosphate linked moiety X)-type motif 4 Nudt4
1449108_at 1.01237 ferredoxin 1 Fdx1
1449109_at 0.51181 suppressor of cytokine signaling 2 Socs2
1449110_at 0.79782 "ras homolog gene family, member B" Rhob

1449111_a_at 0.92292 growth factor receptor bound protein 2 Grb2
 1449112_at 0.97499 "solute carrier family 27 (fatty acid transporter), member 5" Slc27a5
 1449113_at 0.88621 GC-rich promoter binding protein 1-like 1 Gpbp111
 1449114_at 1.02728 "serine/threonine kinase 3 (Ste20, yeast homolog)" Stk3
 1449115_at 1.02566 metal response element binding transcription factor 2 Mtf2
 1449116_a_at 1.07284 deoxythymidylate kinase Dtymk
 1449117_at 0.86503 Jun proto-oncogene related gene d1 Jund1
 1449118_at 0.76431 dihydrolipoamide branched chain transacylase E2 Dbt
 1449119_at 1.03034 ariadne homolog 2 (Drosophila) Arih2
 1449120_a_at 1.03036 pericentriolar material 1 Pcm1
 1449121_at 1.09893 FUS interacting protein (serine-arginine rich) 1 Fusip1
 1449122_at 0.91759 RIKEN cDNA 3110003A22 gene 3110003A22Rik
 1449123_at 1.16970 "inter-alpha trypsin inhibitor, heavy chain 3" Itih3
 1449124_at 0.90504 "ral guanine nucleotide dissociation stimulator,-like 1" Rgl1
 1449125_at 0.91350 "tumor necrosis factor, alpha-induced protein 8-like 1" Tnfaip8l1
 1449126_at 0.99567 zinc finger protein 90 Zfp90
 1449127_at 0.88849 "selectin, platelet (p-selectin) ligand" Selpl
 1449128_at 1.00331 coiled-coil domain containing 43 Ccdc43
 1449129_a_at 1.06678 "Kv channel interacting protein 3, calsenilin" Kcnip3
 1449130_at 0.82716 CD1d1 antigen Cd1d1
 1449131_s_at 0.77252 CD1d1 antigen Cd1d1
 1449132_at 1.02072 "opsin 1 (cone pigments), short-wave-sensitive (color blindness, tritan)"
 Opnlsw
 1449133_at 1.14659 small proline-rich protein 1A Sprr1a
 1449134_s_at 1.06210 Spi-C transcription factor (Spi-1/PU.1 related) Spic
 1449135_at 0.92933 SRY-box containing gene 18 Sox18
 1449136_at 0.96697 eosinophil peroxidase Epx
 1449137_at 0.95874 pyruvate dehydrogenase E1 alpha 1 Pdha1
 1449138_at 1.06289 "splicing factor 3b, subunit 1" Sf3b1
 1449139_at 1.04610 RIKEN cDNA 2310008M10 gene 2310008M10Rik
 1449140_at 1.01318 NudC domain containing 2 Nudcd2
 1449141_at 0.98980 filamin binding LIM protein 1 Fblim1
 1449142_a_at 0.86778 "Yip1 domain family, member 5" Yipf5
 1449143_at 0.99119 receptor transporter protein 4 Rtp4
 1449144_at 0.95269 "guanine nucleotide binding protein, alpha 11" Gna11
 1449145_a_at 1.05321 "caveolin, caveolae protein 1" Cav1
 1449146_at 0.89769 Notch gene homolog 4 (Drosophila) Notch4
 1449147_at 0.94065 carbohydrate (keratan sulfate Gal-6) sulfotransferase 1 Chst1
 1449148_a_at 0.88141 putative homeodomain transcription factor 1 Phtf1
 1449149_at 0.94789 Tnf receptor-associated factor 3 Traf3
 1449150_at 0.94961 RIKEN cDNA A930040G15 gene A930040G15Rik
 1449151_at 1.02494 PCTAIRE-motif protein kinase 3 Pctk3
 1449152_at 1.08973 "cyclin-dependent kinase inhibitor 2B (p15, inhibits CDK4)" Cdkn2b

1449153_at 1.03522 matrix metalloproteinase 12 Mmp12
1449154_at 0.98881 "procollagen, type XI, alpha 1" Col11a1
1449155_at 1.39785 polymerase (RNA) III (DNA directed) polypeptide G Polr3g
1449156_at 1.10287 lymphocyte antigen 9 Ly9
1449157_at 0.87256 "nuclear receptor subfamily 2, group C, member 1" Nr2c1
1449158_at 0.92617 "potassium channel, subfamily K, member 2" Kcnk2
1449159_at 0.96405 "guanine nucleotide binding protein, beta 3" Gnb3
1449160_at 0.98887 natriuretic peptide receptor 1 Npr1
1449161_at 0.75542 endothelin 2 Edn2
1449162_at 1.04110 "processing of precursor 7, ribonuclease P family, (S. cerevisiae)" Pop7
1449163_at 1.11576 single immunoglobulin and toll-interleukin 1 receptor (TIR) domain Sigirr
1449164_at 0.95534 CD68 antigen Cd68
1449165_at 0.89779 calpain 5 Capn5
1449166_at 1.09818 S100 calcium binding protein A14 S100a14
1449167_at 1.05818 erythrocyte protein band 4.1-like 4a Epb4.114a
1449168_a_at 1.09796 A kinase (PRKA) anchor protein 2 Akap2
1449169_at 0.99319 hyaluronan synthase 2 Has2
1449170_at 0.95318 piwi-like homolog 2 (Drosophila) Piwil2
1449171_at 1.04624 Ttk protein kinase Ttk
1449172_a_at 0.94268 lin-7 homolog B (C. elegans) Lin7b
1449173_at 0.82300 "membrane protein, palmitoylated 2 (MAGUK p55 subfamily member 2)"
Mpp2
1449174_at 1.00861 ADP-ribosyltransferase 4 Art4
1449175_at 0.97729 G-protein coupled receptor 65 Gpr65
1449176_a_at 0.90128 deoxycytidine kinase Dck
1449177_at 0.92068 cyclin A1 Ccna1
1449178_at 0.95925 PDZ and LIM domain 3 Pdlim3
1449179_at 0.95949 phosducin Pdc
1449180_at 1.05131 potassium channel modulatory factor 1 Kcmf1
1449181_at 0.94799 ferrochelatase Fech
1449182_at 0.88483 resistin Retn
1449183_at 0.98845 catechol-O-methyltransferase Comt
1449184_at 1.15375 peptidoglycan recognition protein 1 Pglyrp1
1449185_at 1.07363 cone-rod homeobox containing gene Crx
1449186_at 0.92878 BCL2-associated athanogene 4 Bag4
1449187_at 0.96629 "platelet derived growth factor, alpha" Pdgfa
1449188_at 0.79298 midnolin Midn
1449189_at 0.92675 cDNA sequence BC003266 BC003266
1449190_a_at 1.06562 ectonucleoside triphosphate diphosphohydrolase 4 Entpd4
1449191_at 0.99892 WAP four-disulfide core domain 12 Wfdc12
1449192_at 0.98379 activating transcription factor 7 interacting protein Atf7ip
1449193_at 0.96038 CD5 antigen-like Cd5l
1449194_at 1.08072 mitochondrial ribosomal protein S25 Mrps25

1449195_s_at 1.81500 chemokine (C-X-C motif) ligand 16 Cxcl16
 1449196_a_at 0.99969 ribosomal protein S27a /// similar to ribosomal protein S27a /// similar to ubiquitin B precursor /// similar to ribosomal protein S27a /// similar to ribosomal protein S27a /// similar to ribosomal protein S27a Rps27a /// LOC619900 /// LOC625981 /// LOC639567 /// LOC667035 /// LOC677303
 1449197_at 0.97410 RIKEN cDNA 2310040A13 gene 2310040A13Rik
 1449198_a_at 0.85521 "ST3 beta-galactoside alpha-2,3-sialyltransferase 5" St3gal5
 1449199_at 1.60251 "mucin 1, transmembrane" Muc1
 1449200_at 1.09101 nucleoporin 155 Nup155
 1449201_at 1.13391 steroidogenic acute regulatory protein Star
 1449202_at 0.70415 "sema domain, immunoglobulin domain (Ig), transmembrane domain (TM) and short cytoplasmic domain, (semaphorin) 4G" Sema4g
 1449203_at 0.99129 "solute carrier organic anion transporter family, member 1a5" Slco1a5
 1449204_at 1.01059 gap junction membrane channel protein beta 5 Gjb5
 1449205_at 1.04298 ovo-like 2 (Drosophila) Ovol2
 1449206_at 1.11326 synaptophysin-like 2 Sypl2
 1449207_a_at 0.97618 kinesin family member 20A Kif20a
 1449208_at 0.82427 poly (A) polymerase beta (testis specific) Papolb
 1449209_a_at 1.21569 retinol dehydrogenase 11 Rdh11
 1449210_at 1.17068 insulin-like growth factor 2 mRNA binding protein 1 Igf2bp1
 1449211_at 1.03413 bisphosphate 3'-nucleotidase 1 Bpnt1
 1449212_at 1.02167 prolactin induced protein Pip
 1449213_at 1.02018 "L antigen family, member 3" Lage3
 1449214_a_at 1.12748 optic atrophy 1 homolog (human) Opa1
 1449215_at 0.89954 "solute carrier family 22 (organic cation transporter), member 21" Slc22a21
 1449216_at 0.95010 "integrin, alpha E, epithelial-associated" Itgae
 1449217_at 0.90983 caspase 8 associated protein 2 Casp8ap2
 1449218_at 1.04743 "cytochrome c oxidase, subunit VIIIb" Cox8b
 1449219_at 0.84380 fatty acid desaturase 3 Fads3
 1449220_at 0.97385 "GTPase, IMAP family member 3" Gimap3
 1449221_a_at 1.34171 ribosome binding protein 1 Rrbp1
 1449222_at 1.04061 Epstein-Barr virus induced gene 3 Ebi3
 1449223_at 1.02876 "DnaJ (Hsp40) homolog, subfamily B, member 8" Dnajb8
 1449224_at 1.07948 "transient receptor potential cation channel, subfamily M, member 5" Trpm5
 1449225_a_at 0.92943 mitogen-activated protein kinase 8 interacting protein 2 Mapk8ip2
 1449226_at 1.00288 hypermethylated in cancer 1 Hic1
 1449227_at 0.98211 cholesterol 25-hydroxylase Ch25h
 1449228_at 0.94211 SH3-domain GRB2-like 2 Sh3gl2
 1449229_a_at 0.93151 cyclin-dependent kinase-like 2 (CDC2-related kinase) Cdkl2
 1449230_at 0.88809 "iduronidase, alpha-L-" Idua
 1449231_at 0.95472 zinc finger protein 296 Zfp296
 1449232_at 1.09330 GATA binding protein 1 Gata1
 1449233_at 1.06361 "basic helix-loop-helix domain containing, class B, 8" Bhlhb8

1449234_at	1.03267 carbonic anhydrase 15	Car15	
1449235_at	1.00651 "Fas ligand (TNF superfamily, member 6)"		Fasl
1449236_at	1.02721 delta-like 3 (Drosophila)	Dll3	
1449237_at	0.95720 arachidonate lipoxygenase 3	Aloxe3	
1449238_at	1.12484 pregnancy specific glycoprotein 16	Psg16	
1449239_at	0.88938 RIKEN cDNA 1700045I19 gene	1700045I19Rik	
1449240_at	1.05702 G substrate	Gsbs	
1449241_at	1.00602 kelch-like 1 (Drosophila)	Klhl1	
1449242_s_at	1.03951 histidine-rich glycoprotein	Hrg	
1449243_a_at	1.08703 ribosomal protein S19	Rps19	
1449244_at	0.86353 cadherin 2	Cdh2	
1449245_at	1.01385 "glutamate receptor, ionotropic, NMDA2C (epsilon 3)"		Grin2c
1449246_at	1.00642 Rap2 interacting protein	Rap2ip	
1449247_at	0.91708 actin-related protein T2	Actrt2	
1449248_at	0.74489 chloride channel 2	Clcn2	
1449249_at	0.91745 protocadherin 7	Pcdh7	
1449250_at	1.17217 papillary renal cell carcinoma (translocation-associated)		Prcc
1449251_at	0.96288 Norrie disease homolog	Ndph	
1449252_at	0.92627 RIKEN cDNA 9030611O19 gene	9030611O19Rik	
1449253_at	1.00864 structural maintenance of chromosomes 1B	Smc1b	
1449254_at	0.93291 secreted phosphoprotein 1	Spp1	
1449255_a_at	1.03408 ribosomal protein L15	Rpl15	
1449256_a_at	1.00221 "RAB11a, member RAS oncogene family"	Rab11a	
1449257_at	0.93191 "DNA segment, Chr 11, Wayne State University 99, expressed"		D11Wsu99e
1449258_at	1.08123 "DNA segment, Chr 11, Wayne State University 99, expressed"		D11Wsu99e
1449259_at	0.78901 "RAB3D, member RAS oncogene family"	Rab3d	
1449260_at	0.97196 "RAB3D, member RAS oncogene family"	Rab3d	
1449261_at	0.99132 pre B-cell leukemia transcription factor 2	Pbx2	
1449262_s_at	0.86146 lin-7 homolog C (C. elegans)	Lin7c	
1449263_at	1.08196 ubiquitin-fold modifier 1	Ufm1	
1449264_at	0.82358 synaptotagmin XI	Syt11	
1449265_at	1.04754 caspase 1	Casp1	
1449266_at	1.00306 methyl CpG binding protein 2	Mecp2	
1449267_at	0.85462 RIKEN cDNA 3110023E09 gene	3110023E09Rik	
1449268_at	1.01168 glutamine fructose-6-phosphate transaminase 1	Gfpt1	
1449269_at	0.99206 coagulation factor V	F5	
1449270_at	1.07469 plexin domain containing 2	Plxdc2	
1449271_a_at	0.84618 heme binding protein 2	Hebp2	
1449272_at	1.12311 "immunoglobulin superfamily, member 4B"	Igsf4b	
1449273_at	0.87228 cytoplasmic FMR1 interacting protein 2	Cyfi2	
1449274_at	0.91038 RIKEN cDNA 2310038H17 gene	2310038H17Rik	
1449275_at	1.17651 RIKEN cDNA 2310038H17 gene	2310038H17Rik	
1449276_at	0.97171 RIKEN cDNA 1700029P11 gene	1700029P11Rik	

1449277_at 0.94081 chemokine (C-C motif) ligand 19 Ccl19
1449278_at 0.89886 eukaryotic translation initiation factor 2 alpha kinase 3 Eif2ak3
1449279_at 0.96765 glutathione peroxidase 2 Gpx2
1449280_at 0.90547 endothelial cell-specific molecule 1 /// similar to Endothelial cell-specific molecule 1 precursor (ESM-1 secretory protein) (ESM-1) Esm1 /// LOC632677
1449281_at 0.91199 neurturin Nrtn
1449282_at 0.90551 cysteinyl leukotriene receptor 1 Cysltr1
1449283_a_at 1.16238 mitogen-activated protein kinase 12 Mapk12
1449284_at 0.86644 RIKEN cDNA 1700008P20 gene 1700008P20Rik
1449285_at 0.81426 cystatin 9 Cst9
1449286_at 0.98995 netrin G1 Ntng1
1449287_at 0.94727 src-related kinase lacking C-terminal regulatory tyrosine and N-terminal myristylation sites Srms
1449288_at 1.03014 growth differentiation factor 3 Gdf3
1449289_a_at 1.09618 beta-2 microglobulin B2m
1449290_at 0.81246 dihydropyrimidinase-like 5 Dpysl5
1449291_a_at 0.99455 "discoidin, CUB and LCCL domain containing 1" Dcbl1
1449292_at 0.84764 RB1-inducible coiled-coil 1 Rb1cc1
1449293_a_at 1.09642 S-phase kinase-associated protein 2 (p45) Skp2
1449294_at 0.99580 mitochondrial ribosomal protein S15 Mrps15
1449295_at 1.04567 SAP30 binding protein Sap30bp
1449296_a_at 0.93434 cyclic nucleotide phosphodiesterase 1 Cnp1
1449297_at 0.99715 caspase 12 Casp12
1449298_a_at 0.97594 "phosphodiesterase 1A, calmodulin-dependent" Pde1a
1449299_at 0.85437 similar to Low-density lipoprotein receptor-related protein 5 precursor (LRP7) (Lr3) LOC670589
1449300_at 0.92372 CTTNBP2 N-terminal like Ctnnbp2nl
1449301_at 1.04849 "solute carrier family 7, (cationic amino acid transporter, y+ system) member 13" Slc7a13
1449302_at 1.04646 "ATP-binding cassette, sub-family A (ABC1), member 2" Abca2
1449303_at 0.85850 sestrin 3 Sesn3
1449304_at 0.85201 RIKEN cDNA 2310061J03 gene 2310061J03Rik
1449305_at 0.94570 coagulation factor X F10
1449306_at 0.98639 heat shock factor 2 Hsf2
1449307_at 1.15814 dysbindin (dystrobrevin binding protein 1) domain containing 1 Dbn1
1449308_at 1.05876 complement component 6 C6
1449309_at 0.94430 "cytochrome P450, family 8, subfamily b, polypeptide 1" Cyp8b1
1449310_at 0.97733 prostaglandin E receptor 2 (subtype EP2) Ptger2
1449311_at 0.96432 BTB and CNC homology 1 Bach1
1449312_at 0.90670 neuropeptide Y receptor Y5 Npy5r
1449313_at 1.02517 kallikrein 1-related peptidase b5 Klk1b5
1449314_at 1.14202 "zinc finger protein, multitype 2" Zfp2
1449315_at 0.91402 odd Oz/ten-m homolog 3 (Drosophila) Odz3

1449316_at 1.05577 "cytochrome P450, family 4, subfamily f, polypeptide 15" Cyp4f15
 1449317_at 0.92590 CASP8 and FADD-like apoptosis regulator Cflar
 1449318_at 1.27383 "tubulin, gamma 2" Tubg2
 1449319_at 1.05348 R-spondin homolog (Xenopus laevis) Rspo1
 1449320_at 1.01796 probasin Pbsn
 1449321_x_at 0.99735 "serine (or cysteine) peptidase inhibitor, clade A, member 1a /// serine (or cysteine) peptidase inhibitor, clade A, member 1b /// serine (or cysteine) peptidase inhibitor, clade A, member 1c /// serine (or cysteine) peptidase inhibitor, clade A, member 1d /// serine (or cysteine) peptidase inhibitor, clade A, member 1e /// similar to Alpha-1-antitrypsin 1-2 precursor (Serine protease inhibitor 1-2) (Alpha-1 protease inhibitor 2) (Alpha-1-antiproteinase) (AAT)" Serpina1a /// Serpina1b /// Serpina1c /// Serpina1d /// Serpina1e /// LOC675501
 1449322_at 0.97597 protein tyrosine phosphatase 4a1 /// similar to protein tyrosine phosphatase 4a1 /// protein tyrosine phosphatase 4a1-like /// similar to protein tyrosine phosphatase 4a1 /// protein tyrosine phosphatase 4a1 pseudogene /// protein tyrosine phosphatase 4a1-like /// similar to protein tyrosine phosphatase 4a1 /// similar to protein tyrosine phosphatase 4a1 /// similar to protein tyrosine phosphatase 4a1 Ptp4a1 /// LOC269859 /// LOC433406 /// LOC546482 /// LOC627166 /// LOC627200 /// LOC667723 /// LOC669215 /// LOC673579
 1449323_a_at 1.00566 ribosomal protein L3 /// similar to ribosomal protein L3 isoform b /// similar to 60S ribosomal protein L3 (J1 protein) /// similar to 60S ribosomal protein L3 (J1 protein) /// similar to ribosomal protein L3 isoform b Rpl3 /// LOC668219 /// LOC669672 /// LOC674810 /// LOC674874
 1449324_at 1.39869 ERO1-like (*S. cerevisiae*) Ero1l
 1449325_at 1.12407 fatty acid desaturase 2 Fads2
 1449326_x_at 0.97261 serum amyloid A 2 Saa2
 1449327_at 0.99718 RIKEN cDNA 1600015I10 gene 1600015I10Rik
 1449328_at 0.95214 lymphocyte antigen 75 Ly75
 1449329_at 0.93929 zinc finger protein 235 Zfp235
 1449330_at 0.92733 PDZ domain containing 3 Pdzd3
 1449331_a_at 0.81336 receptor-associated protein of the synapse Rapsn
 1449332_at 0.87508 four and a half LIM domains 5 Fhl5
 1449333_at 1.11593 "splicing factor 3a, subunit 1" Sf3a1
 1449334_at 0.70697 tissue inhibitor of metalloproteinase 3 Timp3
 1449335_at 0.73877 tissue inhibitor of metalloproteinase 3 Timp3
 1449336_a_at 1.08580 STE20-like kinase (yeast) Slk
 1449337_at 0.97698 "tryptophan 2,3-dioxygenase" Tdo2
 1449338_at 1.12487 "DNA segment, Chr 10, ERATO Doi 641, expressed" D10Ertd641e
 1449339_at 1.01086 "DNA segment, Chr 10, ERATO Doi 641, expressed" D10Ertd641e
 1449340_at 1.10298 sclerostin domain containing 1 Sostdc1
 1449341_a_at 0.93797 stomatin Stom
 1449342_at 1.02733 "protein tyrosine phosphatase-like (proline instead of catalytic arginine), member b" Ptplb
 1449343_s_at 1.04698 "transcriptional regulator, SIN3A (yeast)" Sin3a
 1449344_s_at 1.03113 RIKEN cDNA 2210409E12 gene 2210409E12Rik
 1449345_at 0.96262 RIKEN cDNA 2810027O19 gene 2810027O19Rik

1449346_s_at 0.90919 RIO kinase 1 (yeast) Riok1
 1449347_a_at 0.98374 X-linked lymphocyte-regulated 4B /// X-linked lymphocyte-regulated 4A /// X-linked lymphocyte-regulated 4E Xlr4b /// Xlr4a /// Xlr4e
 1449348_at 1.15350 "membrane protein, palmitoylated 6 (MAGUK p55 subfamily member 6)" Mpp6
 1449349_at 0.93051 nudix (nucleoside diphosphate linked moiety X)-type motif 1 Nudt1
 1449350_at 0.97134 odd-skipped related 1 (Drosophila) Osr1
 1449351_s_at 0.68483 "platelet-derived growth factor, C polypeptide" Pdgfc
 1449352_at 0.93199 "polymerase (DNA directed), lambda" Poll
 1449353_at 1.14633 zinc finger matrin type 3 Zmat3
 1449354_at 1.03326 "U2 small nuclear ribonucleoprotein auxiliary factor (U2AF) 1, related sequence 1" U2af1-rs1
 1449355_a_at 0.90672 epidermal growth factor receptor pathway substrate 15-like 1 Eps15l1
 1449356_at 1.00632 ankyrin repeat and SOCs box-containing protein 5 Asb5
 1449357_at 0.91331 RIKEN cDNA 2310030G06 gene 2310030G06Rik
 1449358_at 1.06738 "DNA segment, Chr 6, Miriam Meisler 5, expressed" D6Mm5e
 1449359_at 0.92527 paired box gene 1 Pax1
 1449360_at 0.84034 "colony stimulating factor 2 receptor, beta 2, low-affinity (granulocyte-macrophage)" Csf2rb2
 1449361_at 0.97701 T-box 21 Tbx21
 1449362_a_at 1.08771 misshapen-like kinase 1 (zebrafish) Mink1
 1449363_at 0.70112 activating transcription factor 3 Atf3
 1449364_at 0.96933 aurora kinase C Aurkc
 1449365_at 0.82927 "endothelial differentiation, sphingolipid G-protein-coupled receptor, 8" Edg8
 1449366_at 0.82975 matrix metalloproteinase 8 Mmp8
 1449367_at 1.01629 three prime repair exonuclease 2 Trex2
 1449368_at 0.61091 decorin Dcn
 1449369_at 0.90853 "transmembrane protease, serine 2" Tmprss2
 1449370_at 0.70917 SRY-box containing gene 4 /// similar to Transcription factor SOX-4 Sox4 /// LOC672274
 1449371_at 1.01368 histidyl-tRNA synthetase-like Harsl
 1449372_at 0.91090 "DnaJ (Hsp40) homolog, subfamily C, member 3" Dnajc3
 1449373_at 0.96592 "DnaJ (Hsp40) homolog, subfamily C, member 3" Dnajc3
 1449374_at 0.98067 pipercolic acid oxidase Pipox
 1449375_at 0.95873 carboxylesterase 6 Ces6
 1449376_at 0.94493 nicolin 1 Nicn1
 1449377_at 0.84687 nicolin 1 Nicn1
 1449378_at 1.09034 keratin 27 Krt27
 1449379_at 0.88050 kinase insert domain protein receptor Kdr
 1449380_at 0.97900 protein kinase C and casein kinase substrate in neurons 1 Pacsin1
 1449381_a_at 0.92566 protein kinase C and casein kinase substrate in neurons 1 Pacsin1
 1449382_at 1.05703 "solute carrier family 6 (neurotransmitter transporter, betaine/GABA), member

12" Slc6a12

1449383_at	0.87591	adenylosuccinate synthetase like 1	Adssl1
1449384_at	0.96773	RIKEN cDNA 4931417E11 gene	4931417E11Rik
1449385_at	1.09179	hydroxysteroid (17-beta) dehydrogenase 9	Hsd17b9
1449386_at	0.88499	hydroxysteroid (17-beta) dehydrogenase 9	Hsd17b9
1449387_at	0.93055	keratin 33A	Krt33a
1449388_at	1.05081	thrombospondin 4	Thbs4
1449389_at	1.06358	T-cell acute lymphocytic leukemia 1	Tal1
1449390_at	1.29292	G patch domain containing 4	Gpatc4
1449391_at	1.00370	zinc finger protein 37	Zfp37
1449392_at	0.78969	hydroxysteroid (17-beta) dehydrogenase 1	Hsd17b1
1449393_at	1.09357	SH2 domain protein 1A	Sh2d1a
1449394_at	1.16746	"solute carrier organic anion transporter family, member 1b2"	Slco1b2
1449395_at	1.05740	RIKEN cDNA 4921520G13 gene	4921520G13Rik
1449396_at	0.92715	"amine oxidase, copper containing 3"	Aoc3
1449397_at	0.92919	homeo box B2	Hoxb2
1449398_at	1.05903	ribosomal protein L3-like	Rpl3l
1449399_a_at	0.89544	interleukin 1 beta	Il1b
1449400_at	1.00877	citrate synthase like	Csl
1449401_at	1.00251	"complement component 1, q subcomponent, C chain"	C1qc
1449402_at	0.97901	carbohydrate (N-acetylglucosamino) sulfotransferase 7	Chst7
1449403_at	0.88536	phosphodiesterase 9A	Pde9a
1449404_at	0.70750	"phosphatidylinositol-4-phosphate 5-kinase, type II, alpha"	Pip5k2a
1449405_at	0.84027	tensin 1	Tns1
1449406_at	0.90468	cysteine and histidine rich 1	Cyhr1
1449407_at	0.82958	intraflagellar transport 81 homolog (Chlamydomonas)	Ift81
1449408_at	0.94052	junction adhesion molecule 2	Jam2
1449409_at	1.24282	"sulfotransferase family, cytosolic, 1C, member 2"	Sult1c2
1449410_a_at	1.19987	growth arrest specific 5	Gas5
1449411_at	0.98302	Down syndrome cell adhesion molecule	Dscam
1449412_at	0.91569	RIKEN cDNA 1810046J19 gene	1810046J19Rik
1449413_at	1.03645	"Mpv17 transgene, kidney disease mutant-like"	Mpv17l
1449414_at	1.27012	zinc finger protein 53	Zfp53
1449415_at	1.13654	chromodomain helicase DNA binding protein 1-like	Chd1l
1449416_at	0.85131	frizzled homolog 4 (Drosophila)	Fzd4
1449417_at	0.94163	ameloblastin	Ambn
1449418_s_at	1.05694	F-box only protein 36	Fbxo36
1449419_at	0.80184	dedicator of cytokinesis 8	Dock8
1449420_at	0.92155	"phosphodiesterase 1B, Ca ²⁺ -calmodulin dependent"	Pde1b
1449421_a_at	0.93945	"potassium voltage-gated channel, Isk-related subfamily, gene 2"	Kcne2
1449422_at	0.91474	cadherin 4	Cdh4
1449423_at	0.98751	microtubule associated serine/threonine kinase 1	Mast1
1449424_at	0.93714	pleckstrin 2	Plek2

1449425_at 1.02595 wingless-related MMTV integration site 2 Wnt2
 1449426_a_at 1.09170 annexin A10 Anxa10
 1449427_at 1.12790 RIKEN cDNA 1700016G05 gene 1700016G05Rik
 1449428_at 0.91684 claudin 18 Cldn18
 1449429_at 0.82561 FK506 binding protein 1b Fkbp1b
 1449430_a_at 1.05032 trehalase (brush-border membrane glycoprotein) Treh
 1449431_at 1.01700 "transient receptor potential cation channel, subfamily C, member 6" Trpc6
 1449432_a_at 0.89403 membrane metallo-endopeptidase-like 1 Mmell1
 1449433_at 0.98404 "purinergic receptor P2X, ligand-gated ion channel, 5" P2rx5
 1449434_at 1.06536 carbonic anhydrase 3 Car3
 1449435_at 0.88742 "UDP-Gal:betaGlcNAc beta 1,4-galactosyltransferase, polypeptide 3"
 B4galt3
 1449436_s_at 0.94326 "ubiquitin B /// ubiquitin pseudogene /// gene model 1821, (NCBI) /// similar to
 polyubiquitin" Ubb /// LOC218963 /// Gm1821 /// LOC667759
 1449437_at 0.96341 "DNA segment, Chr 6, Wayne State University 163, expressed" D6Wsu163e
 1449438_at 0.98600 dolichol-phosphate (beta-D) mannosyltransferase 1 Dpm1
 1449439_at 0.88368 Kruppel-like factor 7 (ubiquitous) Klf7
 1449440_at 0.92253 lipin 3 Lpin3
 1449441_a_at 0.89282 WW domain binding protein 1 Wbp1
 1449442_at 0.96839 peroxisomal biogenesis factor 11a Pex11a
 1449443_at 0.80159 "2,4-dienoyl CoA reductase 1, mitochondrial" Decr1
 1449444_a_at 0.94297 microfibrillar-associated protein 1 Mfap1
 1449445_x_at 1.02849 microfibrillar-associated protein 1 Mfap1
 1449446_at 0.91473 coiled-coil domain containing 59 Ccdc59
 1449447_at 0.91689 cystatin 10 (chondrocytes) Cst10
 1449448_at 0.98555 "mediator of RNA polymerase II transcription, subunit 9 homolog (yeast)"
 Med9
 1449449_at 0.95203 prostaglandin E synthase Ptges
 1449450_at 1.08148 prostaglandin E synthase Ptges
 1449451_at 0.91782 "serine (or cysteine) peptidase inhibitor, clade B (ovalbumin), member 11"
 Serpinb11
 1449452_a_at 0.97880 glycoprotein 2 (zymogen granule membrane) Gp2
 1449453_at 1.12728 bone marrow stromal cell antigen 1 Bst1
 1449454_at 1.28603 bone marrow stromal cell antigen 1 Bst1
 1449455_at 1.21610 hemopoietic cell kinase Hck
 1449456_a_at 0.96742 "chymase 1, mast cell" Cma1
 1449457_at 0.86868 acyl-CoA thioesterase 12 Acot12
 1449458_at 1.12280 forkhead box I1 Foxo1
 1449459_s_at 1.01565 ankyrin repeat and SOCS box-containing protein 13 Asb13
 1449460_at 1.06837 ankyrin repeat and SOCS box-containing protein 13 Asb13
 1449461_at 0.93541 "retinol binding protein 7, cellular" Rbp7
 1449462_at 1.06267 RIKEN cDNA 3110049J23 gene 3110049J23Rik
 1449463_at 0.92206 kallikrein 1-related peptidase b8 Klk1b8

1449464_at 0.87211 "potassium voltage-gated channel, subfamily Q, member 1" Kcnq1
1449465_at 0.99618 reelin Reln
1449466_at 0.91476 "C-type lectin domain family 3, member b" Clec3b
1449467_at 1.01635 spermiogenesis specific transcript on the Y 1 /// similar to Y-linked testis-specific protein /// similar to Y-linked testis-specific protein /// similar to Y-linked testis-specific protein Ssty1 /// LOC665276 /// LOC665301 /// LOC665687
1449468_at 0.91397 "ST6 (alpha-N-acetyl-neuraminy-2,3-beta-galactosyl-1,3)-N-acetylgalactosaminide alpha-2,6-sialyltransferase 5" St6galnac5
1449469_at 1.09844 polycystic kidney disease 2-like 2 Pkd2l2
1449470_at 1.02076 distal-less homeobox 1 Dlx1
1449471_at 0.80208 "potassium large conductance calcium-activated channel, subfamily M, beta member 4" Kcnmb4
1449472_at 0.97734 G-protein coupled receptor 12 Gpr12
1449473_s_at 1.05350 CD40 antigen Cd40
1449474_a_at 0.93399 nasal embryonic LHRH factor Nelf
1449475_at 0.96141 "ATPase, H+/K+ transporting, nongastric, alpha polypeptide" Atp12a
1449476_at 1.02454 renal tumor antigen Rage
1449477_s_at 0.91372 "solute carrier family 2 (facilitated glucose transporter), member 10" Slc2a10
1449478_at 0.96702 matrix metalloproteinase 7 Mmp7
1449479_at 0.99835 "cytochrome P450, family 2, subfamily b, polypeptide 13" Cyp2b13
1449480_at 1.12377 Sin3-associated polypeptide 18 Sap18
1449481_at 1.09805 "solute carrier family 25 (mitochondrial carrier, adenine nucleotide translocator), member 13" Slc25a13
1449482_at 0.96837 "histone 3, H2ba" Hist3h2ba
1449483_at 1.09656 "polymerase (DNA directed), kappa" Polk
1449484_at 1.03134 stanniocalcin 2 Stc2
1449485_at 0.98466 receptor (TNFRSF)-interacting serine-threonine kinase 1 Ripk1
1449486_at 1.08982 carboxylesterase 1 Ces1
1449487_at 0.95778 coiled-coil domain containing 70 Ccdc70
1449488_at 1.03053 paired-like homeodomain transcription factor 1 Pitx1
1449489_at 0.95212 testis-specific serine kinase 2 Tssk2
1449490_at 0.96499 methyl-CpG binding domain protein 4 Mbd4
1449491_at 0.84962 "caspase recruitment domain family, member 10" Card10
1449492_a_at 1.10408 leukocyte cell-derived chemotaxin 2 Lect2
1449493_at 0.90205 insulin-like 5 Ins5
1449494_at 0.98819 "RAB3C, member RAS oncogene family" Rab3c
1449495_at 0.96209 regenerating islet-derived 3 alpha Reg3a
1449496_at 1.02561 RIKEN cDNA 2010109I03 gene 2010109I03Rik
1449497_at 0.96552 interleukin 12b Il12b
1449498_at 1.05509 macrophage receptor with collagenous structure Marco
1449499_at 0.88596 homeo box A7 Hoxa7
1449500_at 0.98374 "serine (or cysteine) peptidase inhibitor, clade B, member 7" Serpinb7
1449501_a_at 1.04240 granzyme M (lymphocyte met-ase 1) Gzmm

1449502_at 0.87184 deleted in azoospermia-like Dazl
1449503_at 0.97111 karyopherin (importin) alpha 1 Kpna1
1449504_at 0.93791 karyopherin (importin) alpha 1 Kpna1
1449505_at 1.08782 karyopherin (importin) alpha 1 Kpna1
1449506_a_at 1.00012 eukaryotic translation elongation factor 1 delta (guanine nucleotide exchange protein) Eef1d
1449507_a_at 1.20581 "CD47 antigen (Rh-related antigen, integrin-associated signal transducer)" Cd47
1449508_at 1.05916 "interleukin 27 receptor, alpha" Il27ra
1449509_at 0.76451 small EDRK-rich factor 1 Serf1
1449510_at 0.89064 zinc finger protein 467 Zfp467
1449511_a_at 0.89454 single stranded DNA binding protein 4 Ssbp4
1449512_a_at 0.93815 zinc finger protein X-linked Zfx
1449513_at 1.03903 a disintegrin and metallopeptidase domain 24 (testase 1) Adam24
1449514_at 0.87105 G protein-coupled receptor kinase 5 Gprk5
1449515_at 1.12493 zinc finger protein 292 Zfp292
1449516_a_at 0.84692 regulator of G-protein signaling 3 Rgs3
1449517_at 0.92539 glutaminyl-peptide cyclotransferase-like Qpctl
1449518_at 1.00565 glutaminyl-peptide cyclotransferase-like Qpctl
1449519_at 1.10754 growth arrest and DNA-damage-inducible 45 alpha Gadd45a
1449520_at 0.78748 tetratricopeptide repeat domain 28 Ttc28
1449521_at 0.99986 CD93 antigen Cd93
1449522_at 1.05761 unc-5 homolog C (C. elegans) Unc5c
1449523_at 0.96848 B-cell CLL/lymphoma 7C Bcl7c
1449524_at 0.91695 ectodysplasin-A Eda
1449525_at 1.05121 flavin containing monooxygenase 3 Fmo3
1449526_a_at 0.95198 glycerophosphodiester phosphodiesterase domain containing 3 Gdpd3
1449527_at 0.93913 protocadherin beta 7 Pcdhb7
1449528_at 0.91832 c-fos induced growth factor Figf
1449529_s_at 1.00755 prolactin-like protein E Prlpe
1449530_at 0.76100 trichorhinophalangeal syndrome I (human) Trps1
1449531_at 1.02714 leprecan-like 2 Leprel2
1449532_at 0.96619 "cholinergic receptor, nicotinic, gamma polypeptide" Chrng
1449533_at 1.02547 transmembrane protein 100 Tmem100
1449534_at 0.82224 synaptonemal complex protein 3 Sycp3
1449535_at 0.95864 zinc and ring finger 4 Znrf4
1449536_at 0.68862 "potassium intermediate/small conductance calcium-activated channel, subfamily N, member 1" Kcnn1
1449537_at 0.87593 mutS homolog 5 (E. coli) Msh5
1449538_a_at 1.14782 "glucosaminyl (N-acetyl) transferase 1, core 2 /// similar to glucosaminyl (N-acetyl) transferase 1, core 2" Gcnt1 /// LOC635918
1449539_at 0.97921 "membrane-spanning 4-domains, subfamily A, member 11" Ms4a11
1449540_at 0.91720 reproductive homeobox 9 Rhox9

1449541_x_at 1.03784 RIKEN cDNA 1700010I14 gene 1700010I14Rik
1449542_at 0.94385 pre B-cell leukemia transcription factor 1 Pbx1
1449543_at 1.20893 zinc finger protein 371 Zfp371
1449544_a_at 0.52810 "potassium voltage-gated channel, subfamily H (eag-related), member 2"
Kcnh2
1449545_at 0.98293 fibroblast growth factor 18 Fgf18
1449546_a_at 0.98444 zinc finger protein 617 Zfp617
1449547_at 1.05115 ankyrin repeat and SOCS box-containing protein 14 Asb14
1449548_at 0.90230 ephrin B2 Efnb2
1449549_at 0.80802 ephrin B2 Efnb2
1449550_at 1.02204 myosin IC Myo1c
1449551_at 1.11075 myosin IC Myo1c
1449552_at 1.10765 zinc finger RNA binding protein Zfr
1449553_at 1.00342 RIKEN cDNA 2610200G18 gene 2610200G18Rik
1449554_at 0.92971 "transducin-like enhancer of split 3, homolog of Drosophila E(spl)" Tle3
1449555_a_at 1.05947 fetuin beta Fetub
1449556_at 1.74861 "histocompatibility 2, T region locus 23 /// RIKEN cDNA C920025E04 gene ///
similar to H-2 class I histocompatibility antigen, D-37 alpha chain precursor /// similar to H-2 class I
histocompatibility antigen, D-37 alpha chain precursor /// similar to H-2 class I histocompatibility
antigen, D-37 alpha chain precursor" H2-T23 /// C920025E04Rik /// LOC667803 /// LOC672711 ///
LOC674134
1449557_at 1.02226 RIKEN cDNA 1600012F09 gene /// hypothetical LOC624446
1600012F09Rik /// LOC624446
1449558_at 0.88479 coagulation factor VIII F8
1449559_at 1.04665 "homeo box, msh-like 2" Msx2
1449560_at 0.97845 RIKEN cDNA 2310043L02 gene 2310043L02Rik
1449561_at 1.11173 RIKEN cDNA 4921504I05 gene 4921504I05Rik
1449562_at 0.96985 RIKEN cDNA 1700001G17 gene 1700001G17Rik
1449563_at 0.96372 contactin 1 Cntn1
1449564_at 1.07254 tryptase gamma 1 Tpsg1
1449565_at 1.03024 "cytochrome P450, family 2, subfamily g, polypeptide 1" Cyp2g1
1449566_at 1.07342 "NK2 transcription factor related, locus 5 (Drosophila)" Nkx2-5
1449567_at 1.03864 transketolase-like 1 Tk1l
1449568_at 0.89069 klotho beta Klb
1449569_at 1.08073 thrombopoietin Thpo
1449570_at 0.96217 killer cell lectin-like receptor subfamily B member 1C Klr1c
1449571_at 0.96422 thyrotropin releasing hormone receptor Trhr
1449572_at 0.91200 thyrotropin releasing hormone receptor Trhr
1449573_at 1.04641 alpha-kinase 3 Alpk3
1449574_a_at 1.00079 cell division cycle 42 homolog (S. cerevisiae) Cdc42
1449575_a_at 0.94803 "glutathione S-transferase, pi 1" Gstp1
1449576_at 0.98226 "eukaryotic translation initiation factor 1A, Y-linked" Eif1ay
1449577_x_at 0.90864 "tropomyosin 2, beta" Tpm2

1449578_at 1.11608 suppressor of Ty 16 homolog (*S. cerevisiae*) Supt16h
1449579_at 0.72819 Sh3 domain YSC-like 1 Sh3yl1
1449580_s_at 0.91709 "histocompatibility 2, class II, locus Mb1 /// histocompatibility 2, class II, locus Mb2" H2-DMb1 /// H2-DMb2
1449581_at 0.89610 EMI domain containing 1 Emid1
1449582_at 0.90812 caudal type homeo box 1 Cdx1
1449583_at 0.68497 protocadherin beta 20 Pcdhb20
1449584_at 1.00585 "diacylglycerol kinase, gamma" Dgkg
1449585_at 0.96806 interleukin 1 receptor accessory protein Il1rap
1449586_at 0.94453 plakophilin 1 Pkp1
1449587_a_at 1.10653 "mucin 10, submandibular gland salivary mucin" Muc10
1449588_at 1.07005 "ATP-binding cassette, sub-family A (ABC1), member 4" Abca4
1449589_x_at 0.95487 RIKEN cDNA 2610020O08 gene 2610020O08Rik
1449590_a_at 1.03832 muscle and microspikes RAS Mras
1449591_at 1.79757 "caspase 4, apoptosis-related cysteine peptidase" Casp4
1449592_at 1.01036 transcription factor 15 Tcf15
1449593_at 1.05671 --- ---
1449594_at 1.09832 "ATP-binding cassette, sub-family B (MDR/TAP), member 1B" Abcb1b
1449595_at 1.02458 expressed sequence C79490 C79490
1449596_at 0.89740 expressed sequence C78142 C78142
1449597_at 1.06082 similar to odorant binding protein Ia LOC236874
1449598_at 0.99063 Density-regulated protein Denr
1449599_at 0.93344 --- ---
1449600_at 0.98820 "gene model 1683, (NCBI)" Gm1683
1449601_x_at 0.94549 Transcribed locus ---
1449602_at 0.99691 Kelch-like 5 (*Drosophila*) Khlh5
1449603_at 1.11351 expressed sequence AI594671 AI594671
1449604_at 0.93218 "acid phosphatase 1, soluble /// similar to acid phosphatase 1, soluble /// similar to acid phosphatase 1, soluble /// similar to acid phosphatase 1 isoform b /// similar to acid phosphatase 1, soluble /// similar to acid phosphatase 1, soluble /// similar to acid phosphatase 1 isoform b /// similar to acid phosphatase 1 isoform b /// similar to acid phosphatase 1 isoform b" Acp1 /// LOC619997 /// LOC631286 /// LOC635312 /// LOC665717 /// LOC666063 /// LOC668486 /// LOC670469 /// LOC676625
1449605_at 0.94858 Transcribed locus ---
1449606_at 0.80884 --- ---
1449607_at 0.98588 RIKEN cDNA A230070E04 gene A230070E04Rik
1449608_a_at 0.90870 --- ---
1449609_at 0.97650 Transcribed locus ---
1449610_at 0.90515 E1A binding protein p400 Ep400
1449611_at 0.86676 CD82 antigen Cd82
1449612_x_at 0.93964 M-phase phosphoprotein 1 Mphosph1
1449613_at 0.98616 --- ---
1449614_s_at 1.02024 expressed sequence AI314976 /// similar to Protein C6orf130 homolog

AI314976 /// LOC672876

1449615_s_at 0.98745 high density lipoprotein (HDL) binding protein Hdlbp
 1449616_s_at 0.94371 "golgi autoantigen, golgin subfamily a, 3" Golga3
 1449617_at 0.93262 RIKEN cDNA 2900092E17 gene 2900092E17Rik
 1449618_s_at 0.96727 RIKEN cDNA 2900092E17 gene 2900092E17Rik
 1449619_s_at 0.89036 Rho GTPase activating protein 9 Arhgap9
 1449620_s_at 1.06839 "DNA segment, Chr 16, Wayne State University 65, expressed" D16Wsu65e
 1449621_s_at 0.89168 "thrombospondin, type I, domain 1" Thsd1
 1449622_s_at 0.96092 "ATPase, H+ transporting, lysosomal accessory protein 1" Atp6ap1
 1449623_at 1.02178 thioredoxin reductase 3 Txnrd3
 1449624_at 1.00053 "gene model 118, (NCBI)" Gm118
 1449625_at 1.06224 "Transcribed locus, weakly similar to NP_570953.1 CAMP [Mus musculus]" ---
 1449626_s_at 0.93615 acyl-Coenzyme A binding domain containing 4 Acbd4
 1449627_at 0.98016 Oxysterol binding protein-like 6 Osbpl6
 1449628_s_at 0.89854 START domain containing 7 Stard7
 1449629_s_at 1.00489 Small nuclear ribonucleoprotein D3 Snrpd3
 1449630_s_at 1.13001 MAP/microtubule affinity-regulating kinase 1 Mark1
 1449631_at 1.03532 "Enolase 3, beta muscle" Eno3
 1449632_s_at 0.80389 FK506 binding protein 10 Fkbp10
 1449633_s_at 0.95909 "5'-nucleotidase, cytosolic III-like" Nt5c3l
 1449634_a_at 0.91280 ankyrin repeat and sterile alpha motif domain containing 1B /// similar to
 AIDA-1b Anks1b /// LOC544718
 1449635_at 1.00606 PRP19/PSO4 pre-mRNA processing factor 19 homolog (S. cerevisiae)
 Prpf19
 1449637_at 1.01342 cadherin 4 Cdh4
 1449638_at 1.00690 --- ---
 1449639_at 1.00040 RIKEN cDNA 0610040J01 gene 0610040J01Rik
 1449640_at 0.97612 toll-like receptor 7 Tlr7
 1449641_at 1.08086 --- ---
 1449642_at 0.84229 Phosphoglycerate dehydrogenase like 1 Phgdh1
 1449643_s_at 1.04674 basic transcription factor 3 Btf3
 1449644_at 1.05107 similar to T-cell receptor alpha chain V region CTL-F3 precursor
 LOC236118
 1449645_s_at 1.04414 chaperonin subunit 3 (gamma) Cct3
 1449646_s_at 0.86079 tigger transposable element derived 5 Tigd5
 1449647_at 1.04988 Transcribed locus ---
 1449648_s_at 1.09682 RNA polymerase 1-1 Rpo1-1
 1449649_at 0.97498 "ATPase, H+ transporting, lysosomal V1 subunit B2" Atp6v1b2
 1449650_at 1.01344 Ethanolamine kinase 1 Etnk1
 1449651_x_at 1.02278 --- ---
 1449652_at 0.96432 NudC domain containing 1 Nudcd1
 1449653_at 1.67247 Transcribed locus ---
 1449654_s_at 0.95168 expressed sequence C77545 C77545

1449655_a_at 0.96704 CDNA clone IMAGE:5690974 ---
 1449656_at 1.05733 RIKEN cDNA D930048N14 gene D930048N14Rik
 1449657_at 0.96038 "similar to proteasome (prosome, macropain) subunit, beta type 7"
 MGC107702
 1449658_at 0.90844 expressed sequence AA536748 AA536748
 1449659_s_at 1.08724 M-phase phosphoprotein 9 Mphosph9
 1449660_s_at 1.05754 "coronin, actin binding protein 1C" Coro1c
 1449661_at 0.78859 Suppressor of zeste 12 homolog (Drosophila) Suz12
 1449662_at 1.03711 Receptor accessory protein 1 Reep1
 1449663_at 0.98746 --- ---
 1449664_s_at 0.96078 ring finger protein 20 Rnf20
 1449665_at 0.94391 RIKEN cDNA 4833422M21 gene 4833422M21Rik
 1449666_at 0.97287 Attractin like 1 Atrnl1
 1449667_at 0.95128 Similar to Zinc finger protein 124 (HZF-16) LOC626391
 1449668_s_at 1.06539 RIKEN cDNA A730024A03 gene A730024A03Rik
 1449669_at 0.98375 RAB interacting factor Rabif
 1449670_x_at 1.03132 G protein-coupled receptor 137B Gpr137b
 1449671_at 1.06021 --- ---
 1449672_s_at 0.90618 "Tu translation elongation factor, mitochondrial /// similar to Tu translation
 elongation factor, mitochondrial" Tufm /// LOC669832
 1449673_s_at 0.95953 --- ---
 1449674_s_at 1.07541 programmed cell death 6 interacting protein Pdcd6ip
 1449675_at 0.96834 "cyclin B1, related sequence 1 /// similar to G2/mitotic-specific cyclin B1 ///
 cyclin B1 /// similar to G2/mitotic-specific cyclin-B1 /// similar to G2/mitotic-specific cyclin-B1 ///
 similar to G2/mitotic-specific cyclin-B1" Ccnb1-rs1 /// LOC231869 /// Ccnb1 /// LOC434175 ///
 LOC545676 /// LOC672788
 1449676_at 0.76280 "RAB2, member RAS oncogene family" Rab2
 1449677_s_at 0.92332 transmembrane protein 38B Tmem38b
 1449678_at 0.80983 --- ---
 1449679_s_at 0.86427 syntaxin 5A Stx5a
 1449680_at 1.00068 expressed sequence C80678 C80678
 1449681_at 0.85540 Hepatoma-derived growth factor Hdgf
 1449682_s_at 1.08494 "tubulin, beta 2b" Tubb2b
 1449683_x_at 0.92822 "tubulin, beta 2b" Tubb2b
 1449684_at 0.92289 "3-oxoacyl-ACP synthase, mitochondrial" Oxsm
 1449685_s_at 1.02343 "3-oxoacyl-ACP synthase, mitochondrial" Oxsm
 1449686_s_at 0.92912 "sterol carrier protein 2, liver" Scp2
 1449687_at 0.95731 "DNA segment, Chr 10, ERATO Doi 610, expressed" D10Ertd610e
 1449688_at 0.86053 "DNA segment, Chr 10, ERATO Doi 610, expressed" D10Ertd610e
 1449689_at 0.98000 expressed sequence C81600 C81600
 1449690_x_at 0.91166 --- ---
 1449691_at 0.81206 Zinc finger protein 644 Zfp644
 1449692_at 0.88144 "protein kinase C, zeta" Prkcz

1449693_at 1.09393 Mitogen activated protein kinase kinase kinase 7 Map3k7
1449694_s_at 0.96946 COMM domain containing 5 Commd5
1449695_at 0.96540 RIKEN cDNA A830082K12 gene A830082K12Rik
1449696_at 1.02460 Aryl hydrocarbon receptor nuclear translocator Arnt
1449697_s_at 0.93677 mitofusin 1 Mfn1
1449698_at 0.85701 RIKEN cDNA C330027C09 gene C330027C09Rik
1449699_s_at 0.95734 RIKEN cDNA C330027C09 gene C330027C09Rik
1449700_at 0.91813 Immunoglobulin (CD79A) binding protein 1 Igbp1
1449701_at 1.13422 "DNA segment, Chr 6, ERATO Doi 109, expressed" D6Ertd109e
1449702_at 1.05883 "zinc finger, AN1-type domain 2A" Zfand2a
1449703_at 0.83195 "Zinc finger, AN1-type domain 2A" Zfand2a
1449704_at 1.08478 expressed sequence C80171 C80171
1449705_x_at 1.02025 minichromosome maintenance deficient 3 (*S. cerevisiae*) /// similar to DNA replication licensing factor MCM3 (DNA polymerase alpha holoenzyme-associated protein P1) (P1-MCM3) Mcm3 /// LOC671904
1449706_s_at 0.96829 "nuclear receptor subfamily 5, group A, member 2" Nr5a2
1449707_at 0.92091 "nuclear receptor subfamily 5, group A, member 2" Nr5a2
1449708_s_at 0.94476 checkpoint kinase 1 homolog (*S. pombe*) Chek1
1449709_s_at 1.02984 arginine-tRNA-protein transferase 1 Ate1
1449710_s_at 0.92239 "ATP synthase, H⁺ transporting, mitochondrial F1 complex, alpha subunit, isoform 1" Atp5a1
1449711_at 0.92758 "VATPase, H⁺ transporting, lysosomal V1 subunit E1" Atp6v1e1
1449712_s_at 0.96511 "VATPase, H⁺ transporting, lysosomal V1 subunit E1" Atp6v1e1
1449713_at 1.00546 Chromobox homolog 7 Cbx7
1449714_at 0.88789 RIKEN cDNA 5730472N09 gene 5730472N09Rik
1449715_at 0.98368 --- ---
1449716_s_at 1.08744 "nardilysin, N-arginine dibasic convertase, NRD convertase 1" Nrd1
1449717_at 0.87801 --- ---
1449718_s_at 0.88725 RIKEN cDNA 4930453N24 gene 4930453N24Rik
1449719_at 0.91057 RIKEN cDNA 4930453N24 gene 4930453N24Rik
1449720_at 1.02202 --- ---
1449721_at 1.05650 "PREDICTED: Mus musculus hypothetical protein LOC628408 (LOC628408), mRNA" ---
1449722_at 1.01235 Transcribed locus ---
1449723_at 1.04363 "serine (or cysteine) peptidase inhibitor, clade B, member 6e" Serpinb6e
1449724_s_at 1.07385 "DNA segment, Chr 8, ERATO Doi 738, expressed" D8Ertd738e
1449725_at 1.02959 Glutamine fructose-6-phosphate transaminase 1 Gfpt1
1449726_at 0.93553 thioredoxin-like 2 /// similar to thioredoxin-like 2 Txnl2 /// LOC620016
1449727_x_at 0.97463 --- ---
1449728_at 1.00381 Transcribed locus ---
1449729_at 1.09928 fibroblast growth factor 4 Fgf4
1449730_s_at 1.00386 frizzled homolog 3 (*Drosophila*) Fzd3
1449731_s_at 1.48068 "nuclear factor of kappa light chain gene enhancer in B-cells inhibitor, alpha"

Nfkbia

1449732_at 0.88121 zinc finger proliferation 1 Zipr1
1449733_s_at 0.91925 seven in absentia 1A Siah1a
1449734_s_at 0.82311 Bardet-Biedl syndrome 4 homolog (human) Bbs4
1449735_at 0.78343 --- ---
1449736_at 0.96484 "Phosphatidylinositol-4-phosphate 5-kinase, type II, alpha" Pip5k2a
1449737_at 1.03266 Phosphofurin acidic cluster sorting protein 1 Pacs1
1449738_s_at 0.99598 Exosome component 8 Exosc8
1449739_at 0.89015 Phosphatidylserine synthase 1 Ptdss1
1449740_s_at 0.88064 desmoglein 2 Dsg2
1449741_at 0.90078 Desmoglein 2 Dsg2
1449742_at 0.96846 expressed sequence AA522020 AA522020
1449743_a_at 1.01382 CDNA sequence BC021438 BC021438
1449744_at 1.05344 "Solute carrier family 19 (sodium/hydrogen exchanger), member 1" Slc19a1
1449745_at 1.04685 Transcribed locus ---
1449746_s_at 1.15442 GLI pathogenesis-related 1 (glioma) Glipr1
1449747_at 1.03822 "Bin3, bicoid-interacting 3, homolog (Drosophila)" Bcdin3
1449748_at 0.97106 expressed sequence AA517023 AA517023
1449749_s_at 1.06343 "transcription factor B1, mitochondrial" Tfb1m
1449750_at 0.98909 expressed sequence AA407331 AA407331
1449751_at 0.92563 --- ---
1449752_at 1.01160 SplA/ryanodine receptor domain and SOCS box containing 1 Spsb1
1449753_at 0.89747 myosin IE Myo1e
1449754_at 0.99779 expressed sequence AA511261 AA511261
1449755_at 1.43454 "DNA segment, Chr 7, ERATO Doi 183, expressed" D7Ertd183e
1449756_at 1.07009 "DnaJ (Hsp40) homolog, subfamily C, member 17" Dnajc17
1449757_x_at 0.91921 "deoxynucleotidyltransferase, terminal" Dntt
1449758_at 0.87161 --- ---
1449759_at 1.07004 "DNA segment, Chr 11, ERATO Doi 326, expressed" D11Ertd326e
1449760_at 0.99411 Transcribed locus ---
1449761_at 0.98178 expressed sequence AA589418 AA589418
1449762_at 1.04571 Transcribed locus ---
1449763_at 1.07479 Doublesex and mab-3 related transcription factor 3 Dmrt3
1449764_x_at 0.96625 "synovial sarcoma, X breakpoint 2 interacting protein" Ssx2ip
1449765_at 0.95614 expressed sequence C76628 C76628
1449766_at 0.96148 synaptotagmin VI Syt6
1449767_x_at 0.83181 synaptotagmin VI Syt6
1449768_at 0.96594 --- ---
1449769_at 0.97438 --- ---
1449770_x_at 1.03359 "DNA segment, Chr 16, Brigham & Women's Genetics 1494 expressed"
D16Bwg1494e
1449771_at 0.91482 --- ---
1449772_at 0.87024 "TBC1 domain family, member 14" Tbc1d14

1449773_s_at 1.11048 growth arrest and DNA-damage-inducible 45 beta Gadd45b
1449774_at 1.00728 --- ---
1449775_x_at 0.96329 "solute carrier family 35, member A4 /// similar to solute carrier family 35, member A4" Slc35a4 /// LOC545750
1449776_at 1.08849 --- ---
1449777_at 0.93067 Dihydrouridine synthase 4-like (S. cerevisiae) Dus4l
1449778_at 1.03720 --- ---
1449779_at 1.04076 --- ---
1449780_at 0.92372 --- ---
1449781_at 0.91221 --- ---
1449782_at 1.03047 expressed sequence AA517650 AA517650
1449783_at 0.97605 --- ---
1449784_at 1.06195 --- ---
1449785_at 1.04806 expressed sequence AA414993 AA414993
1449786_at 0.95441 "zinc finger, DHHC domain containing 7" Zdhhc7
1449787_at 1.15336 --- ---
1449788_at 0.97916 --- ---
1449789_x_at 0.93799 "lymphocyte antigen 6 complex, locus G6C" Ly6g6c
1449790_at 0.93224 --- ---
1449791_x_at 0.90606 RIKEN cDNA 9130023D20 gene 9130023D20Rik
1449792_at 0.89780 --- ---
1449793_at 0.93211 --- ---
1449794_x_at 0.94730 Cathepsin F Ctsf
1449795_at 1.01810 "DNA segment, Chr 2, ERATO Doi 63, expressed" D2Ertd63e
1449796_at 0.98465 Peripherin 1 Prph1
1449797_x_at 0.87642 Peripherin 1 Prph1
1449798_at 0.98413 Transcribed locus ---
1449799_s_at 1.09558 plakophilin 2 Pkp2
1449800_x_at 0.97623 PHD finger protein 7 Phf7
1449801_at 1.09474 --- ---
1449802_x_at 0.94895 "3 days neonate thymus cDNA, RIKEN full-length enriched library, clone: A630030O14 product:unclassifiable, full insert sequence" ---
1449803_x_at 0.97739 "transporter 1, ATP-binding cassette, sub-family B (MDR/TAP)" Tap1
1449804_at 0.93013 phenylethanolamine-N-methyltransferase Pnmt
1449805_at 1.01309 "Protease, serine, 29" Prss29
1449806_at 0.84630 Transcribed locus ---
1449807_x_at 1.09032 --- ---
1449808_at 1.06404 Transcribed locus ---
1449813_at 0.77502 zinc finger protein 30 Zfp30
1449814_at 0.96559 frequently rearranged in advanced T-cell lymphomas Frat1
1449815_a_at 0.82260 single-stranded DNA binding protein 2 Ssbp2
1449816_at 0.84651 "sulfotransferase family 5A, member 1" Sult5a1
1449817_at 0.94664 "ATP-binding cassette, sub-family B (MDR/TAP), member 11" Abcb11

1449818_at 0.78606 "ATP-binding cassette, sub-family B (MDR/TAP), member 4" Abcb4
1449819_at 0.98840 disrupted meiotic cDNA 1 homolog Dmc1h
1449820_at 1.00187 cortistatin Cort
1449821_a_at 1.07222 RIKEN cDNA 0610016J10 gene /// hypothetical LOC619602
0610016J10Rik /// LOC619602
1449822_at 1.13658 atonal homolog 1 (Drosophila) Atoh1
1449823_at 0.95872 dachshund 2 (Drosophila) Dach2
1449824_at 0.98977 "proteoglycan 4 (megakaryocyte stimulating factor, articular superficial zone
protein)" Prg4
1449825_at 1.11280 actin-related protein T1 Actrt1
1449826_a_at 1.07388 fibroblast growth factor 2 Fgf2
1449827_at 0.95038 aggrecan 1 Agc1
1449828_at 1.01762 prostaglandin F receptor Ptgfr
1449829_at 0.95747 integrin beta 2-like Itgb2l
1449830_at 1.02297 prolactin-like protein J Prlpj
1449831_at 1.00261 Tctex1 domain containing 1 Tctex1d1
1449832_at 0.85633 RIKEN cDNA 1700091H14 gene 1700091H14Rik
1449833_at 0.92531 small proline-rich protein 2F Sprr2f
1449834_at 0.87828 "PDZ domain containing, X chromosome" Pdzx
1449835_at 0.97004 programmed cell death 1 Pcd1
1449836_x_at 1.05574 Bcl2-interacting killer Bik
1449837_at 0.96210 sperm equatorial segment protein 1 Spesp1
1449838_at 0.93591 cysteine-rich secretory protein 3 Crisp3
1449839_at 1.03410 --- ---
1449840_at 0.96292 "syntrophin, basic 2" Sntb2
1449841_at 1.05365 kinesin family member 3A Kif3a
1449842_at 0.93594 RIKEN cDNA 1810059G22 gene 1810059G22Rik
1449843_at 1.03399 "ST8 alpha-N-acetyl-neuraminide alpha-2,8-sialyltransferase 2" St8sia2
1449844_at 0.99356 "solute carrier organic anion transporter family, member 1a1" Slco1a1
1449845_a_at 0.88109 Eph receptor B4 Ephb4
1449846_at 0.93566 "eosinophil-associated, ribonuclease A family, member 2" Ear2
1449847_a_at 1.06998 "procollagen, type IV, alpha 3 (Goodpasture antigen) binding protein"
Col4a3bp
1449848_at 1.00560 "guanine nucleotide binding protein, alpha 14" Gna14
1449849_a_at 1.00142 F-box and leucine-rich repeat protein 6 Fbxl6
1449850_at 0.93107 "signal peptide, CUB domain, EGF-like 1" Scube1
1449851_at 0.89902 period homolog 1 (Drosophila) Per1
1449852_a_at 1.07750 EH-domain containing 4 Ehd4
1449853_at 0.99090 sideroflexin 2 /// similar to Sideroflexin-2 Sfxn2 /// LOC677143
1449854_at 0.84493 "nuclear receptor subfamily 0, group B, member 2" Nr0b2
1449855_s_at 1.11494 ubiquitin carboxyl-terminal esterase L3 (ubiquitin thiolesterase) /// ubiquitin
carboxyl-terminal esterase L4 Uchl3 /// Uchl4
1449856_at 1.02652 regulator of G-protein signaling 18 Rgs18

1449857_at 0.92101 RIKEN cDNA 1200011I18 gene 1200011I18Rik
1449858_at 0.94774 CD86 antigen Cd86
1449859_at 0.85457 golgi transport 1 homolog B (*S. cerevisiae*) Golt1b
1449860_at 0.99501 "HIG1 domain family, member 1B" Higd1b
1449861_at 1.05992 NIMA (never in mitosis gene a)-related expressed kinase 4 Nek4
1449862_a_at 1.10813 phosphatidylinositol 4-kinase type 2 beta Pi4k2b
1449863_a_at 1.05253 distal-less homeobox 5 Dlx5
1449864_at 0.85533 interleukin 4 Il4
1449865_at 0.68454 "sema domain, immunoglobulin domain (Ig), short basic domain, secreted, (semaphorin) 3A" Sema3a
1449866_at 0.98844 synaptotagmin II Syt2
1449867_at 0.90721 homeo box C9 Hoxc9
1449868_at 0.93685 T-box 6 Tbx6
1449869_at 1.05970 pre-B lymphocyte gene 1 Vpreb1
1449870_a_at 1.06841 "ATPase, H⁺ transporting, lysosomal V0 subunit A2" Atp6v0a2
1449871_at 0.76086 T-box18 Tbx18
1449872_at 0.93417 heat shock protein 3 Hspb3
1449873_at 1.11183 bone morphogenetic protein 8a Bmp8a
1449874_at 1.34939 lymphocyte antigen 96 Ly96
1449875_s_at 1.15681 "histocompatibility 2, T region locus 22 /// histocompatibility 2, T region locus 9" H2-T22 /// H2-T9
1449876_at 0.97475 "protein kinase, cGMP-dependent, type I" Prkg1
1449877_s_at 0.77341 kinesin family member C1 Kifc1
1449878_a_at 0.89806 "solute carrier family 12, member 6" Slc12a6
1449879_at 0.97545 RIKEN cDNA 1700008F21 gene 1700008F21Rik
1449880_s_at 1.03709 "bone gamma-carboxyglutamate protein, related sequence 1 /// bone gamma carboxyglutamate protein 1 /// bone gamma-carboxyglutamate protein 2" Bglap-rs1 /// Bglap1 /// Bglap2
1449881_a_at 0.95027 calcium-sensing receptor Casr
1449882_a_at 1.11413 calcium-sensing receptor Casr
1449883_at 0.93689 FXYD domain-containing ion transport regulator 2 Fxyd2
1449884_at 0.93391 EF-hand calcium binding domain 2 Efcab2
1449885_at 0.94469 transmembrane protein 47 Tmem47
1449886_a_at 1.02717 translocase of inner mitochondrial membrane 9 homolog (yeast) Timm9
1449887_at 0.79138 RIKEN cDNA 2310010I16 gene 2310010I16Rik
1449888_at 0.95866 endothelial PAS domain protein 1 Epas1
1449889_a_at 0.97711 OCIA domain containing 1 Ociad1
1449890_at 1.01687 "UDP glucuronosyltransferase 2 family, polypeptide B37" Ugt2b37
1449891_a_at 1.02068 "discoidin, CUB and LCCL domain containing 2" Dcbl2
1449892_at 1.05097 lysozyme-like 1 Lyzl1
1449893_a_at 1.27503 leucine-rich repeats and immunoglobulin-like domains 1 Lrig1
1449894_at 1.03361 leucine rich repeat containing 18 Lrrc18
1449895_at 0.82995 preproacrosin Acr

1449896_at 0.92593 --- ---
1449897_a_at 0.90684 mature T-cell proliferation 1 Mtcp1
1449898_at 0.84239 septin 1 1-Sep
1449899_at 0.87767 "glutamate receptor, ionotropic, NMDA3B" Grin3b
1449900_at 1.00514 galactosylceramidase Galc
1449901_a_at 1.24626 mitogen-activated protein kinase kinase kinase 6 Map3k6
1449902_at 0.99129 RIKEN cDNA 1110058A15 gene 1110058A15Rik
1449903_at 1.09236 cytotoxic and regulatory T cell molecule Crtam
1449904_at 1.00833 dopey family member 2 Dopey2
1449905_at 1.01615 "C-type lectin domain family 4, member f" Clec4f
1449906_at 0.94552 "selectin, platelet" Selp
1449907_at 0.82698 "beta-carotene 15,15'-monooxygenase" Bcmo1
1449908_at 0.83056 gastric inhibitory polypeptide Gip
1449909_at 1.11695 RIKEN cDNA 2010005H15 gene 2010005H15Rik
1449910_at 0.93072 RIKEN cDNA 2210418O10 gene 2210418O10Rik
1449911_at 0.83788 lymphocyte-activation gene 3 Lag3
1449912_at 0.95187 "synovial sarcoma, X member B, breakpoint 1 /// similar to SSXB1 protein"
Ssxb1 /// LOC631002
1449913_at 0.91722 zinc finger protein 2 Zfp2
1449914_at 0.92560 RIB43A domain with coiled-coils 1 Ribc1
1449915_at 1.03360 zinc finger protein 202 Zfp202
1449916_at 1.04343 pre-B-cell leukemia transcription factor 4 Pbx4
1449917_at 0.88223 paired-like homeodomain transcription factor 3 Pitx3
1449918_at 1.01552 CD209g antigen Cd209g
1449919_at 0.97296 keratin associated protein 6-2 Krtap6-2
1449920_at 0.98555 "cytochrome P450, family 19, subfamily a, polypeptide 1" Cyp19a1
1449921_s_at 0.98815 copine VI Cpne6
1449922_at 0.99841 RIKEN cDNA 4933425L06 gene 4933425L06Rik
1449923_at 0.81936 RIKEN cDNA 1700018F24 gene 1700018F24Rik
1449924_at 0.81551 proteoglycan 3 Prg3
1449925_at 1.05540 chemokine (C-X-C motif) receptor 3 Cxcr3
1449926_at 0.84638 "tumor necrosis factor (ligand) superfamily, member 7" Tnfsf7
1449927_at 1.11040 S100 calcium binding protein A5 S100a5
1449928_at 0.95065 dynein light chain Tctex-type 3 Dynlt3
1449929_at 0.88436 dynein light chain Tctex-type 3 Dynlt3
1449930_a_at 1.03270 "signal sequence receptor, beta" Ssr2
1449931_at 0.77115 cytoplasmic polyadenylation element binding protein 4 Cpeb4
1449932_at 0.97477 "casein kinase 1, delta" Csnk1d
1449933_a_at 0.98581 RIKEN cDNA 5730449L18 gene 5730449L18Rik
1449934_at 1.09591 purine rich element binding protein A Pura
1449935_a_at 1.03912 "DnaJ (Hsp40) homolog, subfamily A, member 3" Dnaja3
1449936_at 0.99923 RIKEN cDNA 8430419L09 gene 8430419L09Rik
1449937_at 0.95805 placental protein 11 related Pp11r

1449938_at 1.04042 placental protein 11 related Pp11r
1449939_s_at 0.81588 delta-like 1 homolog (Drosophila) Dlk1
1449940_a_at 0.95547 "eukaryotic translation initiation factor 2B, subunit 4 delta" Eif2b4
1449941_at 1.10120 myosin IE Myo1e
1449942_a_at 0.95943 integrin linked kinase Ilk
1449943_at 0.88693 lunatic fringe gene homolog (Drosophila) Lfng
1449944_a_at 1.22901 "Sec61, alpha subunit 2 (S. cerevisiae)" Sec61a2
1449945_at 1.15758 "peroxisome proliferative activated receptor, gamma, coactivator 1 beta"
Ppargc1b
1449946_a_at 0.98247 zinc finger protein 593 Zfp593
1449947_s_at 1.03808 AT motif binding factor 1 Atbf1
1449948_at 0.96595 ERO1-like beta (S. cerevisiae) Ero1lb
1449949_a_at 1.06824 coxsackievirus and adenovirus receptor Cxadr
1449950_at 1.00877 RIKEN cDNA 5830415L20 gene 5830415L20Rik
1449951_at 1.07002 nuclear factor of kappa light polypeptide gene enhancer in B-cells inhibitor-like
1 Nfkbil1
1449952_s_at 0.95823 "transmembrane protease, serine 8 (intestinal)" Tmprss8
1449953_at 1.03735 integrin beta 3 binding protein (beta3-endonexin) Itgb3bp
1449954_at 0.94549 hyaluronoglucosaminidase 1 /// N-acetyltransferase 6 Hyal1 /// Nat6
1449955_at 1.03133 "calcium channel, voltage-dependent, alpha 1F subunit" Cacna1f
1449956_at 1.07386 "protein kinase C, epsilon" Prkce
1449957_at 0.94801 "protein tyrosine phosphatase, receptor type, V" Ptprv
1449958_a_at 0.90543 fibroblast growth factor 14 Fgf14
1449959_x_at 0.92121 small proline rich-like 9 Sprrl9
1449960_at 0.83376 neuronal pentraxin 2 Nptx2
1449961_at 1.06503 rabphilin 3A Rph3a
1449962_at 1.01836 preferentially expressed antigen in melanoma-like 3 /// expressed sequence
AV320801 /// similar to preferentially expressed antigen in melanoma Pramel3 /// AV320801 ///
LOC666040
1449963_at 1.02562 RIKEN cDNA 2310040M23 gene 2310040M23Rik
1449964_a_at 0.98637 malonyl-CoA decarboxylase Mlycd
1449965_at 1.09264 mast cell protease 8 Mcpt8
1449966_s_at 0.97291 calcium binding protein 39-like Cab39l
1449967_at 1.05009 single-minded homolog 1 (Drosophila) Sim1
1449968_s_at 1.09651 acyl-CoA thioesterase 9 /// acyl-CoA thioesterase 10 Acot9 /// Acot10
1449969_at 0.99802 tropomodulin 4 Tmod4
1449970_at 0.84321 calpain 12 Capn12
1449971_a_at 0.90088 CD209f antigen Cd209f
1449972_s_at 0.83266 zinc finger protein 97 /// cDNA sequence BC018101 Zfp97 /// BC018101
1449973_a_at 1.02104 "proline, glutamic acid and leucine rich protein 1" Pelp1
1449974_at 1.04953 "glyceraldehyde-3-phosphate dehydrogenase, spermatogenic" Gapdhs
1449975_a_at 1.11933 parkin Park2
1449976_a_at 0.95604 G protein-coupled receptor 35 Gpr35

1449977_at 0.96740 early growth response 4 Egr4
1449978_at 0.96965 "zinc finger protein 2, Y linked" Zfy2
1449979_a_at 0.96988 "sparc/osteonectin, cwcv and kazal-like domains proteoglycan 3" Spock3
1449980_a_at 1.00114 "gamma-aminobutyric acid (GABA-A) receptor, subunit delta" Gabrd
1449981_a_at 1.03951 N-acetyltransferase 2 (arylamine N-acetyltransferase) Nat2
1449982_at 0.92788 interleukin 11 Il11
1449983_a_at 1.10266 "NAD(P)H dehydrogenase, quinone 2" Nqo2
1449984_at 1.83016 chemokine (C-X-C motif) ligand 2 Cxcl2
1449985_at 0.93909 RIKEN cDNA 1700072E05 gene 1700072E05Rik
1449986_at 0.95273 RIKEN cDNA 2310034C09 gene 2310034C09Rik
1449987_at 0.96088 anaplastic lymphoma kinase Alk
1449988_at 0.88851 "GTPase, IMAP family member 1" Gimap1
1449989_at 1.09992 mast cell protease 2 Mcpt2
1449990_at 0.84942 interleukin 2 /// similar to Interleukin-2 precursor (IL-2) (T-cell growth factor) (TCGF) Il2 /// LOC630222
1449991_at 1.01119 CD244 natural killer cell receptor 2B4 /// similar to Natural killer cell receptor 2B4 precursor (NKR2B4) (NK cell type I receptor protein 2B4) (CD244 antigen) (Non MHC restricted killing associated) Cd244 /// LOC677008
1449992_at 0.97926 "protease, serine, 29" Prss29
1449993_at 1.19290 cDNA sequence AF119384 AF119384
1449994_at 1.03550 epithelial mitogen Epgn
1449995_at 1.03699 RIKEN cDNA 4933421I07 gene 4933421I07Rik
1449996_a_at 1.11105 "tropomyosin 3, gamma" Tpm3
1449997_at 0.96757 "tropomyosin 3, gamma" Tpm3
1449998_at 1.03022 "NK-3 transcription factor, locus 1 (Drosophila)" Nkx3-1
1449999_a_at 0.82722 "calcium channel, voltage-dependent, alpha2/delta subunit 1" Cacna2d1
1450000_at 1.06163 "NACHT, leucine rich repeat and PYD containing 4C" Nalp4c
1450001_a_at 1.06678 Usher syndrome 1C homolog (human) Ush1c
1450002_at 0.80017 genomic screened homeo box 1 Gsh1
1450003_at 1.03196 "adrenergic receptor, alpha 2b" Adra2b
1450004_at 1.00859 thymic stromal lymphopoietin Tslp
1450005_x_at 0.97401 "EGF-like-domain, multiple 9" Egfl9
1450006_at 0.93803 nuclear receptor coactivator 4 /// similar to nuclear receptor coactivator 4 Ncoa4 /// LOC627557
1450007_at 0.90517 RIKEN cDNA 1500003O03 gene 1500003O03Rik
1450008_a_at 0.98433 "catenin (cadherin associated protein), beta 1" Ctnnb1
1450009_at 1.07220 lactotransferrin Ltf
1450010_at 1.20767 hydroxysteroid (17-beta) dehydrogenase 12 Hsd17b12
1450011_at 1.13464 hydroxysteroid (17-beta) dehydrogenase 12 Hsd17b12
1450012_x_at 1.02074 "3-monooxygenase/tryptophan 5-monooxygenase activation protein, gamma polypeptide" Ywhag
1450013_at 0.92501 RIKEN cDNA 2900073G15 gene 2900073G15Rik
1450014_at 0.86431 claudin 1 Cldn1

1450015_x_at 1.13565 sphingosine-1-phosphate phosphatase 1 Sgpp1
1450016_at 1.17442 cyclin G1 Ccng1
1450017_at 1.09791 cyclin G1 Ccng1
1450018_s_at 0.98044 "solute carrier family 25, member 30" Slc25a30
1450019_at 1.22101 chemokine (C-X3-C) receptor 1 Cx3cr1
1450020_at 0.96980 chemokine (C-X3-C) receptor 1 Cx3cr1
1450021_at 0.92386 ubiquilin 2 Ubqln2
1450022_at 0.99527 GTP binding protein 1 Gtpbp1
1450023_at 0.95078 GTP binding protein 1 Gtpbp1
1450024_at 0.93700 --- ---
1450025_at 0.87906 par-6 partitioning defective 6 homolog gamma (C. elegans) Pard6g
1450026_a_at 0.93369 "UDP-GlcNAc:betaGal beta-1,3-N-acetylglucosaminyltransferase 2" B3gnt2
1450027_at 0.70004 syndecan 3 Sdc3
1450028_a_at 0.91573 LanC (bacterial lantibiotic synthetase component C)-like 2 Lancl2
1450029_s_at 1.11340 integrin alpha 9 Itga9
1450030_at 0.93758 dynactin 4 Dctn4
1450031_at 0.96823 "AF4/FMR2 family, member 4" Aff4
1450032_at 0.96909 "solute carrier organic anion transporter family, member 2a1" Slco2a1
1450033_a_at 0.91120 signal transducer and activator of transcription 1 Stat1
1450034_at 1.21133 signal transducer and activator of transcription 1 Stat1
1450035_a_at 0.98234 PRP40 pre-mRNA processing factor 40 homolog A (yeast) Prpf40a
1450036_at 0.76495 serum/glucocorticoid regulated kinase 3 Sgk3
1450037_at 0.91664 "ubiquitin specific peptidase 9, X chromosome" Usp9x
1450038_s_at 0.97343 "ubiquitin specific peptidase 9, X chromosome" Usp9x
1450039_at 1.09340 "ubiquitin specific peptidase 9, X chromosome" Usp9x
1450040_at 0.85997 tissue inhibitor of metalloproteinase 2 Timp2
1450041_a_at 0.98137 tubby candidate gene Tub
1450042_at 1.00985 aristaless related homeobox gene (Drosophila) Arx
1450043_at 1.01887 frizzled homolog 7 (Drosophila) Fzd7
1450044_at 0.87551 frizzled homolog 7 (Drosophila) Fzd7
1450045_at 1.40421 serine/arginine repetitive matrix 1 Srrm1
1450046_at 1.00226 transmembrane protein 59 Tmem59
1450047_at 0.99244 heparan sulfate 6-O-sulfotransferase 2 Hs6st2
1450048_a_at 0.89627 "isocitrate dehydrogenase 2 (NADP+), mitochondrial" Idh2
1450049_a_at 1.14357 histone cell cycle regulation defective homolog A (S. cerevisiae) Hira
1450050_at 1.11276 histone cell cycle regulation defective homolog A (S. cerevisiae) Hira
1450051_at 0.97876 alpha thalassemia/mental retardation syndrome X-linked homolog (human) Atrx
1450052_at 1.05327 kinesin family member 2A Kif2a
1450053_at 1.06596 Kinesin family member 2A Kif2a
1450054_at 0.88562 adducin 1 (alpha) Add1
1450055_at 1.26657 visinin-like 1 Vsnl1
1450056_at 1.08183 adenomatosis polyposis coli Apc
1450057_at 1.01732 RIKEN cDNA 2610042O14 gene 2610042O14Rik

1450058_at	0.69128	aspartate-beta-hydroxylase	Asph
1450059_at	0.82054	---	---
1450060_at	1.00597	polymeric immunoglobulin receptor	Pigr
1450061_at	0.97111	ectodermal-neural cortex 1	Enc1
1450062_a_at	0.81461	"melanoma antigen, family D, 1"	Maged1
1450063_at	0.93728	formin 2	Fmn2
1450064_at	0.90896	---	---
1450065_at	0.95189	adenylate cyclase 7	Adcy7
1450066_at	0.99029	ubiquitin protein ligase E3 component n-recognin 1	Ubr1
1450067_a_at	0.96803	RIKEN cDNA 1810034K20 gene	1810034K20Rik
1450068_at	0.82480	"bromodomain adjacent to zinc finger domain, 1B"	Baz1b
1450069_a_at	0.98138	"CUG triplet repeat, RNA binding protein 2"	Cugbp2
1450070_s_at	1.00632	p21 (CDKN1A)-activated kinase 1	Pak1
1450071_at	1.07467	"ash1 (absent, small, or homeotic)-like (Drosophila)"	Ash11
1450072_at	1.06258	"ash1 (absent, small, or homeotic)-like (Drosophila)"	Ash11
1450073_at	0.90061	kinesin family member 3B	Kif3b
1450074_at	0.95353	kinesin family member 3B	Kif3b
1450075_at	1.00517	"polymerase (DNA directed), eta (RAD 30 related)"	Polh
1450076_at	0.96623	RIKEN cDNA 4933411K20 gene	4933411K20Rik
1450077_at	0.98161	chromodomain helicase DNA binding protein 1	Chd1
1450078_at	0.97656	Nik related kinase	Nrk
1450079_at	1.03340	Nik related kinase	Nrk
1450080_at	0.94066	CAAX box 1 homolog C (human)	Cxx1c
1450081_x_at	1.09677	glucose phosphate isomerase 1	Gpi1
1450082_s_at	1.30721	ets variant gene 5	Etv5
1450083_at	0.94787	"CCR4-NOT transcription complex, subunit 4"	Cnot4
1450084_s_at	0.85979	influenza virus NS1A binding protein	Ivns1abp
1450085_at	0.89121	---	---
1450086_at	0.93941	glucocorticoid modulatory element binding protein 1	Gmeb1
1450087_a_at	1.30701	nucleolar and coiled-body phosphoprotein 1	Nolc1
1450088_a_at	0.96978	myelin-associated oligodendrocytic basic protein	Mobp
1450089_a_at	0.85957	"signal recognition particle receptor, B subunit"	Srprb
1450090_at	0.98690	zinc finger protein 101	Zfp101
1450091_at	1.00764	immunoglobulin mu binding protein 2	Ighmbp2
1450092_at	0.93334	immunoglobulin mu binding protein 2	Ighmbp2
1450093_s_at	1.04816	zinc finger and BTB domain containing 7a	Zbtb7a
1450094_at	0.95888	RAN binding protein 17	Ranbp17
1450095_a_at	0.89205	"acylphosphatase 1, erythrocyte (common) type"	Acyp1
1450096_at	0.97357	"guanine nucleotide binding protein, alpha 12"	Gna12
1450097_s_at	0.81759	"guanine nucleotide binding protein, alpha 12"	Gna12
1450098_at	1.03844	RIKEN cDNA 0610010D20 gene	0610010D20Rik
1450099_a_at	1.11839	"glucosidase, beta, acid"	Gba
1450100_a_at	0.98172	transcription elongation regulator 1 (CA150)	Tcerg1

1450101_a_at 0.86571 "membrane associated guanylate kinase, WW and PDZ domain containing 3" Magi3

1450102_a_at 0.94814 autocrine motility factor receptor Amfr

1450103_a_at 0.93618 "pleckstrin homology, Sec7 and coiled-coil domains 2" Pscd2

1450104_at 0.77569 a disintegrin and metallopeptidase domain 10 Adam10

1450105_at 0.90209 --- ---

1450106_a_at 0.82011 Ena-vasodilator stimulated phosphoprotein Evl

1450107_a_at 0.92110 renin binding protein Renbp

1450108_at 0.92834 kinesin family member 1A Kif1a

1450109_s_at 1.08688 "ATP-binding cassette, sub-family C (CFTR/MRP), member 2" Abcc2

1450110_at 0.82291 "alcohol dehydrogenase 7 (class IV), mu or sigma polypeptide" Adh7

1450111_a_at 1.12024 nudix (nucleoside diphosphate linked moiety X)-type motif 8 Nudt8

1450112_a_at 1.00422 growth arrest specific 2 Gas2

1450113_at 0.92572 --- ---

1450114_at 0.94446 kinase suppressor of ras 1 Ksr1

1450115_at 0.87157 "guanine nucleotide binding protein, alpha q polypeptide" Gnaq

1450116_at 1.00835 "DNA segment, Chr 3, ERATO Doi 300, expressed" D3Ertd300e

1450117_at 0.67267 transcription factor 3 Tcf3

1450118_a_at 0.81605 "troponin T3, skeletal, fast" Tnnt3

1450119_at 1.35404 dystonin Dst

1450120_at 0.90847 "sodium channel, voltage-gated, type I, alpha" Scn1a

1450121_at 0.93474 "sodium channel, voltage-gated, type I, alpha" Scn1a

1450122_at 0.98433 "protein tyrosine phosphatase, receptor type, G" Ptpg

1450123_at 0.90976 "ryanodine receptor 2, cardiac" Ryr2

1450124_a_at 0.96013 "ATPase, Ca++ transporting, ubiquitous" Atp2a3

1450125_at 1.10278 GATA binding protein 5 Gata5

1450126_at 1.06457 GATA binding protein 5 Gata5

1450127_a_at 1.01080 glucagon receptor Gcgr

1450128_at 0.99451 "phospholipase A2, group IIA (platelets, synovial fluid)" Pla2g2a

1450129_a_at 0.91206 suppressor of cytokine signaling 6 Socs6

1450130_at 0.91199 xenotropic and polytropic retrovirus receptor 1 Xpr1

1450131_a_at 0.93141 B-box and SPRY domain containing Bspry

1450132_at 1.02537 human immunodeficiency virus type I enhancer binding protein 3 Hivep3

1450133_at 0.85493 "UDP glucuronosyltransferase 2 family, polypeptide A3" Ugt2a3

1450134_at 1.06455 lysyl oxidase-like 4 Loxl4

1450135_at 1.03021 frizzled homolog 3 (Drosophila) Fzd3

1450136_at 0.98737 CD38 antigen Cd38

1450137_at 0.93648 protein O-fucosyltransferase 1 Pofut1

1450138_a_at 1.13901 "serine (or cysteine) peptidase inhibitor, clade B, member 6a" Serpinb6a

1450139_at 1.18671 endoplasmic reticulum (ER) to nucleus signalling 2 Ern2

1450140_a_at 1.11983 cyclin-dependent kinase inhibitor 2A Cdkn2a

1450141_at 1.01157 "ATP-binding cassette, sub-family G (WHITE), member 3" Abcg3

1450142_a_at 0.99318 "caspase recruitment domain family, member 14" Card14

1450143_at 1.01646 RAS guanyl releasing protein 1 Rasgrp1
1450144_at 1.08321 phospholipase A2 receptor 1 Pla2r1
1450145_at 0.91606 DNA binding protein with his-thr domain Dbpht1
1450146_at 0.94776 frequenin homolog (Drosophila) Freq
1450147_at 0.96117 neuronal pentraxin receptor Nptxr
1450148_at 1.11619 mucolipin 3 Mcoln3
1450149_a_at 0.94960 "protein phosphatase 1, catalytic subunit, gamma isoform /// similar to protein phosphatase 1 gamma2" Ppp1cc /// LOC434233
1450150_a_at 1.03583 ribosomal protein L13 Rpl13
1450151_at 0.89818 zinc finger protein 316 Zfp316
1450152_at 0.98650 zinc finger protein 316 Zfp316
1450153_at 0.98989 golgi associated PDZ and coiled-coil motif containing Gopc
1450154_at 1.02321 folate hydrolase Folh1
1450155_at 1.05591 integrin alpha 4 Itga4
1450156_a_at 0.97322 hyaluronan mediated motility receptor (RHAMM) Hmnr
1450157_a_at 1.01290 hyaluronan mediated motility receptor (RHAMM) Hmnr
1450158_at 0.98225 rad and gem related GTP binding protein 1 Rem1
1450159_at 0.88634 rad and gem related GTP binding protein 1 Rem1
1450160_at 0.97857 leukemia inhibitory factor Lif
1450161_at 1.00794 inhibitor of kappaB kinase gamma Ikbkg
1450162_at 0.90449 "D4, zinc and double PHD fingers, family 3" Dpf3
1450163_a_at 0.93648 Werner syndrome homolog (human) Wrn
1450164_at 1.02012 achaete-scute complex homolog-like 1 (Drosophila) Ascl1
1450165_at 1.38228 schlafen 2 Slfn2
1450166_at 0.81652 iduronate 2-sulfatase Ids
1450167_at 0.90699 "RAB37, member of RAS oncogene family" Rab37
1450168_at 1.05123 ankyrin repeat domain 12 Ankrd12
1450169_at 0.94192 "solute carrier family 4 (anion exchanger), member 4" Slc4a4
1450170_x_at 1.08781 "Histocompatibility 2, K1, K region" H2-K1
1450171_x_at 1.00232 granzyme E Gzme
1450172_at 0.93737 Pbx/knotted 1 homeobox Pknox1
1450173_at 1.13531 receptor (TNFRSF)-interacting serine-threonine kinase 2 Ripk2
1450174_at 0.91004 "protein tyrosine phosphatase, receptor type, T" Ptptrt
1450175_a_at 1.04360 cathepsin M Ctsm
1450176_at 0.84372 endoplasmic reticulum (ER) to nucleus signalling 1 Ern1
1450177_at 0.94285 "nerve growth factor receptor (TNFR superfamily, member 16)" Ngfr
1450178_at 1.06910 "bromodomain, testis-specific" Brdt
1450179_at 0.88630 sclerostin Sost
1450180_a_at 0.85389 "retinoic acid receptor, alpha" Rara
1450181_at 1.02341 cut-like 2 (Drosophila) Cutl2
1450182_at 0.91151 chloride channel Ka Clnka
1450183_a_at 1.06655 SH2B adaptor protein 3 Sh2b3
1450184_s_at 0.91915 thyrotroph embryonic factor Tef

1450185_a_at 0.98223 "potassium inwardly-rectifying channel, subfamily J, member 15" Kcnj15
 1450186_s_at 0.86057 "GNAS (guanine nucleotide binding protein, alpha stimulating) complex locus" Gnas
 1450187_a_at 0.90519 galactose-1-phosphate uridyl transferase Galt
 1450188_s_at 1.15022 "lipase, endothelial" Lipg
 1450189_at 1.08133 Kell blood group precursor (McLeod phenotype) homolog Xkh
 1450190_at 0.91492 zinc finger protein 106 Zfp106
 1450191_a_at 1.04473 SRY-box containing gene 13 Sox13
 1450192_at 0.91526 luteinizing hormone/choriogonadotropin receptor Lhcgr
 1450193_at 0.86582 "hyperpolarization-activated, cyclic nucleotide-gated K+ 1" Hcn1
 1450194_a_at 0.96187 myeloblastosis oncogene Myb
 1450195_at 0.81514 N-deacetylase/N-sulfotransferase (heparin glucosaminyl) 4 Ndst4
 1450196_s_at 0.94303 "glycogen synthase 1, muscle" Gys1
 1450197_at 1.05812 retinal pigment epithelium 65 Rpe65
 1450198_at 0.97277 dual specificity phosphatase 13 Dusp13
 1450199_a_at 1.15358 stabilin 1 Stab1
 1450200_s_at 0.89083 "colony stimulating factor 2 receptor, beta 1, low-affinity (granulocyte-macrophage) /// colony stimulating factor 2 receptor, beta 2, low-affinity (granulocyte-macrophage)" Csf2rb1 /// Csf2rb2
 1450201_at 1.01760 --- ---
 1450202_at 1.02599 "glutamate receptor, ionotropic, NMDA1 (zeta 1)" Grin1
 1450203_at 1.01544 SET and MYND domain containing 1 Smyd1
 1450204_a_at 0.80398 myoneurin Mynn
 1450205_at 0.88298 --- ---
 1450206_at 0.94979 deleted in liver cancer 1 Dlc1
 1450207_at 1.05583 leukemia inhibitory factor receptor Lifr
 1450208_a_at 1.04538 "engulfment and cell motility 1, ced-12 homolog (C. elegans)" Elmo1
 1450209_at 0.91262 homeo box D4 Hoxd4
 1450210_at 0.98563 bone morphogenetic protein 15 Bmp15
 1450211_at 0.93888 low density lipoprotein-related protein 1B (deleted in tumors) Lrp1b
 1450212_at 1.13071 formin-like 1 Fmn1
 1450213_at 0.93184 phosphodiesterase 7B Pde7b
 1450214_at 0.97965 adenosine A2b receptor Adora2b
 1450215_at 0.98279 recoverin Rcvrn
 1450216_at 1.08386 protocadherin beta 16 Pcdhb16
 1450217_at 0.76172 chemokine (C-C motif) ligand 28 Ccl28
 1450218_at 0.98168 chemokine (C-C motif) ligand 28 Ccl28
 1450219_at 1.01624 5-hydroxytryptamine (serotonin) receptor 1A Htr1a
 1450220_a_at 0.88857 SAM pointed domain containing ets transcription factor Spdef
 1450221_at 1.00373 RIKEN cDNA 4933417A18 gene 4933417A18Rik
 1450222_x_at 0.83903 kallikrein 1-related peptidase b4 Klk1b4
 1450223_at 1.04053 apoptotic peptidase activating factor 1 Apaf1
 1450224_at 0.49426 "procollagen, type IV, alpha 3" Col4a3

1450225_at	0.89090	insulin receptor	Insr
1450226_at	1.05941	prolactin receptor	Prlr
1450227_at	0.95492	ankyrin repeat domain 6	Ankrd6
1450228_a_at	0.94306	"phosphatidylinositol-4-phosphate 5-kinase, type 1 gamma"	Pip5k1c
1450229_at	0.94957	"cofactor required for Sp1 transcriptional activation, subunit 2"	Crsp2
1450230_at	1.03417	"DNA segment, Chr 17, human D6S56E 3"	D17H6S56E-3
1450231_a_at	0.85152	baculoviral IAP repeat-containing 4	Birc4
1450232_at	0.97676	baculoviral IAP repeat-containing 4	Birc4
1450233_a_at	1.02402	radial spokehead-like 1	Rshl1
1450234_at	0.94335	"membrane-spanning 4-domains, subfamily A, member 6C"	Ms4a6c
1450235_at	1.12423	"FYVE, RhoGEF and PH domain containing 3"	Fgd3
1450236_at	0.89716	forkhead box O3a	Foxo3a
1450237_at	0.93161	deoxyribonuclease II beta	Dnase2b
1450238_at	1.04581	"glucosaminyl (N-acetyl) transferase 2, I-branching enzyme"	Gcnt2
1450239_at	1.02062	"glycine receptor, alpha 3 subunit"	Gla3
1450240_a_at	0.99064	synaptotagmin-like 1	Sytl1
1450241_a_at	1.07778	ecotropic viral integration site 2a	Evi2a
1450242_at	0.95523	toll-like receptor 5	Tlr5
1450243_a_at	0.88811	Down syndrome critical region gene 1-like 1	Dscr111
1450244_a_at	1.11720	mitogen activated protein kinase kinase kinase 2	Map4k2
1450245_at	1.11453	"solute carrier family 10, member 2"	Slc10a2
1450246_at	0.98930	fucosyltransferase 2	Fut2
1450247_a_at	0.94546	secretory carrier membrane protein 5	Scamp5
1450248_at	0.96064	a disintegrin and metallopeptidase domain 11	Adam11
1450249_s_at	1.09692	kinesin family member 5A /// kinesin family member 5C	Kif5a /// Kif5c
1450250_at	0.92490	OTU domain containing 7A	Otud7a
1450251_a_at	1.05727	ligand of numb-protein X 1	Lnx1
1450252_at	0.89730	"one cut domain, family member 1"	Onecut1
1450253_a_at	0.98650	mitogen activated protein kinase kinase kinase 4	Map3k4
1450254_at	1.04792	telomerase reverse transcriptase	Tert
1450255_at	1.10417	Cdc42 GTPase-activating protein	Cdgap
1450256_at	1.07874	cerberus 1 homolog (Xenopus laevis)	Cer1
1450257_at	0.97089	cerberus 1 homolog (Xenopus laevis)	Cer1
1450258_a_at	0.89957	"ELAV (embryonic lethal, abnormal vision, Drosophila)-like 4 (Hu antigen D)"	Elavl4
1450259_a_at	0.93565	signal transducer and activator of transcription 5A	Stat5a
1450260_at	0.98863	gastrin releasing peptide receptor	Grpr
1450261_a_at	1.00385	"solute carrier family 10 (sodium/bile acid cotransporter family), member 1"	Slc10a1
1450262_at	0.83913	cardiotrophin-like cytokine factor 1	Clcf1
1450263_at	0.89272	protocadherin beta 5	Pcdhb5
1450264_a_at	1.10634	choline kinase alpha	Chka
1450265_at	0.99970	MAD homolog 9 (Drosophila)	Smad9

1450266_at 0.95147 "sodium channel, voltage-gated, type X, alpha" Scn10a
1450267_at 0.86554 toll-like receptor 8 Tlr8
1450268_at 0.80198 fidgetin Fign
1450269_a_at 1.26799 "phosphofructokinase, liver, B-type" Pfkf
1450270_at 0.84245 protocadherin beta 11 Pcdhb11
1450271_at 1.05568 PTK6 protein tyrosine kinase 6 Ptk6
1450272_at 0.99748 "tumor necrosis factor (ligand) superfamily, member 8" Tnfsf8
1450273_at 0.97788 interleukin 1 receptor-like 2 Il1rl2
1450274_at 0.98861 RIKEN cDNA 1700108E19 gene 1700108E19Rik
1450275_x_at 1.08962 ubiquitin-conjugating enzyme E2F (putative) Ube2f
1450276_a_at 0.81252 scinderin Scin
1450277_at 0.91076 arginine vasopressin receptor 2 Avpr2
1450278_at 0.80437 tachykinin receptor 3 Tacr3
1450279_at 0.96695 prokineticin receptor 1 Prokr1
1450280_a_at 1.02773 retinal pigment epithelium derived rhodopsin homolog Rrh
1450281_a_at 0.96615 RIKEN cDNA 1700021K02 gene 1700021K02Rik
1450282_at 0.87579 fibroblast growth factor 4 Fgf4
1450283_at 0.99946 "ATPase, Cu⁺⁺ transporting, beta polypeptide" Atp7b
1450284_at 1.01926 "phosphodiesterase 3A, cGMP inhibited" Pde3a
1450285_at 1.05634 "ubiquitin-activating enzyme E1, Chr Y 1" Ube1y1
1450286_at 1.01947 natriuretic peptide receptor 3 Npr3
1450287_at 0.96235 neuronal PAS domain protein 3 Npas3
1450288_at 0.55421 cadherin 6 Cdh6
1450289_at 0.90755 forkhead box D2 Foxd2
1450290_at 0.92836 programmed cell death 1 ligand 2 Pcdcl1g2
1450291_s_at 1.00148 "membrane-spanning 4-domains, subfamily A, member 4C" Ms4a4c
1450292_a_at 0.94159 HORMA domain containing 1 Hormad1
1450293_at 0.98562 "melanoma antigen, family B, 3" Mageb3
1450294_a_at 0.91747 "potassium inwardly-rectifying channel, subfamily J, member 6" Kcnj6
1450295_s_at 1.22773 poliovirus receptor Pvr
1450296_at 1.03053 killer cell lectin-like receptor subfamily B member 1A Klrk1a
1450297_at 1.27460 interleukin 6 Il6
1450298_at 0.96388 "tumor necrosis factor (ligand) superfamily, member 14" Tnfsf14
1450299_at 0.96746 "cholinergic receptor, nicotinic, alpha polypeptide 7" Chrna7
1450300_at 0.88164 "gamma-aminobutyric acid (GABA-C) receptor, subunit rho 1" Gabrr1
1450301_at 0.99204 histocompatibility 28 H28
1450302_at 0.89139 RIKEN cDNA 4930470P17 gene 4930470P17Rik
1450303_at 0.99084 ventral anterior homeobox containing gene 2 Vax2
1450304_at 0.97525 "killer cell lectin-like receptor subfamily C, member 2" Klrc2
1450305_at 1.11308 RIKEN cDNA 4930591A17 gene 4930591A17Rik
1450306_at 0.88346 zona pellucida glycoprotein 1 Zp1
1450307_x_at 1.03873 "H2A histone family, member Y3" H2afy3
1450308_a_at 0.96055 5'-3' exoribonuclease 1 Xrn1

1450309_at 1.01033 astrotactin 2 Astn2
1450310_at 0.90675 "glutamate receptor, ionotropic, delta 2 (Grid2) interacting protein 1" Grid2ip
1450311_at 0.96984 "solute carrier family 8 (sodium/calcium exchanger), member 3" Slc8a3
1450312_at 1.00429 cadherin 20 Cdh20
1450313_at 0.91430 "taste receptor, type 1, member 2" Tas1r2
1450314_at 0.84055 DEAQ RNA-dependent ATPase Dqx1
1450315_at 1.07948 "vomeronasal 1 receptor, D14" V1rd14
1450316_at 0.87422 protocadherin beta 13 Pcdhb13
1450317_at 1.02003 "C-type lectin domain family 2, member g" Clec2g
1450318_a_at 1.21084 "purinergic receptor P2Y, G-protein coupled 2" P2ry2
1450319_at 0.96785 "gamma-aminobutyric acid (GABA-A) receptor, subunit beta 2" Gabrb2
1450320_at 0.95167 RIKEN cDNA 4931432E15 gene 4931432E15Rik
1450321_at 1.04987 zinc finger protein 354C Zfp354c
1450322_s_at 0.94866 schlafen 3 /// schlafen 4 Slfn3 /// Slfn4
1450323_at 1.06530 RIKEN cDNA 5730493B19 gene 5730493B19Rik
1450324_a_at 0.90014 RIKEN cDNA 4930521E07 gene 4930521E07Rik
1450325_at 0.87206 angiopoietin 4 Angpt4
1450326_at 1.11929 src homology 2 domain-containing transforming protein C3 Shc3
1450327_at 0.96570 "purinergic receptor P2X-like 1, orphan receptor" P2rx11
1450328_at 0.98701 "Adult male testis cDNA, RIKEN full-length enriched library, clone:1700129L04 product:hypothetical protein, full insert sequence" ---
1450329_a_at 0.95557 "arrestin 3, retinal" Arr3
1450330_at 1.02755 interleukin 10 Il10
1450331_s_at 0.99716 "vomeronasal 2, receptor, 4 /// vomeronasal 2, receptor, 5 /// similar to vomeronasal 2, receptor, 4 /// similar to vomeronasal 2, receptor, 15" V2r4 /// V2r5 /// LOC625605 /// LOC665552
1450332_s_at 0.90165 flavin containing monooxygenase 5 Fmo5
1450333_a_at 0.91017 GATA binding protein 2 Gata2
1450334_at 0.85837 interleukin 21 Il21
1450335_at 0.85002 --- ---
1450336_at 1.08693 non-selective cation channel 1 Nscn1
1450337_a_at 1.07295 NIMA (never in mitosis gene a)-related expressed kinase 8 Nek8
1450338_x_at 0.96852 "vomeronasal 2, receptor, 8 /// vomeronasal 2, receptor, 9" V2r8 /// V2r9
1450339_a_at 0.93318 B-cell leukemia/lymphoma 11B Bcl11b
1450340_a_at 0.86170 chloride channel Kb Clcnkb
1450341_at 1.03990 protocadherin beta 8 Pcdhb8
1450342_at 1.09095 bone morphogenetic protein 8b Bmp8b
1450343_at 0.96225 "vomeronasal 1 receptor, D1" V1rd1
1450344_a_at 1.06827 prostaglandin E receptor 3 (subtype EP3) Ptger3
1450345_at 1.06556 ubiquilin 1 Ubqln1
1450346_at 0.99145 G-protein-coupled receptor 50 Gpr50
1450347_at 0.98087 synaptotagmin X Syt10
1450348_at 1.07526 "solute carrier family 19 (sodium/hydrogen exchanger), member 3" Slc19a3

1450349_at 0.90672 syntaxin 1B1 Stx1b1
 1450350_a_at 0.95799 Jun dimerization protein 2 Jundm2
 1450351_a_at 0.97587 restin (Reed-Steinberg cell-expressed intermediate filament-associated protein)
 Rsn
 1450352_at 1.03741 melatonin receptor 1A Mtnr1a
 1450353_at 0.95436 "solute carrier family 25 (mitochondrial carrier, peroxisomal membrane
 protein), member 17" Slc25a17
 1450354_a_at 1.06576 phosphatidylserine synthase 2 Ptdss2
 1450355_a_at 1.00365 "capping protein (actin filament), gelsolin-like" Capg
 1450356_at 1.00232 thyrotropin releasing hormone receptor 2 Trhr2
 1450357_a_at 0.97714 chemokine (C-C motif) receptor 6 Ccr6
 1450358_at 0.92878 ADP-ribosylation factor-like 4D Arl4d
 1450359_at 1.01386 fucosyltransferase 1 Fut1
 1450360_at 1.02475 "phospholipase C, gamma 1" Plcg1
 1450361_at 0.98347 paired like homeodomain factor 1 Prop1
 1450362_at 1.19685 --- ---
 1450363_at 0.73011 ankyrin repeat and SOCS box-containing protein 7 Asb7
 1450364_a_at 0.94793 hepatitis A virus cellular receptor 1 Havcr1
 1450365_at 0.94273 myozenin 3 Myoz3
 1450366_at 0.83887 "harakiri, BCL2 interacting protein (contains only BH3 domain)" Hrk
 1450367_at 0.96696 --- ---
 1450368_a_at 1.00415 "protein phosphatase 3, regulatory subunit B, alpha isoform (calcineurin B,
 type I)" Ppp3r1
 1450369_at 1.03281 folliculogenesis specific basic helix-loop-helix Figla
 1450370_a_at 1.00196 Kv channel interacting protein 4 Kcnp4
 1450371_at 0.95608 "thyroid stimulating hormone, beta subunit" Tshb
 1450372_a_at 1.14830 ribosomal protein L18 Rpl18
 1450373_at 0.97373 proline-rich protein 15 Prp15
 1450374_at 1.01104 tuberoinfundibular peptide of 39 residues (TIP39) preprohormone Tifp39
 1450375_at 1.00820 persephin Pspn
 1450376_at 0.99366 Max interacting protein 1 Mxi1
 1450377_at 0.80039 similar to thrombospondin 1 LOC640441
 1450378_at 1.29035 TAP binding protein Tapbp
 1450379_at 1.05278 moesin Msn
 1450380_at 0.76944 ependymin related protein 2 (zebrafish) Epdr2
 1450381_a_at 1.06649 B-cell leukemia/lymphoma 6 Bcl6
 1450382_at 0.98118 neurofibromatosis 2 Nf2
 1450383_at 1.33770 low density lipoprotein receptor Ldlr
 1450384_at 1.09018 beta-site APP cleaving enzyme 1 Bace1
 1450385_at 1.05681 karyopherin (importin) alpha 3 Kpna3
 1450386_at 1.09884 karyopherin (importin) alpha 3 Kpna3
 1450387_s_at 1.89535 adenylate kinase 3 alpha-like 1 Ak311
 1450388_s_at 0.88594 twisted gastrulation homolog 1 (Drosophila) Twsg1

1450389_s_at 1.00755 "phosphatidylinositol-4-phosphate 5-kinase, type 1 alpha" Pip5k1a
1450390_x_at 1.08059 ribosomal protein S18 Rps18
1450391_a_at 1.25437 monoglyceride lipase Mgl1
1450392_at 0.73752 "ATP-binding cassette, sub-family A (ABC1), member 1" Abca1
1450393_a_at 0.95091 secretory carrier membrane protein 4 Scamp4
1450394_at 0.94221 golgi phosphoprotein 3 Golph3
1450395_at 0.87923 "solute carrier family 22 (organic cation transporter), member 5" Slc22a5
1450396_at 0.99688 stromal antigen 2 Stag2
1450397_at 1.17901 microtubule-associated protein 1 B Mtap1b
1450398_at 0.91034 "potassium channel, subfamily K, member 5" Kcnk5
1450399_at 1.08238 presenilin 1 Psen1
1450400_at 1.07826 nuclear receptor coactivator 6 interacting protein Ncoa6ip
1450401_at 1.00250 nuclear receptor coactivator 6 interacting protein Ncoa6ip
1450402_at 0.97998 peroxisome proliferator activated receptor binding protein Pparbp
1450403_at 1.43406 signal transducer and activator of transcription 2 Stat2
1450404_at 0.94555 "solute carrier family 23 (nucleobase transporters), member 1" Slc23a1
1450405_at 1.22840 mitochondrial ribosomal protein L19 Mrpl19
1450406_a_at 0.92275 "ST3 beta-galactoside alpha-2,3-sialyltransferase 3" St3gal3
1450407_a_at 1.06567 "acidic (leucine-rich) nuclear phosphoprotein 32 family, member A" Anp32a
1450408_at 0.99168 chloride channel 7 Clcn7
1450409_a_at 0.87703 RIKEN cDNA 4930570C03 gene 4930570C03Rik
1450410_a_at 0.92241 RIKEN cDNA 4930570C03 gene 4930570C03Rik
1450411_at 0.91885 RIKEN cDNA 2900009I07 gene 2900009I07Rik
1450412_at 1.17939 transducin (beta)-like 2 Tbl2
1450413_at 1.04835 "platelet derived growth factor, B polypeptide" Pdgfb
1450414_at 1.13650 "platelet derived growth factor, B polypeptide" Pdgfb
1450415_at 1.04497 "phosphodiesterase 6A, cGMP-specific, rod, alpha" Pde6a
1450416_at 0.87798 chromobox homolog 5 (Drosophila HP1a) Cbx5
1450417_a_at 0.98935 ribosomal protein S20 /// similar to 40S ribosomal protein S20 /// similar to 40S ribosomal protein S20 Rps20 /// LOC245676 /// LOC623568
1450418_a_at 0.84168 "Yip1 domain family, member 4" Yipf4
1450419_at 0.93491 serine hydrolase-like Serhl
1450420_at 1.05441 stromal antigen 1 Stag1
1450421_at 1.01433 transforming growth factor alpha Tgfa
1450422_a_at 0.92305 KDEL (Lys-Asp-Glu-Leu) containing 1 Kdelc1
1450423_s_at 1.12159 brix domain containing 1 Bxdc1
1450424_a_at 1.03249 interleukin 18 binding protein Il18bp
1450425_a_at 1.06291 RIKEN cDNA 2700062C07 gene 2700062C07Rik
1450426_at 0.95292 "cholinergic receptor, nicotinic, alpha polypeptide 6" Chrna6
1450427_at 0.88030 "cholinergic receptor, nicotinic, alpha polypeptide 6" Chrna6
1450428_at 0.80565 LIM homeobox protein 1 Lhx1
1450429_at 0.82625 calpain 6 Capn6
1450430_at 1.01807 "mannose receptor, C type 1" Mrc1

1450431_a_at 0.93871 "neural precursor cell expressed, developmentally down-regulated gene 4"
Nedd4

1450432_s_at 0.94147 MUS81 endonuclease homolog (yeast) Mus81

1450433_at 0.91246 recombining binding protein suppressor of hairless-like (Drosophila) Rbpsuhl

1450434_s_at 1.15893 "phosphate cytidylyltransferase 1, choline, alpha isoform" Pcyt1a

1450435_at 0.86516 L1 cell adhesion molecule L1cam

1450436_s_at 0.86599 "DnaJ (Hsp40) homolog, subfamily B, member 5" Dnajb5

1450437_a_at 0.94289 neural cell adhesion molecule 1 Ncam1

1450438_at 0.83677 neural cell adhesion molecule 1 Ncam1

1450439_at 0.94676 host cell factor C1 Hcfc1

1450440_at 0.99658 glial cell line derived neurotrophic factor family receptor alpha 1 Gfra1

1450441_at 1.02042 leucine rich repeat containing 50 Lrrc50

1450442_at 0.82798 adducin 2 (beta) Add2

1450443_at 1.09414 polypyrimidine tract binding protein 1 Ptbp1

1450444_a_at 0.97642 "nuclear receptor subfamily 1, group H, member 3" Nr1h3

1450445_at 0.96415 "phosphate regulating gene with homologies to endopeptidases on the X
chromosome (hypophosphatemia, vitamin D resistant rickets)" Phex

1450446_a_at 0.93978 suppressor of cytokine signaling 1 Socs1

1450447_at 0.92465 "hepatic nuclear factor 4, alpha" Hnf4a

1450448_at 0.90394 stanniocalcin 1 Stc1

1450449_a_at 0.79238 RIKEN cDNA 2900002H16 gene 2900002H16Rik

1450450_at 1.06831 Down syndrome critical region gene 1-like 2 Dscr112

1450451_at 0.94413 "sparc/osteonectin, cwcv and kazal-like domains proteoglycan 2" Spock2

1450452_a_at 0.92106 nuclear prelamin A recognition factor-like Narfl

1450453_a_at 0.98298 "phosphodiesterase 6G, cGMP-specific, rod, gamma" Pde6g

1450454_at 1.16380 "torsin family 3, member A" Tor3a

1450455_s_at 0.95563 "aldo-keto reductase family 1, member C12" Akr1c12

1450456_at 0.94718 interleukin 21 receptor Il21r

1450457_at 1.01125 Casitas B-lineage lymphoma Cbl

1450458_at 0.92612 nuclear receptor coactivator 2 Ncoa2

1450459_at 1.01112 RIKEN cDNA 2010106G01 gene 2010106G01Rik

1450460_at 1.13582 aquaporin 3 Aqp3

1450461_at 1.07165 "transcription factor 7, T-cell specific" Tcf7

1450462_at 1.06236 corticotropin releasing hormone receptor 2 Crhr2

1450463_at 0.99094 RIKEN cDNA 4833439L19 gene 4833439L19Rik

1450464_at 0.90865 E4F transcription factor 1 E4f1

1450465_at 0.95222 "cyclin-dependent kinase 5, regulatory subunit 2 (p39)" Cdk5r2

1450466_at 0.88327 RIKEN cDNA B230310J22 gene B230310J22Rik

1450467_at 1.02791 bassoon Bsn

1450468_at 0.99953 myocilin Myoc

1450469_at 0.94690 "testis specific protein, Ddc8" Ddc8

1450470_at 0.96070 --- ---

1450471_at 0.84520 MAD homolog 3 (Drosophila) Smad3

1450472_s_at 0.95353 MAD homolog 3 (Drosophila) Smad3
1450473_at 0.99373 protocadherin 12 Pcdh12
1450474_at 1.13781 "serine (or cysteine) peptidase inhibitor, clade B, member 9c" Serpinb9c
1450475_at 0.86973 distal-less homeobox 3 Dlx3
1450476_at 0.91391 cannabinoid receptor 2 (macrophage) Cnr2
1450477_at 1.03499 5-hydroxytryptamine (serotonin) receptor 2C Htr2c
1450478_a_at 1.16744 "protein tyrosine phosphatase, non-receptor type 12" Ptpn12
1450479_x_at 1.06739 "protein tyrosine phosphatase, non-receptor type 12" Ptpn12
1450480_a_at 0.85520 G protein-coupled receptor kinase 6 Gprk6
1450481_at 0.98813 --- ---
1450482_a_at 0.91309 paired-like homeodomain transcription factor 2 Pitx2
1450483_at 0.74183 gap junction membrane channel protein alpha 12 Gja12
1450484_a_at 1.01347 thymidylate kinase family LPS-inducible member Tyki
1450485_at 0.92550 SRY-box containing gene 3 Sox3
1450486_a_at 1.06536 opioid receptor-like 1 Oprl1
1450487_at 0.80650 visual system homeobox 1 homolog (zebrafish) Vsx1
1450488_at 0.92162 chemokine (C-C motif) ligand 24 Ccl24
1450489_at 0.92063 sal-like 1 (Drosophila) Sall1
1450490_at 0.97896 "potassium voltage-gated channel, shaker-related subfamily, member 7" Kcna7
1450491_at 0.89696 prostaglandin E receptor 1 (subtype EP1) Ptger1
1450492_at 0.95937 cyclic nucleotide gated channel beta 3 Cngb3
1450493_at 0.98713 KISS1 receptor Kiss1r
1450494_x_at 0.97742 CEA-related cell adhesion molecule 1 Ceacam1
1450495_a_at 1.07355 "killer cell lectin-like receptor subfamily K, member 1" Klrk1
1450496_a_at 1.04954 RIKEN cDNA 2810433K01 gene 2810433K01Rik
1450497_at 0.96624 adenomatosis polyposis coli 2 Apc2
1450498_at 1.02233 "5,10-methylenetetrahydrofolate reductase" Mthfr
1450499_at 1.06167 ataxin 1 Atnx1
1450500_at 1.02412 U2AF homology motif (UHM) kinase 1 /// similar to U2AF homology motif (UHM) kinase 1 Uhmk1 /// LOC677213
1450501_at 1.52944 integrin alpha 2 Itga2
1450502_at 0.87665 dihydropyrimidinase-like 2 Dpysl2
1450503_at 0.94428 "potassium inwardly-rectifying channel, subfamily J, member 2" Kcnj2
1450504_a_at 1.06269 1-acylglycerol-3-phosphate O-acyltransferase 3 Agpat3
1450505_a_at 1.10369 RIKEN cDNA 1810015C04 gene 1810015C04Rik
1450506_a_at 1.18463 interferon stimulated exonuclease gene 20-like 1 Isg20l1
1450507_at 0.97010 "POU domain, class 2, transcription factor 3" Pou2f3
1450508_at 0.85935 forkhead box N1 Foxn1
1450509_at 0.99964 carbohydrate sulfotransferase 11 Chst11
1450510_a_at 0.87803 "calcium channel, voltage-dependent, P/Q type, alpha 1A subunit" Cacna1a
1450511_at 0.99907 "muscle, skeletal, receptor tyrosine kinase" Musk
1450512_at 0.80793 netrin 4 Ntn4
1450513_at 1.08403 CD33 antigen Cd33

1450514_at 0.93139 zinc finger and SCAN domain containing 2 Zscan2
 1450515_at 0.95055 "potassium inwardly rectifying channel, subfamily J, member 11" Kcnj11
 1450516_a_at 0.94717 "RAB17, member RAS oncogene family" Rab17
 1450517_at 1.20031 T-cell acute lymphocytic leukemia 2 Tal2
 1450518_at 0.99667 "hepatocyte nuclear factor 4, gamma" Hnf4g
 1450519_a_at 0.91938 "protein kinase, cAMP dependent, catalytic, alpha" Prkaca
 1450520_at 0.98231 "calcium channel, voltage-dependent, gamma subunit 3" Cacng3
 1450521_a_at 1.03012 "T-cell receptor gamma chain /// region containing T-cell receptor gamma
 chain; T-cell receptor gamma, variable 4" Tcrp /// LOC670271
 1450522_a_at 1.01744 "H1 histone family, member 0" H1f0
 1450523_at 0.92568 contactin 2 Cntn2
 1450524_at 1.03676 claudin 9 Cldn9
 1450525_at 0.86893 GLI-Kruppel family member GLI3 Gli3
 1450526_at 0.92278 "T-cell leukemia, homeobox 1" Tlx1
 1450527_at 1.12745 somatostatin receptor 1 Sstr1
 1450528_at 0.98963 "UDP-Gal:betaGlcNAc beta 1,3-galactosyltransferase, polypeptide 5"
 B3galt5
 1450529_at 1.13791 "histocompatibility 2, M region locus 9" H2-M9
 1450530_at 1.06909 "UDP-Gal:betaGlcNAc beta 1,3-galactosyltransferase, polypeptide 1"
 B3galt1
 1450531_at 1.05143 "histocompatibility 2, blastocyst" H2-B1
 1450532_at 0.88847 RIKEN cDNA 2310002J15 gene 2310002J15Rik
 1450533_a_at 0.98197 pleiomorphic adenoma gene-like 1 Plagl1
 1450534_x_at 1.18734 MHC (A.CA/J(H-2K-f) class I antigen LOC56628
 1450535_at 0.84774 keratin associated protein 12-1 Krtap12-1
 1450536_s_at 0.95408 keratin associated protein 12-1 Krtap12-1
 1450537_at 1.04588 midline 2 Mid2
 1450538_s_at 0.86621 "mast cell protease 9 /// chymase 2, mast cell" Mcpt9 /// Cma2
 1450539_at 1.04681 keratin associated protein 5-1 Krtap5-1
 1450540_x_at 1.01237 keratin associated protein 5-1 Krtap5-1
 1450541_at 3.07357 plasmacytoma variant translocation 1 Pvt1
 1450542_s_at 0.94755 "melanoma antigen, family A, 1 /// melanoma antigen, family A, 2 ///
 melanoma antigen, family A, 3 /// melanoma antigen, family A, 5 /// melanoma antigen, family A, 6 ///
 melanoma antigen, family A, 8 /// similar to melanoma antigen family A, 8" Magea1 /// Magea2 ///
 Magea3 /// Magea5 /// Magea6 /// Magea8 /// LOC670799
 1450543_at 1.04278 myosin 1H Myo1h
 1450544_at 1.04906 "hypothetical protein, MNCb-4779" AB041550
 1450545_a_at 0.93845 "deoxynucleotidyltransferase, terminal" Dntt
 1450546_at 1.07099 "Alport syndrome, mental retardation, midface hypoplasia and elliptocytosis
 chromosomal region gene 1 homolog (human)" Ammcr1
 1450547_x_at 1.09445 deubiquitinating enzyme 2 Dub2
 1450548_at 1.00324 5-hydroxytryptamine (serotonin) receptor 1F Htr1f
 1450549_s_at 0.92703 "ELK4, member of ETS oncogene family" Elk4

1450550_at 1.11424 interleukin 5 Il5
1450551_x_at 1.00087 "killer cell lectin-like receptor, subfamily A, member 15" Klra15
1450552_at 0.97415 "ATPase, H+ transporting, lysosomal V1 subunit E2" Atp6v1e2
1450553_at 0.93574 homeo box D12 Hoxd12
1450554_at 1.01080 defensin beta 2 Defb2
1450555_at 0.99942 testis expressed gene 13 Tex13
1450556_at 0.78467 spectrin beta 1 Spnb1
1450557_at 0.91309 "sodium channel, voltage-gated, type IV, alpha" Scn4a
1450558_at 1.08916 "vomeronasal 1 receptor, A1" V1ra1
1450559_at 0.95435 RIKEN cDNA 5730559C18 gene 5730559C18Rik
1450560_a_at 0.97441 "protein phosphatase 2, regulatory subunit B (B56), delta isoform" Ppp2r5d
1450561_a_at 0.99543 surfeit gene 1 Surf1
1450562_at 1.17880 "lymphocyte antigen 6 complex, locus F" Ly6f
1450563_at 0.96465 microsomal glutathione S-transferase 3 Mgst3
1450564_x_at 0.93213 interferon alpha 1 Ifna1
1450565_at 0.95713 interleukin 9 Il9
1450566_at 0.97190 interleukin 3 Il3
1450567_a_at 0.88601 "procollagen, type II, alpha 1" Col2a1
1450568_at 1.09210 galanin receptor 1 Galr1
1450569_a_at 1.04159 RNA binding motif protein 14 Rbm14
1450570_a_at 0.97567 CD19 antigen Cd19
1450571_a_at 1.01418 beaded filament structural protein in lens-CP94 Bfsp1
1450572_at 0.97906 "alkaline phosphatase 3, intestine, not Mn requiring" Akp3
1450573_at 1.03634 anti-Mullerian hormone Amh
1450574_at 1.05032 "cytochrome P450, family 11, subfamily b, polypeptide 2" Cyp11b2
1450575_at 1.03091 "cholinergic receptor, muscarinic 4" Chrm4
1450576_a_at 0.99447 "splicing factor 3a, subunit 2" Sf3a2
1450577_at 0.95706 somatostatin receptor 3 Sstr3
1450578_at 1.13135 sex determining region of Chr Y Sry
1450579_x_at 1.03370 sex determining region of Chr Y Sry
1450580_at 0.89058 G protein-coupled receptor 45 Gpr45
1450581_at 0.99376 galanin receptor 3 Galr3
1450582_at 1.18207 "histocompatibility 2, Q region locus 5" H2-Q5
1450583_s_at 0.87450 formyl peptide receptor-like 1 Fprl1
1450584_at 0.95815 homeo box D11 Hoxd11
1450585_at 1.05998 "taste receptor, type 2, member 119" Tas2r119
1450586_at 0.92317 "bradykinin receptor, beta 1" Bdkrb1
1450587_at 0.88685 "histocompatibility 2, M region locus 10.1" H2-M10.1
1450588_at 1.06358 olfactory receptor 67 Olfr67
1450589_at 0.92620 olfactory receptor 749 Olfr749
1450590_at 1.05992 olfactory receptor 159 Olfr159
1450591_at 1.00810 olfactory receptor 154 Olfr154
1450592_at 1.11156 olfactory receptor 69 Olfr69

1450593_at 0.87442 interferon alpha 6 Ifna6
1450594_at 0.89643 "melanoma antigen, family A, 4" Magea4
1450595_at 0.92764 "vomeronasal 1 receptor, C1" V1rc1
1450596_at 0.93896 olfactory receptor 66 Olfr66
1450597_at 0.97233 olfactory receptor 870 Olfr870
1450598_at 0.91966 "vomeronasal 1 receptor, A4" V1ra4
1450599_at 1.05630 "vomeronasal 1 receptor, B3" V1rb3
1450600_at 1.05147 olfactory receptor 1508 Olfr1508
1450601_at 1.07732 "vomeronasal 1 receptor, B8" V1rb8
1450602_at 1.06539 "vomeronasal 1 receptor, C7" V1rc7
1450603_s_at 0.96299 "vomeronasal 1 receptor, A3 /// vomeronasal 1 receptor, A4" V1ra3 /// V1ra4
1450604_at 0.93375 olfactory receptor 140 Olfr140
1450605_at 0.93548 "taste receptor, type 2, member 108" Tas2r108
1450606_at 0.85252 phenylethanolamine-N-methyltransferase Pnmt
1450607_s_at 0.98910 urocortin 2 Ucn2
1450608_at 1.00338 "histone 1, H1t" Hist1h1t
1450609_at 1.03542 lipocalin 8 Lcn8
1450610_at 0.94957 urocortin Ucn
1450611_at 0.96457 orosomucoid 3 Orm3
1450612_a_at 0.88073 phosphatidylethanolamine N-methyltransferase Pemt
1450613_x_at 0.90624 interferon alpha B Ifnab
1450614_x_at 1.11187 interferon alpha 5 Ifna5
1450615_at 0.95404 "eosinophil-associated, ribonuclease A family, member 4" Ear4
1450616_at 0.87754 "eosinophil-associated, ribonuclease A family, member 5" Ear5
1450617_at 1.04869 --- ---
1450618_a_at 1.09784 small proline-rich protein 2A Sprr2a
1450619_x_at 0.93776 cell matrix adhesion regulator Cmar
1450620_at 1.08314 --- ---
1450621_a_at 0.98243 "hemoglobin Y, beta-like embryonic chain" Hbb-y
1450622_at 1.05429 breast cancer anti-estrogen resistance 1 Bcar1
1450623_at 0.96791 "guanine nucleotide binding protein, beta 2" Gnb2
1450624_at 1.09663 betaine-homocysteine methyltransferase Bhmt
1450625_at 1.02547 "procollagen, type V, alpha 2" Col5a2
1450626_at 0.90707 "mannosidase, beta A, lysosomal" Manba
1450627_at 0.71929 progressive ankylosis Ank
1450628_at 1.06160 "solute carrier family 2, (facilitated glucose transporter), member 8" Slc2a8
1450629_at 0.90616 LIM domain and actin binding 1 Lima1
1450630_at 1.11461 queuine tRNA-ribosyltransferase 1 Qtrt1
1450631_x_at 0.98292 "defensin related cryptdin 12 /// defensin related cryptdin 2 /// defensin related
cryptdin 3 /// defensin related cryptdin 6 /// defensin related cryptdin 9 /// defensin related cryptdin 17 ///
defensin related cryptdin 23 /// defensin related cryptdin 24 /// defensin, alpha, pseudogene 1"
Defcr12 /// Defcr2 /// Defcr3 /// Defcr6 /// Defcr9 /// Defcr17 /// Defcr23 /// Defcr24 /// Defa-ps1
1450632_at 1.01822 similar to aplysia ras-related homolog A2 /// similar to aplysia ras-related

homolog A2 LOC433749 /// LOC631743
 1450633_at 0.97345 calmodulin 4 Calm4
 1450634_at 1.05348 "ATPase, H⁺ transporting, lysosomal V1 subunit A" Atp6v1a
 1450635_at 0.78999 alkaline phosphatase 5 Akp5
 1450636_s_at 0.95531 alkaline phosphatase 5 Akp5
 1450637_a_at 0.95261 AE binding protein 1 Aebp1
 1450638_at 1.07005 programmed cell death 5 Pcd5
 1450639_at 0.99574 "solute carrier family 28 (sodium-coupled nucleoside transporter), member 2" ///
 similar to solute carrier family 28 (sodium-coupled nucleoside transporter), member 2" Slc28a2 ///
 LOC381417
 1450640_x_at 0.83515 "ATP synthase, H⁺ transporting, mitochondrial F1F0 complex, subunit e" Atp5k
 1450641_at 0.93679 vimentin Vim
 1450642_at 0.96224 RIKEN cDNA 3110001I20 gene 3110001I20Rik
 1450643_s_at 0.99360 acyl-CoA synthetase long-chain family member 1 Acs11
 1450644_at 0.86189 "zinc finger protein 36, C3H type-like 1" Zfp3611
 1450645_at 0.84199 metallothionein 4 Mt4
 1450646_at 1.51140 "cytochrome P450, family 51" Cyp51
 1450647_at 1.14091 Hermansky-Pudlak syndrome 3 homolog (human) Hps3
 1450648_s_at 0.99989 "histocompatibility 2, class II antigen A, beta 1" H2-Ab1
 1450649_at 0.86758 "guanine nucleotide binding protein (G protein), gamma 10" Gng10
 1450650_at 1.08325 myosin X Myo10
 1450651_at 0.91865 myosin X Myo10
 1450652_at 0.89583 cathepsin K Ctsk
 1450653_at 1.03766 spermatogenic Zip 1 Spz1
 1450654_a_at 0.93611 dehydrolipoyl dihydrolipoyl synthase Dhdds
 1450655_at 0.90516 phosphatase and tensin homolog Pten
 1450656_at 1.01386 "guanine nucleotide binding protein, alpha 13" Gna13
 1450657_at 1.05661 peptidylprolyl isomerase E (cyclophilin E) Ppie
 1450658_at 0.90677 "a disintegrin-like and metalloproteinase (reprolysin type) with thrombospondin
 type 1 motif, 5 (aggrecanase-2)" Adamts5
 1450659_at 0.83616 regulator of G protein signaling 7 Rgs7
 1450660_at 0.95089 6-pyruvoyl-tetrahydropterin synthase Pts
 1450661_x_at 0.97004 nuclear factor I/C Nfic
 1450662_at 0.86843 testis specific protein kinase 1 Tesk1
 1450663_at 0.91644 thrombospondin 2 Thbs2
 1450664_at 0.96014 "GA repeat binding protein, alpha" Gabpa
 1450665_at 0.87699 "GA repeat binding protein, alpha" Gabpa
 1450666_s_at 0.89223 ataxin 10 Atxn10
 1450667_a_at 1.03402 citrate synthase Cs
 1450668_s_at 1.10476 heat shock protein 1 (chaperonin 10) Hspe1
 1450669_at 0.85088 mitogen activated protein kinase kinase kinase 11 Map3k11
 1450670_at 0.95234 dopamine beta hydroxylase Dbh

1450671_at 0.85043 "crystallin, beta A1" Cryba1
1450672_a_at 0.94298 three prime repair exonuclease 1 Trex1
1450673_at 0.95418 "procollagen, type IX, alpha 2" Col9a2
1450674_at 0.93475 cyclin-dependent kinase 5 Cdk5
1450675_at 1.01271 stromal membrane-associated protein 1-like Smap11
1450676_at 0.95374 "transcription elongation factor B (SIII), polypeptide 3" Tceb3
1450677_at 0.91479 checkpoint kinase 1 homolog (S. pombe) Chek1
1450678_at 1.35948 integrin beta 2 Itgb2
1450679_at 1.02053 RIKEN cDNA 5730470L24 gene 5730470L24Rik
1450680_at 1.01688 recombination activating gene 1 Rag1
1450681_at 1.02337 zinc finger protein 143 Zfp143
1450682_at 0.93783 "fatty acid binding protein 6, ileal (gastrotropin)" Fabp6
1450683_at 1.08394 transgelin 3 Tagln3
1450684_at 0.98473 ets variant gene 1 Etv1
1450685_at 0.91350 cAMP-regulated phosphoprotein 19 Arpp19
1450686_at 1.01845 paraoxonase 2 Pon2
1450687_at 0.97135 insulin-like growth factor 2 mRNA binding protein 3 Igf2bp3
1450688_at 0.97031 ral guanine nucleotide dissociation stimulator-like 2 Rgl2
1450689_at 0.94876 "protease, serine, 3 /// protease, serine, 1 (trypsin 1) /// trypsinogen 10 ///
trypsinogen 12" Prss3 /// Prss1 /// LOC436522 /// LOC436523
1450690_at 1.00971 --- ---
1450691_at 0.90515 cask-interacting protein 2 Caskin2
1450692_at 1.05833 kinesin family member 4 Kif4
1450693_at 1.00428 --- ---
1450694_at 1.04338 FK506 binding protein 2 Fkbp2
1450695_at 0.81916 aryl-hydrocarbon receptor Ahr
1450696_at 1.42983 "proteasome (prosome, macropain) subunit, beta type 9 (large multifunctional
peptidase 2)" Psmb9
1450697_at 1.03745 "solute carrier family 30 (zinc transporter), member 7" Slc30a7
1450698_at 0.92904 dual specificity phosphatase 2 Dusp2
1450699_at 0.72724 selenium binding protein 1 Selenbp1
1450700_at 0.71179 CDC42 effector protein (Rho GTPase binding) 3 Cdc42ep3
1450701_a_at 1.06992 "general transcription factor II H, polypeptide 2" Gtf2h2
1450702_at 0.90205 hemochromatosis Hfe
1450703_at 0.85518 "solute carrier family 7 (cationic amino acid transporter, y+ system), member
2" Slc7a2
1450704_at 0.90621 Indian hedgehog Ihh
1450705_at 1.07209 RD RNA-binding protein Rdbp
1450706_a_at 1.00769 ADP-ribosylation factor-like 3 Arl3
1450707_at 0.92633 "double C2, gamma" Doc2g
1450708_at 1.05265 secretogranin II Scg2
1450709_at 0.86032 defensin related cryptdin 5 Defcr5
1450710_at 0.95063 "jumonji, AT rich interactive domain 2" Jarid2

1450711_at 1.07462 bromodomain containing 4 Brd4
1450712_at 1.13444 "potassium inwardly-rectifying channel, subfamily J, member 9" Kcnj9
1450713_at 1.05160 chondroitin sulfate proteoglycan 5 Cspg5
1450714_at 0.93448 antizyme inhibitor 1 Azin1
1450715_at 0.87012 "cytochrome P450, family 1, subfamily a, polypeptide 2" Cyp1a2
1450716_at 0.60719 "a disintegrin-like and metallopeptidase (reprolysin type) with thrombospondin type 1 motif, 1" Adamts1
1450717_at 0.92740 "angiogenin, ribonuclease A family, member 1" Ang1
1450718_at 1.01450 SH2B adaptor protein 2 Sh2b2
1450719_at 1.31637 meprin 1 alpha Mep1a
1450720_at 0.97487 "acid phosphatase 1, soluble" Acp1
1450721_at 1.05941 "acid phosphatase 1, soluble" Acp1
1450722_at 1.15561 nucleoporin 50 Nup50
1450723_at 0.98775 "ISL1 transcription factor, LIM/homeodomain" Isl1
1450724_at 1.03254 "down-regulated by Ctnnb1, a" Drctnnb1a
1450725_s_at 0.95285 carbonic anhydrase 14 Car14
1450726_at 0.87653 N-acylsphingosine amidohydrolase 2 Asah2
1450727_a_at 0.94301 "polymerase (DNA-directed), delta interacting protein 2" Poldip2
1450728_at 0.90742 four jointed box 1 (Drosophila) Fjx1
1450729_at 1.00265 heparan sulfate 2-O-sulfotransferase 1 Hs2st1
1450730_at 0.97191 heparan sulfate 2-O-sulfotransferase 1 Hs2st1
1450731_s_at 1.08791 "tumor necrosis factor receptor superfamily, member 21" Tnfrsf21
1450732_a_at 1.11887 bicaudal D homolog 2 (Drosophila) Bicd2
1450733_at 1.03425 bicaudal D homolog 2 (Drosophila) Bicd2
1450734_at 1.02808 leucine zipper transcription regulator 2 Lztr2
1450735_at 1.12124 RIKEN cDNA 1810003N24 gene 1810003N24Rik
1450736_a_at 0.88600 "hemoglobin Z, beta-like embryonic chain" Hbb-bh1
1450737_at 1.00268 polymerase (RNA) III (DNA directed) polypeptide K Polr3k
1450738_at 1.17100 kinesin family member 21A Kif21a
1450739_at 0.88195 transducin (beta)-like 1X-linked receptor 1 Tbl1xr1
1450740_a_at 0.96541 "microtubule-associated protein, RP/EB family, member 1" Mapre1
1450741_at 0.91955 staufer (RNA binding protein) homolog 1 (Drosophila) Stau1
1450742_at 0.95925 bystin-like Bysl
1450743_s_at 0.97417 "synaptotagmin binding, cytoplasmic RNA interacting protein" Syncrip
1450744_at 0.74000 elongation factor RNA polymerase II 2 Ell2
1450745_at 0.88061 "core 1 UDP-galactose:N-acetylgalactosamine-alpha-R beta 1,3-galactosyltransferase" C1galt1
1450746_at 1.16098 kelch-like ECH-associated protein 1 Keap1
1450747_at 1.10640 kelch-like ECH-associated protein 1 Keap1
1450748_at 1.06131 "sphingomyelin phosphodiesterase 3, neutral" Smpd3
1450749_a_at 1.02754 "nuclear receptor subfamily 4, group A, member 2" Nr4a2
1450750_a_at 0.92184 "nuclear receptor subfamily 4, group A, member 2" Nr4a2
1450751_at 0.92607 zona pellucida binding protein Zpbp

1450752_at 1.11489 "cytochrome c, testis" Cyct
1450753_at 1.01951 natural killer cell group 7 sequence Nkg7
1450754_at 0.95475 "similar to calcium channel, voltage-dependent, alpha 2/delta subunit 2"
LOC669637
1450755_at 1.03144 "platelet-activating factor acetylhydrolase, isoform 1b, alpha2 subunit"
Pafah1b2
1450756_s_at 0.86636 cullin 3 Cul3
1450757_at 1.21918 cadherin 11 Cdh11
1450758_at 1.16491 contactin associated protein-like 2 Cntnap2
1450759_at 0.97990 bone morphogenetic protein 6 Bmp6
1450760_a_at 0.90870 "inhibitor of growth family, member 3" Ing3
1450761_s_at 1.08612 regulating synaptic membrane exocytosis 2 Rims2
1450762_s_at 0.96490 zinc finger protein 191 Zfp191
1450763_x_at 0.97978 wingless-related MMTV integration site 3 Wnt3
1450764_at 0.90321 acyloxyacyl hydrolase Aoah
1450765_a_at 0.97302 "phosphodiesterase 6H, cGMP-specific, cone, gamma" Pde6h
1450766_at 1.12594 "phosphodiesterase 6H, cGMP-specific, cone, gamma" Pde6h
1450767_at 0.82343 "neural precursor cell expressed, developmentally down-regulated gene 9"
Nedd9
1450768_at 1.08060 "discs, large homolog 1 (Drosophila)" Dlgh1
1450769_s_at 0.81145 StAR-related lipid transfer (START) domain containing 5 Stard5
1450770_at 1.15194 RIKEN cDNA 3632451O06 gene 3632451O06Rik
1450771_at 0.97147 fucosyltransferase 9 Fut9
1450772_at 1.04469 wingless-related MMTV integration site 11 Wnt11
1450773_at 0.98262 "potassium voltage-gated channel, Shal-related family, member 2" Kcnd2
1450774_at 0.92926 "lymphocyte antigen 6 complex, locus G6D" Ly6g6d
1450775_at 0.92681 Moloney sarcoma oncogene Mos
1450776_at 0.95410 "1-acylglycerol-3-phosphate O-acyltransferase 6 (lysophosphatidic acid
acyltransferase, zeta)" Agpat6
1450777_at 1.12375 5'-3' exoribonuclease 2 Xrn2
1450778_a_at 1.07808 "RNA U, small nuclear RNA export adaptor" Rnuxa
1450779_at 0.98210 "fatty acid binding protein 7, brain" Fabp7
1450780_s_at 1.10508 high mobility group AT-hook 2 Hmga2
1450781_at 1.17248 high mobility group AT-hook 2 Hmga2
1450782_at 0.86659 wingless-related MMTV integration site 4 Wnt4
1450783_at 1.46021 --- ---
1450784_at 0.76490 reversion-inducing-cysteine-rich protein with kazal motifs Reck
1450785_at 1.20789 thyroid hormone receptor interactor 4 Trip4
1450786_x_at 0.90605 similar to PDZ and LIM domain protein 5 (Enigma homolog) (Enigma-like
PDZ and LIM domains protein) LOC669660
1450787_at 1.00530 chloride channel 5 Clcn5
1450788_at 1.01012 serum amyloid A 1 Saa1
1450789_at 0.99883 "rhopilin, Rho GTPase binding protein 1" Rhpn1

1450790_at 1.08643 thyroglobulin Tgn
1450791_at 0.99389 natriuretic peptide precursor type B Nppb
1450792_at 0.92075 TYRO protein tyrosine kinase binding protein Tyrobp
1450793_at 1.25923 RIKEN cDNA 4930550L24 gene 4930550L24Rik
1450794_at 0.86761 arginine vasopressin Avp
1450795_at 1.03113 luteinizing hormone beta Lhb
1450796_at 0.93427 atonal homolog 7 (Drosophila) Atoh7
1450797_a_at 0.92261 chromobox homolog 1 (Drosophila HP1 beta) Cbx1
1450798_at 0.77284 tenascin XB Tnxb
1450799_at 1.16944 adenylate cyclase activating polypeptide 1 receptor 1 Adcyap1r1
1450800_at 1.00364 synaptotagmin VIII Syt8
1450801_at 0.85372 a disintegrin and metallopeptidase domain 21 Adam21
1450802_at 1.00254 "protease, serine, 28" Prss28
1450803_at 0.95418 neurotrophin 3 Ntf3
1450804_at 0.82190 kinesin family member 5C Kif5c
1450805_at 1.11041 "sarcoglycan, delta (dystrophin-associated glycoprotein)" Sgcd
1450806_at 0.98039 "T-cell leukemia, homeobox 3" Tlx3
1450807_at 0.92549 --- ---
1450808_at 1.07238 formyl peptide receptor 1 Fpr1
1450809_at 0.99604 seminal vesicle antigen-like 2 Sval2
1450810_at 1.08270 follicle stimulating hormone receptor Fshr
1450811_at 0.87034 small proline-rich protein 2J Sprr2j
1450812_at 0.98849 B-cell leukemia/lymphoma 2 related protein A1a /// B-cell leukemia/lymphoma
2 related protein A1b /// B-cell leukemia/lymphoma 2 related protein A1c /// B-cell leukemia/lymphoma
2 related protein A1d Bcl2a1a /// Bcl2a1b /// Bcl2a1c /// Bcl2a1d
1450813_a_at 0.94556 "troponin I, skeletal, slow 1" Tnni1
1450814_a_at 1.07002 importin 4 Ipo4
1450815_s_at 0.98357 coiled-coil-helix-coiled-coil-helix domain containing 2 /// RIKEN cDNA
2410018M08 gene Chchd2 /// 2410018M08Rik
1450816_at 1.00551 "polymerase (DNA directed), gamma 2, accessory subunit" Polg2
1450817_at 1.03175 small optic lobes homolog (Drosophila) Solh
1450818_a_at 0.99551 "NADH dehydrogenase (ubiquinone) 1 alpha subcomplex, 7 (B14.5a)"
Ndufa7
1450819_at 0.82233 poliovirus receptor-related 1 Pvr11
1450820_a_at 0.91727 netrin G2 Ntng2
1450821_at 0.95869 p300/CBP-associated factor Pcaf
1450822_at 0.95664 P lysozyme structural Lzp-s
1450823_at 1.08604 OG9 homeobox gene Og9x
1450824_at 0.93524 patched homolog 1 Ptch1
1450825_at 1.04792 N-acylsphingosine amidohydrolase (alkaline ceramidase) 3 Asah3
1450826_a_at 1.67497 serum amyloid A 3 Saa3
1450827_at 0.97853 kinesin family member 23 Kif23
1450828_at 1.13049 synaptopodin 2 Synpo2

1450829_at 1.31974 "tumor necrosis factor, alpha-induced protein 3" Tnfaip3
1450830_a_at 0.95323 "phosphodiesterase 6C, cGMP specific, cone, alpha prime" Pde6c
1450831_at 1.03777 "POU domain, class 3, transcription factor 2" Pou3f2
1450832_at 0.90903 homeo box C5 Hoxc5
1450833_at 0.89958 "cholinergic receptor, muscarinic 1, CNS" Chrm1
1450834_at 0.94483 fucosyltransferase 4 Fut4
1450835_a_at 0.96576 glial cell line derived neurotrophic factor family receptor alpha 4 Gfra4
1450836_at 1.01436 neurogenin 1 Neurog1
1450837_at 1.14841 proline rich protein HaeIII subfamily 1 Prh1
1450838_x_at 1.17186 ribosomal protein L37 /// similar to ribosomal protein L37 Rpl37 ///
LOC383438
1450839_at 0.47617 "DNA segment, human D4S114" D0H4S114
1450840_a_at 0.95192 ribosomal protein L39 Rpl39
1450841_at 1.12358 "STT3, subunit of the oligosaccharyltransferase complex, homolog A (S. cerevisiae)" Stt3a
1450842_a_at 1.08540 centromere protein A Cenpa
1450843_a_at 0.87286 "serine (or cysteine) peptidase inhibitor, clade H, member 1" Serpinh1
1450844_at 0.99837 syntaxin 6 Stx6
1450845_a_at 1.11918 basic leucine zipper and W2 domains 1 Bzw1
1450846_at 1.09719 basic leucine zipper and W2 domains 1 Bzw1
1450847_at 0.93060 nuclear cap binding protein subunit 2 Ncbp2
1450848_at 0.98377 death associated protein 3 Dap3
1450849_at 1.09457 heterogeneous nuclear ribonucleoprotein U Hnrpu
1450850_at 1.00020 villin 2 Vil2
1450851_at 1.00101 WD repeat domain 1 Wdr1
1450852_s_at 0.90059 coagulation factor II (thrombin) receptor F2r
1450853_at 0.87282 "transducin-like enhancer of split 4, homolog of Drosophila E(spl)" Tle4
1450854_at 1.03740 proliferation-associated 2G4 Pa2g4
1450855_at 0.75464 armadillo repeat gene deleted in velo-cardio-facial syndrome Arvcf
1450856_at 0.80210 armadillo repeat gene deleted in velo-cardio-facial syndrome Arvcf
1450857_a_at 0.89369 "procollagen, type I, alpha 2" Colla2
1450858_a_at 1.00397 "ubiquitin-conjugating enzyme E2D 3 (UBC4/5 homolog, yeast)" Ube2d3
1450859_s_at 0.88200 "ubiquitin-conjugating enzyme E2D 3 (UBC4/5 homolog, yeast)" Ube2d3
1450860_at 1.00779 leucine aminopeptidase 3 Lap3
1450861_at 0.94018 "Fanconi anemia, complementation group C" FancC
1450862_at 0.92043 RAD54 like (S. cerevisiae) Rad54l
1450863_a_at 0.76151 doublecortin and calcium/calmodulin-dependent protein kinase-like 1
Dcamk1l
1450864_at 0.89601 calmodulin 3 Calm3
1450865_s_at 1.00709 mitochondrial ribosomal protein S24 Mrps24
1450866_a_at 1.08358 mitochondrial ribosomal protein L17 Mrpl17
1450867_at 1.04663 mitochondrial ribosomal protein L17 Mrpl17
1450868_at 0.94905 "DNA segment, Chr 8, ERATO Doi 354, expressed" D8ErtD354e

1450869_at 0.52574 fibroblast growth factor 1 Fgf1
1450870_at 0.90444 v-ral simian leukemia viral oncogene homolog A (ras related) Rala
1450871_a_at 1.01237 "branched chain aminotransferase 1, cytosolic" Bcat1
1450872_s_at 0.92954 lysosomal acid lipase 1 Lip1
1450873_at 1.13019 GTP binding protein 4 Gtpbp4
1450874_at 1.05085 matrin 3 Matr3
1450875_at 0.91788 G protein-coupled receptor 37 Gpr37
1450876_at 0.99294 complement component factor h Cfh
1450877_at 0.96882 cDNA sequence BC005537 BC005537
1450878_at 0.97068 sorcin Sri
1450879_at 0.94211 "ATPas, class II, type 9B" Atp9b
1450880_at 1.07810 mitochondrial ribosomal protein L16 Mrpl16
1450881_s_at 1.10005 G protein-coupled receptor 137B Gpr137b
1450882_s_at 1.09476 G protein-coupled receptor 137B /// similar to transmembrane 7 superfamily member 1 /// similar to transmembrane 7 superfamily member 1 Gpr137b /// LOC664862 /// LOC673335
1450883_a_at 0.98209 CD36 antigen Cd36
1450884_at 0.96557 CD36 antigen Cd36
1450885_at 1.18782 "DNA fragmentation factor, alpha subunit" Dffa
1450886_at 0.90618 germ cell-specific gene 2 Gsg2
1450887_at 1.08196 rcd1 (required for cell differentiation) homolog 1 (S. pombe) Rqcd1
1450888_at 1.09174 N-ethylmaleimide sensitive fusion protein attachment protein beta Napb
1450889_at 0.93583 "SWI/SNF related, matrix associated, actin dependent regulator of chromatin, subfamily a, member 3" Smarca3
1450890_a_at 1.00484 abl-interactor 1 Abi1
1450891_at 1.01130 signal recognition particle 19 Srp19
1450892_a_at 1.11169 ubiquitin specific peptidase 4 (proto-oncogene) Usp4
1450893_a_at 0.95412 ubiquitin-associated protein 1 Ubap1
1450894_a_at 0.98666 "adaptor protein complex AP-2, mu1" Ap2m1
1450895_a_at 1.01776 RIKEN cDNA 1810020G14 gene 1810020G14Rik
1450896_at 0.89479 Rho GTPase activating protein 5 Arhgap5
1450897_at 0.92515 Rho GTPase activating protein 5 Arhgap5
1450898_at 1.01800 hippocampus abundant gene transcript 1 Hiat1
1450899_at 0.88978 "neural precursor cell expressed, developmentally down-regulated gene 1" Nedd1
1450900_at 1.15016 expressed sequence AW011752 AW011752
1450901_a_at 1.18735 expressed sequence AW011752 AW011752
1450902_at 1.05471 bromodomain containing 3 Brd3
1450903_at 1.02924 RAD23b homolog (S. cerevisiae) Rad23b
1450904_at 0.97381 transmembrane protein 167 Tmem167
1450905_at 1.06649 plexin C1 Plxnc1
1450906_at 1.01902 plexin C1 Plxnc1
1450907_at 0.99950 signal peptidase complex subunit 2 homolog (S. cerevisiae) Spcs2

1450908_at 1.09698 eukaryotic translation initiation factor 4E /// hypothetical LOC630527 Eif4e ///
LOC630527
1450909_at 0.94203 eukaryotic translation initiation factor 4E Eif4e
1450910_at 0.87767 "CAP, adenylate cyclase-associated protein, 2 (yeast)" Cap2
1450911_at 0.97082 peptidylprolyl isomerase B Ppib
1450912_at 1.04132 "membrane-spanning 4-domains, subfamily A, member 1" Ms4a1
1450913_at 1.01120 "UDP-Gal:betaGlcNAc beta 1,4-galactosyltransferase, polypeptide 6 /// similar
to Beta-1,4-galactosyltransferase 6 (Beta-1,4-GalTase 6) (Beta4Gal-T6) (b4Gal-T6) (UDP-galactose:
beta-N-acetylglucosamine beta-1,4-galactosyltransferase 6) (UDP-Gal:beta-GlcNAc beta-1,4-
galactosyltransferase 6)" B4galt6 /// LOC675709
1450914_at 0.97786 "protein phosphatase 1, regulatory (inhibitor) subunit 14B" Ppp1r14b
1450915_at 0.75666 "adaptor-related protein complex 3, beta 1 subunit" Ap3b1
1450916_at 1.05746 staufer (RNA binding protein) homolog 2 (Drosophila) Stau2
1450917_at 1.80086 myomesin 2 Myom2
1450918_s_at 0.93484 Rous sarcoma oncogene Src
1450919_at 0.88982 "membrane protein, palmitoylated" Mpp1
1450920_at 1.05328 cyclin B2 Ccnb2
1450921_at 0.92062 aprataxin Aptx
1450922_a_at 0.84453 "transforming growth factor, beta 2" Tgfb2
1450923_at 0.85767 "transforming growth factor, beta 2" Tgfb2
1450924_at 1.07189 "hepatoma-derived growth factor, related protein 3" Hdgfrp3
1450925_a_at 1.01343 ribosomal protein S27-like /// similar to 40S ribosomal protein S27-like
protein Rps271 /// LOC667571
1450926_at 1.06284 "ATPase, H+ transporting, lysosomal V1 subunit G1 /// similar to vacuolar H+
ATPase G1 /// similar to vacuolar H+ ATPase G1" Atp6v1g1 /// LOC623391 /// LOC671155
1450927_at 0.81522 "leucine-zipper-like transcriptional regulator, 1" Lztr1
1450928_at 0.61529 inhibitor of DNA binding 4 Id4
1450929_at 0.86877 zinc finger protein 57 Zfp57
1450930_at 0.95999 hippocalcin Hpca
1450931_at 0.84380 dedicator of cytokinesis 9 /// similar to Dedicator of cytokinesis protein 9
(Cdc42 guanine nucleotide exchange factor zizimin 1) Dock9 /// LOC670309
1450932_s_at 1.07581 dedicator of cytokinesis 9 /// similar to Dedicator of cytokinesis protein 9
(Cdc42 guanine nucleotide exchange factor zizimin 1) Dock9 /// LOC670309
1450933_at 1.11772 phosphodiesterase 7A Pde7a
1450934_at 0.86575 eukaryotic translation initiation factor 4A2 Eif4a2
1450935_at 1.01529 "excision repair cross-complementing rodent repair deficiency,
complementation group 5" Ercc5
1450936_a_at 0.99502 deoxyribonuclease 1-like 2 Dnase1l2
1450937_at 0.88536 lin-7 homolog C (C. elegans) Lin7c
1450938_at 0.87438 --- ---
1450939_at 1.00300 ectonucleoside triphosphate diphosphohydrolase 1 Entpd1
1450940_at 0.96891 ganglioside-induced differentiation-associated-protein 1 Gdap1
1450941_at 0.98517 --- ---

1450942_at	0.82854	coiled-coil domain containing 16	Ccdc16
1450943_at	1.00600	RIKEN cDNA 2010012C16 gene	2010012C16Rik
1450944_at	0.75681	chondroitin sulfate proteoglycan 4	Cspg4
1450945_at	0.94068	"protein kinase C, alpha"	Prkca
1450946_at	0.98967	neural retina leucine zipper gene	Nrl
1450947_at	0.96659	RIKEN cDNA 2610528J11 gene	2610528J11Rik
1450948_a_at	1.05262	mitochondrial ribosomal protein L1	Mrpl1
1450949_at	1.06461	katanin p60 (ATPase-containing) subunit A1	Katna1
1450950_at	1.08798	structural maintenace of chromosomes 3	Smc3
1450951_at	1.12435	structural maintenace of chromosomes 3	Smc3
1450952_at	1.10830	phospholamban	Pln
1450953_at	1.00999	WD repeat domain 39	Wdr39
1450954_at	0.80582	YME1-like 1 (<i>S. cerevisiae</i>)	Yme1l1
1450955_s_at	0.82102	sortilin 1	Sort1
1450956_at	0.98005	stearoyl-coenzyme A desaturase 3	Scd3
1450957_a_at	1.03086	sequestosome 1	Sqstm1
1450958_at	0.83787	transmembrane 4 superfamily member 1	Tm4sf1
1450959_at	1.04867	RIKEN cDNA D930014E17 gene	D930014E17Rik
1450960_at	1.02011	insulin-like growth factor binding protein-like 1	Igfbp1
1450961_a_at	1.06814	transcription elongation factor A (SII)-like 6 /// transcription elongation factor A (SII)-like 5 /// transcription elongation factor A (SII)-like 3	Tceal6 /// Tceal5 /// Tceal3
1450962_at	1.02768	pyruvate dehydrogenase E1 alpha 2	Pdha2
1450963_at	1.00284	heterogeneous nuclear ribonucleoprotein F	Hnrpf
1450964_a_at	1.02262	oxysterol binding protein-like 9	Osbpl9
1450965_at	1.05884	testis expressed gene 261	Tex261
1450966_at	0.91561	carnitine O-octanoyltransferase	Crot
1450967_at	0.86457	protein tyrosine phosphatase-like A domain containing 2	Ptplad2
1450968_at	0.95012	"ubiquinol-cytochrome c reductase, Rieske iron-sulfur polypeptide 1"	Uqcrcfs1
1450969_at	0.97671	"propionyl Coenzyme A carboxylase, beta polypeptide"	Pccb
1450970_at	1.03444	"glutamate oxaloacetate transaminase 1, soluble"	Got1
1450971_at	1.03857	growth arrest and DNA-damage-inducible 45 beta	Gadd45b
1450972_at	1.09226	RIKEN cDNA 3110040N11 gene	3110040N11Rik
1450973_s_at	0.76827	mitogen activated protein kinase binding protein 1	Mapkbp1
1450974_at	1.07573	tissue inhibitor of metalloproteinase 4	Timp4
1450975_at	0.92321	"calcium channel, voltage-dependent, gamma subunit 4"	Cacng4
1450976_at	1.05639	N-myc downstream regulated gene 1	Ndrg1
1450977_s_at	0.96764	N-myc downstream regulated gene 1 /// N-myc downstream regulated-like	Ndrg1 /// Ndr1
1450978_at	1.10763	"dishevelled, dsh homolog 1 (<i>Drosophila</i>)"	Dvl1
1450979_at	1.02116	CEA-related cell adhesion molecule 14	Ceacam14
1450980_at	0.91176	GTP binding protein 3	Gtpbp3
1450981_at	0.95790	calponin 2	Cnn2
1450982_at	0.84537	"solute carrier family 9 (sodium/hydrogen exchanger), isoform 3 regulator 1"	

Slc9a3r1

1450983_at 1.11017 A kinase (PRKA) anchor protein 8 Akap8

1450984_at 0.91610 tight junction protein 2 Tjp2

1450985_a_at 0.89757 tight junction protein 2 Tjp2

1450986_at 1.12937 nucleolar protein 5 Nol5

1450987_a_at 0.97948 RIKEN cDNA 2310004I24 gene 2310004I24Rik

1450988_at 1.08645 leucine rich repeat containing G protein coupled receptor 5 Lgr5

1450989_at 1.06596 teratocarcinoma-derived growth factor TdGF1

1450990_at 0.97517 glypican 3 Gpc3

1450991_at 0.92137 "DnaJ (Hsp40) homolog, subfamily B, member 7" Dnajb7

1450992_a_at 0.99978 myeloid ecotropic viral integration site 1 Meis1

1450993_at 0.98695 PAS domain containing serine/threonine kinase Pask

1450994_at 0.96800 Rho-associated coiled-coil containing protein kinase 1 Rock1

1450995_at 0.90617 folate receptor 1 (adult) Folr1

1450996_at 0.94954 follicle stimulating hormone beta Fshb

1450997_at 0.88028 serine/threonine kinase 17b (apoptosis-inducing) Stk17b

1450998_at 0.90641 zinc finger protein 110 Zfp110

1450999_a_at 0.92655 RIKEN cDNA 1700029H14 gene 1700029H14Rik

1451000_at 1.07092 transmembrane protein 126A Tmem126a

1451001_at 0.93533 RIKEN cDNA 1700019N19 gene 1700019N19Rik

1451002_at 0.87896 "aconitase 2, mitochondrial" Aco2

1451003_at 1.02751 mitogen-activated protein kinase kinase kinase 7 interacting protein 2

Map3k7ip2

1451004_at 0.92536 activin receptor IIA Acvr2a

1451005_at 1.06442 SMT3 suppressor of mif two 3 homolog 1 (yeast) Sumo1

1451006_at 1.69564 xanthine dehydrogenase Xdh

1451007_at 0.80508 cyclin M2 Cnnm2

1451008_at 0.91545 "ST8 alpha-N-acetyl-neuraminidase alpha-2,8-sialyltransferase 3" St8sia3

1451009_at 0.92307 ring finger protein 151 Rnf151

1451010_at 1.03524 nucleolar protein 11 Nol11

1451011_at 0.89247 zinc finger protein 358 Zfp358

1451012_a_at 1.08511 cold shock domain protein A CSDA

1451013_at 1.09883 "solute carrier family 29 (nucleoside transporters), member 3" Slc29a3

1451014_at 0.75470 receptor tyrosine kinase-like orphan receptor 1 Ror1

1451015_at 0.96303 transketolase Tkt

1451016_at 1.13261 interferon-related developmental regulator 2 Ifrd2

1451017_at 0.97340 ERGIC and golgi 3 Ergic3

1451018_at 0.95902 leptin receptor overlapping transcript-like 1 Leprotl1

1451019_at 0.90812 cathepsin F Ctsf

1451020_at 1.00657 glycogen synthase kinase 3 beta Gsk3b

1451021_a_at 1.08363 Kruppel-like factor 5 Klf5

1451022_at 0.88657 low density lipoprotein receptor-related protein 6 Lrp6

1451023_at 0.97053 "hyperpolarization-activated, cyclic nucleotide-gated K+ 3" Hcn3

1451024_at	0.93323 "endothelial differentiation, G-protein-coupled receptor 6"	Edg6
1451025_at	0.98701 ADP-ribosylation factor-like 1	Arl1
1451026_at	1.05832 FtsJ homolog 3 (E. coli)	Ftsj3
1451027_at	0.99938 brain-specific angiogenesis inhibitor 1-associated protein 2	Baiap2
1451028_at	1.00603 brain-specific angiogenesis inhibitor 1-associated protein 2	Baiap2
1451029_at	1.32137 Bcl2-like 2	Bcl2l2
1451030_at	0.92861 "aldo-keto reductase family 1, member C21"	Akr1c21
1451031_at	0.91888 secreted frizzled-related sequence protein 4	Sfrp4
1451032_at	1.03877 four and a half LIM domains 4	Fhl4
1451033_a_at	1.00795 "transient receptor potential cation channel, subfamily C, member 4"	Trpc4
1451034_at	0.87583 "zinc finger protein 36, C3H type-like 2"	Zfp36l2
1451035_a_at	0.96679 "aldo-keto reductase family 1, member A4 (aldehyde reductase)"	Akr1a4
1451036_at	1.04732 spastic paraplegia 21 homolog (human)	Spg21
1451037_at	1.05529 "protein tyrosine phosphatase, non-receptor type 9"	Ptpn9
1451038_at	0.91961 apelin	Apln
1451039_at	1.14340 RIKEN cDNA 2610027L16 gene	2610027L16Rik
1451040_at	1.04903 histidyl tRNA synthetase 2	Hars2
1451041_at	1.06685 Rho-associated coiled-coil containing protein kinase 2	Rock2
1451042_a_at	1.05844 myc induced nuclear antigen	Mina
1451043_at	1.08153 NIMA (never in mitosis gene a)-related expressed kinase 6	Nek6
1451044_at	0.85295 survivor of motor neuron protein interacting protein 1	Sip1
1451045_at	0.95973 synaptotagmin XIII	Syt13
1451046_at	0.93965 "zinc finger protein, multitype 1"	Zfpm1
1451047_at	1.11498 integral membrane protein 2A	Itm2a
1451048_at	0.93212 methionine aminopeptidase 2	Metap2
1451049_at	1.00257 B-cell receptor-associated protein 31	Bcap31
1451050_at	1.11139 "5'-nucleotidase, cytosolic III"	Nt5c3
1451051_a_at	0.99987 SCY1-like 1 (S. cerevisiae)	Scyl1
1451052_at	0.95270 component of oligomeric golgi complex 8	Cog8
1451053_a_at	1.05172 transformed mouse 3T3 cell double minute 1	Mdm1
1451054_at	0.97697 orosomucoid 1	Orm1
1451055_at	0.93853 "solute carrier family 45, member 2"	Slc45a2
1451056_at	1.12949 "proteasome (prosome, macropain) 26S subunit, non-ATPase, 7"	Psmd7
1451057_x_at	0.99013 dynamin 2	Dnm2
1451058_at	1.15367 malignant T cell amplified sequence 2	Mcts2
1451059_at	1.02706 zinc finger protein 474	Zfp474
1451060_at	0.89585 G protein-coupled receptor 146	Gpr146
1451061_at	0.97508 RIKEN cDNA 1700018B24 gene	1700018B24Rik
1451062_a_at	0.91717 peroxin 2	Pex2
1451063_at	0.90746 syntaxin binding protein 4	Stxbp4
1451064_a_at	1.16636 phosphoserine aminotransferase 1	Psat1
1451065_a_at	1.10887 DEAD (Asp-Glu-Ala-Asp) box polypeptide 39	Ddx39
1451066_at	0.86657 leukocyte receptor cluster (LRC) member 4	Leng4

1451067_at 0.99482 "small glutamine-rich tetratricopeptide repeat (TPR)-containing, alpha" Sgta
 1451068_s_at 1.17325 ribosomal protein S25 Rps25
 1451069_at 1.11415 proviral integration site 3 Pim3
 1451070_at 0.96940 guanosine diphosphate (GDP) dissociation inhibitor 1 Gdi1
 1451071_a_at 0.85218 "ATPase, Na+/K+ transporting, alpha 1 polypeptide" Atp1a1
 1451072_a_at 1.02288 ring finger protein 4 Rnf4
 1451073_at 0.89671 signal peptide peptidase 3 Spp13
 1451074_at 0.90556 ring finger protein 13 Rnf13
 1451075_s_at 1.00963 "CTD (carboxy-terminal domain, RNA polymerase II, polypeptide A) small phosphatase 2" Ctdsp2
 1451076_s_at 1.12802 dendritic cell protein GA17 Ga17
 1451077_at 1.04889 ribosomal protein L5 /// similar to 60S ribosomal protein L5 /// similar to 60S ribosomal protein L5 /// similar to 60S ribosomal protein L5 /// similar to 60S ribosomal protein L5 Rpl5 /// LOC382740 /// LOC665407 /// LOC668032 /// LOC672748 /// LOC675821
 1451078_at 0.95837 RIKEN cDNA 2510039O18 gene 2510039O18Rik
 1451079_at 1.01700 ADP-dependent glucokinase Adpgk
 1451080_at 0.97634 ubiquitin specific peptidase 1 Usp1
 1451081_a_at 0.95588 transcription factor 25 (basic helix-loop-helix) Tcf25
 1451082_at 1.07866 RIKEN cDNA 1300018I05 gene 1300018I05Rik
 1451083_s_at 1.06212 alanyl-tRNA synthetase Aars
 1451084_at 0.92677 "electron transferring flavoprotein, dehydrogenase" Etfdh
 1451085_at 0.99225 RIKEN cDNA C030006K11 gene C030006K11Rik
 1451086_s_at 0.92650 RAS-related C3 botulinum substrate 1 Rac1
 1451087_at 1.07754 WD repeat domain 36 Wdr36
 1451088_a_at 0.96613 oxidase assembly 1-like Oxa11
 1451089_a_at 1.00424 archain 1 Arcn1
 1451090_a_at 1.09859 "eukaryotic translation initiation factor 2, subunit 3, structural gene X-linked" Eif2s3x
 1451091_at 0.89754 thioredoxin domain containing 5 Txndc5
 1451092_a_at 0.92602 RAN GTPase activating protein 1 Rangap1
 1451093_at 0.98947 polymerase (RNA) II (DNA directed) polypeptide E Polr2e
 1451094_at 0.94713 gamma-glutamyltransferase-like 3 Ggtl3
 1451095_at 1.12110 asparagine synthetase Asns
 1451096_at 0.98776 NADH dehydrogenase (ubiquinone) Fe-S protein 2 Ndufs2
 1451097_at 0.90601 vasodilator-stimulated phosphoprotein Vasp
 1451098_at 0.91259 procollagen (type III) N-endopeptidase Pcoln3
 1451099_at 0.86832 membrane bound C2 domain containing protein Mbc2
 1451100_a_at 1.04079 carnitine deficiency-associated gene expressed in ventricle 3 Cdv3
 1451101_a_at 0.93254 ribosomal protein S28 Rps28
 1451102_at 0.97070 "CCR4-NOT transcription complex, subunit 8" Cnot8
 1451103_at 0.88696 "DNA segment, Chr 14, ERATO Doi 500, expressed" D14Ertd500e
 1451104_a_at 1.06148 U1 small nuclear ribonucleoprotein polypeptide A Snrp70

1451105_at 1.00672 vasohibin 2 Wash2
1451106_at 1.24137 RNA binding motif protein 21 Rbm21
1451107_at 0.92563 "TBC1 domain family, member 22a" Tbc1d22a
1451108_at 0.96463 ring finger protein 185 Rnf185
1451109_a_at 0.91306 "neural precursor cell expressed, developmentally down-regulated gene 4"
Nedd4
1451110_at 1.01157 EGL nine homolog 1 (C. elegans) Egl1
1451111_at 1.09233 nucleoporin 133 Nup133
1451112_s_at 0.98002 death-associated protein Dap
1451113_a_at 0.99880 IK cytokine Ik
1451114_at 0.99382 CKLF-like MARVEL transmembrane domain containing 6 Cmtm6
1451115_at 0.89333 protein inhibitor of activated STAT 3 Pias3
1451116_at 0.95728 acetoacetyl-CoA synthetase Aacs
1451117_a_at 0.96507 target of myb1-like 1 (chicken) Tom111
1451118_a_at 0.95762 RIKEN cDNA 2410018C17 gene 2410018C17Rik
1451119_a_at 0.70570 fibulin 1 Fbln1
1451120_at 1.08599 RNA polymerase 1-3 Rpo1-3
1451121_a_at 0.97187 glioma tumor suppressor candidate region gene 2 Gltscr2
1451122_at 1.47530 isopentenyl-diphosphate delta isomerase Idi1
1451123_at 1.14395 RIKEN cDNA C330016O10 gene C330016O10Rik
1451124_at 1.10962 "superoxide dismutase 1, soluble" Sod1
1451125_at 0.94368 cDNA sequence BC017133 BC017133
1451126_at 0.95422 MAF1 homolog (S. cerevisiae) Maf1
1451127_at 1.02624 expressed sequence AW146242 AW146242
1451128_s_at 1.05530 kinesin family member 22 Kif22
1451129_at 0.86664 calbindin 2 Calb2
1451130_at 1.03066 RIKEN cDNA 2010315L10 gene 2010315L10Rik
1451131_at 0.98238 ADP-ribosylation factor-like 6 interacting protein 1 Arl6ip1
1451132_at 0.80621 pre-B-cell leukemia transcription factor interacting protein 1 Pbxip1
1451133_s_at 0.89260 transmembrane protein 168 Tmem168
1451134_a_at 0.85359 TM2 domain containing 2 Tm2d2
1451135_at 0.99950 general transcription factor IIB Gtf2b
1451136_a_at 1.05983 "eukaryotic translation initiation factor 2B, subunit 2 beta" Eif2b2
1451137_a_at 1.01030 bromodomain containing 8 Brd8
1451138_x_at 0.97205 "mesoderm induction early response 1, family member 2" Mier2
1451139_at 2.22427 "solute carrier family 39 (zinc transporter), member 4" Slc39a4
1451140_s_at 0.91636 "protein kinase, AMP-activated, gamma 2 non-catalytic subunit" Prkag2
1451141_at 1.00753 methyltransferase like 8 Mettl8
1451142_at 1.04588 exocyst complex component 1 Exoc1
1451143_at 1.05002 RIKEN cDNA 1110006G06 gene 1110006G06Rik
1451144_at 1.10883 brix domain containing 2 Bxdc2
1451145_s_at 0.90419 transmembrane protein 111 Tmem111
1451146_at 0.94338 zinc finger protein 386 (Kruppel-like) /// similar to zinc finger protein 386

(Kruppel-like) isoform a Zfp386 /// LOC676952

1451147_x_at 0.72880 "cold shock domain containing C2, RNA binding" Csdc2

1451148_at 0.88793 PTEN induced putative kinase 1 Pink1

1451149_at 1.16758 phosphoglucomutase 2 Pgm2

1451150_at 0.92185 zinc finger protein 410 Zfp410

1451151_s_at 0.92204 zinc finger protein 410 Zfp410

1451152_a_at 0.75240 "ATPase, Na+/K+ transporting, beta 1 polypeptide" Atp1b1

1451153_a_at 0.89891 cysteine and histidine rich 1 Cyhr1

1451154_a_at 1.03023 "CUG triplet repeat, RNA binding protein 2" Cugbp2

1451155_at 0.94343 "CUG triplet repeat, RNA binding protein 2" Cugbp2

1451156_s_at 0.82816 very low density lipoprotein receptor Vldlr

1451157_at 1.01521 ring finger protein 187 Rnf187

1451158_at 0.90450 thyroid hormone receptor interactor 12 Trip12

1451159_at 0.88063 Rho guanine nucleotide exchange factor (GEF) 12 Arhgef12

1451160_s_at 1.20664 poliovirus receptor Pvr

1451161_a_at 1.03425 "EGF-like module containing, mucin-like, hormone receptor-like sequence 1" Emr1

1451162_at 0.98300 heat shock factor binding protein 1 Hsbp1

1451163_at 0.99626 Terf1 (TRF1)-interacting nuclear factor 2 Tinf2

1451164_a_at 1.13739 mitochondrial ribosomal protein S18B Mrps18b

1451165_at 1.00524 limb region 1 like Lmbr1l

1451166_a_at 1.12163 coiled-coil domain containing 101 Ccdc101

1451167_at 1.07838 coiled-coil domain containing 101 Ccdc101

1451168_a_at 1.05950 Rho GDP dissociation inhibitor (GDI) alpha Arhgdia

1451169_at 1.07747 nodal modulator 1 Nomo1

1451170_s_at 1.00593 nodal modulator 1 Nomo1

1451171_at 1.07303 RIKEN cDNA 2310008H04 gene 2310008H04Rik

1451172_at 0.86903 RIKEN cDNA 1200015A19 gene 1200015A19Rik

1451173_at 0.91188 leucine rich repeat containing 49 Lrrc49

1451174_at 1.00222 leucine rich repeat containing 33 Lrrc33

1451175_at 0.94537 signal peptidase complex subunit 3 homolog (S. cerevisiae) Spcs3

1451176_at 0.84585 RIKEN cDNA D430028G21 gene D430028G21Rik

1451177_at 0.81098 "DnaJ (Hsp40) homolog, subfamily B, member 4" Dnajb4

1451178_at 1.03540 mitochondrial ribosomal protein L53 Mrpl53

1451179_a_at 0.94502 quaking Qk

1451180_a_at 0.99646 "5'-nucleotidase, cytosolic III-like" Nt5c3l

1451181_at 0.93069 transmembrane protein 121 Tmem121

1451182_s_at 0.95744 ankyrin repeat domain 54 Ankrd54

1451183_at 1.17284 myosin head domain containing 1 Myohd1

1451184_at 1.16567 heterogeneous nuclear ribonucleoprotein A3 /// similar to heterogeneous nuclear ribonucleoprotein A3 /// similar to heterogeneous nuclear ribonucleoprotein A3 /// similar to heterogeneous nuclear ribonucleoprotein A3 /// hypothetical protein LOC630677 /// similar to heterogeneous nuclear ribonucleoprotein A3 ///

similar to heterogeneous nuclear ribonucleoprotein A3 /// similar to heterogeneous nuclear ribonucleoprotein A3 /// similar to heterogeneous nuclear ribonucleoprotein A3 /// similar to heterogeneous nuclear ribonucleoprotein A3 Hnrpa3 /// LOC545592 /// LOC627828 /// LOC629025 /// LOC630401 /// LOC630677 /// LOC664907 /// LOC668131 /// LOC668559 /// LOC669450 /// LOC674254

1451185_at 1.07223 "splicing factor 3b, subunit 5" Sf3b5
 1451186_at 1.12528 interferon stimulated exonuclease gene 20-like 1 Isg2011
 1451187_at 0.99370 RIKEN cDNA 0610037P05 gene 0610037P05Rik
 1451188_at 1.01028 WD repeat domain 26 Wdr26
 1451189_at 1.05088 "zinc finger, SWIM domain containing 1" Zswim1
 1451190_a_at 0.72737 SH3-binding kinase 1 Sbk1
 1451191_at 0.99833 cellular retinoic acid binding protein II Crabp2
 1451192_a_at 0.96729 tetratricopeptide repeat domain 4 Ttc4
 1451193_x_at 1.02027 tetratricopeptide repeat domain 4 Ttc4
 1451194_at 1.10157 "aldolase 2, B isoform" Aldob
 1451195_a_at 0.89210 thioredoxin domain containing 1 Txndc1
 1451196_at 0.85645 yippee-like 5 (Drosophila) Ypel5
 1451197_s_at 1.04599 GATA zinc finger domain containing 2A Gatad2a
 1451198_at 0.98610 GATA zinc finger domain containing 2A Gatad2a
 1451199_at 1.04844 queuine tRNA-ribosyltransferase domain containing 1 Qtrtd1
 1451200_at 0.89337 kinesin family member 1B Kif1b
 1451201_s_at 1.15389 ribonuclease/angiogenin inhibitor 1 Rnh1
 1451202_at 1.03592 RIKEN cDNA C330007P06 gene C330007P06Rik
 1451203_at 1.03965 myoglobin Mb
 1451204_at 1.06886 "scavenger receptor class A, member 5 (putative)" Scara5
 1451205_at 1.04497 "proteasome (prosome, macropain) subunit, beta type 4" Psmb4
 1451206_s_at 0.78814 "pleckstrin homology, Sec7 and coiled-coil domains, binding protein" Pscdbp
 1451207_at 0.97815 calcium binding atopy-related autoantigen 1 Cbara1
 1451208_at 1.12994 eukaryotic translation termination factor 1 Etf1
 1451209_at 0.92590 longevity assurance homolog 5 (S. cerevisiae) Lass5
 1451210_at 1.04243 phosphatidic acid phosphatase type 2c Ppap2c
 1451211_a_at 0.94057 ligatin Lgtn
 1451212_at 0.96130 coiled-coil domain containing 21 Ccdc21
 1451213_at 0.92342 peroxisomal biogenesis factor 11b Pex11b
 1451214_at 0.96906 kelch repeat and BTB (POZ) domain containing 2 Kbtbd2
 1451215_at 0.82330 --- ---
 1451216_at 1.02358 zinc finger protein 330 Zfp330
 1451217_a_at 0.89485 IMP1 inner mitochondrial membrane peptidase-like (S. cerevisiae) Immp11
 1451218_at 0.96166 "ER degradation enhancer, mannosidase alpha-like 1" Edem1
 1451219_at 0.93644 ORM1-like 1 (S. cerevisiae) Ormdl1
 1451220_at 1.07161 RIKEN cDNA 2310040A13 gene 2310040A13Rik
 1451221_at 0.94922 cDNA sequence BC018507 BC018507
 1451222_at 0.86016 basic transcription factor 3-like 4 Btf3l4

1451223_a_at 1.01014 basic transcription factor 3-like 4 Btf3l4
1451224_at 0.98224 secretory carrier membrane protein 5 Scamp5
1451225_at 0.99429 "protein tyrosine phosphatase, non-receptor type 11" Ptpn11
1451226_at 0.84977 peroxisomal biogenesis factor 6 Pex6
1451227_a_at 0.88363 "solute carrier family 10 (sodium/bile acid cotransporter family), member 3" Slc10a3
1451228_a_at 0.96143 syncollin Syncn
1451229_at 0.64768 histone deacetylase 11 Hdac11
1451230_a_at 0.93764 WW domain binding protein 5 Wbp5
1451231_a_at 1.01410 cullin 2 Cul2
1451232_at 0.90429 CD151 antigen Cd151
1451233_at 1.12946 Tnf receptor-associated factor 2 Traf2
1451234_at 1.00546 cDNA sequence BC021381 BC021381
1451235_at 0.97085 cell cycle exit and neuronal differentiation 1 Cend1
1451236_at 0.97164 "RAS-like, estrogen-regulated, growth-inhibitor" Rerg
1451237_s_at 0.89067 RNA binding motif protein 7 Rbm7
1451238_at 0.88484 RIKEN cDNA 1200003C05 gene 1200003C05Rik
1451239_a_at 0.97377 "solute carrier family 26 (sulfate transporter), member 1" Slc26a1
1451240_a_at 0.96867 glyoxalase 1 Glo1
1451241_at 0.88209 laminin B1 subunit 1 Lamb1-1
1451242_a_at 0.89927 "protein phosphatase 5, catalytic subunit" Ppp5c
1451243_at 1.07625 arginyl aminopeptidase (aminopeptidase B) Rnpep
1451244_a_at 0.85081 zinc finger protein 422 Zfp422
1451245_at 0.90787 leucine rich repeat containing 3B Lrrc3b
1451246_s_at 1.03388 aurora kinase B Aurkb
1451247_at 1.08204 major facilitator superfamily domain containing 1 Mfsd1
1451248_at 1.06587 protein arginine N-methyltransferase 7 Prmt7
1451249_at 1.05767 TRM1 tRNA methyltransferase 1 homolog (S. cerevisiae) Trmt1
1451250_at 0.85608 G protein-regulated inducer of neurite outgrowth 1 Gprin1
1451251_at 1.08660 amyloid beta precursor protein (cytoplasmic tail) binding protein 2 Appbp2
1451252_at 0.96506 interferon regulatory factor 2 binding protein 1 Irf2bp1
1451253_at 0.94897 PX domain containing serine/threonine kinase Pxx
1451254_at 1.06788 "inhibitor of kappa light polypeptide enhancer in B-cells, kinase complex-associated protein" Ikbkap
1451255_at 0.82944 lipolysis stimulated lipoprotein receptor Lsr
1451256_at 1.00843 G protein-coupled receptor 172B Gpr172b
1451257_at 1.07228 acyl-CoA synthetase long-chain family member 6 Acsl6
1451258_at 1.02884 prostate stem cell antigen Psca
1451259_at 0.98668 "REX2, RNA exonuclease 2 homolog (S. cerevisiae)" Rexo2
1451260_at 0.95731 "aldehyde dehydrogenase 1 family, member B1" Aldh1b1
1451261_s_at 0.98836 expressed sequence AW049765 AW049765
1451262_a_at 1.21279 JTV1 gene Jtv1
1451263_a_at 0.99235 "fatty acid binding protein 4, adipocyte" Fabp4

1451264_at 1.05196 FERM domain containing 6 Frmd6
1451265_at 1.12583 coiled-coil domain containing 115 Ccdc115
1451266_at 1.04641 mitochondrial ribosomal protein L50 Mrpl50
1451267_at 0.95978 SHANK-associated RH domain interacting protein Sharpin
1451268_at 0.97521 translocation associated membrane protein 1-like 1 Tram111
1451269_at 0.99569 PDZ domain containing 11 Pdzd11
1451270_at 1.10040 dual specificity phosphatase 18 Dusp18
1451271_a_at 0.90549 acetyl-Coenzyme A acetyltransferase 1 Acat1
1451272_a_at 1.14349 ubiquitin-conjugating enzyme E2F (putative) Ube2f
1451273_x_at 0.97447 cDNA sequence BC025546 BC025546
1451274_at 0.90379 oxoglutarate dehydrogenase (lipoamide) Ogdh
1451275_at 0.96365 RIKEN cDNA E030041M21 gene E030041M21Rik
1451276_at 0.92515 RIKEN cDNA E030041M21 gene E030041M21Rik
1451277_at 1.11645 "zinc binding alcohol dehydrogenase, domain containing 2" Zadh2
1451278_a_at 1.05285 RIKEN cDNA 2610205E22 gene 2610205E22Rik
1451279_at 1.00022 ELKS/RAB6-interacting/CAST family member 1 Erc1
1451280_at 0.81556 "cyclic AMP-regulated phosphoprotein, 21" Arpp21
1451281_at 1.06715 zinc finger protein 96 Zfp96
1451282_at 1.06982 "centaurin, delta 3" Centd3
1451283_at 1.16636 RIKEN cDNA 1810073G14 gene 1810073G14Rik
1451284_at 0.91022 "Yip1 domain family, member 3" Yipf3
1451285_at 1.08188 "fusion, derived from t(12;16) malignant liposarcoma (human)" Fus
1451286_s_at 1.05254 "fusion, derived from t(12;16) malignant liposarcoma (human)" Fus
1451287_s_at 1.02487 RIKEN cDNA 2810003C17 gene 2810003C17Rik
1451288_s_at 0.88946 RIKEN cDNA 1810043G02 gene 1810043G02Rik
1451289_at 1.30975 doublecortin and calcium/calmodulin-dependent protein kinase-like 1
Dcamk1l
1451290_at 0.83911 microtubule-associated protein 1 light chain 3 alpha Map1lc3a
1451291_at 1.10627 oligonucleotide/oligosaccharide-binding fold containing 2B Obfc2b
1451292_at 0.92355 Zinc finger protein 212 Zfp212
1451293_at 1.05083 "RNA, U3 small nucleolar interacting protein 2" Rnu3ip2
1451294_s_at 1.05963 small nuclear ribonucleoprotein E Snrpe
1451295_a_at 0.96022 chromodomain helicase DNA binding protein 4 Chd4
1451296_x_at 1.06903 "poly A binding protein, cytoplasmic 4 /// similar to Poly(A) binding protein,
cytoplasmic 4, isoform 1" Pabpc4 /// LOC432881
1451297_at 1.10657 gulonolactone (L-) oxidase Gulo
1451298_at 0.91705 "pleckstrin homology domain containing, family H (with MyTH4 domain)
member 3" Plekhh3
1451299_at 1.06823 "protein kinase, X-linked" Prkx
1451300_a_at 0.99809 "CHMP family, member 7" Chmp7
1451301_at 1.02722 tropomodulin 2 Tmod2
1451302_at 1.00768 RIKEN cDNA 1110012L19 gene 1110012L19Rik
1451303_at 0.95682 cDNA sequence BC002230 BC002230

1451304_at 1.05978 transmembrane protein 143 Tmem143
1451305_at 0.92621 "PKD2 interactor, golgi and endoplasmic reticulum associated 1" Pgea1
1451306_at 0.85617 cell division cycle associated 7 like Cdca7l
1451307_at 1.05627 mitochondrial ribosomal protein L14 Mrpl14
1451308_at 1.01387 "elongation of very long chain fatty acids (FEN1/Elo2, SUR4/Elo3, yeast)-like 4" Elovl4
1451309_at 0.87715 Rho GTPase activating protein 1 Arhgap1
1451310_a_at 0.87662 cathepsin L Ctsl
1451311_a_at 0.86203 adiponectin receptor 1 Adipor1
1451312_at 0.95734 NADH dehydrogenase (ubiquinone) Fe-S protein 7 Ndufs7
1451313_a_at 0.91814 RIKEN cDNA 1110067D22 gene 1110067D22Rik
1451314_a_at 1.30748 vascular cell adhesion molecule 1 Vcam1
1451315_at 1.09538 transmembrane protein 101 Tmem101
1451316_a_at 0.94577 phosphatidylinositol binding clathrin assembly protein Picalm
1451317_at 0.97010 YTH domain family 2 Ythdf2
1451318_a_at 1.13964 Yamaguchi sarcoma viral (v-yes-1) oncogene homolog /// similar to Yamaguchi sarcoma viral (v-yes-1) oncogene homolog Lyn /// LOC676654
1451319_at 1.04354 SUMO1/sentrin specific peptidase 1 Senp1
1451320_at 0.97269 RIKEN cDNA 3110043J09 gene 3110043J09Rik
1451321_a_at 0.96376 RIKEN cDNA 0610033I05 gene 0610033I05Rik
1451322_at 0.78641 RIKEN cDNA 2310016A09 gene 2310016A09Rik
1451323_at 1.15525 zinc finger protein 7 Zfp7
1451324_s_at 1.00369 forty-two-three domain containing 1 Fyttd1
1451325_at 0.88063 forty-two-three domain containing 1 Fyttd1
1451326_at 0.99160 abhydrolase domain containing 14b Abhd14b
1451327_a_at 1.01987 transmembrane emp24 domain containing 1 Tmed1
1451328_at 0.96721 pecanex-like 3 (Drosophila) Pcnx3
1451329_at 1.06082 nudix (nucleoside diphosphate linked moiety X)-type motif 22 Nudt22
1451330_a_at 1.00228 inositol polyphosphate-5-phosphatase B Inpp5b
1451331_at 0.90871 "protein phosphatase 1, regulatory (inhibitor) subunit 1B" Ppp1r1b
1451332_at 0.99629 zinc finger protein 521 Zfp521
1451333_a_at 1.17988 proacrosin binding protein Acrbp
1451334_at 0.97941 RIKEN cDNA 1810009O10 gene 1810009O10Rik
1451335_at 1.05645 placenta-specific 8 Plac8
1451336_at 1.07518 "lectin, galactose binding, soluble 4" Lgals4
1451337_at 0.96846 "proteasome (prosome, macropain) inhibitor subunit 1" Psmf1
1451338_at 1.18915 nischarin Nisch
1451339_at 1.04142 sulfite oxidase Suox
1451340_at 0.94760 AT rich interactive domain 5A (Mrf1 like) Arid5a
1451341_s_at 1.03530 RIKEN cDNA 1700003N22 gene /// expressed sequence AI840826 1700003N22Rik /// AI840826
1451342_at 1.09074 "spondin 1, (f-spondin) extracellular matrix protein" Spon1
1451343_at 1.04520 vacuolar protein sorting 36 (yeast) Vps36

1451344_at 0.84404 transmembrane protein 119 Tmem119
1451345_at 1.02273 methylthioadenosine phosphorylase Mtap
1451346_at 0.97906 methylthioadenosine phosphorylase Mtap
1451347_at 0.98563 coiled-coil domain containing 95 Ccdc95
1451348_at 0.84747 DEP domain containing 6 Depdc6
1451349_at 0.93528 cDNA sequence BC020077 BC020077
1451350_a_at 0.87127 leptin receptor overlapping transcript Leprot
1451351_at 0.91739 tetratricopeptide repeat domain 13 Ttc13
1451352_s_at 0.94819 metastasis associated 3 Mta3
1451353_at 1.10222 transmembrane 6 superfamily member 1 Tm6sf1
1451354_at 1.05256 FAD-dependent oxidoreductase domain containing 1 Foxred1
1451355_at 0.95775 N-acylsphingosine amidohydrolase 3-like Asah3l
1451356_at 0.98929 "acidic (leucine-rich) nuclear phosphoprotein 32 family, member E" Anp32e
1451357_at 0.99272 RIKEN cDNA E130307M08 gene E130307M08Rik
1451358_a_at 0.95588 Rac GTPase-activating protein 1 Racgap1
1451359_at 0.99751 acyltransferase like 2 Ayt12
1451360_at 1.10542 ERGIC and golgi 2 Ergic2
1451361_a_at 1.04822 patatin-like phospholipase domain containing 7 Pnpla7
1451362_at 0.89191 "RAB7, member RAS oncogene family-like 1" Rab7l1
1451363_a_at 0.96325 DENN/MADD domain containing 2D Dennd2d
1451364_at 0.95476 polymerase (RNA) III (DNA directed) polypeptide G like Polr3gl
1451365_at 1.03288 RNA binding motif protein 19 Rbm19
1451366_at 1.04239 "COP9 (constitutive photomorphogenic) homolog, subunit 6 (Arabidopsis thaliana)" Cops6
1451367_at 0.87425 "COP9 (constitutive photomorphogenic) homolog, subunit 6 (Arabidopsis thaliana)" Cops6
1451368_at 1.02109 "asparagine-linked glycosylation 1 homolog (yeast, beta-1,4-mannosyltransferase)" Alg1
1451369_at 0.95406 COMM domain containing 5 Commd5
1451370_at 1.01232 hypothetical protein LOC622403 LOC622404
1451371_at 0.99881 melanocortin 2 receptor accessory protein Mrap
1451372_a_at 1.03255 ADP-ribosyltransferase 1 Art1
1451373_at 0.94396 "UDP glycosyltransferases 3 family, polypeptide A1 /// similar to UDP glycosyltransferases 3 family, polypeptide A1" Ugt3a1 /// LOC666512
1451374_x_at 0.97829 chemokine-like factor Cklf
1451375_at 1.11947 ets homologous factor Ehf
1451376_at 1.11956 RIKEN cDNA 5730596K20 gene 5730596K20Rik
1451377_a_at 0.98645 "achalasia, adrenocortical insufficiency, alacrimia" Aaas
1451378_at 1.04000 "UTP6, small subunit (SSU) processome component, homolog (yeast)" Utp6
1451379_at 1.05495 "RAB22A, member RAS oncogene family" Rab22a
1451380_at 0.94489 "zinc finger, FYVE domain containing 19" Zfyve19
1451381_at 0.86100 RIKEN cDNA 1810020D17 gene 1810020D17Rik
1451382_at 0.91524 "ChaC, cation transport regulator-like 1 (E. coli)" Chac1

1451383_a_at 1.18700 conserved helix-loop-helix ubiquitous kinase Chuk
1451384_at 0.93290 jumonji domain containing 5 Jmjd5
1451385_at 1.23835 RIKEN cDNA 2310056P07 gene 2310056P07Rik
1451386_at 1.05835 biliverdin reductase B (flavin reductase (NADPH)) Blvrb
1451387_s_at 1.08555 cutA divalent cation tolerance homolog (E. coli) Cuta
1451388_a_at 0.99137 "ATPase, Class VI, type 11B" Atp11b
1451389_at 1.11855 "zinc finger, CSL-type containing 3" Zcs13
1451390_s_at 0.98048 "zinc finger, AN1 type domain 2B" Zfand2b
1451391_at 1.07972 RIKEN cDNA 2700050L05 gene 2700050L05Rik
1451392_at 0.94910 RNA binding motif and ELMO domain 1 Rbed1
1451393_at 1.00382 peroxisome biogenesis factor 26 Pex26
1451394_at 0.95015 dipeptidylpeptidase 6 Dpp6
1451395_at 0.95614 coiled-coil domain containing 92 Ccdc92
1451396_at 1.02811 protein-O-mannosyltransferase 2 Pomt2
1451397_at 1.06050 --- ---
1451398_at 1.02824 carbonyl reductase 4 Cbr4
1451399_at 0.94902 paroxysmal nonkinesinogenic dyskinesia Pnkd
1451400_at 1.09318 cDNA sequence BC023488 BC023488
1451401_a_at 0.94734 RIKEN cDNA 0610009K11 gene 0610009K11Rik
1451402_at 1.01150 ecdysoneless homolog (Drosophila) Ecd
1451403_at 1.05132 cDNA sequence BC024868 BC024868
1451404_at 1.06608 open reading frame 19 ORF19
1451405_at 0.94464 "propionyl-Coenzyme A carboxylase, alpha polypeptide" Pcca
1451406_a_at 0.98470 proprotein convertase subtilisin/kexin type 5 Pcsk5
1451407_at 0.87872 junction adhesion molecule 4 Jam4
1451408_at 1.04404 TruB pseudouridine (psi) synthase homolog 2 (E. coli) Trub2
1451409_at 0.96159 RIKEN cDNA 2210021J22 gene 2210021J22Rik
1451410_a_at 0.94877 cysteine-rich protein 3 Crip3
1451411_at 0.86658 "G protein-coupled receptor, family C, group 5, member B" Gprc5b
1451412_a_at 1.00466 intraflagellar transport 20 homolog (Chlamydomonas) Ift20
1451413_at 0.94669 calpastatin Cast
1451414_at 0.99596 "DNA segment, Chr 13, Wayne State University 177, expressed" D13Wsu177e
1451415_at 1.41143 RIKEN cDNA 1810011O10 gene 1810011O10Rik
1451416_a_at 1.03542 "transglutaminase 1, K polypeptide" Tgm1
1451417_at 0.90409 breast cancer 1 Brca1
1451418_a_at 0.80788 splA/ryanodine receptor domain and SOCS box containing 4 Spsb4
1451419_at 0.75726 splA/ryanodine receptor domain and SOCS box containing 4 Spsb4
1451420_at 0.87231 RIKEN cDNA 2610204L23 gene 2610204L23Rik
1451421_a_at 0.92059 rogdi homolog (Drosophila) Rogdi
1451422_at 0.93546 myosin XVIIIa Myo18a
1451423_at 0.83845 tubulin-specific chaperone c Tbcc
1451424_at 0.88369 "gamma-aminobutyric acid (GABA-A) receptor, pi" Gabrp
1451425_a_at 1.01788 "makorin, ring finger protein, 1" Mkrn1

1451426_at 0.98721 "DNA segment, Chr 11, Lothar Hennighausen 2, expressed" D11Lgp2e
 1451427_a_at 1.00094 EGF-like domain 7 Egfl7
 1451428_x_at 0.97035 EGF-like domain 7 Egfl7
 1451429_at 0.98017 glucose-fructose oxidoreductase domain containing 2 Gfod2
 1451430_at 0.94418 slit homolog 2 (Drosophila) Slit2
 1451431_a_at 0.97403 dysbindin (dystrobrevin binding protein 1) domain containing 2 Dbnidd2
 1451432_x_at 0.85942 "alkB, alkylation repair homolog 4 (E. coli)" Alkbh4
 1451433_at 1.03251 major facilitator superfamily domain containing 3 Mfsd3
 1451434_s_at 1.08941 RIKEN cDNA 5430405G24 gene 5430405G24Rik
 1451435_at 0.79225 cut-like 1 (Drosophila) Cutl1
 1451436_at 1.11046 "sno, strawberry notch homolog 1 (Drosophila)" Sbno1
 1451437_at 1.09483 "zinc finger, DHHC domain containing 20" Zdhhc20
 1451438_s_at 1.02284 "C-type lectin domain family 2, member h" Clec2h
 1451439_at 1.12289 cDNA sequence BC027231 BC027231
 1451440_at 1.03066 chondrolectin Chodl
 1451441_at 1.04536 RIKEN cDNA 2210415F13 gene 2210415F13Rik
 1451442_at 1.03990 coiled-coil domain containing 104 Ccdc104
 1451443_at 1.09379 nuclear factor I/X Nfix
 1451444_s_at 1.01120 RIKEN cDNA 2700008B19 gene 2700008B19Rik
 1451445_at 0.93052 uridine monophosphate synthetase Umps
 1451446_at 0.74579 anthrax toxin receptor 1 Antxr1
 1451447_at 1.03773 CUE domain containing 1 Cuedc1
 1451448_a_at 0.95787 RIKEN cDNA 1110005A03 gene 1110005A03Rik
 1451449_at 0.98776 RIKEN cDNA 4933407N01 gene 4933407N01Rik
 1451450_at 0.63478 RIKEN cDNA 2010011I20 gene 2010011I20Rik
 1451451_at 0.79250 grancalcin Gca
 1451452_a_at 0.83313 regulator of G-protein signaling 16 Rgs16
 1451453_at 1.04971 death-associated kinase 2 Dapk2
 1451454_at 0.97854 protocadherin 20 Pcdh20
 1451455_at 0.98281 cDNA sequence BC051244 BC051244
 1451456_at 1.27048 RIKEN cDNA 6430706D22 gene 6430706D22Rik
 1451457_at 1.50269 "sterol-C5-desaturase (fungal ERG3, delta-5-desaturase) homolog (S. cerevisiae)" Sc5d
 1451458_at 1.17617 transmembrane protein 2 Tmem2
 1451459_at 1.10288 AT hook containing transcription factor 1 Ahctf1
 1451460_a_at 0.98517 "solute carrier family 22 (organic anion transporter), member 7" Slc22a7
 1451461_a_at 1.46468 "aldolase 3, C isoform" Aldoc
 1451462_a_at 1.02881 interferon (alpha and beta) receptor 2 Ifnar2
 1451463_at 0.99324 Rho GTPase activating protein 8 Arhgap8
 1451464_at 0.94628 microfibrillar-associated protein 3 Mfap3
 1451465_at 0.92601 ubiquitin-like 7 (bone marrow stromal cell-derived) Ubl7
 1451466_at 1.08672 "DNA segment, Chr 16, ERATO Doi 472, expressed" D16Ertd472e
 1451467_s_at 0.94213 GTP binding protein 5 Gtpbp5

1451468_s_at 1.16248 exportin 5 Xpo5
1451469_at 0.85936 RIKEN cDNA D530005L17 gene D530005L17Rik
1451470_s_at 1.14929 eukaryotic translation initiation factor 5A Eif5a
1451471_at 0.94623 RIKEN cDNA 3230401I01 gene 3230401I01Rik
1451472_at 0.94095 RIKEN cDNA 3230401I01 gene 3230401I01Rik
1451473_a_at 0.98477 "crystallin, zeta (quinone reductase)-like 1" Cryz11
1451474_a_at 1.09774 "poly (ADP-ribose) polymerase family, member 8" Parp8
1451475_at 0.90389 Plexin D1 Plxd1
1451476_at 1.01328 "zinc finger, DHHC domain containing 8" Zdhhc8
1451477_at 0.93033 RIKEN cDNA 1700029I01 gene 1700029I01Rik
1451478_at 1.16929 angiopoietin-like 7 Angptl7
1451479_a_at 0.70457 transmembrane protein 53 Tmem53
1451480_at 1.07598 E2F transcription factor 4 E2f4
1451481_s_at 0.94545 organic solute transporter alpha Osta
1451482_at 0.88909 cDNA sequence BC027344 BC027344
1451483_s_at 0.99620 RIKEN cDNA 1700054N08 gene 1700054N08Rik
1451484_a_at 0.85036 synapsin I Syn1
1451485_at 0.97481 RIKEN cDNA 3300001P08 gene 3300001P08Rik
1451486_at 0.85534 RIKEN cDNA 1200006F02 gene 1200006F02Rik
1451487_at 1.05661 Rab9 effector protein with kelch motifs Rabepk
1451488_at 0.89893 RIKEN cDNA 1110028A07 gene 1110028A07Rik
1451489_at 0.97070 "solute carrier family 25, member 35" Slc25a35
1451490_at 0.85318 lysophospholipase-like 1 Lyplal1
1451491_at 0.93386 RIKEN cDNA 4930556P03 gene 4930556P03Rik
1451492_at 0.86567 Src-like-adaptor 2 Sla2
1451493_at 0.82002 Nedd4 family interacting protein 1 Ndfip1
1451494_at 0.96165 WW domain containing adaptor with coiled-coil Wac
1451495_at 0.93338 WW domain containing adaptor with coiled-coil Wac
1451496_at 1.05691 metastasis suppressor 1 Mtss1
1451497_at 1.07161 "casein kinase 1, alpha 1" Csnk1a1
1451498_at 0.90398 cDNA sequence BC004853 BC004853
1451499_at 1.07404 Ca²⁺-dependent activator protein for secretion 2 Cadps2
1451500_at 1.04168 Usher syndrome 1C binding protein 1 Ushbp1
1451501_a_at 0.62568 growth hormone receptor Ghr
1451502_at 0.92443 "phospholipase A2, group X" Pla2g10
1451503_at 1.01958 nucleolar protein 3 (apoptosis repressor with CARD domain) Nol3
1451504_at 1.01045 coiled-coil-helix-coiled-coil-helix domain containing 3 Chchd3
1451505_at 0.94683 coiled-coil-helix-coiled-coil-helix domain containing 5 Chchd5
1451506_at 1.03674 myocyte enhancer factor 2C Mef2c
1451507_at 0.92077 myocyte enhancer factor 2C Mef2c
1451508_at 0.99604 RIKEN cDNA 1700108L22 gene 1700108L22Rik
1451509_at 1.01445 "TAF9 RNA polymerase II, TATA box binding protein (TBP)-associated factor" Taf9

1451510_s_at 1.03005 oleoyl-ACP hydrolase Olah
1451511_at 1.00266 3-hydroxyisobutyryl-Coenzyme A hydrolase Hibch
1451512_s_at 0.95743 3-hydroxyisobutyryl-Coenzyme A hydrolase Hibch
1451513_x_at 0.94786 "serine (or cysteine) peptidase inhibitor, clade A, member 1a /// serine (or cysteine) preptidase inhibitor, clade A, member 1b" Serpina1a /// Serpina1b
1451514_at 1.06495 ubiquitin specific peptidase 43 Usp43
1451515_s_at 1.01938 glycine-N-acyltransferase Glyat
1451516_at 1.02609 Ras homolog enriched in brain like 1 Rheb1
1451517_at 0.99606 Rho-related BTB domain containing 2 Rhobtb2
1451518_at 1.00644 zinc finger protein 709 Zfp709
1451519_at 0.91583 ring finger protein 2 Rnf2
1451520_at 1.02567 --- ---
1451521_x_at 0.95984 Williams-Beuren syndrome chromosome region 1 homolog (human) Wbscr1
1451522_s_at 1.02360 leucine-rich repeats and calponin homology (CH) domain containing 4 Lrch4
1451523_a_at 0.94535 MIF4G domain containing Mif4gd
1451524_at 1.06697 F-box and WD-40 domain protein 2 Fbxw2
1451525_at 0.96152 Rho GTPase activating protein 12 Arhgap12
1451526_at 1.03354 Rho GTPase activating protein 12 Arhgap12
1451527_at 0.92463 procollagen C-endopeptidase enhancer 2 Pcolce2
1451528_at 0.88661 cDNA sequence BC025833 BC025833
1451529_at 1.29058 "small glutamine-rich tetratricopeptide repeat (TPR)-containing, beta" Sgfb
1451530_at 0.81477 epidermal growth factor receptor Egfr
1451531_at 1.01311 COBW domain containing 1 Cbwd1
1451532_s_at 1.25562 six transmembrane epithelial antigen of the prostate 1 Steap1
1451533_at 1.08073 cDNA sequence BC022687 BC022687
1451534_at 1.08706 "secretagogin, EF-hand calcium binding protein" Scgn
1451535_at 1.15117 interleukin 31 receptor A Il31ra
1451536_at 0.87649 mitochondrial fission regulator 1 Mtf1
1451537_at 1.04468 chitinase 3-like 1 Chi3l1
1451538_at 0.93406 SRY-box containing gene 9 Sox9
1451539_at 0.67616 BAI1-associated protein 2-like 1 Baiap2l1
1451540_at 1.02462 mannose phosphate isomerase 1 Mpi1
1451541_at 1.00594 BCS1-like (yeast) Bcs1l
1451542_at 1.00587 single-stranded DNA binding protein 2 Ssbp2
1451543_at 0.90793 F-box only protein 21 Fbxo21
1451544_at 1.14768 TAP binding protein-like Tapbp1
1451545_at 0.82838 tudor domain containing 3 Tdrd3
1451546_s_at 1.07123 transmembrane protein 40 Tmem40
1451547_at 1.03612 iodotyrosine deiodinase Iyd
1451548_at 1.01357 --- ---
1451549_at 1.10375 cDNA sequence BC016201 BC016201
1451550_at 0.91190 Eph receptor B3 Ephb3
1451551_at 1.09175 keratin 84 Krt84

1451552_at 1.07361 Similar to lipoyltransferase MGC28431
1451553_at 0.89815 ADP-ribosyltransferase 5 Art5
1451554_a_at 1.00137 anterior pharynx defective 1a homolog (C. elegans) Aph1a
1451555_at 1.17154 neurolysin (metallopeptidase M3 family) Nln
1451556_a_at 1.01262 RIKEN cDNA 2700078E11 gene 2700078E11Rik
1451557_at 0.92367 tyrosine aminotransferase Tat
1451558_at 1.09431 "F-box and WD-40 domain protein 7, archipelago homolog (Drosophila)"
Fbxw7
1451559_a_at 1.03878 dehydrogenase/reductase (SDR family) member 4 Dhrs4
1451560_at 0.96039 proline rich 12 Prr12
1451561_at 0.97636 proline rich 12 Prr12
1451562_at 0.88841 cDNA sequence BC006662 BC006662
1451563_at 0.92900 "EGF-like module containing, mucin-like, hormone receptor-like sequence
4" Emr4
1451564_at 1.40086 "poly (ADP-ribose) polymerase family, member 14" Parp14
1451565_s_at 0.96728 urocanase domain containing 1 Uroc1
1451566_at 0.81251 cDNA sequence BC005471 BC005471
1451567_a_at 1.11063 interferon activated gene 203 Ifi203
1451568_at 1.18092 RIKEN cDNA A630054L15 gene A630054L15Rik
1451569_at 1.04165 "nuclear receptor subfamily 2, group C, member 2" Nr2c2
1451570_a_at 0.95751 RIKEN cDNA 6720467C03 gene 6720467C03Rik
1451571_s_at 0.96085 RIKEN cDNA 9130019O22 gene /// RIKEN cDNA E430018J23 gene ///
RIKEN cDNA 8030466O12 gene /// RIKEN cDNA 6430604K15 gene 9130019O22Rik ///
E430018J23Rik /// 8030466O12Rik /// 6430604K15Rik
1451572_a_at 0.93323 RIKEN cDNA 5230400G24 gene /// hypothetical protein LOC637796
5230400G24Rik /// LOC637796
1451573_a_at 1.12390 syntaxin 4A (placental) Stx4a
1451574_at 0.99008 B-cell CLL/lymphoma 9 Bcl9
1451575_a_at 1.05650 nudix (nucleotide diphosphate linked moiety X)-type motif 3 Nudt3
1451576_at 0.98638 "protein kinase, DNA activated, catalytic polypeptide" Prkdc
1451577_at 1.04615 zinc finger and BTB domain containing 20 Zbtb20
1451578_at 1.06529 coiled-coil domain containing 65 Ccdc65
1451579_at 0.78669 cDNA sequence BC024139 BC024139
1451580_a_at 0.86364 transthyretin Ttr
1451581_at 1.03084 "SAPS domain family, member 2" Saps2
1451582_at 0.95832 tubby like protein 1 Tulp1
1451583_a_at 0.93289 cDNA sequence BC025076 BC025076
1451584_at 1.01285 hepatitis A virus cellular receptor 2 Havcr2
1451585_x_at 0.89785 splA/ryanodine receptor domain and SOCS box containing 2 Spsb2
1451586_at 0.98007 testis enhanced gene transcript Tegt
1451587_a_at 1.03181 "TIP41, TOR signalling pathway regulator-like (S. cerevisiae)" Tipr1
1451588_at 1.08461 RIKEN cDNA 1810022C23 gene 1810022C23Rik
1451589_at 0.84603 opposite strand transcription unit to Stag3 Gats

1451590_at 0.97109 complexin 4 Cplx4
1451591_a_at 0.78012 ephrin B1 Efnb1
1451592_at 0.89929 Myb protein P42POP P42pop
1451593_at 1.06176 "Histocompatibility 2, K1, K region" H2-K1
1451594_s_at 1.04311 "serine (or cysteine) peptidase inhibitor, clade B, member 6c" Serpinb6c
1451595_a_at 1.00497 "potassium voltage-gated channel, subfamily Q, member 2" Kcnq2
1451596_a_at 1.35444 sphingosine kinase 1 Sphk1
1451597_at 1.02283 "transmembrane protease, serine 11d" Tmprss11d
1451598_at 1.04914 paired box gene 4 Pax4
1451599_at 1.11985 sestrin 2 Sesn2
1451600_s_at 1.01440 esterase 31-like /// esterase 31 LOC13909 /// Es31
1451601_a_at 0.70532 cDNA sequence BC011467 BC011467
1451602_at 1.07244 sorting nexin 6 Snx6
1451603_at 0.96446 retbindin Rtbdn
1451604_a_at 1.01568 "activin A receptor, type II-like 1" Acvr11
1451605_at 1.03472 dyslexia susceptibility 1 candidate 1 homolog (human) Dyx1c1
1451606_at 1.13798 RIKEN cDNA A530016L24 gene A530016L24Rik
1451607_at 0.94565 kallikrein 1-related peptidase b21 Klk1b21
1451608_a_at 1.00508 tetraspanin 33 Tspan33
1451609_at 0.95926 tetraspanin 33 Tspan33
1451610_at 1.15578 cDNA sequence BC024561 BC024561
1451611_at 1.01915 HRAS like suppressor 3 Hrasls3
1451612_at 1.08239 metallothionein 1 Mtl
1451613_at 0.99734 hornerin HrnR
1451614_a_at 0.96264 amelogenin X chromosome Amelx
1451615_at 1.09896 cDNA sequence BC026374 BC026374
1451616_at 0.87553 pregnancy specific glycoprotein pseudogene 1 Psg-ps1
1451617_at 0.95401 rhodopsin Rho
1451618_at 1.02789 rhodopsin Rho
1451619_at 0.90006 golgi phosphoprotein 3-like Golph31
1451620_at 1.01218 C1q-like 3 C1ql3
1451621_at 1.08238 RIKEN cDNA 5830417C01 gene 5830417C01Rik
1451622_at 1.01839 LMBR1 domain containing 1 Lmbrd1
1451623_at 0.86216 mitochondrial ribosomal protein L15 Mrpl15
1451624_a_at 1.06233 "phosphatase, orphan 2" Phospho2
1451625_a_at 0.69829 "complement component 8, gamma subunit" C8g
1451626_x_at 1.07493 Transcribed locus ---
1451627_a_at 0.95015 "solute carrier family 1 (glial high affinity glutamate transporter), member 2"
Slc1a2
1451628_a_at 0.88362 "ankyrin 3, epithelial" Ank3
1451629_at 0.93850 limb-bud and heart Lbh
1451630_at 1.05115 tubulin tyrosine ligase Ttl
1451631_at 1.08609 cDNA sequence BC021395 BC021395

1451632_a_at 0.97726 immunoglobulin heavy chain 1a (serum IgG2a) Igh-1a
 1451633_a_at 0.92041 "guanine nucleotide binding protein (G protein), gamma transducing activity polypeptide 1" Gngt1
 1451634_at 1.26881 RIKEN cDNA 2810051F02 gene 2810051F02Rik
 1451635_at 0.96956 RIKEN cDNA D630002G06 gene /// RIKEN cDNA C730048C13 gene /// similar to integral membrane transport protein UST5r D630002G06Rik /// C730048C13Rik /// LOC673838
 1451636_at 0.96759 ankyrin repeat domain 33 Ankrd33
 1451637_a_at 1.18853 "mucin 10, submandibular gland salivary mucin" Muc10
 1451638_s_at 1.00204 armadillo repeat containing 1 /// similar to armadillo repeat-containing protein Armc1 /// LOC634390
 1451639_at 1.04264 "CCAAT/enhancer binding protein (C/EBP), gamma" Cebpg
 1451640_a_at 0.90800 RIKEN cDNA 1500011J06 gene 1500011J06Rik
 1451641_at 1.02710 debranching enzyme homolog 1 (*S. cerevisiae*) Dbr1
 1451642_at 0.93001 kinesin family member 1B Kif1b
 1451643_a_at 1.08079 "RAB4B, member RAS oncogene family" Rab4b
 1451644_a_at 0.94369 "histocompatibility 2, Q region locus 1 /// RIKEN cDNA 0610037M15 gene" H2-Q1 /// 0610037M15Rik
 1451645_at 1.06603 fibrosin 1 Fbs1
 1451646_at 0.98967 PDLIM1 interacting kinase 1 like Pdik11
 1451647_at 0.86436 "solute carrier family 24 (sodium/potassium/calcium exchanger), member 1" Slc24a1
 1451648_a_at 0.94338 folate receptor 2 (fetal) Folr2
 1451649_a_at 1.09338 WD repeat domain 75 Wdr75
 1451650_at 0.99404 D-aspartate oxidase Ddo
 1451651_at 0.94894 V-set and immunoglobulin domain containing 4 Vsig4
 1451652_a_at 0.96257 RIKEN cDNA 5033428A16 gene 5033428A16Rik
 1451653_a_at 1.01860 RIKEN cDNA 4930430E16 gene 4930430E16Rik
 1451654_x_at 1.06047 cDNA sequence BC012278 BC012278
 1451655_at 0.85241 schlafen 8 Slfn8
 1451656_at 0.90233 calmin Clmn
 1451657_a_at 0.88785 cytosolic ovarian carcinoma antigen 1 Coval
 1451658_a_at 0.98964 polymerase (RNA) III (DNA directed) polypeptide C Polr3c
 1451659_at 0.97131 "FYVE, RhoGEF and PH domain containing 4" Fgd4
 1451660_a_at 0.94466 homeo box B6 Hoxb6
 1451661_at 1.05156 A kinase (PRKA) anchor protein 4 Akap4
 1451662_x_at 1.06798 A kinase (PRKA) anchor protein 4 Akap4
 1451663_a_at 0.85236 tripartite motif protein 3 Trim3
 1451664_x_at 1.02861 "killer cell lectin-like receptor, subfamily A, member 1 /// killer cell lectin-like receptor subfamily A, member 12 /// killer cell lectin-like receptor subfamily A, member 13 /// killer cell lectin-like receptor, subfamily A, member 4 /// killer cell lectin-like receptor, subfamily A, member 7 /// killer cell lectin-like receptor, subfamily A, member 15 /// killer cell lectin-like receptor subfamily A, member 23 /// killer cell lectin-like receptor subfamily A, member 20 /// killer cell lectin-like receptor

subfamily A, member 22 /// killer cell lectin-like receptor, subfamily A, member 18 /// similar to killer
 cell lectin-like receptor subfamily A, member 13" Klra1 /// Klra12 /// Klra13 /// Klra4 /// Klra7 ///
 Klra15 /// Klra23 /// Klra20 /// Klra22 /// Klra18 /// LOC673407
 1451665_a_at 1.09679 "adaptor-related protein complex AP-4, sigma 1" Ap4s1
 1451666_at 1.26345 ATP citrate lyase Acly
 1451667_at 0.95112 RIKEN cDNA C530043G21 gene C530043G21Rik
 1451668_at 1.02240 RIKEN cDNA C530043G21 gene C530043G21Rik
 1451669_at 0.96675 "protein phosphatase 1B, magnesium dependent, beta isoform" Ppm1b
 1451670_at 0.84685 RIKEN cDNA 1810048P08 gene 1810048P08Rik
 1451671_at 0.91862 golgi reassembly stacking protein 1 Gorasp1
 1451672_at 0.96173 G protein-coupled receptor kinase 6 Gprk6
 1451673_at 1.02152 "CD8 antigen, alpha chain /// similar to T-cell surface glycoprotein CD8 alpha
 chain precursor (T-cell surface glycoprotein Lyt-2) /// similar to T-cell surface glycoprotein CD8 alpha
 chain precursor (T-cell surface glycoprotein Lyt-2) (CD8a antigen)" Cd8a /// LOC636147 ///
 LOC669166
 1451674_at 0.67673 "solute carrier family 12, member 5" Slc12a5
 1451675_a_at 0.99850 "aminolevulinic acid synthase 2, erythroid" Alas2
 1451676_at 0.98600 Dr1 associated protein 1 (negative cofactor 2 alpha) Drap1
 1451677_at 1.09621 nuclear prelamin A recognition factor Narf
 1451678_at 0.90120 nuclear prelamin A recognition factor Narf
 1451679_at 0.97817 RIKEN cDNA 6530401D17 gene 6530401D17Rik
 1451680_at 0.98936 sulfiredoxin 1 homolog (S. cerevisiae) Srxn1
 1451681_at 0.98531 cDNA sequence BC089597 BC089597
 1451682_at 1.01301 zinc finger protein 142 Zfp142
 1451683_x_at 1.84670 "similar to H-2 class I histocompatibility antigen, L-D alpha chain precursor"
 LOC547343
 1451684_a_at 1.09090 bicaudal D homolog 1 (Drosophila) Bcd1
 1451685_at 1.07418 myeloid/lymphoid or mixed lineage-leukemia translocation to 6 homolog
 (Drosophila) /// similar to myeloid/lymphoid or mixed lineage-leukemia translocation to 6 homolog ///
 similar to myeloid/lymphoid or mixed lineage-leukemia translocation to 6 homolog Mllt6 ///
 LOC639490 /// LOC669933
 1451686_x_at 0.89754 amelogenin X chromosome Amelx
 1451687_a_at 0.85291 transcription factor 2 Tcf2
 1451688_s_at 0.78846 calcium activated nucleotidase 1 Cant1
 1451689_a_at 0.97988 SRY-box containing gene 10 Sox10
 1451690_a_at 1.05018 poliovirus receptor-related 4 Pvr14
 1451691_at 1.02368 endothelin receptor type A Ednra
 1451692_at 1.17457 RIKEN cDNA 2410015B03 gene 2410015B03Rik
 1451693_a_at 1.14063 fibroblast growth factor 12 Fgf12
 1451694_at 0.98478 "tubulin tyrosine ligase-like family, member 3" Ttl13
 1451695_a_at 1.10320 glutathione peroxidase 4 Gpx4
 1451696_at 1.02087 zinc finger protein 64 Zfp64
 1451697_a_at 0.83784 Ellis van Creveld gene homolog (human) Evc

1451698_at 0.99683 proline/serine-rich coiled-coil 1 Psrc1
 1451699_at 1.14686 similar to putative retrovirus-related gag protein /// hypothetical protein
 LOC668631 /// similar to putative retrovirus-related gag protein LOC668468 /// LOC668631 ///
 LOC675654
 1451700_a_at 1.07691 RIKEN cDNA 1110007L15 gene 1110007L15Rik
 1451701_x_at 0.84713 claudin 3 Cldn3
 1451702_at 1.00057 CKLF-like MARVEL transmembrane domain containing 7 Cmtm7
 1451703_s_at 1.08295 --- ---
 1451704_at 1.01938 androgen binding protein delta Abpd
 1451705_a_at 1.04464 "opioid receptor, mu 1" Oprm1
 1451706_a_at 1.09371 "gamma-aminobutyric acid (GABA-A) receptor, subunit alpha 6" Gabra6
 1451707_s_at 0.80073 "solute carrier family 41, member 3" Slc41a3
 1451708_at 1.02379 G protein-coupled receptor 33 Gpr33
 1451709_at 1.08987 "opioid receptor, mu 1" Oprm1
 1451710_at 0.91259 osteoclast associated receptor Oscar
 1451711_at 1.00726 wingless-type MMTV integration site 9B Wnt9b
 1451712_at 1.04301 RIKEN cDNA 8030411F24 gene 8030411F24Rik
 1451713_a_at 0.91293 "Fc receptor, IgE, low affinity II, alpha polypeptide" Fcer2a
 1451714_a_at 1.07555 mitogen activated protein kinase kinase 3 Map2k3
 1451715_at 0.99412 "v-maf musculoaponeurotic fibrosarcoma oncogene family, protein B
 (avian)" Mafb
 1451716_at 0.94228 "v-maf musculoaponeurotic fibrosarcoma oncogene family, protein B
 (avian)" Mafb
 1451717_s_at 1.12668 SUMO/sentrin specific peptidase 2 Senp2
 1451718_at 1.14142 proteolipid protein (myelin) 1 Plp1
 1451719_at 0.95831 "cofactor required for Sp1 transcriptional activation, subunit 6" Crsp6
 1451720_at 0.87646 vacuolar protein sorting 39 (yeast) Vps39
 1451721_a_at 1.11054 "histocompatibility 2, class II antigen A, beta 1" H2-Ab1
 1451722_s_at 0.99549 SET and MYND domain containing 5 Smyd5
 1451723_at 0.89279 "CCR4-NOT transcription complex, subunit 6-like" Cnot6l
 1451724_at 0.90571 ankyrin repeat and MYND domain containing 2 Ankmy2
 1451725_a_at 1.03957 "proteasome (prosome, macropain) 26S subunit, non-ATPase, 4" Psmd4
 1451726_at 0.96577 myotubularin related protein 6 Mtmr6
 1451727_at 0.93820 "DNA segment, Chr 11, ERATO Doi 730, expressed" D11Ert730e
 1451728_at 0.97455 WD repeat domain 13 Wdr13
 1451729_at 0.98612 "bone morphogenetic protein receptor, type 1A" Bmpr1a
 1451730_at 0.96742 zinc finger protein 62 Zfp62
 1451731_at 0.94723 "ATP-binding cassette, sub-family A (ABC1), member 3" Abca3
 1451732_at 1.03095 PRP4 pre-mRNA processing factor 4 homolog B (yeast) Prpf4b
 1451733_at 1.03583 "glucosaminyl (N-acetyl) transferase 2, I-branching enzyme" Gcnt2
 1451734_a_at 0.69918 drebrin 1 Dbn1
 1451735_at 1.01038 ADP-ribosylation factor related protein 1 Arfrp1
 1451736_a_at 0.95050 mitogen activated protein kinase kinase 7 Map2k7

1451737_at 1.10482 "phosphatidylinositol 3-kinase, regulatory subunit, polypeptide 1 (p85 alpha)"
 Pik3r1
 1451738_at 0.91906 O-linked N-acetylglucosamine (GlcNAc) transferase (UDP-N-
 acetylglucosamine:polypeptide-N-acetylglucosaminyl transferase) Ogt
 1451739_at 1.37182 Kruppel-like factor 5 Klf5
 1451740_at 0.87694 polyadenylate binding protein-interacting protein 1 Paip1
 1451741_a_at 1.10238 cyclin-dependent kinase 7 (homolog of Xenopus MO15 cdk-activating
 kinase) Cdk7
 1451742_a_at 0.98888 UDP-glucose pyrophosphorylase 2 Ugp2
 1451743_at 0.91095 "DNA segment, Chr 19, Wayne State University 162, expressed" D19Wsu162e
 1451744_a_at 0.92478 "zinc binding alcohol dehydrogenase, domain containing 1" Zadh1
 1451745_a_at 1.04764 "zinc finger, HIT domain containing 1" Znhit1
 1451746_a_at 0.96279 autophagy-related 12 (yeast) Atg12
 1451747_a_at 0.97068 autophagy-related 12 (yeast) Atg12
 1451748_a_at 1.00452 SET domain containing 3 Setd3
 1451749_at 0.96886 interleukin-1 receptor-associated kinase 4 Irak4
 1451750_at 0.97161 --- ---
 1451751_at 0.86643 DNA-damage-inducible transcript 4-like Ddit4l
 1451752_at 0.88867 forkhead box K1 Foxk1
 1451753_at 0.96685 plexin A2 Plxna2
 1451754_a_at 0.97890 WD repeat domain 45 Wdr45
 1451755_a_at 1.05667 apolipoprotein B editing complex 1 Apobec1
 1451756_at 0.93875 FMS-like tyrosine kinase 1 Flt1
 1451757_at 0.95150 cDNA sequence BC003883 BC003883
 1451758_at 1.17673 laminin gamma 3 Lamc3
 1451759_at 0.97380 mannan-binding lectin serine peptidase 2 Masp2
 1451760_s_at 1.05863 RIKEN cDNA 2010001E11 gene /// expressed sequence AI317395
 2010001E11Rik /// AI317395
 1451761_at 1.01921 homeo box B4 Hoxb4
 1451762_a_at 0.90425 kinesin family member 1B Kif1b
 1451763_at 1.11574 cyclic nucleotide gated channel alpha 1 Cnga1
 1451764_at 1.14352 MARVEL (membrane-associating) domain containing 3 Marveld3
 1451765_a_at 0.85112 ectonucleoside triphosphate diphosphohydrolase 5 Entpd5
 1451766_at 1.25566 cadherin 11 Cdh11
 1451767_at 1.07609 neutrophil cytosolic factor 1 Ncf1
 1451768_a_at 1.08596 "solute carrier family 20, member 2" Slc20a2
 1451769_s_at 0.89671 "protocadherin alpha 4 /// protocadherin alpha 6 /// protocadherin alpha 7 ///
 protocadherin alpha 5 /// protocadherin alpha 11 /// protocadherin alpha 10 /// protocadherin alpha 1 ///
 protocadherin alpha 9 /// protocadherin alpha cluster /// protocadherin alpha 3 /// protocadherin alpha
 12 /// protocadherin alpha 2 /// protocadherin alpha 8 /// protocadherin alpha subfamily C, 1 ///
 protocadherin alpha subfamily C, 2" Pcdha4 /// Pcdha6 /// Pcdha7 /// Pcdha5 /// Pcdha11 ///
 Pcdha10 /// Pcdha1 /// Pcdha9 /// Pcdha@ /// Pcdha3 /// Pcdha12 /// Pcdha2 /// Pcdha8 /// Pcdhac1 ///
 Pcdhac2

1451770_s_at 1.12251 DEAH (Asp-Glu-Ala-His) box polypeptide 9 Dhx9
 1451771_at 0.95487 two pore channel 1 Tpcn1
 1451772_at 1.04602 two pore channel 1 Tpcn1
 1451773_s_at 1.00133 polymerase (RNA) III (DNA directed) polypeptide F Polr3f
 1451774_at 0.89036 similar to keratin associated protein 4-10; keratin associated protein 4.10
 LOC435285
 1451775_s_at 1.06318 "interleukin 13 receptor, alpha 1" Il13ra1
 1451776_s_at 0.94667 homeobox only domain Hod
 1451777_at 0.96784 cDNA sequence BC013672 BC013672
 1451778_at 0.92030 CREB regulated transcription coactivator 3 Crtc3
 1451779_at 1.12797 cDNA sequence BC027061 BC027061
 1451780_at 1.07612 B-cell linker Blnk
 1451781_at 1.00454 "nuclear factor of activated T-cells, cytoplasmic, calcineurin-dependent 2
 interacting protein" Nfatc2ip
 1451782_a_at 0.91778 "solute carrier family 29 (nucleoside transporters), member 1" Slc29a1
 1451783_a_at 0.93543 kinesin-associated protein 3 Kifap3
 1451784_x_at 2.07227 "histocompatibility 2, D region locus 1 /// histocompatibility 2, D region ///
 similar to H-2 class I histocompatibility antigen, L-D alpha chain precursor /// similar to H-2 class I
 histocompatibility antigen, D-B alpha chain precursor (H-2D(B))" H2-D1 /// H2-L ///
 LOC547343 /// LOC636948
 1451785_at 0.96188 retinitis pigmentosa GTPase regulator interacting protein 1 Rpgrip1
 1451786_at 0.95730 mucin 20 Muc20
 1451787_at 0.78785 "cytochrome P450, family 2, subfamily b, polypeptide 10" Cyp2b10
 1451788_at 0.94880 coagulation factor XI F11
 1451789_a_at 0.98648 receptor-like tyrosine kinase Ryk
 1451790_a_at 0.98239 tissue factor pathway inhibitor Tfpi
 1451791_at 1.11295 tissue factor pathway inhibitor Tfpi
 1451792_a_at 0.79263 "praja1, RING-H2 motif containing" Pja1
 1451793_at 0.70895 kelch-like 24 (Drosophila) Klh24
 1451794_at 0.87481 transmembrane and coiled coil domains 3 Tmcc3
 1451795_at 0.87832 target of myb1-like 2 (chicken) Tom112
 1451796_s_at 0.94966 histidine decarboxylase Hdc
 1451797_at 1.24523 "Smg-5 homolog, nonsense mediated mRNA decay factor (C. elegans)"
 Smg5
 1451798_at 1.09355 interleukin 1 receptor antagonist Il1rn
 1451799_at 1.26857 coiled-coil domain containing 25 Ccdc25
 1451800_at 0.95349 GRIP and coiled-coil domain containing 2 Gcc2
 1451801_at 0.96860 triadin Trdn
 1451802_at 1.10581 mast cell protease 6 Mcpt6
 1451803_a_at 1.11141 vascular endothelial growth factor B Vegfb
 1451804_a_at 1.07873 leucine rich repeat containing 16 Lrrc16
 1451805_at 0.90510 pleckstrin homology domain interacting protein Phip
 1451806_at 1.09437 formin binding protein 1 Fbnp1

1451807_at 1.05129 "nuclear receptor subfamily 1, group I, member 2" Nr1i2
 1451808_at 1.01843 "potassium inwardly-rectifying channel, subfamily J, member 4" Kcnj4
 1451809_s_at 0.96383 RWD domain containing 3 Rwdd3
 1451810_at 0.81588 open reading frame 63 ORF63
 1451811_at 1.00131 "calcium channel, voltage-dependent, gamma subunit 6" Cacng6
 1451812_at 1.06742 a disintegrin and metallopeptidase domain 22 Adam22
 1451813_at 0.82590 "opioid receptor, kappa 1" Oprk1
 1451814_a_at 1.18328 "HIV-1 tat interactive protein 2, homolog (human)" Htatip2
 1451815_at 1.03312 "UDP-GlcNAc:betaGal beta-1,3-N-acetylglucosaminyltransferase 4" B3gnt4
 1451816_at 0.84592 zinc finger protein 451 Zfp451
 1451817_at 1.01474 suppressor of variegation 4-20 homolog 1 (Drosophila) Suv420h1
 1451818_at 1.02534 mindbomb homolog 1 (Drosophila) Mib1
 1451819_at 1.13771 "zinc finger, SWIM domain containing 6" Zswim6
 1451820_at 0.88704 "DIRAS family, GTP-binding RAS-like 1" Diras1
 1451821_a_at 1.08664 nuclear antigen Sp100 Sp100
 1451822_a_at 0.99029 secernin 2 Scrn2
 1451823_at 1.25657 chloride channel calcium activated 4 Clca4
 1451824_at 0.96252 "butyrophilin, subfamily 1, member A1" Btn1a1
 1451825_a_at 0.93888 "coatomer protein complex, subunit zeta 1" Copz1
 1451826_at 0.90972 calcium binding protein 5 Cabp5
 1451827_a_at 0.82089 NADPH oxidase 4 Nox4
 1451828_a_at 1.47223 acyl-CoA synthetase long-chain family member 4 Acsl4
 1451829_a_at 0.94879 neurofibromatosis 2 Nf2
 1451830_a_at 1.32393 spectrin beta 2 Spnb2
 1451831_at 0.93076 oxysterol binding protein-like 6 Osbpl6
 1451832_at 1.20287 --- ---
 1451833_a_at 1.01024 "SET domain, bifurcated 1" Setdb1
 1451834_at 0.95685 "calcium channel, voltage-dependent, beta 1 subunit" Cacnb1
 1451835_at 0.93567 SRY-box containing gene 21 Sox21
 1451836_at 1.19677 cDNA sequence BC022651 BC022651
 1451837_at 0.91766 "adaptor-related protein complex 3, beta 2 subunit" Ap3b2
 1451838_a_at 0.94880 membrane targeting (tandem) C2 domain containing 1 Mtac2d1
 1451839_a_at 1.08239 phosphodiesterase 7A Pde7a
 1451840_at 0.77624 Kv channel interacting protein 4 Kcnip4
 1451841_a_at 0.83575 nuclear receptor co-repressor 2 Ncor2
 1451842_a_at 1.03456 "cholinergic receptor, nicotinic, beta polypeptide 3" Chrn3
 1451843_a_at 0.96757 "glycoprotein galactosyltransferase alpha 1, 3" Ggta1
 1451844_at 1.07499 prolactin receptor Prlr
 1451845_a_at 1.07758 peptidyl-tRNA hydrolase 2 Ptrh2
 1451846_at 0.86593 nebulin Nebl
 1451847_s_at 0.84256 AT rich interactive domain 4B (Rbp1 like) Arid4b
 1451848_a_at 0.95962 "crystallin, zeta" Cryz
 1451849_a_at 1.00466 lamin B2 Lmnb2

1451850_at 0.91406 prolactin receptor Prlr
1451851_a_at 1.00509 casein kappa Csn3
1451852_at 0.88330 "2-cell-stage, variable group, member 3" Tcstv3
1451853_at 0.98447 Fukuyama type congenital muscular dystrophy homolog (human) Fcmd
1451854_a_at 1.06513 shroom family member 3 Shroom3
1451855_at 1.04804 B-cell CLL/lymphoma 7A Bcl7a
1451856_at 0.84300 trinucleotide repeat containing 15 Tnrc15
1451857_a_at 0.92417 RIKEN cDNA 5730593N15 gene 5730593N15Rik
1451858_at 0.91614 "MAS-related GPR, member A2 /// similar to MAS-related GPR, member A2"
Mrgpra2 /// LOC668727
1451859_at 1.08836 keratin associated protein 6-1 Krtap6-1
1451860_a_at 1.04277 tripartite motif protein 30 Trim30
1451861_at 0.93087 RIKEN cDNA 4930519N16 gene 4930519N16Rik
1451862_a_at 0.89709 perforin 1 (pore forming protein) Prf1
1451863_at 0.94315 urotensin 2 receptor Uts2r
1451864_at 0.93450 "calcium channel, voltage-dependent, gamma subunit 8" Cacng8
1451865_at 1.05457 coxsackievirus and adenovirus receptor Cxadr
1451866_a_at 0.91284 hepatocyte growth factor Hgf
1451867_x_at 0.96780 Rho GTPase activating protein 6 Arhgap6
1451868_at 1.02485 "potassium inwardly-rectifying channel, subfamily J, member 6" Kcnj6
1451869_at 1.04805 "ATP-binding cassette, sub-family A (ABC1), member 3" Abca3
1451870_a_at 1.33491 bromodomain containing 4 Brd4
1451871_a_at 0.63530 growth hormone receptor Ghr
1451872_a_at 0.95052 neuralized-like homolog (Drosophila) Neurl
1451873_a_at 0.87896 MAD homolog 5 (Drosophila) Smad5
1451874_at 0.93914 --- ---
1451875_at 0.92436 cDNA sequence AF366264 AF366264
1451876_a_at 1.08084 transformation related protein 63 Trp63
1451877_at 1.00084 ectodysplasin-A Eda
1451878_a_at 0.89496 junction-mediating and regulatory protein Jmy
1451879_a_at 1.09068 "solute carrier family 6 (neurotransmitter transporter), member 18" Slc6a18
1451880_at 0.91336 hypothetical LOC638575 LOC638575
1451881_at 0.89819 --- ---
1451882_a_at 0.98363 fibroblast growth factor 8 Fgf8
1451883_at 1.07470 hypothetical LOC620316 LOC620316
1451884_a_at 1.05571 "LSM2 homolog, U6 small nuclear RNA associated (S. cerevisiae)" Lsm2
1451885_at 0.95738 metallothionein 1K LOC574530
1451886_at 0.92556 SPEG complex locus Spieg
1451887_at 1.02995 LPS-responsive beige-like anchor Lrba
1451888_a_at 1.00865 odd Oz/ten-m homolog 4 (Drosophila) Odz4
1451889_at 0.97582 Notch gene homolog 2 (Drosophila) Notch2
1451890_at 0.76323 kinesin family member 13A Kif13a
1451891_a_at 0.91600 dysferlin Dysf

1451892_at 0.88287 klotho Kl
 1451893_s_at 0.90640 "membrane associated guanylate kinase, WW and PDZ domain containing 1"
 Magi1
 1451894_a_at 0.91393 "sodium channel, voltage-gated, type VIII, alpha" Scn8a
 1451895_a_at 1.38554 24-dehydrocholesterol reductase Dhcr24
 1451896_a_at 1.04282 calcium homeostasis endoplasmic reticulum protein Cherp
 1451897_a_at 0.86849 neighbor of Brca1 gene 1 Nbr1
 1451898_a_at 0.88162 "sema domain, transmembrane domain (TM), and cytoplasmic domain,
 (semaphorin) 6C" Sema6c
 1451899_a_at 0.94951 general transcription factor II I repeat domain-containing 1 Gtf2ird1
 1451900_at 0.81770 mitofusin 2 Mfn2
 1451901_at 1.14298 "SWI/SNF related, matrix associated, actin dependent regulator of chromatin,
 subfamily a, member 3" Smarca3
 1451902_at 1.06781 zinc finger protein 758 Zfp758
 1451903_at 1.00999 kynureninase (L-kynurenine hydrolase) Kynu
 1451904_a_at 0.97619 a disintegrin and metallopeptidase domain 33 Adam33
 1451905_a_at 1.04516 myxovirus (influenza virus) resistance 1 Mx1
 1451906_at 1.04764 UBX domain containing 3 Ubx3
 1451907_a_at 1.01678 "pleckstrin homology domain containing, family A (phosphoinositide binding
 specific) member 4" Plekha4
 1451908_a_at 1.02590 SEC14-like 1 (S. cerevisiae) Sec14l1
 1451909_a_at 1.06110 PRP4 pre-mRNA processing factor 4 homolog B (yeast) Prpf4b
 1451910_a_at 0.88671 CD6 antigen Cd6
 1451911_a_at 0.95387 angiotensin I converting enzyme (peptidyl-dipeptidase A) 1 Ace
 1451912_a_at 0.69274 fibroblast growth factor receptor-like 1 Fgfr11
 1451913_a_at 0.94365 hypoxia up-regulated 1 Hyou1
 1451914_a_at 1.07028 adducin 2 (beta) Add2
 1451915_at 0.99636 chaperonin subunit 3 (gamma) Cct3
 1451916_s_at 1.00585 tripartite motif protein 15 Trim15
 1451917_a_at 1.00574 doublecortin and calcium/calmodulin-dependent protein kinase-like 1
 Dcamk11
 1451918_a_at 0.90864 "loss of heterozygosity, 12, chromosomal region 1 homolog (human)"
 Loh12cr1
 1451919_a_at 1.00333 myeloproliferative leukemia virus oncogene Mpl
 1451920_a_at 0.90975 replication factor C 1 Recc1
 1451921_a_at 1.10897 nuclear factor of activated T-cells 5 Nfat5
 1451922_at 0.96520 "lectin, mannose-binding 1 like" Lman11
 1451923_at 0.90997 RIKEN cDNA 3222401M22 gene 3222401M22Rik
 1451924_a_at 0.69794 endothelin 1 Edn1
 1451925_at 1.13264 ectodysplasin-A Eda
 1451926_at 1.09355 "MAS-related GPR, member A4" Mrgpra4
 1451927_a_at 0.96703 mitogen activated protein kinase 14 Mapk14
 1451928_a_at 1.09670 RAD18 homolog (S. cerevisiae) Rad18

1451929_a_at 0.97274 vaccinia related kinase 2 Vrk2
 1451930_at 2.57004 Ephrin A5 Efna5
 1451931_x_at 1.85288 "histocompatibility 2, D region" H2-L
 1451932_a_at 1.02220 ADAMTS-like 4 Adamtsl4
 1451933_a_at 0.95357 cathepsin 7 Cts7
 1451934_at 0.91131 "histocompatibility 2, D region locus 1" H2-D1
 1451935_a_at 0.98430 "serine protease inhibitor, Kunitz type 2" Spint2
 1451936_a_at 0.98344 thioredoxin reductase 2 Txnrd2
 1451937_at 0.99059 "glycine receptor, alpha 4 subunit" Glra4
 1451938_a_at 1.02846 "syntrophin, basic 1" Sntb1
 1451939_a_at 1.03892 sushi-repeat-containing protein SrpX
 1451940_x_at 0.94401 triadin Trdn
 1451941_a_at 1.10402 "Fc receptor, IgG, low affinity IIb" Fcgr2b
 1451942_x_at 0.92960 "killer cell lectin-like receptor subfamily A, member 20" Klra20
 1451943_a_at 0.87090 "protein phosphatase 1A, magnesium dependent, alpha isoform" Ppm1a
 1451944_a_at 0.89884 "tumor necrosis factor (ligand) superfamily, member 11" Tnfsf11
 1451945_at 0.96152 --- ---
 1451946_a_at 0.91468 calcium-binding tyrosine-(Y)-phosphorylation regulated (fibrousheathin 2)
 Cabyr
 1451947_at 0.96861 immunoglobulin heavy chain (V7183 family) Igh-V7183
 1451948_at 0.95514 "gene model 1409, (NCBI)" Gm1409
 1451949_at 1.01825 immunoglobulin heavy chain 4 (serum IgG1) /// immunoglobulin heavy chain 6
 (heavy chain of IgM) /// immunoglobulin heavy chain (V7183 family) /// immunoglobulin heavy chain
 (J558 family) /// similar to immunoglobulin heavy chain variable region /// immunoglobulin heavy chain
 1a (serum IgG2a) /// Immunoglobulin heavy chain (gamma polypeptide) /// Ig heavy chain V region ///
 similar to immunoglobulin mu-chain Igh-4 /// Igh-6 /// Igh-V7183 /// Igh-VJ558 /// LOC238447 ///
 Igh-1a /// Ighg /// LOC380804 /// LOC544903
 1451950_a_at 1.04124 CD80 antigen Cd80
 1451951_at 0.98293 Hypothetical protein LOC625794 LOC625794
 1451952_at 1.18057 prokineticin 2 Prok2
 1451953_at 0.96551 submaxillary gland androgen regulated protein 2 Smr2
 1451954_at 0.92453 --- ---
 1451955_a_at 1.03323 "calcium channel, voltage-dependent, alpha 2/delta subunit 2 /// similar to
 calcium channel, voltage-dependent, alpha 2/delta subunit 2" Cacna2d2 /// LOC669637
 1451956_a_at 1.23691 "opioid receptor, sigma 1" Oprs1
 1451957_at 0.83229 "interleukin 1 family, member 10" Il1f10
 1451958_at 0.93913 similar to Ig heavy chain V region 93G7 precursor /// similar to Ig heavy chain
 V region B1-8/186-2 precursor /// similar to Ig heavy chain V region 3 precursor /// similar to
 monoclonal antibody heavy chain /// similar to Ig heavy chain V region VH558 A1/A4 precursor ///
 similar to monoclonal antibody heavy chain /// similar to Ig heavy chain V region 102 precursor ///
 similar to Ig heavy chain V region B1-8/186-2 precursor /// similar to Ig heavy chain V region VH558
 A1/A4 precursor /// similar to Ig heavy chain V region VH558 A1/A4 precursor /// similar to Ig heavy
 chain V region 3 precursor /// similar to Ig heavy chain V region B1-8/186-2 precursor /// similar to Ig

heavy chain V region VH558 B4 precursor /// similar to Ig heavy chain V region 23 precursor /// similar to Ig heavy chain V region 108A precursor /// similar to Ig heavy chain V region 23 precursor /// similar to Ig heavy chain V region 3 precursor /// similar to Ig heavy chain V region 102 precursor /// similar to Ig heavy chain V region 3 precursor /// similar to Ig heavy chain V region 3 precursor /// similar to Ig heavy chain V region 102 precursor /// similar to Ig heavy chain V region 23 precursor /// similar to Ig heavy chain V region 3 precursor /// similar to Ig heavy chain V region 3 precursor /// similar to Ig heavy chain V region VH558 A1/A4 precursor /// similar to Ig heavy chain V region 145 precursor /// similar to Ig heavy chain V region 93G7 precursor /// similar to Ig heavy chain V region 102 precursor /// similar to Ig heavy chain V region VH558 A1/A4 precursor /// similar to Ig heavy chain V region 3 precursor /// similar to Ig heavy chain V region 93G7 precursor /// similar to Ig heavy chain V region 93G7 precursor /// similar to Ig heavy chain V region 102 precursor LOC211331 /// LOC380823 /// LOC382696 /// VH186.2-D-J-C mu /// LOC434638 /// LOC435333 /// LOC619833 /// LOC619916 /// LOC619994 /// LOC622728 /// LOC629941 /// LOC630549 /// LOC631531 /// LOC633774 /// LOC634123 /// LOC636044 /// LOC636134 /// LOC636244 /// LOC637785 /// LOC637794 /// LOC640207 /// LOC640979 /// LOC641003 /// LOC641114 /// LOC668478 /// LOC668549 /// LOC668577 /// LOC673286 /// LOC673498 /// LOC674018 /// LOC674023 /// LOC674472 /// LOC675659

1451959_a_at 0.94555 vascular endothelial growth factor A Vegfa

1451960_a_at 0.95371 "REV3-like, catalytic subunit of DNA polymerase zeta RAD54 like (S. cerevisiae)" Rev3l

1451961_a_at 0.90706 myelin basic protein Mbp

1451962_at 0.99244 immunoglobulin kappa light chain 17-1A LOC213481

1451963_at 1.03694 Anti-HLA class II antibody gamma F3.3 heavy chain variable region ---

1451964_at 0.93434 melanoma inhibitory activity 2 Mia2

1451965_at 0.92368 Anti-human CD37 antibody WR17 kappa light chain variable region ---

1451966_at 1.02915 melanocortin 2 receptor accessory protein Mrap

1451967_x_at 1.02230 karyopherin (importin) beta 1 Kpnb1

1451968_at 1.00474 X-ray repair complementing defective repair in Chinese hamster cells 5 Xrcc5

1451969_s_at 1.25918 "poly (ADP-ribose) polymerase family, member 3" Parp3

1451970_at 0.96581 RIKEN cDNA E330036I19 gene E330036I19Rik

1451971_at 1.00365 cullin 4A Cul4a

1451972_at 0.78372 glucocorticoid induced transcript 1 Glcci1

1451973_at 0.92362 ubiquitin-activating enzyme E1-domain containing 1 Ube1dc1

1451974_at 0.91286 oxysterol binding protein-like 2 Osbpl2

1451975_at 0.96843 RIKEN cDNA 2810453I06 gene 2810453I06Rik

1451976_s_at 1.01693 CKLF-like MARVEL transmembrane domain containing 2A Cmtm2a

1451977_at 0.99692 dual-specificity tyrosine-(Y)-phosphorylation regulated kinase 1a Dyrk1a

1451978_at 1.08321 lysyl oxidase-like 1 Loxl1

1451979_at 0.96417 v-Ki-ras2 Kirsten rat sarcoma viral oncogene homolog Kras

1451980_at 0.90471 CAS1 domain containing 1 Casd1

1451981_at 1.00179 gene trap ROSA b-geo 22 Gtrgeo22

1451982_at 1.00019 mitogen activated protein kinase kinase 4 Map2k4

1451983_at 0.90312 Iroquois related homeobox 1 (Drosophila) Irx1

1451984_at	0.92822	heterogeneous nuclear ribonucleoprotein U-like 1	Hnrp11
1451985_at	0.94065	leucine-rich repeat kinase 1	Lrrk1
1451986_s_at	0.98243	leucine-rich repeat kinase 1	Lrrk1
1451987_at	0.95085	"arrestin, beta 2"	Arrb2
1451988_s_at	0.96996	chromatin modifying protein 4B	Chmp4b
1451989_a_at	0.87802	"microtubule-associated protein, RP/EB family, member 2"	Mapre2
1451990_at	0.80704	"microtubule-associated protein, RP/EB family, member 2"	Mapre2
1451991_at	0.76073	Eph receptor A7	Epha7
1451992_at	1.22754	"adrenergic receptor kinase, beta 1"	Adrbk1
1451993_at	0.97409	RIKEN cDNA 9130404D08 gene	9130404D08Rik
1451994_s_at	0.97726	RIKEN cDNA 3930401K13 gene	3930401K13Rik
1451995_at	0.93390	"TAF11 RNA polymerase II, TATA box binding protein (TBP)-associated factor"	Taf11
1451996_at	0.93773	TM2 domain containing 1	Tm2d1
1451997_at	0.96245	zinc finger protein 426	Zfp426
1451998_at	1.01796	"taspase, threonine aspartase 1"	Tasp1
1451999_at	1.02645	LIM domain binding 3	Ldb3
1452000_s_at	1.06338	seryl-aminoacyl-tRNA synthetase	Sars
1452001_at	0.84157	"nuclear factor, erythroid derived 2"	Nfe2
1452002_at	0.87696	zinc finger protein 508	Zfp508
1452003_at	0.99875	RNA binding motif protein 14	Rbm14
1452004_at	0.82735	"calcitonin/calcitonin-related polypeptide, alpha"	Calca
1452005_at	0.72705	dihydrolipoamide S-acetyltransferase (E2 component of pyruvate dehydrogenase complex)	Dlat
1452006_x_at	0.91499	zinc finger and BTB domain containing 8 opposite strand	Zbtb8os
1452007_at	0.82166	synaptobrevin like 1	Sybl1
1452008_at	1.43081	RIKEN cDNA 9130422G05 gene	9130422G05Rik
1452009_at	1.08844	RIKEN cDNA 9130422G05 gene	9130422G05Rik
1452010_at	1.00821	"cholinergic receptor, nicotinic, alpha polypeptide 3"	Chrna3
1452011_a_at	1.01050	UDP-glucuronate decarboxylase 1	Uxs1
1452012_a_at	1.23243	exosome component 1	Exosc1
1452013_at	1.05489	"ATPase, class V, type 10A"	Atp10a
1452014_a_at	0.89024	insulin-like growth factor 1	Igf1
1452015_at	1.02262	RIKEN cDNA 6330416G13 gene	6330416G13Rik
1452016_at	0.86004	arachidonate 5-lipoxygenase activating protein	Alox5ap
1452017_at	0.90529	SRY-box containing gene 15	Sox15
1452018_at	0.84681	"NK2 transcription factor related, locus 6 (Drosophila)"	Nkx2-6
1452019_at	0.92338	cysteine and tyrosine-rich protein 1	Cyrr1
1452020_a_at	1.00605	Cd27 binding protein (Hindu God of destruction)	Siva
1452021_a_at	0.89577	hairy and enhancer of split 6 (Drosophila)	Hes6
1452022_at	0.95540	"brain and acute leukemia, cytoplasmic"	Baalc
1452023_at	0.91111	ankyrin repeat and SOCS box-containing 16	Asb16
1452024_a_at	0.92941	LIM domain binding 1	Ldb1

1452025_a_at 0.76510 zinc finger protein 2 Zfp2
1452026_a_at 0.97742 "phospholipase A2, group XIIA" Pla2g12a
1452027_a_at 0.97944 transformation related protein 63 Trp63
1452028_a_at 1.00965 cadherin 23 (otocadherin) Cdh23
1452029_a_at 0.86421 purine-rich element binding protein G Purg
1452030_a_at 0.99066 heterogeneous nuclear ribonucleoprotein R Hnrpr
1452031_at 1.03993 "solute carrier family 1 (glial high affinity glutamate transporter), member 3" Slc1a3
1452032_at 0.93258 "protein kinase, cAMP dependent regulatory, type I, alpha" Prkar1a
1452033_at 0.93473 GLE1 RNA export mediator-like (yeast) Gle11
1452034_at 0.95419 prolyl endopeptidase-like Prepl
1452035_at 0.85864 "procollagen, type IV, alpha 1" Col4a1
1452036_a_at 1.04390 thymopoietin Tmpo
1452037_at 1.01238 mannoside acetylglucosaminyltransferase 2 Mgat2
1452038_at 0.97069 "capping protein (actin filament) muscle Z-line, alpha 1 /// similar to capping protein (actin filament) muscle Z-line, alpha 1" Capza1 /// LOC672255
1452039_a_at 0.94464 Brca1 associated protein 1 Bap1
1452040_a_at 1.01727 cell division cycle associated 3 Cdca3
1452041_at 0.90876 kelch-like 26 (Drosophila) Klhl26
1452042_a_at 0.84356 transmembrane protein 144 Tmem144
1452043_at 1.02736 RIKEN cDNA 2310011J03 gene 2310011J03Rik
1452044_at 0.98691 "actin related protein 2/3 complex, subunit 5-like" Arpc51
1452045_at 0.94498 zinc finger protein 281 Zfp281
1452046_a_at 0.96260 "protein phosphatase 1, catalytic subunit, gamma isoform /// similar to protein phosphatase 1 gamma2" Ppp1cc /// LOC434233
1452047_at 1.09127 calyculin binding protein Cacybp
1452048_at 1.09830 mitochondrial ribosomal protein L12 Mrpl12
1452049_at 1.01206 ribosomal protein L7-like 1 Rpl7l1
1452050_at 0.89059 calcium/calmodulin-dependent protein kinase ID Camk1d
1452051_at 0.94065 ARP3 actin-related protein 3 homolog (yeast) Actr3
1452052_s_at 1.09613 "eukaryotic translation initiation factor 3, subunit 1 alpha" Eif3s1
1452053_a_at 0.91006 transmembrane protein 33 Tmem33
1452054_at 0.87541 ubiquitin-conjugating enzyme E2W (putative) Ube2w
1452055_at 0.91334 "CTD (carboxy-terminal domain, RNA polymerase II, polypeptide A) small phosphatase 1" Ctdsp1
1452056_s_at 1.04109 "protein phosphatase 3, catalytic subunit, alpha isoform" Ppp3ca
1452057_at 1.05898 ARP1 actin-related protein 1 homolog B (yeast) Actr1b
1452058_a_at 0.91547 ring finger protein 11 Rnf11
1452059_at 0.99098 "solute carrier family 35, member F5" Slc35f5
1452060_a_at 0.94828 LIM motif-containing protein kinase 2 Limk2
1452061_s_at 0.87020 spermatid perinuclear RNA binding protein Strbp
1452062_at 0.94811 phosphoribosyl pyrophosphate synthetase-associated protein 2 Prpsap2
1452063_at 0.93329 RIKEN cDNA 2410081M15 gene 2410081M15Rik

1452064_at 0.91018 "cofactor required for Sp1 transcriptional activation, subunit 3" Crsp3
 1452065_at 0.96883 cDNA sequence BC027127 BC027127
 1452066_a_at 0.99080 Nedd4 family interacting protein 2 Ndfip2
 1452067_at 0.93046 N-acylsphingosine amidohydrolase (acid ceramidase)-like Asahl
 1452068_at 0.94167 N-acylsphingosine amidohydrolase (acid ceramidase)-like Asahl
 1452069_a_at 1.06606 THAP domain containing 7 Thap7
 1452070_at 0.95470 death effector domain-containing DNA binding protein 2 Dedd2
 1452071_at 0.92843 "solute carrier family 4 (anion exchanger), member 4" Slc4a4
 1452072_at 0.98470 myc target 1 Myct1
 1452073_at 1.07728 RIKEN cDNA 6720460F02 gene 6720460F02Rik
 1452074_at 0.91863 transmembrane protein 135 Tmem135
 1452075_at 1.22271 RIKEN cDNA 4933435A13 gene 4933435A13Rik
 1452076_at 1.05590 SPRY domain containing 4 Spryd4
 1452077_at 0.96066 "DEAD (Asp-Glu-Ala-Asp) box polypeptide 3, Y-linked" Ddx3y
 1452078_a_at 1.30548 "solute carrier family 11 (proton-coupled divalent metal ion transporters), member 2" Slc11a2
 1452079_s_at 1.02677 "DCUN1D1 DCN1, defective in cullin neddylation 1, domain containing 1 (S. cerevisiae)" Dcun1d1
 1452080_a_at 0.98698 "DCUN1D1 DCN1, defective in cullin neddylation 1, domain containing 1 (S. cerevisiae)" Dcun1d1
 1452081_a_at 0.94622 RIKEN cDNA 9130017N09 gene 9130017N09Rik
 1452082_at 0.91276 RIKEN cDNA 6430548M08 gene 6430548M08Rik
 1452083_a_at 0.85512 "praja1, RING-H2 motif containing" Pja1
 1452084_at 0.99217 "zinc finger, CCHC domain containing 17" Zcchc17
 1452085_at 0.89099 GATA zinc finger domain containing 1 Gatad1
 1452086_at 1.06120 TRM5 tRNA methyltransferase 5 homolog (S. cerevisiae) Trmt5
 1452087_at 0.79281 epithelial stromal interaction 1 (breast) Epsti1
 1452088_at 0.83697 "zinc finger, BED domain containing 3" Zbed3
 1452089_at 1.00040 "calcium channel, voltage-dependent, beta 4 subunit" Cacnb4
 1452090_a_at 0.97405 olfactomedin 3 Olfm3
 1452091_a_at 1.20369 RNA binding motif protein 28 Rbm28
 1452092_at 0.92977 RIKEN cDNA 4631426J05 gene 4631426J05Rik
 1452093_at 0.97498 RIKEN cDNA 2500001K11 gene 2500001K11Rik
 1452094_at 1.06081 "procollagen-proline, 2-oxoglutarate 4-dioxygenase (proline 4-hydroxylase), alpha 1 polypeptide" P4ha1
 1452095_a_at 0.97598 histocompatibility 47 H47
 1452096_s_at 0.98894 RIKEN cDNA D230025D16 gene D230025D16Rik
 1452097_a_at 0.94012 dual specificity phosphatase 7 /// similar to dual specificity phosphatase 7 Dusp7 /// LOC674944
 1452098_at 1.04090 "CTF18, chromosome transmission fidelity factor 18 homolog (S. cerevisiae)" Chtf18
 1452099_at 1.25620 expressed sequence AA408296 AA408296
 1452100_at 0.98556 Dullard homolog (Xenopus laevis) Dullard

1452101_at	0.95337	bleomycin hydrolase	Blmh	
1452102_at	0.93115	"coatamer protein complex, subunit beta 2 (beta prime)"	Copb2	
1452103_at	1.11491	RIKEN cDNA 2600005N12 gene	2600005N12Rik	
1452104_at	0.94367	RIKEN cDNA 2600005N12 gene	2600005N12Rik	
1452105_a_at	0.91376	tuberous sclerosis 2	Tsc2	
1452106_at	0.77801	nephronectin	Npnt	
1452107_s_at	0.71527	Nephronectin	Npnt	
1452108_at	0.95902	insulin-like growth factor I receptor	Igflr	
1452109_at	0.88960	interleukin 17 receptor E	Il17re	
1452110_at	1.06685	5-methyltetrahydrofolate-homocysteine methyltransferase reductase	Mtrr	
1452111_at	1.00419	mitochondrial ribosomal protein S35	Mrps35	
1452112_a_at	0.80317	RNA binding motif protein 4B	Rbm4b	
1452113_a_at	1.22869	"RAB23, member RAS oncogene family"	Rab23	
1452114_s_at	0.68503	insulin-like growth factor binding protein 5	Igfbp5	
1452115_a_at	1.04485	polo-like kinase 4 (Drosophila)	Plk4	
1452116_s_at	1.09244	activating transcription factor 2	Atf2	
1452117_a_at	0.89231	FYN binding protein	Fyb	
1452118_at	1.18755	RIKEN cDNA 2600005C20 gene	2600005C20Rik	
1452119_at	1.09842	RIKEN cDNA 2600005C20 gene	2600005C20Rik	
1452120_at	0.98487	N-acetyltransferase 11	Nat11	
1452121_at	0.87737	F-box only protein 22	Fbxo22	
1452122_at	1.06172	expressed sequence AI314180	AI314180	
1452123_s_at	0.98002	FERM domain containing 4B	Frmd4b	
1452124_at	0.95609	"ankyrin 3, epithelial /// RIKEN cDNA 2900054D09 gene"	Ank3 ///	
		2900054D09Rik		
1452125_at	1.61752	thyroid hormone receptor associated protein 3	Thrap3	
1452126_at	1.09527	zinc finger protein 160	Zfp160	
1452127_a_at	0.88619	"protein tyrosine phosphatase, non-receptor type 13"	Ptpn13	
1452128_a_at	0.98540	"BRCA1/BRCA2-containing complex, subunit 3"	Brcc3	
1452129_at	1.03895	parathyroid hormone receptor 2	Pthr2	
1452130_at	1.00765	thioredoxin domain containing 14	Txndc14	
1452131_at	1.14049	---	---	
1452132_at	1.16178	TLC domain containing 1	Tlcd1	
1452133_at	0.93189	ubiquinol-cytochrome c reductase hinge protein	Uqcrh	
1452134_at	0.83312	RIKEN cDNA 3010001K23 gene	3010001K23Rik	
1452135_at	1.00189	glutathione peroxidase 6	Gpx6	
1452136_at	1.06703	"solute carrier family 5 (sodium/glucose cotransporter), member 9"	Slc5a9	
1452137_at	1.00540	acyl-Coenzyme A binding domain containing 3	Acbd3	
1452138_a_at	0.29340	angiotensin I converting enzyme (peptidyl-dipeptidase A) 2	Ace2	
1452139_at	0.92731	"solute carrier family 35, member C1"	Slc35c1	
1452140_at	0.93575	"TBC1 domain family, member 20"	Tbc1d20	
1452141_a_at	0.52554	"selenoprotein P, plasma, 1"	Sepp1	
1452142_at	1.07414	"solute carrier family 6 (neurotransmitter transporter, GABA), member 1"		

Slc6a1

1452143_at 0.98958 spectrin beta 2 Spnb2
 1452144_a_at 0.96304 mitochondrial ribosomal protein L44 Mrpl44
 1452145_at 1.06665 hexose-6-phosphate dehydrogenase (glucose 1-dehydrogenase) H6pd
 1452146_a_at 0.96409 "COX15 homolog, cytochrome c oxidase assembly protein (yeast)" Cox15
 1452147_at 0.93084 "SEC24 related gene family, member C (S. cerevisiae)" Sec24c
 1452148_at 0.67308 low density lipoprotein receptor-related protein associated protein 1 Lrpap1
 1452149_at 0.87453 ubiquitin protein ligase E3B Ube3b
 1452150_at 0.85101 expressed sequence AU040320 AU040320
 1452151_at 0.95204 cDNA sequence BC021523 BC021523
 1452152_at 1.02955 clathrin interactor 1 Clint1
 1452153_at 0.84576 F-box protein 18 Fbxo18
 1452154_at 1.24998 isoleucine-tRNA synthetase Iars
 1452155_a_at 0.88017 DEAD (Asp-Glu-Ala-Asp) box polypeptide 17 Ddx17
 1452156_a_at 0.88483 nischarin Nisch
 1452157_at 1.10467 glutamyl-prolyl-tRNA synthetase /// similar to Bifunctional aminoacyl-tRNA
 synthetase Eprs /// LOC633677
 1452158_at 1.16911 glutamyl-prolyl-tRNA synthetase /// similar to Bifunctional aminoacyl-tRNA
 synthetase Eprs /// LOC633677
 1452159_at 0.82000 RIKEN cDNA 2310001A20 gene 2310001A20Rik
 1452160_at 1.12058 TCDD-inducible poly(ADP-ribose) polymerase Tiparp
 1452161_at 1.00498 TCDD-inducible poly(ADP-ribose) polymerase Tiparp
 1452162_at 1.04617 WD repeat domain 48 Wdr48
 1452163_at 1.00138 "E26 avian leukemia oncogene 1, 5' domain" Ets1
 1452164_at 0.96042 cDNA sequence BC038286 BC038286
 1452165_at 1.03127 prolactin-like protein K Prlpk
 1452166_a_at 0.90749 keratin 10 Krt10
 1452167_at 1.05476 RIKEN cDNA 2810407C02 gene 2810407C02Rik
 1452168_x_at 0.97548 G1 to S phase transition 1 Gspt1
 1452169_a_at 0.84725 diacylglycerol kinase zeta Dgkz
 1452170_at 0.93360 RIKEN cDNA 2010209O12 gene 2010209O12Rik
 1452171_at 1.00887 glutamate-rich WD repeat containing 1 Grwd1
 1452172_at 0.95348 FAST kinase domains 2 Fastkd2
 1452173_at 0.88534 "hydroxyacyl-Coenzyme A dehydrogenase/3-ketoacyl-Coenzyme A thiolase/
 enoyl-Coenzyme A hydratase (trifunctional protein), alpha subunit" Hadha
 1452174_at 1.06524 sterol regulatory element binding factor 2 Srebf2
 1452175_at 1.02380 RIKEN cDNA 1810026J23 gene 1810026J23Rik
 1452176_at 0.93171 nucleoporin 153 Nup153
 1452177_at 0.97049 "ATP-binding cassette, sub-family F (GCN20), member 3" Abcf3
 1452178_at 0.94271 "plectin 1 /// similar to poly (ADP-ribose) polymerase family, member 10"
 Plec1 /// LOC671535
 1452179_at 0.78970 PHD finger protein 17 Phf17
 1452180_at 0.87787 PHD finger protein 17 Phf17

1452181_at 0.92220 cytoskeleton-associated protein 4 Ckap4
1452182_at 0.80191 UDP-N-acetyl-alpha-D-galactosamine:polypeptide N-acetylgalactosaminyltransferase 2 Galnt2
1452183_a_at 1.06636 "GTL2, imprinted maternally expressed untranslated mRNA" Gtl2
1452184_at 1.03727 "NADH dehydrogenase (ubiquinone) 1 beta subcomplex, 9" Ndufb9
1452185_at 0.92847 importin 8 Ipo8
1452186_at 0.93354 RNA binding motif protein 5 Rbm5
1452187_at 0.90396 RNA binding motif protein 5 Rbm5
1452188_at 1.07970 mastermind like 1 (Drosophila) Maml1
1452189_at 0.98692 WD repeat domain containing 82 Wdr82
1452190_at 0.90924 prolylcarboxypeptidase (angiotensinase C) Prcp
1452191_at 1.03310 prolylcarboxypeptidase (angiotensinase C) Prcp
1452192_at 1.07620 cDNA sequence BC053440 BC053440
1452193_a_at 0.90758 Wiskott-Aldrich syndrome-like (human) Wasl
1452194_at 1.04991 tubulin-specific chaperone d Tbcd
1452195_s_at 0.89033 "Sfi1 homolog, spindle assembly associated (yeast) /// similar to spindle assembly associated Sfi1 homolog isoform b" Sfi1 /// LOC673420
1452196_a_at 1.00868 NCK-associated protein 1 Nckap1
1452197_at 0.92579 structural maintenance of chromosomes 4 Smc4
1452198_at 0.99246 F-box and leucine-rich repeat protein 10 Fbxl10
1452199_at 1.06991 RIKEN cDNA 2700094F01 gene 2700094F01Rik
1452200_at 1.05251 "DNA segment, Chr 11, ERATO Doi 497, expressed" D11Ert497e
1452201_at 0.90577 RIKEN cDNA 2310047B19 gene 2310047B19Rik
1452202_at 0.88198 "phosphodiesterase 2A, cGMP-stimulated" Pde2a
1452203_at 1.23542 RIKEN cDNA 5830411E10 gene 5830411E10Rik
1452204_at 0.97661 ankyrin repeat and SAM domain containing 1 Anks1
1452205_x_at 1.11747 "T-cell receptor beta, joining region /// T-cell receptor beta, variable 13 /// similar to T-cell receptor beta-1 chain C region /// similar to T-cell receptor beta-2 chain C region /// similar to T-cell receptor beta-1 chain C region /// similar to T-cell receptor beta-1 chain C region /// similar to T-cell receptor beta-1 chain C region /// similar to T-cell receptor beta-2 chain C region /// similar to T-cell receptor beta-1 chain C region" Tcrb-J /// Tcrb-V13 /// LOC665500 /// LOC665506 /// LOC669844 /// LOC669847 /// LOC674047 /// LOC674048 /// LOC676782
1452206_at 0.97819 "succinate-Coenzyme A ligase, ADP-forming, beta subunit" Sucla2
1452207_at 0.94056 "Cbp/p300-interacting transactivator, with Glu/Asp-rich carboxy-terminal domain, 2 /// similar to Cbp/p300-interacting transactivator, with Glu/Asp-rich carboxy-terminal domain, 2" Cited2 /// LOC671495
1452208_at 1.21333 PR domain containing 4 Prdm4
1452209_at 0.90089 plakophilin 4 Pkp4
1452210_at 1.06838 DNA2 DNA replication helicase 2-like (yeast) Dna2l
1452211_at 1.12781 "proteasome (prosome, macropain) activator subunit 4" Psme4
1452212_at 0.95353 lamin A Lmna
1452213_at 1.05183 testis expressed gene 2 Tex2
1452214_at 0.90135 SKI-like Skil

1452215_at 1.09521 RIKEN cDNA 9130401M01 gene 9130401M01Rik
1452216_at 0.98774 cDNA sequence BC025519 BC025519
1452217_at 1.02029 AHNAK nucleoprotein (desmoyokin) Ahnak
1452218_at 0.94278 coiled-coil domain containing 117 Ccdc117
1452219_at 1.04631 transmembrane protein 63b Tmem63b
1452220_at 0.93652 dedicator of cyto-kinesis 1 Dock1
1452221_a_at 1.00493 CXXC finger 1 (PHD domain) Cxxc1
1452222_at 0.99735 utrophin Utrn
1452223_s_at 1.11340 granule cell antiserum positive 14 Gcap14
1452224_at 1.10528 microrchidia 3 Morc3
1452225_at 0.95597 RIKEN cDNA 2010106G01 gene 2010106G01Rik
1452226_at 1.25285 regulator of chromosome condensation 2 Rcc2
1452227_at 1.02769 RIKEN cDNA 2310045A20 gene 2310045A20Rik
1452228_at 1.01988 "TBC1 domain family, member 23" Tbc1d23
1452229_at 0.86174 Calmodulin binding transcription activator 2 Camta2
1452230_at 0.95844 "DnaJ (Hsp40) homolog, subfamily C, member 10" Dnajc10
1452231_x_at 1.71246 interferon activated gene 203 Ifi203
1452232_at 1.03017 UDP-N-acetyl-alpha-D-galactosamine: polypeptide N-acetylgalactosaminyltransferase 7 Galnt7
1452233_at 1.16476 "ATP-binding cassette, sub-family C (CFTR/MRP), member 1" Abcc1
1452234_s_at 0.94119 "DNA segment, Chr 16, Brigham & Women's Genetics 1494 expressed" D16Bwg1494e
1452235_at 1.01915 "mannosidase, alpha, class 1B, member 1" Man1b1
1452236_at 1.05726 "ATP-binding cassette, sub-family F (GCN20), member 1" Abcf1
1452237_at 0.98473 HIV-1 Rev binding protein Hrb
1452238_at 1.11893 HIV-1 Rev binding protein Hrb
1452239_at 1.02085 "gene trap ROSA 26, Philippe Soriano" Gt(ROSA)26Sor
1452240_at 0.94022 "bruno-like 4, RNA binding protein (Drosophila)" Brunol4
1452241_at 1.01090 topoisomerase (DNA) II beta binding protein Topbp1
1452242_at 1.01389 centrosomal protein 55 Cep55
1452243_at 1.05557 "potassium inwardly-rectifying channel, subfamily J, member 14" Kcnj14
1452244_at 1.02582 RIKEN cDNA 6330406I15 gene 6330406I15Rik
1452245_at 1.00553 neuro-oncological ventral antigen 1 /// similar to neuro-oncological ventral antigen 1 isoform 2 Nova1 /// LOC664883
1452246_at 1.05922 osteoclast stimulating factor 1 Ostf1
1452247_at 1.04388 "fragile X mental retardation gene 1, autosomal homolog" Fxr1h
1452248_at 0.93491 "pleckstrin homology domain containing, family G (with RhoGef domain) member 5" Plekhg5
1452249_at 0.61716 prickle like 1 (Drosophila) Prickle1
1452250_a_at 0.80507 "procollagen, type VI, alpha 2" Col6a2
1452251_at 0.86364 neurobeachin Nbea
1452252_at 1.22245 "UTP20, small subunit (SSU) processome component, homolog (yeast)" Utp20

1452253_at	1.00586	cysteine rich transmembrane BMP regulator 1 (chordin like)	Crim1
1452254_at	0.99197	myotubularin related protein 9	Mtmr9
1452255_at	1.04749	F-box protein 38	Fbxo38
1452256_at	0.98791	RIKEN cDNA 1110002N22 gene	1110002N22Rik
1452257_at	1.01405	"3-hydroxybutyrate dehydrogenase, type 1"	Bdh1
1452258_at	1.06534	PHD finger protein 20	Phf20
1452259_at	1.05004	PHD finger protein 20	Phf20
1452260_at	1.06272	cell death-inducing DFFA-like effector c	Cidec
1452261_at	1.01869	SNF2 histone linker PHD RING helicase	Shprh
1452262_at	1.20866	"GrpE-like 2, mitochondrial"	Grpel2
1452263_at	1.13176	"solute carrier family 35, member F4"	Slc35f4
1452264_at	0.95279	tensin like C1 domain-containing phosphatase	Tenc1
1452265_at	0.98281	CLIP associating protein 1	Clasp1
1452266_at	1.09506	LAS1-like (<i>S. cerevisiae</i>)	Las1l
1452267_at	0.90820	FLYWCH-type zinc finger 1	Flywch1
1452268_at	1.05936	RIKEN cDNA 2810485I05 gene	2810485I05Rik
1452269_at	0.97144	spectrin beta 3	Spnb3
1452270_s_at	1.03317	cubilin (intrinsic factor-cobalamin receptor)	Cubn
1452271_at	0.77150	xenotropic and polytropic retrovirus receptor 1	Xpr1
1452272_a_at	1.12208	"growth factor, erv1 (<i>S. cerevisiae</i>)-like (augmenter of liver regeneration)"	Gfer
1452273_at	1.01368	expressed sequence AA409316	AA409316
1452274_at	0.97350	"zinc finger, AN1-type domain 3"	Zfand3
1452275_at	0.88559	"zinc finger, AN1-type domain 3"	Zfand3
1452276_at	0.95110	"SWI/SNF-related, matrix-associated actin-dependent regulator of chromatin, subfamily a, containing DEAD/H box 1"	Smarcad1
1452277_at	1.02212	arylsulfatase G	Arsg
1452278_a_at	0.89418	"HECT domain and ankyrin repeat containing, E3 ubiquitin protein ligase 1"	Hace1
1452279_at	0.93665	complement factor properdin	Cfp
1452280_at	1.02693	"FERM, RhoGEF (Arhgef) and pleckstrin domain protein 1 (chondrocyte-derived)"	Farp1
1452281_at	0.91334	Son of sevenless homolog 2 (<i>Drosophila</i>)	Sos2
1452282_at	1.05178	"cofactor required for Sp1 transcriptional activation, subunit 7"	Crsp7
1452283_at	0.83466	Ras association (RalGDS/AF-6) domain family 8	Rassf8
1452284_at	0.94596	"protein tyrosine phosphatase, receptor type Z, polypeptide 1"	Ptprz1
1452285_a_at	1.02730	"eukaryotic translation initiation factor 3, subunit 5 (epsilon)"	Eif3s5
1452286_at	0.88010	RIKEN cDNA 5033405K12 gene	5033405K12Rik
1452287_at	0.80088	"Mus musculus, clone IMAGE:4527696, mRNA"	---
1452288_at	1.05498	expressed sequence BB128963	BB128963
1452289_a_at	1.08464	ring finger protein 135	Rnf135
1452290_at	0.87493	transmembrane protein 106B	Tmem106b
1452291_at	0.77836	"centaurin, delta 1"	Centd1

1452292_at 0.94932 "adaptor-related protein complex 2, beta 1 subunit" Ap2b1
1452293_at 1.01793 sorting nexin 19 Snx19
1452294_at 0.90894 protocadherin 1 Pcdh1
1452295_at 0.67518 similar to Nedd4 WW binding protein 4 /// similar to Nedd4 WW binding protein 4 LOC637870 /// LOC676013
1452296_at 1.05808 slit homolog 3 (Drosophila) Slit3
1452297_at 0.88698 cyclin T2 Ccnt2
1452298_a_at 0.88636 myosin Vb Myo5b
1452299_at 0.97553 WW domain containing E3 ubiquitin protein ligase 1 Wwp1
1452300_at 0.90150 RIKEN cDNA 2410005O16 gene 2410005O16Rik
1452301_at 1.17074 "aldehyde dehydrogenase 3 family, member B1" Aldh3b1
1452302_at 0.88318 Rho guanine nucleotide exchange factor (GEF) 10 Arhgef10
1452303_at 0.94839 Rho guanine nucleotide exchange factor (GEF) 10 Arhgef10
1452304_a_at 0.96084 Rho guanine nucleotide exchange factor (GEF) 5 Arhgef5
1452305_s_at 0.95311 RIKEN cDNA 2610510J17 gene 2610510J17Rik
1452306_at 1.01136 "zinc finger, FYVE domain containing 26" Zfyve26
1452307_at 0.94355 Cdk5 and Abl enzyme substrate 2 Cables2
1452308_a_at 0.93331 "ATPase, Na⁺/K⁺ transporting, alpha 2 polypeptide" Atp1a2
1452309_at 1.06643 cingulin-like 1 /// similar to cingulin-like 1 isoform 1 Cgnl1 /// LOC677485
1452310_at 1.13778 "transcriptional adaptor 2 (ADA2 homolog, yeast)-like" Tada2l
1452311_at 0.85861 dimethylglycine dehydrogenase precursor Dmgdh
1452312_at 1.00693 RIKEN cDNA 2810002D19 gene 2810002D19Rik
1452313_at 0.98802 RIKEN cDNA 5930416I19 gene 5930416I19Rik
1452314_at 1.00688 kinesin family member 11 Kif11
1452315_at 1.03595 kinesin family member 11 Kif11
1452316_at 0.99318 integrator complex subunit 12 Ints12
1452317_at 0.92365 Homeo box B9 Hoxb9
1452318_a_at 0.87973 heat shock protein 1B Hspa1b
1452319_at 1.18725 zinc finger protein 82 Zfp82
1452320_at 0.92868 low density lipoprotein receptor-related protein 2 Lrp2
1452321_at 1.18121 bromodomain and WD repeat domain containing 1 Brwd1
1452322_a_at 0.94499 bromodomain and WD repeat domain containing 1 Brwd1
1452323_at 0.91317 SPRY domain containing 3 Spryd3
1452324_at 1.07331 plasmacytoma variant translocation 1 Pvt1
1452325_at 0.96105 transformation related protein 73 Trp73
1452326_at 0.95109 RIKEN cDNA 1810010G06 gene 1810010G06Rik
1452327_at 0.85005 IQ motif and Sec7 domain 1 Iqsec1
1452328_s_at 0.87729 "praja 2, RING-H2 motif containing" Pja2
1452329_at 1.03954 "pleckstrin homology domain containing, family N member 1" Plekhn1
1452330_a_at 0.76045 matrix-remodelling associated 8 Mxra8
1452331_s_at 1.01631 glutamine and serine rich 1 Qser1
1452332_at 1.00328 coiled-coil domain containing 85A Ccdc85a
1452333_at 0.88275 "SWI/SNF related, matrix associated, actin dependent regulator of chromatin,

subfamily a, member 2" Smarca2

1452334_at 1.02606 centromere protein F Cenpf
 1452335_at 0.80766 RIKEN cDNA 2810423E13 gene 2810423E13Rik
 1452336_at 0.94007 cDNA sequence BC027382 BC027382
 1452337_at 0.88399 RIKEN cDNA 4930427A07 gene 4930427A07Rik
 1452338_s_at 0.96273 intersectin 1 (SH3 domain protein 1A) Itsn1
 1452339_at 0.86467 "a disintegrin-like and metallopeptidase (reprolysin type) with thrombospondin type 1 motif, 7" Adamts7
 1452340_at 1.01167 RIKEN cDNA 6820424L24 gene 6820424L24Rik
 1452341_at 1.04324 "enoyl Coenzyme A hydratase, short chain, 1, mitochondrial" Echs1
 1452342_at 0.92533 "amyloid beta (A4) precursor protein-binding, family B, member 2" Apbb2
 1452343_at 0.89093 "DNA segment, Chr 18, ERATO Doi 653, expressed" D18Ertd653e
 1452344_at 1.26995 synaptojanin 2 Synj2
 1452345_at 1.01506 leiomodulin 2 (cardiac) Lmod2
 1452346_at 0.78718 "UDP-GlcNAc:betaGal beta-1,3-N-acetylglucosaminyltransferase 1" B3gnt1
 1452347_at 0.83784 myocyte enhancer factor 2A Mef2a
 1452348_s_at 1.38213 interferon activated gene 203 /// interferon activated gene 204 /// similar to interferon activated gene 205 /// interferon activated gene 205 /// myeloid cell nuclear differentiation antigen /// similar to Interferon-activatable protein 205 (IFI-205) (D3 protein) /// similar to Interferon-activatable protein 204 (Ifi-204) (Interferon-inducible protein p204) /// similar to Interferon-activatable protein 204 (Ifi-204) (Interferon-inducible protein p204) Ifi203 /// Ifi204 /// LOC192690 /// Ifi205 /// Mnda /// LOC640890 /// LOC672547 /// LOC677184
 1452349_x_at 1.26962 interferon activated gene 205 /// myeloid cell nuclear differentiation antigen Ifi205 /// Mnda
 1452350_at 1.37181 bromodomain containing 8 Brd8
 1452351_at 1.07910 "RIKEN cDNA C030027K23 gene /// poly (ADP-ribose) polymerase family, member 4" C030027K23Rik /// Parp4
 1452352_at 0.83907 cytotoxic T lymphocyte-associated protein 2 beta Ctla2b
 1452353_at 0.84434 G protein-coupled receptor 155 Gpr155
 1452354_at 0.93435 RIKEN cDNA 2810459M11 gene 2810459M11Rik
 1452355_at 0.79772 RIKEN cDNA 3322402L07 gene 3322402L07Rik
 1452356_at 1.03867 IQ motif containing C Iqcc
 1452357_at 0.90901 "glycoprotein Ib, beta polypeptide /// septin 5" Gp1bb /// Sept5
 1452358_at 1.00416 retinoic acid induced 2 Rai2
 1452359_at 0.85288 expressed sequence AA536743 AA536743
 1452360_a_at 0.89054 "jumonji, AT rich interactive domain 1A (Rbp2 like)" Jarid1a
 1452361_at 0.93566 ring finger protein 20 Rnf20
 1452362_at 0.96706 tripartite motif protein 16 Trim16
 1452363_a_at 0.88058 "ATPase, Ca++ transporting, cardiac muscle, slow twitch 2" Atp2a2
 1452364_at 1.06411 suppressor of zeste 12 homolog (Drosophila) Suz12
 1452365_at 0.97587 RIKEN cDNA 4732435N03 gene 4732435N03Rik
 1452366_at 0.72782 RIKEN cDNA 4732435N03 gene 4732435N03Rik
 1452367_at 1.08195 "coronin, actin binding protein 2A" Coro2a

1452368_at 1.13203 breakpoint cluster region homolog Bcr
 1452369_at 0.99286 "membrane associated guanylate kinase, WW and PDZ domain containing 1" Magi1
 1452370_s_at 0.96499 RIKEN cDNA B230208H17 gene B230208H17Rik
 1452371_at 1.00355 "splicing factor, arginine/serine-rich 11" Sfrs11
 1452372_at 0.81175 BSD domain containing 1 Bsdcl
 1452373_at 0.96931 RIKEN cDNA 1700081L11 gene 1700081L11Rik
 1452374_at 1.09484 zinc finger protein 322a Zfp322a
 1452375_at 0.99289 "aldehyde dehydrogenase 4 family, member A1" Aldh4a1
 1452376_at 0.95534 zinc finger protein 444 Zfp444
 1452377_at 1.05885 myeloid/lymphoid or mixed-lineage leukemia 1 Mll1
 1452378_at 1.43387 metastasis associated lung adenocarcinoma transcript 1 (non-coding RNA) Malat1
 1452379_at 1.06451 autism susceptibility candidate 2 Aut2
 1452380_at 0.88571 Eph receptor A7 Epha7
 1452381_at 0.70974 cAMP responsive element binding protein 3-like 2 Creb3l2
 1452382_at 0.99942 "dynamamin 3, opposite strand" Dnm3os
 1452383_at 0.87386 --- ---
 1452384_at 0.87501 ectonucleotide pyrophosphatase/phosphodiesterase 3 Enpp3
 1452385_at 1.08497 ubiquitin specific peptidase 53 Usp53
 1452386_at 1.00385 sal-like 3 (Drosophila) Sall3
 1452387_a_at 0.99952 angiomin like 2 Amotl2
 1452388_at 0.89494 heat shock protein 1A Hspa1a
 1452389_at 0.91465 "tumor necrosis factor receptor superfamily, member 7" Tnfrsf7
 1452390_at 0.88910 EST AA881470 AA881470
 1452391_at 0.96119 coxsackievirus and adenovirus receptor Cxadr
 1452392_a_at 1.02156 "WD repeat domain, phosphoinositide interacting 1" Wipi1
 1452393_at 1.29450 expressed sequence AI597013 AI597013
 1452394_at 1.08914 cysteinyl-tRNA synthetase Cars
 1452395_at 0.95761 "mediator of RNA polymerase II transcription, subunit 19 homolog (yeast)" Med19
 1452396_at 1.03837 "splicing factor, arginine/serine-rich 15 /// similar to splicing factor, arginine/serine-rich 15" Sfrs15 /// LOC670368
 1452397_at 0.91319 RIKEN cDNA 2810474O19 gene 2810474O19Rik
 1452398_at 0.84367 "phospholipase C, epsilon 1" Plce1
 1452399_at 1.01886 regulator of G-protein signaling 6 Rgs6
 1452400_a_at 1.01579 "homeo box A11, opposite strand transcript" Hoxa11os
 1452401_at 0.96223 Wilms' tumour 1-associating protein Wtap
 1452402_at 1.06225 Ubiquitin carboxyl-terminal esterase L3 (ubiquitin thiolesterase) Uchl3
 1452403_a_at 0.88019 otoconin 90 Oc90
 1452404_at 1.08820 phosphatase and actin regulator 2 Phacr2
 1452405_x_at 0.85187 T-cell receptor alpha chain /// RIKEN cDNA A430107P09 gene TcrA /// A430107P09Rik

1452406_x_at 1.13120 erythroid differentiation regulator 1 Erdr1
1452407_at 0.81796 sperm associated antigen 4 Spag4
1452408_at 1.08092 "G protein-coupled receptor 31, D17Leh66c region" Gpr31c
1452409_at 0.88044 glioma tumor suppressor candidate region gene 2 Gltscr2
1452410_a_at 0.89172 feline sarcoma oncogene Fes
1452411_at 0.96830 leucine rich repeat containing 1 Lrrc1
1452412_at 1.09961 homeo box C8 Hoxc8
1452413_at 1.01797 RIKEN cDNA C230081A13 gene C230081A13Rik
1452414_s_at 1.01109 coiled-coil domain containing 86 Ccdc86
1452415_at 1.05923 "actinin, alpha 1" Actn1
1452416_at 1.19152 "interleukin 6 receptor, alpha" Il6ra
1452417_x_at 0.97518 "immunoglobulin kappa chain, constant region /// immunoglobulin kappa chain variable 28 (V28) /// RIKEN cDNA 2010205A11 gene /// immunoglobulin kappa chain variable 8 (V8)-16" Igk-C /// Igk-V28 /// 2010205A11Rik /// Igk-V8-16
1452418_at 1.58365 RIKEN cDNA 1200016E24 gene 1200016E24Rik
1452419_at 0.94233 HEAT repeat containing 1 Heatr1
1452420_at 1.11510 serine/threonine kinase 32B Stk32b
1452421_at 0.89816 homeo box A3 Hoxa3
1452422_a_at 1.10595 U2 small nuclear ribonucleoprotein B Snrpb2
1452423_at 0.98534 piccolo (presynaptic cytomatrix protein) Pclo
1452424_at 0.96226 G protein-coupled receptor 23 Gpr23
1452425_at 0.92636 "tumor necrosis factor receptor superfamily, member 14 (herpesvirus entry mediator)" Tnfrsf14
1452426_x_at 1.08598 Zinc finger protein 236 /// CDNA clone IMAGE:4009820 Zfp236
1452427_s_at 0.95578 protein tyrosine phosphatase-like A domain containing 1 Ptplad1
1452428_a_at 1.20654 beta-2 microglobulin B2m
1452429_s_at 1.03486 "ATP-binding cassette, sub-family F (GCN20), member 1" Abcf1
1452430_s_at 1.04287 "splicing factor, arginine/serine-rich 1 (ASF/SF2)" Sfrs1
1452431_s_at 0.94685 "histocompatibility 2, class II antigen A, alpha /// histocompatibility 2, class II antigen E alpha" H2-Aa /// H2-Ea
1452432_at 1.00261 tissue factor pathway inhibitor Tfpi
1452433_at 1.22004 --- ---
1452434_s_at 1.11110 DiGeorge syndrome critical region gene 6 Dgcr6
1452435_at 0.80828 WD repeat domain 22 Wdr22
1452436_at 1.04731 lysyl oxidase-like 2 Loxl2
1452437_at 0.92028 RIKEN cDNA 1110021J02 gene 1110021J02Rik
1452438_s_at 1.09591 "TAF4A RNA polymerase II, TATA box binding protein (TBP)-associated factor" Taf4a
1452439_s_at 1.04458 "splicing factor, arginine/serine-rich 2 (SC-35)" Sfrs2
1452440_at 1.00826 "tumor necrosis factor (ligand) superfamily, member 12 /// tumor necrosis factor (ligand) superfamily, member 12-member 13" Tnfsf12 /// Tnfsf12-tnfsf13
1452441_at 0.88639 PHD finger protein 3 Phf3
1452442_at 1.07347 Ubiquitin specific peptidase 13 (isopeptidase T-3) Usp13

1452443_s_at 0.97146 helicase with zinc finger domain Helz
1452444_at 0.98435 N-ethylmaleimide sensitive fusion protein attachment protein beta Napb
1452445_at 1.13911 "solute carrier family 41, member 2" Slc41a2
1452446_a_at 0.93275 RIKEN cDNA 2010008E23 gene 2010008E23Rik
1452447_at 0.93106 RIKEN cDNA 0610007P08 gene 0610007P08Rik
1452448_at 0.97668 aquarius Aqr
1452449_at 0.91344 homeobox containing 1 Hmbox1
1452450_at 0.77884 "Atpase, class I, type 8B, member 2" Atp8b2
1452451_at 0.93443 "brain expressed, associated with Nedd4" Bean
1452452_at 0.91819 "PREDICTED: Mus musculus hypothetical protein LOC626156 (LOC626156), mRNA" ---
1452453_a_at 0.99923 calcium/calmodulin-dependent protein kinase II alpha Camk2a
1452454_at 1.21636 SDA1 domain containing 1 Sdad1
1452455_at 1.00291 "fer-1-like 3, myoferlin (C. elegans)" Fer113
1452456_at 0.94958 nuclear receptor interacting protein 2 Nrip2
1452457_a_at 0.90860 "protein kinase, AMP-activated, beta 1 non-catalytic subunit" Prkab1
1452458_s_at 1.12937 peptidylprolyl isomerase (cyclophilin) like 5 Ppil5
1452459_at 0.95593 "asp (abnormal spindle)-like, microcephaly associated (Drosophila)" Aspm
1452460_at 0.92293 ankyrin repeat domain 26 /// similar to ankyrin repeat domain 26
Ankrd26 /// LOC669838
1452461_a_at 0.95452 "N-acetylglucosamine-1-phosphate transferase, alpha and beta subunits"
Gnptab
1452462_a_at 0.96499 Btg3 associated nuclear protein Banp
1452463_x_at 0.92331 Immunoglobulin kappa chain complex ---
1452464_a_at 0.91076 methionine aminopeptidase-like 1 Metap11
1452465_at 1.01101 "myosin, heavy polypeptide 1, skeletal muscle, adult" Myh1
1452466_a_at 0.99387 RNA binding motif protein 6 Rbm6
1452467_at 1.25130 methylmalonic aciduria (cobalamin deficiency) type B homolog (human)
Mmab
1452468_s_at 1.05257 methylmalonic aciduria (cobalamin deficiency) type B homolog (human)
Mmab
1452469_a_at 0.89774 smoothelin Smtn
1452470_at 0.97947 centrosomal protein 350 Cep350
1452471_at 1.30121 interleukin 17 receptor D Il17rd
1452472_at 0.98894 receptor transporter protein 3 Rtp3
1452473_at 1.02824 RIKEN cDNA E130201N16 gene E130201N16Rik
1452474_a_at 1.05306 ADP-ribosyltransferase 3 Art3
1452475_at 0.96722 proprotein convertase subtilisin/kexin type 5 Pcsk5
1452476_at 0.82867 "calcium channel, voltage-dependent, beta 2 subunit" Cacnb2
1452477_at 1.11259 RIKEN cDNA 3110057O12 gene 3110057O12Rik
1452478_at 0.83907 alpha-kinase 2 Alpk2
1452479_at 0.94615 "coatamer protein complex, subunit gamma 2, antisense 2" Copg2as2
1452480_at 0.96382 cDNA sequence BC024760 BC024760

1452481_at 0.88905 "phospholipase C, beta 2" Plcb2
1452482_at 1.05684 v-erb-b2 erythroblastic leukemia viral oncogene homolog 3 (avian) Erbb3
1452483_a_at 1.18999 CD44 antigen Cd44
1452484_at 0.86815 carbonic anhydrase 7 Car7
1452485_at 0.89002 "phosphatase, orphan 1" Phospho1
1452486_a_at 0.87937 "crystallin, alpha A" Cryaa
1452487_x_at 0.87796 paired-Ig-like receptor A2 Pira2
1452488_at 0.95296 --- ---
1452489_at 1.06686 vacuolar protein sorting 11 (yeast) Vps11
1452490_a_at 1.09210 "adaptor protein complex AP-2, alpha 2 subunit" Ap2a2
1452491_at 0.94455 RIKEN cDNA 2900060N18 gene 2900060N18Rik
1452492_a_at 0.99247 "solute carrier family 37 (glycerol-3-phosphate transporter), member 2" Slc37a2
1452493_s_at 1.00670 homeo box B8 Hoxb8
1452494_s_at 0.86019 "solute carrier organic anion transporter family, member 1b2" Slco1b2
1452495_at 1.08214 a disintegrin and metallopeptidase domain 28 Adam28
1452496_at 0.89215 "Atpase, class VI, type 11C" Atp11c
1452497_a_at 1.14680 "nuclear factor of activated T-cells, cytoplasmic, calcineurin-dependent 3" Nfatc3
1452498_at 1.04691 G patch domain containing 2 Gpatc2
1452499_a_at 0.96379 kinesin family member 2A Kif2a
1452500_at 1.32259 "trimethyllysine hydroxylase, epsilon" Tmlhe
1452501_at 0.99276 "cytochrome P450, family 2, subfamily c, polypeptide 38" Cyp2c38
1452502_at 1.00589 small EDRK-rich factor 1 Serf1
1452503_a_at 0.97246 bromodomain and WD repeat domain containing 1 Brwd1
1452504_s_at 0.90135 "chitobiase, di-N-acetyl-" Ctbs
1452505_at 1.01639 "laminin, gamma 2" Lamc2
1452506_a_at 1.04517 integrin-linked kinase-associated serine/threonine phosphatase 2C Ilkap
1452507_at 0.98383 distal-less homeobox 6 Dlx6
1452508_x_at 1.05004 RIKEN cDNA 2610009E16 gene 2610009E16Rik
1452509_at 0.90472 "ubiquitin specific peptidase 9, Y chromosome" Usp9y
1452510_at 1.03455 paired box gene 7 Pax7
1452511_at 0.79516 SRY-box containing gene 5 Sox5
1452512_a_at 0.96963 "ankyrin 1, erythroid" Ank1
1452513_a_at 0.96041 "Fanconi anemia, complementation group A" Fanca
1452514_a_at 0.97912 kit oncogene Kit
1452515_a_at 0.92133 xylosyltransferase II Xylt2
1452516_at 1.05482 sperm motility kinase 2-like /// RIKEN cDNA 4931427P12 gene
LOC545814 /// 4931427P12Rik
1452517_at 1.11220 "pleckstrin homology domain containing, family H (with MyTH4 domain)
member 1" Plekhh1
1452518_a_at 1.05247 IL2-inducible T-cell kinase Itk
1452519_a_at 0.70540 zinc finger protein 36 Zfp36

1452520_a_at 1.04098 "cholinergic receptor, nicotinic, gamma polypeptide" Chrng
1452521_a_at 1.11232 "plasminogen activator, urokinase receptor" Plaur
1452522_at 0.94072 stomatin Stom
1452523_a_at 1.05407 RIKEN cDNA 4930527D15 gene 4930527D15Rik
1452524_a_at 0.96138 "ELAV (embryonic lethal, abnormal vision, Drosophila)-like 3 (Hu antigen C)" Elavl3
1452525_a_at 1.09774 neurofibromatosis 1 Nf1
1452526_a_at 0.94046 paired box gene 6 Pax6
1452527_a_at 1.09269 "purinergic receptor P2X, ligand-gated ion channel 4" P2rx4
1452528_a_at 0.93117 "NK2 transcription factor related, locus 3 (Drosophila)" Nkx2-3
1452529_a_at 0.95439 cAMP responsive element binding protein 1 Creb1
1452530_a_at 1.06654 runt related transcription factor 1 Runx1
1452531_at 1.02679 runt related transcription factor 1 Runx1
1452532_x_at 0.97331 CEA-related cell adhesion molecule 1 Ceacam1
1452533_at 0.84887 ryanodine receptor 3 Ryr3
1452534_a_at 0.98512 high mobility group box 2 /// similar to High mobility group protein 2 (HMG-2) /// high mobility group box 2 /// high mobility group box 2 Hmgb2 /// LOC433785 /// LOC678110 /// LOC678341
1452535_at 0.92379 Immunoglobulin heavy chain 1a (serum IgG2a) Igh-1a
1452536_s_at 1.00678 immunoglobulin kappa chain variable 1 (V1) Igh-V1
1452537_at 0.88576 immunoglobulin kappa chain variable 1 (V1) Igh-V1
1452538_at 1.08010 Immunoglobulin heavy chain 1a (serum IgG2a) Igh-1a
1452539_a_at 1.00874 CD247 antigen Cd247
1452540_a_at 1.10045 "histone 1, H2bc /// histone 1, H2be /// histone 1, H2bl /// histone 1, H2bm /// histone 1, H2bp /// similar to Histone H2B F (H2B 291A) /// similar to Histone H2B F (H2B 291A) /// similar to Histone H2B F (H2B 291A)" Hist1h2bc /// Hist1h2be /// Hist1h2bl /// Hist1h2bm /// Hist1h2bp /// LOC665596 /// LOC665622 /// LOC671645
1452541_at 0.92168 erythrocyte protein band 4.1-like 2 Epb4.112
1452542_x_at 0.87423 transcription factor E2a Tcfe2a
1452543_a_at 1.04714 "secretoglobin, family 1A, member 1 (uteroglobin)" Scgb1a1
1452544_x_at 2.03087 "histocompatibility 2, D region locus 1" H2-D1
1452545_a_at 0.91819 integrin beta 1 (fibronectin receptor beta) Itgb1
1452546_x_at 1.02334 defensin beta 11 Defb11
1452547_s_at 0.97198 "histocompatibility 2, T region locus 18 /// similar to histocompatibility 2, T region locus 3 /// similar to histocompatibility 2, T region locus 3" H2-T18 /// LOC633417 /// LOC674370
1452548_x_at 0.99487 "histocompatibility 2, T region locus 18 /// histocompatibility 2, T region locus 3 /// similar to histocompatibility 2, T region locus 3 /// MHC class I antigen /// similar to histocompatibility 2, T region locus 3 /// similar to histocompatibility 2, T region locus 3" H2-T18 /// H2-T3 /// H2-Tw3 /// H2-T3-like /// LOC633417 /// LOC674370
1452549_at 1.02969 "gene model 1549, (NCBI)" Gm1549
1452550_a_at 0.92947 fragile X mental retardation syndrome 1 homolog Fmr1
1452551_at 1.24285 --- ---

1452552_at 0.98463 neoplastic progression 2 Npn2
1452553_at 0.99274 "Mus musculus, clone IMAGE:5068294" ---
1452554_at 0.89071 "Strain DBA/2J thiopurine S-methyltransferase (Tpmt) pseudogene, mRNA sequence" ---
1452555_at 0.91220 hypothetical protein LOC667141 /// hypothetical protein LOC671961 /// hypothetical protein LOC675420 LOC667141 /// LOC671961 /// LOC675420
1452556_at 0.88043 "Envelope protein, mRNA sequence" ---
1452557_a_at 1.07021 --- ---
1452558_at 0.90166 "T-cell receptor gamma-chain mRNA V4-J1 region, 3' end" ---
1452559_at 1.02730 --- ---
1452560_a_at 1.01257 nuclear transcription factor-Y alpha Nfya
1452561_at 0.92307 Latrophilin 3 Lphn3
1452562_at 0.88593 "jumonji, AT rich interactive domain 1D (Rbp2 like)" Jarid1d
1452563_a_at 0.91968 "jumonji, AT rich interactive domain 1D (Rbp2 like)" Jarid1d
1452564_at 0.94395 oncofetal antigen Ofa
1452565_x_at 0.96571 hypothetical protein LOC641050 LOC641050
1452566_at 1.04434 --- ---
1452567_at 0.98935 granule cell antiserum positive 8 Gcap8
1452568_at 1.07435 granule cell antiserum positive 8 Gcap8
1452569_at 1.00063 "solute carrier organic anion transporter family, member 1b2" Slco1b2
1452570_at 0.93478 "DNA segment, KIST 5" D0Kist5
1452571_at 1.01328 "tubulin alpha, related sequence 1" Tuba-rs1
1452572_at 0.89620 calcium/calmodulin-dependent protein kinase IV Camk4
1452573_a_at 0.91486 stretch responsive protein 278 Sr278
1452574_x_at 0.92188 Immunoglobulin heavy chain (J558 family) Igh-VJ558
1452575_at 1.03533 keratin associated protein 6-3 Krtap6-3
1452576_at 0.97290 Immunoglobulin heavy chain complex Igh
1452577_at 1.04728 Immunoglobulin heavy chain complex Igh
1452578_at 1.15024 runt related transcription factor 1 Runx1
1452579_at 0.91660 NifU-like N-terminal domain containing Nifun
1452580_a_at 0.99627 mitochondrial ribosomal protein L21 Mrpl21
1452581_at 1.03491 RIKEN cDNA 2610304G08 gene 2610304G08Rik
1452582_at 0.84596 galactose mutarotase Galm
1452583_s_at 0.90212 galactose mutarotase Galm
1452584_at 1.02904 RIKEN cDNA 1500032L24 gene 1500032L24Rik
1452585_at 1.06865 mitochondrial ribosomal protein S28 Mrps28
1452586_at 0.91318 anaphase promoting complex subunit 13 Anapc13
1452587_at 1.09955 ARP2 actin-related protein 2 homolog (yeast) Actr2
1452588_at 1.03855 zinc finger protein 688 Zfp688
1452589_at 0.91520 PTK7 protein tyrosine kinase 7 Ptk7
1452590_a_at 1.04071 placenta specific 9 Plac9
1452591_a_at 1.00850 RIKEN cDNA 2410018G20 gene 2410018G20Rik
1452592_at 1.02205 microsomal glutathione S-transferase 2 Mgst2

1452593_a_at 1.06205 "transcription elongation factor B (SIII), polypeptide 1" Tceb1
 1452594_at 1.11986 dual specificity phosphatase 11 (RNA/RNP complex 1-interacting) Dusp11
 1452595_at 1.14209 "a disintegrin-like and metallopeptidase (reprolysin type) with thrombospondin
 type 1 motif, 4" Adamts4
 1452596_at 0.90220 polymerase (RNA) II (DNA directed) polypeptide K Polr2k
 1452597_at 0.92480 RIKEN cDNA 2310061C15 gene 2310061C15Rik
 1452598_at 1.04549 GINS complex subunit 1 (Psf1 homolog) Gins1
 1452599_s_at 1.24875 expressed sequence AI413582 AI413582
 1452600_at 0.99178 "TAF6-like RNA polymerase II, p300/CBP-associated factor (PCAF)-
 associated factor" Taf6l
 1452601_a_at 0.97466 acyl-Coenzyme A binding domain containing 6 Acbd6
 1452602_a_at 0.95312 RIKEN cDNA 1700001C19 gene 1700001C19Rik
 1452603_at 0.99770 RIKEN cDNA 5330431N19 gene 5330431N19Rik
 1452604_at 1.06416 serologically defined colon cancer antigen 13 Stard13
 1452605_at 0.90221 threonine synthase-like 1 (bacterial) Thns11
 1452606_at 1.20813 meiotic nuclear divisions 1 homolog (*S. cerevisiae*) Mnd1
 1452607_at 0.97104 RIKEN cDNA 2610030H06 gene 2610030H06Rik
 1452608_at 0.91449 c-myc binding protein Mycbp
 1452609_at 0.63983 RIKEN cDNA 1190005I06 gene 1190005I06Rik
 1452610_at 0.99907 zinc finger protein 99 Zfp99
 1452611_at 0.96523 zinc finger protein 294 Zfp294
 1452612_at 1.12315 zinc finger protein 294 Zfp294
 1452613_at 0.88262 "ubiquinol-cytochrome c reductase, complex III subunit VII" Uqcrcq
 1452614_at 1.28977 "gene model 566, (NCBI)" Gm566
 1452615_s_at 0.87079 tRNA phosphotransferase 1 Trpt1
 1452616_s_at 1.11506 single-stranded DNA binding protein 1 Ssbp1
 1452617_at 0.93237 single-stranded DNA binding protein 1 Ssbp1
 1452618_at 0.98961 CDK5 regulatory subunit associated protein 2 Cdk5rap2
 1452619_a_at 0.85158 ATP/GTP binding protein-like 3 Agbl3
 1452620_at 0.94435 phosphoenolpyruvate carboxykinase 2 (mitochondrial) Pck2
 1452621_at 0.91939 pterin 4 alpha carbinolamine dehydratase/dimerization cofactor of hepatocyte
 nuclear factor 1 alpha (TCF1) 2 Pcbd2
 1452622_a_at 0.98225 TNFRSF1A-associated via death domain Tradd
 1452623_at 0.87259 zinc finger protein 759 Zfp759
 1452624_at 1.05355 leucine rich repeat transmembrane neuronal 1 Lrrtm1
 1452625_at 0.94266 potassium channel tetramerisation domain containing 2 Kctd2
 1452626_a_at 0.86179 RIKEN cDNA 1810014F10 gene 1810014F10Rik
 1452627_at 0.99603 SUMO/sentrin specific peptidase 6 Senp6
 1452628_at 0.95247 BCL2-associated athanogene 5 Bag5
 1452629_at 1.11895 scaffold attachment factor B2 Safb2
 1452630_at 1.01043 triggering receptor expressed on myeloid cells-like 1 Trem11
 1452631_at 0.88823 RUN and FYVE domain-containing 2 Rufy2
 1452632_at 0.97847 hypothetical protein LOC637741 LOC637741

1452633_s_at 1.01615 hypothetical protein LOC637741 LOC637741
1452634_at 0.87074 oocyte specific homeobox 3 Obox3
1452635_x_at 1.01649 Josephin domain containing 3 Josd3
1452636_x_at 0.96317 GTP binding protein 5 Gtpbp5
1452637_a_at 1.18535 bolA-like 1 (E. coli) Bolal
1452638_s_at 0.92073 dynamin 1-like Dnm1l
1452639_at 0.99040 ectonucleotide pyrophosphatase/phosphodiesterase 4 Enpp4
1452640_at 1.03539 RIKEN cDNA 3110007F17 gene /// hypothetical LOC546368 /// hypothetical
LOC625591 3110007F17Rik /// LOC546368 /// LOC625591
1452641_at 0.94667 cDNA sequence BC028663 BC028663
1452642_at 1.04905 Transmembrane protein 16F Tmem16f
1452643_at 1.08392 grainyhead-like 2 (Drosophila) Grhl2
1452644_at 1.00513 methyl-CpG binding domain protein 3-like 2 Mbd3l2
1452645_x_at 0.95074 "killer cell lectin-like receptor subfamily A, member 22" Klra22
1452646_at 0.79621 tumor protein p53 inducible nuclear protein 2 Trp53inp2
1452647_a_at 1.00895 DPH2 homolog (S. cerevisiae) Dph2
1452648_at 1.07586 transforming growth factor beta regulated gene 1 Tbrg1
1452649_at 0.93767 reticulon 4 Rtn4
1452650_at 1.02135 tripartite motif-containing 62 Trim62
1452651_a_at 0.97086 "myosin, light polypeptide 1" Myl1
1452652_at 1.09991 transmembrane protein 158 Tmem158
1452653_at 0.97491 "solute carrier family 25 (mitochondrial carrier, glutamate), member 22"
Slc25a22
1452654_at 0.88428 --- ---
1452655_at 0.71408 "zinc finger, DHHC domain containing 2" Zdhhc2
1452656_at 0.88479 "zinc finger, DHHC domain containing 2" Zdhhc2
1452657_at 0.92374 "adaptor-related protein complex 1, sigma 2 subunit" Ap1s2
1452658_at 1.07723 "KRR1, small subunit (SSU) processome component, homolog (yeast)" Krr1
1452659_at 0.93580 DEK oncogene (DNA binding) Dek
1452660_s_at 0.93320 kelch-like 7 (Drosophila) Klhl7
1452661_at 1.09130 transferrin receptor Tfrc
1452662_a_at 1.09475 "eukaryotic translation initiation factor 2, subunit 1 alpha" Eif2s1
1452663_at 0.98431 SV2 related protein Svop
1452664_a_at 0.83838 transmembrane 7 superfamily member 3 Tm7sf3
1452665_at 0.94923 tetratricopeptide repeat domain 27 Ttc27
1452666_a_at 0.90387 transmembrane and coiled-coil domains 2 Tmcc2
1452667_at 0.93192 "RAB2B, member RAS oncogene family" Rab2b
1452668_x_at 0.95104 "RAB2B, member RAS oncogene family" Rab2b
1452669_at 0.88693 RIKEN cDNA 2810012G03 gene 2810012G03Rik
1452670_at 0.65477 "myosin, light polypeptide 9, regulatory" Myl9
1452671_s_at 0.87073 "lectin, mannose-binding, 1" Lman1
1452672_at 1.01886 THO complex 5 Thoc5
1452673_at 0.93620 RAN binding protein 3 Ranbp3

1452674_a_at 0.99195 "eukaryotic translation initiation factor 3, subunit 12" Eif3s12
1452675_at 1.02425 RNA binding motif protein 22 Rbm22
1452676_a_at 0.99515 polyribonucleotide nucleotidyltransferase 1 Pnpt1
1452677_at 0.96604 polyribonucleotide nucleotidyltransferase 1 /// similar to polyribonucleotide nucleotidyltransferase 1 /// similar to polyribonucleotide nucleotidyltransferase 1 Pnpt1 ///
LOC635551 /// LOC641190
1452678_a_at 1.04482 cysteine conjugate-beta lyase 1 Ccbl1
1452679_at 1.14925 "tubulin, beta 2b" Tubb2b
1452680_at 1.08266 small nuclear ribonucleoprotein D2 Snrpd2
1452681_at 1.01818 deoxythymidylate kinase Dtymk
1452682_at 1.03393 RIKEN cDNA 4632404H22 gene 4632404H22Rik
1452683_at 1.07010 "DnaJ (Hsp40) homolog, subfamily C, member 8 /// similar to DnaJ (Hsp40) homolog, subfamily C, member 8 (predicted)" Dnajc8 /// LOC669266
1452684_at 0.95089 AKT1 substrate 1 (proline-rich) Akt1s1
1452685_at 0.97181 RIKEN cDNA 0610009O20 gene 0610009O20Rik
1452686_s_at 0.91787 "DNA segment, Chr 4, ERATO Doi 196, expressed" D4ErtD196e
1452687_at 1.03052 "dihydrouridine synthase 2-like (SMM1, S. cerevisiae)" Dus2l
1452688_at 0.93418 PRP39 pre-mRNA processing factor 39 homolog (yeast) Prpf39
1452689_at 0.97593 zinc finger protein 512 Zfp512
1452690_at 1.10022 KH-type splicing regulatory protein Khgrp
1452691_at 0.97992 RNA binding motif protein 17 Rbm17
1452692_a_at 0.97843 NADH dehydrogenase (ubiquinone) flavoprotein 2 Ndufv2
1452693_at 0.91297 DEAH (Asp-Glu-Ala-His) box polypeptide 35 Dhx35
1452694_at 0.83174 inositol hexaphosphate kinase 1 Ihpk1
1452695_at 0.96285 RIKEN cDNA 1810033M07 gene 1810033M07Rik
1452696_a_at 1.09272 RIKEN cDNA 4933439C10 gene 4933439C10Rik
1452697_at 1.07933 "CTD (carboxy-terminal domain, RNA polymerase II, polypeptide A) phosphatase, subunit 1" Ctdp1
1452698_at 0.97864 "Ts translation elongation factor, mitochondrial" Tsfm
1452699_at 0.95842 methyltransferase 11 domain containing 1 Mett11d1
1452700_s_at 0.82136 kelch repeat and BTB (POZ) domain containing 7 Kbtbd7
1452701_x_at 1.00654 ubiquitin A-52 residue ribosomal protein fusion product 1 /// similar to ubiquitin A-52 residue ribosomal protein fusion product 1 /// similar to ubiquitin A-52 residue ribosomal protein fusion product 1 /// similar to ubiquitin A-52 residue ribosomal protein fusion product 1 Uba52 /// LOC666586 /// LOC669193 /// LOC676687
1452702_at 1.00891 chloride channel 7 Clcn7
1452703_at 0.91453 RIKEN cDNA 4631427C17 gene 4631427C17Rik
1452704_at 1.00624 RIKEN cDNA 1200015F23 gene 1200015F23Rik
1452705_at 1.04853 RIKEN cDNA 2210010A19 gene 2210010A19Rik
1452706_a_at 1.01298 RIKEN cDNA 1700027J05 gene 1700027J05Rik
1452707_at 0.74537 RIKEN cDNA 4631423F02 gene 4631423F02Rik
1452708_a_at 0.97423 Luc7 homolog (S. cerevisiae)-like Luc7l
1452709_at 1.00649 "polymerase (DNA-directed), delta interacting protein 3" Poldip3

1452710_at 1.08340 RNA pseudouridylate synthase domain containing 4 Rpusd4
1452711_at 0.80064 spastic paraplegia 3A homolog (human) Spg3a
1452712_at 0.95692 heterogeneous nuclear ribonucleoprotein A3 Hnrpa3
1452713_a_at 1.03547 WD repeat domain 57 (U5 snRNP specific) Wdr57
1452714_at 1.00098 "tetratricopeptide repeat, ankyrin repeat and coiled-coil containing 1" Tanc1
1452715_at 1.00783 RIKEN cDNA 2310022K01 gene 2310022K01Rik
1452716_at 0.98562 RIKEN cDNA 5730469M10 gene 5730469M10Rik
1452717_at 1.06646 "solute carrier family 25 (mitochondrial carrier, phosphate carrier), member 24" Slc25a24
1452718_at 1.02834 "E3 ubiquitin protein ligase, HECT domain containing, 1" Edd1
1452719_at 0.93155 --- ---
1452720_a_at 0.97401 FIP1 like 1 (S. cerevisiae) Fip111
1452721_a_at 1.07514 coiled-coil domain containing 53 Ccdc53
1452722_a_at 0.96702 cullin 5 Cul5
1452723_at 1.08084 Kruppel-like factor 9 Klf9
1452724_at 0.94039 "protein phosphatase 1, regulatory (inhibitor) subunit 16A" Ppp1r16a
1452725_a_at 1.03589 "ribonuclease H2, large subunit" Rnaseh2a
1452726_a_at 1.04466 NOP17 Nop17
1452727_at 0.90766 R3H domain containing 2 R3hdm2
1452728_at 0.87005 kin of IRRE like 3 (Drosophila) Kirrel3
1452729_at 0.99083 dolichyl-phosphate mannosyltransferase polypeptide 3 Dpm3
1452730_at 0.96193 RIKEN cDNA 1110033J19 gene 1110033J19Rik
1452731_x_at 1.03516 hypothetical protein LOC544988 /// RIKEN cDNA B930046C15 gene ///
hypothetical protein LOC545007 /// RIKEN cDNA 2610042L04 gene /// similar to Spetex-2F protein ///
similar to Spetex-2H protein /// similar to Spetex-2H protein /// hypothetical protein LOC664825 ///
similar to Spetex-2F protein /// similar to Spetex-2H protein /// similar to Spetex-2H protein /// similar to
Spetex-2H protein /// hypothetical protein LOC666772 /// hypothetical protein LOC671957
LOC544988 /// B930046C15Rik /// LOC545007 /// 2610042L04Rik /// LOC622916 /// LOC622931 ///
LOC638853 /// LOC664825 /// LOC666193 /// LOC666329 /// LOC666442 /// LOC666684 ///
LOC666772 /// LOC671957
1452732_at 1.03240 RIKEN cDNA 2300003P22 gene 2300003P22Rik
1452733_at 0.93346 pantothenate kinase 2 (Hallervorden-Spatz syndrome) Pank2
1452734_at 0.93884 ribonuclease T2 Rnaset2
1452735_at 1.06931 PEST proteolytic signal containing nuclear protein Pcnp
1452736_at 0.82372 R3H domain and coiled-coil containing 1 R3hcc1
1452737_at 0.90344 RIKEN cDNA 2810008M24 gene 2810008M24Rik
1452738_at 0.91438 stomatin-like 1 Stoml1
1452739_at 1.06188 F-box only protein 7 Fbxo7
1452740_at 0.83229 "myosin, heavy polypeptide 10, non-muscle" Myh10
1452741_s_at 1.01762 "glycerol phosphate dehydrogenase 2, mitochondrial" Gpd2
1452742_at 0.83732 "trafficking protein, kinesin binding 1" Trak1
1452743_at 0.97303 --- ---
1452744_at 0.86168 RIKEN cDNA 2210016F16 gene 2210016F16Rik

1452745_at 0.86273 RIKEN cDNA 1810044A24 gene 1810044A24Rik
1452746_at 0.91375 ATPase type 13A2 Atp13a2
1452747_at 1.00348 ATPase type 13A2 /// similar to ATPase type 13A2 Atp13a2 /// LOC676620
1452748_at 0.97950 X-ray repair complementing defective repair in Chinese hamster cells 3 Xrcc3
1452749_at 0.91911 PAP associated domain containing 1 Papd1
1452750_at 0.79752 RIKEN cDNA 5530601H04 gene 5530601H04Rik
1452751_at 0.96500 early B-cell factor 3 Ebf3
1452752_at 0.74790 leprecan 1 Lepre1
1452753_at 1.08945 forkhead box K2 Foxk2
1452754_at 0.93337 cysteine-rich with EGF-like domains 2 Creld2
1452755_at 1.00733 --- ---
1452756_at 0.85392 RIKEN cDNA 0610009J22 gene 0610009J22Rik
1452757_s_at 0.99033 "hemoglobin alpha, adult chain 1" Hba-a1
1452758_s_at 0.99742 "eukaryotic translation initiation factor 4, gamma 2" Eif4g2
1452759_s_at 0.97454 "PTPRF interacting protein, binding protein 1 (liprin beta 1)" Ppfibp1
1452760_at 0.98777 RIKEN cDNA 2510012J08 gene 2510012J08Rik
1452761_a_at 0.92057 RIKEN cDNA 8430436O14 gene 8430436O14Rik
1452762_at 0.80551 RIKEN cDNA 8430436O14 gene 8430436O14Rik
1452763_at 0.86665 non imprinted in Prader-Willi/Angelman syndrome 1 homolog (human)
Nipa1
1452764_at 0.88974 suppressor of cytokine signaling 6 Socs6
1452765_at 0.94867 "similar to solute carrier family 39 (zinc transporter), member 9" LOC674308
1452766_at 1.02324 RIKEN cDNA 2900041A09 gene 2900041A09Rik
1452767_at 1.09529 ribosome binding protein 1 Rrbp1
1452768_at 0.93966 testis expressed gene 261 Tex261
1452769_at 0.99512 RIKEN cDNA 3732413I11 gene 3732413I11Rik
1452770_at 0.96150 "vitamin K epoxide reductase complex, subunit 1" Vkorc1
1452771_s_at 1.20081 acyl-CoA synthetase long-chain family member 3 Acsl3
1452772_at 0.84783 "tankyrase, TRF1-interacting ankyrin-related ADP-ribose polymerase 2" Tnks2
1452773_at 0.91045 RIKEN cDNA 5730494N06 gene 5730494N06Rik
1452774_at 0.99727 heterogeneous nuclear ribonucleoprotein A3 Hnrpa3
1452775_at 1.05403 cysteine and glycine-rich protein 2 binding protein Csrp2bp
1452776_a_at 1.01784 negative regulator of ubiquitin-like proteins 1 Nub1
1452777_a_at 1.01201 negative regulator of ubiquitin-like proteins 1 Nub1
1452778_x_at 1.02278 nucleosome assembly protein 1-like 1 Nap11
1452779_at 1.08897 RIKEN cDNA 3110006E14 gene 3110006E14Rik
1452780_at 0.99117 "general transcription factor IIIC, polypeptide 2, beta" Gtf3c2
1452781_a_at 1.01333 "general transcription factor IIIC, polypeptide 2, beta" Gtf3c2
1452782_a_at 1.00523 thioredoxin 2 Txn2
1452783_at 1.00798 fibronectin type III domain containing 3B Fndc3b
1452784_at 1.01063 integrin alpha V Itgav
1452785_at 1.02222 RIKEN cDNA 1700034H14 gene 1700034H14Rik
1452786_at 0.85609 tRNA 5-methylaminomethyl-2-thiouridylate methyltransferase Trmu

1452787_at 1.10736 protein arginine N-methyltransferase 1 Prmt1
 1452788_at 0.91249 "protein phosphatase 2, regulatory subunit B (B56), epsilon isoform" Ppp2r5e
 1452789_at 0.81386 stannin Snn
 1452790_x_at 0.90530 "NADH dehydrogenase (ubiquinone) 1 alpha subcomplex, 3" Ndufa3
 1452791_at 0.95389 "coenzyme Q2 homolog, prenyltransferase (yeast)" Coq2
 1452792_at 0.97209 DAZ interacting protein 1 Dzip1
 1452793_at 1.00689 DAZ interacting protein 1 Dzip1
 1452794_x_at 1.01326 "spermatogenesis associated glutamate (E)-rich protein 1, pseudogene 1 ///
 similar to Speer1-ps1 protein /// Speer1-like /// similar to spermatogenesis associated glutamate (E)-rich
 protein 2" Speer1-ps1 /// LOC545728 /// LOC545732 /// LOC667901
 1452795_at 1.16087 RIKEN cDNA 1110008B24 gene 1110008B24Rik
 1452796_at 0.99214 differentially expressed in FDCP 6 Def6
 1452797_at 1.02960 FAST kinase domains 3 Fastkd3
 1452798_s_at 0.94253 mitogen activated protein kinase 1 interacting protein 1 Mapk1ip1
 1452799_at 0.95927 RIKEN cDNA 2310009E04 gene 2310009E04Rik
 1452800_a_at 0.93917 RIKEN cDNA 0610008C08 gene /// hypothetical LOC621156 /// hypothetical
 protein LOC631337 0610008C08Rik /// LOC621156 /// LOC631337
 1452801_at 1.01013 "phosphatidylinositol glycan anchor biosynthesis, class K" Pigk
 1452802_at 0.92799 RIKEN cDNA 4933411K16 gene 4933411K16Rik
 1452803_at 1.00518 GLI pathogenesis-related 2 Glipr2
 1452804_at 0.93650 RIKEN cDNA 1700010A17 gene 1700010A17Rik
 1452805_at 1.03451 "DNA segment, Chr 11, Wayne State University 47, expressed" D11Wsu47e
 1452806_at 0.97191 RIKEN cDNA 1500016O10 gene 1500016O10Rik
 1452807_s_at 1.11464 RIKEN cDNA 1500016O10 gene 1500016O10Rik
 1452808_at 1.01472 snurportin 1 Snupn
 1452809_at 1.08263 RIKEN cDNA 9030607L17 gene 9030607L17Rik
 1452810_at 0.91319 integrator complex subunit 10 /// similar to integrator complex subunit 10
 Ints10 /// LOC674888
 1452811_at 1.01209 5-aminoimidazole-4-carboxamide ribonucleotide formyltransferase/IMP
 cyclohydrolase Atic
 1452812_at 0.76968 latrophilin 1 Lphn1
 1452813_a_at 0.93960 RIKEN cDNA 5033428A16 gene 5033428A16Rik
 1452814_at 1.15561 copine III Cpne3
 1452815_at 1.07360 "purinergic receptor P2Y, G-protein coupled 10" P2ry10
 1452816_at 1.05335 myeloid leukemia factor 1 interacting protein Mlf1ip
 1452817_at 0.83807 SET and MYND domain containing 3 Smyd3
 1452818_at 0.90665 "transcription termination factor, RNA polymerase II" Ttf2
 1452819_at 0.98605 latrophilin 3 Lphn3
 1452820_at 0.92693 bromodomain and WD repeat domain containing 2 Brwd2
 1452821_at 0.93168 Tial1 cytotoxic granule-associated RNA binding protein-like 1 Tial1
 1452822_at 1.02546 integrator complex subunit 7 Ints7
 1452823_at 0.92863 glutathione S-transferase kappa 1 Gstk1
 1452824_at 1.00163 "DNA segment, Chr 1, Brigham & Women's Genetics 0212 expressed"

D1Bwg0212e

1452825_at 0.95883 RIKEN cDNA 5330410G16 gene 5330410G16Rik
 1452826_s_at 1.06563 F-box and leucine-rich repeat protein 20 Fbxl20
 1452827_at 0.96635 RIKEN cDNA 1500009C09 gene 1500009C09Rik
 1452828_at 0.83949 F-box only protein 21 Fbxo21
 1452829_at 1.22652 "carbamoyl-phosphate synthetase 2, aspartate transcarbamylase, and dihydroorotase" Cad
 1452830_s_at 1.09157 "carbamoyl-phosphate synthetase 2, aspartate transcarbamylase, and dihydroorotase" Cad
 1452831_s_at 1.03835 phosphoribosyl pyrophosphate amidotransferase Ppat
 1452832_s_at 0.88476 CDP-diacylglycerol synthase (phosphatidate cytidyltransferase) 2 Cds2
 1452833_at 0.99257 Rap guanine nucleotide exchange factor (GEF) 2 Rapgef2
 1452834_at 0.88224 RIKEN cDNA 2600010E01 gene 2600010E01Rik
 1452835_a_at 0.97859 polymerase (RNA) mitochondrial (DNA directed) Polrmt
 1452836_at 1.15158 lipin 2 Lpin2
 1452837_at 1.15780 lipin 2 Lpin2
 1452838_at 1.11924 DEAD (Asp-Glu-Ala-Asp) box polypeptide 10 Ddx10
 1452839_at 1.12781 DPH5 homolog (*S. cerevisiae*) Dph5
 1452840_at 1.01501 RIKEN cDNA 1500009L16 gene 1500009L16Rik
 1452841_at 0.96971 phosphoglucomutase 2-like 1 Pgm2l1
 1452842_at 1.08868 "heat shock protein, alpha-crystallin-related, B9" Hspb9
 1452843_at 0.99504 interleukin 6 signal transducer Il6st
 1452844_at 0.99980 "POU domain, class 6, transcription factor 1" Pou6f1
 1452845_at 0.97003 "hypoxia-inducible factor 1, alpha subunit inhibitor" Hif1an
 1452846_at 1.08369 "protein tyrosine phosphatase, receptor type, f polypeptide (PTPRF), interacting protein (liprin), alpha 4" Ppfia4
 1452847_at 0.92668 RIKEN cDNA 2410008K03 gene 2410008K03Rik
 1452848_at 1.00900 G protein-coupled receptor 178 Gpr178
 1452849_at 0.82529 "transcriptional regulator, SIN3B (yeast)" Sin3b
 1452850_s_at 1.01746 breast cancer metastasis-suppressor 1-like Brms1l
 1452851_at 0.96474 trinucleotide repeat containing 4 Tnrc4
 1452852_at 1.11638 TWIST neighbor Twistnb
 1452853_at 1.04053 carbohydrate kinase-like Carkl
 1452854_at 1.01476 --- ---
 1452855_at 0.96141 "lymphocyte antigen 6 complex, locus K" Ly6k
 1452856_at 0.85121 HCF-binding transcription factor Zhangfei Zf
 1452857_at 0.87024 --- ---
 1452858_at 0.99874 "ELAV (embryonic lethal, abnormal vision, *Drosophila*)-like 1 (Hu antigen R)" Elavl1
 1452859_at 0.93697 RIKEN cDNA 1200016B10 gene 1200016B10Rik
 1452860_at 1.00056 F-box and leucine-rich repeat protein 17 Fbxl17
 1452861_at 0.73626 RIKEN cDNA 2010300C02 gene /// similar to Y59A8B.19
 2010300C02Rik /// LOC639555

1452862_at 0.85595 ras responsive element binding protein 1 Rreb1
1452863_at 1.05333 RIKEN cDNA 1700003F12 gene 1700003F12Rik
1452864_at 0.82427 "mediator of RNA polymerase II transcription, subunit 12 homolog (yeast)-like" Med12l
1452865_at 1.01100 leucine rich repeat containing 27 Lrrc27
1452866_at 1.10632 asparaginyl-tRNA synthetase Nars
1452867_at 1.14526 "procollagen, type IV, alpha 3 (Goodpasture antigen) binding protein" Col4a3bp
1452868_at 1.02897 ubiquitin specific peptidase 24 Usp24
1452869_at 1.15862 PRP38 pre-mRNA processing factor 38 (yeast) domain containing B Prpf38b
1452870_at 1.11001 --- ---
1452871_at 0.93688 nei endonuclease VIII-like 1 (E. coli) Neil1
1452872_at 0.89560 RIKEN cDNA 2900054D09 gene 2900054D09Rik
1452873_at 0.96828 RIKEN cDNA 5830415F09 gene 5830415F09Rik
1452874_at 0.90039 RIKEN cDNA 2510003E04 gene 2510003E04Rik
1452875_at 0.97833 HD domain containing 3 Hddc3
1452876_x_at 0.95374 RIKEN cDNA 2610044O15 gene 2610044O15Rik
1452877_at 0.95354 RIKEN cDNA 2700029M09 gene 2700029M09Rik
1452878_at 0.91288 "protein kinase C, epsilon" Prkce
1452879_at 1.00731 synaptopodin 2 Synpo2
1452880_at 1.10504 "zinc finger, HIT type 3" Znhit3
1452881_at 0.94353 RIKEN cDNA 4833427B12 gene 4833427B12Rik
1452882_at 0.88574 progesterone receptor membrane component 2 Pgrmc2
1452883_a_at 0.98378 RIKEN cDNA 2310002J21 gene 2310002J21Rik
1452884_at 1.07929 "splicing factor, arginine/serine-rich 2, interacting protein" Sfrs2ip
1452885_at 0.94651 "splicing factor, arginine/serine-rich 2, interacting protein" Sfrs2ip
1452886_at 0.87578 "0 day neonate cerebellum cDNA, RIKEN full-length enriched library, clone: C230071G16 product:unclassifiable, full insert sequence" ---
1452887_at 1.05559 TNF receptor-associated factor 3 interacting protein 1 Traf3ip1
1452888_at 0.90315 RIKEN cDNA 1110034G24 gene 1110034G24Rik
1452889_at 0.90371 RIKEN cDNA 2310007H09 gene 2310007H09Rik
1452890_at 0.93035 tubulin tyrosine ligase like family 5 Ttl5
1452891_at 1.12231 coiled-coil domain containing 51 Ccdc51
1452892_at 1.00328 serine/threonine kinase 33 Stk33
1452893_s_at 1.06888 RIKEN cDNA 1110065P19 gene /// RIKEN cDNA 2310040A07 gene 1110065P19Rik /// 2310040A07Rik
1452894_at 1.04766 "ELAV (embryonic lethal, abnormal vision, Drosophila)-like 4 (Hu antigen D)" Elavl4
1452895_at 1.00257 F-box protein 45 Fbxo45
1452896_at 1.07248 gene trap locus 3 Gtl3
1452897_at 1.02877 cell division cycle 2-like 5 (cholinesterase-related cell division controller) Cdc215
1452898_at 1.05026 --- ---

1452899_at 0.86111 RNA imprinted and accumulated in nucleus Rian
 1452900_at 1.11688 "similar to Proline oxidase, mitochondrial precursor (Proline dehydrogenase)"
 LOC671160
 1452901_at 0.94477 cAMP responsive element binding protein 1 Creb1
 1452902_at 0.94860 dehydrogenase/reductase (SDR family) member 13 Dhrs13
 1452903_at 0.99436 RIKEN cDNA 6230427J02 gene 6230427J02Rik
 1452904_at 0.91455 RIKEN cDNA 1700026L06 gene 1700026L06Rik
 1452905_at 0.98030 "GTL2, imprinted maternally expressed untranslated mRNA /// latrophilin 1"
 Gtl2 /// Lphn1
 1452906_at 0.88310 "GTL2, imprinted maternally expressed untranslated mRNA" Gtl2
 1452907_at 1.14651 galactosylceramidase Galc
 1452908_at 0.90804 DIP2 disco-interacting protein 2 homolog A (Drosophila) Dip2a
 1452909_at 0.89690 RIKEN cDNA 2410127E18 gene 2410127E18Rik
 1452910_at 1.01914 Bcl6 interacting corepressor Bcor
 1452911_at 1.00747 "sprouty protein with EVH-1 domain 1, related sequence" Spred1
 1452912_at 0.96174 RIKEN cDNA 2600005O03 gene 2600005O03Rik
 1452913_at 0.86523 Purkinje cell protein 4-like 1 Pcp4l1
 1452914_at 1.09977 RIKEN cDNA 2410024N18 gene 2410024N18Rik
 1452915_at 0.97189 "protein kinase, cAMP dependent regulatory, type II alpha" Prkar2a
 1452916_at 1.00584 WW domain binding protein 7 Wbp7
 1452917_at 0.94415 replication factor C (activator 1) 5 Rfc5
 1452918_at 0.92391 "DNA segment, Chr 19, ERATO Doi 737, expressed" D19Ertd737e
 1452919_a_at 1.24265 RIKEN cDNA 1700012G19 gene 1700012G19Rik
 1452920_a_at 0.93406 peptidylprolyl isomerase (cyclophilin)-like 2 Ppil2
 1452921_at 0.79485 RIKEN cDNA B130050I23 gene B130050I23Rik
 1452922_at 0.94508 "protein phosphatase 1, regulatory subunit 3D" Ppp1r3d
 1452923_at 0.97047 Ndufa12-like Ndufa12l
 1452924_at 0.83839 RIKEN cDNA 2310007D09 gene 2310007D09Rik
 1452925_a_at 0.92296 membrane-associated ring finger (C3HC4) 5 5-Mar
 1452926_at 0.98556 tetratricopeptide repeat domain 14 Ttc14
 1452927_x_at 1.14837 triosephosphate isomerase 1 Tpi1
 1452928_at 0.90306 "ABI gene family, member 3" Abi3
 1452929_at 0.88337 restin (Reed-Steinberg cell-expressed intermediate filament-associated protein)
 Rsn
 1452930_at 1.01114 dedicator of cyto-kinesis 3 Dock3
 1452931_at 1.10874 RIKEN cDNA 2810408M09 gene 2810408M09Rik
 1452932_at 0.94208 zinc finger and BTB domain containing 7a Zbtb7a
 1452933_at 1.03072 ankyrin repeat domain 39 Ankrd39
 1452934_at 0.94849 transmembrane channel-like gene family 5 Tmc5
 1452935_at 0.93782 hypothetical protein LOC546143 LOC546143
 1452936_at 0.99704 cartilage acidic protein 1 Crtac1
 1452937_s_at 1.15379 coiled coil domain containing 28B Ccdc28b
 1452938_at 1.11860 ankyrin repeat and sterile alpha motif domain containing 1B /// similar to AIDA-

1b Anks1b /// LOC544718

1452939_a_at	1.03568	"phosphatidylinositol transfer protein, cytoplasmic 1"	Pitpnc1
1452940_x_at	0.99130	"phosphatidylinositol transfer protein, cytoplasmic 1"	Pitpnc1
1452941_at	1.02098	RIKEN cDNA 0610038F07 gene	0610038F07Rik
1452942_at	0.90867	transmembrane protein 65	Tmem65
1452943_at	0.86148	transmembrane protein 65	Tmem65
1452944_at	0.89117	arylformamidase	Afmid
1452945_at	0.79691	RIKEN cDNA 2610020C11 gene	2610020C11Rik
1452946_a_at	0.92932	RIKEN cDNA 3222401M22 gene	3222401M22Rik
1452947_at	0.80155	"G protein-coupled receptor, family C, group 5, member C"	Gprc5c
1452948_at	0.77889	"tumor necrosis factor, alpha-induced protein 8-like 2"	Tnfaip8l2
1452949_at	1.11604	polymerase (RNA) III (DNA directed) polypeptide B	Polr3b
1452950_at	1.13336	methyltransferase 5 domain containing 1	Mett5d1
1452951_at	0.92410	RIKEN cDNA 2410089E03 gene	2410089E03Rik
1452952_at	1.05371	RIKEN cDNA 9030418K01 gene	9030418K01Rik
1452953_at	0.92075	"family with sequence similarity 18, member B"	Fam18b
1452954_at	1.15970	ubiquitin-conjugating enzyme E2C	Ube2c
1452955_at	1.03723	"general transcription factor IIH, polypeptide 5"	Gtf2h5
1452956_a_at	1.15851	"DNA segment, Chr 12, ERATO Doi 647, expressed"	D12Ertd647e
1452957_at	1.04830	keratin associated protein 3-3	Krtap3-3
1452958_at	1.02615	aspartate beta-hydroxylase domain containing 2	Asphd2
1452959_a_at	1.02453	calpain 10	Capn10
1452960_at	0.87587	SCY1-like 3 (<i>S. cerevisiae</i>)	Scyl3
1452961_at	1.01341	RIKEN cDNA 1200009O22 gene	1200009O22Rik
1452962_at	0.97204	transmembrane protein 25	Tmem25
1452963_at	1.09575	RIKEN cDNA 9530077C05 gene	9530077C05Rik
1452964_at	0.83710	"tubulin tyrosine ligase-like family, member 11"	Ttll11
1452965_at	0.90329	"ankyrin repeat domain 13 family, member D"	Ankrd13d
1452966_at	0.91674	RIKEN cDNA 9130430L19 gene	9130430L19Rik
1452967_at	1.10435	RIKEN cDNA 1700006H03 gene	1700006H03Rik
1452968_at	1.04886	collagen triple helix repeat containing 1	Cthrc1
1452969_at	0.93980	"ATPase, Ca ⁺⁺ transporting, plasma membrane 1"	Atp2b1
1452970_at	0.86948	"zinc finger, MYM-type 2"	Zmym2
1452971_at	1.01376	UPF3 regulator of nonsense transcripts homolog A (yeast)	Upf3a
1452972_at	1.12610	tetratricopeptide repeat domain 32	Ttc32
1452973_at	1.01542	protein phosphatase 1K (PP2C domain containing)	Ppm1k
1452974_at	0.93832	nucleolar protein 8	Nol8
1452975_at	1.02871	alanine-glyoxylate aminotransferase 2-like 1	Agxt2l1
1452976_a_at	0.99115	"solute carrier family 9 (sodium/hydrogen exchanger), isoform 3 regulator 2"	Slc9a3r2
1452977_at	0.82209	zinc fingers and homeoboxes 3	Zhx3
1452978_at	1.07974	transmembrane protein 138	Tmem138
1452979_at	0.89712	RIKEN cDNA 2610110G12 gene	2610110G12Rik

1452980_at 0.89757 RIKEN cDNA 2810468N07 gene 2810468N07Rik
1452981_at 0.95514 contactin 1 Cntn1
1452982_at 0.99191 insulin-like growth factor I receptor Igflr
1452983_at 1.00704 centrosomal protein 57 Cep57
1452984_at 0.89012 RIKEN cDNA 5730405I09 gene 5730405I09Rik
1452985_at 0.97277 uveal autoantigen with coiled-coil domains and ankyrin repeats Uaca
1452986_at 0.93987 "homogentisate 1, 2-dioxygenase" Hgd
1452987_at 1.61776 Josephin domain containing 3 Jsd3
1452988_at 0.88013 RIKEN cDNA 2610306M01 gene 2610306M01Rik
1452989_at 1.04377 RIKEN cDNA 2900009J20 gene 2900009J20Rik
1452990_at 0.94556 metal response element binding transcription factor 1 Mtf1
1452991_at 0.79885 RIKEN cDNA 2810013C04 gene 2810013C04Rik
1452992_at 0.86909 similar to cell division cycle 26 /// similar to cell division cycle 26 /// similar to
cell division cycle 26 LOC668450 /// LOC672690 /// LOC675848
1452993_at 0.98004 RIKEN cDNA 5430416O09 gene 5430416O09Rik
1452994_at 0.98394 "phosphatidylinositol-3-phosphate/phosphatidylinositol 5-kinase, type III"
Pip5k3
1452995_at 0.81628 nudix (nucleoside diphosphate linked moiety X)-type motif 17 Nudt17
1452996_a_at 1.02498 "apoptosis, caspase activation inhibitor" Aven
1452997_at 0.94098 RIKEN cDNA 2610005L07 gene 2610005L07Rik
1452998_at 1.03475 RIKEN cDNA 2010200O16 gene 2010200O16Rik
1452999_at 1.01294 survival motor neuron domain containing 1 Smndc1
1453000_at 0.92194 calmodulin regulated spectrin-associated protein 1-like 1 Camsap111
1453001_at 1.07520 deleted in lymphocytic leukemia 8 Dleu8
1453002_at 0.85371 SRY-box containing gene 11 Sox11
1453003_at 0.75352 "sortilin-related receptor, LDLR class A repeats-containing" Sorl1
1453004_at 0.93234 RIKEN cDNA 3110004L20 gene 3110004L20Rik
1453005_at 1.03901 PRP31 pre-mRNA processing factor 31 homolog (yeast) Prpf31
1453006_at 1.15905 RIKEN cDNA 2610306H15 gene 2610306H15Rik
1453007_at 1.00779 RIKEN cDNA 3110082I17 gene 3110082I17Rik
1453008_at 1.06865 RIKEN cDNA 2300002D11 gene 2300002D11Rik
1453009_at 0.98723 carboxypeptidase M Cpm
1453010_at 1.08479 IWS1 homolog (S. cerevisiae) Iws1
1453011_at 0.77015 "3-hydroxybutyrate dehydrogenase, type 2" Bdh2
1453012_at 0.70709 TSC22 domain family 2 Tsc22d2
1453013_at 1.09955 zinc finger protein 740 Zfp740
1453014_a_at 0.81690 SEC31-like 1 (S. cerevisiae) Sec3111
1453015_at 0.90441 RIKEN cDNA 5830471E12 gene 5830471E12Rik
1453016_at 0.92117 RIKEN cDNA 2900042B11 gene 2900042B11Rik
1453017_at 1.05677 "DNA segment, Chr 5, ERATO Doi 585, expressed" D5Ertd585e
1453018_at 0.98132 nuclear VCP-like Nvl
1453019_at 1.07087 nuclear VCP-like Nvl
1453020_at 1.11728 RIKEN cDNA 1810048J11 gene 1810048J11Rik

1453021_at	1.04681	syntaxin binding protein 5 (tomosyn)	Stxbp5
1453022_at	0.82285	GPI-anchored HDL-binding protein 1	Gpihbp1
1453023_at	1.05969	ankyrin repeat and KH domain containing 1	Ankhd1
1453024_at	0.94023	WD repeat domain 37	Wdr37
1453025_at	1.09469	RIKEN cDNA 2900006F19 gene	2900006F19Rik
1453026_at	0.91855	cDNA sequence BC061039	BC061039
1453027_at	1.03247	"discs, large (Drosophila) homolog-associated protein 1"	Dlgap1
1453028_at	1.16764	RIKEN cDNA 4631424J17 gene	4631424J17Rik
1453029_s_at	0.98651	RIKEN cDNA 1110059H15 gene	1110059H15Rik
1453030_at	0.94887	male-specific lethal 2-like 1 (Drosophila)	Msl2l1
1453031_at	1.12709	RIKEN cDNA 0610009B10 gene	0610009B10Rik
1453032_at	0.97083	RIKEN cDNA 2610109B12 gene	2610109B12Rik
1453033_at	0.96786	RIKEN cDNA 2010003O02 gene	2010003O02Rik
1453034_at	0.86591	zinc finger protein 251	Zfp251
1453035_at	0.98178	limb and neural patterns	Lnp
1453036_at	0.89573	RIKEN cDNA 5430439M09 gene	5430439M09Rik
1453037_at	0.93076	---	---
1453038_at	1.00401	RIKEN cDNA 4930422G04 gene	4930422G04Rik
1453039_at	0.98532	zinc finger protein 335	Zfp335
1453040_at	0.73191	RIKEN cDNA 2810402A17 gene	2810402A17Rik
1453041_at	0.97392	Trp53 inducible protein 5	Trp53i5
1453042_at	1.10283	ankyrin repeat and sterile alpha motif domain containing 4B	Anks4b
1453043_at	1.06440	RIKEN cDNA 0610012H03 gene	0610012H03Rik
1453044_at	0.99606	RIKEN cDNA C030014I23 gene	C030014I23Rik
1453045_at	1.13539	RIKEN cDNA 4921537D05 gene	4921537D05Rik
1453046_at	0.92283	"similar to diacylglycerol kinase, eta isoform 2"	LOC623215
1453047_at	1.09420	"benzodiazapine receptor, peripheral-like 1"	Bzrp11
1453048_at	1.04008	NHL repeat containing 2	Nhlrc2
1453049_at	0.96151	RIKEN cDNA 6620401M08 gene	6620401M08Rik
1453050_at	0.96432	GINS complex subunit 3 (Psf3 homolog)	Gins3
1453051_at	0.80898	zinc finger with KRAB and SCAN domains 1	Zkscan1
1453052_at	1.21749	RIKEN cDNA 3110005G23 gene	3110005G23Rik
1453053_at	0.80670	RIKEN cDNA 2610036L11 gene	2610036L11Rik
1453054_at	0.89634	secretory carrier membrane protein 1	Scamp1
1453055_at	0.50822	"sema domain, transmembrane domain (TM), and cytoplasmic domain, (semaphorin) 6D"	Sema6d
1453056_at	1.07432	"solute carrier family 16 (monocarboxylic acid transporters), member 13"	Slc16a13
1453057_at	1.02158	transmembrane protein 162	Tmem162
1453058_at	0.90177	WD repeat domain 5B	Wdr5b
1453059_at	1.02137	RIKEN cDNA 2310046A06 gene /// RIKEN cDNA 2810047F03 gene	2310046A06Rik /// 2810047F03Rik
1453060_at	1.01323	regulator of G-protein signaling 8	Rgs8

1453061_at	0.84818	RIKEN cDNA 2610018O07 gene	2610018O07Rik
1453062_at	0.80085	RIKEN cDNA A930026I22 gene	A930026I22Rik
1453063_at	0.95485	"clathrin, light polypeptide (Lcb)"	Cltb
1453064_at	0.84901	RIKEN cDNA 5730466H23 gene	5730466H23Rik
1453065_at	0.86134	"aldhehyde dehydrogenase family 5, subfamily A1"	Aldh5a1
1453066_at	1.00060	RIKEN cDNA 2510022D24 gene	2510022D24Rik
1453067_at	1.02372	RIKEN cDNA 2610040C18 gene	2610040C18Rik
1453068_at	0.97819	"PR domain containing 2, with ZNF domain"	Prdm2
1453069_at	1.18288	"phosphatidylinositol 3-kinase, catalytic, beta polypeptide"	Pik3cb
1453070_at	0.95893	protocadherin 17	Pcdh17
1453071_s_at	1.12220	KDEL (Lys-Asp-Glu-Leu) containing 2	Kdelc2
1453072_at	1.03157	G protein-coupled receptor 160	Gpr160
1453073_at	1.03481	RIKEN cDNA 5830403F22 gene	5830403F22Rik
1453074_at	0.91868	dual specificity phosphatase 23	Dusp23
1453075_at	0.90320	RIKEN cDNA 1600012P17 gene	1600012P17Rik
1453076_at	0.91666	RIKEN cDNA 9130211I03 gene	9130211I03Rik
1453077_a_at	1.06231	"small nuclear RNA activating complex, polypeptide 3"	Snopc3
1453078_at	0.76278	RIKEN cDNA 8430408J07 gene	8430408J07Rik
1453079_at	0.88018	RIKEN cDNA 5730494M16 gene	5730494M16Rik
1453080_at	0.71833	"RIKEN cDNA 9130022K13 gene /// similar to apolipoprotein L, 3 (predicted)"	9130022K13Rik /// LOC672691
1453081_at	0.82986	RIKEN cDNA 2410022M11 gene	2410022M11Rik
1453082_at	1.01636	RIKEN cDNA 2310022L06 gene	2310022L06Rik
1453083_at	1.10948	RIKEN cDNA 6430701C03 gene	6430701C03Rik
1453084_s_at	1.11164	"collagen, type XXII, alpha 1"	Col22a1
1453085_at	1.06720	"serine (or cysteine) peptidase inhibitor, clade A, member 1f"	Serpina1f
1453086_at	0.93632	RIKEN cDNA 6330408A02 gene	6330408A02Rik
1453087_at	1.03404	RIKEN cDNA 6330403L08 gene	6330403L08Rik
1453088_at	0.90718	RIKEN cDNA 6430628N08 gene	6430628N08Rik
1453089_at	1.08423	RIKEN cDNA 3110079O15 gene	3110079O15Rik
1453090_x_at	1.01859	oocyte specific homeobox 1 /// similar to oocyte specific homeobox 2 Obox1 /// LOC666466	
1453091_s_at	0.95315	LETM1 domain containing 1	Letmd1
1453092_at	0.92134	RIKEN cDNA 2300002G24 gene	2300002G24Rik
1453093_at	1.00044	"RasGEF domain family, member 1C"	Rasgef1c
1453094_at	0.77453	checkpoint suppressor 1	Ches1
1453095_at	0.88299	"RAB10, member RAS oncogene family"	Rab10
1453096_x_at	1.08038	ribosomal protein L27 /// similar to ribosomal protein L27 /// similar to ribosomal protein L27 /// similar to ribosomal protein L27 /// similar to ribosomal protein L27	Rpl27 /// LOC621100 /// LOC622236 /// LOC669376 /// LOC672370 /// LOC675088
1453097_a_at	1.00981	"upstream binding transcription factor, RNA polymerase I"	Ubtf
1453098_at	0.90961	"glutamate receptor, ionotropic, AMPA2 (alpha 2)"	Gria2

1453099_at	1.12718	"casein kinase 2, alpha prime polypeptide"	Csnk2a2
1453100_at	1.14204	"casein kinase 2, alpha prime polypeptide"	Csnk2a2
1453101_at	0.98785	Kelch-like 25 (Drosophila)	Klh125
1453102_at	0.79378	fibronectin leucine rich transmembrane protein 3	Flrt3
1453103_at	1.01654	actin-binding LIM protein 1	Ablim1
1453104_at	0.99876	mitogen activated protein kinase 1	Mapk1
1453105_at	1.13876	zinc finger protein 263	Zfp263
1453106_a_at	1.10629	RNA (guanine-7-) methyltransferase	Rnmt
1453107_s_at	1.09777	forkhead box M1 /// phosphatidylethanolamine binding protein 1 /// RIKEN cDNA 4933413G19 gene	Foxm1 /// Pebp1 /// 4933413G19Rik
1453108_at	0.94321	arylsulfatase K	Arsk
1453109_at	0.76620	arylsulfatase K	Arsk
1453110_at	1.07083	WD repeat domain 51B	Wdr51b
1453111_a_at	0.89978	"DNA segment, Chr 11, ERATO Doi 333, expressed"	D11ErtD333e
1453112_a_at	1.00426	leucine rich repeat containing 18	Lrrc18
1453113_at	0.98563	"WD repeat, SAM and U-box domain containing 1"	Wdsub1
1453114_at	1.14596	nucleolar protein 9	Nol9
1453115_at	1.00529	ornithine decarboxylase antizyme 3	Oaz3
1453116_at	0.94919	zinc finger protein 689	Zfp689
1453117_at	0.99060	surfeit gene 2	Surf2
1453118_s_at	1.02064	ribosomal protein L22	Rpl22
1453119_at	0.94346	OTU domain containing 1	Otud1
1453120_at	0.87224	thioredoxin domain containing 13	Txndc13
1453121_at	1.12578	tektin 4	Tekt4
1453122_at	1.01626	RIKEN cDNA 4921533L14 gene	4921533L14Rik
1453123_at	0.93136	"splicing factor 3b, subunit 2"	Sf3b2
1453124_at	1.25634	transportin 3	Tnpo3
1453125_at	0.89038	SRY-box containing gene 11	Sox11
1453126_at	1.02770	leucine rich repeat and fibronectin type III domain containing 2	Lrfn2
1453127_at	0.94657	protein phosphatase 1J	Ppm1j
1453128_at	0.95354	extra cellular link domain-containing 1	Xlkd1
1453129_a_at	0.79508	regulator of G-protein signaling 12	Rgs12
1453130_at	1.06542	E1A binding protein p400	Ep400
1453131_at	0.93721	CD300 antigen like family member G	Cd300lg
1453132_a_at	0.99263	RIKEN cDNA 1810036H07 gene	1810036H07Rik
1453133_at	1.01430	"solute carrier family 25 (mitochondrial carrier; adenine nucleotide translocator), member 31"	Slc25a31
1453134_at	1.02225	"phosphatidylinositol 3-kinase, catalytic, alpha polypeptide"	Pik3ca
1453135_at	0.83378	fibronectin type III domain containing 5	Fndc5
1453136_at	0.94116	F-box protein 30	Fbxo30
1453137_at	0.92127	F-box protein 30	Fbxo30
1453138_at	0.91313	RNA pseudouridylate synthase domain containing 2	Rpusd2
1453139_at	0.72082	nudix (nucleoside diphosphate linked moiety X)-type motif 12	Nudt12

1453140_at	1.16481	RIKEN cDNA 9030612M13	gene	9030612M13Rik
1453141_at	0.95387	RIKEN cDNA 0610009L18	gene	0610009L18Rik
1453142_at	0.92335	RIKEN cDNA 4921507P07	gene	4921507P07Rik
1453143_at	1.01623	RIKEN cDNA 4921509F24	gene	4921509F24Rik
1453144_at	1.01602	RIKEN cDNA 4933439C20	gene	4933439C20Rik
1453145_at	1.03136	RIKEN cDNA 4933439C20	gene	4933439C20Rik
1453146_at	1.10575	GTPase activating protein and VPS9 domains 1		Gapvd1
1453147_at	1.01088	polymerase (RNA) III (DNA directed) polypeptide E		Polr3e
1453148_at	1.02848	"sema domain, immunoglobulin domain (Ig), short basic domain, secreted, (semaphorin) 3D"		Sema3d
1453149_at	1.10134	"solute carrier family 25, member 32"		Slc25a32
1453150_at	1.05114	RIKEN cDNA 5830406J20	gene	5830406J20Rik
1453151_at	0.92163	"solute carrier family 26, member 3"		Slc26a3
1453152_at	1.00087	MAM domain containing 2		Mamdc2
1453153_at	0.91840	lines homolog 2 (Drosophila)		Lins2
1453154_at	0.97792	RIKEN cDNA 1700029M20	gene	1700029M20Rik
1453155_at	1.03475	transmembrane protein 50A		Tmem50a
1453156_s_at	0.90175	"zinc binding alcohol dehydrogenase, domain containing 1"		Zadh1
1453157_at	0.97397	RIKEN cDNA 1700022A21	gene	1700022A21Rik
1453158_at	0.92856	zinc finger protein 622		Zfp622
1453159_at	0.98217	EF-hand domain (C-terminal) containing 1		Efhc1
1453160_at	1.07563	thyroid hormone receptor associated protein 1		Thrap1
1453161_at	1.10978	thyroid hormone receptor associated protein 1		Thrap1
1453162_at	1.16924	"UTP11-like, U3 small nucleolar ribonucleoprotein, (yeast)"		Utp11
1453163_at	0.84607	"protein phosphatase 1, regulatory (inhibitor) subunit 12A"		Ppp1r12a
1453164_a_at	0.80630	phosphatidylserine synthase 2		Ptdss2
1453165_at	1.00339	methyltransferase like 4		Mettl4
1453166_at	0.94339	RIKEN cDNA D130073L02	gene	D130073L02Rik
1453167_at	0.81437	RIKEN cDNA 1700013G24	gene	1700013G24Rik
1453168_at	0.79090	RIKEN cDNA 1700029J07	gene	1700029J07Rik
1453169_a_at	1.07136	"general transcription factor II H, polypeptide 1"		Gtf2h1
1453170_at	0.80495	SFT2 domain containing 3		Sft2d3
1453171_s_at	1.14318	"protein phosphatase 1A, magnesium dependent, alpha isoform"		Ppm1a
1453172_at	1.08257	"stress 70 protein chaperone, microsomal-associated, human homolog"		Stch
1453173_at	0.87912	RIKEN cDNA 2310005E10	gene	2310005E10Rik
1453174_at	0.95357	RIKEN cDNA 2310076G13	gene	2310076G13Rik
1453175_at	1.27151	zinc finger and BTB domain containing 25		Zbtb25
1453176_a_at	1.01316	RIKEN cDNA 4933404M02	gene	4933404M02Rik
1453177_at	0.98914	RIKEN cDNA 4631426E05	gene	4631426E05Rik
1453178_at	1.07912	IQ motif containing H		Iqch
1453179_at	1.07717	"phytoceramidase, alkaline"		Phca
1453180_at	0.93640	RIKEN cDNA 6530404N21	gene	6530404N21Rik
1453181_x_at	1.07067	phospholipid scramblase 1		Plscr1

1453182_at 1.06774 sphingomyelin phosphodiesterase 4 Smpd4
1453183_at 0.99825 RIKEN cDNA 1110034A24 gene 1110034A24Rik
1453184_at 0.86839 RIKEN cDNA 2310040C09 gene 2310040C09Rik
1453185_at 0.99290 RIKEN cDNA 5730406M06 gene 5730406M06Rik
1453186_at 1.00628 RIKEN cDNA 5730406M06 gene 5730406M06Rik
1453187_at 0.79150 OCIA domain containing 2 Ociad2
1453188_at 0.81580 RIKEN cDNA 6230424C14 gene 6230424C14Rik
1453189_at 1.52390 ubiquitin-conjugating enzyme E2I Ube2i
1453190_at 1.02198 ubiquitin specific peptidase 19 Usp19
1453191_at 1.14919 "procollagen, type XXVII, alpha 1" Col27a1
1453192_at 0.98249 RIKEN cDNA 3110032G18 gene 3110032G18Rik
1453193_s_at 0.99512 kinesin family member 12 Kif12
1453194_at 0.69354 RIKEN cDNA 4930432K21 gene 4930432K21Rik
1453195_at 1.00410 serologically defined colon cancer antigen 3 Sdccag3
1453196_a_at 1.36496 2'-5' oligoadenylate synthetase-like 2 Oasl2
1453197_at 0.91544 "mediator of RNA polymerase II transcription, subunit 25 homolog (yeast)"
Med25
1453198_at 1.07440 "zinc finger protein 422, related sequence 1" Zfp422-rs1
1453199_at 1.10231 acyl-Coenzyme A binding domain containing 6 Acbd6
1453200_at 0.91087 retinoic acid induced 1 Rai1
1453201_at 0.99565 RIKEN cDNA 4632411J06 gene 4632411J06Rik
1453202_at 1.02662 RIKEN cDNA E330016A19 gene E330016A19Rik
1453203_at 1.06384 RIKEN cDNA 1700011K15 gene 1700011K15Rik
1453204_at 0.95160 RIKEN cDNA 4933431K14 gene 4933431K14Rik
1453205_at 0.99721 DPY30 domain containing 1 Dydc1
1453206_at 1.02269 "acyl-Coenzyme A dehydrogenase family, member 9" Acad9
1453207_at 0.99038 RIKEN cDNA 2900053A13 gene 2900053A13Rik
1453208_at 1.00029 RIKEN cDNA 2700089E24 gene 2700089E24Rik
1453209_at 1.09272 "BRF1 homolog, subunit of RNA polymerase III transcription initiation factor
IIIB (S. cerevisiae)" Brf1
1453210_at 1.04988 RIKEN cDNA 5730507C01 gene 5730507C01Rik
1453211_at 1.09525 coiled-coil domain containing 89 Ccdc89
1453212_at 1.07732 zinc finger protein 383 Zfp383
1453213_at 0.98587 spermatogenesis associated 9 Spata9
1453214_at 1.20248 leucine rich repeat containing 15 Lrrc15
1453215_at 0.95755 "ribonuclease, RNase A family, 10 (non-active)" Rnase10
1453216_at 1.03522 "glutamic pyruvic transaminase 1, soluble" Gpt1
1453217_at 0.92179 "similar to Xlr-related, meiosis regulated /// similar to Xlr-related, meiosis
regulated /// similar to Xlr-related, meiosis regulated /// similar to Xlr-related, meiosis regulated ///
similar to Xlr-related, meiosis regulated /// similar to Xlr-related, meiosis regulated /// similar to Xlr-
related, meiosis regulated /// similar to Xlr-related, meiosis regulated /// similar to Xlr-related, meiosis
regulated /// similar to Xlr-related, meiosis regulated /// similar to Xlr-related, meiosis regulated"
LOC625508 /// LOC625802 /// LOC625895 /// LOC666096 /// LOC666120 /// LOC666122 ///

LOC666135 /// LOC666151 /// LOC666161 /// LOC666171 /// LOC666179
 1453218_at 0.97264 RIKEN cDNA 1110014K05 gene 1110014K05Rik
 1453219_a_at 0.99598 RIKEN cDNA 2410004F06 gene 2410004F06Rik
 1453220_at 0.96225 RIKEN cDNA 4432416J03 gene 4432416J03Rik
 1453221_at 1.02072 golgi associated PDZ and coiled-coil motif containing Gopc
 1453222_at 0.87809 genetic suppressor element 1 Gse1
 1453223_s_at 0.91985 developmental pluripotency associated 2 Dppa2
 1453224_at 0.86083 "zinc finger, AN1-type domain 5" Zfand5
 1453225_at 1.03331 RIKEN cDNA A930038C07 gene A930038C07Rik
 1453226_at 0.96178 RIKEN cDNA 3000004C01 gene 3000004C01Rik
 1453227_at 0.94103 Rho-related BTB domain containing 3 Rhobtb3
 1453228_at 1.17826 syntaxin 11 Stx11
 1453229_s_at 0.92509 ubiquinol-cytochrome c reductase hinge protein Uqcrh
 1453230_at 1.02829 zinc finger protein 74 Zfp74
 1453231_at 1.00891 cyclin-dependent kinase-like 1 (CDC2-related kinase) Cdkl1
 1453232_at 1.02853 calreticulin 3 Calr3
 1453233_s_at 1.17742 calreticulin 3 Calr3
 1453234_at 0.97370 RIKEN cDNA 1300002K09 gene 1300002K09Rik
 1453235_at 0.91296 RIKEN cDNA 1700012F11 gene 1700012F11Rik
 1453236_at 0.96695 RIKEN cDNA 1700054F22 gene 1700054F22Rik
 1453237_at 0.96486 glutaminyl-tRNA synthetase Qars
 1453238_s_at 1.65718 RIKEN cDNA 3930401B19 gene /// RIKEN cDNA 1200016E24 gene ///
 RIKEN cDNA A130040M12 gene /// RIKEN cDNA E430024C06 gene 3930401B19Rik ///
 1200016E24Rik /// A130040M12Rik /// E430024C06Rik
 1453239_a_at 1.00292 ankyrin repeat domain 22 Ankrd22
 1453240_a_at 0.97170 granule cell antiserum positive 14 Gcap14
 1453241_a_at 0.91608 RIKEN cDNA 2810047C21 gene 2810047C21Rik
 1453242_x_at 1.09400 RIKEN cDNA 2810047C21 gene 2810047C21Rik
 1453243_at 1.11085 RIKEN cDNA 0610030E20 gene 0610030E20Rik
 1453244_at 1.06164 RIKEN cDNA 5830416P10 gene 5830416P10Rik
 1453245_at 1.05761 RIKEN cDNA 9130024F11 gene 9130024F11Rik
 1453246_at 1.06485 "RAB39B, member RAS oncogene family" Rab39b
 1453247_at 1.00330 zinc fingerprotein 618 /// similar to zinc finger protein 618 Zfp618 ///
 LOC667396
 1453248_at 0.95224 RIKEN cDNA 2810039B14 gene 2810039B14Rik
 1453249_a_at 0.90260 testis expressed gene 21 Tex21
 1453250_at 1.11187 RIKEN cDNA 4921511C04 gene 4921511C04Rik
 1453251_at 1.11982 DEAH (Asp-Glu-Ala-His) box polypeptide 30 Dhx30
 1453252_at 1.10050 dihydrouridine synthase 4-like (S. cerevisiae) Dus4l
 1453253_a_at 0.96901 RNA pseudouridylate synthase domain containing 1 Rpusd1
 1453254_at 0.90005 RIKEN cDNA 1700034O15 gene 1700034O15Rik
 1453255_at 1.09409 "solute carrier family 43, member 1" Slc43a1
 1453256_at 1.07511 polymerase (RNA) III (DNA directed) polypeptide C Polr3c

1453257_at 1.13943 "1-acylglycerol-3-phosphate O-acyltransferase 5 (lysophosphatidic acid acyltransferase, epsilon)" Agpat5

1453258_at 0.92063 --- ---

1453259_at 0.97953 RIKEN cDNA 3830422K02 gene 3830422K02Rik

1453260_a_at 1.09906 "protein phosphatase 2 (formerly 2A), regulatory subunit B (PR 52), alpha isoform" Ppp2r2a

1453261_at 0.98751 RIKEN cDNA 2610035D17 gene 2610035D17Rik

1453262_at 0.98863 RIKEN cDNA 2810032G03 gene 2810032G03Rik

1453263_at 0.94627 "MAK10 homolog, amino-acid N-acetyltransferase subunit, (*S. cerevisiae*)" Mak10

1453264_at 0.84540 MARVEL (membrane-associating) domain containing 3 Marveld3

1453265_at 0.99820 RIKEN cDNA 4930579C15 gene 4930579C15Rik

1453266_at 0.94684 --- ---

1453267_at 0.83498 AT motif binding factor 1 Atbf1

1453268_at 0.97250 RIKEN cDNA 5730409G07 gene 5730409G07Rik

1453269_at 0.79186 unc-5 homolog B (*C. elegans*) Unc5b

1453270_a_at 0.94771 PHD finger protein 14 Phf14

1453271_at 0.85600 PHD finger protein 14 Phf14

1453272_at 1.12806 RIKEN cDNA 4930579F01 gene 4930579F01Rik

1453273_at 0.89624 "potassium channel, subfamily V, member 1" Kcnv1

1453274_at 0.83469 coiled-coil domain containing 103 Ccdc103

1453275_at 0.99018 RIKEN cDNA 2310002L13 gene 2310002L13Rik

1453276_at 0.98963 kinesin family member 13B Kif13b

1453277_at 0.92139 RIKEN cDNA 3021401N23 gene 3021401N23Rik

1453278_a_at 0.99309 restin-like 2 Rsnl2

1453279_x_at 1.09503 keratin 76 Krt76

1453280_at 1.08424 "melanoma antigen, family B, 5 /// similar to Melanoma-associated antigen B3 (MAGE-B3 antigen)" Mageb5 /// LOC629542

1453281_at 0.98760 phosphatidylinositol 3-kinase catalytic delta polypeptide Pik3cd

1453282_at 1.11833 coxsackievirus and adenovirus receptor Cxadr

1453283_at 0.99186 phosphoglucomutase 1 Pgm1

1453284_at 0.94204 RIKEN cDNA 4930405M20 gene 4930405M20Rik

1453285_at 0.91061 transmembrane protein 88 Tmem88

1453286_at 0.93701 plexin A2 Plxna2

1453287_at 1.05808 RIKEN cDNA 5730557B15 gene 5730557B15Rik

1453288_at 1.05718 activating transcription factor 6 Atf6

1453289_at 0.99163 "eukaryotic translation initiation factor 2C, 4" Eif2c4

1453290_at 0.96871 high mobility group box 2-like 1 Hmgb2l1

1453291_at 1.11085 high mobility group box 2-like 1 Hmgb2l1

1453292_at 1.06974 RIKEN cDNA 4921501E09 gene 4921501E09Rik

1453293_a_at 0.88047 RIKEN cDNA 2810408A11 gene 2810408A11Rik

1453294_at 0.92338 RIKEN cDNA 1700012B15 gene 1700012B15Rik

1453295_at 0.90900 additional sex combs like 2 (*Drosophila*) Asxl2

1453296_at 1.08078 transmembrane protein 103 Tmem103
1453297_at 0.95698 RIKEN cDNA 4933417O08 gene 4933417O08Rik
1453298_at 1.56690 "protein tyrosine phosphatase, non-receptor type 21" Ptpn21
1453299_a_at 1.25726 purine-nucleoside phosphorylase /// similar to purine-nucleoside phosphorylase /// similar to purine-nucleoside phosphorylase Pnp /// LOC545044 /// LOC667034
1453300_at 0.98909 "solute carrier family 35, member D2 /// similar to UDP-glucuronic acid/UDP-N-acetylgalactosamine transporter" Slc35d2 /// LOC674340
1453301_a_at 0.87744 "alkB, alkylation repair homolog 4 (E. coli)" Alkbh4
1453302_at 1.17372 transmembrane protein 81 Tmem81
1453303_at 0.75634 RIKEN cDNA 4833417J20 gene 4833417J20Rik
1453304_s_at 1.02439 "lymphocyte antigen 6 complex, locus E" Ly6e
1453305_at 0.96112 IQ motif containing D Iqcd
1453306_at 0.89677 RIKEN cDNA 6330531I01 gene 6330531I01Rik
1453307_a_at 0.87778 anaphase-promoting complex subunit 5 Anapc5
1453308_at 1.02578 RIKEN cDNA 8430429K09 gene 8430429K09Rik
1453309_at 1.07205 RIKEN cDNA 9330179D12 gene 9330179D12Rik
1453310_at 0.88056 peptidylprolyl isomerase (cyclophilin)-like 6 Ppil6
1453311_at 1.91038 RIKEN cDNA 2310008B10 gene 2310008B10Rik
1453312_at 0.88714 IQ motif and WD repeats 1 Iqwd1
1453313_at 0.99474 sestrin 3 Sesn3
1453314_x_at 0.99138 RIKEN cDNA 2610039C10 gene 2610039C10Rik
1453315_at 0.90793 within bgcn homolog (Drosophila) Wibg
1453316_at 1.02699 RIKEN cDNA 4933401F05 gene 4933401F05Rik
1453317_a_at 0.81324 "KH domain containing, RNA binding, signal transduction associated 3" Khdrbs3
1453318_at 0.88938 transmembrane protein 102 Tmem102
1453319_at 0.90462 cell division cycle and apoptosis regulator 1 Ccar1
1453320_at 0.99127 RIKEN cDNA 1700027A23 gene 1700027A23Rik
1453321_at 0.94758 fibronectin type III domain containing 1 Fndc1
1453322_at 1.11657 WD repeat domain 33 Wdr33
1453323_at 1.11316 RIKEN cDNA 2900079G21 gene 2900079G21Rik
1453324_at 0.85759 RIKEN cDNA 6330509M23 gene 6330509M23Rik
1453325_at 1.00165 "SAPS domain family, member 2" Saps2
1453326_at 0.96991 RIKEN cDNA 3300001A09 gene /// similar to 3300001A09Rik protein 3300001A09Rik /// LOC244911
1453327_at 0.80730 keratin 24 Krt24
1453328_at 0.98225 RIKEN cDNA 2700008G24 gene 2700008G24Rik
1453329_s_at 1.06144 T-cell lymphoma breakpoint associated target 1 Tcba1
1453330_at 0.97132 RIKEN cDNA 0610010D24 gene 0610010D24Rik
1453331_at 1.04717 RIKEN cDNA 1700013H16 gene 1700013H16Rik
1453332_at 0.97823 RIKEN cDNA 2410002O22 gene 2410002O22Rik
1453333_at 0.97010 RIKEN cDNA 1700012P22 gene 1700012P22Rik
1453334_at 0.94953 RIKEN cDNA B230216N24 gene B230216N24Rik

1453335_a_at 0.91983 spermatogenesis associated 3 Spata3
1453336_at 1.09112 coiled-coil domain containing 98 Ccdc98
1453337_at 0.86358 RIKEN cDNA 1700013E18 gene 1700013E18Rik
1453338_at 1.00503 signal transducing adaptor molecule (SH3 domain and ITAM motif) 2 Stam2
1453339_at 0.82857 RIKEN cDNA 1700008I05 gene 1700008I05Rik
1453340_at 0.98062 RIKEN cDNA 1700057D03 gene 1700057D03Rik
1453341_a_at 0.95200 RIKEN cDNA 4930425N13 gene 4930425N13Rik
1453342_at 1.03812 cell division cycle 40 homolog (yeast) Cdc40
1453343_s_at 0.88096 vaccinia related kinase 2 Vrk2
1453344_at 1.15687 RIKEN cDNA 1810032O08 gene 1810032O08Rik
1453345_at 0.98431 NIPA-like domain containing 1 Npal1
1453346_at 0.95269 RIKEN cDNA 1700011L22 gene 1700011L22Rik
1453347_at 1.24740 RIKEN cDNA 4930589P08 gene 4930589P08Rik
1453348_at 1.05974 RIKEN cDNA 1700085D22 gene 1700085D22Rik
1453349_at 1.05239 "castor homolog 1, zinc finger (Drosophila)" Casz1
1453350_at 1.01656 RIKEN cDNA 2010005J08 gene 2010005J08Rik
1453351_at 1.02236 T-box 20 Tbx20
1453352_at 1.03557 RIKEN cDNA 9030605H24 gene 9030605H24Rik
1453353_at 1.00376 RIKEN cDNA 1700047I16 gene 1700047I16Rik
1453354_at 1.02307 NADH dehydrogenase (ubiquinone) Fe-S protein 1 Ndufs1
1453355_at 0.91193 WNK lysine deficient protein kinase 2 Wnk2
1453356_at 1.43193 RIKEN cDNA 5730453I16 gene 5730453I16Rik
1453357_at 0.88352 tudor domain containing 9 Tdrd9
1453358_s_at 1.04483 RIKEN cDNA 5830467E07 gene /// RIKEN cDNA C730024G19 gene
5830467E07Rik /// C730024G19Rik
1453359_at 1.66953 exosome component 1 Exosc1
1453360_a_at 1.00237 testis expressed gene 9 Tex9
1453361_at 1.71781 "helicase, lymphoid specific" Hells
1453362_x_at 1.06155 ribosomal protein S24 Rps24
1453363_at 1.11430 "calcium channel, voltage-dependent, gamma subunit 2" Cacng2
1453364_x_at 0.97423 RIKEN cDNA 9130019O22 gene 9130019O22Rik
1453365_at 0.93318 RAB GTPase activating protein 1-like Rabgap11
1453366_at 1.16085 tudor and KH domain containing protein Tdrkh
1453367_a_at 0.87937 abhydrolase domain containing 12 Abhd12
1453368_at 1.04838 RIKEN cDNA 2310003H01 gene 2310003H01Rik
1453369_a_at 0.97462 FUN14 domain containing 1 Fundc1
1453370_at 1.09446 trophinin associated protein Troap
1453371_at 1.12481 RIKEN cDNA 4930535B03 gene 4930535B03Rik
1453372_at 0.86246 "DnaJ (Hsp40) homolog, subfamily C, member 1" Dnajc1
1453373_at 1.00656 RIKEN cDNA 6330532G10 gene 6330532G10Rik
1453374_at 0.70448 "zinc finger, AN1-type domain 5" Zfand5
1453375_at 0.89496 RIKEN cDNA 4930422N03 gene 4930422N03Rik
1453376_at 0.97500 RIKEN cDNA 4921524J06 gene 4921524J06Rik

1453377_at	1.04708 SH2 domain containing 4A	Sh2d4a
1453378_at	0.92644 "zinc finger protein, subfamily 1A, 3 (Aiolos)"	Zfpn1a3
1453379_at	0.98651 RIKEN cDNA 4930519P11 gene	4930519P11Rik
1453380_a_at	1.05510 XRCC6 binding protein 1	Xrcc6bp1
1453381_at	0.86921 RIKEN cDNA C030033M12 gene	C030033M12Rik
1453382_at	0.97763 F-box protein 42	Fbxo42
1453383_at	0.94867 RIKEN cDNA 1700022F17 gene	1700022F17Rik
1453384_at	1.10991 RIKEN cDNA 4632404N19 gene	4632404N19Rik
1453385_at	1.04747 RIKEN cDNA 1110015C02 gene	1110015C02Rik
1453386_at	0.78153 tumor suppressor candidate 1	Tusc1
1453387_at	1.05836 RIKEN cDNA 4833432E10 gene	4833432E10Rik
1453388_at	1.04380 RIKEN cDNA 4933416M07 gene	4933416M07Rik
1453389_a_at	0.93226 SH2B adaptor protein 2	Sh2b2
1453390_at	1.06234 RIKEN cDNA 4930428F12 gene	4930428F12Rik
1453391_at	1.01739 "spermatogenesis associated glutamate (E)-rich protein 7, pseudogene 1"	Speer7-ps1
1453392_at	1.08342 RIKEN cDNA 1810054D07 gene	1810054D07Rik
1453393_a_at	1.00815 carbohydrate (chondroitin 6/keratan) sulfotransferase 4	Chst4
1453394_at	0.97786 RIKEN cDNA 2410017I17 gene	2410017I17Rik
1453395_at	0.92588 RIKEN cDNA 5330403D14 gene	5330403D14Rik
1453396_at	1.00507 RIKEN cDNA 1700123D08 gene	1700123D08Rik
1453397_at	1.00863 RIKEN cDNA 9130016M20 gene	9130016M20Rik
1453398_at	1.11448 RIKEN cDNA A930009A15 gene	A930009A15Rik
1453399_at	1.75167 Cyclin T2	Ccnt2
1453400_at	0.91260 RIKEN cDNA 4933428D01 gene	4933428D01Rik
1453401_at	0.93919 RIKEN cDNA 4930428O21 gene	4930428O21Rik
1453402_at	1.09048 RIKEN cDNA 6430500C12 gene	6430500C12Rik
1453403_at	0.90374 RIKEN cDNA 2210019I11 gene	2210019I11Rik
1453404_at	1.01834 RIKEN cDNA 2900002H16 gene	2900002H16Rik
1453405_at	0.93766 "phospholipase A2, group IVD"	Pla2g4d
1453406_a_at	0.96888 "RAB28, member RAS oncogene family"	Rab28
1453407_at	0.95889 RIKEN cDNA 1700018L24 gene	1700018L24Rik
1453408_s_at	1.00392 RIKEN cDNA 4930519G04 gene	4930519G04Rik
1453409_at	0.93816 cell growth regulator with ring finger domain 1	Cgrrf1
1453410_at	0.89765 angiopoietin-like 4	Angptl4
1453411_at	0.92107 RIKEN cDNA 2010110K16 gene	2010110K16Rik
1453412_a_at	0.98373 SEC14-like 1 (S. cerevisiae)	Sec14l1
1453413_at	1.01999 "GNAS (guanine nucleotide binding protein, alpha stimulating) complex locus"	Gnas
1453414_at	0.89886 yippee-like 2 (Drosophila)	Ypel2
1453415_at	0.89967 zinc finger CCCH type containing 6	Zc3h6
1453416_at	1.10819 growth arrest-specific 2 like 3	Gas2l3
1453417_at	1.16908 RIKEN cDNA 4933407K13 gene	4933407K13Rik

1453418_at	0.96667	"procollagen, type XXIV, alpha 1"	Col24a1
1453419_at	0.91947	RIKEN cDNA 2900078C09 gene	2900078C09Rik
1453420_at	1.26780	RIKEN cDNA 1700013G10 gene	1700013G10Rik
1453421_at	0.81581	serine racemase Srr	
1453422_a_at	1.21083	RIKEN cDNA 1110020G09 gene	1110020G09Rik
1453423_at	0.85698	RIKEN cDNA 4933409L14 gene	4933409L14Rik
1453424_at	1.19703	FYVE and coiled-coil domain containing 1	Fyco1
1453425_at	1.02742	coiled-coil domain containing 83	Ccdc83
1453426_a_at	0.90734	WD repeat and FYVE domain containing 1	Wdfy1
1453427_at	0.99317	"casein kinase 2, alpha 1 polypeptide"	Csnk2a1
1453428_at	1.03450	RIKEN cDNA 2700045P11 gene	2700045P11Rik
1453429_at	0.87679	RIKEN cDNA 9530057J20 gene	/// hypothetical protein LOC665756 ///
hypothetical protein LOC665787		9530057J20Rik	/// LOC665756 /// LOC665787
1453430_at	0.98160	RIKEN cDNA 2010315L10 gene	2010315L10Rik
1453431_at	0.96863	RIKEN cDNA 2610206C17 gene	2610206C17Rik
1453432_at	1.08885	RIKEN cDNA 1700007G11 gene	1700007G11Rik
1453433_at	1.13539	RIKEN cDNA 2600001A11 gene	2600001A11Rik
1453434_at	1.05342	RIKEN cDNA 1110019D14 gene	1110019D14Rik
1453435_a_at	1.01016	flavin containing monooxygenase 2	Fmo2
1453436_at	1.01811	RIKEN cDNA 4921516I12 gene	4921516I12Rik
1453437_at	0.99853	RIKEN cDNA 2810407D09 gene	2810407D09Rik
1453438_x_at	1.15582	gasdermin 2	Gsdm2
1453439_at	0.94442	RIKEN cDNA 5430427O21 gene	5430427O21Rik
1453440_at	0.97333	RIKEN cDNA 4921539E11 gene	4921539E11Rik
1453441_at	0.97193	RIKEN cDNA 4921522P10 gene	4921522P10Rik
1453442_at	0.84904	RIKEN cDNA 2310043M15 gene	2310043M15Rik
1453443_at	1.02344	RIKEN cDNA 1110015O18 gene	1110015O18Rik
1453444_at	0.97989	RIKEN cDNA 5730437C12 gene	5730437C12Rik
1453445_at	1.15780	RIKEN cDNA 9030601B04 gene	9030601B04Rik
1453446_at	0.83395	RIKEN cDNA 6430411K18 gene	6430411K18Rik
1453447_at	1.14125	RIKEN cDNA 1700109H08 gene	1700109H08Rik
1453448_at	1.87381	RIKEN cDNA 2310067E19 gene	2310067E19Rik
1453449_at	1.10526	---	---
1453450_at	0.96554	RIKEN cDNA C030006K11 gene	C030006K11Rik
1453451_at	1.36107	RIKEN cDNA 6820402I19 gene	6820402I19Rik
1453452_at	0.90042	RIKEN cDNA 4930517J16 gene	4930517J16Rik
1453453_at	0.86844	"RIKEN cDNA 1700080O16 gene	/// similar to melanoma antigen family A, 5 /// similar to melanoma antigen family A, 5" 1700080O16Rik
		/// LOC665895	/// LOC671223
		/// LOC676479	
1453454_at	0.98925	RIKEN cDNA 1700012E03 gene	1700012E03Rik
1453455_at	0.83977	RIKEN cDNA 1810059M14 gene	1810059M14Rik
1453456_at	0.97984	RIKEN cDNA 2900084O13 gene	2900084O13Rik
1453457_at	1.39458	RIKEN cDNA 2900070H08 gene	2900070H08Rik

1453458_at 1.19879 RIKEN cDNA 9430019C24 gene 9430019C24Rik
1453459_at 1.07629 RIKEN cDNA 1700028I16 gene 1700028I16Rik
1453460_at 0.90943 RIKEN cDNA 1700012I11 gene 1700012I11Rik
1453461_at 1.06063 fractured callus expressed transcript 1 Fxc1
1453462_at 1.03682 RIKEN cDNA 1110067M19 gene 1110067M19Rik
1453463_at 0.98095 RIKEN cDNA 1700019B03 gene 1700019B03Rik
1453464_at 1.01453 RIKEN cDNA D530014G21 gene D530014G21Rik
1453465_x_at 0.91990 "protein phosphatase 1, regulatory (inhibitor) subunit 14c" Ppp1r14c
1453466_at 1.01352 ribosomal protein S6 Rps6
1453467_s_at 1.06613 ribosomal protein S15a Rps15a
1453468_at 0.97295 RIKEN cDNA 4930430F08 gene 4930430F08Rik
1453469_at 1.07915 RIKEN cDNA 1810073G21 gene 1810073G21Rik
1453470_a_at 1.14096 "guanine nucleotide binding protein, alpha 13" Gna13
1453471_at 1.10436 RIKEN cDNA 4833421E05 gene 4833421E05Rik
1453472_a_at 1.05074 SLAM family member 7 Slamf7
1453473_a_at 0.97000 dynein light chain Tctex-type 1 /// similar to Cytoplasmic dynein light chain (T-complex testis-specific protein 1) (TCTEX-1) Dynlt1 /// LOC671261
1453474_at 0.95162 RIKEN cDNA 1300007F04 gene 1300007F04Rik
1453475_at 1.34695 RIKEN cDNA 4930445K14 gene 4930445K14Rik
1453476_at 1.01269 RIKEN cDNA 1700060J05 gene 1700060J05Rik
1453477_at 0.94573 zinc finger CCCH type containing 6 Zc3h6
1453478_at 1.00884 "POU domain, class 3, transcription factor 2" Pou3f2
1453479_at 0.91829 RIKEN cDNA 1700086P04 gene 1700086P04Rik
1453480_at 1.08486 RIKEN cDNA D630002J15 gene D630002J15Rik
1453481_at 0.64460 "zinc finger, DHHC domain containing 2" Zdhhc2
1453482_at 1.00632 RIKEN cDNA 2310024N18 gene 2310024N18Rik
1453483_at 1.00143 RIKEN cDNA 1700023E05 gene 1700023E05Rik
1453484_at 0.87821 RIKEN cDNA 1700081D17 gene 1700081D17Rik
1453485_s_at 1.06222 RIKEN cDNA 1110005A03 gene 1110005A03Rik
1453486_a_at 0.99408 "signal peptide, CUB domain, EGF-like 2" Scube2
1453487_at 1.05603 dihydrodiol dehydrogenase (dimeric) Dhdh
1453488_at 1.06910 RIKEN cDNA 4930458D05 gene 4930458D05Rik
1453489_at 0.69831 RIKEN cDNA 1700020O03 gene 1700020O03Rik
1453490_at 0.96076 spindle assembly 6 homolog (C. elegans) Sass6
1453491_at 0.91671 RIKEN cDNA 4833409A17 gene 4833409A17Rik
1453492_at 0.98945 transmembrane protein 67 Tmem67
1453493_at 0.97694 RIKEN cDNA 4921528O07 gene /// similar to RIKEN cDNA 4921528O07 gene /// hypothetical protein LOC640258 4921528O07Rik /// LOC625202 /// LOC640258
1453494_at 0.83538 RIKEN cDNA 4921513H07 gene 4921513H07Rik
1453495_at 0.97935 kallikrein 12 Klk12
1453496_at 0.98197 RIKEN cDNA 9430073N08 gene 9430073N08Rik
1453497_a_at 1.07392 "phosphatidylinositol glycan anchor biosynthesis, class A" Piga
1453498_x_at 1.03013 STEAP family member 3 Steap3

1453499_at	0.94964	D-aspartate oxidase	Ddo	
1453500_at	1.09697	"cytochrome P450, family 2, subfamily u, polypeptide 1"	Cyp2u1	
1453501_at	1.00282	homeo box B1	Hoxb1	
1453502_at	0.84906	RIKEN cDNA 2210408I21	gene	2210408I21Rik
1453503_at	1.12888	RIKEN cDNA 9230117E20	gene	9230117E20Rik
1453504_at	1.21641	"TAF15 RNA polymerase II, TATA box binding protein (TBP)-associated factor"	Taf15	
1453505_a_at	0.86184	eukaryotic translation initiation factor 2 alpha kinase 3		Eif2ak3
1453506_at	1.07814	RIKEN cDNA 4930500O09	gene	4930500O09Rik
1453507_at	0.86729	methyl-CpG binding domain protein 5	Mbd5	
1453508_at	1.08325	RIKEN cDNA 1700104B16	gene	1700104B16Rik
1453509_at	0.85761	Ly6/Plaur domain containing 2	Lypd2	
1453510_s_at	0.88962	RIKEN cDNA 4930589M24	gene	4930589M24Rik
1453511_at	0.99329	RIKEN cDNA 2310007B03	gene	2310007B03Rik
1453512_at	2.60858	RIKEN cDNA 5830407P18	gene	5830407P18Rik
1453513_at	1.19893	---	---	
1453514_at	1.04242	glycoprotein m6b	Gpm6b	
1453515_at	1.05791	RIKEN cDNA 4930511A21	gene	4930511A21Rik
1453516_at	0.98992	RIKEN cDNA 4931431F19	gene	4931431F19Rik
1453517_at	1.01672	methionine aminopeptidase-like 1	Metap11	
1453518_at	0.96166	PDZ domain containing 11	Pdzd11	
1453519_at	1.02371	RIKEN cDNA 2610037D02	gene	2610037D02Rik
1453520_at	1.02170	tetratricopeptide repeat domain 21B	Ttc21b	
1453521_at	0.87438	RIKEN cDNA 4933405L10	gene	4933405L10Rik
1453522_at	0.97358	RIKEN cDNA 6530401N04	gene	6530401N04Rik
1453523_at	1.02707	RIKEN cDNA A030006P16	gene	A030006P16Rik
1453524_at	1.29062	RIKEN cDNA 5530401D11	gene	5530401D11Rik
1453525_at	1.10461	RIKEN cDNA 9130413E14	gene	9130413E14Rik
1453526_at	0.99674	RIKEN cDNA 4930503E14	gene	4930503E14Rik
1453527_a_at	0.90511	neuralized-like homolog (Drosophila)	Neurl	
1453528_at	0.93830	leukotriene A4 hydrolase	Lta4h	
1453529_at	0.91620	RIKEN cDNA 6330418B08	gene	6330418B08Rik
1453530_at	0.98573	---	---	
1453531_at	1.06422	zinc finger protein 248	Zfp248	
1453532_at	0.86914	RIKEN cDNA 4933415A04	gene	4933415A04Rik
1453533_at	0.90555	RIKEN cDNA 4933403O08	gene	4933403O08Rik
1453534_at	1.19486	RIKEN cDNA 2810004I08	gene	2810004I08Rik
1453535_at	0.97729	olfactory receptor 78	Olf78	
1453536_at	1.00989	RIKEN cDNA 1700034K16	gene	1700034K16Rik
1453537_a_at	1.02356	WD repeat domain 17	Wdr17	
1453538_at	1.03695	RIKEN cDNA A330102K23	gene	A330102K23Rik
1453539_at	0.92143	"alkB, alkylation repair homolog 8 (E. coli)"	Alkbh8	
1453540_at	1.08288	RIKEN cDNA 5430404G13	gene	5430404G13Rik

1453541_at	1.70673	"phosphofructokinase, liver, B-type"	Pfkl
1453542_at	0.99525	RIKEN cDNA 2610007B07	gene 2610007B07Rik
1453543_at	1.05404	RIKEN cDNA 4930556H04	gene 4930556H04Rik
1453544_at	0.99170	DMRT-like family C1c /// similar to doublesex and mab-3 related transcription factor 8.1 isoform a	Dmr1c1c /// LOC668357
1453545_at	1.02351	RIKEN cDNA 5730492I20	gene 5730492I20Rik
1453546_at	0.87693	RIKEN cDNA 1700029I08	gene 1700029I08Rik
1453547_at	1.08767	RIKEN cDNA 1810046K07	gene 1810046K07Rik
1453548_at	0.95705	---	---
1453549_at	1.09706	RIKEN cDNA 4930500G05	gene 4930500G05Rik
1453550_a_at	0.93959	male sterility domain containing 2	Mlstd2
1453551_at	0.95299	"polymerase (DNA directed), theta"	Polq
1453552_at	1.07887	RIKEN cDNA 2310014F07	gene 2310014F07Rik
1453553_at	1.01593	---	---
1453554_a_at	0.98228	WD repeat domain 33	Wdr33
1453555_at	0.98838	RIKEN cDNA 3830408D07	gene 3830408D07Rik
1453556_x_at	0.93217	CD99 antigen	Cd99
1453557_at	0.90596	RIKEN cDNA 4930562F07	gene 4930562F07Rik
1453558_at	0.87416	RIKEN cDNA 4930504H06	gene 4930504H06Rik
1453559_a_at	0.90939	Sell (suppressor of lin-12) 1 homolog (C. elegans)	Sellh
1453560_at	0.98315	feminization 1 homolog a (C. elegans)	Fem1a
1453561_x_at	1.00955	RIKEN cDNA 8430431K14	gene 8430431K14Rik
1453562_a_at	0.91285	NmrA-like family domain containing 1	Nmr1l
1453563_at	1.13154	NmrA-like family domain containing 1	Nmr1l
1453564_a_at	1.07836	vacuolar protein sorting 24 (yeast)	Vps24
1453565_at	1.04429	"NADH dehydrogenase (ubiquinone) 1, alpha/beta subcomplex, 1"	Ndufab1
1453566_at	0.99616	RIKEN cDNA 3200001K10	gene 3200001K10Rik
1453567_s_at	0.99863	RIKEN cDNA 2810441K11	gene 2810441K11Rik
1453568_at	0.94033	RIKEN cDNA 2310032F03	gene 2310032F03Rik
1453569_s_at	0.83646	MAP/microtubule affinity-regulating kinase 3	Mark3
1453570_x_at	1.02789	blocked early in transport 1 homolog (S. cerevisiae)-like	Bet1l
1453571_at	0.83806	DEP domain containing 6	Depdc6
1453572_a_at	0.97386	proteolipid protein 2	Plp2
1453573_at	1.56369	"histone1, H3d"	Hist1h3d
1453574_at	1.01410	"hemoglobin alpha, adult chain 1"	Hba-a1
1453575_at	0.98240	RIKEN cDNA 1200003I10	gene 1200003I10Rik
1453576_at	1.07848	Nipped-B homolog (Drosophila)	Nipbl
1453577_at	1.18501	RIKEN cDNA 2610018I03	gene 2610018I03Rik
1453578_at	0.92756	phosphotriesterase related	Pter
1453579_at	1.01663	PHD finger protein 12	Phf12
1453580_at	1.04506	RIKEN cDNA 5630401D24	gene 5630401D24Rik
1453581_at	0.96615	centrosomal protein 170	Cep170
1453582_at	1.79652	choline kinase alpha	Chka

1453583_at	1.25370	"zinc finger, ZZ-type with EF hand domain 1"	Zzef1	
1453584_at	0.99026	"serine/threonine kinase 36 (fused homolog, Drosophila)"		Stk36
1453585_at	0.94131	RIKEN cDNA 1600010M07 gene	1600010M07Rik	
1453586_at	0.98438	ectonucleoside triphosphate diphosphohydrolase 1		Entpd1
1453587_at	0.97886	gamma-glutamyltransferase 6	Ggt6	
1453588_at	1.07763	carbonic anhydrase 3	Car3	
1453589_a_at	0.97020	RIKEN cDNA 2610005L07 gene	2610005L07Rik	
1453590_at	0.99704	ADP-ribosylation factor-like 5B	Arl5b	
1453591_at	0.89007	RIKEN cDNA 5730437N04 gene	5730437N04Rik	
1453592_at	1.00463	leucine rich repeat containing 39	Lrrc39	
1453593_at	0.74324	RIKEN cDNA 1700110N18 gene	1700110N18Rik	
1453594_at	0.91711	RIKEN cDNA 4930451G09 gene	4930451G09Rik	
1453595_at	0.98854	RIKEN cDNA 2900064B18 gene	2900064B18Rik	
1453596_at	1.16852	inhibitor of DNA binding 2	Id2	
1453597_at	0.95762	RIKEN cDNA 4933425O20 gene	4933425O20Rik	
1453598_at	1.05107	RIKEN cDNA 3110082D06 gene	3110082D06Rik	
1453599_at	1.11972	tripartite motif-containing 71	Trim71	
1453600_at	0.94523	coiled-coil domain containing 18	Ccdc18	
1453601_at	1.09315	zinc finger protein 710	Zfp710	
1453602_at	0.91492	RIKEN cDNA 2810002I04 gene	2810002I04Rik	
1453603_at	0.97820	RIKEN cDNA 2700022O18 gene	2700022O18Rik	
1453604_a_at	0.97586	Hbs1-like (S. cerevisiae)	Hbs1l	
1453605_s_at	0.90443	coiled-coil domain containing 91	Ccdc91	
1453606_at	1.03289	RIKEN cDNA 4931403G20 gene	4931403G20Rik	
1453607_at	1.00334	microfibrillar-associated protein 3-like	Mfap3l	
1453608_at	1.48874	RIKEN cDNA 2310020J12 gene	2310020J12Rik	
1453609_s_at	1.07550	coiled-coil domain containing 9	Ccdc9	
1453610_at	0.90729	RIKEN cDNA 4931406B18 gene	4931406B18Rik	
1453611_at	1.05937	retinoblastoma binding protein 6	Rbbp6	
1453612_at	0.85454	NIMA (never in mitosis gene a)-related expressed kinase 1		Nek1
1453613_at	0.91363	RIKEN cDNA A130006I12 gene	A130006I12Rik	
1453614_a_at	0.88113	"nuclear factor, erythroid derived 2, like 3"	Nfe2l3	
1453615_at	0.97913	---	---	
1453616_at	0.92522	RIKEN cDNA 3830408C21 gene	3830408C21Rik	
1453617_at	0.91108	"zinc finger, RAN-binding domain containing 3"	Zranb3	
1453618_at	0.84618	RIKEN cDNA 4933424N20 gene	4933424N20Rik	
1453619_at	0.88645	EF hand calcium binding domain 1	Efcab1	
1453620_at	0.80131	RIKEN cDNA 3010022N24 gene	3010022N24Rik	
1453621_at	1.07892	THO complex 7 homolog (Drosophila) /// similar to Ngg1 interacting factor 3 like 1 binding protein 1 isoform 1	Thoc7 /// LOC670839	
1453622_s_at	0.84531	myeloid/lymphoid or mixed lineage-leukemia translocation to 3 homolog (Drosophila)	Mllt3	
1453623_a_at	1.08993	RAD23a homolog (S. cerevisiae)	Rad23a	

1453624_at	1.18373	RIKEN cDNA C430046K18 gene	C430046K18Rik
1453625_at	1.01791	RIKEN cDNA 2210407P21 gene	2210407P21Rik
1453626_at	1.10904	RIKEN cDNA 3930402G23 gene	3930402G23Rik
1453627_at	1.20821	RIKEN cDNA 2610020P09 gene	2610020P09Rik
1453628_s_at	0.86883	leucine rich repeat containing 2	Lrrc2
1453629_at	0.99495	RIKEN cDNA 4930478L05 gene	4930478L05Rik
1453630_at	0.94574	UDP-N-acetyl-alpha-D-galactosamine:polypeptide N-acetylgalactosaminyltransferase 14	Galnt14
1453631_at	0.95671	syntaxin 8	Stx8
1453632_at	0.93856	RIKEN cDNA 4930538K18 gene	4930538K18Rik
1453633_a_at	1.02415	ring finger protein 41	Rnf41
1453634_a_at	1.02234	endoplasmic reticulum protein 29	Erp29
1453635_at	1.03864	RIKEN cDNA 4932416N17 gene	4932416N17Rik
1453636_at	1.40928	polycomb group ring finger 5	Pcgf5
1453637_at	1.03752	RIKEN cDNA 5430410E06 gene	5430410E06Rik
1453638_at	0.92262	RIKEN cDNA 9030420J04 gene	9030420J04Rik
1453639_s_at	0.75156	RIKEN cDNA 0610037D15 gene	0610037D15Rik
1453640_at	0.98915	RIKEN cDNA 2210019G11 gene	2210019G11Rik
1453641_at	1.15372	breast carcinoma amplified sequence 1	Bcas1
1453642_at	0.97959	RIKEN cDNA 6530411M01 gene	6530411M01Rik
1453643_at	1.17343	RIKEN cDNA 1700001L19 gene	1700001L19Rik
1453644_at	0.84109	odorant binding protein 1a	Obp1a
1453645_at	0.85711	RIKEN cDNA 2700046A07 gene	2700046A07Rik
1453646_at	1.37022	RIKEN cDNA 3110005L21 gene	3110005L21Rik
1453647_at	0.95618	RIKEN cDNA 9530059J11 gene	9530059J11Rik
1453648_at	0.98483	sushi domain containing 3	Susd3
1453649_at	0.90969	RIKEN cDNA C030039L03 gene	C030039L03Rik
1453650_at	0.96087	RIKEN cDNA 4930413F20 gene	4930413F20Rik
1453651_a_at	1.06923	brain protein 44-like	Brp44l
1453652_at	0.89415	RIKEN cDNA 4933400F21 gene	4933400F21Rik
1453653_at	1.02239	RIKEN cDNA 6330576A10 gene	6330576A10Rik
1453654_at	0.87816	"CCR4-NOT transcription complex, subunit 10"	Cnot10
1453655_at	0.95088	RIKEN cDNA 2900005J15 gene	2900005J15Rik
1453656_a_at	0.98090	RIKEN cDNA 4933431G14 gene	4933431G14Rik
1453657_at	0.88944	RIKEN cDNA 2310065F04 gene	2310065F04Rik
1453658_at	0.94664	RIKEN cDNA 5830477G23 gene	5830477G23Rik
1453659_at	1.03042	centromere protein O	Cenpo
1453660_at	1.10983	RIKEN cDNA 5430420F09 gene	5430420F09Rik
1453661_at	0.87505	RIKEN cDNA 9130403I23 gene	9130403I23Rik
1453662_at	1.05811	hypothetical LOC77413	LOC77413
1453663_at	0.99304	RIKEN cDNA 4932442E05 gene	4932442E05Rik
1453664_at	0.73886	RIKEN cDNA 9130017K11 gene	9130017K11Rik
1453665_at	1.10324	RIKEN cDNA 4930529M08 gene	4930529M08Rik

1453666_at	0.98159 ring finger protein 32	Rnf32	
1453667_at	1.02238 RIKEN cDNA 1700065D16	gene	1700065D16Rik
1453668_at	0.74776 RIKEN cDNA 3110052M02	gene	3110052M02Rik
1453669_at	0.97616 RIKEN cDNA 4930578C19	gene	4930578C19Rik
1453670_at	1.01511 ubiquitin specific peptidase 50	Usp50	
1453671_at	0.95552 RIKEN cDNA 9430041J12	gene	9430041J12Rik
1453672_at	0.95940 RIKEN cDNA 2810038L03	gene	2810038L03Rik
1453673_at	0.95389 ATPase type 13A4	Atp13a4	
1453674_at	1.19439 "glucosaminyl (N-acetyl) transferase 2, I-branching enzyme"	Gcnt2	
1453675_at	0.88327 "solute carrier family 16 (monocarboxylic acid transporters), member 10"	Slc16a10	
1453676_at	0.90103 RIKEN cDNA A930041D05	gene	A930041D05Rik
1453677_a_at	0.96273 "Der1-like domain family, member 3"	Derl3	
1453678_at	0.88780 methyl-CpG binding domain protein 1	Mbd1	
1453679_at	0.99580 RIKEN cDNA 1700100I10	gene	1700100I10Rik
1453680_at	1.10158 RIKEN cDNA 8030456M14	gene	8030456M14Rik
1453681_at	1.35766 ATPase inhibitory factor 1	Atpif1	
1453682_at	0.87288 "GA repeat binding protein, beta 2"	Gabpb2	
1453683_a_at	1.01403 centrosomal protein 55	Cep55	
1453684_s_at	1.02805 RIKEN cDNA 2610312B22	gene	2610312B22Rik
1453685_at	0.89777 nudix (nucleoside diphosphate linked moiety X)-type motif 7	Nudt7	
1453686_x_at	1.03009 nephronophthisis 1 (juvenile) homolog (human)	Nphp1	
1453687_at	0.87099 "solute carrier family 4 (anion exchanger), member 3"	Slc4a3	
1453688_at	1.18885 "CWF19-like 2, cell cycle control (S. pombe)"	Cwf19l2	
1453689_at	0.93941 "Fanconi anemia, complementation group E"	Fance	
1453690_at	0.93311 "membrane protein, palmitoylated 7 (MAGUK p55 subfamily member 7)"	Mpp7	
1453691_at	0.92625 RIKEN cDNA 4921528I07	gene	4921528I07Rik
1453692_at	0.89394 RIKEN cDNA 4930581F22	gene	4930581F22Rik
1453693_at	0.97993 WD repeat domain 20	Wdr20	
1453694_at	1.05996 RIKEN cDNA 4631402N15	gene	4631402N15Rik
1453695_at	0.98580 RIKEN cDNA B430211C08	gene	B430211C08Rik
1453696_at	0.95796 RIKEN cDNA 4921508A21	gene	4921508A21Rik
1453697_at	0.94842 RIKEN cDNA 6530402F18	gene	6530402F18Rik
1453698_at	1.02080 RIKEN cDNA 6030451C04	gene	6030451C04Rik
1453699_at	0.88020 RIKEN cDNA 4930448D08	gene	4930448D08Rik
1453700_s_at	1.00310 "RIKEN cDNA 4933403O03	gene /// similar to double homeobox, 4"	4933403O03Rik /// LOC245263
1453701_at	1.06112 RIKEN cDNA 4921515E04	gene	4921515E04Rik
1453702_at	0.93487 coiled-coil domain containing 105	Ccdc105	
1453703_at	0.99203 "dynein, axonemal, heavy polypeptide 17"	Dnah17	
1453704_at	1.08962 --- ---		
1453705_at	0.97095 RIKEN cDNA B230110C06	gene	B230110C06Rik

1453706_at	1.32041	RIKEN cDNA 2900042A17 gene	2900042A17Rik
1453707_at	0.92422	RIKEN cDNA 4930480G23 gene	4930480G23Rik
1453708_a_at	1.07985	glutathione S-transferase omega 2	Gsto2
1453709_at	1.00417	RIKEN cDNA 3426406O18 gene	3426406O18Rik
1453710_at	0.88949	transmembrane protein 116	Tmem116
1453711_at	1.05734	"R-spondin family, member 4"	Rspo4
1453712_a_at	0.93231	mitogen activated protein kinase kinase 5	Map2k5
1453713_s_at	1.07807	RIKEN cDNA 4930546H06 gene	4930546H06Rik
1453714_a_at	0.94531	shroom family member 3	Shroom3
1453715_at	0.95109	synaptic vesicle glycoprotein 2c	Sv2c
1453716_at	0.92917	RIKEN cDNA 4930480K23 gene	4930480K23Rik
1453717_s_at	1.28161	"tubulin tyrosine ligase-like family, member 10"	Ttl10
1453718_at	0.93050	BCL2-like 12 (proline rich)	Bcl2l12
1453719_at	0.95945	RIKEN cDNA 4930506C21 gene	4930506C21Rik
1453720_at	0.91332	ring finger protein 157	Rnf157
1453721_a_at	1.05934	"solute carrier family 31, member 2"	Slc31a2
1453722_s_at	1.05588	"splicing factor, arginine/serine-rich 1 (ASF/SF2)"	Sfrs1
1453723_x_at	0.98462	"ubiquitin B /// gene model 1821, (NCBI)"	Ubb /// Gm1821
1453724_a_at	0.87068	"serine (or cysteine) peptidase inhibitor, clade F, member 1"	Serpinf1
1453725_a_at	1.09629	mitochondrial ribosomal protein S7	Mrps7
1453726_s_at	1.02641	RIKEN cDNA 2810407C02 gene	2810407C02Rik
1453727_at	0.98868	RIKEN cDNA 2610101J03 gene	2610101J03Rik
1453728_a_at	0.99921	mitochondrial ribosomal protein S17	Mrps17
1453729_a_at	1.04501	ribosomal protein L37	Rpl37
1453730_at	0.81364	sterile alpha motif domain containing 8	Samd8
1453731_a_at	1.02850	transmembrane protein 77	Tmem77
1453732_at	0.93792	RIKEN cDNA 6230416A05 gene	6230416A05Rik
1453733_a_at	1.05435	RIKEN cDNA 4933434E20 gene	4933434E20Rik
1453734_at	0.89844	alpha thalassemia/mental retardation syndrome X-linked homolog (human)	Atrx
1453735_at	0.92910	RIKEN cDNA 5730455P16 gene	5730455P16Rik
1453736_s_at	0.80557	RIKEN cDNA B230219D22 gene	B230219D22Rik
1453737_at	1.08881	RIKEN cDNA 5730509C05 gene	5730509C05Rik
1453738_at	0.89192	RIKEN cDNA C330018D20 gene	C330018D20Rik
1453739_at	1.01961	transmembrane protein 126B	Tmem126b
1453740_a_at	0.94419	cyclin L2	Ccnl2
1453741_x_at	1.07044	angel homolog 2 (Drosophila)	Angel2
1453742_at	1.02933	vacuolar protein sorting 33A (yeast)	Vps33a
1453743_x_at	1.01571	PHD finger protein 14	Phf14
1453744_a_at	0.91120	ankyrin repeat domain 40	Ankrd40
1453745_at	1.04963	RIKEN cDNA 2700038G22 gene	2700038G22Rik
1453746_at	0.98017	formin binding protein 1	Fnbp1
1453747_at	0.91084	RIKEN cDNA 2810021J22 gene	2810021J22Rik
1453748_a_at	1.21543	kinesin family member 23	Kif23

1453749_at	0.84893	RIKEN cDNA 2610507I01 gene	2610507I01Rik	
1453750_x_at	0.91862	"phosphatidylinositol transfer protein, cytoplasmic 1"	Pitpnc1	
1453751_at	0.94128	DEAH (Asp-Glu-Ala-His) box polypeptide 38	Dhx38	
1453752_at	1.04910	ribosomal protein L17	Rpl17	
1453753_at	2.17948	denticleless homolog (Drosophila)	Dtl	
1453754_at	1.04040	S100P binding protein	S100pbp	
1453755_at	1.19752	U7 snRNP-specific Sm-like protein LSM11	Lsm11	
1453756_at	0.86739	RIKEN cDNA 2900075N08 gene	2900075N08Rik	
1453757_at	0.96053	hect domain and RLD 5	Herc5	
1453758_at	1.10860	RIKEN cDNA 2810027O19 gene	2810027O19Rik	
1453759_at	0.95407	NOL1/NOP2/Sun domain family 2	Nsun2	
1453760_at	1.12304	mesoderm induction early response 1 homolog (Xenopus laevis)	Mier1	
1453761_at	1.08268	PHD finger protein 6	Phf6	
1453762_at	0.97557	vacuolar protein sorting 26 homolog B (yeast)	Vps26b	
1453763_at	0.91804	thioredoxin domain containing 11	Txndc11	
1453764_at	0.96961	RIKEN cDNA 3110021N24 gene	3110021N24Rik	
1453765_at	1.01609	RIKEN cDNA A330106J01 gene	A330106J01Rik	
1453766_a_at	1.08829	RIKEN cDNA 4931407K02 gene	4931407K02Rik	
1453767_a_at	0.99565	"5',3'-nucleotidase, mitochondrial"	Nt5m	
1453768_a_at	1.18968	RIKEN cDNA 5430432M24 gene	5430432M24Rik	
1453769_at	0.98091	cytoskeleton associated protein 2-like	Ckap2l	
1453770_at	1.01294	carboxypeptidase A4	Cpa4	
1453771_at	1.00506	"GULP, engulfment adaptor PTB domain containing 1"	Gulp1	
1453772_at	0.95953	synaptogyrin 1	Syngr1	
1453773_at	0.97469	RIKEN cDNA 4931406I20 gene	4931406I20Rik	
1453774_at	0.98546	RIKEN cDNA 2810002O09 gene	2810002O09Rik	
1453775_at	1.14328	RIKEN cDNA 4921505C17 gene	4921505C17Rik	
1453776_at	1.45656	RIKEN cDNA 5730407K14 gene	5730407K14Rik	
1453777_a_at	1.09671	N-deacetylase/N-sulfotransferase (heparan glucosaminyl) 3	Ndst3	
1453778_at	0.92704	RIKEN cDNA 2810407C02 gene	2810407C02Rik	
1453779_at	1.14756	RIKEN cDNA 1700020L24 gene	1700020L24Rik	
1453780_at	1.00319	---	---	
1453781_at	0.84022	---	---	
1453782_at	1.14033	RIKEN cDNA 3021401C12 gene	3021401C12Rik	
1453783_at	0.96246	purine rich element binding protein A	Pura	
1453784_at	1.03995	integrin-linked kinase-associated serine/threonine phosphatase 2C	Ilkap	
1453785_at	1.01941	tafazzin	Taz	
1453786_at	0.95296	---	---	
1453787_at	0.85074	thioredoxin domain containing 13	Txndc13	
1453788_at	0.99118	---	---	
1453789_at	1.02627	RIKEN cDNA 4933440N22 gene	4933440N22Rik	
1453790_at	1.02871	RIKEN cDNA 5730403B10 gene	5730403B10Rik	
1453791_at	1.09604	RIKEN cDNA 2810449C10 gene	2810449C10Rik	

1453792_at	1.03035	RIKEN cDNA A930003G23 gene	A930003G23Rik
1453793_at	0.90041	RIKEN cDNA 1700026J12 gene	1700026J12Rik
1453794_at	1.06244	fer-1-like 4 (C. elegans) Fer114	
1453795_at	0.95390	fumarylacetoacetate hydrolase domain containing 2A	Fahd2a
1453796_a_at	1.00399	ERGIC and golgi 2 Ergic2	
1453797_at	0.89702	"phosphatase, orphan 2" Phospho2	
1453798_at	0.83437	coiled-coil domain containing 93	Ccdc93
1453799_at	1.00854	RIKEN cDNA 9430038I01 gene	9430038I01Rik
1453800_at	1.06746	RIKEN cDNA 2500002L14 gene	2500002L14Rik
1453801_at	0.95179	thioesterase superfamily member 5	Them5
1453802_at	0.88646	RIKEN cDNA 3732412D22 gene	3732412D22Rik
1453803_at	0.93558	cytotoxic granule-associated RNA binding protein 1	Tia1
1453804_a_at	1.03008	"origin recognition complex, subunit 4-like (S. cerevisiae)"	Orc4l
1453805_at	0.97687	"Fucosidase, alpha-L- 1, tissue"	Fuca1
1453806_at	0.94238	"NADH dehydrogenase (ubiquinone) 1 beta subcomplex, 2"	Ndufb2
1453807_at	0.98889	RIKEN cDNA 6330563C09 gene	6330563C09Rik
1453808_at	0.83738	RIKEN cDNA 2210416O15 gene	2210416O15Rik
1453809_at	1.08975	Ly6/Plaur domain containing 5	Lypd5
1453810_at	1.37426	aftiphilin	Aftph
1453811_at	0.93275	heat shock protein 9A	Hspa9a
1453812_at	0.97491	janus kinase and microtubule interacting protein 2	Jakmip2
1453813_at	1.00270	CD200 receptor 3	Cd200r3
1453814_at	1.04555	B-cell CLL/lymphoma 11A (zinc finger protein)	Bcl11a
1453815_at	0.88599	spindle pole body component 25 homolog (S. cerevisiae)	Spbc25
1453816_at	0.97934	RIKEN cDNA 5730427N09 gene	5730427N09Rik
1453817_at	0.96874	"ATP-binding cassette, sub-family A (ABC1), member 6"	Abca6
1453818_a_at	0.94353	RIKEN cDNA 9030025P20 gene	9030025P20Rik
1453819_x_at	1.04360	syntaxin 18	Stx18
1453820_at	0.89312	RIKEN cDNA A930005F02 gene /// suppressor of hairy wing homolog 4 (Drosophila) A930005F02Rik /// Suhw4	
1453821_at	1.07730	HemK methyltransferase family member 2	Hemk2
1453822_at	1.10857	RIKEN cDNA 1700010H15 gene	1700010H15Rik
1453823_a_at	1.01223	RIKEN cDNA A930028L21 gene	A930028L21Rik
1453824_at	1.04472	RIKEN cDNA A930028L21 gene	A930028L21Rik
1453825_at	1.01711	"Mammary gland RCB-0526 Jyg-MC(A) cDNA, RIKEN full-length enriched library, clone:G830038F10 product:unclassifiable, full insert sequence" ---	
1453826_at	0.70879	RIKEN cDNA 1810008K04 gene	1810008K04Rik
1453827_at	0.91437	RIKEN cDNA 1110035H17 gene	1110035H17Rik
1453828_at	0.89194	RIKEN cDNA 4932422M17 gene	4932422M17Rik
1453829_at	0.92603	RIKEN cDNA 2310007J06 gene	2310007J06Rik
1453830_at	0.93040	---	---
1453831_at	1.11686	transmembrane protein 138	Tmem138
1453832_at	1.00429	RIKEN cDNA 5330430P22 gene	5330430P22Rik

1453833_a_at 1.09668 ribonuclease H1 Rnaseh1
1453834_at 0.93272 RIKEN cDNA 1500002C15 gene 1500002C15Rik
1453835_at 1.01742 RIKEN cDNA 4930505M18 gene 4930505M18Rik
1453836_a_at 1.08776 monoglyceride lipase Mgl1
1453837_at 1.01263 RIKEN cDNA 6330500D04 gene 6330500D04Rik
1453838_at 1.04426 RIKEN cDNA 4930471G03 gene 4930471G03Rik
1453839_a_at 1.01778 peptidase inhibitor 16 Pi16
1453840_at 1.56286 "poly A binding protein, cytoplasmic 1" Pabpc1
1453841_at 0.94699 RIKEN cDNA 2310050P20 gene 2310050P20Rik
1453842_at 0.83378 RIKEN cDNA 4930448K12 gene 4930448K12Rik
1453843_at 1.02154 THAP domain containing 6 Thap6
1453844_at 0.95915 chitinase 1 (chitotriosidase) Chit1
1453845_at 1.04487 RIKEN cDNA 4733401D01 gene 4733401D01Rik
1453846_at 0.96207 RIKEN cDNA 2810013C04 gene 2810013C04Rik
1453847_at 1.02830 RIKEN cDNA A930037O16 gene A930037O16Rik
1453848_s_at 0.89131 "zinc finger, BED domain containing 3" Zbed3
1453849_s_at 1.03881 heterogeneous nuclear ribonucleoprotein A/B Hnrpab
1453850_at 1.05733 RIKEN cDNA 1500002I01 gene 1500002I01Rik
1453851_a_at 0.94868 growth arrest and DNA-damage-inducible 45 gamma Gadd45g
1453852_at 1.28372 RIKEN cDNA 8430408E17 gene 8430408E17Rik
1453853_a_at 1.01325 Rho guanine nucleotide exchange factor (GEF) 12 Arhgef12
1453854_at 1.07970 protein prenyltransferase alpha subunit repeat containing 1 /// similar to protein prenyltransferase alpha subunit repeat containing 1 Ptar1 /// LOC676290
1453855_at 0.89970 RIKEN cDNA 1810057P16 gene 1810057P16Rik
1453856_at 0.87238 BTB (POZ) domain containing 4 Btbd4
1453857_at 1.00988 ring finger protein 180 Rnf180
1453858_at 1.09124 "solute carrier family 35 (UDP-galactose transporter), member 2" Slc35a2
1453859_at 0.90895 RIKEN cDNA 2410003P15 gene 2410003P15Rik
1453860_s_at 0.75313 "nuclear receptor subfamily 3, group C, member 1" Nr3c1
1453861_at 0.90580 RIKEN cDNA C030014K22 gene C030014K22Rik
1453862_at 0.96126 RIKEN cDNA 4930465K10 gene 4930465K10Rik
1453863_at 1.10957 RIKEN cDNA 3300002P09 gene 3300002P09Rik
1453864_at 1.18636 retinol dehydrogenase 14 (all-trans and 9-cis) Rdh14
1453865_a_at 1.00057 OTU domain containing 5 Otud5
1453866_a_at 1.21700 Kell blood group precursor (McLeod phenotype) homolog Xkh
1453867_at 0.98189 F-box and WD-40 domain protein 2 Fbxw2
1453868_at 1.02534 coiled-coil domain containing 11 Ccdc11
1453869_at 1.10536 hypothetical protein LOC328277 LOC328277
1453870_at 0.93269 RIKEN cDNA 1810018F18 gene 1810018F18Rik
1453871_at 0.99340 "solute carrier family 6 (neurotransmitter transporter, GABA), member 11" Slc6a11
1453872_at 0.96557 doublesex and mab-3 related transcription factor like family C2 Dmrtd2
1453873_at 0.94553 RIKEN cDNA 4933421H06 gene 4933421H06Rik

1453874_at 0.95655 RIKEN cDNA 4933401B06 gene 4933401B06Rik
 1453875_at 0.99293 "solute carrier family 7 (cationic amino acid transporter, y+ system), member 15" Slc7a15
 1453876_at 1.03156 hypothetical LOC621121 /// hypothetical protein LOC631862 /// RIKEN cDNA 1700120B22 gene LOC621121 /// LOC631862 /// 1700120B22Rik
 1453877_at 0.87622 RIKEN cDNA 4930467D21 gene 4930467D21Rik
 1453878_at 1.07874 ubiquitin protein ligase E3 component n-recognin 2 Ubr2
 1453879_at 1.08832 RIKEN cDNA 1700016H13 gene 1700016H13Rik
 1453880_s_at 0.90497 RIKEN cDNA 1700041C02 gene 1700041C02Rik
 1453881_x_at 1.05096 centrin 1 Cetn1
 1453882_at 0.99178 RIKEN cDNA 1700034F02 gene 1700034F02Rik
 1453883_at 0.90434 RIKEN cDNA 4933425B07 gene 4933425B07Rik
 1453884_at 0.95523 RIKEN cDNA 4930402K13 gene 4930402K13Rik
 1453885_at 1.02096 "solute carrier family 24 (sodium/potassium/calcium exchanger), member 2" Slc24a2
 1453886_a_at 0.99358 "solute carrier family 25 (mitochondrial carrier, phosphate carrier), member 26" Slc25a26
 1453887_a_at 1.06715 T-cell lymphoma invasion and metastasis 1 Tiam1
 1453888_at 0.96438 copine IV Cpne4
 1453889_at 1.11128 RIKEN cDNA 4930461C15 gene 4930461C15Rik
 1453890_at 0.90684 --- ---
 1453891_at 0.99144 polymerase I and transcript release factor Ptrf
 1453892_at 1.11351 RIKEN cDNA A930007I19 gene A930007I19Rik
 1453893_at 1.00534 RIKEN cDNA 4933412O06 gene 4933412O06Rik
 1453894_at 1.07839 RIKEN cDNA 4933434C23 gene 4933434C23Rik
 1453895_at 1.12991 RIKEN cDNA C330026N13 gene C330026N13Rik
 1453896_at 1.01610 PDZ domain containing 2 Pdzd2
 1453897_at 1.09401 RIKEN cDNA C030014A21 gene C030014A21Rik
 1453898_at 1.08623 integrin beta 1 binding protein 3 Itgb1bp3
 1453899_at 0.93029 --- ---
 1453900_at 0.82365 RIKEN cDNA 1700018A14 gene 1700018A14Rik
 1453901_at 0.88043 "aldolase 1, A isoform, pseudogene 2" Aldoa-ps2
 1453902_at 1.08877 eukaryotic translation initiation factor 2 alpha kinase 1 Eif2ak1
 1453903_at 0.89907 RIKEN cDNA 4930503B20 gene 4930503B20Rik
 1453904_at 0.97338 RIKEN cDNA 4930528G09 gene 4930528G09Rik
 1453905_at 0.95136 RIKEN cDNA 5033415L01 gene 5033415L01Rik
 1453906_at 1.28785 thyroid hormone receptor associated protein 2 Thrap2
 1453907_at 1.02904 RIKEN cDNA C030041B07 gene C030041B07Rik
 1453908_at 1.09079 "protein tyrosine phosphatase, receptor type, B" Ptprb
 1453909_at 0.88695 RIKEN cDNA 4930505N22 gene 4930505N22Rik
 1453910_at 0.96058 RIKEN cDNA 4930500A05 gene 4930500A05Rik
 1453911_at 0.87928 --- ---
 1453912_at 0.94468 zinc finger protein 558 Zfp558

1453913_a_at 1.09819 "transporter 2, ATP-binding cassette, sub-family B (MDR/TAP)" Tap2
 1453914_at 1.01141 heat shock protein 1 Hspb1
 1453915_a_at 0.95364 "solute carrier family 37 (glycerol-3-phosphate transporter), member 3"
 Slc37a3
 1453916_at 1.14887 RIKEN cDNA 4933431K23 gene 4933431K23Rik
 1453917_at 0.98526 interferon induced transmembrane protein 7 Ifitm7
 1453918_at 0.91393 RIKEN cDNA 4933426F18 gene 4933426F18Rik
 1453919_at 1.03131 RIKEN cDNA 2610028H24 gene 2610028H24Rik
 1453920_a_at 1.08084 motile sperm domain containing 2 Mospd2
 1453921_at 0.89640 RIKEN cDNA 4930429F11 gene 4930429F11Rik
 1453922_at 1.09078 RIKEN cDNA 4921539H07 gene 4921539H07Rik
 1453923_at 0.93804 coiled-coil domain containing 83 Ccdc83
 1453924_a_at 0.93643 prostaglandin F receptor Ptgfr
 1453925_at 0.96935 RIKEN cDNA 4930429F24 gene 4930429F24Rik
 1453926_at 1.09443 RAD54 like (*S. cerevisiae*) Rad54l
 1453927_at 0.98832 mitochondrial ribosomal protein L47 Mrpl47
 1453928_a_at 1.06719 Sjogren syndrome antigen B Ssb
 1453929_at 0.97738 ring finger protein 24 Rnf24
 1453930_at 1.05165 RIKEN cDNA 1110015M06 gene 1110015M06Rik
 1453931_at 0.95280 "procollagen, type XIV, alpha 1" Col14a1
 1453932_at 0.76450 RIKEN cDNA 4931433A01 gene 4931433A01Rik
 1453933_at 0.97888 RIKEN cDNA 4933435E02 gene 4933435E02Rik
 1453934_at 0.97312 RIKEN cDNA 4930535C22 gene 4930535C22Rik
 1453935_a_at 0.94223 coiled-coil domain containing 7 Ccdc7
 1453936_at 0.93477 --- ---
 1453937_at 1.00115 RIKEN cDNA 4933407G14 gene 4933407G14Rik
 1453938_at 1.02880 RIKEN cDNA 9330198N18 gene 9330198N18Rik
 1453939_x_at 1.17958 hypothetical protein LOC677168 LOC677168
 1453940_at 1.08379 RIKEN cDNA 2810404M03 gene 2810404M03Rik
 1453941_at 1.01452 RIKEN cDNA 4930579P08 gene 4930579P08Rik
 1453942_at 0.97301 RIKEN cDNA 1700123I01 gene 1700123I01Rik
 1453943_a_at 0.88354 "acid phosphatase, prostate" Acpp
 1453944_at 0.96426 RIKEN cDNA 5430435K18 gene 5430435K18Rik
 1453945_at 0.98801 RIKEN cDNA 2310005E17 gene 2310005E17Rik
 1453946_a_at 0.97439 serologically defined colon cancer antigen 8 Sdccag8
 1453947_at 1.12974 kinesin family member 24 Kif24
 1453948_at 1.00455 RIKEN cDNA 4930529K09 gene 4930529K09Rik
 1453949_s_at 0.98993 lysophospholipase 1 Lypl1
 1453950_a_at 0.82695 X-ray repair complementing defective repair in Chinese hamster cells 2 Xrcc2
 1453951_a_at 0.90914 "DNA segment, Chr 19, ERATO Doi 652, expressed /// similar to DNA
 segment, Chr 19, ERATO Doi 652, expressed" D19Ertd652e /// LOC677118
 1453952_at 0.87842 RIKEN cDNA 4930572G02 gene 4930572G02Rik
 1453953_at 1.03136 RIKEN cDNA 9130015A21 gene 9130015A21Rik

1453954_a_at	0.99817	mitochondrial ribosomal protein S5	Mrps5	
1453955_a_at	0.84452	RIKEN cDNA 4930432H15	gene	4930432H15Rik
1453956_a_at	0.91902	PFTAIRE protein kinase 1	Pftk1	
1453957_a_at	1.01823	insulin-like growth factor 2 mRNA binding protein 3	Igf2bp3	
1453958_at	0.95888	RIKEN cDNA 2610203C22	gene	2610203C22Rik
1453959_at	0.93213	RIKEN cDNA 1700065O13	gene	1700065O13Rik
1453960_a_at	0.98599	"capping protein (actin filament) muscle Z-line, beta"	Capzb	
1453961_a_at	0.89567	motile sperm domain containing 3	Mospd3	
1453962_at	0.99090	profilin 3	Pfn3	
1453963_at	1.17923	RIKEN cDNA 1600014C23	gene	1600014C23Rik
1453964_at	0.87511	RIKEN cDNA 1700021A07	gene	1700021A07Rik
1453965_at	1.06085	RIKEN cDNA 4930432O09	gene	4930432O09Rik
1453966_at	0.95064	RIKEN cDNA 2610008G14	gene	2610008G14Rik
1453967_at	0.96472	RIKEN cDNA 1700093J21	gene	1700093J21Rik
1453968_at	1.12432	RIKEN cDNA 4833439F03	gene	4833439F03Rik
1453969_at	0.95405	prolactin-like protein C 4	Prlpc4	
1453970_at	0.99341	RIKEN cDNA 4833417A11	gene	4833417A11Rik
1453971_at	0.93547	RIKEN cDNA 2610015P09	gene	2610015P09Rik
1453972_x_at	1.00205	peroxisome proliferator activated receptor binding protein	Pparbp	
1453973_s_at	0.95454	RIKEN cDNA 2900057H02	gene	2900057H02Rik
1453974_at	0.92038	RIKEN cDNA 1700023A20	gene	1700023A20Rik
1453975_a_at	1.13441	RIKEN cDNA 1700029M20	gene	1700029M20Rik
1453976_at	1.81780	RIKEN cDNA 4432414F05	gene	4432414F05Rik
1453977_at	0.98105	exocyst complex component 4	Exoc4	
1453978_at	0.97980	RIKEN cDNA 4921522H14	gene	4921522H14Rik
1453979_at	0.95539	RNA binding motif protein 12B	Rbm12b	
1453980_at	0.98145	RIKEN cDNA 5330434G04	gene	5330434G04Rik
1453981_at	1.02376	RIKEN cDNA 2900009I07	gene	2900009I07Rik
1453982_at	0.92150	transformed mouse 3T3 cell double minute 4	Mdm4	
1453983_a_at	1.10080	methyltransferase 10 domain containing	Mett10d	
1453984_at	1.05240	"phosphatidylinositol 3-kinase, regulatory subunit, polypeptide 2 (p85 beta) /// interferon gamma inducible protein 30"	Pik3r2 /// Ifi30	
1453985_at	1.04239	RIKEN cDNA 0610007P08	gene	0610007P08Rik
1453986_at	0.84483	"ligase III, DNA, ATP-dependent"	Lig3	
1453987_at	1.07173	BTB (POZ) domain containing 5	Btbd5	
1453988_a_at	1.05326	insulin degrading enzyme	Ide	
1453989_at	1.09484	syntaxin binding protein 4	Stxbp4	
1453990_at	1.05375	RIKEN cDNA 4930554G24	gene	4930554G24Rik
1453991_at	0.91572	centromere protein O	Cenpo	
1453992_at	1.00471	zona pellucida binding protein	Zbp	
1453993_a_at	1.02501	"BCL2/adenovirus E1B interacting protein 1, NIP2"	Bnip2	
1453994_at	1.05189	RIKEN cDNA C230094A16	gene	C230094A16Rik
1453995_a_at	0.95874	HpaII tiny fragments locus 9c	Htf9c	

1453996_a_at 0.94809 cathepsin 3 Cts3
1453997_a_at 1.02757 nestin Nes
1453998_at 1.04347 stromal membrane-associated protein 1 Smap1
1453999_at 1.09624 similar to Nucleolar preribosomal-associated protein 1 LOC670473
1454000_s_at 0.84539 RIKEN cDNA 4930578N16 gene 4930578N16Rik
1454001_at 0.92340 --- ---
1454002_at 1.01002 --- ---
1454003_at 1.01170 AFG3(ATPase family gene 3)-like 2 (yeast) Afg3l2
1454004_at 0.89172 RIKEN cDNA 5033414K04 gene 5033414K04Rik
1454005_at 0.84632 flavin containing monooxygenase 2 Fmo2
1454006_a_at 0.99822 UBX domain containing 6 Ubxd6
1454007_a_at 1.00767 zinc finger protein 142 Zfp142
1454008_at 0.94841 protein arginine N-methyltransferase 3 Prmt3
1454009_at 1.03330 RIKEN cDNA 1700008O03 gene 1700008O03Rik
1454010_a_at 1.02484 RIKEN cDNA 1700025E21 gene 1700025E21Rik
1454011_a_at 0.99263 replication protein A2 Rpa2
1454012_a_at 0.81688 "poly (ADP-ribose) polymerase family, member 6" Parp6
1454013_at 0.94277 RIKEN cDNA 1810062O18 gene 1810062O18Rik
1454014_a_at 0.86137 McKusick-Kaufman syndrome protein Mkks
1454015_a_at 1.41384 cadherin 13 Cdh13
1454016_at 0.95242 --- ---
1454017_at 0.93548 RIKEN cDNA 4921509O07 gene 4921509O07Rik
1454018_at 1.06747 tousled-like kinase 2 (Arabidopsis) Tlk2
1454019_at 0.83839 --- ---
1454020_at 0.94043 RIKEN cDNA 4930554H23 gene 4930554H23Rik
1454021_a_at 1.09077 exosome component 10 Exosc10
1454022_at 1.20602 Eph receptor B2 Ephb2
1454023_a_at 0.83172 "DNA segment, Chr 1, Brigham & Women's Genetics 1363 expressed"
D1Bwg1363e
1454024_at 1.09833 leucine rich repeat containing 9 Lrrc9
1454025_at 0.98670 D-aspartate oxidase Ddo
1454026_a_at 1.10615 regulator of G-protein signaling 3 Rgs3
1454027_at 0.88586 RIKEN cDNA 4930448N21 gene 4930448N21Rik
1454028_at 0.96325 RIKEN cDNA 4931402H11 gene 4931402H11Rik
1454029_at 1.05108 RIKEN cDNA 4930431F12 gene 4930431F12Rik
1454030_at 1.63188 "SAPS domain family, member 3" Saps3
1454031_at 1.09172 RIKEN cDNA 2310029O18 gene 2310029O18Rik
1454032_at 0.88431 neuropilin (NRP) and tolloid (TLL)-like 2 Neto2
1454033_at 1.01883 RIKEN cDNA 1700007B13 gene 1700007B13Rik
1454034_a_at 0.98806 ubiquitin specific peptidase 21 Usp21
1454035_at 1.02456 RIKEN cDNA 3110009E22 gene 3110009E22Rik
1454036_a_at 1.14284 ubiquitin specific peptidase 15 Usp15
1454037_a_at 0.96846 FMS-like tyrosine kinase 1 Flt1

1454038_at	0.91944	RIKEN cDNA 4833403I15 gene	4833403I15Rik
1454039_at	0.99523	RIKEN cDNA 4921517D16 gene	4921517D16Rik
1454040_at	1.07003	RIKEN cDNA 5730591J02 gene	5730591J02Rik
1454041_at	0.89616	"DNA segment, Chr 14, ERATO Doi 581, expressed"	D14ErtD581e
1454042_a_at	1.08849	serine/arginine-rich protein specific kinase 1	Srpk1
1454043_a_at	0.95038	"potassium voltage-gated channel, shaker-related subfamily, beta member 1"	Kcnab1
1454044_a_at	0.89557	peroxisomal biogenesis factor 3	Pex3
1454045_a_at	1.04112	phosphatidylglycerophosphate synthase 1	Pgs1
1454046_x_at	1.11608	phosphatidylglycerophosphate synthase 1	Pgs1
1454047_a_at	0.97755	RIKEN cDNA 2410017P07 gene	2410017P07Rik
1454048_a_at	0.97196	RIKEN cDNA 4931408A02 gene	/// similar to Protein C21orf63 homolog precursor 4931408A02Rik
1454049_at	0.95091	RIKEN cDNA 4933426G20 gene	4933426G20Rik
1454050_at	0.94692	RIKEN cDNA 4921521D15 gene	4921521D15Rik
1454051_at	1.01104	RIKEN cDNA 2600011E07 gene	2600011E07Rik
1454052_at	0.88385	HORMA domain containing 2	Hormad2
1454053_at	0.89616	RIKEN cDNA 5730527A11 gene	5730527A11Rik
1454054_at	1.05721	RIKEN cDNA 4930459I23 gene	4930459I23Rik
1454055_at	0.86329	RIKEN cDNA 1600029O15 gene	1600029O15Rik
1454056_at	1.06109	RIKEN cDNA 9030407P20 gene	9030407P20Rik
1454057_at	0.95642	RIKEN cDNA 4930470B04 gene	4930470B04Rik
1454058_at	0.97508	RIKEN cDNA 4931406E20 gene	4931406E20Rik
1454059_at	0.97920	RIKEN cDNA 4930500F04 gene	4930500F04Rik
1454060_a_at	1.04458	neuroblastoma ras oncogene	Nras
1454061_at	1.20470	THUMP domain containing 3	Thumpd3
1454062_at	1.03044	RIKEN cDNA 4933402D24 gene	4933402D24Rik
1454063_at	0.90332	RIKEN cDNA 4933412E24 gene	4933412E24Rik
1454064_a_at	0.93500	ring finger protein 138	Rnf138
1454065_at	0.95367	RIKEN cDNA 4930562D21 gene	4930562D21Rik
1454066_at	0.93707	myc induced nuclear antigen	Mina
1454067_a_at	0.91637	RIKEN cDNA 4931406C07 gene	4931406C07Rik
1454068_at	1.00921	RIKEN cDNA 4930470H14 gene	4930470H14Rik
1454069_at	1.00082	RIKEN cDNA 9030409C19 gene	9030409C19Rik
1454070_a_at	0.92548	DDHD domain containing 1	Ddhd1
1454071_at	1.06143	"lymphocyte antigen 6 complex, locus K"	Ly6k
1454072_at	0.87109	RIKEN cDNA 4731417B20 gene	4731417B20Rik
1454073_at	1.23000	RIKEN cDNA 4930455D15 gene	4930455D15Rik
1454074_a_at	0.95568	RIKEN cDNA 1500011J06 gene	1500011J06Rik
1454075_s_at	1.03657	nudix (nucleoside diphosphate linked moiety X)-type motif 13	Nudt13
1454076_at	1.06678	leucine rich repeat containing 17	Lrrc17
1454077_at	0.99266	vesicle transport through interaction with t-SNAREs homolog 1A (yeast)	Vti1a
1454078_a_at	0.88374	galactose-3-O-sulfotransferase 1	Gal3st1

1454079_at	1.06564	RIKEN cDNA 4933400F03 gene	4933400F03Rik
1454080_at	1.10832	RIKEN cDNA 4933428M03 gene	4933428M03Rik
1454081_at	1.04534	---	---
1454082_a_at	0.90655	GIY-YIG domain containing 2	Giyd2
1454083_at	1.00652	RIKEN cDNA 4930440C22 gene	4930440C22Rik
1454084_a_at	0.92967	SUMO/sentrin specific peptidase 8	Senp8
1454085_at	0.94709	RIKEN cDNA 4933409D19 gene	4933409D19Rik
1454086_a_at	0.84707	LIM domain only 2	Lmo2
1454087_at	0.88023	zinc fingerprotein 618	Zfp618
1454088_at	0.96169	RIKEN cDNA 5330411O13 gene	5330411O13Rik
1454089_at	1.01110	"apolipoprotein B mRNA editing enzyme, catalytic polypeptide-like 4 (putative)"	Apobec4
1454090_at	1.33520	"prenyl (solanesyl) diphosphate synthase, subunit 1"	Pdss1
1454091_at	0.89748	pregnancy-specific glycoprotein 21	Psg21
1454092_a_at	0.92332	"general transcription factor IIIH, polypeptide 3"	Gtf2h3
1454093_at	0.95135	RIKEN cDNA 4933436I20 gene	4933436I20Rik
1454094_at	1.04345	RIKEN cDNA 4930471E19 gene	4930471E19Rik
1454095_at	1.08929	RIKEN cDNA 4930579H20 gene	4930579H20Rik
1454096_a_at	0.89679	RIKEN cDNA 4930555I21 gene	4930555I21Rik
1454097_at	1.15053	"zinc finger, CW type with PWWP domain 2"	Zcwpw2
1454098_at	1.12446	olfactory receptor 112	Olf112
1454099_at	0.86599	RIKEN cDNA 4930593A02 gene	4930593A02Rik
1454100_at	0.96073	RIKEN cDNA 4930480E11 gene	4930480E11Rik
1454101_at	0.91115	RIKEN cDNA 4933401P06 gene	4933401P06Rik
1454102_at	0.86769	RIKEN cDNA 5031425E22 gene	5031425E22Rik
1454103_at	0.86087	dynein heavy chain domain 3	Dnhd3
1454104_a_at	0.95021	"solute carrier family 16 (monocarboxylic acid transporters), member 9"	Slc16a9
1454105_at	1.04459	---	---
1454106_a_at	0.84959	CXXC finger 1 (PHD domain)	Cxxc1
1454107_a_at	0.88405	kinesin family member 2A	Kif2a
1454108_at	1.01125	RIKEN cDNA 4930524O08 gene	4930524O08Rik
1454109_a_at	1.43731	phosphatidylserine receptor	Ptdsr
1454110_at	1.00081	RIKEN cDNA 4930408K08 gene	4930408K08Rik
1454111_at	0.92832	RIKEN cDNA 4930404A10 gene	4930404A10Rik
1454112_a_at	1.13770	centrosomal protein 27	Cep27
1454113_at	0.97096	RIKEN cDNA 4930472D12 gene	4930472D12Rik
1454114_a_at	1.02169	RIKEN cDNA 4933425K02 gene	4933425K02Rik
1454115_at	1.05055	RIKEN cDNA 4930572J10 gene	4930572J10Rik
1454116_a_at	0.98159	MTERF domain containing 1	Mterfd1
1454117_at	0.88812	RIKEN cDNA 5330439B14 gene	5330439B14Rik
1454118_at	0.96169	RIKEN cDNA 5830410O09 gene	5830410O09Rik
1454119_at	0.96493	MAP kinase-interacting serine/threonine kinase 1	Mknk1

1454120_a_at	1.21400	polycomb group ring finger 6	Pcgf6	
1454121_x_at	1.05297	coiled-coil domain containing 18	Ccdc18	
1454122_at	0.92432	RIKEN cDNA 1810011H11 gene	1810011H11Rik	
1454123_at	0.99999	RIKEN cDNA 4933406K04 gene	4933406K04Rik	
1454124_at	0.93913	RIKEN cDNA 1700067K01 gene	1700067K01Rik	
1454125_a_at	0.98751	RIKEN cDNA 4930563E19 gene	4930563E19Rik	
1454126_at	0.96556	RIKEN cDNA 0710001A04 gene	0710001A04Rik	
1454127_at	0.98327	RIKEN cDNA 1010001I08 gene	1010001I08Rik	
1454128_at	0.86288	RIKEN cDNA 4930413E15 gene	4930413E15Rik	
1454129_at	0.98690	RIKEN cDNA 4933416M06 gene	4933416M06Rik	
1454130_at	1.01886	RIKEN cDNA 4930432J01 gene	4930432J01Rik	
1454131_at	1.11852	RIKEN cDNA 4931413K12 gene	4931413K12Rik	
1454132_at	1.13415	RIKEN cDNA 7630402F16 gene	7630402F16Rik	
1454133_s_at	1.13263	RIKEN cDNA 4930523O13 gene	4930523O13Rik	
1454134_at	0.96846	RIKEN cDNA 5730412P04 gene	5730412P04Rik	
1454135_at	1.02054	RIKEN cDNA 4930458K08 gene	4930458K08Rik	
1454136_a_at	0.93833	RIKEN cDNA 4921524J17 gene	4921524J17Rik	
1454137_s_at	0.88736	hemochromatosis type 2 (juvenile) (human homolog)	Hfe2	
1454138_a_at	1.05566	serine threonine kinase 31	Stk31	
1454139_at	0.93133	RIKEN cDNA 4930405N21 gene	4930405N21Rik	
1454140_at	0.97268	Neural cell adhesion molecule 1	Ncam1	
1454141_at	1.37482	RIKEN cDNA 5730416O20 gene	5730416O20Rik	
1454142_a_at	1.07207	PWP1 homolog (<i>S. cerevisiae</i>)	Pwp1	
1454143_at	0.99292	RIKEN cDNA 4930442H23 gene	4930442H23Rik	
1454144_a_at	0.98484	cyclin C	Ccnc	
1454145_at	0.94031	phospholipase A2 receptor 1	Pla2r1	
1454146_at	0.85868	RIKEN cDNA 9530036M11 gene	9530036M11Rik	
1454147_at	1.01723	RIKEN cDNA 2810429I04 gene	2810429I04Rik	
1454148_at	1.02396	RIKEN cDNA 4930443B20 gene	4930443B20Rik	
1454149_a_at	1.11509	cyclin L2	Ccnl2	
1454150_at	0.95358	RIKEN cDNA 4930453O03 gene	4930453O03Rik	
1454151_at	1.00121	transmembrane protein 167	Tmem167	
1454152_a_at	1.02760	"G protein-coupled receptor kinase 2, groucho gene related (<i>Drosophila</i>)"	Gprk21	
1454153_at	0.86993	RIKEN cDNA 4930556J24 gene	4930556J24Rik	
1454154_at	0.95104	RIKEN cDNA 4933416I08 gene	4933416I08Rik	
1454155_at	0.96352	RIKEN cDNA 4930500H12 gene	4930500H12Rik	
1454156_at	0.93256	RIKEN cDNA A230104H11 gene	A230104H11Rik	
1454157_a_at	1.00553	"phospholipase A2, group IID"	Pla2g2d	
1454158_at	1.15227	"membrane protein, palmitoylated 7 (MAGUK p55 subfamily member 7)"	Mpp7	
1454159_a_at	0.99863	insulin-like growth factor binding protein 2	Igfbp2	
1454160_at	0.94190	RIKEN cDNA 4930483C13 gene	4930483C13Rik	

1454161_s_at	1.19073	RIKEN cDNA 0610007P14	gene	0610007P14Rik
1454162_at	0.94220	RIKEN cDNA 4930518P08	gene	4930518P08Rik
1454163_at	1.14700	RIKEN cDNA 5730420D15	gene	5730420D15Rik
1454164_at	0.97506	RIKEN cDNA 4921523L03	gene	4921523L03Rik
1454165_at	0.98078	RIKEN cDNA 1700025K04	gene	1700025K04Rik
1454166_at	1.01445	RIKEN cDNA 1700040N02	gene	1700040N02Rik
1454167_at	0.95776	"nuclear receptor subfamily 2, group C, member 2"		Nr2c2
1454168_a_at	1.04127	ClpB caseinolytic peptidase B homolog (E. coli)	ClpB	
1454169_a_at	0.89763	epithelial stromal interaction 1 (breast)	Epsti1	
1454170_at	0.94659	RIKEN cDNA E130101E03	gene	E130101E03Rik
1454171_x_at	1.02737	RIKEN cDNA 9530053H05	gene	9530053H05Rik
1454172_at	1.25996	BCL6 co-repressor-like 1	Bcorl1	
1454173_at	0.93462	SH2 domain containing 4B	Sh2d4b	
1454174_a_at	1.03516	RIKEN cDNA C330007P06	gene	C330007P06Rik
1454175_at	1.09025	RIKEN cDNA 4930517N10	gene	4930517N10Rik
1454176_at	1.00672	"excision repair cross-complementing rodent repair deficiency, complementation group 8"	Ercc8	
1454177_at	1.01567	RIKEN cDNA 4930597G03	gene	4930597G03Rik
1454178_at	0.93021	RIKEN cDNA 2610027F03	gene	2610027F03Rik
1454179_at	1.00618	RIKEN cDNA 4933406K04	gene	4933406K04Rik
1454180_at	0.97691	armadillo repeat containing 2	Armc2	
1454181_at	1.05108	RIKEN cDNA 4930519B02	gene	4930519B02Rik
1454182_at	1.54345	RIKEN cDNA 5430417C01	gene	5430417C01Rik
1454183_at	0.98321	RIKEN cDNA 3100004P22	gene	3100004P22Rik
1454184_a_at	0.97134	inhibitor of kappaB kinase beta	Ikbkb	
1454185_at	1.08277	RIKEN cDNA 4933436E23	gene	4933436E23Rik
1454186_a_at	0.80022	RIKEN cDNA 2310051N18	gene	2310051N18Rik
1454187_at	1.01101	polycystic kidney disease 1 like 2	Pkd1l2	
1454188_at	0.99350	RIKEN cDNA 4921508D12	gene	4921508D12Rik
1454189_at	1.03731	RIKEN cDNA 4930557J02	gene	4930557J02Rik
1454190_at	1.15376	RIKEN cDNA 9630013K17	gene	9630013K17Rik
1454191_at	0.91221	RIKEN cDNA A030007E19	gene	A030007E19Rik
1454192_at	0.91944	RIKEN cDNA 4930579C15	gene	4930579C15Rik
1454193_at	0.90069	RIKEN cDNA 5430401H09	gene	5430401H09Rik
1454194_at	1.06443	RIKEN cDNA 9330118I20	gene	9330118I20Rik
1454195_at	0.95390	RIKEN cDNA 4933433G19	gene	4933433G19Rik
1454196_at	0.89715	RIKEN cDNA 4930568A13	gene	4930568A13Rik
1454197_a_at	1.05049	coiled-coil domain containing 86	Ccdc86	
1454198_a_at	1.00108	EF hand calcium binding domain 1	Efcab1	
1454199_at	0.95277	RIKEN cDNA 1700014N06	gene	1700014N06Rik
1454200_at	1.05374	zinc finger homeobox 1b	Zfx1b	
1454201_a_at	0.89714	RIKEN cDNA 5730405I09	gene	5730405I09Rik
1454202_a_at	1.02094	RIKEN cDNA 1700061J05	gene	1700061J05Rik

1454203_at 0.82509 RIKEN cDNA 1700061J05 gene 1700061J05Rik
1454204_at 1.01796 RIKEN cDNA 4930442G10 gene 4930442G10Rik
1454205_at 1.27382 RIKEN cDNA 4930448A20 gene 4930448A20Rik
1454206_a_at 0.88261 a disintegrin and metallopeptidase domain 15 (metargidin) Adam15
1454207_at 1.05711 RIKEN cDNA C430039J16 gene C430039J16Rik
1454208_at 0.86730 RIKEN cDNA 4930483O08 gene 4930483O08Rik
1454209_at 0.95355 RIKEN cDNA 1500016O10 gene 1500016O10Rik
1454210_at 0.90367 RIKEN cDNA 4933428M09 gene 4933428M09Rik
1454211_a_at 1.03208 shroom family member 3 Shroom3
1454212_x_at 0.99468 gasdermin domain containing 2 /// RIKEN cDNA 4933426G20 gene ///
RIKEN cDNA 1700065I16 gene /// similar to gasdermin 1 Gsdmdc2 /// 4933426G20Rik ///
1700065I16Rik /// LOC665769
1454213_at 1.10252 armadillo repeat containing 9 Armc9
1454214_a_at 1.11478 RIKEN cDNA 2410019A14 gene 2410019A14Rik
1454215_at 0.92573 RIKEN cDNA 2410007B07 gene 2410007B07Rik
1454216_at 0.88708 RIKEN cDNA 4930408F14 gene 4930408F14Rik
1454217_at 0.98913 RIKEN cDNA 4930556N09 gene 4930556N09Rik
1454218_at 1.08443 RIKEN cDNA 4930405D01 gene 4930405D01Rik
1454219_at 0.99161 "DnaJ (Hsp40) homolog, subfamily C, member 2" Dnajc2
1454220_at 0.97070 RIKEN cDNA 9330185C12 gene 9330185C12Rik
1454221_a_at 0.98265 kinesin family member 2C Kif2c
1454222_a_at 1.01933 coiled-coil domain containing 13 Ccdc13
1454223_at 0.97249 RIKEN cDNA 4933423P22 gene 4933423P22Rik
1454224_at 1.07542 RIKEN cDNA 2010300F17 gene 2010300F17Rik
1454225_s_at 0.91451 "DNA segment, Chr 3, ERATO Doi 751, expressed" D3Ertd751e
1454226_at 0.94232 RIKEN cDNA 4931403M11 gene 4931403M11Rik
1454227_at 0.99770 "HIV-1 tat interactive protein 2, homolog (human)" Htatip2
1454228_a_at 1.05111 coiled-coil domain containing 77 Ccdc77
1454229_a_at 0.94178 exonuclease domain containing 1 Exod1
1454230_a_at 0.93086 "solute carrier family 25, member 27" Slc25a27
1454231_a_at 1.01108 retinitis pigmentosa GTPase regulator interacting protein 1 Rpgrip1
1454232_at 0.89609 RIKEN cDNA 9430027B09 gene 9430027B09Rik
1454233_at 1.00608 RIKEN cDNA 2310006M14 gene 2310006M14Rik
1454234_at 0.87431 Williams-Beuren syndrome chromosome region 28 (human) Wbscr28
1454235_a_at 1.12137 "inhibitor of growth family, member 5" Ing5
1454236_a_at 1.11176 RIKEN cDNA C030004A17 gene C030004A17Rik
1454237_at 1.05317 spermatogenesis associated 9 Spata9
1454238_a_at 0.91954 RIKEN cDNA 1700010H22 gene 1700010H22Rik
1454239_at 0.97048 RIKEN cDNA 4930578G10 gene 4930578G10Rik
1454240_at 1.03763 "nuclear factor, erythroid derived 2, like 3" Nfe2l3
1454241_at 1.04674 RWD domain containing 1 Rwdd1
1454242_at 0.73161 RIKEN cDNA 2310079G19 gene 2310079G19Rik
1454243_at 1.33174 intestinal cell kinase Ick

1454244_at	0.93375	minichromosome maintenance deficient 10 (<i>S. cerevisiae</i>)	Mcm10
1454245_at	0.97205	CUB and Sushi multiple domains 3	Csmd3
1454246_at	1.06683	"ST3 beta-galactoside alpha-2,3-sialyltransferase 6"	St3gal6
1454247_a_at	1.05222	glycoprotein A33 (transmembrane)	Gpa33
1454248_at	0.99384	RIKEN cDNA 1700041E20 gene	1700041E20Rik
1454249_at	0.99395	RIKEN cDNA D530049N12 gene	D530049N12Rik
1454250_at	0.88687	SEC15-like 2 (<i>S. cerevisiae</i>)	Sec15l2
1454251_at	0.98822	RIKEN cDNA 1700034J04 gene	1700034J04Rik
1454252_at	0.87438	RIKEN cDNA 8030453O22 gene	8030453O22Rik
1454253_at	0.89395	RIKEN cDNA 2810452K22 gene	2810452K22Rik
1454254_s_at	0.97213	RIKEN cDNA 1600029D21 gene	1600029D21Rik
1454255_at	1.12044	RIKEN cDNA 5430434F05 gene	5430434F05Rik
1454256_s_at	0.90621	RIKEN cDNA 1700113I22 gene	1700113I22Rik
1454257_at	1.16581	"serine palmitoyltransferase, long chain base subunit 2"	Sptlc2
1454258_at	0.98085	RIKEN cDNA 1110008E08 gene	1110008E08Rik
1454259_s_at	0.96033	RIKEN cDNA 4931433A01 gene	4931433A01Rik
1454260_at	0.87745	U2AF homology motif (UHM) kinase 1	Uhmk1
1454261_at	0.83003	RIKEN cDNA E130304I02 gene	E130304I02Rik
1454262_at	0.89215	RIKEN cDNA 1700124P09 gene	1700124P09Rik
1454263_at	0.92721	goosecoid-like	Gscl
1454264_at	1.02578	RIKEN cDNA 2310046K23 gene	2310046K23Rik
1454265_a_at	1.04946	"ATP-binding cassette, sub-family B (MDR/TAP), member 10"	Abcb10
1454266_at	1.05941	claudin domain containing 2	Cldnd2
1454267_a_at	1.03431	"protease, serine, 32"	Prss32
1454268_a_at	1.22094	"cytochrome b-245, alpha polypeptide"	Cyba
1454269_s_at	0.96624	RIKEN cDNA 4930519L02 gene	4930519L02Rik
1454270_at	0.92432	galectin-related inter-fiber protein	Grifin
1454271_at	0.98595	RIKEN cDNA 1810013D15 gene	1810013D15Rik
1454272_at	0.95117	RIKEN cDNA 9630015K15 gene	9630015K15Rik
1454273_at	1.00066	RIKEN cDNA 1700025H01 gene	1700025H01Rik
1454274_at	1.13040	RIKEN cDNA 4930571C24 gene	4930571C24Rik
1454275_at	1.00892	---	---
1454276_at	0.87545	RIKEN cDNA 1700021O21 gene	1700021O21Rik
1454277_at	0.91658	RIKEN cDNA 1700049J03 gene	1700049J03Rik
1454278_at	1.06464	RIKEN cDNA A430105D02 gene	A430105D02Rik
1454279_at	1.04697	ornithine decarboxylase antizyme 3	Oaz3
1454280_at	1.17224	RIKEN cDNA 4931406H21 gene	4931406H21Rik
1454281_at	0.97091	RIKEN cDNA 4930506C02 gene	4930506C02Rik
1454282_at	0.91437	RIKEN cDNA 1700084C06 gene	1700084C06Rik
1454283_at	1.01110	RIKEN cDNA 9130023D20 gene	9130023D20Rik
1454284_at	1.21711	"solute carrier family 25 (mitochondrial carrier, phosphate carrier), member 25"	Slc25a25
1454285_at	1.02495	RIKEN cDNA 2900042E19 gene	2900042E19Rik

1454286_at	1.99186	RIKEN cDNA 1110004M10	gene	1110004M10Rik
1454287_at	0.90318	RIKEN cDNA 4931428A05	gene	4931428A05Rik
1454288_at	1.12282	RIKEN cDNA 5830468K08	gene	5830468K08Rik
1454289_at	0.91266	zinc fingers and homeoboxes 3		Zhx3
1454290_at	1.07743	RIKEN cDNA 4930573O21	gene	4930573O21Rik
1454291_at	0.93375	RIKEN cDNA 4933428P19	gene	4933428P19Rik
1454292_at	0.92214	germ cell-less homolog 1 (Drosophila)		Gmcl1
1454293_at	0.92435	retinoblastoma-like 1 (p107)		Rbl1
1454294_at	0.89318	RIKEN cDNA 2310061A09	gene	2310061A09Rik
1454295_at	1.19695	nuclear receptor interacting protein 1		Nrip1
1454296_at	0.90679	RIKEN cDNA 4631402F24	gene	4631402F24Rik
1454297_at	1.08739	RIKEN cDNA 4631402F24	gene	4631402F24Rik
1454298_at	1.11684	RIKEN cDNA 4932430A15	gene	4932430A15Rik
1454299_at	1.18582	RIKEN cDNA 4833422B07	gene	4833422B07Rik
1454300_at	0.92969	RIKEN cDNA 4933424C09	gene	4933424C09Rik
1454301_at	1.87688	RIKEN cDNA 2900073C17	gene	2900073C17Rik
1454302_at	1.13655	RIKEN cDNA 5730410E19	gene	5730410E19Rik
1454303_at	0.97174	epsin 2		Epn2
1454304_at	0.92998	epsin 2		Epn2
1454305_at	1.00993	chromobox homolog 3 (Drosophila HP1 gamma)		Cbx3
1454306_at	0.92446	similar to RIKEN cDNA 1190002C06	gene	LOC672445
1454307_at	1.05590	RIKEN cDNA 1810058N15	gene	1810058N15Rik
1454308_at	1.06697	similar to development- and differentiation-enhancing factor 2; PYK2 C terminus-associated protein	/// similar to development- and differentiation-enhancing factor 2	
1454309_at	1.10319	RIKEN cDNA 2810002N01	gene	2810002N01Rik
1454310_at	0.92898	RIKEN cDNA 5830490A12	gene	5830490A12Rik
1454311_at	1.11407	---	---	
1454312_at	1.01647	RIKEN cDNA 2900097C17	gene	2900097C17Rik
1454313_at	1.05982	epidermal growth factor receptor		Egfr
1454314_at	0.88712	RIKEN cDNA 1700031K17	gene	1700031K17Rik
1454315_at	1.03792	RIKEN cDNA 3300002A11	gene	3300002A11Rik
1454316_at	0.96387	RIKEN cDNA 5830426C09	gene	5830426C09Rik
1454317_at	0.83350	a disintegrin and metallopeptidase domain 1b		Adam1b
1454318_at	1.04051	RIKEN cDNA 2810403D21	gene	2810403D21Rik
1454319_at	0.86196	RIKEN cDNA 4930402H05	gene	4930402H05Rik
1454320_at	0.96133	RIKEN cDNA 5830420C07	gene	5830420C07Rik
1454321_at	0.81854	RIKEN cDNA 1500005C15	gene	1500005C15Rik
1454322_at	1.03002	RIKEN cDNA 4933404M09	gene	4933404M09Rik
1454323_at	1.01958	RIKEN cDNA 5830433D23	gene	5830433D23Rik
1454324_at	1.03390	RIKEN cDNA 4930487N04	gene	4930487N04Rik
1454325_at	1.02505	RIKEN cDNA 4930487N04	gene	4930487N04Rik
1454326_at	1.01409	RIKEN cDNA 1700017H01	gene	1700017H01Rik

1454327_at	0.91813	RIKEN cDNA 4933430H06	gene	4933430H06Rik
1454328_at	0.88559	RIKEN cDNA 2900060K15	gene	2900060K15Rik
1454329_at	1.03407	RIKEN cDNA 4933415J04	gene	4933415J04Rik
1454330_at	1.03799	RIKEN cDNA 4933415J04	gene	4933415J04Rik
1454331_at	1.12728	RIKEN cDNA 9430092D12	gene	9430092D12Rik
1454332_at	0.97153	RIKEN cDNA 6030442H21	gene	6030442H21Rik
1454333_at	0.93857	"crystallin, gamma A"	Cryga	
1454334_at	0.97367	RIKEN cDNA 2810409C01	gene	2810409C01Rik
1454335_at	0.92838	RIKEN cDNA 9130214F15	gene	9130214F15Rik
1454336_at	0.93390	Cyclin C	Ccnc	
1454337_at	1.02452	RIKEN cDNA 6330571C24	gene	6330571C24Rik
1454338_at	1.06336	BAT2 domain containing 1	Bat2d	
1454339_at	0.94995	RIKEN cDNA 4930551I15	gene	4930551I15Rik
1454340_at	0.96279	RIKEN cDNA 9530025H10	gene	9530025H10Rik
1454341_at	0.94412	RIKEN cDNA 4933404I11	gene	4933404I11Rik
1454342_at	1.08581	RIKEN cDNA C030007D22	gene	C030007D22Rik
1454343_at	1.09887	RIKEN cDNA 6430710M23	gene	6430710M23Rik
1454344_at	1.04341	"solute carrier family 44, member 1"	Slc44a1	
1454345_at	1.09812	RIKEN cDNA 1700081H04	gene	1700081H04Rik
1454346_at	1.03577	RIKEN cDNA 4930540E01	gene	4930540E01Rik
1454347_at	0.98677	RIKEN cDNA 4930540E01	gene	4930540E01Rik
1454348_at	1.04644	RIKEN cDNA 4933407C09	gene	4933407C09Rik
1454349_at	0.97036	RIKEN cDNA 2810487C13	gene	2810487C13Rik
1454350_at	0.98683	PDZ domain containing 6	Pdzd6	
1454351_at	0.90219	RIKEN cDNA 4933435C09	gene	4933435C09Rik
1454352_at	0.93247	RIKEN cDNA A930028O11	gene	A930028O11Rik
1454353_at	1.03737	RIKEN cDNA 4833420D23	gene	4833420D23Rik
1454354_at	1.16506	RIKEN cDNA 8030476L19	gene	8030476L19Rik
1454355_at	0.92388	RIKEN cDNA 1810021M19	gene	1810021M19Rik
1454356_at	0.89829	RIKEN cDNA 1810021M19	gene	1810021M19Rik
1454357_at	0.92334	RIKEN cDNA 3110067C02	gene	3110067C02Rik
1454358_at	1.04470	RIKEN cDNA 2900060N12	gene	2900060N12Rik
1454359_at	0.85530	RIKEN cDNA 9530023I19	gene	9530023I19Rik
1454360_at	1.13466	RIKEN cDNA 4930418C01	gene	4930418C01Rik
1454361_at	0.95505	RIKEN cDNA 9230106L01	gene	9230106L01Rik
1454362_at	0.98000	RIKEN cDNA 8030431A06	gene	8030431A06Rik
1454363_at	1.00796	RIKEN cDNA 9430099H24	gene	9430099H24Rik
1454364_at	1.02935	RIKEN cDNA 9530025L08	gene	9530025L08Rik
1454365_at	0.95724	RIKEN cDNA 4930554P06	gene	4930554P06Rik
1454366_at	0.91163	RIKEN cDNA 4833419A21	gene	4833419A21Rik
1454367_at	1.00783	RIKEN cDNA 4930413G21	gene	4930413G21Rik
1454368_at	1.01417	"proteasome (prosome, macropain) 26S subunit, non-ATPase, 7"	Psmid7	
1454369_a_at	1.05458	"nuclear factor of activated T-cells, cytoplasmic, calcineurin-dependent 4"		

Nfatc4

1454370_at	0.96721	RIKEN cDNA 4930557B21 gene	4930557B21Rik
1454371_at	0.94674	synaptophysin-like 2 Sypl2	
1454372_at	0.83842	CD80 antigen Cd80	
1454373_x_at	0.98631	ubiquitin C Ubc	
1454374_at	0.99911	RIKEN cDNA 5730458M16 gene	5730458M16Rik
1454375_at	1.05233	RIKEN cDNA 5730458M16 gene	5730458M16Rik
1454376_at	0.93686	RIKEN cDNA 4833410I11 gene	4833410I11Rik
1454377_at	0.99553	RIKEN cDNA 2700080J24 gene	2700080J24Rik
1454378_at	1.15121	"Tyrosine 3-monooxygenase/tryptophan 5-monooxygenase activation protein, theta polypeptide" Ywhaq	
1454379_at	1.05555	RIKEN cDNA 5730405O12 gene	5730405O12Rik
1454380_at	0.88055	phosphatase and actin regulator 4 Phactr4	
1454381_at	1.00498	RIKEN cDNA 5033428C03 gene	5033428C03Rik
1454382_at	1.03787	sterile alpha motif domain containing 4 Samd4	
1454383_at	1.03249	RIKEN cDNA 5430434G16 gene	5430434G16Rik
1454384_at	1.05423	RIKEN cDNA 2900057C01 gene	2900057C01Rik
1454385_at	0.89986	RIKEN cDNA 1700018P08 gene	1700018P08Rik
1454386_at	1.02000	RIKEN cDNA 6430500D05 gene	6430500D05Rik
1454387_at	0.84978	pannexin 3 Panx3	
1454388_at	1.57853	RIKEN cDNA 2900046F13 gene	2900046F13Rik
1454389_at	0.87383	Leucine-rich repeats and calponin homology (CH) domain containing 4 Lrch4	Lrch4
1454390_at	0.95942	leucine-rich repeats and calponin homology (CH) domain containing 4 Lrch4	Lrch4
1454391_at	0.93580	RIKEN cDNA 6030442K20 gene	6030442K20Rik
1454392_at	1.07526	RIKEN cDNA 9230112J17 gene	9230112J17Rik
1454393_at	1.12154	RIKEN cDNA 2310047C04 gene	2310047C04Rik
1454394_at	1.26210	RIKEN cDNA 4932432N04 gene	4932432N04Rik
1454395_at	0.95903	RIKEN cDNA 4632404M16 gene	4632404M16Rik
1454396_at	0.94623	RIKEN cDNA 4632409D06 gene	4632409D06Rik
1454397_at	1.11881	RIKEN cDNA 4632418H02 gene	4632418H02Rik
1454398_at	0.99567	---	---
1454399_at	1.02462	RIKEN cDNA 2010003H20 gene	2010003H20Rik
1454400_at	1.01905	RIKEN cDNA 4632432E15 gene	4632432E15Rik
1454401_at	0.93415	RIKEN cDNA 5730488B01 gene	5730488B01Rik
1454402_at	0.88608	RIKEN cDNA 3110048L19 gene	3110048L19Rik
1454403_at	0.91558	RIKEN cDNA 4930466I24 gene	4930466I24Rik
1454404_at	1.09614	RIKEN cDNA 2610011E03 gene	2610011E03Rik
1454405_at	0.96271	RIKEN cDNA 4930453J04 gene	4930453J04Rik
1454406_at	0.96213	RIKEN cDNA 4930453J04 gene	4930453J04Rik
1454407_at	0.91935	RIKEN cDNA 4930435N07 gene	4930435N07Rik
1454408_at	0.94981	RIKEN cDNA 4930544L18 gene	4930544L18Rik
1454409_at	0.94053	RIKEN cDNA 4833408G04 gene	4833408G04Rik
1454410_at	0.89213	RIKEN cDNA 9130414P19 gene	9130414P19Rik

1454411_at	0.96011 RIKEN cDNA 4933406F09 gene	4933406F09Rik	
1454412_at	1.06173 RIKEN cDNA 2810410P21 gene	2810410P21Rik	
1454413_at	0.95666 RIKEN cDNA 4933424H11 gene	4933424H11Rik	
1454414_at	1.00861 BTB (POZ) domain containing 7	Btbd7	
1454415_at	0.98674 scaffold attachment factor B	Safb	
1454416_at	1.05219 RIKEN cDNA 5730575I04 gene	5730575I04Rik	
1454417_at	1.04009 RIKEN cDNA 4933408M05 gene	4933408M05Rik	
1454418_at	0.97843 RIKEN cDNA 5330421C15 gene	5330421C15Rik	
1454419_at	1.11657 RIKEN cDNA 4930537H20 gene	4930537H20Rik	
1454420_at	0.89998 RIKEN cDNA 5730420F10 gene	5730420F10Rik	
1454421_at	0.99115 RIKEN cDNA 4930423D22 gene	4930423D22Rik	
1454422_at	1.08073 RIKEN cDNA 4930456J16 gene	4930456J16Rik	
1454423_at	0.98649 RIKEN cDNA 4930538D17 gene	4930538D17Rik	
1454424_at	0.98815 RIKEN cDNA 2610040L17 gene	2610040L17Rik	
1454425_at	0.98004 RIKEN cDNA 5830495A06 gene	5830495A06Rik	
1454426_at	1.03558 RIKEN cDNA 4930519E07 gene	4930519E07Rik	
1454427_at	0.92962 --- ---		
1454428_at	0.99952 RIKEN cDNA 4930467D19 gene	4930467D19Rik	
1454429_at	1.05565 RIKEN cDNA 5830462O15 gene	5830462O15Rik	
1454430_at	0.93603 phosphodiesterase 11A	Pde11a	
1454431_at	1.07659 RIKEN cDNA 9030613N10 gene	9030613N10Rik	
1454432_at	0.91857 RIKEN cDNA 4933423N03 gene	4933423N03Rik	
1454433_at	0.88682 RIKEN cDNA 6330526H18 gene	6330526H18Rik	
1454434_at	0.91966 RIKEN cDNA 9030607J07 gene	9030607J07Rik	
1454435_at	0.85094 RIKEN cDNA 5530400N10 gene	5530400N10Rik	
1454436_at	0.88897 RIKEN cDNA 4921506L19 gene	4921506L19Rik	
1454437_at	0.98470 --- ---		
1454438_at	1.09211 "RAB27A, member RAS oncogene family"	Rab27a	
1454439_at	1.03973 RIKEN cDNA 4833419G08 gene	4833419G08Rik	
1454440_at	0.99116 RIKEN cDNA 4930421C12 gene	4930421C12Rik	
1454441_at	0.81431 "BCL2/adenovirus E1B interacting protein 1, NIP2"		Bnip2
1454442_at	0.92774 "BCL2/adenovirus E1B interacting protein 1, NIP2"		Bnip2
1454443_at	0.89314 RIKEN cDNA 4833411I10 gene	4833411I10Rik	
1454444_at	0.99442 RIKEN cDNA 5730406E14 gene	5730406E14Rik	
1454445_at	1.08296 RIKEN cDNA 5430430K15 gene	5430430K15Rik	
1454446_at	0.87216 RIKEN cDNA 4930564B12 gene	4930564B12Rik	
1454447_at	1.01417 RIKEN cDNA 5830434F19 gene	5830434F19Rik	
1454448_at	0.90048 RIKEN cDNA 2900018E21 gene	2900018E21Rik	
1454449_at	1.30844 Rab9 effector protein with kelch motifs	Rabepk	
1454450_at	1.02690 RIKEN cDNA 4930572C08 gene	4930572C08Rik	
1454451_at	1.04559 RIKEN cDNA 5330430B06 gene	5330430B06Rik	
1454452_at	1.06935 RIKEN cDNA 4933428L01 gene	4933428L01Rik	
1454453_at	1.01840 RIKEN cDNA 2610011I18 gene	2610011I18Rik	

1454454_at	0.82008 --- ---	
1454455_at	1.02479 RIKEN cDNA 6530413G14 gene	6530413G14Rik
1454456_at	0.91920 RIKEN cDNA 6530413G14 gene	6530413G14Rik
1454457_at	0.96002 RIKEN cDNA 4930439G18 gene	4930439G18Rik
1454458_at	0.99313 RIKEN cDNA 6230426I18 gene	6230426I18Rik
1454459_at	0.95098 RIKEN cDNA 4933423L19 gene	4933423L19Rik
1454460_at	0.93995 RIKEN cDNA 5730433N10 gene	5730433N10Rik
1454461_at	1.03093 RIKEN cDNA 5830433M15 gene	5830433M15Rik
1454462_at	0.96369 RIKEN cDNA 6430605C03 gene	6430605C03Rik
1454463_at	0.91347 RIKEN cDNA 5430402O13 gene	5430402O13Rik
1454464_at	0.96031 RIKEN cDNA 2700078F05 gene	2700078F05Rik
1454465_at	0.78341 RIKEN cDNA 5830435N06 gene	5830435N06Rik
1454466_at	1.02112 RIKEN cDNA 4933407I18 gene	4933407I18Rik
1454467_at	0.93063 similar to Vinculin CG3299-PA ///	similar to Vinculin CG3299-PA
LOC668894 ///	LOC671420	
1454468_at	0.87732 RIKEN cDNA 5430434N17 gene	5430434N17Rik
1454469_at	1.07162 RIKEN cDNA 5430434N17 gene	5430434N17Rik
1454470_at	0.87433 RIKEN cDNA 4930404F17 gene	4930404F17Rik
1454471_at	0.99763 RIKEN cDNA 3110062G12 gene	3110062G12Rik
1454472_at	1.10143 RIKEN cDNA 2900092N22 gene	2900092N22Rik
1454473_at	0.92770 RIKEN cDNA 4933433M23 gene	4933433M23Rik
1454474_at	1.02485 RIKEN cDNA 4933433M23 gene	4933433M23Rik
1454475_at	0.91447 RIKEN cDNA 4921524M04 gene	4921524M04Rik
1454476_at	1.03322 RIKEN cDNA 4933428L12 gene	4933428L12Rik
1454477_at	0.88433 RIKEN cDNA 5530401A10 gene	5530401A10Rik
1454478_at	1.06431 RIKEN cDNA 3100002H20 gene	3100002H20Rik
1454479_at	0.91808 --- ---	
1454480_at	0.98098 RIKEN cDNA 4933439N06 gene	4933439N06Rik
1454481_at	0.99132 macrophage migration inhibitory factor Mif	
1454482_at	0.94728 RIKEN cDNA 5830474E16 gene	5830474E16Rik
1454483_at	1.06073 RIKEN cDNA 6330590E21 gene	6330590E21Rik
1454484_at	0.87811 RIKEN cDNA 4933424L07 gene	4933424L07Rik
1454485_at	0.95341 RIKEN cDNA 4930456G14 gene	4930456G14Rik
1454486_at	0.97818 RIKEN cDNA 9530002O20 gene	9530002O20Rik
1454487_at	1.17858 RIKEN cDNA 5830490A04 gene	5830490A04Rik
1454488_at	1.00829 RIKEN cDNA 4930534I15 gene	4930534I15Rik
1454489_at	0.94222 RIKEN cDNA 4930513L20 gene	4930513L20Rik
1454490_at	1.13072 RIKEN cDNA 4930423C22 gene	4930423C22Rik
1454491_at	0.90557 RIKEN cDNA 2900040J22 gene	2900040J22Rik
1454492_at	1.13348 RIKEN cDNA 8430418B16 gene	8430418B16Rik
1454493_at	0.97585 RIKEN cDNA 5033425B01 gene	5033425B01Rik
1454494_at	1.10281 RIKEN cDNA 5830448L01 gene	5830448L01Rik
1454495_at	1.05284 RIKEN cDNA 4930432H08 gene	4930432H08Rik

1454496_at	0.86502 RIKEN cDNA 5031415H12 gene	5031415H12Rik
1454497_at	1.11944 RIKEN cDNA 5230401M06 gene	5230401M06Rik
1454498_at	1.04761 RIKEN cDNA 9230118N17 gene	9230118N17Rik
1454499_at	0.98317 RIKEN cDNA 4933431C10 gene	4933431C10Rik
1454500_at	0.95991 RIKEN cDNA 6330419E04 gene	6330419E04Rik
1454501_at	0.91301 RIKEN cDNA 6330419E04 gene	6330419E04Rik
1454502_at	0.96762 RIKEN cDNA 2900022B07 gene	2900022B07Rik
1454503_at	0.86443 RIKEN cDNA 9030201C23 gene	9030201C23Rik
1454504_at	0.89509 Kv channel-interacting protein 1	Kcnip1
1454505_at	1.06325 RIKEN cDNA 6330562C20 gene	6330562C20Rik
1454506_at	0.94542 --- ---	
1454507_at	1.05134 RIKEN cDNA 8430432A02 gene	8430432A02Rik
1454508_at	1.02475 RIKEN cDNA 4930485E13 gene	4930485E13Rik
1454509_at	0.91505 RIKEN cDNA 4930444E06 gene	4930444E06Rik
1454510_at	0.85694 RIKEN cDNA 2900034C19 gene	2900034C19Rik
1454511_at	0.99032 RIKEN cDNA 9430018C23 gene	9430018C23Rik
1454512_at	1.10913 RIKEN cDNA 9430018C23 gene	9430018C23Rik
1454513_at	1.00727 RIKEN cDNA 9430031J08 gene	9430031J08Rik
1454514_at	0.98435 RIKEN cDNA 5730415C11 gene	5730415C11Rik
1454515_at	0.89126 RIKEN cDNA 5430425E15 gene	5430425E15Rik
1454516_at	0.98642 RIKEN cDNA 4930424E08 gene	4930424E08Rik
1454517_at	1.08416 RIKEN cDNA C030011I16 gene	C030011I16Rik
1454518_at	1.12937 RIKEN cDNA 4930429L21 gene	4930429L21Rik
1454519_at	0.88238 RIKEN cDNA 4930429L21 gene	4930429L21Rik
1454520_at	0.96729 RIKEN cDNA 5830415B17 gene	5830415B17Rik
1454521_at	0.99981 RIKEN cDNA 2810421E14 gene	2810421E14Rik
1454522_at	0.97396 RIKEN cDNA 4930573G07 gene	4930573G07Rik
1454523_at	0.98860 RIKEN cDNA C530007A02 gene	C530007A02Rik
1454524_at	1.04859 RIKEN cDNA 2310075M01 gene	2310075M01Rik
1454525_at	0.94905 selenoprotein 15-Sep	
1454526_at	1.06179 RIKEN cDNA 9430014F16 gene	9430014F16Rik
1454527_at	1.03796 RIKEN cDNA 8430437B07 gene	8430437B07Rik
1454528_at	1.01980 RIKEN cDNA A930011E06 gene	A930011E06Rik
1454529_at	1.01539 RIKEN cDNA 4930518C04 gene	4930518C04Rik
1454530_at	0.81734 --- ---	
1454531_at	0.92653 RIKEN cDNA C030043A13 gene	C030043A13Rik
1454532_at	1.06895 RIKEN cDNA C030043A13 gene	C030043A13Rik
1454533_at	0.97507 RIKEN cDNA 9430085L16 gene	9430085L16Rik
1454534_at	1.02562 "inhibitor of growth family, member 1" Ing1	
1454535_at	0.98597 RIKEN cDNA A430110A21 gene	A430110A21Rik
1454536_at	0.94126 RIKEN cDNA 4933401B01 gene	4933401B01Rik
1454537_at	0.91448 RIKEN cDNA 4930556A12 gene	4930556A12Rik
1454538_at	0.90866 RIKEN cDNA D730048A03 gene	D730048A03Rik

1454539_at	0.99382	RIKEN cDNA 4933417G07	gene	4933417G07Rik
1454540_at	0.94887	RIKEN cDNA 4930504E06	gene	4930504E06Rik
1454541_at	0.93948	RIKEN cDNA 4921522E08	gene	4921522E08Rik
1454542_at	0.96391	RIKEN cDNA 9330179C17	gene	9330179C17Rik
1454543_at	0.98333	serine/threonine kinase 40	Stk40	
1454544_at	0.97971	RIKEN cDNA 9530078B04	gene	9530078B04Rik
1454545_at	1.05097	RIKEN cDNA 4933403L11	gene	4933403L11Rik
1454546_at	1.00071	RIKEN cDNA 4933403L11	gene	4933403L11Rik
1454547_at	0.98968	RIKEN cDNA 2310045N14	gene	2310045N14Rik
1454548_at	0.94359	RIKEN cDNA 1700039M15	gene	1700039M15Rik
1454549_at	0.91606	RIKEN cDNA 2900005I04	gene	2900005I04Rik
1454550_at	0.96859	RIKEN cDNA 4930478M09	gene	4930478M09Rik
1454551_at	1.17077	RIKEN cDNA 9530034D02	gene	9530034D02Rik
1454552_at	0.94257	RIKEN cDNA 4930456K20	gene	4930456K20Rik
1454553_at	1.10932	RIKEN cDNA 4930509E22	gene	4930509E22Rik
1454554_at	1.02840	RIKEN cDNA 9330159N22	gene	9330159N22Rik
1454555_at	0.99370	RIKEN cDNA 4930472F24	gene	4930472F24Rik
1454556_at	1.05134	RIKEN cDNA 2700057C20	gene	2700057C20Rik
1454557_at	1.10071	RIKEN cDNA 6720454L07	gene	6720454L07Rik
1454558_at	1.06222	RIKEN cDNA 5430416B10	gene	5430416B10Rik
1454559_at	1.02003	RIKEN cDNA 5430437A18	gene	5430437A18Rik
1454560_at	1.02768	RIKEN cDNA 4930519N06	gene	4930519N06Rik
1454561_at	1.00997	RIKEN cDNA 9430087B13	gene	9430087B13Rik
1454562_at	1.03276	"Adult male testis cDNA, RIKEN full-length enriched library, clone:4930590N04 product:hypothetical protein, full insert sequence" ---		
1454563_at	1.11212	RIKEN cDNA 4930573C08	gene	4930573C08Rik
1454564_at	0.91242	RIKEN cDNA 4933401J01	gene	4933401J01Rik
1454565_at	0.93929	RIKEN cDNA C030046M01	gene	C030046M01Rik
1454566_at	0.95080	RIKEN cDNA 4933439C20	gene ///	RIKEN cDNA D030013I16 gene 4933439C20Rik /// D030013I16Rik
1454567_at	1.08358	RIKEN cDNA 4933404K13	gene	4933404K13Rik
1454568_at	1.08694	RIKEN cDNA 6030460B20	gene	6030460B20Rik
1454569_at	1.03516	RIKEN cDNA 4930480K02	gene	4930480K02Rik
1454570_at	1.02459	RIKEN cDNA 5830432F11	gene	5830432F11Rik
1454571_at	1.08631	RIKEN cDNA A930036A04	gene	A930036A04Rik
1454572_at	0.87866	RIKEN cDNA 2810414N06	gene	2810414N06Rik
1454573_at	0.88206	RIKEN cDNA 4933412L11	gene	4933412L11Rik
1454574_at	1.01795	RIKEN cDNA C430049A07	gene	C430049A07Rik
1454575_at	0.98722	RIKEN cDNA 5730422E09	gene	5730422E09Rik
1454576_at	0.93832	RIKEN cDNA A230102O21	gene	A230102O21Rik
1454577_at	1.10248	similar to cell division cycle 20 homolog	LOC238896	
1454578_at	1.02191	RIKEN cDNA 6030458A19	gene	6030458A19Rik
1454579_at	0.90785	RIKEN cDNA 4930474A20	gene	4930474A20Rik

1454580_at	1.03456	RIKEN cDNA 5430427N15	gene	5430427N15Rik
1454581_at	1.02892	RIKEN cDNA 5330425B07	gene	5330425B07Rik
1454582_at	0.93355	RIKEN cDNA D530037P16	gene	D530037P16Rik
1454583_at	0.93878	RIKEN cDNA 2810034D10	gene	2810034D10Rik
1454584_at	1.10364	RIKEN cDNA 2810034D10	gene	2810034D10Rik
1454585_at	1.09292	RIKEN cDNA 1110065P19	gene	1110065P19Rik
1454586_at	1.04503	RIKEN cDNA 1500002K03	gene	1500002K03Rik
1454587_at	0.88796	RIKEN cDNA 9330199C07	gene	9330199C07Rik
1454588_at	0.83088	RIKEN cDNA 9430006E15	gene	9430006E15Rik
1454589_at	0.85159	RIKEN cDNA 9430006E15	gene	9430006E15Rik
1454590_at	1.06344	RIKEN cDNA A930018O16	gene	A930018O16Rik
1454591_at	0.94153	microtubule associated serine/threonine kinase family member 4		Mast4
1454592_at	1.58845	RIKEN cDNA 9430012M22	gene	9430012M22Rik
1454593_at	0.91221	"FYVE, RhoGEF and PH domain containing 3"		Fgd3
1454594_at	0.98883	RIKEN cDNA 2900078I11	gene	2900078I11Rik
1454595_at	0.93443	RIKEN cDNA 4930451E06	gene	4930451E06Rik
1454596_at	0.89839	RIKEN cDNA A730089K16	gene	A730089K16Rik
1454597_at	0.97406	RIKEN cDNA 4930423O20	gene	4930423O20Rik
1454598_at	0.88824	RIKEN cDNA 4930425F17	gene	4930425F17Rik
1454599_at	0.96518	RIKEN cDNA 4930425F17	gene	4930425F17Rik
1454600_at	1.00937	RIKEN cDNA 9530086P17	gene	9530086P17Rik
1454601_at	1.01644	RIKEN cDNA 8030497O21	gene	8030497O21Rik
1454602_s_at	1.01457	"CCR4-NOT transcription complex, subunit 2"		Cnot2
1454603_a_at	1.01946	"CCR4-NOT transcription complex, subunit 2"		Cnot2
1454604_s_at	0.79969	tetraspanin 12		Tspan12
1454605_a_at	1.04570	phosphatidylinositol 4-kinase type 2 alpha		Pi4k2a
1454606_at	1.26604	RIKEN cDNA 4933426M11	gene	4933426M11Rik
1454607_s_at	1.06657	phosphoserine aminotransferase 1		Psat1
1454608_x_at	0.44837	transthyretin		Ttr
1454609_x_at	0.88111	RIKEN cDNA 6430527G18	gene	6430527G18Rik
1454610_at	0.94656	septin 7		7-Sep
1454611_a_at	1.00250	calmodulin 1		Calm1
1454612_at	0.98525	ring finger and KH domain containing 2		Rkhd2
1454613_at	0.33584	dihydropyrimidinase-like 3		Dpysl3
1454614_at	0.87785	RIKEN cDNA 1810013D10	gene	1810013D10Rik
1454615_x_at	0.90014	signal recognition particle 14		Srp14
1454616_at	0.87136	RIKEN cDNA 5730410I19	gene	5730410I19Rik
1454617_at	0.59081	arrestin domain containing 3		Arrdc3
1454618_at	0.91908	---		---
1454619_at	0.96840	transmembrane protein 153		Tmem153
1454620_x_at	1.08719	ribosomal protein S6 /// similar to 40S ribosomal protein S6 /// similar to 40S ribosomal protein S6 /// similar to 40S ribosomal protein S6 /// similar to 40S ribosomal protein S6 /// similar to 40S ribosomal protein S6 /// similar to 40S ribosomal protein S6 /// similar to 40S ribosomal protein S6		

protein S6 /// similar to 40S ribosomal protein S6 /// similar to 40S ribosomal protein S6 /// similar to 40S ribosomal protein S6 /// similar to 40S ribosomal protein S6 /// similar to 40S ribosomal protein S6 /// similar to 40S ribosomal protein S6 /// similar to 40S ribosomal protein S6 /// similar to 40S ribosomal protein S6 /// similar to 40S ribosomal protein S6 /// similar to 40S ribosomal protein S6
 Rps6 /// LOC434404 /// LOC545640 /// LOC626259 /// LOC629575 /// LOC638497 /// LOC639124 /// LOC639593 /// LOC639750 /// LOC666577 /// LOC667717 /// LOC667739 /// LOC667922 /// LOC671306 /// LOC671432 /// LOC672192 /// LOC672620 /// LOC676999
 1454621_at 0.85203 RIKEN cDNA 5730472N09 gene 5730472N09Rik
 1454622_at 0.98610 "solute carrier family 38, member 5" Slc38a5
 1454623_at 1.12336 "carboxypeptidase A2, pancreatic" Cpa2
 1454624_at 0.86702 DENN/MADD domain containing 2A Dennd2a
 1454625_at 1.00173 PHD finger protein 6 Phf6
 1454626_at 1.02218 "clathrin, heavy polypeptide (Hc)" Cltc
 1454627_a_at 1.07375 ribosomal protein L29 /// similar to 60S ribosomal protein L29 /// similar to 60S ribosomal protein L29 /// similar to 60S ribosomal protein L29 /// similar to 60S ribosomal protein L29 /// similar to 60S ribosomal protein L29 /// similar to 60S ribosomal protein L29 /// similar to 60S ribosomal protein L29 /// similar to 60S ribosomal protein L29 /// similar to 60S ribosomal protein L29 /// hypothetical protein LOC669999 /// similar to 60S ribosomal protein L29 /// similar to 60S ribosomal protein L29 Rpl29 /// LOC433350 /// LOC433941 /// LOC622707 /// LOC626772 /// LOC665032 /// LOC666642 /// LOC667466 /// LOC669999 /// LOC670334 /// LOC675846
 1454628_at 1.11053 RIKEN cDNA A930037G23 gene A930037G23Rik
 1454629_at 0.96331 RIKEN cDNA A930037G23 gene A930037G23Rik
 1454630_at 0.85554 sterile alpha motif domain containing 14 Samd14
 1454631_at 0.99219 RIKEN cDNA 6330549H03 gene 6330549H03Rik
 1454632_at 0.95794 RIKEN cDNA 6330442E10 gene 6330442E10Rik
 1454633_at 0.86077 ethanolamine kinase 1 Etnk1
 1454634_at 1.09764 fucokinase Fuk
 1454635_at 0.95514 F-box and leucine-rich repeat protein 3 Fbxl3
 1454636_at 0.94481 chromobox homolog 5 (Drosophila HP1a) Cbx5
 1454637_at 0.88041 kelch-like 8 (Drosophila) Klhl8
 1454638_a_at 0.92525 phenylalanine hydroxylase Pah
 1454639_x_at 1.52570 ribosomal protein L41 /// RIKEN cDNA 2210411K19 gene Rpl41 /// 2210411K19Rik
 1454640_at 0.88124 coiled-coil-helix-coiled-coil-helix domain containing 7 Chchd7
 1454641_at 0.93270 CGG triplet repeat binding protein 1 Cggbp1
 1454642_a_at 1.04438 COMM domain containing 3 Commd3
 1454643_at 1.02932 ubiquitin associated protein 2-like Ubap2l
 1454644_at 1.12392 RIKEN cDNA 6330569M22 gene 6330569M22Rik
 1454645_at 0.90352 "mahogunin, ring finger 1" Mgrn1
 1454646_at 0.77968 t-complex 11 (mouse) like 2 Tcp112
 1454647_at 0.83085 "acyl-Coenzyme A dehydrogenase family, member 11" Acad11
 1454648_s_at 1.15155 "DNA segment, Chr 10, Wayne State University 102, expressed" D10Wsu102e

1454649_at 1.02286 steroid 5 alpha-reductase 1 Srd5a1
1454650_at 0.91839 tripartite motif-containing 35 Trim35
1454651_x_at 0.99679 myelin basic protein Mbp
1454652_at 0.92654 "zinc finger, RAN-binding domain containing 2" Zranb2
1454653_at 0.97486 copine family member IX Cpne9
1454654_at 0.88343 disrupted in renal carcinoma 2 (human) Dirc2
1454655_at 0.94633 "diacylglycerol kinase, delta" Dgkd
1454656_at 0.94270 spermatogenesis associated 13 Spata13
1454657_s_at 0.99298 "MAK10 homolog, amino-acid N-acetyltransferase subunit, (S. cerevisiae)" Mak10
1454658_at 0.96652 ilvB (bacterial acetolactate synthase)-like Ilvbl
1454659_at 1.05938 dCMP deaminase Dctd
1454660_at 0.95817 RIKEN cDNA 1100001E04 gene 1100001E04Rik
1454661_at 0.94698 "ATP synthase, H+ transporting, mitochondrial F0 complex, subunit c (subunit 9), isoform 3" Atp5g3
1454662_at 0.91631 RIKEN cDNA A230106M15 gene A230106M15Rik
1454663_at 1.12592 eukaryotic translation initiation factor 5 Eif5
1454664_a_at 1.08542 eukaryotic translation initiation factor 5 Eif5
1454665_at 0.83017 interferon regulatory factor 2 binding protein 2 /// similar to interferon regulatory factor 2 binding protein 2 Irf2bp2 /// LOC672960
1454666_at 0.91794 Kruppel-like factor 3 (basic) Klf3
1454667_at 1.09242 sorting nexin 8 Snx8
1454668_at 1.00412 "zinc finger, UBR1 type 1" Zubr1
1454669_at 1.04763 transmembrane protein 11 Tmem11
1454670_at 1.03868 arginine glutamic acid dipeptide (RE) repeats Rere
1454671_at 1.11499 insulin induced gene 1 Insig1
1454672_at 1.03365 --- ---
1454673_at 0.94984 --- ---
1454674_at 1.04080 fasciculation and elongation protein zeta 1 (zygin I) Fez1
1454675_at 0.73293 thyroid hormone receptor alpha Thra
1454676_s_at 0.83289 toll-like receptor adaptor molecule 1 Ticam1
1454677_at 0.86155 tissue inhibitor of metalloproteinase 2 Timp2
1454678_s_at 0.91704 RIKEN cDNA A130022J15 gene A130022J15Rik
1454679_at 1.06297 "DNA segment, Chr 8, ERATO Doi 457, expressed" D8Ertd457e
1454680_at 0.87542 "DNA segment, Chr 5, ERATO Doi 579, expressed" D5Ertd579e
1454681_at 1.16533 RNA binding motif protein 35A Rbm35a
1454682_at 1.01602 RIKEN cDNA A430005L14 gene A430005L14Rik
1454683_at 0.88456 "splicing factor, arginine/serine-rich 8" Sfrs8
1454684_at 0.94567 Bardet-Biedl syndrome 7 Bbs7
1454685_at 1.00516 G protein-coupled receptor 146 Gpr146
1454686_at 0.92472 RIKEN cDNA C330011F01 gene /// RIKEN cDNA 6430706D22 gene C330011F01Rik /// 6430706D22Rik
1454687_at 1.03651 leucine rich repeat and fibronectin type III domain containing 5 Lfn5

1454688_x_at 1.07113 transmembrane emp24-like trafficking protein 10 (yeast) /// similar to
 transmembrane trafficking protein Tmed10 /// LOC634748
 1454689_at 0.95261 serine/arginine repetitive matrix 1 Srrm1
 1454690_at 0.98717 inhibitor of kappaB kinase gamma Ikbkg
 1454691_at 0.94773 neurexin I Nrnx1
 1454692_x_at 0.93446 Heterogeneous nuclear ribonucleoprotein K Hnrpk
 1454693_at 0.87454 Histone deacetylase 4 Hdac4
 1454694_a_at 1.05752 topoisomerase (DNA) II alpha Top2a
 1454695_at 0.97352 WD repeat domain 18 Wdr18
 1454696_at 1.02048 "guanine nucleotide binding protein, beta 1" Gnb1
 1454697_at 0.86251 translocation protein 1 Tloc1
 1454698_at 0.94678 protein tyrosine phosphatase-like A domain containing 1 Ptplad1
 1454699_at 0.73686 sestrin 1 Sesn1
 1454700_at 0.95336 leucine rich repeat and fibronectin type III domain containing 4 Lrnf4
 1454701_at 1.01946 RIKEN cDNA 4930503L19 gene 4930503L19Rik
 1454702_at 0.99293 RIKEN cDNA 4930503L19 gene 4930503L19Rik
 1454703_x_at 1.14413 "small nucleolar RNA, C/D box 22" Snord22
 1454704_at 0.98709 "scavenger receptor class B, member 2" Scarb2
 1454705_at 1.00163 "DNA segment, Chr 15, ERATO Doi 621, expressed" D15Ertd621e
 1454706_at 1.11647 UV radiation resistance associated gene Uvrag
 1454707_at 0.90530 RIKEN cDNA 2310035C23 gene 2310035C23Rik
 1454708_at 1.08585 actin-binding LIM protein 1 Ablim1
 1454709_at 0.77531 transmembrane protein 64 Tmem64
 1454710_at 0.95879 "serine peptidase inhibitor, Kazal type 2" Spink2
 1454711_at 0.98115 triple functional domain (PTPRF interacting) Trio
 1454712_at 0.94519 mitochondrial carrier triple repeat 1 Mcart1
 1454713_s_at 0.94492 histidine decarboxylase Hdc
 1454714_x_at 1.13564 3-phosphoglycerate dehydrogenase /// similar to 3-phosphoglycerate
 dehydrogenase /// similar to 3-phosphoglycerate dehydrogenase /// similar to 3-phosphoglycerate
 dehydrogenase /// similar to 3-phosphoglycerate dehydrogenase Phgdh /// LOC668771 ///
 LOC670155 /// LOC671972 /// LOC673015
 1454715_at 0.95415 RIKEN cDNA 0710005M24 gene 0710005M24Rik
 1454716_x_at 0.99161 "cytochrome c oxidase, subunit Vb" Cox5b
 1454717_at 1.05827 Ankyrin repeat domain 27 (VPS9 domain) Ankrd27
 1454718_at 1.02021 N-acetylglucosamine-1-phosphodiester alpha-N-acetylglucosaminidase
 Nagpa
 1454719_at 0.94901 rhomboid domain containing 3 Rhbdd3
 1454720_at 0.93394 "amyloid beta (A4) precursor protein-binding, family A, member 3 /// heparan
 sulfate (glucosamine) 3-O-sulfotransferase 5" Apba3 /// Hs3st5
 1454721_at 0.97538 RIKEN cDNA 1110018G07 gene 1110018G07Rik
 1454722_at 0.85571 phosphatase and tensin homolog Pten
 1454723_at 1.01456 RIKEN cDNA 1110033M05 gene 1110033M05Rik
 1454724_x_at 0.93026 RIKEN cDNA 5730446C15 gene 5730446C15Rik

1454725_at 1.02712 transformer 2 alpha homolog (Drosophila) Tra2a
1454726_s_at 0.82947 expressed sequence AW456874 AW456874
1454727_at 0.96960 expressed sequence AI173486 AI173486
1454728_s_at 0.66441 "ATPase, aminophospholipid transporter (APLT), class I, type 8A, member 1" Atp8a1
1454729_at 1.04800 transmembrane protein 108 Tmem108
1454730_at 1.02096 RIKEN cDNA 4932414K18 gene 4932414K18Rik
1454731_at 1.12352 myosin X Myo10
1454732_at 1.00554 microtubule-associated protein 1S Mtap1s
1454733_at 0.94236 caspase recruitment domain 4 Card4
1454734_at 0.92217 --- ---
1454735_at 1.01940 outer dense fiber of sperm tails 2 Odf2
1454736_at 1.02615 ankyrin repeat domain 57 Ankrd57
1454737_at 1.26637 dual specificity phosphatase 9 Dusp9
1454738_x_at 0.84547 peroxisomal biogenesis factor 6 Pex6
1454739_at 1.06508 cell division cycle 27 homolog (S. cerevisiae) Cdc27
1454740_at 1.09092 mindbomb homolog 1 (Drosophila) Mib1
1454741_s_at 0.94819 transmembrane protein 164 Tmem164
1454742_at 1.02512 "RasGEF domain family, member 1B" Rasgef1b
1454743_at 0.99732 nucleoporin 205 Nup205
1454744_at 1.08561 RIKEN cDNA F630043A04 gene F630043A04Rik
1454745_at 1.06963 Rho GTPase activating protein 29 Arhgap29
1454746_at 0.97692 "pleckstrin homology domain containing, family M (with RUN domain) member 1" Plekhm1
1454747_a_at 0.93739 kelch domain containing 3 Klhdc3
1454748_at 0.81184 nicotinate phosphoribosyltransferase domain containing 1 Naprt1
1454749_at 0.93839 pericentrin (kendrin) Pcnt
1454750_a_at 1.04391 cDNA sequence BC057552 BC057552
1454751_at 1.04147 RIKEN cDNA G430022H21 gene G430022H21Rik
1454752_at 0.95297 similar to RNA binding motif protein 24 /// similar to RNA binding motif protein 24 /// similar to RNA binding motif protein 24 LOC380843 /// LOC638266 /// LOC666794
1454753_at 0.99471 arginyl aminopeptidase (aminopeptidase B)-like 1 Rnpepl1
1454754_a_at 0.93310 angio-associated migratory protein Aamp
1454755_at 1.09990 "inositol 1,4,5-trisphosphate 3-kinase C" Itpkc
1454756_at 1.08541 leucine-rich repeats and calponin homology (CH) domain containing 3 Lrch3
1454757_s_at 1.15711 "DNA segment, Chr 12, ERATO Doi 647, expressed" D12Ertd647e
1454758_a_at 0.83600 "TSC22 domain family, member 1" Tsc22d1
1454759_at 0.93384 G protein-coupled receptor kinase-interactor 1 Git1
1454760_at 0.91134 HIV TAT specific factor 1 Htatsf1
1454761_at 0.91388 cDNA sequence BC005764 BC005764
1454762_at 0.89751 X Kell blood group precursor related X linked Xkrx
1454763_at 0.88523 ankyrin repeat domain 17 Ankrd17
1454764_s_at 1.00123 "solute carrier family 38, member 1" Slc38a1

1454765_at 0.94584 "general transcription factor IIIC, polypeptide 3" Gtf3c3
1454766_at 0.95306 RIKEN cDNA C730024G19 gene C730024G19Rik
1454767_at 0.86698 "DNA segment, Chr 2, Brigham & Women's Genetics 1423 expressed"
D2Bwg1423e
1454768_at 1.00596 "potassium voltage-gated channel, subfamily F, member 1" Kcnf1
1454769_at 0.97555 TatD DNase domain containing 2 Tatdn2
1454770_at 0.90865 cholecystokinin B receptor Cckbr
1454771_at 0.96512 --- ---
1454772_at 1.07021 activating signal cointegrator 1 complex subunit 3-like 1 Ascc3l1
1454773_at 0.85162 retinoid X receptor alpha Rxra
1454774_at 1.00735 zinc finger protein 445 Zfp445
1454775_at 0.96649 histone deacetylase 10 Hdac10
1454776_at 0.97586 euchromatic histone methyltransferase 1 Ehmt1
1454777_at 0.97272 "solute carrier organic anion transporter family, member 2b1" Slco2b1
1454778_x_at 0.94376 Cytotoxic granule-associated RNA binding protein 1 Tia1
1454779_s_at 1.05420 RIKEN cDNA C230052I12 gene C230052I12Rik
1454780_at 1.01523 UDP-N-acetyl-alpha-D-galactosamine:polypeptide N-
acetylgalactosaminyltransferase-like 4 Galntl4
1454781_x_at 0.96929 COMM domain containing 9 Commd9
1454782_at 0.88317 brain-specific angiogenesis inhibitor 3 Bai3
1454783_at 1.18374 "interleukin 13 receptor, alpha 1" Il13ra1
1454784_at 1.00354 heparan sulfate (glucosamine) 3-O-sulfotransferase 2 Hs3st2
1454785_at 1.00579 dual specificity phosphatase 11 (RNA/RNP complex 1-interacting) Dusp11
1454786_at 0.88892 RIKEN cDNA 5031439G07 gene 5031439G07Rik
1454787_at 1.08817 "zinc finger, DHHC domain containing 9" Zdhhc9
1454788_at 0.91959 similar to ADP-ribosylation factor-like protein 7 LOC632433
1454789_x_at 0.91643 PRP6 pre-mRNA splicing factor 6 homolog (yeast) Prpf6
1454790_at 0.94672 WD repeat domain 35 Wdr35
1454791_a_at 1.01115 retinoblastoma binding protein 4 Rbbp4
1454792_s_at 0.99021 selenophosphate synthetase 1 Sephs1
1454793_x_at 0.90345 DEAD (Asp-Glu-Ala-Asp) box polypeptide 5 Ddx5
1454794_at 0.95035 spastin Spast
1454795_at 0.90620 Cobl-like 1 Cobll1
1454796_at 0.78905 "DNA segment, Chr 5, ERATO Doi 40, expressed" D5Ertd40e
1454797_at 1.20124 transmembrane protein 55b Tmem55b
1454798_at 1.08800 "Arginine-rich, mutated in early stage tumors" Armet
1454799_at 1.02142 RIKEN cDNA A230097K15 gene A230097K15Rik
1454800_at 0.82499 MORN repeat containing 2 Morn2
1454801_at 0.98049 ankyrin repeat domain 28 Ankrd28
1454802_x_at 1.08182 ariadne homolog 2 (Drosophila) Arih2
1454803_a_at 0.69504 histone deacetylase 11 Hdac11
1454804_at 1.00800 nuclear protein localization 4 homolog (S. cerevisiae) Nploc4
1454805_at 0.90848 Wilms' tumour 1-associating protein Wtap

1454806_at 1.03519 "DNA segment, Chr 12, ERATO Doi 553, expressed" D12Erttd553e
1454807_a_at 1.04679 sorting nexin 12 Snx12
1454808_at 0.96639 EF hand domain family A1 Efha1
1454809_at 0.92449 nuclear receptor coactivator 7 Ncoa7
1454810_s_at 0.90260 expressed sequence AI315068 AI315068
1454811_a_at 0.91894 serine incorporator 1 Serinc1
1454812_at 0.99913 RIKEN cDNA 5730601F06 gene 5730601F06Rik
1454813_at 0.92921 coiled-coil domain containing 72 Ccdc72
1454814_s_at 0.91782 expressed sequence AU021838 AU021838
1454815_at 0.98966 SERTA domain containing 2 Sertad2
1454816_at 1.04585 retinitis pigmentosa 2 homolog (human) Rp2h
1454817_at 1.05418 "UTP18, small subunit (SSU) processome component, homolog (yeast)"
Utp18
1454818_at 0.97975 glucocorticoid modulatory element binding protein 2 Gmeb2
1454819_at 0.94921 --- ---
1454820_at 0.99749 cDNA sequence BC037034 BC037034
1454821_at 0.95038 "beta-1,3-glucuronyltransferase 1 (glucuronosyltransferase P)" B3gat1
1454822_x_at 0.43087 adenomatosis polyposis coli down-regulated 1 Apcdd1
1454823_at 1.02716 WD repeat domain 37 Wdr37
1454824_s_at 0.58213 mitochondrial tumor suppressor 1 Mtus1
1454825_at 0.83930 RIKEN cDNA 1110014N23 gene 1110014N23Rik
1454826_at 1.02520 zinc finger and BTB domain containing 11 Zbtb11
1454827_at 1.00804 pogo transposable element with ZNF domain Pogz
1454828_at 0.98719 expressed sequence AI790205 AI790205
1454829_at 0.98814 RUN domain containing 1 Rundc1
1454830_at 1.05040 fibrillin 2 Fbn2
1454831_at 1.01637 forkhead box N2 Foxn2
1454832_at 0.73489 phosphatase and actin regulator 1 Phactr1
1454833_at 1.19915 ribosomal protein L35 Rpl35
1454834_at 1.07502 nuclear factor I/B Nfib
1454835_at 0.97055 EPM2A (laforin) interacting protein 1 Epm2aip1
1454836_at 0.84768 transmembrane protein 18 Tmem18
1454837_at 0.92710 "ceroid-lipofuscinosis, neuronal 6" Cln6
1454838_s_at 1.14969 expressed sequence AW548124 AW548124
1454839_a_at 1.02475 coiled-coil domain containing 84 Ccdc84
1454840_at 0.95729 methylcrotonoyl-Coenzyme A carboxylase 2 (beta) Mccc2
1454841_at 1.09671 RIKEN cDNA 4921511H13 gene 4921511H13Rik
1454842_a_at 1.12445 "UDP-GalNAc:betaGlcNAc beta 1,3-galactosaminyltransferase, polypeptide
2" B3galnt2
1454843_at 0.85189 phosphoribosyl pyrophosphate synthetase 2 Prps2
1454844_at 0.95908 melanin-concentrating hormone receptor 1 Mchr1
1454845_x_at 1.00919 melanin-concentrating hormone receptor 1 Mchr1
1454846_at 1.00175 "UTP15, U3 small nucleolar ribonucleoprotein, homolog (yeast)" Utp15

1454847_at 1.15778 lipoma HMGIC fusion partner-like 2 Lhfpl2
1454848_at 0.86060 "protein phosphatase 1, regulatory (inhibitor) subunit 12C" Ppp1r12c
1454849_x_at 0.83062 clusterin Clu
1454850_at 0.98741 "TBC1 domain family, member 10c" Tbc1d10c
1454851_at 0.94965 "nuclear receptor subfamily 2, group C, member 2" Nr2c2
1454852_at 0.86615 trans-acting transcription factor 1 Sp1
1454853_s_at 0.95693 expressed sequence AW146154 /// expressed sequence AI987944
AW146154 /// AI987944
1454854_at 1.01545 organic solute transporter beta Ostb
1454855_at 0.79338 "membrane associated guanylate kinase, WW and PDZ domain containing 2"
Magi2
1454856_x_at 1.05911 ribosomal protein L35 Rpl35
1454857_at 0.89446 ring finger protein 122 Rnf122
1454858_x_at 0.91221 methyltransferase like 7A Mettl7a
1454859_a_at 1.01634 ribosomal protein L23 Rpl23
1454860_x_at 0.90575 defender against cell death 1 Dad1
1454861_at 0.96521 taxilin alpha Txlna
1454862_at 1.06254 "pleckstrin homology-like domain, family B, member 2" Phldb2
1454863_at 0.96209 ankyrin repeat domain 11 Ankrd11
1454864_at 0.89714 zinc finger protein 592 Zfp592
1454865_at 1.03748 "solute carrier family 9 (sodium/hydrogen exchanger), member 8" Slc9a8
1454866_s_at 1.02806 chloride intracellular channel 6 Clic6
1454867_at 0.55667 meningioma 1 Mn1
1454868_at 1.00453 "DNA segment, Chr 4, ERATO Doi 429, expressed" D4Ertd429e
1454869_at 0.89231 WD repeat domain 40B Wdr40b
1454870_x_at 0.98851 G protein-coupled receptor 172B Gpr172b
1454871_at 0.91075 RNA binding motif protein 15B Rbm15b
1454872_at 0.92465 RIKEN cDNA B230308N11 gene B230308N11Rik
1454873_at 0.94073 RIKEN cDNA C130032F08 gene C130032F08Rik
1454874_at 0.97754 BTB (POZ) domain containing 7 Btbd7
1454875_a_at 0.91944 retinoblastoma binding protein 4 Rbbp4
1454876_at 1.06207 "RAB23, member RAS oncogene family" Rab23
1454877_at 1.08827 SERTA domain containing 4 Sertad4
1454878_at 1.08156 RIKEN cDNA 2310047C04 gene 2310047C04Rik
1454879_s_at 0.96152 RIKEN cDNA 1700052N19 gene 1700052N19Rik
1454880_s_at 0.55572 Bcl2 modifying factor Bmf
1454881_s_at 1.20166 uroplakin 3B Upk3b
1454882_at 0.86297 l(3)mbt-like 3 (Drosophila) L3mbtl3
1454883_at 0.96044 expressed sequence AI987692 AI987692
1454884_at 0.89197 BTB (POZ) domain containing 4 Btbd4
1454885_at 0.98354 cDNA sequence BC063263 BC063263
1454886_x_at 1.02747 --- ---
1454887_at 1.01480 p21 (CDKN1A)-activated kinase 2 Pak2

1454888_at 1.08168 prefoldin 4 Pfdn4
1454889_x_at 0.81861 transmembrane and coiled coil domains 3 Tmcc3
1454890_at 0.92090 angiomotin Amot
1454891_at 1.13871 CDP-diacylglycerol synthase (phosphatidate cytidyltransferase) 2 Cds2
1454892_at 1.01738 "phosphatidylinositol transfer protein, beta" Pitpnb
1454893_at 1.03814 RIKEN cDNA 1110013L07 gene 1110013L07Rik
1454894_at 0.91509 SMAD specific E3 ubiquitin protein ligase 2 Smurf2
1454895_at 0.99449 "cytochrome b, ascorbate dependent 3" Cybasc3
1454896_at 0.95709 --- ---
1454897_at 0.99367 RIKEN cDNA 6330509M05 gene 6330509M05Rik
1454898_s_at 0.81656 RIKEN cDNA 4833421E05 gene 4833421E05Rik
1454899_at 0.95356 LIM domain containing preferred translocation partner in lipoma Lpp
1454900_s_at 0.94792 "pam, highwire, rpm 1" Phr1
1454901_at 0.74843 Hypothetical protein 6430570G24 6430570G24
1454902_at 0.83409 "protein kinase C, zeta" Prkcz
1454903_at 0.93383 "nerve growth factor receptor (TNFR superfamily, member 16)" Ngfr
1454904_at 1.27952 X-linked myotubular myopathy gene 1 Mtm1
1454905_at 1.02470 inhibitor of Bruton agammaglobulinemia tyrosine kinase Ibk
1454906_at 0.95909 "retinoic acid receptor, beta" Rarb
1454907_at 0.96665 --- ---
1454908_x_at 0.85630 small EDRK-rich factor 1 Serf1
1454909_at 0.88581 "transforming, acidic coiled-coil containing protein 1" Tacc1
1454910_at 0.93691 "13 days embryo heart cDNA, RIKEN full-length enriched library, clone: D330008M10 product:hypothetical protein, full insert sequence" ---
1454911_at 1.10044 hypothetical protein LOC382106 LOC382106
1454912_at 1.03595 pre-B-cell leukemia transcription factor 4 Pbx4
1454913_at 1.05027 RIKEN cDNA 9930104L06 gene 9930104L06Rik
1454914_at 1.09117 RIKEN cDNA 2610101N10 gene 2610101N10Rik
1454915_at 1.12422 RAB3 GTPase activating protein subunit 2 Rab3gap2
1454916_s_at 0.98797 ADP-ribosylation factor interacting protein 1 Arfip1
1454917_at 0.97364 expressed sequence AU045404 AU045404
1454918_at 1.04128 alkylglycerone phosphate synthase Agps
1454919_at 0.99057 Expressed sequence AU044698 AU044698
1454920_at 0.88874 "ubiquitin-like, containing PHD and RING finger domains 2" Uhrf2
1454921_at 1.03508 "gene model 561, (NCBI)" Gm561
1454922_at 1.07010 expressed sequence AI553587 AI553587
1454923_at 1.07748 expressed sequence AW214353 AW214353
1454924_at 1.07625 fucosyltransferase 10 Fut10
1454925_x_at 1.03708 "malate dehydrogenase 1, NAD (soluble)" Mdh1
1454926_at 1.06309 RIKEN cDNA 4930544G21 gene 4930544G21Rik
1454927_at 0.91799 zinc finger protein 41 Zfp41
1454928_at 1.06322 scaffold attachment factor B Safb
1454929_s_at 0.98330 scaffold attachment factor B Safb

1454930_at 1.11818 leucine rich repeat containing 35 Lrrc35
1454931_at 0.94195 CREBBP/EP300 inhibitory protein 2 Cri2
1454932_at 1.00792 REST corepressor 1 Rcor1
1454933_at 0.94787 RIKEN cDNA 2610027C15 gene 2610027C15Rik
1454934_at 0.89979 protein phosphatase 1F (PP2C domain containing) Ppm1f
1454935_at 0.98814 RIKEN cDNA D930001I22 gene D930001I22Rik
1454936_at 0.94219 valyl-tRNA synthetase 2-like Vars2l
1454937_at 1.11710 RIKEN cDNA B630005N14 gene B630005N14Rik
1454938_at 1.00431 sorting nexin 13 Snx13
1454939_at 0.83751 PHD finger protein 20-like 1 Phf20l1
1454940_at 0.95384 "proteasome (prosome, macropain) subunit, alpha type 5 /// similar to Proteasome subunit alpha type 5 (Proteasome zeta chain) (Macropain zeta chain) (Multicatalytic endopeptidase complex zeta chain) /// similar to Proteasome subunit alpha type 5 (Proteasome zeta chain) (Macropain zeta chain) (Multicatalytic endopeptidase complex zeta chain) /// similar to Proteasome subunit alpha type 5 (Proteasome zeta chain) (Macropain zeta chain) (Multicatalytic endopeptidase complex zeta chain) /// similar to Proteasome subunit alpha type 5 (Proteasome zeta chain) (Macropain zeta chain) (Multicatalytic endopeptidase complex zeta chain)" Psma5 /// LOC545687 /// LOC666974 /// LOC671726 /// LOC676113
1454941_at 0.94913 N-myristoyltransferase 1 Nmt1
1454942_at 0.87252 niban protein Niban
1454943_a_at 1.05113 PAX interacting (with transcription-activation domain) protein 1 Paxip1
1454944_at 1.00020 hypermethylated in cancer 2 Hic2
1454945_at 0.91414 leucine-rich repeats and IQ motif containing 2 Lrriq2
1454946_at 0.83734 myeloblastosis oncogene-like 2 Mybl2
1454947_a_at 0.99282 ubiquitin-like domain containing CTD phosphatase 1 /// similar to ubiquitin-like domain containing CTD phosphatase 1 Ublcp1 /// LOC665689
1454948_at 1.13333 ubiquitin specific peptidase 7 Usp7
1454949_at 1.00935 Ubiquitin specific peptidase 7 Usp7
1454950_at 1.03872 RIKEN cDNA B930006L02 gene B930006L02Rik
1454951_at 0.79679 zinc finger protein 606 Zfp606
1454952_s_at 1.07184 RIKEN cDNA B130055D15 gene B130055D15Rik
1454953_at 0.97066 ring finger protein 157 Rnf157
1454954_at 0.94102 WD repeat domain 61 Wdr61
1454955_at 1.14352 importin 7 Ipo7
1454956_at 1.10507 "ribosomal protein S6 kinase, polypeptide 1" Rps6kb1
1454957_at 1.16609 NIN1/RPN12 binding protein 1 homolog (S. cerevisiae) Nob1
1454958_at 0.99550 glycogen synthase kinase 3 beta Gsk3b
1454959_s_at 0.57643 "guanine nucleotide binding protein, alpha inhibiting 1" Gnai1
1454960_at 1.02381 MAD homolog 3 (Drosophila) Smad3
1454961_at 1.08935 synaptojanin 1 Synj1
1454962_at 1.15502 spire homolog 1 (Drosophila) Spire1
1454963_at 0.99440 RIKEN cDNA E430028B21 gene E430028B21Rik
1454964_at 0.91199 cDNA sequence BC021395 BC021395

1454965_at	0.98852	RIKEN cDNA D430039N05 gene	D430039N05Rik
1454966_at	0.99488	integrin alpha 8	Itga8
1454967_at	1.04089	Transcribed locus	---
1454968_at	0.93540	RIKEN cDNA 1110034A24 gene	1110034A24Rik
1454969_at	0.84277	LY6/PLAUR domain containing 6	Lypd6
1454970_at	1.00903	---	---
1454971_x_at	0.82802	"TSC22 domain family, member 1"	Tsc22d1
1454972_at	0.99683	"ataxia, cerebellar, Cayman type homolog (human)"	Atcay
1454973_at	0.94105	activating transcription factor 7 interacting protein	Atf7ip
1454974_at	0.93729	netrin 1	Ntn1
1454975_at	1.19043	proline/serine-rich coiled-coil 2	Psrc2
1454976_at	0.87298	"superoxide dismutase 2, mitochondrial"	Sod2
1454977_at	0.86336	expressed sequence AU020772	AU020772
1454978_at	0.84772	tweety homolog 3 (Drosophila)	Ttyh3
1454979_at	1.04222	diaphanous homolog 1 (Drosophila) /// similar to Protein diaphanous homolog 1 (Diaphanous-related formin-1) (DRF1) (mDIA1) (p140mDIA) Diap1 /// LOC671545	
1454980_at	0.98630	RIKEN cDNA 4930402E16 gene	4930402E16Rik
1454981_at	0.90578	"DNA segment, Chr 4, ERATO Doi 22, expressed"	D4Ertd22e
1454982_at	1.11859	ADP-ribosylation factor guanine nucleotide-exchange factor 2 (brefeldin A-inhibited) /// similar to ADP-ribosylation factor guanine nucleotide-exchange factor 2 (brefeldin A-inhibited)	Arfgef2 /// LOC670796
1454983_at	1.08800	expressed sequence AI465301	AI465301
1454984_at	0.80129	expressed sequence AW061234	AW061234
1454985_at	0.98582	RIKEN cDNA D030051N19 gene	D030051N19Rik
1454986_at	1.09137	zinc finger protein 668	Zfp668
1454987_a_at	0.85136	H2-K region expressed gene 6	H2-Ke6
1454988_s_at	0.96492	"RAB22A, member RAS oncogene family"	Rab22a
1454989_at	0.98246	centrosome and spindle pole associated protein 1	Cspp1
1454990_at	0.90202	AT rich interactive domain 2 (Arid-rfx like)	Arid2
1454991_at	1.00625	"solute carrier family 7 (cationic amino acid transporter, y+ system), member 1"	Slc7a1
1454992_at	1.11874	"solute carrier family 7 (cationic amino acid transporter, y+ system), member 1"	Slc7a1
1454993_a_at	0.91106	"splicing factor, arginine/serine-rich 3 (SRp20)"	Sfrs3
1454994_at	1.04907	kelch-like 20 (Drosophila)	Klhl20
1454995_at	1.09788	dimethylarginine dimethylaminohydrolase 1	Ddah1
1454996_at	1.03974	hydroxysteroid dehydrogenase like 1	Hsd11
1454997_at	0.86862	methionine sulfoxide reductase B3	Msrb3
1454998_at	1.03081	RIKEN cDNA 1200011I18 gene	1200011I18Rik
1454999_at	1.01033	PHD finger protein 21B	Phf21b
1455000_at	0.99532	G protein-coupled receptor 68	Gpr68
1455001_x_at	1.05719	ribosomal protein L13a	Rpl13a
1455002_at	0.96000	protein tyrosine phosphatase 4a1	Ptp4a1

1455003_at 0.93148 "THAP domain containing, apoptosis associated protein 1" Thap1
1455004_at 0.88854 "DNA segment, Chr 2, Wayne State University 81, expressed" D2Wsu81e
1455005_s_at 0.87258 "solute carrier family 5 (sodium/glucose cotransporter), member 2" Slc5a2
1455006_at 0.94396 RIKEN cDNA 2310016M24 gene 2310016M24Rik
1455007_s_at 0.99809 glutamic pyruvate transaminase (alanine aminotransferase) 2 Gpt2
1455008_at 0.80729 "guanine nucleotide binding protein, alpha 12" Gna12
1455009_at 0.87694 carboxypeptidase D Cpd
1455010_at 1.05832 RIKEN cDNA 1500012F01 gene 1500012F01Rik
1455011_at 1.33819 StAR-related lipid transfer (START) domain containing 4 Stard4
1455012_s_at 0.90038 tripartite motif protein 37 Trim37
1455013_at 0.96181 ariadne homolog 2 (Drosophila) Arih2
1455014_at 0.80950 expressed sequence AV009015 AV009015
1455015_at 0.88996 "TBC1 domain family, member 9" Tbc1d9
1455016_at 1.09922 PRP38 pre-mRNA processing factor 38 (yeast) domain containing B Prpf38b
1455017_a_at 0.90993 "zinc finger, MYM-type 3" Zmym3
1455018_at 1.05456 lemur tyrosine kinase 2 Lmtk2
1455019_x_at 0.84205 cytoskeleton-associated protein 4 Ckap4
1455020_at 0.91826 sorting nexin 25 Snx25
1455021_at 1.55425 "gamma-aminobutyric acid (GABA-B) receptor, 1" Gabbr1
1455022_at 1.13055 "striatin, calmodulin binding protein" Strn
1455023_at 1.00727 expressed sequence N28178 N28178
1455024_at 0.98552 tousled-like kinase 1 Tlk1
1455025_at 0.88246 progesterin and adipoQ receptor family member IX Paqr9
1455026_at 0.98228 "sno, strawberry notch homolog 1 (Drosophila)" Sbno1
1455027_at 0.93227 RUN and FYVE domain containing 3 Rufy3
1455028_at 1.07123 microtubule-associated protein tau Mapt
1455029_at 1.06333 kinesin family member 21A Kif21a
1455030_at 1.03581 "protein tyrosine phosphatase, receptor type, J" Ptpnj
1455031_at 0.95077 --- ---
1455032_at 1.12761 RIKEN cDNA 9630037P07 gene 9630037P07Rik
1455033_at 1.11332 RIKEN cDNA B430201A12 gene B430201A12Rik
1455034_at 0.74762 "Nuclear receptor subfamily 4, group A, member 2" Nr4a2
1455035_s_at 1.22621 nucleolar protein 5A Nol5a
1455036_s_at 0.93433 "NADH dehydrogenase (ubiquinone) 1, subcomplex unknown, 2 /// similar to
NADH dehydrogenase (ubiquinone) 1, subcomplex unknown, 2" Ndufc2 /// LOC675851
1455037_at 0.86075 plexin A2 Plxna2
1455038_at 1.15573 similar to melanoma antigen LOC632764
1455039_a_at 1.02823 "transcriptional regulator, SIN3B (yeast)" Sin3b
1455040_s_at 0.90271 RIKEN cDNA 1110062M06 gene 1110062M06Rik
1455041_at 0.89928 "oocyte maturation, beta" Omt2b
1455042_at 0.97060 Transducin (beta)-like 1 X-linked Tbl1x
1455043_at 1.11599 transportin 1 Tnpol
1455044_at 0.78417 transmembrane protein 44 Tmem44

1455045_at 0.81894 serine racemase Srr
1455046_a_at 0.86484 pogo transposable element with ZNF domain Pogz
1455047_at 0.94181 F-box only protein 3 Fbxo3
1455048_at 0.95505 "immunoglobulin superfamily, member 3" Igsf3
1455049_at 0.89870 "immunoglobulin superfamily, member 3" Igsf3
1455050_at 1.00236 RIKEN cDNA E130203B14 gene E130203B14Rik
1455051_at 0.97311 ring finger protein 31 Rnf31
1455052_a_at 0.94921 RIKEN cDNA 2410129H14 gene 2410129H14Rik
1455053_a_at 0.89913 "DCUN1D1 DCN1, defective in cullin neddylation 1, domain containing 1 (S. cerevisiae)" Dcun1d1
1455054_a_at 0.99641 "testis derived transcript 3, pseudogene /// DCUN1D1 DCN1, defective in cullin neddylation 1, domain containing 1 (S. cerevisiae)" Tes3-ps /// Dcun1d1
1455055_at 0.88811 RIKEN cDNA 4930562D19 gene 4930562D19Rik
1455056_at 1.06559 LIM domain only 7 Lmo7
1455057_at 0.97516 guanine monphosphate synthetase Gmps
1455058_at 0.92273 myotubularin related protein 9 Mtmr9
1455059_at 0.79990 RIKEN cDNA 9430093I07 gene 9430093I07Rik
1455060_at 0.96543 Ras-GTPase-activating protein SH3-domain binding protein RP23-336J1.4
1455061_a_at 0.89415 acetyl-Coenzyme A acyltransferase 2 (mitochondrial 3-oxoacyl-Coenzyme A thiolase) Acaa2
1455062_at 1.03116 expressed sequence AU017455 AU017455
1455063_at 0.90826 RIKEN cDNA 4921513D11 gene /// expressed sequence AW061290
4921513D11Rik /// AW061290
1455064_at 0.85210 "RAB36, member RAS oncogene family" Rab36
1455065_x_at 1.08106 glucosamine-6-phosphate deaminase 1 /// similar to Glucosamine-6-phosphate isomerase (Glucosamine-6-phosphate deaminase) (GNPDA) (GlcN6P deaminase) (Oscillin) /// similar to Glucosamine-6-phosphate isomerase (Glucosamine-6-phosphate deaminase) (GNPDA) (GlcN6P deaminase) (Oscillin) Gnpda1 /// LOC231914 /// LOC667410
1455066_s_at 0.89532 RIKEN cDNA 9130229H14 gene 9130229H14Rik
1455067_at 1.03473 pleckstrin and Sec7 domain containing 4 Psd4
1455068_at 1.04590 pleckstrin and Sec7 domain containing 4 Psd4
1455069_x_at 0.97419 "solute carrier family 25 (mitochondrial carrier, adenine nucleotide translocator), member 4" Slc25a4
1455070_at 1.14171 DCP2 decapping enzyme homolog (S. cerevisiae) Dcp2
1455071_at 0.96278 Vzinc finger and BTB domain containing 7B Zbtb7b
1455072_at 0.95679 centrosomal protein 350 Cep350
1455073_at 0.81362 cytidine and dCMP deaminase domain containing 1 Cdadc1
1455074_at 1.10785 --- ---
1455075_at 0.95616 "phosphatidylinositol glycan anchor biosynthesis, class V" Pigv
1455076_a_at 0.90175 RIKEN cDNA 4933424B01 gene 4933424B01Rik
1455077_a_at 0.95870 mortality factor 4 like 1 /// similar to mortality factor 4 like 1 isoform b /// RIKEN cDNA A730098P11 gene /// similar to mortality factor 4 like 1 /// similar to mortality factor 4 like 1 (predicted) Morf4l1 /// LOC433955 /// A730098P11Rik /// LOC627352 /// LOC635663

1455078_at 1.03641 "Heat shock protein 90kDa alpha (cytosolic), class A member 1" Hsp90aa1
 1455079_at 0.83993 "DCN1, defective in cullin neddylation 1, domain containing 4 (S. cerevisiae)"
 Dcun1d4
 1455080_at 0.81273 "protein phosphatase 1, regulatory (inhibitor) subunit 16B" Ppp1r16b
 1455081_at 0.91788 thioredoxin-like 4B Txnl4b
 1455082_at 0.72236 Casitas B-lineage lymphoma b Cblb
 1455083_at 0.89985 "Atpase, class VI, type 11C" Atp11c
 1455084_x_at 1.10008 serine hydroxymethyl transferase 2 (mitochondrial) Shmt2
 1455085_at 0.93466 RIKEN cDNA 1700086L19 gene 1700086L19Rik
 1455086_at 0.94574 Ubiquitin carboxyl-terminal esterase L5 Uchl5
 1455087_at 0.89127 "DNA segment, Chr 7, ERATO Doi 715, expressed" D7Ertd715e
 1455088_at 0.94277 similar to Ser/Thr protein kinase PAR-1A /// similar to serine/threonine
 kinase /// similar to serine/threonine kinase /// similar to serine/threonine kinase /// similar to serine/
 threonine kinase /// similar to serine/threonine kinase /// similar to serine/threonine kinase /// similar to
 serine/threonine kinase /// similar to KP78a CG6715-PA /// similar to MAP/microtubule affinity-
 regulating kinase 1 LOC381936 /// LOC545926 /// LOC620736 /// LOC628664 /// LOC667151 ///
 LOC667378 /// LOC667522 /// LOC667570 /// LOC672121 /// LOC672131
 1455089_at 1.11257 "guanine nucleotide binding protein (G protein), gamma 12" Gng12
 1455090_at 0.68764 angiopoietin-like 2 Angptl2
 1455091_at 0.86243 RIKEN cDNA 3222402P14 gene 3222402P14Rik
 1455092_at 1.20582 zinc finger protein 207 Zfp207
 1455093_a_at 0.97536 alpha-2-HS-glycoprotein Ahsg
 1455094_s_at 0.97452 "ubiquitin-conjugating enzyme E2G 1 (UBC7 homolog, C. elegans)" Ube2g1
 1455095_at 0.97574 "histone 2, H2be" Hist2h2be
 1455096_at 1.05576 --- ---
 1455097_at 0.96106 WD repeat domain 70 Wdr70
 1455098_a_at 1.02814 vitronectin Vtn
 1455099_at 0.98861 monoacylglycerol O-acyltransferase 2 Mogat2
 1455100_at 0.97866 "aldo-keto reductase family 1, member D1" Akr1d1
 1455101_at 1.11686 "Mus musculus, clone IMAGE:2647821, mRNA" ---
 1455102_at 1.11902 RIKEN cDNA D330037H05 gene /// similar to La-related protein 4 (La
 ribonucleoprotein domain family member 4) /// similar to La-related protein 4 (La ribonucleoprotein
 domain family member 4) D330037H05Rik /// LOC673040 /// LOC677019
 1455103_at 1.01157 DEAD (Asp-Glu-Ala-Asp) box polypeptide 46 Ddx46
 1455104_at 0.79049 MAX dimerization protein 1 Mxd1
 1455105_at 1.03269 "protein tyrosine phosphatase, non-receptor type 12" Ptpn12
 1455106_a_at 0.76915 "creatine kinase, brain" Ckb
 1455107_at 1.20579 serine/arginine-rich protein specific kinase 1 Srpk1
 1455108_at 0.94787 eukaryotic translation initiation factor 4E member 2 Eif4e2
 1455109_at 0.85219 transducin (beta)-like 1X-linked receptor 1 Tbl1xr1
 1455110_at 1.06515 "GA repeat binding protein, beta 2" Gabpb2
 1455111_at 0.83505 "Yip1 domain family, member 6" Yipf6
 1455112_at 0.85565 apoptosis-inducing factor (AIF)-like mitochondrion-associated inducer of death

Amid

1455113_at 1.10923 armadillo repeat containing 8 Armc8
 1455114_at 1.01809 uracil DNA glycosylase 2 Ung2
 1455115_a_at 0.95926 crumbs homolog 3 (Drosophila) Crb3
 1455116_at 1.06273 "tumor necrosis factor receptor superfamily, member 19-like /// similar to GrpE protein homolog 2, mitochondrial precursor (Mt-GrpE#2)" Tnfrsf191 /// LOC671976
 1455117_at 0.92066 minichromosome maintenance deficient domain containing 1 Mcmdc1
 1455118_at 1.10366 "DNA segment, Chr 9, ERATO Doi 402, expressed" D9Ert402e
 1455119_at 1.09571 phosphatidic acid phosphatase type 2 domain containing 1A Ppapdc1a
 1455120_at 1.21072 glyoxalase domain containing 1 Gloxd1
 1455121_at 0.99124 ligand dependent nuclear receptor corepressor Lcor
 1455122_at 0.93109 patched domain containing 2 Ptchd2
 1455123_at 1.00359 suppression of tumorigenicity 18 St18
 1455124_at 0.86969 tripartite motif containing 68 Trim68
 1455125_at 0.94140 "centaurin, beta 1" Centb1
 1455126_x_at 1.00495 RIKEN cDNA 2310028O11 gene 2310028O11Rik
 1455127_at 1.03398 "X Kell blood group precursor-related family, member 5" Xkr5
 1455128_x_at 0.99131 trinucleotide repeat containing 6a Tnrc6a
 1455129_at 1.07224 Metadherin Mtdh
 1455130_at 0.90835 "SPT2, Suppressor of Ty, domain containing 1 (S. cerevisiae)" Spty2d1
 1455131_at 1.09184 optic atrophy 3 (human) Opa3
 1455132_at 0.98723 RIKEN cDNA A430107D22 gene A430107D22Rik
 1455133_s_at 0.98375 expressed sequence AI848100 AI848100
 1455134_at 1.08946 RIKEN cDNA A630051L19 gene A630051L19Rik
 1455135_at 0.81074 RIKEN cDNA 2610014I16 gene 2610014I16Rik
 1455136_at 0.98194 "ATPase, Na⁺/K⁺ transporting, alpha 2 polypeptide" Atp1a2
 1455137_at 0.58413 Rap guanine nucleotide exchange factor (GEF) 5 Rapgef5
 1455138_x_at 1.01555 "cofilin 1, non-muscle" Cfl1
 1455139_at 0.71931 expressed sequence AI851716 AI851716
 1455140_at 0.71861 RIKEN cDNA A330068P14 gene A330068P14Rik
 1455141_at 0.91063 trinucleotide repeat containing 6a Tnrc6a
 1455142_at 0.92585 suppressor of cytokine signaling 4 Socs4
 1455143_at 1.02527 neuroligin 2 Nlgn2
 1455144_s_at 0.81636 expressed sequence AU040829 AU040829
 1455145_at 0.90353 "14, 17 days embryo head cDNA, RIKEN full-length enriched library, clone:3222402J19 product:unclassifiable, full insert sequence" ---
 1455146_at 0.92265 RIKEN cDNA C230082I21 gene C230082I21Rik
 1455147_at 1.03342 "PREDICTED: Mus musculus similar to KIAA1913 (LOC622645), mRNA" ---
 1455148_at 1.09406 transmembrane protein 130 Tmem130
 1455149_at 1.08256 SH3 domain containing ring finger 1 Sh3rf1
 1455150_at 0.95231 "13 days embryo forelimb cDNA, RIKEN full-length enriched library, clone:5930438H13 product:unclassifiable, full insert sequence" ---
 1455151_at 0.96500 A kinase (PRKA) anchor protein (yotiao) 9 Akap9

1455152_at 1.01272 expressed sequence AI462493 AI462493
1455153_at 1.01472 zinc finger protein 236 Zfp236
1455154_at 1.05517 GLI-Kruppel family member GLI3 Gli3
1455155_at 1.10105 "LSM14 homolog B (SCD6, S. cerevisiae)" Lsm14b
1455156_at 1.04254 "striatin, calmodulin binding protein" Strn
1455157_a_at 0.98300 cDNA sequence BC039210 BC039210
1455158_at 0.95367 integrin alpha 3 Itga3
1455159_at 0.96377 RIKEN cDNA 2900057D21 gene 2900057D21Rik
1455160_at 0.57677 RIKEN cDNA 2610203C20 gene 2610203C20Rik
1455161_at 1.09617 expressed sequence AI504432 AI504432
1455162_at 1.28898 RIKEN cDNA 4922503N01 gene 4922503N01Rik
1455163_at 0.94674 GUF1 GTPase homolog (S. cerevisiae) Guf1
1455164_at 1.04838 Cdc42 GTPase-activating protein Cdgap
1455165_at 1.00945 RAR-related orphan receptor alpha Rora
1455166_at 0.97005 expressed sequence AW146430 AW146430
1455167_at 0.92194 "cytochrome c oxidase, subunit VIIIc" Cox8c
1455168_a_at 1.04405 "guanine nucleotide binding protein (G protein), beta polypeptide 2 like 1"
Gnb211
1455169_at 0.89952 RAB11 family interacting protein 2 (class I) Rab11fip2
1455170_at 1.09505 RIKEN cDNA 2810001G20 gene 2810001G20Rik
1455171_at 0.94321 suppressor of variegation 4-20 homolog 1 (Drosophila) Suv420h1
1455172_at 1.01303 expressed sequence AU020094 AU020094
1455173_at 1.18866 G1 to S phase transition 1 Gspt1
1455174_at 1.09568 ribosomal protein S19 binding protein 1 Rps19bp1
1455175_at 0.88337 PHD finger protein 13 Phf13
1455176_a_at 0.93829 synaptotagmin XI Syt11
1455177_at 0.84327 Abelson helper integration site Ahi1
1455178_at 0.83505 RUN and TBC1 domain containing 1 Rutbc1
1455179_at 0.99909 "membrane protein, palmitoylated 7 (MAGUK p55 subfamily member 7)"
Mpp7
1455180_at 0.70679 expressed sequence AA407270 AA407270
1455181_at 1.24853 RAS p21 protein activator 2 Rasa2
1455182_at 1.04853 kinesin family member 1B Kif1b
1455183_at 0.96163 serine/threonine kinase 38 like Stk38l
1455184_at 1.13366 "MOB1, Mps One Binder kinase activator-like 1A (yeast)" Mobk1a
1455185_s_at 1.11895 --- ---
1455186_a_at 1.07321 RIKEN cDNA 1190003J15 gene 1190003J15Rik
1455187_at 1.06720 zinc finger and BTB domain containing 40 /// similar to zinc finger and BTB
domain containing 40 Zbtb40 /// LOC638050
1455188_at 0.92348 Eph receptor B1 Ephb1
1455189_at 0.91496 tripartite motif protein 33 /// RIKEN cDNA 8030451N04 gene Trim33 ///
8030451N04Rik
1455190_at 0.92624 "guanine nucleotide binding protein (G protein), gamma 7 subunit" Gng7

1455191_x_at	1.09385	"Phosphatidylinositol-4-phosphate 5-kinase, type 1 beta"	Pip5k1b
1455192_at	0.93236	RIKEN cDNA A230078I05 gene	A230078I05Rik
1455193_at	0.95686	zinc finger and BTB domain containing 8	Zbtb8
1455194_at	0.79297	mitogen-activated protein kinase 8 interacting protein 2	Mapk8ip2
1455195_at	1.34520	ribosomal protein S24	Rps24
1455196_s_at	0.89706	expressed sequence AA987161	AA987161
1455197_at	1.14904	Rho family GTPase 1	Rnd1
1455198_a_at	0.95271	"protein phosphatase 2 (formerly 2A), regulatory subunit B, alpha"	Ppp2r3a
1455199_at	0.98289	expressed sequence AI429214	AI429214
1455200_at	0.85927	p21 (CDKN1A)-activated kinase 6	Pak6
1455201_x_at	1.11310	apolipoprotein A-I	Apoa1
1455202_at	0.83013	"gene model 50, (NCBI)"	Gm50
1455203_at	1.01261	RIKEN cDNA A930003A15 gene	A930003A15Rik
1455204_at	1.01794	"phosphatidylinositol transfer protein, cytoplasmic 1"	Pitpnc1
1455205_a_at	0.80296	ubiquitin specific peptidase 19	Usp19
1455206_at	0.96568	Hypothetical protein C130006E23	C130006E23
1455207_at	1.12103	RIKEN cDNA 2410017P09 gene	2410017P09Rik
1455208_at	0.98078	peroxisome biogenesis factor 19	Pex19
1455209_at	0.94743	"Phosphoenolpyruvate carboxykinase 1, cytosolic"	Pck1
1455210_at	1.01221	zinc fingers and homeoboxes protein 2	Zhx2
1455211_a_at	1.12243	translocase of inner mitochondrial membrane 13 homolog (yeast)	Timm13
1455212_at	0.97386	WD repeat domain 24	Wdr24
1455213_at	0.95747	RIKEN cDNA 4930488E11 gene	4930488E11Rik
1455214_at	0.68622	microphthalmia-associated transcription factor	Mitf
1455215_at	1.09702	RIKEN cDNA C530028O21 gene	C530028O21Rik
1455216_at	1.20765	progesterin and adipoQ receptor family member VI	Paqr6
1455217_at	0.91261	leucine-rich repeats and immunoglobulin-like domains 2	Lrig2
1455218_at	0.95291	RIKEN cDNA 6330503K22 gene	6330503K22Rik
1455219_at	0.96745	RIKEN cDNA 1110030E23 gene	1110030E23Rik
1455220_at	0.91496	frequently rearranged in advanced T-cell lymphomas 2	Frat2
1455221_at	1.02760	"ATP-binding cassette, sub-family G (WHITE), member 1"	Abcg1
1455222_a_at	1.00668	upstream binding protein 1	Ubp1
1455223_at	1.11355	insulin-like growth factor 2 mRNA binding protein 1	Igf2bp1
1455224_at	0.89288	angiopoietin-like 1	Angptl1
1455225_at	1.03986	synaptic nuclear envelope 1	Syne1
1455226_at	0.77583	spectrin beta 1	Spnb1
1455227_at	0.95685	arylacetamide deacetylase-like 1	Aadacl1
1455228_at	1.12053	Wolf-Hirschhorn syndrome candidate 1 (human)	Whsc1
1455229_x_at	1.06896	phosphatidylglycerophosphate synthase 1	Pgs1
1455230_at	1.01171	"calcium channel, voltage-dependent, gamma subunit 4"	Cacng4
1455231_s_at	1.01528	---	---
1455232_at	1.00958	camello-like 2	Cml2
1455233_at	0.94885	mitochondrial ribosomal protein S11	Mrps11

1455234_at 0.88574 "UDP-Gal:betaGlcNAc beta 1,3-galactosyltransferase, polypeptide 1"
 B3galt1
 1455235_x_at 1.01181 lactate dehydrogenase B Ldhb
 1455236_x_at 0.97428 small EDRK-rich factor 2 Serf2
 1455237_at 1.01889 ubiquitin specific peptidase 36 Usp36
 1455238_at 0.70338 melanoma associated antigen (mutated) 1-like 1 Mum111
 1455239_at 0.85357 RIKEN cDNA 6330512M04 gene 6330512M04Rik
 1455240_x_at 1.01532 hypothetical protein LOC666185 /// hypothetical protein LOC666222 ///
 hypothetical protein LOC673030 LOC666185 /// LOC666222 /// LOC673030
 1455241_at 0.98776 cDNA sequence BC037703 BC037703
 1455242_at 0.93922 forkhead box P1 Foxp1
 1455243_at 0.99480 "bromodomain and PHD finger containing, 3" Brpf3
 1455244_at 0.94636 dishevelled associated activator of morphogenesis 1 Daam1
 1455245_x_at 1.04500 ribosomal protein L13 Rpl13
 1455246_at 1.12925 "SWI/SNF related, matrix associated, actin dependent regulator of chromatin,
 subfamily c, member 1" Smarcc1
 1455247_at 1.02061 angiomin-1-like 1 Amotl1
 1455248_at 1.01127 RIKEN cDNA 0610011N22 gene 0610011N22Rik
 1455249_at 0.88823 "10 days neonate skin cDNA, RIKEN full-length enriched library,
 clone:4732480G12 product:unclassifiable, full insert sequence" ---
 1455250_at 0.83573 SH3-domain binding protein 4 Sh3bp4
 1455251_at 0.74334 integrin alpha 1 Itga1
 1455252_at 1.08679 tuberous sclerosis 1 Tsc1
 1455253_at 0.86534 RIKEN cDNA 4930535B03 gene 4930535B03Rik
 1455254_at 0.93366 RIKEN cDNA 4833420G11 gene 4833420G11Rik
 1455255_at 0.99778 RIKEN cDNA 4833420G11 gene 4833420G11Rik
 1455256_at 0.59979 TRAF2 and NCK interacting kinase /// similar to [Segment 1 of 2] Traf2 and
 NCK-interacting kinase /// region containing TRAF2 and NCK interacting kinase; RIKEN cDNA
 C630040K21 gene Tnik /// LOC665113 /// LOC674765
 1455257_at 0.99241 integrin beta 3 Itgb3
 1455258_at 1.00940 "potassium voltage gated channel, Shaw-related subfamily, member 2" Kcnc2
 1455259_a_at 1.17471 RIKEN cDNA D530033C11 gene D530033C11Rik
 1455260_at 0.86179 ligand dependent nuclear receptor corepressor-like Lcorl
 1455261_at 1.14363 "PREDICTED: Mus musculus similar to Hemoglobin alpha subunit
 (Hemoglobin alpha chain) (Alpha-globin) (LOC383229), mRNA" ---
 1455262_at 0.81662 "thrombospondin, type I, domain containing 4" Thsd4
 1455263_at 1.12678 RIKEN cDNA 9030625A04 gene 9030625A04Rik
 1455264_at 0.96307 C1q domain containing 2 C1qdc2
 1455265_a_at 0.99964 regulator of G-protein signaling 16 Rgs16
 1455266_at 0.92186 kinesin family member 5C Kif5c
 1455267_at 1.00901 estrogen-related receptor gamma Esrrg
 1455268_at 0.83397 "zinc finger, CSL domain containing 2" Zcs12
 1455269_a_at 0.93930 "coronin, actin binding protein 1A" Coro1a

1455270_at	1.02485	---	---	
1455271_at	1.08256	hypothetical protein LOC620695	LOC620695	
1455272_at	0.96405	"glutamate receptor, metabotropic 5"	Grm5	
1455273_at	1.04582	---	---	
1455274_at	1.01697	Nonagouti	a	
1455275_at	1.06791	RIKEN cDNA E530001K10 gene	E530001K10Rik	
1455276_x_at	0.89430	RIKEN cDNA 1110004E09 gene	1110004E09Rik	
1455277_at	1.18217	Hedgehog-interacting protein	Hhip	
1455278_at	0.99465	WD repeat domain 37	Wdr37	
1455279_at	0.93272	"gene model 1060, (NCBI)"	Gm1060	
1455280_at	0.98445	Fras1 related extracellular matrix protein 1	Frem1	
1455281_at	0.91041	WD repeat domain 33	Wdr33	
1455282_x_at	1.04345	aminolevulinic acid synthase 1	Alas1	
1455283_x_at	0.95029	NADH dehydrogenase (ubiquinone) Fe-S protein 8	Ndufs8	
1455284_x_at	1.34308	"phosphatidylinositol glycan anchor biosynthesis, class X"	Pigx	
1455285_at	1.07233	"solute carrier family 31, member 1"	Slc31a1	
1455286_at	0.95958	BTB (POZ) domain containing 1	Btbd1	
1455287_at	1.18602	cyclin-dependent kinase 6	Cdk6	
1455288_at	0.91560	RIKEN cDNA 1110036O03 gene	1110036O03Rik	
1455289_at	0.90231	ankyrin repeat domain 13b	Ankrd13b	
1455290_at	0.93473	zinc and ring finger 2	Znrf2	
1455291_s_at	0.99624	zinc and ring finger 2	Znrf2	
1455292_x_at	0.84808	Regulator of sex limited protein-Slp 1	Rsl1	
1455293_at	0.95205	"Leo1, Paf1/RNA polymerase II complex component, homolog (S. cerevisiae)"	Leo1	
1455294_at	1.03084	RIKEN cDNA 1110029L17 gene	1110029L17Rik	
1455295_at	0.96661	cDNA sequence BC031853	BC031853	
1455296_at	0.76131	adenylate cyclase 5	Adcy5	
1455297_at	1.00455	"spindlin family, member 2"	Spin2	
1455298_at	0.92162	Transcribed locus	---	
1455299_at	0.87908	RIKEN cDNA 1700110N18 gene	1700110N18Rik	
1455300_at	0.97509	RIKEN cDNA E130014J05 gene	E130014J05Rik	
1455301_at	0.88138	expressed sequence BQ952480	BQ952480	
1455302_at	0.99136	vacuolar protein sorting 29 (S. pombe)	Vps29	
1455303_at	1.05870	---	---	
1455304_at	1.09867	unc-13 homolog C (C. elegans)	Unc13c	
1455305_x_at	1.02863	heterogeneous nuclear ribonucleoprotein A1	Hnrpa1	
1455306_at	0.88830	cDNA sequence BC037112	BC037112	
1455307_at	0.88730	cDNA sequence BC037112	BC037112	
1455308_at	0.89976	transmembrane protein 16F	Tmem16f	
1455309_at	0.89668	transmembrane protein 16F	Tmem16f	
1455310_at	1.01005	RNA binding motif protein 16	Rbm16	
1455311_at	0.94403	DiGeorge syndrome critical region gene 8	Dgcr8	

1455312_at 1.01768 polyhomeotic-like 3 (Drosophila) Phc3
1455313_at 0.96182 actin-binding LIM protein 2 Ablim2
1455314_at 0.96109 LIM domain containing preferred translocation partner in lipoma Lpp
1455315_at 0.86899 tumor protein p53 inducible protein 13 Trp53i13
1455316_x_at 1.15032 "Solute carrier family 15 (H+/peptide transporter), member 2 /// Potassium voltage-gated channel, subfamily H (eag-related), member 6 /// 17 days pregnant adult female amnion cDNA, RIKEN full-length enriched library, clone:I920040L03 product:unclassifiable, full insert sequence" Slc15a2 /// Kcnh6
1455317_at 1.01053 enhancer of polycomb homolog 2 (Drosophila) Epc2
1455318_at 0.95866 T-cell immunoglobulin and mucin domain containing 4 Timd4
1455319_x_at 1.07796 ribosomal protein S8 Rps8
1455320_at 1.25985 --- ---
1455321_at 1.06859 DDHD domain containing 1 Ddhd1
1455322_at 1.04256 tetratricopeptide repeat domain 21B Ttc21b
1455323_at 0.82653 RB-associated KRAB repressor Rbak
1455324_at 0.96986 hypothetical LOC433022 LOC433022
1455325_at 0.95595 RIKEN cDNA A230057G18 gene A230057G18Rik
1455326_at 0.99147 RIKEN cDNA 4932416N17 gene 4932416N17Rik
1455327_at 1.12712 SUMO/sentrin specific peptidase 2 Senp2
1455328_at 0.96739 "amiloride-sensitive cation channel 2, neuronal" Accn2
1455329_at 1.11202 --- ---
1455330_at 0.98822 nucleolar protein 9 Nol9
1455331_at 0.93329 RIKEN cDNA 9430067K14 gene 9430067K14Rik
1455332_x_at 0.95811 "Fc receptor, IgG, low affinity IIb" Fcgr2b
1455333_at 0.96529 tensin 3 Tns3
1455334_at 1.00418 RIKEN cDNA 9830167H18 gene 9830167H18Rik
1455335_at 0.88765 X-ray repair complementing defective repair in Chinese hamster cells 2 Xrcc2
1455336_at 0.90013 "THAP domain containing, apoptosis associated protein 2" Thap2
1455337_at 1.02485 "FYVE, RhoGEF and PH domain containing 4" Fgd4
1455338_at 1.10988 "alpha 1,4-galactosyltransferase" A4galt
1455339_at 1.11019 RIKEN cDNA C430014M02 gene C430014M02Rik
1455340_at 1.02900 Expressed sequence AI852444 AI852444
1455341_at 1.09217 RIKEN cDNA 2010003J03 gene 2010003J03Rik
1455342_at 1.05045 RIKEN cDNA A230083H22 gene A230083H22Rik
1455343_at 0.89931 "pleckstrin homology domain containing, family A member 7" Plekha7
1455344_at 1.08793 ephrin A3 Efna3
1455345_at 1.19483 PHD finger protein 15 Phf15
1455346_at 0.64775 mannan-binding lectin serine peptidase 1 Masp1
1455347_at 0.98139 regulator of telomere elongation helicase 1 Rtel1
1455348_x_at 1.12203 ribosomal protein L29 /// similar to 60S ribosomal protein L29 /// similar to 60S ribosomal protein L29 /// similar to 60S ribosomal protein L29 /// similar to 60S ribosomal protein L29 /// similar to 60S ribosomal protein L29 /// similar to 60S ribosomal protein L29 /// similar to 60S ribosomal protein L29 /// hypothetical protein LOC669999 /// similar to 60S ribosomal protein L29 ///

similar to 60S ribosomal protein L29 Rpl29 /// LOC433350 /// LOC433941 /// LOC622707 ///
 LOC626772 /// LOC665032 /// LOC666642 /// LOC667466 /// LOC669999 /// LOC670334 ///
 LOC675846

1455349_at 1.03490 RAS related protein 1b Rap1b
 1455350_at 1.03349 transmembrane protein 62 Tmem62
 1455351_at 0.91450 expressed sequence AU023006 AU023006
 1455352_at 1.02571 expressed sequence AU023006 AU023006
 1455353_at 0.90854 transmembrane and coiled coil domains 1 Tmcc1
 1455354_at 0.96346 WD repeat domain 22 Wdr22
 1455355_at 1.04162 RIKEN cDNA 6030408C04 gene 6030408C04Rik
 1455356_at 0.94008 calmodulin regulated spectrin-associated protein 1 Camsap1
 1455357_x_at 1.10037 translocase of outer mitochondrial membrane 20 homolog (yeast) Tomm20
 1455358_at 1.00275 Ataxin 2 binding protein 1 A2bp1
 1455359_at 0.96901 "protein tyrosine phosphatase, non-receptor type 14" Ptpn14
 1455360_at 1.02348 RIKEN cDNA 4833412L08 gene 4833412L08Rik
 1455361_at 0.98998 "diacylglycerol kinase, beta" Dgkb
 1455362_at 0.97160 --- ---
 1455363_at 1.15982 brain-specific angiogenesis inhibitor 1 Bai1
 1455364_a_at 1.06099 ribosomal protein S7 /// similar to 40S ribosomal protein S7 (S8) Rps7 ///
 LOC640050

1455365_at 0.99234 cadherin 8 Cdh8
 1455366_at 0.96001 mitochondrial ribosomal protein L52 Mrpl52
 1455367_at 0.99708 dead end homolog 1 (zebrafish) Dnd1
 1455368_at 1.04990 "zinc finger, DHHC domain containing 3" Zdhhc3
 1455369_at 0.91129 "amyloid beta (A4) precursor protein binding, family A, member 1" Apba1
 1455370_at 1.14327 RIKEN cDNA A630023P12 gene A630023P12Rik
 1455371_at 0.94457 regulatory factor X-associated ankyrin-containing protein Rfxank
 1455372_at 0.73093 cytoplasmic polyadenylation element binding protein 3 Cpeb3
 1455373_at 1.07647 Transcribed locus ---
 1455374_at 0.89769 --- ---
 1455375_at 0.86735 similar to Protein enabled homolog (NPC-derived proline-rich protein 1)
 (NDPP-1) LOC669787

1455376_at 0.77468 RIKEN cDNA 1300010F03 gene 1300010F03Rik
 1455377_at 0.97528 RIKEN cDNA 4921517B04 gene 4921517B04Rik
 1455378_at 0.89043 cDNA sequence BC057371 BC057371
 1455379_at 1.05790 "dynein, axonemal, light intermediate polypeptide 1" Dnali1
 1455380_at 1.01002 prenylcysteine oxidase 1 like Pcyox11
 1455381_at 0.84202 RIKEN cDNA 4921513D23 gene 4921513D23Rik
 1455382_at 1.00662 ZW10 interactor Zwint
 1455383_at 0.95787 hypothetical protein C730043O17 C730043O17
 1455384_x_at 0.99285 RIKEN cDNA D030056L22 gene D030056L22Rik
 1455385_at 1.11275 exocyst complex component 6 Exoc6
 1455386_at 1.26827 Transcribed locus ---

1455387_at 1.09805 nuclear fragile X mental retardation protein interacting protein 2 Nufip2
1455388_at 0.88416 protein-L-isoaspartate (D-aspartate) O-methyltransferase domain containing 1 Pcmd1
1455389_s_at 1.01445 RIKEN cDNA 2310051F07 gene 2310051F07Rik
1455390_at 0.92620 "RIKEN cDNA 1500005P14 gene /// alkB, alkylation repair homolog 6 (E. coli)" 1500005P14Rik /// Alkbh6
1455391_at 1.08529 RAD23a homolog (S. cerevisiae) Rad23a
1455392_at 1.07736 RIKEN cDNA 9630019K15 gene 9630019K15Rik
1455393_at 1.12626 --- ---
1455394_at 1.01203 protein inhibitor of activated STAT 4 Pias4
1455395_at 1.06090 "3-oxoacyl-ACP synthase, mitochondrial" Oxsm
1455396_at 0.93132 "ATPase, class I, type 8B, member 1" Atp8b1
1455397_at 0.98471 CASK interacting protein 1 Caskin1
1455398_at 0.73983 "leucine rich repeat containing 8 family, member C" Lrrc8c
1455399_at 0.79971 connector enhancer of kinase suppressor of Ras 1 /// similar to connector enhancer of kinase suppressor of Ras 1 Cnksr1 /// LOC637116
1455400_at 1.04172 dimethylarginine dimethylaminohydrolase 1 Ddah1
1455401_at 1.00776 "calcium/calmodulin-dependent protein kinase kinase 2, beta" Camkk2
1455402_at 0.85022 suppressor of cytokine signaling 7 Socs7
1455403_at 0.98424 "mannosidase, endo-alpha" Manea
1455404_at 0.99289 junctophilin 2 Jph2
1455405_at 0.83710 proline-serine-threonine phosphatase-interacting protein 2 Pstpip2
1455406_at 0.95708 "0 day neonate head cDNA, RIKEN full-length enriched library, clone:4833431M13 product:unclassifiable, full insert sequence" ---
1455407_at 0.91979 zinc finger protein 236 Zfp236
1455408_at 0.90828 "phospholipase A2, group IVF" Pla2g4f
1455409_at 1.24010 spire homolog 1 (Drosophila) Spire1
1455410_at 0.95878 Fas apoptotic inhibitory molecule 2 Faim2
1455411_at 0.92648 aquaporin 11 Aqp11
1455412_at 0.86936 --- ---
1455413_at 0.93594 zinc finger protein 11 Zfp11
1455414_at 0.97182 tetratricopeptide repeat domain 16 Ttc16
1455415_at 1.07660 RIKEN cDNA A730056A06 gene A730056A06Rik
1455416_at 0.96544 Riken cDNA C130021I20 gene C130021I20Rik
1455417_at 0.90982 "potassium inwardly rectifying channel, subfamily J, member 11" Kcnj11
1455418_at 0.80449 Transcribed locus ---
1455419_at 0.97890 RIKEN cDNA D630045J12 gene D630045J12Rik
1455420_at 1.03862 RAD23b homolog (S. cerevisiae) Rad23b
1455421_x_at 0.95287 RIKEN cDNA 6330503C03 gene 6330503C03Rik
1455422_x_at 1.16456 septin 4 4-Sep
1455423_at 0.98741 Nur77 downstream gene 1 /// similar to chromosome 6 open reading frame 148 Ndg1 /// LOC623189
1455424_at 0.93943 WNK lysine deficient protein kinase 2 Wnk2

1455425_at 1.04914 expressed sequence BB001228 BB001228
1455426_at 0.94282 --- ---
1455427_at 0.96683 angiopoietin 4 Angpt4
1455428_at 0.89356 RIKEN cDNA A930008G19 gene A930008G19Rik
1455429_at 0.91132 expressed sequence AI450540 AI450540
1455430_at 1.00993 RIKEN cDNA 3200001K10 gene 3200001K10Rik
1455431_at 1.02759 "solute carrier family 5 (sodium/glucose cotransporter), member 1" Slc5a1
1455432_at 0.92102 TAO kinase 1 Taok1
1455433_at 1.12343 hypothetical protein LOC383216 MGC107415
1455434_a_at 1.08477 kinectin 1 Ktn1
1455435_s_at 1.08420 --- ---
1455436_at 1.02448 "DIRAS family, GTP-binding RAS-like 2" Diras2
1455437_at 0.98382 cDNA sequence BC033915 BC033915
1455438_at 0.81554 peroxisomal membrane protein 4 Pxmp4
1455439_a_at 0.98493 "lectin, galactose binding, soluble 1" Lgals1
1455440_at 1.18206 chloride channel 6 Clcn6
1455441_at 0.87869 --- ---
1455442_at 0.96460 "solute carrier family 6 (neurotransmitter transporter), member 19" Slc6a19
1455443_at 1.01191 ganglioside-induced differentiation-associated protein 1-like 1 Gdap111
1455444_at 0.94229 "gamma-aminobutyric acid (GABA-A) receptor, subunit alpha 2" Gabra2
1455445_at 0.69005 cerebellin 3 precursor protein Cbln3
1455446_x_at 0.94104 "acyl-Coenzyme A dehydrogenase, short/branched chain" Acadsb
1455447_at 1.07772 RIKEN cDNA D430019H16 gene D430019H16Rik
1455448_at 1.00198 5'-nucleotidase domain containing 3 Nt5dc3
1455449_at 1.02064 "gene model 468, (NCBI)" Gm468
1455450_at 0.93578 "protein tyrosine phosphatase, non-receptor type 3" Ptpn3
1455451_at 0.95966 expressed sequence AI449310 AI449310
1455452_x_at 1.09275 expressed sequence AI449310 AI449310
1455453_at 1.15490 gem (nuclear organelle) associated protein 5 Gemin5
1455454_at 0.86385 "aldo-keto reductase family 1, member C19" Akr1c19
1455455_at 0.87295 glycosyltransferase 28 domain containing 2 Glt28d2
1455456_a_at 0.93279 translocase of inner mitochondrial membrane 50 homolog (yeast) Timm50
1455457_at 0.94650 "cytochrome P450, family 2, subfamily c, polypeptide 50 /// cytochrome P450" Cyp2c50 /// Cyp2c54
1455458_x_at 0.99130 GDP-mannose pyrophosphorylase A Gmppa
1455459_at 1.04588 PR domain containing 15 Prdm15
1455460_at 0.99314 RIKEN cDNA 6430547I21 gene 6430547I21Rik
1455461_at 0.96971 RIKEN cDNA A530017D24 gene A530017D24Rik
1455462_at 1.07004 adenylate cyclase 2 Adcy2
1455463_at 0.97283 phytanoyl-CoA hydroxylase interacting protein Phyhip
1455464_x_at 1.48893 --- ---
1455465_at 0.97869 Sorting nexin 12 Snx12
1455466_at 0.97169 G protein-coupled receptor 133 /// similar to G protein-coupled receptor 133

Gpr133 /// LOC669205

1455467_at 0.76584 RIKEN cDNA C330043M08 gene C330043M08Rik
 1455468_at 0.95424 "0 day neonate kidney cDNA, RIKEN full-length enriched library, clone: D630029P09 product:unclassifiable, full insert sequence" ---
 1455469_at 0.93692 "solute carrier family 6 (neurotransmitter transporter, L-proline), member 7" Slc6a7
 1455470_x_at 1.05821 LIM and SH3 protein 1 Lasp1
 1455471_at 0.97597 RIKEN cDNA 1500012D20 gene 1500012D20Rik
 1455472_at 1.04251 RIKEN cDNA A630071D13 gene A630071D13Rik
 1455473_at 1.00626 ubiquitin specific peptidase 13 (isopeptidase T-3) Usp13
 1455474_at 1.20717 "DNA segment, Chr 6, Wayne State University 116, expressed" D6Wsu116e
 1455475_at 1.04247 RIKEN cDNA 3110057O12 gene 3110057O12Rik
 1455476_a_at 0.87067 genetic suppressor element 1 Gsel
 1455477_s_at 1.03773 PDZK1 interacting protein 1 Pdzk1ip1
 1455478_at 1.23334 "protein tyrosine phosphatase, receptor type, f polypeptide (PTPRF), interacting protein, alpha 1" Ppfia1
 1455479_a_at 1.04777 "ubiquitin-conjugating enzyme E2D 3 (UBC4/5 homolog, yeast)" Ube2d3
 1455480_s_at 1.05693 "ubiquitin-conjugating enzyme E2D 3 (UBC4/5 homolog, yeast)" Ube2d3
 1455481_at 0.88761 iduronate 2-sulfatase Ids
 1455482_at 1.08330 "adaptor protein complex AP-2, alpha 2 subunit" Ap2a2
 1455483_at 0.92429 zinc finger protein 148 Zfp148
 1455484_at 1.04018 RIKEN cDNA 2600009P04 gene 2600009P04Rik
 1455485_x_at 1.00415 ribosomal protein L13a Rpl13a
 1455486_at 0.87994 protein inhibitor of activated STAT 1 Pias1
 1455487_at 0.93693 RIKEN cDNA 2600014M03 gene 2600014M03Rik
 1455488_at 1.07197 RIKEN cDNA 6230416J20 gene 6230416J20Rik
 1455489_at 1.01294 leucine rich repeat transmembrane neuronal 2 Lrrtm2
 1455490_at 0.94727 "upstream binding transcription factor, RNA polymerase I" Ubtf
 1455491_at 1.01074 heterogeneous nuclear ribonucleoprotein H3 /// similar to heterogeneous nuclear ribonucleoprotein H3 isoform a Hnrph3 /// LOC669773
 1455492_at 0.98589 RIKEN cDNA B330016D10 gene B330016D10Rik
 1455493_at 1.11159 synaptic nuclear envelope 1 Syne1
 1455494_at 0.92578 "procollagen, type I, alpha 1" Coll1a1
 1455495_at 1.00586 Expressed sequence AA536808 AA536808
 1455496_at 1.09417 phosphoribosylformylglycinamide synthase (FGAR amidotransferase) Pfas
 1455497_at 0.92392 leukocyte receptor cluster (LRC) member 9 Leng9
 1455498_at 1.08387 G-protein-coupled receptor 50 Gpr50
 1455499_at 0.99742 neurexin II Nrnx2
 1455500_at 1.01572 "DNA segment, Chr 11, ERATO Doi 759, expressed /// similar to chromosome 17 open reading frame 27" D11Ertd759e /// LOC672513
 1455501_at 1.05999 "solute carrier family 2 (facilitated glucose transporter), member 12" Slc2a12
 1455502_at 0.98685 MAP-kinase activating death domain Madd
 1455503_at 0.92560 "zinc finger protein 85, related sequence 1 /// zinc finger protein 273"

Zfp85-rs1 /// Zfp273

1455504_a_at	1.00852	"makorin, ring finger protein, 1"	Mkrn1
1455505_at	0.96264	GATA zinc finger domain containing 2A	Gatad2a
1455506_at	1.02331	"solute carrier family 25, member 34"	Slc25a34
1455507_s_at	0.89939	"DNA segment, Chr 8, ERATO Doi 587, expressed"	D8Ert587e
1455508_at	1.00497	RIKEN cDNA A530082C11 gene	A530082C11Rik
1455509_at	0.82132	cDNA sequence AB182283	AB182283
1455510_at	1.70333	Speckle-type POZ protein	Spop
1455511_at	0.77496	selenophosphate synthetase 1	Sephs1
1455512_at	0.83008	"gene model 879, (NCBI)"	Gm879
1455513_at	1.23514	"TAF1 RNA polymerase II, TATA box binding protein (TBP)-associated factor"	Taf1
1455514_at	1.05436	"potassium voltage-gated channel, Shal-related family, member 1"	Kcnd1
1455515_at	0.98653	RIKEN cDNA 1810041L15 gene	1810041L15Rik
1455516_at	0.93516	Transcribed locus	---
1455517_at	1.07111	RNA binding motif protein 4	Rbm4
1455518_at	0.93074	"Transcribed locus, weakly similar to XP_232745.3 PREDICTED: hypothetical protein XP_232745 [Rattus norvegicus]"	---
1455519_at	1.01858	desmoglein 1 beta	Dsg1b
1455520_at	0.98093	hypothetical protein LOC629605	LOC629605
1455521_at	0.96668	Kruppel-like factor 12	Klf12
1455522_at	0.88404	Rho guanine nucleotide exchange factor (GEF) 15	Arhgef15
1455523_at	0.92935	"cleavage stimulation factor, 3' pre-RNA subunit 2"	Cstf2
1455524_at	1.08203	RIKEN cDNA 2810046L04 gene	2810046L04Rik
1455525_at	0.98837	endonuclease G-like 1	Endog1
1455526_at	1.09719	"DIRAS family, GTP-binding RAS-like 1"	Diras1
1455527_at	1.12161	RIKEN cDNA E430002D04 gene	E430002D04Rik
1455528_at	0.76046	"Transcribed locus, weakly similar to NP_665733.1 Lipogenin [Rattus norvegicus]"	---
1455529_at	0.84857	RIKEN cDNA 2700083E18 gene	2700083E18Rik
1455530_at	0.96113	Immunoglobulin heavy chain 1a (serum IgG2a)	Igh-1a
1455531_at	1.06564	RIKEN cDNA A930031D07 gene	A930031D07Rik
1455532_at	1.05475	ataxin 7-like 2	Atxn7l2
1455533_at	1.08677	RIKEN cDNA A930025D01 gene	A930025D01Rik
1455534_s_at	0.94047	oxysterol binding protein-like 11	Osbpl11
1455535_at	0.85704	RIKEN cDNA A730017D01 gene	A730017D01Rik
1455536_at	0.97052	RIKEN cDNA A630023A22 gene	A630023A22Rik
1455537_at	0.92293	RIKEN cDNA 6430547I21 gene	6430547I21Rik
1455538_at	1.08897	RIKEN cDNA 6330403M23 gene	6330403M23Rik
1455539_at	1.03058	---	---
1455540_at	1.00972	carbamoyl-phosphate synthetase 1	Cps1
1455541_a_at	0.98173	RIKEN cDNA 4430402I18 gene	4430402I18Rik
1455542_at	0.99696	RIKEN cDNA C630043F03 gene	C630043F03Rik

1455543_at 0.96963 kelch-like 18 (Drosophila) Klhl18
1455544_at 0.83375 "zinc finger, RAN-binding domain containing 3" Zranb3
1455545_at 0.96501 RIKEN cDNA 1110065P20 gene 1110065P20Rik
1455546_s_at 1.01685 "splicing factor 3a, subunit 2" Sf3a2
1455547_at 0.90018 zinc finger CCCH type containing 7B Zc3h7b
1455548_at 0.85663 "discs, large homolog-associated protein 4 (Drosophila)" Dlgap4
1455549_at 0.97985 SEC14 and spectrin domains 1 Sestd1
1455550_x_at 0.86547 serine incorporator 3 Serinc3
1455551_at 0.98458 UEV and lactate/malate dehydrogenase domains Uevld
1455552_at 1.07715 "small nuclear RNA activating complex, polypeptide 4" Snapc4
1455553_at 0.87971 RIKEN cDNA 5330401P04 gene 5330401P04Rik
1455554_at 0.87285 RIKEN cDNA A830039N20 gene A830039N20Rik
1455555_at 0.94071 Expressed sequence AV071699 AV071699
1455556_at 0.96966 Notch gene homolog 2 (Drosophila) Notch2
1455557_at 1.06028 hypothetical LOC553095 LOC553095
1455558_at 0.78575 "gene model 114, (NCBI)" Gm114
1455559_at 1.07978 RIKEN cDNA E330039K12 gene E330039K12Rik
1455560_at 0.98511 Zinc finger protein 653 Zfp653
1455561_at 0.97272 carnosine dipeptidase 1 (metallopeptidase M20 family) Cndp1
1455562_at 0.99446 SRY-box containing gene 12 Sox12
1455563_at 1.01923 DEAD (Asp-Glu-Ala-Asp) box polypeptide 49 Ddx49
1455564_at 1.07734 breakpoint cluster region homolog Bcr
1455565_at 1.52752 B-cell CLL/lymphoma 9 Bcl9
1455566_s_at 1.04494 RIKEN cDNA 2810022L02 gene 2810022L02Rik
1455567_at 1.11819 "Cdc2-related kinase, arginine/serine-rich" Crkrs
1455568_at 0.96717 RIKEN cDNA 2310015A05 gene 2310015A05Rik
1455569_at 0.88527 CDNA clone IMAGE:1548559 ---
1455570_x_at 0.84680 "calponin 3, acidic" Cnn3
1455571_x_at 1.03802 calmodulin 1 Calm1
1455572_x_at 1.04764 ribosomal protein S18 Rps18
1455573_at 0.86559 keratin 14 Krt14
1455574_at 1.05071 "Smg-7 homolog, nonsense mediated mRNA decay factor (C. elegans)"
Smg7
1455575_at 0.91472 eukaryotic translation initiation factor 4E binding protein 2 Eif4ebp2
1455576_at 0.78650 RIKEN cDNA 5830482F20 gene 5830482F20Rik
1455577_at 1.08250 Chemokine (C-C motif) ligand 28 Ccl28
1455578_x_at 1.01234 ribosomal protein L41 Rpl41
1455579_at 1.03028 casein alpha s2-like A Csn1s2a
1455580_at 1.10026 USP6 N-terminal like Usp6nl
1455581_x_at 0.87826 hypothetical protein 9530028C05 9530028C05
1455582_at 0.65954 Transcribed locus ---
1455583_at 0.87719 glucosamine Gne
1455584_at 0.86342 stromal cell derived factor 4 Sdf4

1455585_at 1.00098 ring finger protein 168 Rnf168
1455586_at 0.93550 ring finger protein 168 Rnf168
1455587_at 0.94250 cDNA sequence BC030183 BC030183
1455588_at 0.82238 cDNA sequence BC034664 BC034664
1455589_at 1.00595 RIKEN cDNA 1700100M05 gene 1700100M05Rik
1455590_at 1.03564 "NAD(P)H dehydrogenase, quinone 2" Nqo2
1455591_at 0.98923 --- ---
1455592_at 1.01603 "vang-like 2 (van gogh, Drosophila)" Vangl2
1455593_at 1.09198 apolipoprotein B Apob
1455594_at 0.78637 exocyst complex component 3 Exoc3
1455595_at 0.73327 "UDP glucuronosyltransferase 2 family, polypeptide B36" Ugt2b36
1455596_a_at 1.02409 cDNA sequence BC029214 BC029214
1455597_at 1.00540 Expressed sequence AI585793 AI585793
1455598_at 0.94041 ubiquitin specific peptidase 30 Usp30
1455599_at 0.91816 glucose-fructose oxidoreductase domain containing 1 Gfod1
1455600_at 1.16324 ribosomal protein S3 Rps3
1455601_at 0.99491 "0 day neonate thymus cDNA, RIKEN full-length enriched library, clone:
A430079C03 product:unclassifiable, full insert sequence" ---
1455602_x_at 0.83484 RIKEN cDNA 1110028C15 gene 1110028C15Rik
1455603_at 0.86850 Transcribed locus ---
1455604_at 0.68628 frizzled homolog 5 (Drosophila) Fzd5
1455605_at 1.16000 RUN and FYVE domain containing 3 Rufy3
1455606_at 0.96186 cDNA sequence BC004022 BC004022
1455607_at 0.97880 R-spondin 3 homolog (Xenopus laevis) Rspo3
1455608_at 0.95452 sodium channel and clathrin linker 1 Sclt1
1455609_at 0.94193 RIKEN cDNA C030025P15 gene C030025P15Rik
1455610_at 0.81465 desmuslin Dmn
1455611_at 0.82386 protein inhibitor of activated STAT 1 Pias1
1455612_at 0.84629 expressed sequence AI848218 AI848218
1455613_at 0.85542 RIKEN cDNA E130308A19 gene E130308A19Rik
1455614_at 0.90891 translocase of outer mitochondrial membrane 40 homolog-like (yeast)
Tomm40l
1455615_at 1.02578 chemokine-like factor Cklf
1455616_at 1.30950 expressed sequence AI586002 AI586002
1455617_at 0.88928 LMBR1 domain containing 1 Lmbrd1
1455618_x_at 0.95013 tetraspanin 33 Tspan33
1455619_at 1.04978 cDNA sequence BC062258 BC062258
1455620_at 0.89637 "PREDICTED: Mus musculus similar to heparan sulfate D-glucosaminyl 3-O-
sulfotransferase 4 (LOC628805), mRNA" ---
1455621_at 0.94952 cDNA sequence BC066107 BC066107
1455622_at 1.03933 podocalyxin-like 2 Podxl2
1455623_at 0.86013 RIKEN cDNA A930031D07 gene A930031D07Rik
1455624_at 0.96053 --- ---

1455625_at	0.78053	---	---	
1455626_at	0.92855	homeo box A9	Hoxa9	
1455627_at	1.19361	"procollagen, type VIII, alpha 1"	Col8a1	
1455628_at	1.04061	RIKEN cDNA 6430543G08 gene	6430543G08Rik	
1455629_at	0.96510	dopamine receptor D1A	Drd1a	
1455630_at	0.94470	RIKEN cDNA 1700010H15 gene	1700010H15Rik	
1455631_at	0.91004	histocompatibility 13	H13	
1455632_at	0.96448	"Guanine nucleotide binding protein, beta 5"	Gnb5	
1455633_at	0.79775	zinc finger protein 647	Zfp647	
1455634_at	0.94829	---	---	
1455635_at	1.03277	Transcribed locus	---	
1455636_at	0.95244	limbic system-associated membrane protein	Lsamp	
1455637_x_at	0.95984	Similar to hypothetical protein FLJ38281	LOC245305	
1455638_at	0.72521	zinc finger protein 319	Zfp319	
1455639_at	0.95155	Transcribed locus	---	
1455640_a_at	0.95910	thioredoxin 2	Txn2	
1455641_at	0.96640	RIKEN cDNA 1600014C10 gene	1600014C10Rik	
1455642_a_at	0.79109	tetraspanin 17	Tspan17	
1455643_s_at	1.10101	"TSR1, 20S rRNA accumulation, homolog (yeast)"	Tsr1	
1455644_at	0.99148	vacuolar protein sorting 53 (yeast)	Vps53	
1455645_at	0.77983	RIKEN cDNA 8030451F13 gene	8030451F13Rik	
1455646_at	0.79885	RIKEN cDNA 2010004M13 gene	2010004M13Rik	
1455647_at	0.89341	androgen receptor	Ar	
1455648_at	1.02617	similar to reduced expression 2	/// similar to reduced expression 2	
LOC278757	///	LOC631624		
1455649_at	0.83763	tetratricopeptide repeat domain 9	Ttc9	
1455650_at	0.91578	Transcribed locus	---	
1455651_at	0.94271	telomeric repeat binding factor 2	Terf2	
1455652_at	0.99874	expressed sequence AW124694	AW124694	
1455653_at	0.96277	cyclin J	Ccnj	
1455654_at	0.72903	RIKEN cDNA 6430706D22 gene	6430706D22Rik	
1455655_a_at	1.13314	TAR DNA binding protein	Tardbp	
1455656_at	0.96581	B and T lymphocyte associated	Btla	
1455657_at	1.17047	RIKEN cDNA 2610207I05 gene	2610207I05Rik	
1455658_at	1.05510	CGG triplet repeat binding protein 1	Cggbp1	
1455659_at	1.01953	---	---	
1455660_at	0.82685	"colony stimulating factor 2 receptor, beta 1, low-affinity (granulocyte-macrophage)"	Csf2rb1	
1455661_at	1.04683	"0 day neonate thymus cDNA, RIKEN full-length enriched library, clone: A430106B04 product:hypothetical protein, full insert sequence"	---	
1455662_x_at	1.08266	ribosomal protein S17	Rps17	
1455663_at	1.08959	olfactomedin-like 1	Olfml1	
1455664_at	0.69246	reticulon 4 receptor-like 1	Rtn4rl1	

1455665_at 1.01600 "LON peptidase N-terminal domain and ring finger 1 /// similar to CG32369-PB, isoform B" Lonrf1 /// LOC631639
 1455666_at 0.98713 RIKEN cDNA A030001O10 gene A230042K10Rik
 1455667_at 0.81312 prolactin regulatory element binding Preb
 1455668_at 0.96353 Wolf-Hirschhorn syndrome candidate 1-like 1 (human) Whsc111
 1455669_at 0.79039 WD repeat domain 19 Wdr19
 1455670_at 1.13472 "15 days embryo head cDNA, RIKEN full-length enriched library, clone: D930003N06 product:unclassifiable, full insert sequence" ---
 1455671_at 0.97713 COMM domain containing 8 Commd8
 1455672_s_at 0.95948 complexin 2 Cplx2
 1455673_at 0.95856 "alkB, alkylation repair homolog 2 (E. coli)" Alkbh2
 1455674_at 1.04266 WD repeat domain 76 Wdr76
 1455675_a_at 1.04279 Tial1 cytotoxic granule-associated RNA binding protein-like 1 Tial1
 1455676_x_at 1.17334 Tial1 cytotoxic granule-associated RNA binding protein-like 1 Tial1
 1455677_s_at 0.88758 chloride channel Ka /// similar to Chloride channel protein ClC-Ka (Chloride channel Ka) (ClC-K1) Clnka /// LOC639048
 1455678_at 0.94354 "sema domain, immunoglobulin domain (Ig), transmembrane domain (TM) and short cytoplasmic domain, (semaphorin) 4B" Sema4b
 1455679_at 0.89603 RIKEN cDNA 5830411E10 gene 5830411E10Rik
 1455680_at 1.03204 RIKEN cDNA 9630025H16 gene 9630025H16Rik
 1455681_at 0.92389 zinc finger protein 369 Zfp369
 1455682_at 1.19101 expressed sequence AA536808 AA536808
 1455683_a_at 0.96075 "TBC1 domain family, member 8" Tbc1d8
 1455684_at 0.97933 mitogen-activated protein kinase kinase kinase 9 Map3k9
 1455685_at 1.14122 "microtubule associated monooxygenase, calponin and LIM domain containing 2" Mical2
 1455686_at 0.89003 --- ---
 1455687_at 0.91781 intestinal cell kinase Ick
 1455688_at 1.21062 "discoidin domain receptor family, member 2" Ddr2
 1455689_at 0.95864 frizzled homolog 10 (Drosophila) Fzd10
 1455690_at 0.94847 --- ---
 1455691_at 1.15968 "cytochrome P450, family 21, subfamily a, polypeptide 1" Cyp21a1
 1455692_x_at 1.04083 RIKEN cDNA 1700097N02 gene 1700097N02Rik
 1455693_x_at 1.06245 ribosomal protein S6 /// similar to 40S ribosomal protein S6 /// similar to 40S ribosomal protein S6 /// similar to 40S ribosomal protein S6 /// similar to 40S ribosomal protein S6 /// similar to 40S ribosomal protein S6 /// similar to 40S ribosomal protein S6 /// similar to 40S ribosomal protein S6 Rps6 /// LOC434404 /// LOC629575 /// LOC638497 /// LOC666577 /// LOC667739 /// LOC667922 /// LOC671306 /// LOC671432 /// LOC672192
 1455694_at 0.94606 neurobeachin-like 2 Nbeal2
 1455695_at 1.04816 "ST8 alpha-N-acetyl-neuraminide alpha-2,8-sialyltransferase 1" St8sia1
 1455696_a_at 0.95835 PRP4 pre-mRNA processing factor 4 homolog B (yeast) Prpf4b
 1455697_at 0.98622 "PREDICTED: Mus musculus hypothetical protein LOC629064 (LOC629064),

mRNA" ---

1455698_at 0.80135 translocation protein 1 Tloc1
 1455699_at 1.01769 Transcribed locus ---
 1455700_at 0.89214 MTERF domain containing 3 Mterfd3
 1455701_at 0.98025 sorting nexin 26 Snx26
 1455702_at 0.89611 --- ---
 1455703_at 0.89472 thymoma viral proto-oncogene 2 Akt2
 1455704_at 0.80201 X Kell blood group precursor related family member 8 homolog Xkr8
 1455705_at 0.96341 transmembrane protein 161A Tmem161a
 1455706_at 0.76581 syntaxin binding protein 4 Stxbp4
 1455707_at 1.04424 RIKEN cDNA C130037N17 gene C130037N17Rik
 1455708_at 0.96919 Tropomodulin 3 Tmod3
 1455709_at 1.80597 RIKEN cDNA 4930422I22 gene 4930422I22Rik
 1455710_x_at 0.95107 mature T-cell proliferation 1 Mtcp1
 1455711_at 1.01019 deltex 4 homolog (Drosophila) Dtx4
 1455712_at 1.01932 "histone 3, H2a" Hist3h2a
 1455713_x_at 1.09274 prohibitin 2 Phb2
 1455714_at 0.98047 "gene model 691, (NCBI)" Gm691
 1455715_at 0.79868 RIKEN cDNA 2610044O15 gene 2610044O15Rik
 1455716_at 1.12467 "Adult male corpora quadrigemina cDNA, RIKEN full-length enriched library, clone:B230342J21 product:unclassifiable, full insert sequence" ---
 1455717_s_at 0.70313 dishevelled associated activator of morphogenesis 2 Daam2
 1455718_at 1.00556 spermatogenesis associated 21 Spata21
 1455719_at 0.94550 "tubulin, beta 5" Tubb5
 1455720_at 0.78439 "a disintegrin-like and metallopeptidase (reprolysin type) with thrombospondin type 1 motif, 2" Adamts2
 1455721_at 0.93403 G1 to S phase transition 2 Gspt2
 1455722_at 0.98252 --- ---
 1455723_at 0.84089 "DNA segment, Chr 1, ERATO Doi 448, expressed" D1Ertd448e
 1455724_at 0.88614 "2 days pregnant adult female oviduct cDNA, RIKEN full-length enriched library, clone:E230030F22 product:unclassifiable, full insert sequence" ---
 1455725_a_at 1.03516 "H3 histone, family 3B" H3f3b
 1455726_at 1.08452 "gene model 71, (NCBI)" Gm71
 1455727_at 1.17829 "U2 small nuclear ribonucleoprotein auxiliary factor (U2AF) 1, related sequence 2" U2af1-rs2
 1455728_at 0.96609 phosphatase and tensin homolog Pten
 1455729_at 0.93388 "guanine nucleotide binding protein, alpha q polypeptide" Gnaq
 1455730_at 1.11216 "discs, large homolog 7 (Drosophila)" Dlg7
 1455731_at 0.90935 "solute carrier family 29 (nucleoside transporters), member 3" Slc29a3
 1455732_at 1.18208 RIKEN cDNA 1700025G04 gene 1700025G04Rik
 1455733_at 1.29616 TAO kinase 3 Taok3
 1455734_at 0.86800 cereblon Crbn
 1455735_at 0.97923 "adaptor-related protein complex AP-1, sigma 3" Ap1s3

1455736_at 0.75139 "myosin binding protein C, fast-type" Mybpc2
1455737_at 0.88470 RIKEN cDNA C030002B11 gene C030002B11Rik
1455738_at 1.13063 coiled-coil domain containing 55 Ccdc55
1455739_at 0.84951 hypothetical protein 6430704N06 6430704N06
1455740_at 1.43844 heterogeneous nuclear ribonucleoprotein A1 Hnrpa1
1455741_a_at 0.95895 endothelin converting enzyme 1 Ece1
1455742_x_at 0.98487 mortality factor 4 like 1 /// RIKEN cDNA A730098P11 gene /// similar to
mortality factor 4 like 1 /// similar to mortality factor 4 like 1 (predicted) Morf411 /// A730098P11Rik ///
LOC627352 /// LOC635663
1455743_at 1.03670 olfactomedin-like 2A Olfml2a
1455744_at 1.26825 G protein-coupled receptor 178 Gpr178
1455745_at 0.92617 "ceroid-lipofuscinosis, neuronal 8" Cln8
1455746_at 0.86562 kinesin family member 13A Kif13a
1455747_at 1.00770 gamma-glutamyltransferase-like activity 1 Ggta1
1455748_at 0.88478 G protein-coupled receptor 178 Gpr178
1455749_x_at 1.08255 "NADH dehydrogenase (ubiquinone) 1 alpha subcomplex, 7 (B14.5a)"
Ndufa7
1455750_at 0.86733 cDNA sequence BC053994 BC053994
1455751_at 0.93700 cullin associated and neddylation disassociated 1 Cand1
1455752_a_at 0.99880 transmembrane protein 134 Tmem134
1455753_at 1.07331 RIKEN cDNA C630035N08 gene C630035N08Rik
1455754_at 0.84919 LIM domain only 3 Lmo3
1455755_at 1.04305 "gene model 88, (NCBI)" Gm88
1455756_at 0.89409 RIKEN cDNA F730014I05 gene F730014I05Rik
1455757_at 1.00853 "DNA segment, Chr 3, ERATO Doi 254, expressed" D3Ertd254e
1455758_at 0.99947 "protein kinase C, gamma" Prkcc
1455759_a_at 1.06500 ankyrin repeat and KH domain containing 1 Ankhd1
1455760_at 1.17448 "solute carrier family 9 (sodium/hydrogen exchanger), member 5" Slc9a5
1455761_at 1.06773 --- ---
1455762_at 0.97540 RIKEN cDNA C330002I19 gene C330002I19Rik
1455763_at 0.86850 ring finger protein 41 Rnf41
1455764_at 1.02181 RIKEN cDNA B930007L02 gene B930007L02Rik
1455765_a_at 0.96429 "ATP-binding cassette, sub-family C (CFTR/MRP), member 8" Abcc8
1455766_at 0.87398 "gamma-aminobutyric acid (GABA-A) receptor, subunit alpha 1" Gabra1
1455767_x_at 1.01224 ribosomal protein L21 Rpl21
1455768_at 0.92488 Niemann Pick type C2 Npc2
1455769_at 1.02672 "DCN1, defective in cullin neddylation 1, domain containing 5 (S. cerevisiae)"
Dcun1d5
1455770_at 0.99737 Transcribed locus ---
1455771_at 0.98190 benzodiazapine receptor associated protein 1 Bzrap1
1455772_at 0.95183 progesterone receptor Pgr
1455773_at 1.54736 Phosphatidylinositol binding clathrin assembly protein Picalm
1455774_at 0.94604 Jun proto-oncogene related gene d1 Jund1

1455775_at 0.95233 RIKEN cDNA 4932415D10 gene 4932415D10Rik
1455776_x_at 0.97279 bolA-like 2 (E. coli) Bola2
1455777_x_at 0.91033 hydroxysteroid (17-beta) dehydrogenase 4 Hsd17b4
1455778_at 0.83792 zinc finger protein 192 Zfp192
1455779_at 1.05916 Transcribed locus ---
1455780_at 0.90013 "tubulin tyrosine ligase-like family, member 6" Ttl6
1455781_at 0.78822 cDNA sequence BC027231 BC027231
1455782_at 1.10393 sterile alpha motif domain containing 7 Samd7
1455783_at 0.93206 --- ---
1455784_at 0.99664 secretory blood group 1 Sec1
1455785_at 1.04842 Transcribed locus ---
1455786_at 0.86715 RIKEN cDNA 2610036F08 gene 2610036F08Rik
1455787_x_at 0.96420 multiple inositol polyphosphate histidine phosphatase 1 Minpp1
1455788_x_at 0.98673 "Polymerase (DNA-directed), delta interacting protein 3" Poldip3
1455789_x_at 0.99490 similar to heat shock protein 8 LOC621284
1455790_at 1.07304 E2F transcription factor 2 E2f2
1455791_at 0.98923 chromatin licensing and DNA replication factor 1 Cdt1
1455792_x_at 1.06279 necdin Ndn
1455793_at 0.82356 cDNA sequence BC035537 BC035537
1455794_at 1.47815 RIKEN cDNA D130058I21 gene D130058I21Rik
1455795_at 0.84279 squamous cell carcinoma antigen recognized by T cells 2 Sart2
1455796_x_at 0.85699 olfactomedin 1 Olfm1
1455797_x_at 0.97284 ligatin Lgtn
1455798_at 1.16374 galactokinase 2 Galk2
1455799_at 0.85785 RAR-related orphan receptor beta Rorb
1455800_x_at 1.05036 sorting and assembly machinery component 50 homolog (S. cerevisiae)
Samm50
1455801_x_at 1.04430 tubulin-specific chaperone d Tbcd
1455802_x_at 1.04141 anterior gradient 2 (Xenopus laevis) Agr2
1455803_at 1.02284 "solute carrier organic anion transporter family, member 4a1" Slco4a1
1455804_x_at 0.87370 3-oxoacid CoA transferase 1 Oxct1
1455805_x_at 1.00247 coiled-coil domain containing 22 Ccdc22
1455806_x_at 1.04002 "NADH dehydrogenase (ubiquinone) 1 alpha subcomplex, 12 /// similar to
13kDa differentiation-associated protein" Ndufa12 /// LOC669626
1455807_at 1.07656 "testis-specific protein, Y-encoded-like 5" Tspyl5
1455808_at 0.95611 RIKEN cDNA 4922502D21 gene 4922502D21Rik
1455809_x_at 0.99109 resistance to inhibitors of cholinesterase 8 homolog (C. elegans) Ric8
1455810_a_at 0.89100 RIKEN cDNA E130112L23 gene /// similar to quattro E130112L23Rik ///
LOC671456
1455811_at 0.98240 "eukaryotic translation initiation factor 3, subunit 6 interacting protein"
Eif3s6ip
1455812_x_at 0.77826 vasorin Vasn
1455813_at 0.97430 similar to oogenesis MGC91194

1455814_x_at 1.11228 DEAD (Asp-Glu-Ala-Asp) box polypeptide 39 Ddx39
1455815_a_at 0.87928 "tyrosine 3-monooxygenase/tryptophan 5-monooxygenase activation protein, beta polypeptide" Ywhab
1455816_a_at 0.96485 potassium channel tetramerisation domain containing 3 Kctd3
1455817_x_at 0.97653 similar to Zinc finger X-linked protein ZXDB LOC668166
1455818_at 1.02311 RIKEN cDNA 4930427A07 gene 4930427A07Rik
1455819_at 0.94482 ROD1 regulator of differentiation 1 (S. pombe) Rod1
1455820_x_at 0.98819 "scavenger receptor class B, member 1" Scarb1
1455821_x_at 1.03618 "complement component 1, q subcomponent binding protein" C1qbp
1455822_x_at 1.04470 surfeit gene 4 Surf4
1455823_at 0.91320 Bardet-Biedl syndrome 4 homolog (human) Bbs4
1455824_x_at 1.07583 "STT3, subunit of the oligosaccharyltransferase complex, homolog A (S. cerevisiae)" Stt3a
1455825_s_at 0.86820 ligand of numb-protein X 1 Lnx1
1455826_a_at 0.93001 beta-site APP cleaving enzyme 1 Bace1
1455827_at 1.02915 muscleblind-like 2 Mbnl2
1455828_at 0.91132 G protein-regulated inducer of neurite outgrowth 1 Gprin1
1455829_at 1.05788 ubiquitin specific peptidase 14 Usp14
1455830_s_at 1.03008 expressed sequence C87414 /// similar to PRAME family member 9 /// similar to PRAME family member 9 C87414 /// LOC664950 /// LOC665592
1455831_at 2.08452 "fusion, derived from t(12;16) malignant liposarcoma (human)" Fus
1455832_a_at 1.09826 uridine monophosphate synthetase Umps
1455833_at 1.01127 expressed sequence AU041783 AU041783
1455834_x_at 1.19828 "transforming, acidic coiled-coil containing protein 3" Tacc3
1455835_at 0.94885 Ribosomal protein S16 Rps16
1455836_at 1.00691 poly (A) polymerase alpha Papola
1455837_at 1.06146 tripartite motif-containing 36 Trim36
1455838_at 0.84329 schlafen like 1 Slfn1
1455839_at 1.01853 UDP-glucose ceramide glucosyltransferase-like 1 Ugcgl1
1455840_at 0.62781 Rap guanine nucleotide exchange factor (GEF) 5 Rapgef5
1455841_s_at 0.95799 glutamate-rich WD repeat containing 1 Grwd1
1455842_x_at 0.95118 ubiquitin specific peptidase 15 Usp15
1455843_at 0.79360 fucosyltransferase 4 Fut4
1455844_at 0.87506 Cdk5 and Abl enzyme substrate 1 Cables1
1455845_at 0.92941 cDNA sequence BC030477 BC030477
1455846_at 0.88467 aquaporin 11 Aqp11
1455847_at 0.97935 p21 (CDKN1A)-activated kinase 7 Pak7
1455848_at 1.06635 "transmembrane protease, serine 11a" Tmprss11a
1455849_at 0.98086 --- ---
1455850_at 0.97904 RIKEN cDNA 2310003H01 gene 2310003H01Rik
1455851_at 1.00487 bone morphogenetic protein 5 Bmp5
1455852_at 1.13869 RIKEN cDNA 4833432M17 gene 4833432M17Rik
1455853_x_at 0.82707 tetraspanin 31 Tspan31

1455854_a_at 1.03530 slingshot homolog 1 (Drosophila) Ssh1
1455855_x_at 0.96703 heterogeneous nuclear ribonucleoprotein A/B Hnrpab
1455856_at 0.95279 --- ---
1455857_a_at 0.95821 "RAB2B, member RAS oncogene family" Rab2b
1455858_x_at 1.03262 pro-opiomelanocortin-alpha Pomc1
1455859_at 1.69998 RIKEN cDNA A330021E22 gene A330021E22Rik
1455860_at 0.96868 "phosphatidylinositol glycan anchor biosynthesis, class H" Pigh
1455861_at 1.07638 erythrocyte protein band 4.1-like 5 Epb4.115
1455862_at 1.05548 RIKEN cDNA 9630054F20 gene 9630054F20Rik
1455863_at 0.98134 spermatogenesis associated 5-like 1 Spata51
1455864_at 1.01536 tripartite motif-containing 75 Trim75
1455865_at 0.94695 insulinoma-associated 1 Insm1
1455866_x_at 1.11748 thioredoxin-like 1 Txnl1
1455867_at 0.73941 SRY-box containing gene 4 Sox4
1455868_a_at 0.87431 "tubulin, gamma complex associated protein 2" Tubgcp2
1455869_at 1.01068 "Calcium/calmodulin-dependent protein kinase II, beta" Camk2b
1455870_at 0.99505 A kinase (PRKA) anchor protein 2 Akap2
1455871_s_at 0.80785 Tax1 (human T-cell leukemia virus type I) binding protein 3 /// ribosomal
protein L13 Tax1bp3 /// Rp113
1455872_at 0.74457 cDNA sequence BC065085 BC065085
1455873_a_at 0.99725 vacuolar protein sorting 18 (yeast) Vps18
1455874_at 0.91031 RIKEN cDNA 1810059G22 gene 1810059G22Rik
1455875_x_at 0.88956 transmembrane 9 superfamily member 2 Tm9sf2
1455876_at 1.21074 "solute carrier family 4, sodium bicarbonate cotransporter, member 7" Slc4a7
1455877_a_at 0.99721 nanos homolog 1 (Drosophila) Nanos1
1455878_at 1.18829 RIKEN cDNA 2700023E23 gene 2700023E23Rik
1455879_at 0.87014 GCN1 general control of amino-acid synthesis 1-like 1 (yeast) Gcn111
1455880_s_at 1.00562 "beclin 1 (coiled-coil, myosin-like BCL2-interacting protein)" Becn1
1455881_at 0.87482 immediate early response 5-like Ier51
1455882_x_at 0.92414 von Willebrand factor C domain containing 2 Vwc2
1455883_a_at 0.97122 leucine rich repeat transmembrane neuronal 1 Lrrtm1
1455884_at 0.88478 dipeptidylpeptidase 9 Dpp9
1455885_at 0.84917 RIKEN cDNA 6530401C20 gene 6530401C20Rik
1455886_at 1.08597 Casitas B-lineage lymphoma Cbl
1455887_at 0.95246 "asparagine-linked glycosylation 8 homolog (yeast, alpha-1,3-
glucosyltransferase)" Alg8
1455888_at 0.80566 leucine rich repeat neuronal 6C Lrrn6c
1455889_at 1.07690 male sterility domain containing 1 Mlstd1
1455890_x_at 0.94886 small nuclear ribonucleoprotein N /// similar to small nuclear
ribonucleoparticle-associated protein Snrpn /// LOC545062
1455891_at 0.91200 RIKEN cDNA 4933425K02 gene 4933425K02Rik
1455892_x_at 0.97721 similar to melanoma antigen /// similar to melanoma antigen /// similar to
melanoma antigen /// similar to melanoma antigen /// similar to melanoma antigen /// similar to

melanoma antigen /// similar to melanoma antigen LOC622147 /// LOC668227 /// LOC668269 ///
 LOC669176 /// LOC669658 /// LOC669821 /// LOC676636
 1455893_at 1.06414 R-spondin 2 homolog (*Xenopus laevis*) Rspo2
 1455894_at 0.92968 peptidyl-tRNA hydrolase 2 Ptrh2
 1455895_x_at 1.08123 RIKEN cDNA 1110012M11 gene 1110012M11Rik
 1455896_a_at 0.97982 "potassium channel, subfamily K, member 1" Kcnk1
 1455897_x_at 0.99372 high mobility group nucleosomal binding domain 1 Hmgn1
 1455898_x_at 0.86554 "solute carrier family 2 (facilitated glucose transporter), member 3" Slc2a3
 1455899_x_at 0.96412 suppressor of cytokine signaling 3 Socs3
 1455900_x_at 0.94918 "transglutaminase 2, C polypeptide" Tgm2
 1455901_at 0.84902 choline phosphotransferase 1 Chpt1
 1455902_x_at 0.99213 "Ras homolog gene family, member f" Rhof
 1455903_at 0.90417 "TBC1 domain family, member 4" Tbc1d4
 1455904_at 1.13409 growth arrest specific 5 Gas5
 1455905_at 0.94934 RIKEN cDNA 2610507B11 gene 2610507B11Rik
 1455906_at 0.78998 RIKEN cDNA 6030446N20 gene 6030446N20Rik
 1455907_x_at 0.98073 paired-like homeobox 2b Phox2b
 1455908_a_at 0.97019 serine carboxypeptidase 1 Scpep1
 1455909_at 1.14309 RIKEN cDNA 9530002K18 gene 9530002K18Rik
 1455910_at 0.97810 kelch domain containing 3 Klhdc3
 1455911_x_at 0.91768 "NADH dehydrogenase (ubiquinone) 1 beta subcomplex, 11" Ndufb11
 1455912_x_at 0.86989 Unc-45 homolog A (*C. elegans*) Unc45a
 1455913_x_at 0.80215 transthyretin Ttr
 1455914_at 0.89401 expressed sequence AI987944 AI987944
 1455915_at 0.82637 UDP-N-acetyl-alpha-D-galactosamine:polypeptide N-
 acetylgalactosaminyltransferase 4 Galnt4
 1455916_at 1.10014 expressed sequence AI931714 AI931714
 1455917_at 0.94258 "neurotrophic tyrosine kinase, receptor, type 3" Ntrk3
 1455918_at 1.07105 "adrenergic receptor, beta 3" Adrb3
 1455919_at 1.00308 "Gene model 879, (NCBI)" Gm879
 1455920_x_at 1.13892 lysozyme-like 6 Lyzl6
 1455921_at 0.95538 Transcribed locus ---
 1455922_at 0.98225 RAB3 GTPase activating protein subunit 1 Rab3gap1
 1455923_at 0.90082 potassium channel tetramerisation domain containing 8 Kctd8
 1455924_at 0.87381 "RAB6B, member RAS oncogene family" Rab6b
 1455925_at 0.99535 PR domain containing 8 Prdm8
 1455926_at 0.94114 "LSM6 homolog, U6 small nuclear RNA associated (*S. cerevisiae*)" Lsm6
 1455927_x_at 1.01130 --- ---
 1455928_x_at 0.84063 "leucine-zipper-like transcriptional regulator, 1" Lztr1
 1455929_x_at 0.93260 "protein phosphatase 2 (formerly 2A), regulatory subunit A (PR 65), alpha
 isoform" Ppp2r1a
 1455930_at 0.76991 hypothetical protein LOC673503 LOC673503
 1455931_at 1.05289 "cholinergic receptor, nicotinic, alpha polypeptide 3" Chrna3

1455932_at	0.95177	Metadherin	Mtdh	
1455933_at	1.26851	transformer 2 alpha homolog (Drosophila)	Tra2a	
1455934_at	0.92420	"NADH dehydrogenase (ubiquinone) 1 beta subcomplex, 9"	Ndufb9	
1455935_at	0.93843	RIKEN cDNA 2410131K14 gene	2410131K14Rik	
1455936_a_at	0.92929	RNA binding protein gene with multiple splicing	Rbpms	
1455937_at	1.03451	leucine rich repeat transmembrane neuronal 4	Lrrtm4	
1455938_x_at	0.98275	RAD21 homolog (S. pombe)	Rad21	
1455939_x_at	0.93780	signal recognition particle 14	Srp14	
1455940_x_at	0.88068	WD repeat domain 6	Wdr6	
1455941_s_at	0.95059	mitogen activated protein kinase kinase 5	Map2k5	
1455942_at	1.10462	F-box and leucine-rich repeat protein 11	Fbxl11	
1455943_at	1.01393	zinc finger protein 451	Zfp451	
1455944_at	0.93130	zinc finger protein 516	Zfp516	
1455945_at	0.94329	RIKEN cDNA A530094I17 gene	A530094I17Rik	
1455946_x_at	1.03560	"thymosin, beta 10"	Tmsb10	
1455947_at	0.94802	cDNA sequence AK162044	AK162044	
1455948_x_at	1.03558	matrilin 3	Matn3	
1455949_at	0.89920	"ATPase, Na ⁺ /K ⁺ transporting, beta 1 polypeptide"	Atp1b1	
1455950_x_at	1.01124	ribosomal protein L35	Rpl35	
1455951_at	1.11440	methionine-tRNA synthetase	Mars	
1455952_at	0.93804	ADP-ribosylhydrolase like 1	Adprh1	
1455953_x_at	1.10029	phosphoseryl-tRNA kinase	Pstk	
1455954_x_at	0.87599	GPI anchor attachment protein 1	Gpaal	
1455955_s_at	0.96724	sorting nexin 17	Snx17	
1455956_x_at	1.03728	cyclin D2	Ccnd2	
1455957_x_at	0.94791	CEA-related cell adhesion molecule 11	Ceacam11	
1455958_s_at	1.13629	PTC7 protein phosphatase homolog (S. cerevisiae)	Pptc7	
1455959_s_at	0.89723	---	---	
1455960_at	0.88712	multiple EGF-like-domains 9	Megf9	
1455961_at	0.80409	---	---	
1455962_at	1.06054	hedgehog acyltransferase	Hhat	
1455963_at	0.91673	RIKEN cDNA 6332401O19 gene	6332401O19Rik	
1455964_at	1.07635	"Cdc2-related kinase, arginine/serine-rich"	Crkrs	
1455965_at	1.08417	---	---	
1455966_s_at	0.91970	nudix (nucleoside diphosphate linked moiety X)-type motif 21	Nudt21	
1455967_at	1.00424	sorbin and SH3 domain containing 1	Sorbs1	
1455968_x_at	0.96392	transmembrane emp24 domain trafficking protein 2	Tmed2	
1455969_at	1.05877	nudix (nucleoside diphosphate linked moiety X)-type motif 15	Nudt15	
1455970_at	0.79581	"Phosphodiesterase 5A, cGMP-specific"	Pde5a	
1455971_at	0.92926	RIKEN cDNA 2810032E02 gene	2810032E02Rik	
1455972_x_at	0.87639	"L-3-hydroxyacyl-Coenzyme A dehydrogenase, short chain"	Hadhsc	
1455973_at	1.02762	hypothetical protein LOC626870	RP23-219P13.3	
1455974_at	1.01549	RIKEN cDNA 1110049F12 gene	1110049F12Rik	

1455975_x_at 1.01564 zinc finger protein 313 Zfp313
1455976_x_at 1.16964 diazepam binding inhibitor Dbi
1455977_x_at 0.94243 kallikrein 1 /// kallikrein 1-related peptidase b5 Klk1 /// Klk1b5
1455978_a_at 1.02183 matrilin 2 Matn2
1455979_at 0.92891 AT rich interactive domain 1B (Swi1 like) Arid1b
1455980_a_at 1.54129 growth arrest-specific 2 like 3 Gas2l3
1455981_at 1.03152 ribosomal protein S6 /// similar to 40S ribosomal protein S6 /// similar to 40S
ribosomal protein S6 /// similar to 40S ribosomal protein S6 /// similar to 40S ribosomal protein S6 ///
similar to 40S ribosomal protein S6 /// similar to 40S ribosomal protein S6 /// similar to 40S ribosomal
protein S6 /// similar to 40S ribosomal protein S6 /// similar to 40S ribosomal protein S6 /// similar to
40S ribosomal protein S6 /// similar to 40S ribosomal protein S6 /// similar to 40S ribosomal protein
S6 /// similar to 40S ribosomal protein S6 /// similar to 40S ribosomal protein S6 /// similar to 40S
ribosomal protein S6 Rps6 /// LOC214738 /// LOC384383 /// LOC545640 /// LOC626259 ///
LOC629575 /// LOC638497 /// LOC639593 /// LOC639750 /// LOC666577 /// LOC667739 ///
LOC667922 /// LOC671306 /// LOC671432 /// LOC672620 /// LOC675985
1455982_at 0.98159 jumonji domain containing 4 Jmjd4
1455983_at 0.99546 cell division cycle associated 2 Cdca2
1455984_at 0.98708 retinoic acid induced 17 Rai17
1455985_x_at 1.02570 serine hydroxymethyl transferase 2 (mitochondrial) Shmt2
1455986_at 0.91178 "zinc finger, DHHC domain containing 17" Zdhhc17
1455987_at 1.22334 Sec61 alpha 1 subunit (S. cerevisiae) Sec61a1
1455988_a_at 1.00525 chaperonin subunit 6a (zeta) Cct6a
1455989_at 1.21042 gap junction membrane channel protein alpha 12 Gja12
1455990_at 1.06651 kinesin family member 23 Kif23
1455991_at 1.10351 cysteine conjugate-beta lyase 2 Ccbl2
1455992_at 0.83553 vestigial like 4 (Drosophila) Vgll4
1455993_at 0.77371 --- ---
1455994_x_at 1.13111 "elongation of very long chain fatty acids (FEN1/Elo2, SUR4/Elo3, yeast)-like
1" Elovl1
1455995_at 0.90150 "DNA segment, Chr 10, Brigham & Women's Genetics 1379 expressed"
D10Bwg1379e
1455996_x_at 1.01106 proline-rich acidic protein 1 Prap1
1455997_a_at 0.89810 ubiquinol-cytochrome c reductase binding protein Uqcrb
1455998_at 1.17462 similar to Zinc finger BED domain containing protein 4 /// similar to Zinc finger
BED domain containing protein 4 /// similar to Zinc finger BED domain containing protein 4
LOC667118 /// LOC669797 /// LOC675760
1455999_at 0.98148 RIKEN cDNA C330003B14 gene C330003B14Rik
1456000_at 1.02769 DEAD (Asp-Glu-Ala-Asp) box polypeptide 18 Ddx18
1456001_at 0.93197 similar to late cornified envelope protein LOC545548
1456002_at 1.00735 "xeroderma pigmentosum, complementation group A" Xpa
1456003_a_at 0.92136 "solute carrier family 1 (glutamate/neutral amino acid transporter), member 4"
Slc1a4
1456004_x_at 1.00616 islet amyloid polypeptide Iapp

1456005_a_at 0.80579 BCL2-like 11 (apoptosis facilitator) Bcl2l11
1456006_at 0.87188 BCL2-like 11 (apoptosis facilitator) Bcl2l11
1456007_at 0.95216 ADP-ribosylation factor 3 Arf3
1456008_at 0.79718 "leucine rich repeat containing 8 family, member E" Lrrc8e
1456009_x_at 1.00202 topoisomerase (DNA) III beta Top3b
1456010_x_at 0.96659 hairy and enhancer of split 5 (Drosophila) Hes5
1456011_x_at 0.90933 acetyl-Coenzyme A acyltransferase 1A Acaa1a
1456012_x_at 0.99404 ribonuclease T2 Rnaset2
1456013_x_at 0.83745 "solute carrier family 35, member A4" Slc35a4
1456014_s_at 1.05475 cDNA sequence BC032204 BC032204
1456015_x_at 0.92660 NADH dehydrogenase (ubiquinone) flavoprotein 1 Ndufv1
1456016_x_at 0.87082 CDNA sequence BC005624 BC005624
1456017_x_at 1.02587 similar to oocyte specific homeobox 2 LOC666466
1456018_at 1.00540 expressed sequence AL022779 AL022779
1456019_at 0.93490 "CWF19-like 2, cell cycle control (S. pombe)" Cwf19l2
1456020_at 0.99318 SH3 domain and tetratricopeptide repeats 2 Sh3tc2
1456021_at 0.95956 activating transcription factor 6 Atf6
1456022_at 0.73438 homeodomain interacting protein kinase 2 Hipk2
1456023_at 0.93863 RIKEN cDNA 2210010B09 gene 2210010B09Rik
1456024_at 0.90632 general transcription factor III C 1 Gtf3c1
1456025_at 0.82695 doublecortin and CaM kinase-like 2 Dcamk12
1456026_at 0.86551 RIKEN cDNA 8030451K01 gene 8030451K01Rik
1456027_at 1.01867 RNA binding motif protein 41 Rbm41
1456028_x_at 1.05070 Myristoylated alanine rich protein kinase C substrate Marcks
1456029_a_at 1.04673 tetratricopeptide repeat domain 7B Ttc7b
1456030_at 0.97330 Kruppel-like factor 13 Klf13
1456031_at 0.86526 "Tubulin, beta 2c" Tubb2c
1456032_x_at 1.02308 "H2A histone family, member Z /// similar to H2A histone family, member Z /// similar to H2A histone family, member Z" H2afz /// LOC666634 /// LOC675899
1456033_at 0.87596 T-box 4 Tbx4
1456034_at 1.03030 tetratricopeptide repeat domain 18 Ttc18
1456035_at 0.99867 nuclear RNA export factor 3 Nxf3
1456036_x_at 1.21144 glutathione S-transferase omega 1 Gsto1
1456037_x_at 1.01273 prolactin regulatory element binding Preb
1456038_at 1.08321 F-box and leucine-rich repeat protein 4 Fbxl4
1456039_at 0.91909 sorting nexin 14 Snx14
1456040_at 0.98059 "splicing factor 3b, subunit 2" Sf3b2
1456041_at 0.94716 sorting nexin 16 Snx16
1456042_s_at 1.01739 "Crm, cramped-like (Drosophila)" Cramp11
1456043_at 0.90537 ubiquitin specific peptidase 22 Usp22
1456044_at 1.09880 "neural precursor cell expressed, developmentally down-regulated gene 4-like" Nedd4l
1456045_at 0.82612 RIKEN cDNA 1700106N22 gene 1700106N22Rik

1456046_at 0.93662 CD93 antigen Cd93
1456047_at 1.01817 "phospholipase A2, group IVB (cytosolic)" Pla2g4b
1456048_at 0.96735 cytoplasmic polyadenylation element binding protein 3 Cpeb3
1456049_at 1.20783 v-ral simian leukemia viral oncogene homolog A (ras related) Rala
1456050_at 1.24387 expressed sequence C80998 C80998
1456051_at 1.05946 dopamine receptor D1A Drd1a
1456052_at 0.93430 cyclin-dependent kinase-like 3 Cdkl3
1456053_at 0.98566 Transcribed locus ---
1456054_a_at 1.02469 pumilio 1 (Drosophila) Pum1
1456055_x_at 0.97680 "polymerase (DNA directed), delta 1, catalytic subunit" Pold1
1456056_a_at 1.05138 "DNA segment, Chr 6, Wayne State University 116, expressed" D6Wsu116e
1456057_x_at 0.94097 transmembrane protein 109 Tmem109
1456058_at 1.04405 RNA binding motif protein 27 Rbm27
1456059_at 1.08944 "proteasome (prosome, macropain) 26S subunit, non-ATPase, 11" Psmd11
1456060_at 0.72710 avian musculoaponeurotic fibrosarcoma (v-maf) AS42 oncogene homolog
Maf
1456061_at 1.05331 "GTPase, IMAP family member 8" Gimap8
1456062_at 1.04865 natriuretic peptide precursor type A Nppa
1456063_at 0.79481 open reading frame 34 ORF34
1456064_at 1.07429 expressed sequence AI504432 AI504432
1456065_at 0.92869 "ubiquitin associated and SH3 domain containing, A" Ubash3a
1456066_a_at 1.29038 RNA polymerase 1-4 Rpo1-4
1456067_at 1.03975 GLI-Kruppel family member GLI3 Gli3
1456068_at 1.04124 neurofascin Nfasc
1456069_at 1.13218 dystrobrevin alpha Dtna
1456070_at 1.03077 RIKEN cDNA 5430405N12 gene 5430405N12Rik
1456071_a_at 0.95907 "cytochrome c, somatic /// similar to Cytochrome c, somatic /// similar to
Cytochrome c, somatic" Cyps /// LOC670717 /// LOC672195
1456072_at 1.16365 "protein phosphatase 1, regulatory (inhibitor) subunit 9A" Ppp1r9a
1456073_s_at 1.14435 pannexin 3 Panx3
1456074_at 1.05239 orphan short chain dehydrogenase/reductase Sdro
1456075_at 1.00163 protein kinase D2 Prkd2
1456076_at 0.98335 defensin beta 19 Defb19
1456077_x_at 1.11097 --- ---
1456078_x_at 0.91709 "tubulin, beta 2c" Tubb2c
1456079_x_at 1.20850 apurinic/aprimidinic endonuclease 1 Apex1
1456080_a_at 0.83807 serine incorporator 3 Serinc3
1456081_a_at 1.31831 acetoacetyl-CoA synthetase Aacs
1456082_x_at 0.94208 chaperonin subunit 4 (delta) Cct4
1456083_x_at 0.96478 "eukaryotic translation initiation factor 3, subunit 8" Eif3s8
1456084_x_at 0.69538 fibromodulin Fmod
1456085_x_at 0.88253 CD151 antigen Cd151
1456086_x_at 0.88186 similar to polyglutamine binding protein 1 LOC631302

1456087_at 0.82991 nuclear factor I/A Nfia
1456088_at 1.13807 baculoviral IAP repeat-containing 4 Birc4
1456089_at 0.91349 tripartite motif protein 23 Trim23
1456090_at 1.04661 "pyruvate dehydrogenase complex, component X" Pdhx
1456091_at 1.08691 SEC22 vesicle trafficking protein-like C (S. cerevisiae) Sec22c
1456092_at 1.07458 potassium channel tetramerisation domain containing 7 Kctd7
1456093_at 0.87008 zinc finger protein 536 Zfp536
1456094_at 0.95475 ubiquitin specific peptidase 36 Usp36
1456095_at 1.04621 tyrosinase Tyr
1456096_at 0.63718 RIKEN cDNA 6430573F11 gene 6430573F11Rik
1456097_a_at 0.83491 integrin beta 3 binding protein (beta3-endonexin) Itgb3bp
1456098_a_at 0.83674 "engulfment and cell motility 2, ced-12 homolog (C. elegans)" Elmo2
1456099_at 1.06727 RIKEN cDNA D930017J03 gene D930017J03Rik
1456100_at 1.04383 cDNA sequence BC089491 BC089491
1456101_at 1.14718 hypothetical protein LOC640072 /// hypothetical protein LOC677194
LOC640072 /// LOC677194
1456102_a_at 1.04810 cullin 5 Cul5
1456103_at 0.97966 promyelocytic leukemia Pml
1456104_at 0.97798 "proteasome (prosome, macropain) 26S subunit, non-ATPase, 11" Psmd11
1456105_at 0.99391 membrane associated DNA binding protein Mnab
1456106_x_at 0.87991 serologically defined colon cancer antigen 3 Sdccag3
1456107_x_at 1.01903 elongation factor Tu GTP binding domain containing 2 Eftud2
1456108_x_at 0.94843 zinc finger protein 179 Zfp179
1456109_a_at 1.02821 mitochondrial ribosomal protein S15 Mrps15
1456110_at 0.90235 ankyrin repeat domain 11 Ankrd11
1456111_at 0.86984 RIKEN cDNA D930028F11 gene D930028F11Rik
1456112_at 1.01591 translocated promoter region Tpr
1456113_at 1.08195 ornithine aminotransferase-like 1 Oat11
1456114_at 1.00591 CDP-diacylglycerol synthase 1 Cds1
1456115_at 0.82686 PHD finger protein 3 Phf3
1456116_at 0.93597 "catenin (cadherin associated protein), delta 2" Ctnnd2
1456117_at 1.13597 RIKEN cDNA 2600005C20 gene 2600005C20Rik
1456118_at 1.05773 methyltransferase like 2 Mettl2
1456119_at 0.95569 "glutamate receptor, metabotropic 5" Grm5
1456120_at 0.99084 RIKEN cDNA 3110001I20 gene 3110001I20Rik
1456121_at 0.99405 leucine-rich repeats and IQ motif containing 2 Lrriq2
1456122_at 1.43417 --- ---
1456123_at 0.92078 expressed sequence AI481121 AI481121
1456124_x_at 0.91255 seminal vesicle secretion 5 Svs5
1456125_a_at 0.93815 "dynein light chain LC8-type 1 /// similar to dynein, cytoplasmic, light peptide /// similar to dynein, cytoplasmic, light peptide /// similar to dynein, cytoplasmic, light peptide" Dynl11 /// LOC627788 /// LOC637840 /// LOC672375
1456126_at 0.95813 mucosa associated lymphoid tissue lymphoma translocation gene 1 Malt1

1456127_at 0.89541 RIKEN cDNA 9630008K15 gene 9630008K15Rik
1456128_at 0.88340 "ATP synthase, H⁺ transporting, mitochondrial F₀ complex, subunit c (subunit 9), isoform 2" Atp5g2
1456129_at 1.03687 RIKEN cDNA E030013G06 gene E030013G06Rik
1456130_at 0.79934 hypothetical LOC553091 LOC553091
1456131_x_at 0.93978 dystroglycan 1 Dag1
1456132_x_at 0.98478 "TSC22 domain family, member 1" Tsc22d1
1456133_x_at 0.69660 integrin beta 5 Itgb5
1456134_x_at 0.95272 Yip1 interacting factor homolog A (S. cerevisiae) Yif1a
1456135_s_at 0.93170 paxillin Pxn
1456136_at 1.65013 Kinesin family member C1 Kifc1
1456137_at 0.94408 neurexin III Nrnx3
1456138_at 1.05437 LY6/PLAUR domain containing 6 Lypd6
1456139_at 2.04741 antisense Igf2r RNA Air
1456140_at 0.97760 similar to zinc finger protein of the cerebellum 5 LOC624168
1456141_x_at 0.94165 "BRCA1/BRCA2-containing complex, subunit 3" Brcc3
1456142_x_at 0.97527 mortality factor 4 like 1 /// similar to mortality factor 4 like 1 isoform b /// RIKEN cDNA A730098P11 gene /// similar to mortality factor 4 like 1 /// similar to mortality factor 4 like 1 (predicted) Morf4l1 /// LOC433955 /// A730098P11Rik /// LOC627352 /// LOC635663
1456143_at 1.11456 protein kinase C binding protein 1 Prkcbp1
1456144_at 1.07308 region containing RIKEN cDNA 9630020C08 gene; neuron navigator 3 LOC676640
1456145_at 1.13505 "deleted in lymphocytic leukemia, 2" Dleu2
1456146_at 0.92866 oxysterol binding protein-like 10 Osbpl10
1456147_at 0.93099 "ST8 alpha-N-acetyl-neuraminidase alpha-2,8-sialyltransferase 6" St8sia6
1456148_a_at 0.96787 RIKEN cDNA 1110014K08 gene 1110014K08Rik
1456149_at 1.06863 "DNA segment, Chr 11, Brigham & Women's Genetics 0517 expressed" D11Bwg0517e
1456150_at 1.09565 RIKEN cDNA A630082K20 gene A630082K20Rik
1456151_at 1.02580 zinc finger protein 358 Zfp358
1456152_at 0.96963 progesterin and adipoQ receptor family member V Paqr5
1456153_at 1.07720 slingshot homolog 2 (Drosophila) Ssh2
1456154_at 1.02896 Zinc finger protein 444 Zfp444
1456155_x_at 1.00564 "fucosidase, alpha-L- 1, tissue" Fuca1
1456156_at 1.08730 leptin receptor Lepr
1456157_at 1.02129 "DNA segment, Chr 14, ERATO Doi 209, expressed" D14Ertd209e
1456158_at 1.05866 --- ---
1456159_at 3.36743 SET domain containing 5 Setd5
1456160_at 1.12200 Nucleoredoxin Nxn
1456161_at 0.89562 RIKEN cDNA 0610040B10 gene 0610040B10Rik
1456162_x_at 1.04262 adducin 3 (gamma) Add3
1456163_at 1.14208 RIKEN cDNA 2700049P18 gene 2700049P18Rik
1456164_at 0.96846 expressed sequence AW011738 AW011738

1456165_at	1.04516	RIKEN cDNA 4921521F21 gene	4921521F21Rik
1456166_at	1.10824	---	---
1456167_at	1.03189	similar to PHD finger protein 19	LOC241051
1456168_at	1.08898	Cyclin J-like Ccnjl	
1456169_at	1.00568	similar to KAT protein	LOC226654
1456170_x_at	0.95388	calreticulin Calr	
1456171_at	0.96419	ring finger protein 190	Rnf190
1456172_at	1.08202	Thioredoxin domain containing 5	Txndc5
1456173_at	0.83588	"Interleukin 10 receptor, alpha"	Il10ra
1456174_x_at	1.11678	N-myc downstream regulated gene 1	Ndr1
1456175_a_at	0.93771	"coatamer protein complex, subunit beta 2 (beta prime)"	Copb2
1456176_x_at	0.82952	"DNA segment, Chr 11, ERATO Doi 333, expressed"	D11Ert333e
1456177_x_at	0.87850	zinc finger protein 706	Zfp706
1456178_at	0.94265	"BMP and activin membrane-bound inhibitor, pseudogene (Xenopus laevis)"	
Bambi-ps1			
1456179_at	0.93427	---	---
1456180_at	0.88751	similar to RNA binding motif protein 24 /// similar to RNA binding motif protein 24 /// similar to RNA binding motif protein 24	LOC380843 /// LOC638266 /// LOC666794
1456181_at	1.04849	RIKEN cDNA 9530020G05 gene	9530020G05Rik
1456182_x_at	0.91043	melanoma antigen	Mela
1456183_at	1.04892	oogenesis 4	Oog4
1456184_at	0.89575	transmembrane protein 63c	Tmem63c
1456185_at	0.98388	"NACHT, LRR and PYD containing protein 9a"	Nalp9a
1456186_at	0.96329	RIKEN cDNA 8030443D09 gene	8030443D09Rik
1456187_at	0.99748	"solute carrier family 7 (cationic amino acid transporter, y+ system), member 14"	Slc7a14
1456188_at	1.00662	"seminal vesicle protein, secretion 2"	Svs2
1456189_x_at	0.89300	latent transforming growth factor beta binding protein 3	Ltbp3
1456190_a_at	1.04962	Acyl-CoA synthetase medium-chain family member 2	Acsm2
1456191_x_at	0.98905	protein kinase C and casein kinase substrate in neurons 3	Pacsin3
1456192_x_at	1.11364	cysteine-rich perinuclear theca 3	Cypt3
1456193_x_at	1.03780	"glutathione peroxidase 4 /// similar to Phospholipid hydroperoxide glutathione peroxidase, mitochondrial precursor (PHGPx) (GPX-4)"	Gpx4 /// LOC671271
1456194_a_at	1.07917	"Parkinson disease (autosomal recessive, early onset) 7"	Park7
1456195_x_at	0.66059	integrin beta 5	Itgb5
1456196_x_at	1.02124	FK506 binding protein 1a	Fkbp1a
1456197_x_at	0.95637	Transcribed locus	---
1456198_at	0.94137	RIKEN cDNA 1810007D17 gene	1810007D17Rik
1456199_x_at	0.87341	RAN binding protein 9	Ranbp9
1456200_at	1.30008	inositol polyphosphate multikinase	Ipmk
1456201_at	1.23712	RIKEN cDNA 4632427E13 gene	4632427E13Rik
1456202_at	0.98256	cDNA sequence BC094219	BC094219
1456203_at	0.99358	RIKEN cDNA 1110020A10 gene	1110020A10Rik

1456204_at 0.97496 RIKEN cDNA 2010107H07 gene 2010107H07Rik
1456205_x_at 1.01049 tubulin cofactor a Tbca
1456206_at 0.91208 "AlkB, alkylation repair homolog 3 (E. coli)" Alkbh3
1456207_at 1.08048 mature T-cell proliferation 1 Mtcp1
1456208_at 1.05946 RIKEN cDNA A530057A03 gene A530057A03Rik
1456209_x_at 1.03260 --- ---
1456210_at 1.38571 RIKEN cDNA 5430407P10 gene 5430407P10Rik
1456211_at 0.99468 "NACHT, leucine rich repeat and PYD containing 10" Nalp10
1456212_x_at 0.96743 suppressor of cytokine signaling 3 Socs3
1456213_x_at 0.98824 glutaminyl-tRNA synthetase Qars
1456214_at 0.90057 protocadherin 7 Pcdh7
1456215_at 0.93295 "transcription factor A, mitochondrial" Tfam
1456216_at 1.66426 "Casein kinase 1, alpha 1" Csnk1a1
1456217_at 0.82756 "DNA segment, Chr 7, Wayne State University 128, expressed" D7Wsu128e
1456218_at 0.87597 sorting nexin 22 Snx22
1456219_at 0.99290 similar to zinc finger protein of the cerebellum 5 LOC624168
1456220_at 0.76036 F-box and leucine-rich repeat protein 7 Fbxl7
1456221_at 0.92675 Cadherin 8 Cdh8
1456222_at 0.98752 BTB (POZ) domain containing 1 Btbd1
1456223_at 1.75879 F-box and leucine-rich repeat protein 5 Fbxl5
1456224_x_at 0.93771 cancer antigen 1 Cage1
1456225_x_at 1.05817 tribbles homolog 3 (Drosophila) Trib3
1456226_x_at 0.86678 "discoidin domain receptor family, member 1" Ddr1
1456227_x_at 0.98975 retinoblastoma binding protein 7 Rbbp7
1456228_x_at 0.91807 myelin basic protein Mbp
1456229_at 0.98064 homeo box B3 Hoxb3
1456230_at 0.97766 RIKEN cDNA 4933422H20 gene 4933422H20Rik
1456231_at 0.98062 "phospholipase A2, group III" Pla2g3
1456232_at 1.25733 glucocorticoid modulatory element binding protein 1 Gmeb1
1456233_at 0.95973 Transcribed locus ---
1456234_at 1.05743 "LIM-domain containing, protein kinase" Limk1
1456235_at 0.89167 expressed sequence BB165335 BB165335
1456236_s_at 1.08917 COMM domain containing 10 Commd10
1456237_x_at 0.99454 CDNA sequence BC053401 BC053401
1456238_at 1.02770 similar to Zinc finger protein 133 LOC668917
1456239_at 0.99398 fibroblast growth factor 17 Fgf17
1456240_x_at 0.96074 cell division cycle associated 4 Cdca4
1456241_a_at 0.86961 RIKEN cDNA 1810073N04 gene 1810073N04Rik
1456242_at 1.02984 embryonic stem cell- and germ cell-specific protein Esgp
1456243_x_at 0.89201 myeloid cell leukemia sequence 1 Mcl1
1456244_x_at 1.01170 thioredoxin-like 2 Txnl2
1456245_x_at 0.97423 vesicle-associated membrane protein 3 Vamp3
1456246_x_at 0.93173 trophoblast specific protein beta Tpbpb

1456247_x_at 1.00691 proteolipid protein 2 /// similar to Proteolipid protein 2 /// similar to Proteolipid protein 2 /// similar to Proteolipid protein 2 Plp2 /// LOC620648 /// LOC669344 /// LOC672630
 1456248_at 0.90044 RIKEN cDNA 2310002A05 gene /// hypothetical protein LOC630971
 2310002A05Rik /// LOC630971
 1456249_x_at 1.00742 Synaptophysin Syp
 1456250_x_at 1.01821 "transforming growth factor, beta induced" Tgfb1
 1456251_x_at 0.89525 translocator protein Tspo
 1456252_x_at 0.98206 "adaptor-related protein complex 1, sigma 2 subunit" Ap1s2
 1456253_s_at 0.82015 "pleckstrin homology domain containing, family N member 1 /// kelch-like 17 (Drosophila)" Plekhn1 /// Klhl17
 1456254_at 1.03352 "Inositol polyphosphate-4-phosphatase, type II" Inpp4b
 1456255_at 0.94807 expressed sequence AI314180 AI314180
 1456256_at 1.40733 eukaryotic translation initiation factor 5 Eif5
 1456257_at 1.38612 RIKEN cDNA C130065N10 gene C130065N10Rik
 1456258_at 0.93865 empty spiracles homolog 2 (Drosophila) Emx2
 1456259_at 0.77212 similar to integrator complex subunit 10 LOC674888
 1456260_at 2.30973 retinoblastoma binding protein 4 Rbbp4
 1456261_at 0.96089 SH3-domain kinase binding protein 1 Sh3kbp1
 1456262_at 1.24597 RNA binding motif protein 5 Rbm5
 1456263_at 0.86297 Notch gene homolog 4 (Drosophila) Notch4
 1456264_at 0.96194 COMM domain containing 7 /// similar to COMM domain containing protein 7 /// similar to COMM domain-containing protein 7 Commd7 /// LOC631742 /// LOC674161
 1456265_at 0.88533 hypothetical protein LOC624866 LOC624866
 1456266_at 1.13956 ribosomal protein L30 Rpl30
 1456267_at 0.88767 expressed sequence AI451606 AI451606
 1456268_at 1.08776 CDC28 protein kinase 1b Cks1b
 1456269_at 1.00296 preferentially expressed antigen in melanoma like 6 Prmel6
 1456270_s_at 1.08016 preferentially expressed antigen in melanoma like 6 Prmel6
 1456271_at 0.82276 "gene model 944, (NCBI)" Gm944
 1456272_at 2.61285 SET domain containing 5 Setd5
 1456273_x_at 0.78169 thiopurine methyltransferase Tpmt
 1456274_at 1.01111 RIKEN cDNA C230071H18 gene C230071H18Rik
 1456275_at 0.93393 mitochondrial ribosomal protein L21 /// hypothetical gene supported by AK003285; NM_172252 Mrpl21 /// LOC384939
 1456276_at 1.02596 --- ---
 1456277_at 0.93220 RIKEN cDNA 7530414M10 gene 7530414M10Rik
 1456278_x_at 0.94594 G protein-coupled receptor 172B Gpr172b
 1456279_a_at 0.88765 B-cell receptor-associated protein 31 Bcap31
 1456280_at 1.13208 claspin homolog (Xenopus laevis) Clspn
 1456281_at 1.15138 F-box and leucine-rich repeat protein 18 Fbxl18
 1456282_at 1.15423 CDNA clone IMAGE:30014192 ---
 1456283_at 0.93747 neuropilin (NRP) and tolloid (TLL)-like 1 Neto1
 1456284_at 1.14186 "gene model 905, (NCBI)" Gm905

1456285_at	0.95876	"Polymerase (DNA directed), alpha 1"	Pola1
1456286_at	1.13509	Component of oligomeric golgi complex 3	Cog3
1456287_at	0.88784	expressed sequence BB236558	BB236558
1456288_at	0.91756	schlafen 5	Slfn5
1456289_at	0.86673	hepcidin antimicrobial peptide 2 /// hepcidin antimicrobial peptide 1	Hamp2 /// Hamp1
1456290_x_at	0.96298	cerebral cavernous malformation 2 homolog (human)	Ccm2
1456291_x_at	0.91310	scleraxis	Scx
1456292_a_at	0.99985	vimentin	Vim
1456293_s_at	1.05621	cyclin H /// similar to cyclin H	Ccnh /// LOC672655
1456294_at	0.84818	RIKEN cDNA 0610013E23 gene	0610013E23Rik
1456295_at	0.70641	RIKEN cDNA B230114P17 gene	B230114P17Rik
1456296_at	1.58459	RIKEN cDNA 5830418K08 gene	5830418K08Rik
1456297_at	1.01064	thymic stromal cotransporter	Tscot
1456298_at	1.00576	---	---
1456299_at	1.09173	RIKEN cDNA E330012B07 gene	E330012B07Rik
1456300_at	1.04155	ilvB (bacterial acetolactate synthase)-like	Ilvbl
1456301_at	1.07821	homeo box C5	Hoxc5
1456302_at	0.88216	peroxisomal biogenesis factor 6	Pex6
1456303_at	2.89388	PHD finger protein 14	Phf14
1456304_at	0.92551	"gene model 996, (NCBI)"	Gm996
1456305_x_at	0.90924	similar to oocyte specific homeobox 2	LOC666466
1456306_a_at	1.02931	uromodulin	Umod
1456307_s_at	1.03607	adenylate cyclase 7	Adcy7
1456308_x_at	0.99175	tripartite motif protein 28	Trim28
1456309_x_at	1.06062	LIM and SH3 protein 1	Lasp1
1456310_a_at	0.99136	RIKEN cDNA 2610002J02 gene	2610002J02Rik
1456311_x_at	1.13262	establishment of cohesion 1 homolog 2 (S. cerevisiae)	Esco2
1456312_x_at	0.86934	gelsolin	Gsn
1456313_x_at	0.99393	mitochondrial ribosomal protein L28	Mrpl28
1456314_x_at	0.92846	CDK2 (cyclin-dependent kinase 2)-associated protein 1	Cdk2ap1
1456315_a_at	0.93702	"protein tyrosine phosphatase-like (proline instead of catalytic arginine), member a"	Ptpla
1456316_a_at	0.84784	acyl-Coenzyme A binding domain containing 3	Acbd3
1456317_at	0.96291	"aldehyde dehydrogenase 3 family, member B2"	Aldh3b2
1456318_at	1.17034	"C-type lectin domain family 1, member a"	Clec1a
1456319_at	1.18617	hypothetical protein LOC665081	LOC665081
1456320_at	1.01677	cDNA sequence BC049806	BC049806
1456321_at	0.84216	NIPA-like domain containing 1	Npal1
1456322_at	0.98051	---	---
1456323_at	1.01817	protein O-fucosyltransferase 1	Pofut1
1456324_at	0.79800	zinc finger protein 748	Zfp748
1456325_at	0.88089	centrosomal protein 68	Cep68

1456326_at 0.94100 "gene model 784, (NCBI)" Gm784
1456327_at 0.99752 Transcribed locus ---
1456328_at 1.04045 B-cell scaffold protein with ankyrin repeats 1 Bank1
1456329_at 1.07849 protogenin homolog (Gallus gallus) Prtg
1456330_at 0.99135 polycomb group ring finger 2 Pcgf2
1456331_at 1.01813 RIKEN cDNA 9130023H24 gene 9130023H24Rik
1456332_at 1.07797 transmembrane protein 17 Tmem17
1456333_a_at 0.88066 Rho GTPase activating protein 6 Arhgap6
1456334_s_at 1.03402 RIKEN cDNA A230106D06 gene A230106D06Rik
1456335_at 0.97555 "gene model 106, (NCBI)" Gm106
1456336_at 0.91985 RIKEN cDNA A330102K23 gene A330102K23Rik
1456337_at 0.91487 "centaurin, delta 1" Centd1
1456338_at 1.06235 RIKEN cDNA 9130023D20 gene 9130023D20Rik
1456339_at 0.91466 RIKEN cDNA 2810410D24 gene 2810410D24Rik
1456340_at 1.20766 RIKEN cDNA 2610205E22 gene 2610205E22Rik
1456341_a_at 0.72225 Kruppel-like factor 9 Klf9
1456342_at 1.00421 paired box gene 6 Pax6
1456343_at 1.00748 "solute carrier family 35, member F1" Slc35f1
1456344_at 0.98109 Tenascin C Tnc
1456345_at 0.94637 RIKEN cDNA A630057N01 gene A630057N01Rik
1456346_at 1.04850 Dynamin 1 Dnm1
1456347_at 1.97527 Leucyl/cystinyl aminopeptidase Lnpep
1456348_x_at 1.06095 progesterin and adipoQ receptor family member V Paqr5
1456349_x_at 1.04390 SMT3 suppressor of mif two 3 homolog 1 (yeast) Sumo1
1456350_at 1.29151 zinc finger protein 451 Zfp451
1456351_at 1.03349 bromodomain containing 8 Brd8
1456352_a_at 1.00808 "splicing factor 3b, subunit 2" Sf3b2
1456353_at 1.23382 suppressor of Ty 4 homolog 1 (S. cerevisiae) Supt4h1
1456354_at 0.83206 "cholinergic receptor, nicotinic, alpha polypeptide 4" Chrna4
1456355_s_at 1.00538 RIKEN cDNA 2810002G02 gene /// region containing RIKEN cDNA
2810002G02 gene; tuftelin interacting protein 11 2810002G02Rik /// LOC671202
1456356_at 1.04157 RIKEN cDNA B230217O12 gene B230217O12Rik
1456357_at 1.10263 RIKEN cDNA A930041I02 gene A930041I02Rik
1456358_at 1.33334 Ets variant gene 3 Etv3
1456359_at 0.99738 peptidylprolyl isomerase domain and WD repeat containing 1 Ppwd1
1456360_at 0.99819 "DCUN1D1 DCN1, defective in cullin neddylation 1, domain containing 1 (S.
cerevisiae)" Dcun1d1
1456361_at 0.92774 RIKEN cDNA 2600003E23 gene 2600003E23Rik
1456362_at 0.92665 "cytochrome P450, family 11, subfamily b, polypeptide 1" Cyp11b1
1456363_at 1.04028 Suppression of tumorigenicity 7-like St7l
1456364_at 0.94079 RIKEN cDNA C230057M02 gene C230057M02Rik
1456365_at 0.96601 similar to ubiquitin A-52 residue ribosomal protein fusion product 1 /// similar
to ubiquitin A-52 residue ribosomal protein fusion product 1 LOC629750 /// LOC639383

1456366_at 0.96137 RIKEN cDNA 1700015F17 gene 1700015F17Rik
1456367_at 1.00588 fucosyltransferase 8 Fut8
1456368_at 1.07305 a disintegrin and metalloprotease domain 4b Adam4b
1456369_at 1.13664 protection of telomeres 1 Pot1
1456370_s_at 0.96382 RIKEN cDNA 0610037L13 gene 0610037L13Rik
1456371_a_at 0.85009 shroom family member 1 Shroom1
1456372_at 1.08775 Integral membrane protein 2B Itm2b
1456373_x_at 1.01568 ribosomal protein S20 Rps20
1456374_x_at 1.03973 "eukaryotic translation initiation factor 3, subunit 8" Eif3s8
1456375_x_at 0.96264 tripartite motif protein 27 Trim27
1456376_at 1.12466 Matrix metalloproteinase 24 Mmp24
1456377_x_at 0.98585 LIM domain containing 2 /// similar to epithelial protein lost in neoplasm
Limd2 /// LOC632329
1456378_s_at 0.93598 F-box and leucine-rich repeat protein 20 Fbxl20
1456379_x_at 0.90727 delta/notch-like EGF-related receptor Dner
1456380_x_at 0.88969 "calponin 3, acidic" Cnn3
1456381_x_at 0.85563 myeloid cell leukemia sequence 1 Mcl1
1456382_at 1.06411 "ATPase family, AAA domain containing 1" Atad1
1456383_at 1.03072 ribosomal L1 domain containing 1 Rsl1d1
1456384_at 1.00397 neuroligin 3 Nlgn3
1456385_x_at 1.01972 UBX domain containing 3 Ubx3
1456386_at 1.68643 "RNA-binding region (RNP1, RRM) containing 2" Rnpc2
1456387_at 0.95543 nucleolar protein 4 Nol4
1456388_at 0.92617 "ATPase, class VI, type 11A" Atp11a
1456389_at 0.96782 zinc finger homeobox 1b Zfx1b
1456390_at 1.23812 --- ---
1456391_at 0.92929 tudor domain containing 5 Tdrd5
1456392_at 1.08153 neuronal growth regulator 1 Negr1
1456393_at 0.79157 RIKEN cDNA 2310002J21 gene 2310002J21Rik
1456394_at 0.98064 RAD50 interactor 1 Rint1
1456395_at 0.77438 "peroxisome proliferative activated receptor, gamma, coactivator 1 alpha"
Ppargc1a
1456396_at 1.18171 --- ---
1456397_at 0.96952 cadherin 4 Cdh4
1456398_at 0.94723 taurine upregulated gene 1 Tug1
1456399_at 0.98522 --- ---
1456400_at 0.94188 "telomeric repeat binding factor 2, interacting protein" Terf2ip
1456401_at 0.80879 "calcium channel, voltage-dependent, beta 2 subunit" Cacnb2
1456402_at 1.01816 RIKEN cDNA A330076H08 gene A330076H08Rik
1456403_at 0.93732 phosphoprotein associated with glycosphingolipid microdomains 1 Pag1
1456404_at 0.57131 "a disintegrin-like and metalloproteinase (reprolysin type) with thrombospondin
type 1 motif, 5 (aggrecanase-2)" Adamts5
1456405_at 0.89965 death inducer-obliterator 1 Dido1

1456406_at 1.08296 --- ---
1456407_a_at 1.16808 tousled-like kinase 1 Tlk1
1456408_x_at 1.37614 RIKEN cDNA 4933439C10 gene 4933439C10Rik
1456409_at 1.01826 --- ---
1456410_at 1.14028 expressed sequence AU045128 AU045128
1456411_at 1.07243 RCC1 domain containing 1 Rccd1
1456412_a_at 1.02146 putative phosphatase RP23-136K12.4
1456413_at 1.25197 phosphodiesterase 4D interacting protein (myomegalin) Pde4dip
1456414_at 0.90327 "In vitro fertilized eggs cDNA, RIKEN full-length enriched library, clone:7420414K24 product:unclassifiable, full insert sequence" ---
1456415_at 0.90788 zinc finger protein 451 Zfp451
1456416_at 0.95381 "core-binding factor, runt domain, alpha subunit 2, translocated to, 2 homolog (human)" Cbfa2t2h
1456417_at 0.90743 zinc finger protein of the cerebellum 4 Zic4
1456418_at 0.99542 "PREDICTED: Mus musculus similar to Inward rectifier potassium channel 13 (Potassium channel, inwardly rectifying, subfamily J, member 13) (Inward rectifier K(+)) channel Kir7.1) (LOC620405), mRNA" ---
1456419_at 1.09753 RIKEN cDNA 5730455P16 gene 5730455P16Rik
1456420_at 0.82984 AT rich interactive domain 4A (Rbp1 like) Arid4a
1456421_at 0.99440 RIKEN cDNA 4930578M01 gene 4930578M01Rik
1456422_at 0.83504 --- ---
1456423_at 0.77746 methyl-CpG binding domain protein 5 Mbd5
1456424_s_at 0.93517 phospholipid transfer protein Pltp
1456425_at 0.97776 RIKEN cDNA 9230115E21 gene 9230115E21Rik
1456426_at 1.09962 --- ---
1456427_at 0.96048 "glycoprotein Ib, beta polypeptide" Gp1bb
1456428_at 1.08973 Chemokine (C-X-C motif) ligand 15 Cxcl15
1456429_at 1.24537 mucosa associated lymphoid tissue lymphoma translocation gene 1 Malt1
1456430_at 1.37612 coiled-coil domain containing 39 Ccdc39
1456431_at 0.95162 zinc finger protein 64 Zfp64
1456432_at 0.95142 GRB2-related adaptor protein 2 Grap2
1456433_at 1.04033 regulator of chromosome condensation (RCC1) and BTB (POZ) domain containing protein 1 Rcbtb1
1456434_x_at 0.84685 heat shock protein 8 Hspb8
1456435_at 1.00003 MORN repeat containing 1 Morn1
1456436_x_at 1.01397 ribosomal protein S20 Rps20
1456437_x_at 1.45212 "complement component 1, r subcomponent /// similar to Complement C1r-B subcomponent precursor (Complement component 1, r-B subcomponent) /// similar to complement component 1, r subcomponent" C1r /// LOC667277 /// LOC677405
1456438_x_at 0.96070 ribophorin I Rpn1
1456439_x_at 0.88468 "microtubule associated monooxygenase, calponin and LIM domain containing 1" Mical1
1456440_s_at 0.99215 "ST8 alpha-N-acetyl-neuraminide alpha-2,8-sialyltransferase 6" St8sia6

1456441_at 1.15363 coiled-coil domain containing 87 Ccdc87
1456442_at 0.85350 RAB3A interacting protein (rabin3)-like 1 Rab3il1
1456443_at 1.02018 RIKEN cDNA 4930404N11 gene 4930404N11Rik
1456444_at 0.97531 F-box protein 41 Fbxo41
1456445_at 1.13566 RIKEN cDNA 4930563D23 gene 4930563D23Rik
1456446_at 1.18986 RIKEN cDNA 4930523C07 gene 4930523C07Rik
1456447_at 0.80046 --- ---
1456448_at 1.03680 RIKEN cDNA 2610301B20 gene 2610301B20Rik
1456449_at 2.30845 Suppressor of Ty 16 homolog (S. cerevisiae) Supt16h
1456450_at 1.22842 "Cystinosis, nephropathic" Ctns
1456451_at 0.87889 CEA-related cell adhesion molecule 15 Ceacam15
1456452_at 1.03153 cDNA sequence BC107230 BC107230
1456453_at 1.48194 Ena-vasodilator stimulated phosphoprotein Evl
1456454_at 0.98508 cartilage homeo protein 1 Cart1
1456455_at 0.98299 mortality factor 4 like 1 /// similar to mortality factor 4 like 1 isoform b ///
similar to mortality factor 4 like 1 /// similar to mortality factor 4 like 1 (predicted) Morf4l1 ///
LOC433955 /// LOC627352 /// LOC635663
1456456_x_at 0.95705 Melanoma antigen Mela
1456457_at 1.08509 "Solute carrier family 32 (GABA vesicular transporter), member 1" Slc32a1
1456458_at 0.94705 --- ---
1456459_x_at 0.99869 coiled-coil domain containing 53 Ccdc53
1456460_at 1.06887 cDNA sequence BC068281 BC068281
1456461_at 1.07549 Ewing sarcoma breakpoint region 1 Ewsr1
1456462_x_at 0.85884 "protein phosphatase 1, catalytic subunit, beta isoform" Ppp1cb
1456463_at 0.99342 RIKEN cDNA A230106M20 gene A230106M20Rik
1456464_x_at 0.93711 --- ---
1456465_at 1.07868 RIKEN cDNA B430110G05 gene B430110G05Rik
1456466_x_at 0.91242 ataxin 10 Atxn10
1456467_s_at 0.88752 nemo like kinase Nlk
1456468_x_at 0.84236 RIKEN cDNA 1110012N22 gene 1110012N22Rik
1456469_x_at 0.91138 LIM and SH3 protein 1 Lasp1
1456470_x_at 0.87751 "tubulin, beta 2c" Tubb2c
1456471_x_at 0.91633 3-phosphoglycerate dehydrogenase /// similar to 3-phosphoglycerate
dehydrogenase /// similar to 3-phosphoglycerate dehydrogenase /// similar to 3-phosphoglycerate
dehydrogenase /// similar to 3-phosphoglycerate dehydrogenase /// similar to 3-phosphoglycerate
dehydrogenase /// similar to 3-phosphoglycerate dehydrogenase /// similar to 3-phosphoglycerate
dehydrogenase /// similar to 3-phosphoglycerate dehydrogenase /// similar to 3-phosphoglycerate
dehydrogenase /// similar to 3-phosphoglycerate dehydrogenase /// similar to 3-phosphoglycerate
dehydrogenase /// similar to 3-phosphoglycerate dehydrogenase /// similar to 3-phosphoglycerate
dehydrogenase /// similar to 3-phosphoglycerate dehydrogenase Phgdh /// LOC382931 ///
LOC384524 /// LOC385344 /// LOC547171 /// LOC627427 /// LOC666422 /// LOC666875 ///
LOC668771 /// LOC669671 /// LOC669985 /// LOC670155 /// LOC670187 /// LOC671102 ///

LOC671972 /// LOC673015 /// LOC675010

1456472_at 0.97681 armadillo repeat containing 4 Armc4
 1456473_x_at 0.88988 ADP-ribosylation factor 2 Arf2
 1456474_at 0.98960 --- ---
 1456475_s_at 0.91567 "protein kinase, cAMP dependent regulatory, type II beta" Prkar2b
 1456476_at 1.03601 ataxin 2-like Atxn2l
 1456477_at 1.05063 cyclin T1 Ccnt1
 1456478_at 1.13679 phosphoglucomutase 2-like 1 Pgm2l1
 1456479_at 0.89195 sorting nexin family member 30 Snx30
 1456480_at 0.61416 furry homolog (Drosophila) Fry
 1456481_at 0.98533 "DNA segment, Chr 9, ERATO Doi 280, expressed" D9Ert280e
 1456482_at 0.84447 "phosphatidylinositol 3 kinase, regulatory subunit, polypeptide 3 (p55)"
 Pik3r3
 1456483_at 0.80717 zinc finger protein 9 Zfp9
 1456484_at 0.85414 dehydrogenase/reductase (SDR family) X chromosome Dhrrx
 1456485_at 0.94676 nuclear protein in the AT region Npat
 1456486_at 0.91194 zinc finger protein 574 Zfp574
 1456487_at 0.75843 adenylate cyclase 1 Adcy1
 1456488_at 1.04958 WD repeat domain 33 Wdr33
 1456489_at 1.02159 cleavage and polyadenylation factor subunit homolog (S. cerevisiae) Pcf11
 1456490_at 0.96970 CDK2-associated protein 2 Cdk2ap2
 1456491_at 0.91683 Expressed sequence AI606861 AI606861
 1456492_at 1.14800 RIKEN cDNA 9130404D08 gene 9130404D08Rik
 1456493_at 1.13792 RIKEN cDNA 5031415C07 gene /// RIKEN cDNA 5830484A20 gene ///
 similar to Sp110 nuclear body protein /// similar to Sp110 nuclear body protein /// similar to Sp110
 nuclear body protein /// similar to Sp110 nuclear body protein /// similar to Sp110 nuclear body
 protein /// similar to Sp110 nuclear body protein /// similar to Sp110 nuclear body protein /// similar to
 Sp110 nuclear body protein /// similar to Sp110 nuclear body protein /// similar to Sp110 nuclear body
 protein 5031415C07Rik /// 5830484A20Rik /// LOC619671 /// MGC118608 /// LOC671287 ///
 LOC673121 /// LOC673432 /// LOC673556 /// LOC673850 /// LOC673913 /// LOC673955 ///
 LOC673970
 1456494_a_at 1.02850 tripartite motif protein 30 /// expressed sequence AI451617 Trim30 ///
 AI451617
 1456495_s_at 0.87107 oxysterol binding protein-like 6 Osbpl6
 1456496_at 1.03549 Death-associated kinase 3 Dapk3
 1456497_x_at 1.01170 ribosomal protein S10 Rps10
 1456498_at 0.93976 integrin alpha 4 Itga4
 1456499_at 0.74458 hypothetical LOC552912 LOC552912
 1456500_at 0.79866 anterior pharynx defective 1b homolog (C. elegans) Aph1b
 1456501_at 0.90763 RIKEN cDNA 2610528E23 gene 2610528E23Rik
 1456502_at 0.95798 dual-specificity tyrosine-(Y)-phosphorylation regulated kinase 2 Dyrk2
 1456503_at 1.19031 nucleoporin 214 Nup214
 1456504_at 1.01198 zinc finger protein 182 Zfp182

1456505_at 0.90425 Braf transforming gene Braf
1456506_at 1.21876 PRP38 pre-mRNA processing factor 38 (yeast) domain containing B Prpf38b
1456507_at 1.10440 zinc finger protein 454 Zfp454
1456508_at 1.21450 RIKEN cDNA A030001H23 gene A030001H23Rik
1456509_at 0.92689 RIKEN cDNA 1110032F04 gene 1110032F04Rik
1456510_x_at 0.92013 UbiE-YGHL1 fusion protein Ubie
1456511_x_at 1.04519 ES cell-expressed Ras Eras
1456512_at 1.01116 PDZ domain containing RING finger 4 Pdzn4
1456513_at 1.14306 Early B-cell factor 3 Ebf3
1456514_at 2.92346 --- ---
1456515_s_at 0.92052 transcription factor-like 5 (basic helix-loop-helix) Tcf15
1456516_x_at 1.09768 UDP-N-acetylglucosamine pyrophosphorylase 1 /// similar to UDP-N-acetylhexosamine pyrophosphorylase Uap1 /// LOC640502
1456517_at 0.91808 transmembrane protein 44 Tmem44
1456518_at 0.89742 RIKEN cDNA 4930422I07 gene 4930422I07Rik
1456519_at 1.13087 Expressed sequence AW493845 AW493845
1456520_at 1.00006 RIKEN cDNA 9530033F24 gene 9530033F24Rik
1456521_at 0.86936 Transcribed locus ---
1456522_at 0.89318 --- ---
1456523_at 1.05661 expressed sequence C77713 C77713
1456524_at 1.11242 neuregulin 1 Nrg1
1456525_at 0.99319 EST AA881470 AA881470
1456526_at 1.76864 RIKEN cDNA C130034I24 gene C130034I24Rik
1456527_at 0.96302 "HECT, C2 and WW domain containing E3 ubiquitin protein ligase 1"
Hecw1
1456528_x_at 1.19094 nucleolin Ncl
1456529_at 0.96351 --- ---
1456530_x_at 1.12909 "elongation of very long chain fatty acids (FEN1/Elo2, SUR4/Elo3, yeast)-like 1" Elovl1
1456531_x_at 1.02109 PRP19/PSO4 pre-mRNA processing factor 19 homolog (S. cerevisiae)
Prpf19
1456532_at 1.06140 "Platelet-derived growth factor, D polypeptide" Pdgfd
1456533_at 1.43519 dpy-19-like 1 (C. elegans) Dpy1911
1456534_at 0.86615 "Mannosidase, alpha, class 1A, member 2" Man1a2
1456535_at 0.90075 vacuolar protein sorting 13C (yeast) Vps13c
1456536_at 0.82929 THAP domain containing 6 Thap6
1456537_at 1.02060 RIKEN cDNA A930033H14 gene A930033H14Rik
1456538_at 0.74660 serologically defined colon cancer antigen 8 Sdccag8
1456539_at 1.05419 --- ---
1456540_s_at 0.99872 myotubularin related protein 6 Mtmr6
1456541_x_at 0.94571 "ATPase family, AAA domain containing 3A" Atad3a
1456542_s_at 1.03937 glutaminyl-tRNA synthase (glutamine-hydrolyzing)-like 1 Qrs11
1456543_at 0.73386 prokineticin receptor 1 Prokr1

1456544_at 1.06257 --- ---
1456545_at 0.99611 interleukin 18 receptor accessory protein Il18rap
1456546_at 0.90426 RIKEN cDNA 1700097N02 gene 1700097N02Rik
1456547_at 2.45211 Sloan-Kettering viral oncogene homolog Ski
1456548_at 1.45795 Metadherin Mtdh
1456549_at 1.07980 "gene model 550, (NCBI)" Gm550
1456550_at 1.10856 --- ---
1456551_at 0.83463 "Aldehyde dehydrogenase 1 family, member L1" Aldh111
1456552_at 1.07135 "2 days pregnant adult female ovary cDNA, RIKEN full-length enriched library, clone:E330015D05 product:hypothetical protein, full insert sequence" ---
1456553_at 1.91596 "Zinc finger, CCHC domain containing 11" Zcchc11
1456554_at 0.93144 RIKEN cDNA 7530414M10 gene 7530414M10Rik
1456555_at 1.06467 coiled-coil domain containing 67 Ccdc67
1456556_at 1.13060 RIKEN cDNA A230009B12 gene A230009B12Rik
1456557_at 0.89205 RIKEN cDNA 1700041C02 gene 1700041C02Rik
1456558_s_at 0.90526 expressed sequence C87977 C87977
1456559_at 1.00803 empty spiracles homolog 2 (Drosophila) opposite strand Emx2os
1456560_at 1.06355 "gene model 93, (NCBI)" Gm93
1456561_s_at 0.83679 Kruppel-like factor 17 Klf17
1456562_x_at 1.03942 KIAA0415 protein KIAA0415
1456563_at 1.05402 RIKEN cDNA 4933429F08 gene 4933429F08Rik
1456564_at 1.01894 RIKEN cDNA C030046I01 gene C030046I01Rik
1456565_s_at 0.93657 mitogen activated protein kinase kinase kinase 12 Map3k12
1456566_x_at 1.10054 RNA binding motif protein 14 Rbm14
1456567_x_at 0.91199 granulin Grn
1456568_at 0.96275 Gelsolin Gsn
1456569_x_at 0.91875 gelsolin Gsn
1456570_at 1.00677 --- ---
1456571_at 1.03306 "decapping enzyme, scavenger" Dcps
1456572_x_at 0.99607 chaperonin subunit 4 (delta) Cct4
1456573_x_at 0.87566 nicotinamide nucleotide transhydrogenase Nnt
1456574_at 0.94902 EST AA407452 AA407452
1456575_at 1.01874 Necdin Ndn
1456576_x_at 1.02724 "CCR4-NOT transcription complex, subunit 2" Cnot2
1456577_x_at 0.98389 pitrilysin metallopeptidase 1 Pitrm1
1456578_x_at 1.02495 LIM and SH3 protein 1 Lasp1
1456579_x_at 1.02562 RIKEN cDNA 1200003M09 gene 1200003M09Rik
1456580_s_at 0.92538 "ATP synthase, H+ transporting, mitochondrial F1 complex, delta subunit" Atp5d
1456581_x_at 0.96988 guanosine diphosphate (GDP) dissociation inhibitor 2 Gdi2
1456582_x_at 0.90461 RIKEN cDNA 5230400G24 gene /// hypothetical protein LOC637796
5230400G24Rik /// LOC637796
1456583_x_at 0.89791 transmembrane protein 144 Tmem144

1456584_x_at 1.06650 3-phosphoglycerate dehydrogenase /// similar to 3-phosphoglycerate dehydrogenase /// similar to 3-phosphoglycerate dehydrogenase /// similar to 3-phosphoglycerate dehydrogenase /// similar to 3-phosphoglycerate dehydrogenase /// similar to 3-phosphoglycerate dehydrogenase /// similar to 3-phosphoglycerate dehydrogenase /// similar to 3-phosphoglycerate dehydrogenase /// similar to 3-phosphoglycerate dehydrogenase /// similar to 3-phosphoglycerate dehydrogenase /// similar to 3-phosphoglycerate dehydrogenase /// similar to 3-phosphoglycerate dehydrogenase Phgdh /// LOC382931 /// LOC384524 /// LOC385344 /// LOC627427 /// LOC666422 /// LOC666875 /// LOC668771 /// LOC669671 /// LOC669985 /// LOC670155 /// LOC670187 /// LOC671102 /// LOC671972 /// LOC673015 /// LOC675010
 1456585_x_at 0.93439 RIKEN cDNA E130309D02 gene E130309D02Rik
 1456586_x_at 1.10878 major vault protein Mvp
 1456587_x_at 0.89885 RIKEN cDNA 2010005J08 gene 2010005J08Rik
 1456588_x_at 0.94392 "cytochrome c oxidase, subunit Vb" Cox5b
 1456589_x_at 0.98461 allantoicase Allc
 1456590_x_at 0.99085 "aldo-keto reductase family 1, member B3 (aldose reductase)" Akr1b3
 1456591_x_at 0.94961 A kinase (PRKA) anchor protein 3 Akap3
 1456592_at 0.91006 Vacuolar protein sorting 13A (yeast) Vps13a
 1456593_at 1.09481 Synaptotagmin-like 2 Sytl2
 1456594_at 1.00731 --- ---
 1456595_x_at 1.33677 growth hormone Gh
 1456596_at 0.98308 RIKEN cDNA 6430550H21 gene 6430550H21Rik
 1456597_at 1.04210 HEAT repeat containing 3 Heatr3
 1456598_at 1.05891 similar to PRAME family member 8 /// expressed sequence AU018829 /// similar to PRAME family member 8 /// similar to PRAME family member 8 /// similar to PRAME family member 8 /// similar to PRAME family member 8 /// similar to PRAME family member 8 /// similar to PRAME family member 8 /// similar to PRAME family member 8 /// similar to PRAME family member 8 /// similar to PRAME family member 8 /// similar to PRAME family member 8 /// similar to PRAME family member 8 /// similar to PRAME family member 8 /// similar to PRAME family member 8 /// similar to PRAME family member 8 /// similar to PRAME family member 8 /// similar to PRAME family member 8 LOC545761 /// AU018829 /// LOC546201 /// LOC619753 /// LOC621019 /// LOC622792 /// LOC664993 /// LOC665003 /// LOC665558 /// LOC665616 /// LOC665711 /// LOC665718 /// LOC665723 /// LOC665802 /// LOC666058 /// LOC666140
 1456599_at 0.88725 nuclear transport factor 2-like export factor 2 Nxt2
 1456600_a_at 0.96631 ring finger protein 7 Rnf7
 1456601_x_at 1.04314 FXYP domain-containing ion transport regulator 2 Fxyd2
 1456602_at 0.81430 RIKEN cDNA 4932417I16 gene 4932417I16Rik
 1456603_at 0.90520 RIKEN cDNA 1500005K14 gene 1500005K14Rik
 1456604_a_at 0.93610 protein-L-isoaspartate (D-aspartate) O-methyltransferase 1 Pcmt1
 1456605_at 1.06150 "CCAAT/enhancer binding protein (C/EBP), delta" Cebpδ
 1456606_a_at 0.94190 carbohydrate sulfotransferase 11 /// phosphatase and actin regulator 1 Chst11 /// Phactr1

1456607_at 0.93178 valosin containing protein (p97)/p47 complex interacting protein 1 Vcpi1
1456608_at 1.06457 --- ---
1456609_at 0.82626 calcium/calmodulin-dependent protein kinase II inhibitor 1 Camk2n1
1456610_at 0.92893 jumonji domain containing 3 Jmjd3
1456611_at 0.77669 RIKEN cDNA D430015B01 gene D430015B01Rik
1456612_at 1.06616 Interleukin 17D II17d
1456613_at 0.90128 eukaryotic translation initiation factor 4E binding protein 2 Eif4ebp2
1456614_at 0.97704 ACN9 homolog (S. cerevisiae) Acn9
1456615_a_at 1.02198 fetal Alzheimer antigen Falz
1456616_a_at 0.95816 basigin Bsg
1456617_a_at 1.15093 "eukaryotic translation initiation factor 2, subunit 2 (beta)" Eif2s2
1456618_at 0.97365 MAP/microtubule affinity-regulating kinase 4 Mark4
1456619_at 0.90255 "Lipase, member H" Liph
1456620_at 1.17613 "N-acetylglucosamine-1-phosphate transferase, alpha and beta subunits" Gnptab
1456621_at 0.79782 RIKEN cDNA D430015B01 gene D430015B01Rik
1456622_at 0.83523 calmodulin regulated spectrin-associated protein 1 Camsap1
1456623_at 1.11381 "tropomyosin 1, alpha" Tpm1
1456624_at 0.78963 "WD repeat domain, phosphoinositide interacting 1" Wipi1
1456625_at 0.91531 --- ---
1456626_a_at 1.19957 RIKEN cDNA 1110005A23 gene /// similar to cytokine induced protein 29
kDa /// similar to cytokine induced protein 29 kDa 1110005A23Rik /// LOC625193 /// LOC636633
1456627_at 0.99970 Ubiquilin 2 Ubqln2
1456628_x_at 1.02139 ribosomal protein S24 Rps24
1456629_at 1.15143 ankyrin repeat domain 47 Ankrd47
1456630_x_at 0.94828 Son cell proliferation protein Son
1456631_at 1.18366 RIKEN cDNA 6430706D22 gene 6430706D22Rik
1456632_at 0.94111 B-cell CLL/lymphoma 11A (zinc finger protein) Bcl11a
1456633_at 0.98365 "transient receptor potential cation channel, subfamily M, member 3" Trpm3
1456634_at 0.84453 RIKEN cDNA 9830001H06 gene 9830001H06Rik
1456635_at 1.04444 --- ---
1456636_at 0.75191 RIKEN cDNA 4930414L22 gene 4930414L22Rik
1456637_at 1.15634 leucine rich repeat transmembrane neuronal 2 Lrrtm2
1456638_at 1.05180 WD repeat domain 59 Wdr59
1456639_at 1.07785 zinc finger protein 398 Zfp398
1456640_at 1.01514 SH3 domain containing ring finger 2 Sh3rf2
1456641_at 0.64939 RIKEN cDNA 1190007F08 gene 1190007F08Rik
1456642_x_at 1.05123 S100 calcium binding protein A10 (calpactin) S100a10
1456643_at 1.10915 RIKEN cDNA 9230114K14 gene 9230114K14Rik
1456644_at 0.96592 RIKEN cDNA 5730528L13 gene 5730528L13Rik
1456645_at 1.01663 WD repeat domain 25 Wdr25
1456646_at 1.06427 peroxisome biogenesis factor 10 Pex10
1456647_a_at 0.92426 eyes absent 4 homolog (Drosophila) Eya4
1456648_at 1.03330 "Mannosidase 1, alpha" Man1a

1456649_at 1.02387 Sorting nexin 15 Snx15
1456650_at 1.16495 "6 days neonate skin cDNA, RIKEN full-length enriched library, clone: A030010B11 product:unclassifiable, full insert sequence" ---
1456651_a_at 1.02109 translocated promoter region Tpr
1456652_at 2.27758 denticleless homolog (Drosophila) Dtl
1456653_a_at 1.26167 methylenetetrahydrofolate dehydrogenase (NADP+ dependent) 1-like Mthfd11
1456654_at 1.01298 progesterin and adipoQ receptor family member V Paqr5
1456655_at 2.38263 Exostoses (multiple) 1 Ext1
1456656_at 0.93419 lin-7 homolog A (C. elegans) Lin7a
1456657_at 1.48517 expressed sequence AA415437 AA415437
1456658_at 1.05462 "actin, alpha 2, smooth muscle, aorta" Acta2
1456659_at 1.90289 hypothetical LOC552902 LOC552902
1456660_a_at 1.02605 RIKEN cDNA 0610010F05 gene 0610010F05Rik
1456661_at 2.09722 "Jumonji, AT rich interactive domain 2" Jarid2
1456662_at 1.67978 Hypothetical protein LOC546058 LOC546058
1456663_x_at 0.87628 TM2 domain containing 2 Tm2d2
1456664_x_at 0.99325 heterogeneous nuclear ribonucleoprotein F /// similar to heterogeneous nuclear ribonucleoprotein F /// similar to heterogeneous nuclear ribonucleoprotein F /// similar to heterogeneous nuclear ribonucleoprotein F Hnrpf /// LOC637008 /// LOC665008 /// LOC672616
1456665_at 0.76433 eyes absent 4 homolog (Drosophila) Eya4
1456666_at 0.88995 RIKEN cDNA 2610020C07 gene 2610020C07Rik
1456667_at 1.62503 Huntington disease gene homolog Hdh
1456668_at 1.61868 "Zinc finger, CCHC domain containing 11" Zcchc11
1456669_at 1.03872 Transcribed locus ---
1456670_at 0.99806 RIKEN cDNA A930007A09 gene A930007A09Rik
1456671_at 0.98508 transforming growth factor beta regulated gene 3 Tbrg3
1456672_at 1.08900 "CDNA, clone:Y1G0143G21, strand:unspecified" ---
1456673_at 1.04658 expressed sequence AL022943 AL022943
1456674_at 1.04179 RIKEN cDNA E130016E03 gene E130016E03Rik
1456675_at 0.88559 expressed sequence AA536749 AA536749
1456676_a_at 0.95848 "6-phosphofructo-2-kinase/fructose-2,6-biphosphatase 3" Pfkfb3
1456677_at 1.25815 Hect domain and RLD 4 Herc4
1456678_at 1.02423 Src family associated phosphoprotein 1 Scap1
1456679_at 1.10952 Oligonucleotide/oligosaccharide-binding fold containing 1 Obfc1
1456680_at 1.01364 "UDP-GlcNAc:betaGal beta-1,3-N-acetylglucosaminyltransferase 6 (core 3 synthase)" B3gnt6
1456681_at 0.97428 expressed sequence AI848149 AI848149
1456682_at 0.95471 LON peptidase N-terminal domain and ring finger 2 Lonrf2
1456683_at 1.59129 "SAFB-like, transcription modulator" Sltm
1456684_at 1.02476 transmembrane protein 74 Tmem74
1456685_at 0.97216 neuron specific gene family member 2 Nsg2
1456686_at 1.38329 Zinc finger homeobox 1a Zfhx1a

1456687_at 1.11408 Retinitis pigmentosa 9 homolog (human) Rp9h
1456688_at 1.11018 Topoisomerase (DNA) II beta Top2b
1456689_at 1.40137 Ring finger protein 10 Rnf10
1456690_at 1.00036 --- ---
1456691_s_at 1.01863 steroid 5 alpha-reductase 2-like Srd5a2l
1456692_at 0.81068 Bromodomain containing 1 Brd1
1456693_at 0.96527 RIKEN cDNA 4933406F09 gene 4933406F09Rik
1456694_x_at 1.09814 "protein tyrosine phosphatase, non-receptor type 6" Ptpn6
1456695_x_at 0.90474 anaphase-promoting complex subunit 5 Anapc5
1456696_x_at 1.20026 "RMI1, RecQ mediated genome instability 1, homolog (S. cerevisiae)" Rmi1
1456697_x_at 0.90920 cyclin D binding myb-like transcription factor 1 Dmtf1
1456698_s_at 1.01128 heterogeneous nuclear ribonucleoprotein D-like Hnrpdl
1456699_s_at 0.98185 RIKEN cDNA A730098D12 gene A730098D12Rik
1456700_x_at 0.92873 "0 day neonate cerebellum cDNA, RIKEN full-length enriched library, clone: C230070N19 product:myristoylated alanine rich protein kinase C substrate, full insert sequence" ---
1456701_at 0.99531 RIKEN cDNA B230208H17 gene B230208H17Rik
1456702_x_at 0.91349 "methionine adenosyltransferase II, alpha" Mat2a
1456703_at 1.14982 "DNA segment, Chr 5, ERATO Doi 301, expressed" D5Ertd301e
1456704_at 1.13334 RIKEN cDNA 9530010C24 gene 9530010C24Rik
1456705_at 0.98067 Zinc finger protein 532 Zfp532
1456706_at 2.06969 RIKEN cDNA 4833441D16 gene 4833441D16Rik
1456707_at 0.98990 "Adult male corpora quadrigemina cDNA, RIKEN full-length enriched library, clone:B230387P15 product:unclassifiable, full insert sequence" ---
1456708_at 1.02850 RIKEN cDNA 2610028E06 gene 2610028E06Rik
1456709_at 1.00667 oligophrenin 1 Ophn1
1456710_at 0.97613 Protocadherin beta 11 Pcdhb11
1456711_at 1.07667 RIKEN cDNA 4932425I24 gene 4932425I24Rik
1456712_at 1.06972 ligand dependent nuclear receptor corepressor-like Lcorl
1456713_at 1.48594 "polymerase (DNA directed), alpha 2" Pola2
1456714_at 0.94680 WW domain containing E3 ubiquitin protein ligase 2 Wwp2
1456715_at 1.40549 "Pam, highwire, rpm 1" Phr1
1456716_s_at 1.01683 RIKEN cDNA 3110002H16 gene 3110002H16Rik
1456717_at 1.97277 TEA domain family member 1 Tead1
1456718_at 0.94400 transmembrane protein 56 Tmem56
1456719_at 0.85719 "2 cells egg cDNA, RIKEN full-length enriched library, clone:B020004L17 product:unclassifiable, full insert sequence" ---
1456720_at 1.07892 RIKEN cDNA 1500041B16 gene 1500041B16Rik
1456721_at 0.65966 "gene model 837, (NCBI)" Gm837
1456722_at 0.96042 chordin-like 1 Chrdl1
1456723_at 1.00084 Proline rich 14 Prr14
1456724_x_at 0.93816 WD repeat domain 79 Wdr79
1456725_x_at 0.98489 villin 2 Vil2
1456726_x_at 0.92749 glutaminyl-tRNA synthetase Qars

1456727_a_at 0.98918 "casein kinase 1, delta" Csnk1d
1456728_x_at 1.00208 aconitase 1 Aco1
1456729_x_at 0.98605 regulator of telomere elongation helicase 1 Rtel1
1456730_x_at 0.99435 actin-like 6A Actl6a
1456731_x_at 0.93951 Polymerase (RNA) III (DNA directed) polypeptide K Polr3k
1456732_at 0.90278 Glycerol-3-phosphate dehydrogenase 1 (soluble) Gpd1
1456733_x_at 0.82900 "serine (or cysteine) peptidase inhibitor, clade H, member 1" Serpinh1
1456734_at 1.04121 thioredoxin 1 Txn1
1456735_x_at 0.59954 acid phosphatase-like 2 Acpl2
1456736_x_at 0.90644 RIKEN cDNA 5230400G24 gene 5230400G24Rik
1456737_x_at 0.93654 acetyl-Coenzyme A acyltransferase 1A Acaa1a
1456738_s_at 1.14005 brain protein 16 Brp16
1456739_x_at 1.02354 "armadillo repeat containing, X-linked 2" Armcx2
1456740_x_at 0.90305 component of oligomeric golgi complex 1 Cog1
1456741_s_at 0.94214 glycoprotein m6a Gpm6a
1456742_x_at 1.02335 transmembrane 9 superfamily member 2 Tm9sf2
1456743_x_at 0.96984 mortality factor 4 like 2 /// similar to Mortality factor 4-like protein 2 (MORF-related gene X protein) (Transcription factor-like protein MRGX) (Sid 393) /// similar to Mortality factor 4-like protein 2 (MORF-related gene X protein) (Transcription factor-like protein MRGX) (Sid 393) Morf4l2 /// LOC433261 /// LOC677515
1456744_x_at 1.10917 folliculin Flcn
1456745_x_at 0.93367 ERGIC and golgi 3 Ergic3
1456746_a_at 0.86050 Cd99 antigen-like 2 Cd99l2
1456747_x_at 0.95796 Cd99 antigen-like 2 Cd99l2
1456748_a_at 0.94920 4-nitrophenylphosphatase domain and non-neuronal SNAP25-like protein homolog 1 (C. elegans) Nipsnap1
1456749_at 1.00575 Transcribed locus ---
1456750_at 0.93131 RIKEN cDNA B230303O12 gene B230303O12Rik
1456751_x_at 0.92625 hypothetical protein A530021J07 A530021J07
1456752_at 1.20393 "13 days embryo heart cDNA, RIKEN full-length enriched library, clone: D330038P10 product:Mus musculus Nespas mRNA, full insert sequence" ---
1456753_at 0.86945 --- ---
1456754_at 1.00465 "zinc finger, FYVE domain containing 20" Zfyve20
1456755_at 0.92935 "trafficking protein, kinesin binding 1" Trak1
1456756_at 0.87533 RIKEN cDNA E430016P22 gene E430016P22Rik
1456757_at 1.06732 RIKEN cDNA G630024C07 gene G630024C07Rik
1456758_at 2.63128 RIKEN cDNA 9930017N22 gene 9930017N22Rik
1456759_at 1.02802 "16 days neonate cerebellum cDNA, RIKEN full-length enriched library, clone:9630048A22 product:unclassifiable, full insert sequence" ---
1456760_at 0.87461 CDNA clone IMAGE:6813709 ---
1456761_at 1.53580 RIKEN cDNA D630030B22 gene D630030B22Rik
1456762_at 0.89174 --- ---
1456763_at 0.75563 expressed sequence AA536749 AA536749

1456764_at	1.02666	"solute carrier family 35, member F3"	Slc35f3
1456765_at	0.91332	hypothetical protein 6430511F03	6430511F03
1456766_at	0.89770	RIKEN cDNA C330001K17 gene	C330001K17Rik
1456767_at	0.81869	leucine rich repeat and fibronectin type III domain containing 3	Lrn3
1456768_a_at	0.95612	multimerin 2	Mmrn2
1456769_at	1.31098	dual specificity phosphatase 3 (vaccinia virus phosphatase VH1-related)	Dusp3
1456770_at	0.96059	zinc finger protein 14	Zfp14
1456771_at	1.07546	zyg-11 homolog B (C. elegans)-like	Zyg11bl
1456772_at	0.93447	neutrophil cytosolic factor 1	Ncf1
1456773_at	1.12746	nucleoporin like 2	Nupl2
1456774_at	0.92475	"protein phosphatase 1, regulatory (inhibitor) subunit 13 like"	Ppp1r13l
1456775_at	1.13745	integrator complex subunit 8	Ints8
1456776_at	1.02106	spermatogenesis associated 2	Spata2
1456777_at	1.00006	maltase-glucoamylase	Mgam
1456778_at	1.49751	Neuropilin 2	Nrp2
1456779_a_at	1.05993	RIKEN cDNA 4930414L22 gene	4930414L22Rik
1456780_at	1.34462	Beta-transducin repeat containing protein	Btrc
1456781_at	1.03870	"10 days neonate cortex cDNA, RIKEN full-length enriched library, clone: A830044L07 product:unclassifiable, full insert sequence"	---
1456782_at	1.15031	"transcriptional adaptor 1 (HFI1 homolog, yeast) like"	Tada1l
1456783_at	0.95151	RIKEN cDNA 9330107J05 gene	9330107J05Rik
1456784_at	1.06162	hypothetical protein LOC623781	LOC623781
1456785_at	0.85872	"cofactor required for Sp1 transcriptional activation, subunit 2"	Crsp2
1456786_at	1.09510	LIM domain binding 2	Ldb2
1456787_at	0.91310	---	---
1456788_at	0.89178	mitochondrial ribosomal protein L54	Mrpl54
1456789_at	0.90471	Zinc finger protein 462	Zfp462
1456790_at	1.02514	EST AA407452	AA407452
1456791_at	1.04564	EST AA407452	AA407452
1456792_at	1.01768	"TBC1 domain family, member 2"	Tbc1d2
1456793_at	1.08770	cytokine like 1	Cyt1
1456794_at	1.07175	RIKEN cDNA E230016M11 gene	E230016M11Rik
1456795_at	1.06545	RIKEN cDNA D330027G24 gene	D330027G24Rik
1456796_at	0.99599	snail homolog 3 (Drosophila)	Snai3
1456797_at	0.93754	"clathrin, light polypeptide (Lcb)"	Cltb
1456798_at	0.90292	RIKEN cDNA 9330118A15 gene	9330118A15Rik
1456799_at	0.97576	RIKEN cDNA E130110O22 gene	E130110O22Rik
1456800_a_at	0.91201	RIKEN cDNA D130029J02 gene	D130029J02Rik
1456801_at	0.97475	"16 days neonate thymus cDNA, RIKEN full-length enriched library, clone: A130037H12 product:unclassifiable, full insert sequence"	---
1456802_at	1.00075	ring finger protein 133	Rnf133
1456803_at	1.26173	Polymerase (RNA) III (DNA directed) polypeptide C	Polr3c
1456804_at	0.96720	hypothetical protein LOC627821	Av381130

1456805_at 1.04012 RIKEN cDNA C030010B13 gene C030010B13Rik
 1456806_at 1.57906 RIKEN cDNA A130010C12 gene A130010C12Rik
 1456807_at 0.92916 "protein phosphatase 1, regulatory (inhibitor) subunit 3E" Ppp1r3e
 1456808_at 2.58224 RIKEN cDNA 4933426M11 gene 4933426M11Rik
 1456809_at 0.97857 "Adult male thymus cDNA, RIKEN full-length enriched library,
 clone:5830435O18 product:unclassifiable, full insert sequence" ---
 1456810_at 2.08128 vacuolar protein sorting 54 (yeast) Vps54
 1456811_at 0.68854 CXXC finger 4 Cxxc4
 1456812_at 0.92084 "ATP-binding cassette, sub-family D (ALD), member 2" Abcd2
 1456813_at 1.00777 "Adult male hypothalamus cDNA, RIKEN full-length enriched library, clone:
 A230069E07 product:unclassifiable, full insert sequence" ---
 1456814_at 0.94076 RIKEN cDNA 1110014J01 gene 1110014J01Rik
 1456815_at 0.93895 forkhead box N1 Foxn1
 1456816_at 1.27692 Retinoic acid induced 17 Rai17
 1456817_at 0.98614 "Adult retina cDNA, RIKEN full-length enriched library, clone:A930002A02
 product:unclassifiable, full insert sequence" ---
 1456818_at 0.96407 serine/threonine kinase 32A Stk32a
 1456819_at 1.07100 RIKEN cDNA G630049C14 gene G630049C14Rik
 1456820_at 1.01864 "9 days embryo whole body cDNA, RIKEN full-length enriched library, clone:
 D030040D24 product:unclassifiable, full insert sequence" ---
 1456821_at 1.23779 RIKEN cDNA 4932439K10 gene 4932439K10Rik
 1456822_at 1.56901 RAD23b homolog (S. cerevisiae) Rad23b
 1456823_at 0.89916 "gene model 70, (NCBI)" Gm70
 1456824_at 0.95976 zinc finger protein 612 Zfp612
 1456825_at 0.92562 WD repeat domain 33 Wdr33
 1456826_at 0.98622 TAO kinase 2 Taok2
 1456827_at 0.82716 expressed sequence AA987161 AA987161
 1456828_at 1.05816 RIKEN cDNA C030023E24 gene C030023E24Rik
 1456829_at 0.83044 paraneoplastic antigen MA3 Pnma3
 1456830_at 1.20203 "protein phosphatase 1, regulatory (inhibitor) subunit 2" Ppp1r2
 1456831_at 0.99847 AT rich interactive domain 5B (Mrf1 like) Arid5b
 1456832_at 1.47498 Alpha thalassemia/mental retardation syndrome X-linked homolog (human)
 Atrx
 1456833_at 0.96649 G protein-coupled receptor 17 Gpr17
 1456834_at 1.00863 IBR domain containing 2 Ibrdc2
 1456835_at 0.90992 Hypothetical protein LOC624549 LOC624549
 1456836_at 1.03921 IL2-inducible T-cell kinase Itk
 1456837_at 0.85899 RIKEN cDNA A830007L07 gene A830007L07Rik
 1456838_at 0.91297 cDNA sequence BC072620 BC072620
 1456839_at 0.96665 GRAM domain containing 1B Gramd1b
 1456840_at 1.09193 --- ---
 1456841_at 1.12690 "PREDICTED: Mus musculus similar to Kinesin-like protein KIF2C (Mitotic
 centromere-associated kinesin) (MCAK) (LOC242914), mRNA" ---

1456842_at 0.94362 "PREDICTED: Mus musculus hypothetical protein LOC628013 (LOC628013), mRNA" ---

1456843_at 1.31473 Yamaguchi sarcoma viral (v-yes) oncogene homolog 1 Yes1

1456844_at 1.07228 "calcium/calmodulin-dependent protein kinase II, delta" Camk2d

1456845_at 0.90510 Oxidase assembly 1-like Oxa1l

1456846_at 0.92800 similar to RP58 protein LOC382639

1456847_at 1.00834 CD160 antigen Cd160

1456848_at 1.13689 --- ---

1456849_at 1.44014 --- ---

1456850_at 0.93719 --- ---

1456851_at 0.98117 Forkhead box P1 Foxp1

1456852_at 1.02064 polyadenylate binding protein-interacting protein 1 Paip1

1456853_at 0.91674 Inhibitor of four 1 (Mif1) ---

1456854_at 1.45652 neuralized-like homolog (Drosophila) Neurl

1456855_at 1.03537 --- ---

1456856_at 0.97486 "protein tyrosine phosphatase, receptor type, f polypeptide (PTPRF), interacting protein (liprin), alpha 2" Ppfia2

1456857_at 1.19719 RIKEN cDNA 1500011B03 gene 1500011B03Rik

1456858_at 0.95078 G protein-coupled receptor 149 Gpr149

1456859_at 0.99951 limb and neural patterns Lnp

1456860_at 0.97784 cell division cycle 26 Cdc26

1456861_at 1.02787 "Activated spleen cDNA, RIKEN full-length enriched library, clone: F830206P12 product:hypothetical Cytochrome c heme-binding site containing protein, full insert sequence" ---

1456862_at 1.19384 sine oculis-related homeobox 4 homolog (Drosophila) Six4

1456863_at 1.09457 Eph receptor A4 Epha4

1456864_at 1.86939 --- ---

1456865_x_at 0.95191 RRS1 ribosome biogenesis regulator homolog (S. cerevisiae) Rrs1

1456866_x_at 1.04991 RIKEN cDNA 1700027D21 gene 1700027D21Rik

1456867_x_at 0.92627 ERGIC and golgi 3 Ergic3

1456868_at 1.29440 RIKEN cDNA 2900073G15 gene 2900073G15Rik

1456869_at 0.91409 RIKEN cDNA 2210018M03 gene 2210018M03Rik

1456870_at 0.94191 RIKEN cDNA A430107D22 gene A430107D22Rik

1456871_a_at 1.03067 PHD finger protein 20-like 1 Phf20l1

1456872_a_at 0.91880 RIKEN cDNA D230010M03 gene D230010M03Rik

1456873_at 0.88477 chloride intracellular channel 5 Clic5

1456874_at 0.91719 fibronectin leucine rich transmembrane protein 2 Flrt2

1456875_at 1.00671 RIKEN cDNA 1810013L24 gene 1810013L24Rik

1456876_at 0.96986 --- ---

1456877_at 0.80889 dual oxidase 1 /// similar to dual oxidase 1 Duox1 /// LOC640578

1456878_at 0.62619 expressed sequence AI646023 AI646023

1456879_at 1.01119 RIKEN cDNA C130022K22 gene C130022K22Rik

1456880_at 1.56481 "Splicing factor, arginine/serine-rich 8" Sfrs8

1456881_at 0.85335 neurobeachin-like 2 Nbeal2
1456882_at 0.95638 SRY-box containing gene 12 Sox12
1456883_at 0.96035 storkhead box 1 Stox1
1456884_at 0.86267 Hypothetical LOC67660 LOC67660
1456885_at 0.98741 Transcribed locus ---
1456886_at 1.03521 Brain protein 44-like Brp44l
1456887_at 0.93021 chemokine-like receptor 1 Cmklr1
1456888_at 1.27658 "6-phosphofructo-2-kinase/fructose-2,6-biphosphatase 4" Pfkfb4
1456889_at 0.97309 "PREDICTED: Mus musculus similar to aspartoacylase-3 (LOC629627), mRNA" ---
1456890_at 1.03840 DEAD (Asp-Glu-Ala-Asp) box polypeptide 58 Ddx58
1456891_at 1.12365 DENN/MADD domain containing 2C Dennd2c
1456892_at 0.95485 RIKEN cDNA D030036P13 gene D030036P13Rik
1456893_at 0.88011 "Adult male liver tumor cDNA, RIKEN full-length enriched library, clone: C730013D23 product:unclassifiable, full insert sequence" ---
1456894_at 1.09356 RIKEN cDNA C130039O16 gene C130039O16Rik
1456895_at 1.01874 CD209b antigen Cd209b
1456896_at 1.18490 RIKEN cDNA 6720462K09 gene 6720462K09Rik
1456897_at 1.17168 Stam binding protein like 1 Stambpl1
1456898_at 1.00261 purine rich element binding protein A Pura
1456899_at 0.96430 RIKEN cDNA 1500004F05 gene 1500004F05Rik
1456900_at 0.95108 --- ---
1456901_at 0.99016 "a disintegrin-like and metallopeptidase (reprolysin type) with thrombospondin type 1 motif, 20" Adamts20
1456902_at 0.93131 Special AT-rich sequence binding protein 1 Satb1
1456903_at 1.00212 pentraxin related gene Ptx3
1456904_at 1.06751 Transcribed locus ---
1456905_at 0.91842 coiled-coil domain containing 24 Ccdc24
1456906_at 1.51346 RIKEN cDNA 4833432P19 gene 4833432P19Rik
1456907_at 0.97101 chemokine (C-X-C motif) ligand 9 Cxcl9
1456908_at 1.08459 cDNA sequence BC023202 BC023202
1456909_at 1.93693 similar to Glucose-6-phosphate isomerase (GPI) (Phosphoglucose isomerase) (PGI) (Phosphohexose isomerase) (PHI) (Neuroleukin) (NLK) LOC676974
1456910_at 0.95523 Transcribed locus ---
1456911_at 0.84840 CLIP associating protein 2 Clasp2
1456912_at 0.92507 zinc finger protein 691 Zfp691
1456913_at 1.52236 Tropomodulin 3 Tmod3
1456914_at 1.35113 --- ---
1456915_at 1.02264 RIKEN cDNA D930005D10 gene D930005D10Rik
1456916_at 1.18718 Nuclear receptor-binding SET-domain protein 1 Nsd1
1456917_at 1.48453 ADP-ribosylation factor guanine nucleotide-exchange factor 1(brefeldin A-inhibited) Arfgef1
1456918_at 2.57374 Ras association (RalGDS/AF-6) and pleckstrin homology domains 1 Raph1

1456919_at 1.09808 RIKEN cDNA A930013B10 gene A930013B10Rik
1456920_at 1.21398 --- ---
1456921_at 1.01014 Ataxin 2 binding protein 1 A2bp1
1456922_at 1.07382 RIKEN cDNA 4933437K13 gene 4933437K13Rik
1456923_at 0.97717 "transient receptor potential cation channel, subfamily M, member 3" Trpm3
1456924_at 1.10143 expressed sequence AW492981 AW492981
1456925_at 0.82356 --- ---
1456926_at 0.87139 regulating synaptic membrane exocytosis 4 Rims4
1456927_at 1.78582 microtubule associated serine/threonine kinase 2 Mast2
1456928_at 1.04268 --- ---
1456929_at 0.98165 CDNA sequence BC042782 BC042782
1456930_at 0.86387 calmodulin regulated spectrin-associated protein 1 Camsap1
1456931_at 0.94919 expressed sequence C85363 C85363
1456932_at 0.97943 Hypothetical protein B430201F14 B430201F14
1456933_at 1.12941 "Smg-7 homolog, nonsense mediated mRNA decay factor (C. elegans)"
Smg7
1456934_at 1.07872 --- ---
1456935_at 0.89636 expressed sequence AI606473 AI606473
1456936_at 1.08359 calcium binding protein 4 Cabp4
1456937_at 1.06106 cadherin-like 26 Cdh26
1456938_at 1.01087 "SWI/SNF related, matrix associated, actin dependent regulator of chromatin,
subfamily b, member 1" Smarcb1
1456939_at 1.00737 RIKEN cDNA 1700129I04 gene 1700129I04Rik
1456940_at 0.96968 "solute carrier family 43, member 2" Slc43a2
1456941_at 0.99304 telomerase reverse transcriptase Tert
1456942_x_at 1.14012 hypothetical protein LOC668339 LOC668339
1456943_a_at 1.00913 "Adult male corpora quadrigemina cDNA, RIKEN full-length enriched library,
clone:B230215D24 product:unclassifiable, full insert sequence" ---
1456944_at 1.13135 "potassium voltage gated channel, Shaw-related subfamily, member 1" Kcnc1
1456945_at 0.98693 nudix (nucleoside diphosphate linked moiety X)-type motif 6 Nudt6
1456946_at 0.92383 RIKEN cDNA 4831416G18 gene 4831416G18Rik
1456947_at 1.41388 "Platelet-activating factor acetylhydrolase, isoform 1b, beta1 subunit" Pafah1b1
1456948_at 1.11225 "adaptor-related protein complex AP-4, epsilon 1" Ap4e1
1456949_at 0.97843 FCH and double SH3 domains 2 Fchsd2
1456950_at 0.78232 Alstrom syndrome 1 homolog (human) Alms1
1456951_at 1.04332 "Adult male testis cDNA, RIKEN full-length enriched library,
clone:4921513I01 product:unclassifiable, full insert sequence" ---
1456952_at 1.74396 AT rich interactive domain 1B (Swi1 like) Arid1b
1456953_at 1.01040 "12 days embryo spinal ganglion cDNA, RIKEN full-length enriched library,
clone:D130062H10 product:unclassifiable, full insert sequence" ---
1456954_at 1.11443 "potassium voltage-gated channel, shaker-related, subfamily, member 6" Kcna6
1456955_at 1.63058 --- ---
1456956_at 2.27848 "zinc finger protein, subfamily 1A, 2 (Helios)" Zfpn1a2

1456957_at 1.09240 --- ---
1456958_at 1.04116 RIKEN cDNA C230072F16 gene C230072F16Rik
1456959_at 0.92063 RIKEN cDNA 1700034H15 gene 1700034H15Rik
1456960_at 3.16749 Adenosine kinase Adk
1456961_at 0.85251 enhancer of yellow 2 homolog (Drosophila) Eny2
1456962_at 1.18817 contactin 2 Cntn2
1456963_at 0.96175 RIKEN cDNA 1110039B18 gene 1110039B18Rik
1456964_at 0.99418 RNA binding motif protein 12 Rbm12
1456965_at 0.91103 --- ---
1456966_at 0.97172 CDNA sequence BC028528 BC028528
1456967_at 0.95690 tripartite motif-containing 66 Trim66
1456968_at 1.04917 Actinin alpha 2 Actn2
1456969_at 1.01216 RIKEN cDNA 9530092O11 gene 9530092O11Rik
1456970_at 0.80257 "Cadherin 7, type 2" Cdh7
1456971_at 1.01155 "solute carrier family 1 (glutamate transporter), member 7" Slc1a7
1456972_at 1.10174 expressed sequence C80256 C80256
1456973_at 1.41332 AT rich interactive domain 5B (Mrf1 like) Arid5b
1456974_at 0.99113 "one cut domain, family member 1" Onecut1
1456975_at 1.49408 TAO kinase 1 Taok1
1456976_at 1.03925 wingless-related MMTV integration site 5A Wnt5a
1456977_at 0.91773 --- ---
1456978_s_at 0.93208 "DNA segment, Chr 19, ERATO Doi 652, expressed" D19Ertd652e
1456979_at 0.98595 Zinc fingers and homeoboxes 3 Zhx3
1456980_at 1.01009 RIKEN cDNA 9830134C10 gene 9830134C10Rik
1456981_at 0.79363 transmembrane channel-like gene family 7 Tmc7
1456982_at 0.84339 RIKEN cDNA 6430562O15 gene 6430562O15Rik
1456983_at 2.32714 Trichorhinophalangeal syndrome I (human) Trps1
1456984_at 0.95893 sex comb on midleg-like 2 (Drosophila) Scml2
1456985_at 1.32789 "Deoxynucleotidyltransferase, terminal, interacting protein 1" Dnttip1
1456986_at 1.05291 Zinc finger and BTB domain containing 16 Zbtb16
1456987_at 1.07825 RIKEN cDNA 4932418E24 gene 4932418E24Rik
1456988_at 1.00086 expressed sequence AI848258 AI848258
1456989_at 1.01332 oxoglutarate (alpha-ketoglutarate) receptor 1 Oxgr1
1456990_at 0.95187 "0 day neonate thymus cDNA, RIKEN full-length enriched library, clone:
A430015G07 product:unclassifiable, full insert sequence" ---
1456991_at 2.72829 Cobl-like 1 Cobll1
1456992_at 1.33400 ubiquitin protein ligase E3C Ube3c
1456993_at 1.30544 "DNA segment, Chr 2, ERATO Doi 640, expressed" D2Ertd640e
1456994_at 0.96741 Ras association (RalGDS/AF-6) domain family 1 Rassf1
1456995_at 1.01467 RIKEN cDNA 2700045P11 gene 2700045P11Rik
1456996_at 1.10087 RIKEN cDNA 1190005F20 gene 1190005F20Rik
1456997_at 1.14002 Nuclear receptor coactivator 5 Ncoa5
1456998_at 1.00271 "7 days neonate cerebellum cDNA, RIKEN full-length enriched library, clone:

A730081K04 product:unclassifiable, full insert sequence" ---

1456999_at 1.07472 "WD repeat domain, phosphoinositide interacting 2" Wipi2

1457000_at 0.96547 cDNA sequence BC060267 BC060267

1457001_at 1.02714 centromere protein K Cenpk

1457002_at 1.03929 zinc finger protein 408 Zfp408

1457003_at 0.94002 expressed sequence AW490526 AW490526

1457004_at 0.91736 "DNA segment, Chr 15, Wayne State University 169, expressed" D15Wsu169e

1457005_at 1.03682 mitochondrial methionyl-tRNA formyltransferase Mtfmt

1457006_at 1.03731 RIKEN cDNA 1700106J12 gene 1700106J12Rik

1457007_at 1.58068 Nuclear receptor co-repressor 1 Ncor1

1457008_at 0.94151 "cholinergic receptor, nicotinic, beta polypeptide 4" Chrn4

1457009_at 0.95543 Rho-related BTB domain containing 3 Rhobtb3

1457010_at 1.13802 RIKEN cDNA 3010001F23 gene 3010001F23Rik

1457011_at 1.02095 Transcribed locus ---

1457012_at 1.02043 developing brain homeobox 1 Dbx1

1457013_at 0.93293 GTPase activating RANGAP domain-like 1 Garnl1

1457014_x_at 0.93891 "Solute carrier family 16 (monocarboxylic acid transporters), member 8" Slc16a8

1457015_at 1.08591 RIKEN cDNA A230070E04 gene A230070E04Rik

1457016_at 0.86362 Transcribed locus ---

1457017_at 0.98192 --- ---

1457018_at 0.97231 "Potassium large conductance calcium-activated channel, subfamily M, alpha member 1" Kcnma1

1457019_s_at 1.01721 retinol dehydrogenase 14 (all-trans and 9-cis) Rdh14

1457020_at 1.40530 GC-rich promoter binding protein 1-like 1 Gpbp111

1457021_x_at 0.99278 anti-Mullerian hormone type 2 receptor Amhr2

1457022_at 1.02579 "Dystrophin, muscular dystrophy" Dmd

1457023_at 0.90010 RIKEN cDNA 5830411G16 gene 5830411G16Rik

1457024_x_at 0.86118 "solute carrier family 35 (UDP-galactose transporter), member 2" Slc35a2

1457025_at 0.93522 RIKEN cDNA 4833413O15 gene 4833413O15Rik

1457026_at 1.01096 "lipase, member H" Liph

1457027_at 0.95492 dehydrogenase E1 and transketolase domain containing 1 Dhktd1

1457028_at 0.72950 RIKEN cDNA A430033K04 gene A430033K04Rik

1457029_at 0.99494 RIKEN cDNA C030010B13 gene C030010B13Rik

1457030_at 1.06037 miRNA containing gene Mirg

1457031_at 0.98223 fibronectin type III and SPRY domain containing 2 Fsd2

1457032_at 0.95947 adenylate kinase 5 Ak5

1457033_at 0.78214 "gene model 397, (NCBI) /// Mus musculus sequence XM_486399 /// similar to zinc finger and SCAN domain containing 4 /// similar to zinc finger and SCAN domain containing 4 /// similar to zinc finger and SCAN domain containing 4" Gm397 /// XM_486399 /// LOC545913 /// LOC632758 /// LOC665902 /// LOC665913

1457034_at 0.96141 "DNA segment, Chr 14, Abbott 1 expressed" D14Abble

1457035_at 0.98021 expressed sequence AI607873 AI607873

1457036_at 1.74042 similar to upstream transcription factor 2 LOC639063
1457037_at 0.97983 "gene model 1103, (NCBI)" Gm1103
1457038_at 0.94687 Fras1 related extracellular matrix protein 2 Frem2
1457039_at 1.01402 "cat eye syndrome chromosome region, candidate 2 homolog (human)"
Cecr2
1457040_at 0.92144 "leucine-rich repeat LGI family, member 2" Lgi2
1457041_at 1.13184 RIKEN cDNA 1700026B20 gene 1700026B20Rik
1457042_at 0.96485 EST AI256396 AI256396
1457043_at 0.97305 "beta 1,3-galactosyltransferase-like" B3galt1
1457044_at 1.02662 RIKEN cDNA 4732474O15 gene 4732474O15Rik
1457045_at 1.11795 UDP-N-acetyl-alpha-D-galactosamine:polypeptide N-
acetylgalactosaminyltransferase 13 Galnt13
1457046_s_at 0.95925 expressed sequence C77370 C77370
1457047_at 1.13619 U1 small nuclear ribonucleoprotein polypeptide A Snrp70
1457048_at 0.92789 G protein-coupled receptor 103 Gpr103
1457049_at 0.96630 "VATPase, H+ transporting, lysosomal V1 subunit E1" Atp6v1e1
1457050_at 1.02047 RIKEN cDNA 1700129O19 gene 1700129O19Rik
1457051_at 0.96543 tripartite motif protein 27 Trim27
1457052_at 0.94401 expressed sequence AW536275 AW536275
1457053_at 1.33599 CDNA sequence BC016423 BC016423
1457054_a_at 0.86141 RIKEN cDNA D130020L05 gene D130020L05Rik
1457055_at 1.05311 Nucleotide binding protein 1 Nubp1
1457056_at 0.90431 Formiminotransferase cyclodeaminase Ftcd
1457057_at 0.96456 RIKEN cDNA 1700042G07 gene 1700042G07Rik
1457058_at 0.53329 "a disintegrin-like and metallopeptidase (reprolysin type) with thrombospondin
type 1 motif, 2" Adamts2
1457059_at 1.04182 zinc finger protein 438 Zfp438
1457060_at 0.88615 ankyrin repeat and sterile alpha motif domain containing 1B /// similar to AIDA-
1b Anks1b /// LOC544718
1457061_at 0.96970 glucocorticoid induced transcript 1 Glcci1
1457062_at 1.64401 RIKEN cDNA 1700081L11 gene 1700081L11Rik
1457063_at 1.03240 RIKEN cDNA 9530033F24 gene 9530033F24Rik
1457064_at 1.71280 RIKEN cDNA D030002E05 gene D030002E05Rik
1457065_at 1.38295 uroplakin 1B Upk1b
1457066_at 0.94776 "ATP-binding cassette, sub-family C (CFTR/MRP), member 8" Abcc8
1457067_at 0.98816 Vacuolar protein sorting 4b (yeast) Vps4b
1457068_at 0.93269 N-acetylated alpha-linked acidic dipeptidase-like 1 Naalad11
1457069_at 0.99498 activating signal cointegrator 1 complex subunit 3 Ascc3
1457070_at 0.95607 "Ubiquitin-like, containing PHD and RING finger domains 2" Uhrf2
1457071_x_at 1.03560 protein arginine N-methyltransferase 6 Prmt6
1457072_at 0.96900 RIKEN cDNA 2410127E16 gene /// B-cell CLL/lymphoma 11A (zinc finger
protein) 2410127E16Rik /// Bcl11a
1457073_at 1.38343 O-linked N-acetylglucosamine (GlcNAc) transferase (UDP-N-

acetylglucosamine:polypeptide-N-acetylglucosaminyl transferase) Ogt
1457074_at 1.24174 kinesin family member 24 Kif24
1457075_at 0.99666 RIKEN cDNA A830031M15 gene A830031M15Rik
1457076_at 0.89677 "gene model 693, (NCBI) /// hypothetical protein LOC670665" Gm693 ///
LOC670665
1457077_at 1.01754 "2 days neonate sympathetic ganglion cDNA, RIKEN full-length enriched
library, clone:7120457A19 product:unclassifiable, full insert sequence" ---
1457078_at 1.08756 zinc finger protein 473 Zfp473
1457079_at 1.08697 CDNA clone IMAGE:30014192 ---
1457080_at 1.01014 Transcribed locus ---
1457081_at 1.73255 RIKEN cDNA D630038D15 gene D630038D15Rik
1457082_at 0.91191 forkhead box P3 Foxp3
1457083_at 0.94851 PRP31 pre-mRNA processing factor 31 homolog (yeast) Prpf31
1457084_at 1.14950 "NACHT, LRR and PYD containing protein 9b" Nalp9b
1457085_at 1.06088 hypothetical protein LOC665934 /// hypothetical protein LOC669392 ///
hypothetical protein LOC676468 LOC665934 /// LOC669392 /// LOC676468
1457086_at 0.93263 hypothetical LOC434147 LOC434147
1457087_at 0.75545 RIKEN cDNA 2900083I11 gene 2900083I11Rik
1457088_at 1.09698 pallidin Pldn
1457089_at 1.41474 --- ---
1457090_at 1.06590 "0 day neonate head cDNA, RIKEN full-length enriched library,
clone:4831433L12 product:unclassifiable, full insert sequence" ---
1457091_at 0.91796 RIKEN cDNA 4930526D03 gene 4930526D03Rik
1457092_at 0.86455 RIKEN cDNA C630007B19 gene C630007B19Rik
1457093_at 1.16097 GTPase activating protein and VPS9 domains 1 Gapvd1
1457094_at 1.12031 src homology 2 domain-containing transforming protein E She
1457095_at 0.86148 "Pleckstrin homology, Sec7 and coiled-coil domains 1" Pscd1
1457096_at 0.96554 RIKEN cDNA 6430520M22 gene 6430520M22Rik
1457097_at 0.81460 src family associated phosphoprotein 2 Scap2
1457098_at 0.91561 "DNA segment, Chr 2, ERATO Doi 624, expressed" D2Ertd624e
1457099_at 0.93078 RIKEN cDNA D030011O10 gene D030011O10Rik
1457100_at 1.10146 expressed sequence AW552889 AW552889
1457101_at 0.90285 "0 day neonate thymus cDNA, RIKEN full-length enriched library, clone:
A430035F24 product:unclassifiable, full insert sequence" ---
1457102_at 0.94162 RIKEN cDNA A030001D16 gene A030001D16Rik
1457103_at 0.98721 expressed sequence AU046084 AU046084
1457104_at 1.02778 RIKEN cDNA D930036F22 gene D930036F22Rik
1457105_at 0.89906 polycystic kidney disease 2-like 1 Pkd2l1
1457106_at 0.97929 thyroid adenoma associated Thada
1457107_at 1.87285 N-myristoyltransferase 2 Nmt2
1457108_at 1.07713 "Interleukin 1 receptor, type II" Il1r2
1457109_x_at 0.98745 Transcribed locus ---
1457110_at 1.03800 pantothenate kinase 1 Pank1

1457111_at 2.01043 expressed sequence AA415038 AA415038
1457112_at 0.89932 hypothetical protein A230106N23 A230106N23
1457113_at 2.27984 "nuclear autoantigenic sperm protein (histone-binding) /// DNA segment, Chr 4, ERATO Doi 767, expressed" Nasp /// D4ErtD767e
1457114_at 0.80214 "torsin family 1, member B" Tor1b
1457115_at 1.16403 "Adult male hippocampus cDNA, RIKEN full-length enriched library, clone: C630007G15 product:hypothetical protein, full insert sequence" ---
1457116_at 1.18622 "REX1, RNA exonuclease 1 homolog (S. cerevisiae)" Rexo1
1457117_at 1.05038 "nuclear factor, erythroid derived 2, like 2" Nfe2l2
1457118_at 0.86602 "SHC (Src homology 2 domain containing) family, member 4" Shc4
1457119_at 0.90183 --- ---
1457120_at 1.05625 IL2-inducible T-cell kinase Itk
1457121_at 0.85062 expressed sequence AW822216 AW822216
1457122_at 1.07500 "Adaptor protein complex AP-2, mu1" Ap2m1
1457123_at 1.04617 neuregulin 4 Nrg4
1457124_at 1.06324 "Adult male corpora quadrigemina cDNA, RIKEN full-length enriched library, clone:B230352I09 product:unclassifiable, full insert sequence" ---
1457125_at 1.02640 "Adult male testis cDNA, RIKEN full-length enriched library, clone:4930515K13 product:unclassifiable, full insert sequence" ---
1457126_at 0.92077 "myosin, light polypeptide 4" Myl4
1457127_at 1.12362 defensin beta 42 Defb42
1457128_at 1.04059 --- ---
1457129_at 1.06539 Zinc finger protein 458 Zfp458
1457130_at 0.95786 RIKEN cDNA 2900035I09 gene 2900035I09Rik
1457131_at 0.92737 furry homolog (Drosophila) Fry
1457132_at 0.96833 Transcribed locus ---
1457133_at 0.95264 Transcribed locus ---
1457134_at 1.00476 cytoplasmic FMR1 interacting protein 1 Cyfip1
1457135_at 1.23983 fat tumor suppressor homolog 2 (Drosophila) Fath2
1457136_at 0.98429 "splicing factor, arginine/serine-rich 1 (ASF/SF2)" Sfrs1
1457137_at 0.96755 Transcribed locus ---
1457138_x_at 1.04420 Opticin Optc
1457139_at 0.92230 autism susceptibility candidate 2 Aut2
1457140_s_at 1.06896 RIKEN cDNA 4632411J06 gene 4632411J06Rik
1457141_at 1.01187 Aquaporin 11 Aqp11
1457142_at 0.93137 EF hand calcium binding protein 1 Efcfbp1
1457143_at 0.83995 "PREDICTED: Mus musculus hypothetical protein LOC625121 (LOC625121), mRNA" ---
1457144_at 0.96048 "sodium channel, voltage-gated, type II, beta" Scn2b
1457145_at 1.35950 "pleckstrin homology domain containing, family G (with RhoGef domain) member 4" Plekhg4
1457146_at 0.91885 dedicator of cytokinesis 4 Dock4
1457147_at 1.39331 cDNA sequence BC026657 BC026657

1457148_at 0.87811 CUB and Sushi multiple domains 2 Csm2
1457149_at 0.94821 tetratricopeptide repeat domain 22 Ttc22
1457150_at 1.20360 expressed sequence AI428301 AI428301
1457151_at 0.97262 "Mus musculus, clone IMAGE:1527712, mRNA" ---
1457152_at 1.00087 --- ---
1457153_at 0.95699 hypothetical LOC552882 LOC552882
1457154_at 1.26927 Membrane-associated ring finger (C3HC4) 6 6-Mar
1457155_at 1.01666 "aldehyde dehydrogenase 2, mitochondrial" Aldh2
1457156_at 0.96684 TRH-degrading enzyme Trhde
1457157_at 1.00720 "phospholipase C, eta 1" Plch1
1457158_at 1.11641 RIKEN cDNA 0610039J04 gene 0610039J04Rik
1457159_at 0.89679 "gene model 101, (NCBI)" Gm101
1457160_at 0.82949 four jointed box 1 (Drosophila) Fjx1
1457161_at 2.17954 RIKEN cDNA 9530029O12 gene 9530029O12Rik
1457162_at 1.09752 low density lipoprotein receptor adaptor protein 1 Ldlrap1
1457163_at 1.03762 RIKEN cDNA D730035F11 gene D730035F11Rik
1457164_at 1.04246 "transient receptor potential cation channel, subfamily A, member 1" Trpa1
1457165_at 1.02210 --- ---
1457166_at 1.11158 Expressed sequence AA536749 AA536749
1457167_at 0.98976 "cofactor required for Sp1 transcriptional activation, subunit 2" Crsp2
1457168_at 1.24709 expressed sequence AI225934 AI225934
1457169_at 1.13560 "Protein phosphatase 1, regulatory (inhibitor) subunit 12B" Ppp1r12b
1457170_at 0.87909 leucine rich repeat containing 36 Lrrc36
1457171_at 1.05047 --- ---
1457172_at 1.07668 Rho guanine nucleotide exchange factor (GEF) 15 Arhgef15
1457173_at 1.64315 "Tyrosine 3-monooxygenase/tryptophan 5-monooxygenase activation protein, epsilon polypeptide" Ywhae
1457174_at 1.28822 expressed sequence AU015680 AU015680
1457175_at 1.36234 Numb gene homolog (Drosophila) Numb
1457176_at 1.04534 GTPase activating RANGAP domain-like 1 Garnl1
1457177_at 1.08809 RAR-related orphan receptor alpha Rora
1457178_at 1.16153 RAB GTPase activating protein 1-like Rabgap11
1457179_at 0.98204 "RIKEN cDNA 4930406H16 gene /// NACHT, leucine rich repeat and PYD containing 4E" 4930406H16Rik /// Nalp4e
1457180_at 0.96240 DIP2 disco-interacting protein 2 homolog C (Drosophila) Dip2c
1457181_at 1.76171 "DNA segment, Chr 15, Wayne State University 75, expressed" D15Wsu75e
1457182_at 0.90009 mitogen activated protein kinase kinase 7 Map2k7
1457183_at 1.13349 "solute carrier family 6 (neurotransmitter transporter, GABA), member 1" Slc6a1
1457184_at 1.17556 FERM domain containing 6 Frmd6
1457185_at 0.96040 --- ---
1457186_at 1.11765 transducin (beta)-like 1X-linked receptor 1 Tbl1xr1
1457187_at 0.96981 RIKEN cDNA A630071L07 gene A630071L07Rik

1457188_at 1.18531 Rho guanine nucleotide exchange factor (GEF) 11 Arhgef11
1457189_at 1.43205 "Inositol 1,4,5-triphosphate receptor 1" Itpr1
1457190_at 0.98680 "12 days embryo eyeball cDNA, RIKEN full-length enriched library, clone:
D230005M19 product:unclassifiable, full insert sequence" ---
1457191_at 1.86037 --- ---
1457192_at 0.88001 --- ---
1457193_at 1.00737 myeloid/lymphoid or mixed-lineage leukemia 3 Mll3
1457194_at 1.18416 --- ---
1457195_at 1.12812 "pleckstrin homology domain containing, family M (with RUN domain)
member 1" Plekhm1
1457196_at 0.90973 Transcribed locus ---
1457197_at 1.01619 "protein kinase, cAMP dependent, catalytic, beta" Prkacb
1457198_at 0.85372 neuropilin 1 Nrp1
1457199_at 1.14954 --- ---
1457200_at 1.01413 "serine (or cysteine) peptidase inhibitor, clade B, member 9d" Serpinb9d
1457201_at 0.99073 Arsenic (+3 oxidation state) methyltransferase As3mt
1457202_at 1.09778 --- ---
1457203_at 1.08229 Transcribed locus ---
1457204_at 0.97288 "Zinc finger, DHHC domain containing 2" Zdhhc2
1457205_at 1.04584 N-ethylmaleimide sensitive fusion protein Nsf
1457206_at 0.96754 Zinc finger protein 521 Zfp521
1457207_at 0.97351 Signaling molecule ATTP Attp
1457208_at 0.96181 "nuclear transcription factor, X-box binding-like 1" Nfxl1
1457209_at 1.02298 DDHD domain containing 2 Ddhd2
1457210_at 0.97707 RIKEN cDNA 1700092E19 gene 1700092E19Rik
1457211_at 0.99450 Glutaminase Gls
1457212_at 0.91896 Transcribed locus ---
1457213_a_at 1.00446 "diacylglycerol kinase, eta" Dgkh
1457214_at 1.28227 Nuclear receptor coactivator 3 Ncoa3
1457215_at 0.92044 "Adult male hippocampus cDNA, RIKEN full-length enriched library, clone:
C630028A20 product:unclassifiable, full insert sequence" ---
1457216_at 1.30650 RING CCCH (C3H) domains 1 Rc3h1
1457217_at 0.94696 Paralemmin Palm
1457218_at 1.17296 RIKEN cDNA 6430510M02 gene 6430510M02Rik
1457219_at 1.00030 rho/rac guanine nucleotide exchange factor (GEF) 18 Arhgef18
1457220_at 0.90917 --- ---
1457221_at 0.96681 "Immunoglobulin superfamily, member 11" Igsf11
1457222_at 1.08498 cAMP responsive element binding protein 5 Creb5
1457223_at 1.04583 ELMO domain containing 1 Elmod1
1457224_at 0.96861 RIKEN cDNA 4631422O05 gene 4631422O05Rik
1457225_at 0.78021 olfactory receptor 288 Olfr288
1457226_at 0.93810 --- ---
1457227_at 0.93415 expressed sequence AI843755 AI843755

1457228_x_at 0.96068 GLE1 RNA export mediator-like (yeast Gle11
1457229_at 1.03575 G-protein coupled receptor 173 Gpr173
1457230_at 1.07669 glucosamine-6-phosphate deaminase 2 Gnpda2
1457231_at 1.87432 "Hypoxia inducible factor 1, alpha subunit" Hif1a
1457232_at 0.84139 F-box and leucine-rich repeat protein 21 Fbxl21
1457233_at 1.21644 "DnaJ (Hsp40) homolog, subfamily A, member 2" Dnaja2
1457234_at 1.01519 Transcribed locus ---
1457235_at 0.94735 low density lipoprotein-related protein 1B (deleted in tumors) Lrp1b
1457236_at 1.09353 G protein-coupled receptor 62 Gpr62
1457237_at 0.94094 muscleblind-like 2 Mbnl2
1457238_at 1.02315 Transcribed locus ---
1457239_at 0.93834 Spleen tyrosine kinase Syk
1457240_at 1.12128 cyclin-dependent kinase 9 (CDC2-related kinase) Cdk9
1457241_at 0.86890 Leucine rich repeat containing 48 Lrrc48
1457242_at 0.91804 U box domain containing 5 Ubox5
1457243_at 1.04734 RIKEN cDNA 1110032O16 gene 1110032O16Rik
1457244_at 0.96007 "DNA segment, Chr 6, ERATO Doi 474, expressed" D6Ertd474e
1457245_at 0.99205 disrupted in renal carcinoma 2 (human) Dirc2
1457246_at 1.00391 retinoic acid induced 16 Rai16
1457247_at 1.10950 expressed sequence AI465495 AI465495
1457248_x_at 1.43264 hydroxysteroid (17-beta) dehydrogenase 7 Hsd17b7
1457249_at 1.28462 Ubiquitin associated protein 2-like Ubap2l
1457250_x_at 1.12973 "Nuclear receptor subfamily 1, group D, member 2" Nr1d2
1457251_x_at 1.07737 Adenylate cyclase 1 Adcy1
1457252_x_at 0.92459 phospholipase D2 Pld2
1457253_at 1.11705 tripartite motif-containing 40 Trim40
1457254_x_at 1.08014 RIKEN cDNA 6330442E10 gene 6330442E10Rik
1457255_x_at 1.00963 Kinesin family member 5C Kif5c
1457256_x_at 0.89904 patched homolog 2 /// similar to Protein patched homolog 2 (PTC2)
Ptch2 /// LOC674483
1457257_x_at 0.84507 "0 day neonate lung cDNA, RIKEN full-length enriched library, clone:
E030046E07 product:unclassifiable, full insert sequence" ---
1457258_at 0.92273 "PREDICTED: Mus musculus hypothetical LOC381645 (LOC381645),
mRNA" ---
1457259_at 1.03628 --- ---
1457260_at 0.93026 RIKEN cDNA 7530403E16 gene 7530403E16Rik
1457261_at 0.84318 RIKEN cDNA A930025H08 gene A930025H08Rik
1457262_at 1.09461 RIKEN cDNA 2610207I05 gene 2610207I05Rik
1457263_at 0.92046 "Serine (or cysteine) peptidase inhibitor, clade A, member 3K" Serpina3k
1457264_at 1.67996 PHD finger protein 20-like 1 Phf20l1
1457265_at 0.88512 RIKEN cDNA B230333C21 gene B230333C21Rik
1457266_at 1.18540 expressed sequence AW322671 AW322671
1457267_at 1.02027 --- ---

1457268_at 0.93721 "DOT1-like, histone H3 methyltransferase (*S. cerevisiae*)" Dot11
 1457269_at 1.02314 "gene model 1123, (NCBI) /// similar to coxsackie-adenovirus receptor-like"
 Gm1123 /// LOC674752
 1457270_at 1.07850 RIKEN cDNA B230343A10 gene B230343A10Rik
 1457271_at 0.87253 "gene model 131, (NCBI) /// similar to prochymosin" Gm131 /// LOC634449
 1457272_at 1.04374 Transcribed locus ---
 1457273_at 0.94553 similar to odd Oz/ten-m homolog 2 LOC669490
 1457274_at 1.01000 hypothetical protein E330017N17 E330017N17
 1457275_at 0.81025 desmuslin Dmn
 1457276_at 1.05825 SNF1-like kinase 2 Snf1lk2
 1457277_at 0.87441 cDNA sequence BC038925 BC038925
 1457278_at 0.95528 RIKEN cDNA 6720489N17 gene 6720489N17Rik
 1457279_at 1.01269 "Adult male corpora quadrigemina cDNA, RIKEN full-length enriched library,
 clone:B230347A18 product:unclassifiable, full insert sequence" ---
 1457280_at 1.03080 --- ---
 1457281_at 1.04606 RIKEN cDNA 4930461P20 gene 4930461P20Rik
 1457282_x_at 1.25194 "Tubulin, gamma complex associated protein 5" Tubgcp5
 1457283_at 1.03026 --- ---
 1457284_at 1.20832 CDNA clone IMAGE:5256843 ---
 1457285_at 0.89812 zinc finger protein 187 Zfp187
 1457286_at 0.96777 Histidine triad nucleotide binding protein 1 Hint1
 1457287_at 0.89807 Transcribed locus ---
 1457288_at 1.03596 "ankyrin 3, epithelial" Ank3
 1457289_at 0.92141 "nuclear receptor subfamily 2, group E, member 1" Nr2e1
 1457290_at 1.07854 mab-21-like 1 (*C. elegans*) Mab21l1
 1457291_at 1.07931 "Adult male hypothalamus cDNA, RIKEN full-length enriched library, clone:
 A230080O15 product:unclassifiable, full insert sequence" ---
 1457292_at 1.77525 Josephin domain containing 3 Josd3
 1457293_at 0.78799 Zinc finger and BTB domain containing 4 Zbtb4
 1457294_at 1.27535 Transmembrane and tetratricopeptide repeat containing 3 Tmtc3
 1457295_at 1.06574 trinucleotide repeat containing 6a Tnrc6a
 1457296_at 1.05123 "cartilage intermediate layer protein, nucleotide pyrophosphohydrolase" Cilp
 1457297_at 1.59660 "LysM, putative peptidoglycan-binding, domain containing 4" Lysmd4
 1457298_at 1.01723 --- ---
 1457299_at 0.90984 "glutamate receptor, metabotropic 4" Grm4
 1457300_at 0.92629 aftiphilin Aftph
 1457301_at 1.01623 expressed sequence AI931714 AI931714
 1457302_at 1.38304 "Solute carrier family 20, member 2" Slc20a2
 1457303_at 0.93644 Ankyrin repeat and sterile alpha motif domain containing 3 Anks3
 1457304_at 1.02810 "DNA segment, Chr 13, ERATO Doi 787, expressed" D13Ertd787e
 1457305_at 0.82386 Transcribed locus ---
 1457306_at 0.93777 Transcribed locus ---
 1457307_at 1.09384 RIKEN cDNA A330102K04 gene A330102K04Rik

1457308_at 1.05778 Ubiquitin specific petidase 45 Usp45
1457309_at 0.97510 expressed sequence AU015228 AU015228
1457310_x_at 0.86421 "scratch homolog 2, zinc finger protein (Drosophila)" Scrt2
1457311_at 1.00311 calcium/calmodulin-dependent protein kinase II alpha Camk2a
1457312_at 1.11712 NHL repeat containing 2 Nhlrc2
1457313_at 1.06818 oculocerebrorenal syndrome of Lowe Oclrl
1457314_at 1.02591 LINE-1 type transposase domain containing 1 L1td1
1457315_at 0.99766 RIKEN cDNA A230051G13 gene A230051G13Rik
1457316_at 1.26683 microtubule-associated protein 6 Mtap6
1457317_at 1.67048 --- ---
1457318_at 1.00729 RIKEN cDNA A330008L17 gene A330008L17Rik
1457319_at 1.07210 RIKEN cDNA A130038J17 gene A130038J17Rik
1457320_at 0.82991 Wings apart-like homolog (Drosophila) Wapal
1457321_at 0.83911 hypothetical protein LOC667963 LOC667963
1457322_at 1.05374 "Complement component 1, q subcomponent, beta polypeptide" C1qb
1457323_at 0.90570 hypothetical protein LOC632191 LOC632191
1457324_at 1.06635 "Opioid receptor, sigma 1" Oprs1
1457325_at 1.14172 "K+ voltage-gated channel, subfamily S, 2" Kcns2
1457326_at 0.90570 similar to zinc finger protein 616 /// similar to zinc finger protein 420
LOC545191 /// LOC631233
1457327_at 1.04104 SNF2 histone linker PHD RING helicase Shprh
1457328_at 0.81234 "adrenergic receptor, alpha 2c" Adra2c
1457329_at 0.92388 adenylate cyclase 1 Adcy1
1457330_at 0.93185 "GPC PDZ domain containing family, member 2" Gipc2
1457331_at 0.97341 Retbindin Rtbdn
1457332_at 0.93864 protein arginine N-methyltransferase 3 Prmt3
1457333_at 1.01300 sidekick homolog 2 (chicken) Sdk2
1457334_at 1.48656 RIKEN cDNA C130057M05 gene C130057M05Rik
1457335_at 0.95125 "Mammary gland RCB-0527 Jyg-MC(B) cDNA, RIKEN full-length enriched library, clone:G930025P03 product:unclassifiable, full insert sequence" ---
1457336_at 1.04309 R3H domain 1 (binds single-stranded nucleic acids) R3hdm1
1457337_at 1.02512 --- ---
1457338_at 1.04281 "Protein phosphatase 1, regulatory (inhibitor) subunit 12B" Ppp1r12b
1457339_at 1.15489 Cerebral cavernous malformation 2 homolog (human) Ccm2
1457340_at 1.00297 RUN and FYVE domain containing 1 Rufy1
1457341_at 0.93680 "Transcribed locus, moderately similar to XP_579758.1 PREDICTED: hypothetical protein XP_579758 [Rattus norvegicus]" ---
1457342_at 1.00043 "zinc finger protein, subfamily 1A, 4 (Eos)" Zfpn1a4
1457343_at 1.14552 "Striatin, calmodulin binding protein 3" Strn3
1457344_at 1.02491 neuropilin (NRP) and tolloid (TLL)-like 2 Neto2
1457345_at 1.25248 Polymerase (RNA) III (DNA directed) polypeptide B Polr3b
1457346_at 1.00211 Transcribed locus ---
1457347_at 1.07129 "ryanodine receptor 1, skeletal muscle" Ryr1

1457348_at 0.98265 ADP-ribosylation factor 4 Arf4
1457349_at 0.69205 similar to Tetratricopeptide repeat protein 6 (TPR repeat protein 6)
LOC639426
1457350_at 0.87932 period homolog 2 (Drosophila) Per2
1457351_at 0.87556 "TAF2 RNA polymerase II, TATA box binding protein (TBP)-associated factor" Taf2
1457352_x_at 0.97300 RIKEN cDNA 9430071P14 gene 9430071P14Rik
1457353_at 0.93937 Translocase of outer mitochondrial membrane 70 homolog A (yeast) Tomm70a
1457354_at 1.03491 keratin 222 Krt222
1457355_at 0.96026 --- ---
1457356_at 1.87580 Insulin-like growth factor 2 receptor Igf2r
1457357_at 1.68887 tousled-like kinase 2 (Arabidopsis) Tlk2
1457358_at 1.70311 --- ---
1457359_at 0.86452 "inositol polyphosphate-4-phosphatase, type II" Inpp4b
1457360_at 0.95761 Craniofacial development protein 1 Cfdp1
1457361_at 0.98065 RIKEN cDNA C630007C17 gene C630007C17Rik
1457362_at 1.02603 Agouti related protein Agrp
1457363_at 0.93729 hypothetical protein LOC654469 LOC654469
1457364_at 1.01513 --- ---
1457365_at 1.00846 --- ---
1457366_at 1.11856 cell division cycle 40 homolog (yeast) Cdc40
1457367_at 1.14626 "Amylase 2, pancreatic" Amy2
1457368_at 0.93899 RIKEN cDNA 2310033K02 gene 2310033K02Rik
1457369_at 1.31340 Phosphatase and actin regulator 4 Phactr4
1457370_at 1.21325 RAB GTPase activating protein 1 Rabgap1
1457371_at 1.02030 RIKEN cDNA 5230400G24 gene 5230400G24Rik
1457372_at 0.97915 --- ---
1457373_at 0.94259 Transcribed locus ---
1457374_at 1.96538 "neural precursor cell expressed, developmentally down-regulated gene 4-like" Nedd4l
1457375_at 0.93645 "PREDICTED: Mus musculus hypothetical protein LOC627707 (LOC627707), mRNA" ---
1457376_at 1.09978 Integrin alpha 4 Itga4
1457377_at 1.04113 L(3)mbt-like 3 (Drosophila) L3mbtl3
1457378_at 0.98307 Mitochondrial ribosomal protein L54 Mrpl54
1457379_at 1.11308 expressed sequence C78549 C78549
1457380_at 1.19669 expressed sequence C85504 C85504
1457381_at 0.99063 Angiopoietin 1 Angpt1
1457382_at 1.01534 "PREDICTED: Mus musculus similar to Heterogeneous nuclear ribonucleoprotein A1 (Helix-destabilizing protein) (Single-strand binding protein) (hnRNP core protein A1) (HDP) (LOC628392), mRNA" ---
1457383_at 1.31723 RIKEN cDNA 1500011J06 gene 1500011J06Rik

1457384_at 0.95982 "0 day neonate head cDNA, RIKEN full-length enriched library, clone:4833438M18 product:unclassifiable, full insert sequence" ---

1457385_at 1.00393 --- ---

1457386_at 0.93557 RIKEN cDNA 9530010C24 gene 9530010C24Rik

1457387_at 0.96842 Replication initiator 1 Repin1

1457388_at 0.86543 RIKEN cDNA A830080L01 gene A830080L01Rik

1457389_at 0.90287 RIKEN cDNA 9830166K06 gene 9830166K06Rik

1457390_at 1.46608 PRP3 pre-mRNA processing factor 3 homolog (yeast) Prpf3

1457391_at 0.83149 Vesicle-associated membrane protein 3 Vamp3

1457392_at 1.34269 "Adult retina cDNA, RIKEN full-length enriched library, clone:A930031I16 product:unclassifiable, full insert sequence" ---

1457393_at 0.98722 RIKEN cDNA 2700008B19 gene 2700008B19Rik

1457394_at 0.95705 RIKEN cDNA 2900002K06 gene 2900002K06Rik

1457395_at 0.94556 RIKEN cDNA 5330421F07 gene 5330421F07Rik

1457396_at 0.95756 --- ---

1457397_at 1.00937 Kruppel-like factor 14 Klf14

1457398_at 1.96767 Guanosine diphosphate (GDP) dissociation inhibitor 2 Gdi2

1457399_at 1.00749 "ELAV (embryonic lethal, abnormal vision, Drosophila)-like 4 (Hu antigen D)" Elavl4

1457400_at 0.77798 sorting nexin 19 Snx19

1457401_at 0.93712 "dynein, axonemal, heavy chain 9" Dnahc9

1457402_at 1.00773 Sulfatase 1 Sulf1

1457403_at 1.00707 RIKEN cDNA 9130409I23 gene 9130409I23Rik

1457404_at 1.14539 "Nuclear factor of kappa light polypeptide gene enhancer in B-cells inhibitor, zeta" Nfkbiz

1457405_at 0.99533 Dynamin 3 Dnm3

1457406_at 0.95865 "Adult male brain UNDEFINED_CELL_LINE cDNA, RIKEN full-length enriched library, clone:M5C1091A08 product:unclassifiable, full insert sequence" ---

1457407_at 1.17219 roundabout homolog 1 (Drosophila) Robo1

1457408_at 1.02049 "Phosphodiesterase 4A, cAMP specific" Pde4a

1457409_at 0.99265 fucosyltransferase 9 Fut9

1457410_at 1.86199 Rho GTPase activating protein 5 Arhgap5

1457411_at 0.98911 --- ---

1457412_at 1.09182 "sodium channel, voltage-gated, type VIII, alpha" Scn8a

1457413_at 0.92266 Receptor tyrosine kinase-like orphan receptor 1 Ror1

1457414_at 1.02379 "9.5 days embryo parthenogenote cDNA, RIKEN full-length enriched library, clone:B130039H18 product:unclassifiable, full insert sequence" ---

1457415_a_at 1.07255 RIKEN cDNA 4930513N10 gene 4930513N10Rik

1457416_at 0.98649 Methionine aminopeptidase 2 Metap2

1457417_at 1.03680 --- ---

1457418_at 0.92293 "Spermatogenesis associated glutamate (E)-rich protein 1, pseudogene 1" Speer1-ps1

1457419_s_at 0.95052 "Spermatogenesis associated glutamate (E)-rich protein 1, pseudogene 1"

Speer1-ps1

1457420_at 1.02679 lemur tyrosine kinase 3 Lmtk3
1457421_at 1.01830 RIKEN cDNA 1700063D05 gene 1700063D05Rik
1457422_at 1.08692 RIKEN cDNA 9030205A07 Gene 9030205A07Rik
1457423_at 0.92978 similar to voltage gated channel like 1 LOC675405
1457424_at 1.00024 eyes absent 1 homolog (Drosophila) Eya1
1457425_at 0.95293 RIKEN cDNA C130038G02 gene C130038G02Rik
1457426_at 1.04121 RIKEN cDNA 1700048O20 gene 1700048O20Rik
1457427_at 1.02996 Adherens junction associated protein 1 Ajap1
1457428_at 1.01631 hypothetical protein C230040D14 C230040D14
1457429_s_at 0.94957 "gene model 106, (NCBI)" Gm106
1457430_at 0.96764 "Sparc/osteonectin, cwcv and kazal-like domains proteoglycan 2" Spock2
1457431_at 0.99082 Transcribed locus ---
1457432_at 0.93649 prospero-related homeobox 1 Prox1
1457433_x_at 1.08877 Zinc finger protein 120 Zfp120
1457434_s_at 0.92559 "protein tyrosine phosphatase-like (proline instead of catalytic arginine), member a" Ptpla
1457435_x_at 2.41394 myomesin 2 Myom2
1457436_at 0.96471 "TBC1 domain family, member 7" Tbc1d7
1457437_at 0.87260 RIKEN cDNA C630035N08 gene C630035N08Rik
1457438_at 1.04962 --- ---
1457439_at 0.98929 Cyclin T2 Ccnt2
1457440_at 1.02346 somatostatin receptor 4 Sstr4
1457441_at 0.84817 early B-cell factor 1 Ebf1
1457442_at 0.97640 expressed sequence AW125324 AW125324
1457443_at 0.97197 --- ---
1457444_at 1.00178 "Adult male aorta and vein cDNA, RIKEN full-length enriched library, clone: A530012F11 product:unclassifiable, full insert sequence" ---
1457445_at 1.45227 trichorhinophalangeal syndrome I (human) Trps1
1457446_at 0.98249 opioid binding protein/cell adhesion molecule-like Opcml
1457447_at 1.27355 Retinoblastoma 1 Rb1
1457448_at 0.95268 patatin-like phospholipase domain containing 1 Pnpla1
1457449_at 1.43213 Longevity assurance homolog 6 (S. cerevisiae) Lass6
1457450_at 1.10466 RIKEN cDNA A830091I15 gene A830091I15Rik
1457451_at 1.06178 --- ---
1457452_at 1.00186 expressed sequence AA407782 AA407782
1457453_at 1.05953 RIKEN cDNA 1700057G04 gene 1700057G04Rik
1457454_at 2.28274 ubiquitin specific peptidase 47 Usp47
1457455_at 1.56944 suppressor of hairy wing homolog 4 (Drosophila) Suhw4
1457456_at 1.01470 mitogen activated protein kinase kinase kinase 10 Map3k10
1457457_at 0.91047 RIKEN cDNA 5033405K12 gene 5033405K12Rik
1457458_at 1.89450 cDNA sequence BC057627 BC057627
1457459_at 1.64594 expressed sequence AU014973 AU014973

1457460_at 1.08139 acyl-CoA synthetase short-chain family member 2 Acss2
1457461_at 0.94554 RIKEN cDNA A630039O03 gene A630039O03Rik
1457462_at 0.94071 --- ---
1457463_at 1.04347 Down syndrome cell adhesion molecule-like 1 Dscam1
1457464_at 1.67540 hypothetical LOC403342 C130015C19
1457465_at 1.02591 shroom family member 4 Shroom4
1457466_at 1.04081 expressed sequence AA409368 AA409368
1457467_at 0.97536 Pbx/knotted 1 homeobox Pknox1
1457468_at 2.48293 "Tankyrase, TRF1-interacting ankyrin-related ADP-ribose polymerase 2"
Tnks2
1457469_at 0.80879 --- ---
1457470_at 1.27658 "DNA segment, Chr 3, ERATO Doi 452, expressed" D3Ertd452e
1457471_at 0.81326 interleukin 17 receptor C Il17rc
1457472_at 0.87384 trinucleotide repeat containing 15 Tnrc15
1457473_at 0.94436 chromodomain helicase DNA binding protein 1 Chd1
1457474_at 1.07748 Cartilage acidic protein 1 Crtac1
1457475_at 1.16219 expressed sequence C80993 C80993
1457476_at 1.04503 "DNA segment, Chr 8, ERATO Doi 362, expressed" D8Ertd362e
1457477_at 1.00121 Muscleblind-like 2 Mbnl2
1457478_at 0.93083 Spermatid perinuclear RNA binding protein Strbp
1457479_at 1.67426 --- ---
1457480_at 1.53447 Polycomb group ring finger 3 Pcgf3
1457481_at 0.94272 Transcribed locus ---
1457482_at 0.95843 "jumonji, AT rich interactive domain 1B (Rbp2 like)" Jarid1b
1457483_at 1.66434 AT rich interactive domain 5B (Mrf1 like) Arid5b
1457484_at 1.17411 hypothetical D930050J11 D930050J11
1457485_at 1.07182 F-box and WD-40 domain protein 11 Fbxw11
1457486_at 1.05953 "DNA segment, Chr 9, University of California at Los Angeles 1" D9Ucla1
1457487_at 0.89117 "Adult male liver tumor cDNA, RIKEN full-length enriched library, clone:
C730014I02 product:unclassifiable, full insert sequence" ---
1457488_at 1.01579 RIKEN cDNA B230339M05 gene B230339M05Rik
1457489_at 1.82215 Eukaryotic translation initiation factor 4E Eif4e
1457490_at 0.98104 RIKEN cDNA B230218L05 gene B230218L05Rik
1457491_at 1.08006 --- ---
1457492_at 2.19780 Triple functional domain (PTPRF interacting) Trio
1457493_at 1.13980 phosphatase and tensin homolog Pten
1457494_at 1.06743 "Adult male corpus striatum cDNA, RIKEN full-length enriched library, clone:
C030004I10 product:unclassifiable, full insert sequence" ---
1457495_at 1.01469 RIKEN cDNA 2900052N01 gene 2900052N01Rik
1457496_at 1.01173 "Adult male olfactory brain cDNA, RIKEN full-length enriched library,
clone:6430514C12 product:unclassifiable, full insert sequence" ---
1457497_at 0.96386 Synaptotagmin I Syt1
1457498_at 1.07238 "mannosidase 2, alpha 2" Man2a2

1457499_at	0.86815 WW domain containing E3 ubiquitin protein ligase 2	Wwp2
1457500_at	1.42009 DEP domain containing 5 Depdc5	
1457501_at	1.00944 hypothetical protein LOC622665	LOC622665
1457502_at	0.97138 Quaking Qk	
1457503_at	1.05607 RIKEN cDNA 2310035C23 gene	2310035C23Rik
1457504_at	1.02467 --- ---	
1457505_at	1.22994 Utrophin	Utrn
1457506_at	0.92703 "Amyotrophic lateral sclerosis 2 (juvenile) chromosome region, candidate 19 (human)"	Als2cr19
1457507_at	1.03171 Neuregulin 3	Nrg3
1457508_at	2.32744 RIKEN cDNA C430003N24 gene	C430003N24Rik
1457509_at	1.34842 "sno, strawberry notch homolog 1 (Drosophila)"	Sbno1
1457510_at	1.43781 PHD finger protein 14	Phf14
1457511_at	1.09225 regulator of G-protein signaling 12	Rgs12
1457512_at	0.97908 "16 days neonate thymus cDNA, RIKEN full-length enriched library, clone: A130094L10 product:unclassifiable, full insert sequence"	---
1457513_at	1.17293 RIKEN cDNA 6330569M22 gene	6330569M22Rik
1457514_at	1.09050 YLP motif containing 1	Ylpm1
1457515_at	1.15290 Homeodomain interacting protein kinase 1	Hipk1
1457516_at	0.94497 RIKEN cDNA 1110033M05 gene	1110033M05Rik
1457517_at	1.08103 RIKEN cDNA A730046G19 gene	A730046G19Rik
1457518_at	1.00347 RIKEN cDNA C630004H02 gene	C630004H02Rik
1457519_at	1.11342 Transcribed locus	---
1457520_at	0.93989 Grainyhead-like 2 (Drosophila)	Grhl2
1457521_at	0.89927 --- ---	
1457522_at	1.99939 RIKEN cDNA 1110034C04 gene	1110034C04Rik
1457523_at	1.03403 RIKEN cDNA 9530056K15 gene	9530056K15Rik
1457524_at	1.19521 Transcription factor 12	Tcf12
1457525_at	0.95360 "Activated spleen cDNA, RIKEN full-length enriched library, clone: F830103I17 product:unclassifiable, full insert sequence"	---
1457526_at	0.95183 growth factor receptor bound protein 14	Grb14
1457527_at	0.91703 --- ---	
1457528_at	1.77750 "solute carrier family 4, sodium bicarbonate cotransporter, member 7"	Slc4a7
1457529_x_at	1.02051 RIKEN cDNA 2810021G02 gene	2810021G02Rik
1457530_at	1.02471 Transcribed locus	---
1457531_a_at	0.98708 RIKEN cDNA A930017K11 gene	A930017K11Rik
1457532_at	1.00806 GTPase activating RANGAP domain-like 1	Garnl1
1457533_at	0.99462 --- ---	
1457534_at	1.49978 Ubiquitin associated protein 2-like	Ubap2l
1457535_at	1.15162 RIKEN cDNA 4932438A13 gene	4932438A13Rik
1457536_at	1.08025 glypican 5	Gpc5
1457537_at	1.09338 "RAB1, member RAS oncogene family"	Rab1
1457538_at	0.89291 RIKEN cDNA 5930422O12 gene	5930422O12Rik

1457539_at 1.06867 "DNA segment, Chr 10, ERATO Doi 709, expressed" D10Ertd709e
 1457540_at 1.22602 "Solute carrier family 25 (mitochondrial carrier, phosphate carrier), member 26" Slc25a26
 1457541_at 1.14006 A kinase (PRKA) anchor protein 14 Akap14
 1457542_at 1.08046 nucleoporin 133 Nup133
 1457543_at 0.93979 "NOL1/NOP2/Sun domain family, member 7" Nsun7
 1457544_at 0.93066 ubiquitin-conjugating enzyme E2W (putative) Ube2w
 1457545_at 1.01819 RIKEN cDNA 9530036O11 gene 9530036O11Rik
 1457546_at 1.02657 "CCR4-NOT transcription complex, subunit 1" Cnot1
 1457547_at 1.01958 "Adult male spinal cord cDNA, RIKEN full-length enriched library, clone: A330081M13 product:unclassifiable, full insert sequence" ---
 1457548_at 1.25735 "A disintegrin-like and metallopeptidase (reprolysin type) with thrombospondin type 1 motif, 6" Adamts6
 1457549_at 1.07513 Cyclin D3 Ccnd3
 1457550_at 0.92240 RIKEN cDNA 9530059O14 gene 9530059O14Rik
 1457551_at 1.32438 "Activin A receptor, type 1" Acvr1
 1457552_at 1.10516 zinc finger protein 295 Zfp295
 1457553_at 2.39585 "CUG triplet repeat, RNA binding protein 2" Cugbp2
 1457554_at 0.83311 apolipoprotein B Apob
 1457555_at 0.97979 G protein-coupled receptor 151 Gpr151
 1457556_at 0.84811 "PREDICTED: Mus musculus hypothetical LOC433662 (LOC433662), mRNA" ---
 1457557_at 1.13245 RIKEN cDNA A330076H08 gene A330076H08Rik
 1457558_at 1.08296 RIKEN cDNA A330050F15 gene A330050F15Rik
 1457559_at 1.22887 WW domain containing adaptor with coiled-coil Wac
 1457560_at 1.24242 Single-stranded DNA binding protein 3 Ssbp3
 1457561_at 1.06950 integrin alpha 5 (fibronectin receptor alpha) Itga5
 1457562_at 1.21809 "ribosomal protein S6 kinase, polypeptide 1" Rps6kb1
 1457563_at 0.88078 epidermal growth factor receptor Egfr
 1457564_at 1.08793 "DNA fragmentation factor, alpha subunit" Dffa
 1457565_at 1.00207 RIKEN cDNA C130052G03 gene C130052G03Rik
 1457566_at 0.97186 RIKEN cDNA A830058L05 gene A830058L05Rik
 1457567_at 1.00005 N-acetylglucosamine kinase Nagk
 1457568_at 1.00774 hypothetical protein C230004L04 C230004L04
 1457569_at 1.88047 "DNA segment, Chr 8, ERATO Doi 51, expressed" D8Ertd51e
 1457570_at 0.88611 RIKEN cDNA A230001M10 gene A230001M10Rik
 1457571_at 0.73263 zinc finger protein 68 Zfp68
 1457572_at 1.09603 "Membrane associated guanylate kinase, WW and PDZ domain containing 1" Magi1
 1457573_at 1.03961 Topoisomerase (DNA) I Top1
 1457574_at 0.96238 similar to F-box and WD-40 domain protein FBXW14 /// similar to F-box and WD-40 domain protein FBXW14 LOC635668 /// LOC665428
 1457575_at 1.08516 expressed sequence AU021128 AU021128

1457576_at 0.79751 --- ---
 1457577_at 0.91522 --- ---
 1457578_at 1.09028 "Peroxisome proliferative activated receptor, gamma, coactivator 1 beta"
 Ppargc1b
 1457579_at 1.41859 "DNA segment, Chr 11, ERATO Doi 717, expressed" D11Ertd717e
 1457580_at 1.13635 chromodomain helicase DNA binding protein 8 Chd8
 1457581_at 1.00031 UBX domain containing 3 Ubx3
 1457582_at 0.94831 "Ubiquitously transcribed tetratricopeptide repeat gene, Y chromosome" Uty
 1457583_at 0.92802 "ST6 (alpha-N-acetyl-neuraminyl-2,3-beta-galactosyl-1,3)-N-
 acetylglactosaminide alpha-2,6-sialyltransferase 3" St6galnac3
 1457584_at 1.01492 expressed sequence AI848100 AI848100
 1457585_at 0.89994 transmembrane protein 24 Tmem24
 1457586_at 1.36471 RAS p21 protein activator 2 Rasa2
 1457587_at 0.98088 Dendritic cell protein GA17 Ga17
 1457588_at 1.54509 expressed sequence C76213 C76213
 1457589_at 0.87668 FAT tumor suppressor homolog 3 (Drosophila) Fat3
 1457590_at 1.16845 "Purinergic receptor P2X, ligand-gated ion channel, 1" P2rx1
 1457591_at 1.18187 Nucleolar protein 10 Nol10
 1457592_x_at 0.93886 RIKEN cDNA 2210018M03 gene 2210018M03Rik
 1457593_at 1.19020 RIKEN cDNA 2610202C22 gene 2610202C22Rik
 1457594_at 0.92269 Transcribed locus ---
 1457595_at 0.93012 "2 cells egg cDNA, RIKEN full-length enriched library, clone:B020028D20
 product:hypothetical protein, full insert sequence" ---
 1457596_at 0.98113 Son cell proliferation protein Son
 1457597_at 0.80020 Transcribed locus ---
 1457598_at 1.18724 Thioredoxin-like 2 Txnl2
 1457599_at 0.92444 RIKEN cDNA 2310047H23 gene 2310047H23Rik
 1457600_x_at 1.11380 Naked cuticle 2 homolog (Drosophila) Nkd2
 1457601_at 1.03692 "Adult male liver tumor cDNA, RIKEN full-length enriched library, clone:
 C730010C06 product:unclassifiable, full insert sequence" ---
 1457602_at 0.96042 "16 days neonate thymus cDNA, RIKEN full-length enriched library, clone:
 A130043M03 product:unclassifiable, full insert sequence" ---
 1457603_at 0.98412 "protein phosphatase 1, regulatory (inhibitor) subunit 9A" Ppp1r9a
 1457604_x_at 0.83160 "cytochrome P450, family 11, subfamily a, polypeptide 1" Cyp11a1
 1457605_at 1.13538 Ataxin 1 Atxn1
 1457606_x_at 0.96208 expressed sequence AU015228 AU015228
 1457607_at 0.89163 expressed sequence AW046457 AW046457
 1457608_at 0.91710 "GTPase, IMAP family member 8" Gimap8
 1457609_at 1.01534 "Adenosine deaminase, RNA-specific, B2" Adarb2
 1457610_at 0.95017 Expressed sequence AL033326 AL033326
 1457611_x_at 0.99308 RIKEN cDNA 2010309E21 gene 2010309E21Rik
 1457612_at 1.14557 similar to glyceraldehyde-3-phosphate dehydrogenase /// similar to
 glyceraldehyde-3-phosphate dehydrogenase (phosphorylating)-like /// similar to glyceraldehyde-3-

1457629_at 1.04197 cDNA sequence BC060631 BC060631
1457630_at 1.12364 Wolf-Hirschhorn syndrome candidate 1-like 1 (human) Whsc111
1457631_at 1.08424 RIKEN cDNA 2310047N11 gene 2310047N11Rik
1457632_s_at 1.01402 myeloid ecotropic viral integration site-related gene 1 Mrg1
1457633_x_at 0.96904 "cytochrome c oxidase, subunit VI a, polypeptide 2" Cox6a2
1457634_at 1.15800 RIKEN cDNA 4930488E11 gene /// similar to thymosin beta-like protein ///
similar to thymosin beta-like protein /// similar to thymosin beta-like protein 4930488E11Rik ///
LOC666244 /// LOC669681 /// LOC674935
1457635_s_at 0.85044 "nuclear receptor subfamily 3, group C, member 1" Nr3c1
1457636_x_at 1.01304 --- ---
1457637_at 1.11254 Prickle like 1 (Drosophila) Prickle1
1457638_x_at 0.88978 replication factor C (activator 1) 2 Rfc2
1457639_at 1.01331 "ATPase, H⁺ transporting, lysosomal V1 subunit H" Atp6v1h
1457640_x_at 0.94804 Transcribed locus ---
1457641_at 1.26250 CREB binding protein Crebbp
1457642_at 1.13431 RIKEN cDNA 5730507N06 gene 5730507N06Rik
1457643_x_at 0.99333 RIKEN cDNA 2610316D01 gene 2610316D01Rik
1457644_s_at 6.88001 chemokine (C-X-C motif) ligand 1 Cxcl1
1457645_at 1.01599 RIKEN cDNA C130079G13 gene C130079G13Rik
1457646_at 0.99514 "homeo box A11, opposite strand transcript" Hoxa11os
1457647_x_at 0.97439 RIKEN cDNA 1600023N17 gene 1600023N17Rik
1457648_x_at 1.12902 cDNA sequence BC004044 BC004044
1457649_x_at 0.99975 expressed sequence AI848149 AI848149
1457650_x_at 1.05312 autophagy-related 4A (yeast) /// similar to Cysteine protease ATG4A
(Autophagy-related protein 4 homolog A) (Autophagin-2) (Autophagy-related cysteine endopeptidase 2)
(AUT-like 2 cysteine endopeptidase) /// similar to Cysteine protease ATG4A (Autophagy-related protein
4 homolog A) (Autophagin-2) (Autophagy-related cysteine endopeptidase 2) (AUT-like 2 cysteine
endopeptidase) Atg4a /// LOC666476 /// LOC675173
1457651_x_at 0.76769 rad and gem related GTP binding protein 2 Rem2
1457652_x_at 0.90854 Prostaglandin-endoperoxide synthase 1 Ptgs1
1457653_at 1.02482 RIKEN cDNA A630042L21 gene A630042L21Rik
1457654_at 0.99328 "Transcribed locus, moderately similar to NP_598437.1 O-sialoglycoprotein
endopeptidase [Mus musculus]" ---
1457655_x_at 0.92245 Rho GTPase activating protein 27 Arhgap27
1457656_s_at 0.96687 RIKEN cDNA C230085N15 gene C230085N15Rik
1457657_at 1.20275 --- ---
1457658_x_at 1.03820 annexin A4 Anxa4
1457659_x_at 0.95429 outer dense fiber of sperm tails 1 Odf1
1457660_x_at 1.11876 RIKEN cDNA 5430416O09 gene 5430416O09Rik
1457661_at 1.07958 RIKEN cDNA B930036G03 gene B930036G03Rik
1457662_x_at 0.80013 thiamine pyrophosphokinase Tpk1
1457663_at 1.03604 RIKEN cDNA 1700006F04 gene 1700006F04Rik
1457664_x_at 0.88246 Complement component 2 (within H-2S) C2

1457665_x_at	0.99169	"cytochrome P450, family 27, subfamily a, polypeptide 1"	Cyp27a1
1457666_s_at	1.39055	interferon activated gene 202B	Ifi202b
1457667_x_at	0.92583	dual-specificity tyrosine-(Y)-phosphorylation regulated kinase 2	Dyrk2
1457668_x_at	1.00940	EBNA1 binding protein 2	Ebna1bp2
1457669_x_at	0.88743	replication factor C (activator 1) 2	Rfc2
1457670_s_at	0.96859	lamin A	Lmna
1457671_at	1.00921	RIKEN cDNA 9330120H11 gene	9330120H11Rik
1457672_at	0.78989	chromodomain helicase DNA binding protein 9	Chd9
1457673_at	1.17146	RIKEN cDNA 2610005L07 gene	2610005L07Rik
1457674_at	1.04855	vacuolar protein sorting 13A (yeast)	Vps13a
1457675_at	1.05121	RIKEN cDNA 2510002D24 gene	2510002D24Rik
1457676_at	1.12290	toll-interleukin 1 receptor (TIR) domain-containing adaptor protein	Tirap
1457677_at	0.90416	CDNA sequence BC057593	BC057593
1457678_at	1.12695	RIKEN cDNA 2310035C23 gene	2310035C23Rik
1457679_at	1.03086	"DNA cross-link repair 1C, PSO2 homolog (S. cerevisiae)"	Dclre1c
1457680_a_at	0.99514	transmembrane protein 69	Tmem69
1457681_at	1.00004	RIKEN cDNA 2610301F02 gene	2610301F02Rik
1457682_at	0.96963	RIKEN cDNA 9030420J04 gene	9030420J04Rik
1457683_at	1.00799	"glutamate receptor, ionotropic, kainate 2 (beta 2)"	Grik2
1457684_at	0.86876	RIKEN cDNA 1700037H04 gene	1700037H04Rik
1457685_at	0.83986	RIKEN cDNA D330034E10 gene	D330034E10Rik
1457686_at	0.97259	RIKEN cDNA B230333C21 gene	B230333C21Rik
1457687_at	0.87368	B-cell leukemia/lymphoma 2	Bcl2
1457688_at	0.81561	zinc finger protein 398	Zfp398
1457689_at	1.04893	SET binding factor 2	Sbf2
1457690_at	1.03788	"kalirin, RhoGEF kinase"	Kalrn
1457691_at	0.90493	Epstein-Barr virus induced gene 2	Ebi2
1457692_at	1.22956	---	---
1457693_a_at	1.09617	RIKEN cDNA 6430537H07 gene	6430537H07Rik
1457694_at	0.97858	RIKEN cDNA 6430537H07 gene	6430537H07Rik
1457695_at	1.08748	AP1 gamma subunit binding protein 1	Ap1gbp1
1457696_at	0.92958	Rab interacting lysosomal protein	Rilp
1457697_at	1.05796	RIKEN cDNA A330021E22 gene	A330021E22Rik
1457698_at	0.91626	expressed sequence AI314831	AI314831
1457699_at	1.07486	RIKEN cDNA E330009J07 gene	E330009J07Rik
1457700_at	1.10743	RIKEN cDNA 4732474A20 gene	4732474A20Rik
1457701_at	0.93290	transmembrane protein 136	Tmem136
1457702_at	0.91678	G-protein coupled receptor 12	Gpr12
1457703_at	0.98879	expressed sequence BE686333	BE686333
1457704_at	0.93681	zinc finger protein 533	Zfp533
1457705_at	0.97214	RIKEN cDNA 5330421C15 gene	5330421C15Rik
1457706_at	0.89262	cDNA sequence BC030336	BC030336
1457707_at	1.05585	"gene model 489, (NCBI)"	Gm489

1457708_at 0.98170 methyl-CpG binding domain protein 4 Mbd4
 1457709_a_at 1.02964 RIKEN cDNA A930005H10 gene A930005H10Rik
 1457710_at 0.91333 zinc fingers and homeoboxes 3 Zhx3
 1457711_at 1.11797 RAN binding protein 3 Ranbp3
 1457712_at 1.33591 chromodomain helicase DNA binding protein 8 Chd8
 1457713_at 0.94760 "Excision repair cross-complementing rodent repair deficiency,
 complementation group 5" Ercc5
 1457714_at 1.00872 RIKEN cDNA 2310005N03 gene 2310005N03Rik
 1457715_at 0.94906 RIKEN cDNA 1010001B22 gene 1010001B22Rik
 1457716_at 0.99191 OTU domain containing 7B Otud7b
 1457717_at 0.96281 expressed sequence AI987986 AI987986
 1457718_at 0.98809 RIKEN cDNA D030022P06 gene D030022P06Rik
 1457719_at 0.95562 Transcribed locus ---
 1457720_at 0.98106 --- ---
 1457721_at 1.01191 Expressed sequence AI118064 AI118064
 1457722_at 0.99388 killer cell lectin-like receptor subfamily B member 1F Klrb1f
 1457723_at 1.92499 Paraspeckle protein 1 Pspc1
 1457724_at 0.88598 cathepsin L Ctsl
 1457725_at 0.98188 "Membrane-spanning 4-domains, subfamily A, member 4B" Ms4a4b
 1457726_at 1.04018 ribosomal protein S15a Rps15a
 1457727_at 1.27863 hypothetical protein LOC544864 LOC544864
 1457728_at 0.99193 6-phosphogluconolactonase Pgl5
 1457729_at 1.03581 Transcribed locus ---
 1457730_at 0.96185 "Fanconi anemia, complementation group M" Fancm
 1457731_at 1.11696 "small nuclear RNA activating complex, polypeptide 3" Snapc3
 1457732_at 1.09567 Protein-L-isoaspartate (D-aspartate) O-methyltransferase domain containing 2
 Pcmt2
 1457733_at 0.97510 Transcribed locus ---
 1457734_at 1.03669 "POU domain, class 3, transcription factor 2" Pou3f2
 1457735_at 0.95434 Transcribed locus ---
 1457736_at 0.85601 vacuolar protein sorting 37D (yeast) Vps37d
 1457737_at 1.00854 hypothetical protein E130012K09 E130012K09
 1457738_at 1.04079 "LSM14 homolog A (SCD6, S. cerevisiae)" Lsm14a
 1457739_at 1.08837 "fascin homolog 1, actin bundling protein (Strongylocentrotus purpuratus)"
 Fscn1
 1457740_at 1.22152 Aryl hydrocarbon receptor nuclear translocator-like Arntl
 1457741_at 1.02048 --- ---
 1457742_at 0.98832 "DNA segment, Chr 9, Wayne State University 74, expressed" D9Wsu74e
 1457743_at 1.03563 "DNA segment, Chr 13, Brigham & Women's Genetics 1146 expressed"
 D13Bwg1146e
 1457744_at 1.04695 DEAD (Asp-Glu-Ala-Asp) box polypeptide 46 Ddx46
 1457745_at 1.00669 G protein-coupled receptor 4 Gpr4
 1457746_at 1.94666 RIKEN cDNA 2610028H07 gene 2610028H07Rik

1457747_at 0.95779 RIKEN cDNA 9330109K16 gene 9330109K16Rik
1457748_at 1.05352 pogo transposable element with ZNF domain Pogz
1457749_at 1.03650 coiled-coil and C2 domain containing 1B Cc2d1b
1457750_at 1.01812 "DNA segment, Chr 16, ERATO Doi 780, expressed" D16Ertd780e
1457751_at 1.05523 RIKEN cDNA 4832420A03 gene 4832420A03Rik
1457752_at 0.96879 expressed sequence AU014876 AU014876
1457753_at 1.00142 toll-like receptor 13 Tlr13
1457754_at 1.04785 RIKEN cDNA 4930430F08 gene 4930430F08Rik
1457755_at 1.05101 "guanine nucleotide binding protein (G protein), gamma 8 subunit" Gng8
1457756_at 0.91071 zinc finger protein 192 Zfp192
1457757_at 1.05732 similar to granulosa cell HMG-box protein-1 LOC269389
1457758_at 0.95281 enhancer of yellow 2 homolog (Drosophila) Eny2
1457759_at 1.00166 RIKEN cDNA A630081D01 gene A630081D01Rik
1457760_at 1.37518 RIKEN cDNA A930004J17 gene A930004J17Rik
1457761_at 0.92289 expressed sequence C79607 C79607
1457762_at 1.11515 Tetratricopeptide repeat domain 15 Ttc15
1457763_at 0.96115 "Gamma-aminobutyric acid (GABA-A) receptor, subunit delta" Gabrd
1457764_at 0.93556 RIKEN cDNA A530058O07 gene A530058O07Rik
1457765_at 2.43631 "Proteasome (prosome, macropain) 26S subunit, non-ATPase, 14" Psm14
1457766_at 0.97141 WAP four-disulfide core domain 6A Wfdc6a
1457767_at 0.90761 Nischarin Nisch
1457768_at 1.62196 prune homolog (Drosophila) Prune
1457769_at 0.92226 histocompatibility 60 H60
1457770_at 0.89020 "solute carrier family 39 (zinc transporter), member 14" Slc39a14
1457771_at 0.96914 "Protein tyrosine phosphatase, receptor type, O" Ptpro
1457772_at 0.99799 "Adult male aorta and vein cDNA, RIKEN full-length enriched library, clone:
A530055C18 product:unclassifiable, full insert sequence" ---
1457773_at 0.97959 SLAM family member 6 Slamf6
1457774_at 1.07275 RIKEN cDNA 2310079N02 gene 2310079N02Rik
1457775_at 0.99857 saccin Sacs
1457776_at 1.01665 "DNA segment, Chr 9, ERATO Doi 720, expressed" D9Ertd720e
1457777_at 1.05810 gasdermin 1 Gsdm1
1457778_at 0.96544 storkhead box 2 Stox2
1457779_at 0.76349 RIKEN cDNA 1110046J04 gene 1110046J04Rik
1457780_at 0.99026 syntaxin 11 Stx11
1457781_at 1.91062 KCNQ1 overlapping transcript 1 Kcnq1ot1
1457782_at 1.29791 talin 1 Tln1
1457783_at 1.21332 "RAB12, member RAS oncogene family" Rab12
1457784_at 0.92389 --- ---
1457785_at 0.95900 "Zinc finger, SWIM domain containing 3" Zswim3
1457786_at 1.13286 Sialic acid binding Ig-like lectin H Siglech
1457787_at 0.91262 --- ---
1457788_at 1.53070 Ras association (RalGDS/AF-6) domain family 1 Rassf1

1457789_at 1.13593 "ceroid lipofuscinosis, neuronal 3, juvenile (Batten, Spielmeier-Vogt disease)"
Cln3

1457790_at 0.88288 Ankyrin repeat and SOCS box-containing protein 3 Asb3

1457791_at 1.13552 RIKEN cDNA B830008H07 gene B830008H07Rik

1457792_at 1.13901 Limb region 1 like Lmbr11

1457793_a_at 1.43094 Wolf-Hirschhorn syndrome candidate 1-like 1 (human) Whsc111

1457794_at 1.20713 Wolf-Hirschhorn syndrome candidate 1-like 1 (human) Whsc111

1457795_at 1.05638 Secretory carrier membrane protein 4 Scamp4

1457796_at 1.03799 Ubiquitin protein ligase E3 component n-recognin 1 Ubr1

1457797_at 0.96529 expressed sequence AI605517 AI605517

1457798_at 1.08031 RIKEN cDNA C130039O16 gene C130039O16Rik

1457799_at 1.05593 "potassium voltage-gated channel, delayed-rectifier, subfamily S, member 3"
Kcns3

1457800_at 1.07422 RIKEN cDNA 2900019G14 gene 2900019G14Rik

1457801_at 1.01172 RIKEN cDNA 9930024M15 gene 9930024M15Rik

1457802_at 0.92733 RIKEN cDNA B930012P20 gene B930012P20Rik

1457803_at 1.32867 "Protein kinase, AMP-activated, gamma 1 non-catalytic subunit" Prkag1

1457804_at 0.95053 Syntaxin 8 Stx8

1457805_at 1.13388 --- ---

1457806_at 1.46675 dedicator of cyto-kinesis 1 Dock1

1457807_at 1.04328 Glucosamine-phosphate N-acetyltransferase 1 Gnpnat1

1457808_at 0.96254 hypothetical LOC432842 LOC432842

1457809_at 1.12589 p300/CBP-associated factor /// similar to p300/CBP-associated factor Pcaf ///
LOC330129

1457810_at 0.93373 zinc finger and BTB domain containing 37 Zbtb37

1457811_at 0.94149 --- ---

1457812_at 1.00639 transformation related protein 53 binding protein 1 Trp53bp1

1457813_at 1.35680 Mitochondrial ribosomal protein S18A Mrps18a

1457814_at 2.35972 RIKEN cDNA 1110014D18 gene 1110014D18Rik

1457815_at 1.20264 Nucleoporin 214 Nup214

1457816_at 0.98069 caspase 9 Casp9

1457817_at 1.12088 Breast carcinoma amplified sequence 3 Bcas3

1457818_at 1.09975 "General transcription factor II H, polypeptide 2" Gtf2h2

1457819_at 1.22606 "sodium channel, voltage-gated, type IX, alpha /// similar to sodium channel
25" Scn9a /// LOC671835

1457820_at 0.92002 --- ---

1457821_at 1.18826 RIKEN cDNA 2610511M17 gene 2610511M17Rik

1457822_at 0.92616 --- ---

1457823_at 0.82212 cysteine rich protein 61 Cyr61

1457824_at 1.25560 "Adult male bone cDNA, RIKEN full-length enriched library,
clone:9830165G24 product:unclassifiable, full insert sequence" ---

1457825_x_at 0.67516 transcobalamin 2 Tcn2

1457826_a_at 0.93300 RIKEN cDNA 1810043H04 gene 1810043H04Rik

1457827_at 1.03007 arylsulfatase J Arsj
1457828_at 1.04990 signal transducing adaptor molecule (SH3 domain and ITAM motif) 1 Stam
1457829_at 0.94245 calmegin Clgn
1457830_at 1.12390 Neuroepithelial cell transforming gene 1 Net1
1457831_at 0.98783 tripartite motif-containing 61 Trim61
1457832_at 1.06029 "7 days neonate cerebellum cDNA, RIKEN full-length enriched library, clone:
A730031J22 product:unclassifiable, full insert sequence" ---
1457833_at 0.95910 expressed sequence AI450241 AI450241
1457834_at 0.90328 YY1 transcription factor Yy1
1457835_at 1.11308 hypothetical protein C920025J10 C920025J10
1457836_at 1.30240 RIKEN cDNA 2600014M03 gene 2600014M03Rik
1457837_at 0.78615 cDNA sequence BC033606 /// similar to phospholipase B BC033606 ///
LOC665270
1457838_at 1.16742 cell division cycle 45 homolog (S. cerevisiae)-like Cdc451
1457839_at 1.50990 DEAH (Asp-Glu-Ala-His) box polypeptide 40 Dhx40
1457840_at 0.85615 plexin A4 Plxna4
1457841_at 1.01172 "In vitro fertilized eggs cDNA, RIKEN full-length enriched library,
clone:7420405G17 product:unclassifiable, full insert sequence" ---
1457842_at 2.94993 Zinc finger protein 292 Zfp292
1457843_at 0.98009 LY6/PLAUR domain containing 6 Lypd6
1457844_a_at 0.95763 RIKEN cDNA 8430427H17 gene 8430427H17Rik
1457845_at 1.37176 RIKEN cDNA 1110059P08 gene 1110059P08Rik
1457846_at 1.04397 "COX11 homolog, cytochrome c oxidase assembly protein (yeast)" Cox11
1457847_at 1.01423 Regulatory factor X domain containing 2 homolog (human) Rfxdc2
1457848_at 1.24988 RIKEN cDNA A930016P21 gene A930016P21Rik
1457849_at 1.02496 UDP-N-acetyl-alpha-D-galactosamine:polypeptide N-
acetylgalactosaminyltransferase 4 Galnt4
1457850_at 1.00174 "Solute carrier family 7 (cationic amino acid transporter, y+ system), member
14" Slc7a14
1457851_at 0.90607 "DNA segment, Chr 10, ERATO Doi 276, expressed" D10Ertd276e
1457852_at 1.19181 Latent transforming growth factor beta binding protein 1 Ltbp1
1457853_at 1.07676 CDNA sequence BC021785 BC021785
1457854_at 1.02307 expressed sequence AU018877 AU018877
1457855_at 0.95331 "retinol binding protein 3, interstitial" Rbp3
1457856_at 0.99645 --- ---
1457857_at 1.08121 --- ---
1457858_at 0.89411 similar to RNA and export factor binding protein 2 LOC226957
1457859_at 1.04860 ADP-ribosylation factor-like 15 Arl15
1457860_at 0.98659 --- ---
1457861_at 0.98778 RIKEN cDNA A230009B12 gene A230009B12Rik
1457862_at 1.02222 --- ---
1457863_at 1.02003 --- ---
1457864_at 0.98210 RAB11 family interacting protein 3 (class II) Rab11fip3

1457865_at 0.95858 "Pam, highwire, rpm 1" Phr1
1457866_at 0.94939 expressed sequence C77717 C77717
1457867_at 0.71229 sphingosine-1-phosphate phosphatase 2 Sgpp2
1457868_at 0.97340 "Solute carrier family 2 (facilitated glucose transporter), member 13" Slc2a13
1457869_at 1.31064 --- ---
1457870_at 1.02280 Transcribed locus ---
1457871_at 0.98073 collectin sub-family member 10 Colec10
1457872_at 0.94950 --- ---
1457873_at 1.00331 "11 days embryo gonad cDNA, RIKEN full-length enriched library, clone:7030407I18 product:unclassifiable, full insert sequence" ---
1457874_at 1.15842 "Adult male testis cDNA, RIKEN full-length enriched library, clone:4930550J05 product:unclassifiable, full insert sequence" ---
1457875_at 0.89586 host cell factor C1 Hcfc1
1457876_at 0.87251 "DNA segment, Chr 9, ERATO Doi 496, expressed" D9Ertd496e
1457877_at 0.99040 Estrogen receptor 1 (alpha) Esr1
1457878_at 0.88269 RIKEN cDNA C430042M11 gene C430042M11Rik
1457879_at 0.95887 RIKEN cDNA A230107O07 gene A230107O07Rik
1457880_at 0.92141 Transcribed locus ---
1457881_at 0.74648 oxysterol binding protein-like 6 Osbpl6
1457882_at 1.06430 RIKEN cDNA 5730466H23 gene 5730466H23Rik
1457883_at 1.00484 "Adult male aorta and vein cDNA, RIKEN full-length enriched library, clone: A530095B21 product:unclassifiable, full insert sequence" ---
1457884_at 1.12305 RIKEN cDNA 4732496C06 gene 4732496C06Rik
1457885_at 1.21130 "Protein kinase, AMP-activated, gamma 2 non-catalytic subunit" Prkag2
1457886_at 0.93702 hypothetical LOC552873 LOC552873
1457887_at 0.84008 SCO-spondin Sspo
1457888_at 1.77256 Polycomb group ring finger 5 Pcgf5
1457889_at 1.66586 expressed sequence AU014972 AU014972
1457890_at 0.91881 "DNA segment, Chr 6, ERATO Doi 234, expressed" D6Ertd234e
1457891_at 0.98047 RIKEN cDNA C230011H18 gene C230011H18Rik
1457892_at 0.93266 Transcribed locus ---
1457893_at 0.90657 Protein tyrosine phosphatase-like A domain containing 2 Ptplad2
1457894_at 0.94616 --- ---
1457895_at 0.90092 kinesin family member 18A Kif18a
1457896_at 1.00637 estrogen-related receptor gamma Esrrg
1457897_at 0.80700 Transcribed locus ---
1457898_at 0.92602 Dendritic cell protein GA17 Ga17
1457899_at 1.07585 "kalirin, RhoGEF kinase" Kalrn
1457900_at 0.93801 activating signal cointegrator 1 complex subunit 3 Ascc3
1457901_at 1.05021 "DNA segment, Chr 4, ERATO Doi 76, expressed" D4Ertd76e
1457902_at 0.86626 "Transcribed locus, moderately similar to XP_576460.1 PREDICTED: similar to hypothetical protein PB402898.00.0 [Rattus norvegicus]" ---
1457903_at 0.85756 Heme binding protein 1 Hebp1

1457904_at 0.93061 carbonic anhydrase 8 Car8
1457905_at 0.93241 "DNA segment, Chr 7, ERATO Doi 59, expressed" D7Ertd59e
1457906_at 0.99747 RNA binding motif protein 13 Rbm13
1457907_at 0.90678 NIMA (never in mitosis gene a)-related expressed kinase 7 Nek7
1457908_at 1.93275 Zinc finger protein 407 Zfp407
1457909_at 0.94671 DNA2 DNA replication helicase 2-like (yeast) Dna2l
1457910_at 1.38004 Formin-like 2 Fmnl2
1457911_at 1.12281 RIKEN cDNA A930002I21 gene A930002I21Rik
1457912_at 1.00345 RIKEN cDNA E130317F20 gene E130317F20Rik
1457913_at 1.41427 RIKEN cDNA 5730601F06 gene 5730601F06Rik
1457914_at 1.12268 "Adult retina cDNA, RIKEN full-length enriched library, clone:A930016I24
product:unclassifiable, full insert sequence" ---
1457915_at 1.15556 RIKEN cDNA 4833442J19 gene 4833442J19Rik
1457916_at 0.95447 --- ---
1457917_at 1.00212 Lymphocyte protein tyrosine kinase Lck
1457918_at 0.98738 "Tumor necrosis factor, alpha-induced protein 2" Tnfaip2
1457919_at 0.88860 RIKEN cDNA D030054H15 gene D030054H15Rik
1457920_at 1.01803 RIKEN cDNA C130026L21 gene C130026L21Rik
1457921_at 0.79749 RIKEN cDNA 1810037C20 gene 1810037C20Rik
1457922_at 1.36495 --- ---
1457923_at 0.98414 RIKEN cDNA C130045F17 gene C130045F17Rik
1457924_at 0.90800 muscleblind-like 1 (Drosophila) Mbnl1
1457925_at 1.06988 RIKEN cDNA C530049I24 gene C530049I24Rik
1457926_at 0.99248 arginyl aminopeptidase (aminopeptidase B)-like 1 Rnpepl1
1457927_at 1.04449 "transient receptor potential cation channel, subfamily M, member 3" Trpm3
1457928_at 0.98780 RIKEN cDNA 4930435E12 gene 4930435E12Rik
1457929_at 0.91108 transformed mouse 3T3 cell double minute 2 Mdm2
1457930_at 1.38669 "jumonji, AT rich interactive domain 1C (Rbp2 like)" Jarid1c
1457931_at 1.00473 --- ---
1457932_at 0.89684 Human immunodeficiency virus type I enhancer binding protein 2 Hivep2
1457933_at 1.05583 "Gene model 1964, (NCBI)" Gm1964
1457934_at 0.95808 RNA binding motif protein 12 Rbm12
1457935_at 1.23868 Teashirt zinc finger family member 2 Tshz2
1457936_at 0.94890 mitogen activated protein kinase 8 Mapk8
1457937_at 1.03297 RIKEN cDNA A430088P11 gene A430088P11Rik
1457938_at 0.96559 Nuclear receptor co-repressor 1 Ncor1
1457939_at 1.03266 RIKEN cDNA 4930578N16 gene 4930578N16Rik
1457940_at 0.89026 Ribosomal protein S17 Rps17
1457941_at 1.02560 "PREDICTED: Mus musculus similar to serine-threonine protein kinase pim-2
isoform 1 (LOC622647), mRNA" ---
1457942_at 0.88440 Transcribed locus ---
1457943_at 1.01196 RIKEN cDNA 1700011J10 gene 1700011J10Rik
1457944_at 1.74298 Ets variant gene 6 (TEL oncogene) Etv6

1457945_at 1.00300 "Eukaryotic translation initiation factor 2, subunit 3, structural gene Y-linked" Eif2s3y
1457946_at 0.86885 OG9 homeobox gene Og9x
1457947_at 0.99018 DEAD (Asp-Glu-Ala-Asp) box polypeptide 49 Ddx49
1457948_at 1.07436 Transcribed locus ---
1457949_at 0.88155 Transcribed locus ---
1457950_at 0.75444 Latrophilin 3 Lphn3
1457951_at 1.03311 RIKEN cDNA 4930434E21 gene 4930434E21Rik
1457952_at 0.98939 CD80 antigen Cd80
1457953_at 0.92003 "PREDICTED: Mus musculus hypothetical protein LOC629238 (LOC629238), mRNA" ---
1457954_at 0.95402 --- ---
1457955_at 1.08292 "Golgi associated, gamma adaptin ear containing, ARF binding protein 2" Gga2
1457956_at 0.97800 RIKEN cDNA 2310058O09 gene 2310058O09Rik
1457957_at 0.96445 Transcribed locus ---
1457958_at 1.28660 Ankyrin repeat domain 26 Ankrd26
1457959_at 0.99663 Transcribed locus ---
1457960_at 0.93965 Transcribed locus ---
1457961_at 0.80973 --- ---
1457962_at 1.00858 --- ---
1457963_at 0.87985 Transcribed locus ---
1457964_at 1.04696 RIKEN cDNA 1810044D09 gene 1810044D09Rik
1457965_at 1.01331 Transcribed locus ---
1457966_at 1.39350 F-box and leucine-rich repeat protein 7 Fbxl7
1457967_at 0.80169 RIKEN cDNA A030003K21 gene A030003K21Rik
1457968_at 1.12948 Transmembrane protein 23 Tmem23
1457969_at 1.12483 RAB interacting factor Rabif
1457970_at 1.17048 --- ---
1457971_at 1.09504 Centrosome and spindle pole associated protein 1 Csppl
1457972_at 0.88728 Expressed sequence C87436 C87436
1457973_at 1.04927 Protocadherin 9 Pcdh9
1457974_at 1.10057 --- ---
1457975_at 0.95100 GTP binding protein 3 Gtpbp3
1457976_at 1.13594 RIKEN cDNA 2010002M12 gene 2010002M12Rik
1457977_at 0.85833 RIKEN cDNA 2310005N01 gene 2310005N01Rik
1457978_at 1.04082 "Centaurin, gamma 2" Centg2
1457979_at 1.01137 "Adult male hypothalamus cDNA, RIKEN full-length enriched library, clone: A230017C18 product:unclassifiable, full insert sequence" ---
1457980_x_at 0.91502 uridine-cytidine kinase 2 Uck2
1457981_x_at 1.01384 GINS complex subunit 1 (Psf1 homolog) Gins1
1457982_at 0.89711 RIKEN cDNA 1700052M18 gene 1700052M18Rik
1457983_s_at 0.90837 RWD domain containing 4A Rwdd4a
1457984_at 0.93336 corticotropin releasing hormone Crh

1457985_at 0.91716 calmin Clmn
1457986_at 1.06821 Transcribed locus ---
1457987_at 1.40076 RIKEN cDNA 6030458C11 gene 6030458C11Rik
1457988_at 3.05369 SEC63-like (S. cerevisiae) Sec63
1457989_at 0.99691 "solute carrier family 4, sodium bicarbonate transporter-like, member 11"
Slc4a11
1457990_at 1.00595 ankyrin repeat and sterile alpha motif domain containing 1B /// similar to AIDA-
1b Anks1b /// LOC544718
1457991_at 1.22052 "ATP-binding cassette, sub-family D (ALD), member 3" Abcd3
1457992_at 1.04925 RNA terminal phosphate cyclase-like 1 Rcl1
1457993_at 1.07484 --- ---
1457994_at 0.97437 RIKEN cDNA 4930563J15 gene 4930563J15Rik
1457995_at 1.01675 RIKEN cDNA F730014I05 gene F730014I05Rik
1457996_at 1.06055 "Tumor necrosis factor, alpha-induced protein 8" Tnfaip8
1457997_at 1.07819 leukocyte receptor cluster (LRC) member 4 Leng4
1457998_at 1.07705 Transcribed locus ---
1457999_at 0.92055 RIKEN cDNA 2410005O16 gene 2410005O16Rik
1458000_at 1.02240 desmoglein 1 alpha Dsg1a
1458001_at 1.01043 --- ---
1458002_at 0.99550 Mitogen activated protein kinase 10 Mapk10
1458003_at 1.34339 zinc finger protein 398 Zfp398
1458004_at 0.94023 paired related homeobox 1 Prrx1
1458005_at 1.05503 --- ---
1458006_at 1.06157 Carboxypeptidase M Cpm
1458007_at 1.04855 myosin IB Myo1b
1458008_at 1.08662 "Adult male corpora quadrigemina cDNA, RIKEN full-length enriched library,
clone:B230326J17 product:unclassifiable, full insert sequence" ---
1458009_at 0.92837 Transcribed locus ---
1458010_at 1.00844 --- ---
1458011_s_at 0.97805 "DNA segment, Chr 4, ERATO Doi 117, expressed" D4ErtD117e
1458012_at 1.00603 dachshund 2 (Drosophila) Dach2
1458013_at 1.06620 RNA binding motif protein 27 Rbm27
1458014_at 1.02743 "AF4/FMR2 family, member 2" Aff2
1458015_at 0.97670 multiple EGF-like-domains 11 Megf11
1458016_at 1.01313 --- ---
1458017_at 1.34894 PHD finger protein 20 Phf20
1458018_at 1.07955 Protein phosphatase 1H (PP2C domain containing) Ppm1h
1458019_at 1.04168 --- ---
1458020_at 0.96173 Reelin Reln
1458021_at 1.25390 Sorting nexin 15 Snx15
1458022_at 0.99161 RIKEN cDNA D830039M14 gene D830039M14Rik
1458023_at 1.07464 "DNA segment, Chr X, ERATO Doi 697, expressed" DXErtD697e
1458024_at 1.13599 hypothetical protein B930018B01 B930018B01

1458025_at	1.00195	---	---	
1458026_at	0.98185	---	---	
1458027_at	1.04920	mitochondrial ribosomal protein L17	Mrpl17	
1458028_at	1.17135	Transcribed locus	---	
1458029_at	0.93820	---	---	
1458030_at	0.91989	---	---	
1458031_at	1.24376	RIKEN cDNA B230315F11 gene	B230315F11Rik	
1458032_at	0.96956	Tumor protein D52-like 2	Tpd52l2	
1458033_at	1.03093	Ubiquitin protein ligase E3 component n-recogin 1	Ubr1	
1458034_at	0.95580	RIKEN cDNA 1200016B10 gene	1200016B10Rik	
1458035_at	0.87115	Sphingosine-1-phosphate phosphatase 2	Sgpp2	
1458036_at	1.01006	Expressed sequence C77576	C77576	
1458037_at	1.06098	Neurocalcin delta	Ncald	
1458038_at	1.23116	---	---	
1458039_at	1.14595	Nuclear receptor coactivator 3	Ncoa3	
1458040_at	0.94643	"DNA segment, Chr 7, Wayne State University 130, expressed"	D7Wsu130e	
1458041_at	1.01175	expressed sequence C80360	C80360	
1458042_at	1.05297	---	---	
1458043_at	1.13339	BTB (POZ) domain containing 9	Btbd9	
1458044_at	0.96845	camello-like 1	Cml1	
1458045_at	0.95300	Odd Oz/ten-m homolog 4 (Drosophila)	Odz4	
1458046_at	0.93772	"0 day neonate kidney cDNA, RIKEN full-length enriched library, clone: D630027M10 product:unclassifiable, full insert sequence"	---	
1458047_at	1.00135	"tumor necrosis factor (ligand) superfamily, member 13b"	Tnfsf13b	
1458048_at	0.86889	hypothetical protein 6720406K03	6720406K03	
1458049_at	0.99662	---	---	
1458050_at	1.16251	---	---	
1458051_at	1.35483	RIKEN cDNA A230048O21 gene	A230048O21Rik	
1458052_at	0.99906	Transcribed locus	---	
1458053_at	1.21614	abl-interactor 2	Abi2	
1458054_at	2.15608	Exostoses (multiple) 1	Ext1	
1458055_at	0.95999	Desmocollin 3	Dsc3	
1458056_at	2.72913	"splicing factor, arginine/serine-rich 12"	Sfrs12	
1458057_at	2.22517	Caldesmon 1	Cald1	
1458058_at	1.11979	RIKEN cDNA 7030407E18 gene	7030407E18Rik	
1458059_at	1.00960	---	---	
1458060_at	0.94171	"10 days lactation, adult female mammary gland cDNA, RIKEN full-length enriched library, clone:D730006C24 product:unclassifiable, full insert sequence"	---	
1458061_at	0.99688	hypothetical LOC402769	LOC402769	
1458062_at	1.00326	Mitogen activated protein kinase 10	Mapk10	
1458063_at	0.96126	RIKEN cDNA 4930518F22 gene	4930518F22Rik	
1458064_at	0.99985	Ring finger protein 34	Rnf34	
1458065_at	1.10771	---	---	

1458066_at 0.90358 --- ---
1458067_at 1.03476 RIKEN cDNA A430041B07 gene A430041B07Rik
1458068_at 1.13323 RIKEN cDNA 4932417H02 gene 4932417H02Rik
1458069_at 2.18535 "TBC1 domain family, member 5" Tbc1d5
1458070_at 1.02335 V-set domain containing T cell activation inhibitor 1 Vtcn1
1458071_at 0.91348 Transcribed locus ---
1458072_at 1.04328 "12 days embryo spinal ganglion cDNA, RIKEN full-length enriched library, clone:D130082N20 product:unclassifiable, full insert sequence" ---
1458073_at 1.03988 RIKEN cDNA 9330156P08 gene 9330156P08Rik
1458074_at 1.03294 Regulator of sex-limitation candidate 24 Rslcan24
1458075_at 2.47950 Dystonin Dst
1458076_at 1.14920 --- ---
1458077_at 1.25315 "Bromodomain adjacent to zinc finger domain, 2B" Baz2b
1458078_at 1.61777 chromodomain helicase DNA binding protein 9 Chd9
1458079_at 0.97944 ubiquitin specific peptidase 40 Usp40
1458080_at 0.89996 senataxin Setx
1458081_at 0.92951 "13 days embryo heart cDNA, RIKEN full-length enriched library, clone: D330025O06 product:unclassifiable, full insert sequence" ---
1458082_at 0.90523 Cd200 antigen Cd200
1458083_at 1.22096 RIKEN cDNA 4932438A13 gene 4932438A13Rik
1458084_at 0.90234 "Zinc finger, DHHC domain containing 17" Zdhhc17
1458085_at 0.97014 --- ---
1458086_at 1.06807 RIKEN cDNA 4732471D19 gene 4732471D19Rik
1458087_at 0.95526 SH3 and cysteine rich domain 3 Stac3
1458088_at 0.90931 Ubiquitin-associated protein 2 Ubap2
1458089_at 1.07687 FK506 binding protein 5 Fkbp5
1458090_at 1.04412 RIKEN cDNA 4930504E06 gene 4930504E06Rik
1458091_at 0.97883 Dipeptidylpeptidase 10 Dpp10
1458092_at 1.00111 "Adaptor-related protein complex 3, mu 1 subunit" Ap3m1
1458093_at 0.93049 5-methyltetrahydrofolate-homocysteine methyltransferase reductase Mtrr
1458094_at 1.15290 Zinc finger protein 407 Zfp407
1458095_at 0.82556 "GTL2, imprinted maternally expressed untranslated mRNA" Gtl2
1458096_at 1.04140 Nance-Horan syndrome (human) Nhs
1458097_at 3.22608 Cobl-like 1 Cobll1
1458098_at 1.03497 "Adult male medulla oblongata cDNA, RIKEN full-length enriched library, clone:6330559K07 product:unclassifiable, full insert sequence" ---
1458099_at 2.59060 --- ---
1458100_at 0.89373 expressed sequence C87482 C87482
1458101_at 0.94274 leucine rich repeat containing 56 Lrrc56
1458102_at 0.93849 Plexin A2 Plxna2
1458103_at 0.99210 nuclear receptor co-repressor 1 Ncor1
1458104_a_at 0.89341 cDNA sequence BC038167 BC038167
1458105_at 0.88031 --- ---

1458106_at	1.07539	---	---	
1458107_at	0.87401	---	---	
1458108_at	1.10507	Pentatricopeptide repeat domain 2	Ptcd2	
1458109_at	1.02253	Hyaluronan and proteoglycan link protein 1	Hapln1	
1458110_at	1.26362	RIKEN cDNA D430030G11 gene	D430030G11Rik	
1458111_at	1.44216	RIKEN cDNA C530043G21 gene	C530043G21Rik	
1458112_at	0.88970	"adenosine deaminase, RNA-specific, B2"	Adarb2	
1458113_at	1.05740	RIKEN cDNA 9530019H20 gene	9530019H20Rik	
1458114_at	0.95738	sterile alpha motif domain containing 12	Samd12	
1458115_at	1.05537	RIKEN cDNA 4930512H18 gene	4930512H18Rik	
1458116_at	0.98706	RIKEN cDNA 9530079D04 gene	9530079D04Rik	
1458117_at	0.86739	GTPase activating RANGAP domain-like 1	Garnl1	
1458118_at	1.07766	"DNA segment, Chr 2, ERATO Doi 391, expressed"	D2Ertd391e	
1458119_at	0.97310	"solute carrier family 25, member 27"	Slc25a27	
1458120_at	0.86152	cadherin 18 /// RIKEN cDNA B230220E17 gene	Cdh18 /// B230220E17Rik	
1458121_at	1.04207	RIKEN cDNA A430107O13 gene	A430107O13Rik	
1458122_at	1.11814	"7 days neonate cerebellum cDNA, RIKEN full-length enriched library, clone: A730016A03 product:unclassifiable, full insert sequence"	---	
1458123_at	1.00782	RIKEN cDNA 9630002A11 gene	9630002A11Rik	
1458124_at	1.06999	IQ motif containing GTPase activating protein 1	Iqgap1	
1458125_at	1.26911	Expressed sequence C80913	C80913	
1458126_at	1.38905	RIKEN cDNA D030041H20 gene	D030041H20Rik	
1458127_at	1.00893	Cell adhesion molecule-related/down-regulated by oncogenes	Cdon	
1458128_at	1.04685	RIKEN cDNA 1110028C15 gene	1110028C15Rik	
1458129_at	0.91494	RAR-related orphan receptor alpha	Rora	
1458130_at	1.02940	RIKEN cDNA 1110001A07 gene	1110001A07Rik	
1458131_at	0.93081	---	---	
1458132_at	1.06006	cyclin M2	Cnm2	
1458133_at	0.94891	---	---	
1458134_at	1.40716	Glutathione transferase zeta 1 (maleylacetoacetate isomerase)	Gstz1	
1458135_at	1.31024	RIKEN cDNA G630014P10 gene	G630014P10Rik	
1458136_at	0.97665	Musashi homolog 2 (Drosophila)	Msi2	
1458137_at	0.90720	Transcribed locus	---	
1458138_at	0.94189	Ring finger protein 190	Rnf190	
1458139_at	0.97974	---	---	
1458140_at	1.02381	slit homolog 2 (Drosophila)	Slit2	
1458141_at	0.95927	---	---	
1458142_at	1.07928	"zinc finger, DHHC domain containing 9"	Zdhhc9	
1458143_at	0.98313	"ADAM metallopeptidase with thrombospondin type 1 motif, 3"	Adamts3	
1458144_at	1.07219	CDNA sequence BC031353	BC031353	
1458145_at	1.21903	RAR-related orphan receptor alpha	Rora	
1458146_at	0.86107	RIKEN cDNA D030011O10 gene	D030011O10Rik	
1458147_at	1.00664	MAM domain containing 1	Mamdc1	

1458148_at	0.96020	RIKEN cDNA D230007K08 gene	D230007K08Rik
1458149_at	1.08296	---	---
1458150_at	0.98625	RIKEN cDNA D030051N19 gene	D030051N19Rik
1458151_at	1.20692	RIKEN cDNA 4833444G19 gene	4833444G19Rik
1458152_at	1.01736	"electron transferring flavoprotein, dehydrogenase"	Etfdh
1458153_at	0.89157	Peroxisomal biogenesis factor 14	Pex14
1458154_at	0.97851	Transcribed locus	---
1458155_at	0.93046	RAB GTPase activating protein 1-like	Rabgap11
1458156_at	2.16126	RIKEN cDNA E230012J19 gene	E230012J19Rik
1458157_at	1.10875	NIMA (never in mitosis gene a)-related expressed kinase 8	Nek8
1458158_at	1.06810	WD repeat domain 67	Wdr67
1458159_at	1.78911	"Guanine nucleotide binding protein, alpha q polypeptide"	Gnaq
1458160_at	0.93202	"Adult male testis cDNA, RIKEN full-length enriched library, clone:4933400K21 product:unclassifiable, full insert sequence"	---
1458161_at	1.65853	KCNQ1 overlapping transcript 1	Kcnq1ot1
1458162_at	0.99033	"Adult male hippocampus cDNA, RIKEN full-length enriched library, clone: C630034F03 product:unclassifiable, full insert sequence"	---
1458163_at	1.02688	cDNA sequence BC066028	BC066028
1458164_at	0.93653	hypothetical LOC552913	LOC552913
1458165_at	0.99192	importin 7	Ipo7
1458166_at	1.12600	PHD finger protein 21A	Phf21a
1458167_at	0.97375	---	---
1458168_at	1.05626	"Adult male adrenal gland cDNA, RIKEN full-length enriched library, clone:7330440N08 product:hypothetical protein, full insert sequence"	---
1458169_at	0.90271	"gene model 700, (NCBI) /// similar to Motilin receptor (G-protein coupled receptor 38) /// similar to platelet endothelial aggregation receptor 1"	Gm700 /// LOC668482 /// LOC675947
1458170_at	1.02586	---	---
1458171_at	0.97592	"solute carrier organic anion transporter family, member 2b1"	Slco2b1
1458172_at	1.16154	"malic enzyme 2, NAD(+)-dependent, mitochondrial"	Me2
1458173_at	1.07783	"Centaurin, delta 1"	Centd1
1458174_at	0.97122	coiled-coil domain containing 52	Ccdc52
1458175_at	0.83637	"spondin 2, extracellular matrix protein"	Spon2
1458176_at	1.48556	Period homolog 3 (Drosophila)	Per3
1458177_at	0.99146	emopamil binding protein-like	Ebpl
1458178_at	1.02804	RIKEN cDNA 1700012H17 gene	1700012H17Rik
1458179_at	1.42261	---	---
1458180_at	1.29179	---	---
1458181_at	1.05933	RIKEN cDNA 9030624J02 gene	9030624J02Rik
1458182_at	1.64195	RNA binding motif protein 18	Rbm18
1458183_at	1.16462	RIKEN cDNA 2810039B14 gene	2810039B14Rik
1458184_at	0.98594	RIKEN cDNA D030020J04 gene	D030020J04Rik
1458185_at	1.19309	THAP domain containing 4	Thap4

1458186_at 1.00170 Transcribed locus ---
1458187_at 1.16047 zinc finger protein 142 Zfp142
1458188_at 0.89947 Dihydropyrimidinase-like 2 Dpysl2
1458189_at 0.95246 EMI domain containing 2 Emid2
1458190_at 0.99174 Rho GTPase activating protein 4 Arhgap4
1458191_at 0.81336 forkhead box P2 Foxp2
1458192_at 1.01301 RIKEN cDNA C130065N10 gene C130065N10Rik
1458193_at 0.96645 peripheral myelin protein 2 Pmp2
1458194_at 0.96959 --- ---
1458195_at 1.19205 vacuolar protein sorting 13 D (yeast) Vps13d
1458196_at 1.01974 Glucosamine-phosphate N-acetyltransferase 1 Gnpnat1
1458197_x_at 0.91251 MAP kinase-activated protein kinase 5 /// a disintegrin and metallopeptidase
domain 1a Mapkapk5 /// Adam1a
1458198_at 1.04518 expressed sequence C81508 C81508
1458199_at 1.34263 Activating signal cointegrator 1 complex subunit 2 Ascc2
1458200_at 1.54603 BTB (POZ) domain containing 14B Btbd14b
1458201_at 1.02102 --- ---
1458202_at 0.85751 RIKEN cDNA 6330500D04 gene 6330500D04Rik
1458203_at 1.25340 Spire homolog 1 (Drosophila) Spire1
1458204_at 1.00768 secreted acidic cysteine rich glycoprotein Sparc
1458205_at 1.10833 --- ---
1458206_at 0.98175 --- ---
1458207_at 1.00797 "Zinc finger, BED domain containing 3" Zbed3
1458208_s_at 1.13186 --- ---
1458209_at 1.04587 Kinase non-catalytic C-lobe domain (KIND) containing 1 Kndc1
1458210_at 1.01686 RIKEN cDNA 9130404H23 gene 9130404H23Rik
1458211_at 0.96421 NIMA (never in mitosis gene a)-related expressed kinase 1 Nek1
1458212_at 1.02509 "Phosphodiesterase 4D, cAMP specific" Pde4d
1458213_at 1.10655 Nuclear factor of activated T-cells 5 Nfat5
1458214_at 0.91965 RIKEN cDNA A730012O14 gene A730012O14Rik
1458215_at 1.28256 "SET domain, bifurcated 1" Setdb1
1458216_at 1.03655 Transcribed locus ---
1458217_s_at 1.08505 RIKEN cDNA 2210018M11 gene 2210018M11Rik
1458218_s_at 1.02356 phosphodiesterase 7A Pde7a
1458219_at 0.89331 breast carcinoma amplified sequence 2 Bcas2
1458220_at 1.07364 deleted in liver cancer 1 Dlc1
1458221_at 1.02256 RIKEN cDNA G630014P10 gene G630014P10Rik
1458222_at 1.34940 Transcribed locus ---
1458223_at 0.82744 transformed mouse 3T3 cell double minute 4 /// similar to transformed mouse
3T3 cell double minute 4 /// similar to transformed mouse 3T3 cell double minute 4 Mdm4 ///
LOC546376 /// LOC630948
1458224_at 1.15136 Leucine rich repeat containing 2 Lrrc2
1458225_at 1.13997 FK506 binding protein 14 Fkbp14

1458226_at 0.93260 "filamin, beta" Flnb
 1458227_at 0.98449 --- ---
 1458228_at 1.02055 Opioid binding protein/cell adhesion molecule-like Opcml
 1458229_at 0.99359 roundabout homolog 2 (Drosophila) Robo2
 1458230_at 0.98997 "Glutamate receptor, ionotropic, AMPA4 (alpha 4)" Gria4
 1458231_at 0.91562 "PREDICTED: Mus musculus similar to Zinc finger protein 195 (LOC625519), mRNA" ---
 1458232_at 0.97100 dickkopf homolog 1 (Xenopus laevis) Dkk1
 1458233_at 1.43252 RIKEN cDNA 9030227G01 gene 9030227G01Rik
 1458234_at 0.81666 RIKEN cDNA A730052K04 gene A730052K04Rik
 1458235_at 1.17955 A kinase (PRKA) anchor protein (gravin) 12 Akap12
 1458236_at 0.97959 "12 days embryo eyeball cDNA, RIKEN full-length enriched library, clone: D230015O18 product:hypothetical protein, full insert sequence" ---
 1458237_at 0.95554 --- ---
 1458238_at 1.06157 AT rich interactive domain 5B (Mrf1 like) Arid5b
 1458239_at 1.04744 --- ---
 1458240_at 1.08071 "Membrane associated guanylate kinase, WW and PDZ domain containing 1" Magi1
 1458241_at 1.14798 High mobility group AT-hook 2 Hmga2
 1458242_at 0.93452 Gephyrin Gphn
 1458243_at 1.00348 --- ---
 1458244_at 0.86477 transmembrane and coiled-coil domains 5 Tmco5
 1458245_at 1.01017 similar to sushi domain containing 1 LOC634731
 1458246_at 1.07279 --- ---
 1458247_s_at 1.00167 dynactin 5 Dctn5
 1458248_at 0.84545 --- ---
 1458249_at 1.13996 --- ---
 1458250_at 1.17463 RIKEN cDNA 1200015N20 gene 1200015N20Rik
 1458251_at 0.96129 expressed sequence AU022166 AU022166
 1458252_at 1.04074 "Mus musculus, Similar to RIKEN cDNA B430319H21 gene, clone IMAGE:3991101, mRNA" ---
 1458253_at 0.88651 "Adult male bone cDNA, RIKEN full-length enriched library, clone:9830128M12 product:unclassifiable, full insert sequence" ---
 1458254_at 1.06321 RIKEN cDNA D330027H18 gene D330027H18Rik
 1458255_at 0.99846 WD repeat domain 33 Wdr33
 1458256_at 0.98729 RIKEN cDNA 5830446M03 gene 5830446M03Rik
 1458257_at 1.37697 --- ---
 1458258_at 0.98560 --- ---
 1458259_x_at 1.00110 "COP9 (constitutive photomorphogenic) homolog, subunit 3 (Arabidopsis thaliana)" Cops3
 1458260_at 0.99624 Espin Espn
 1458261_at 1.23793 Casitas B-lineage lymphoma-like 1 Cbl1
 1458262_at 0.91132 Vexonuclease 3'-5' domain-like 1 Exd1

1458263_at 1.39576 "CUG triplet repeat, RNA binding protein 2" Cugbp2
1458264_at 0.99319 RIKEN cDNA 4930412F12 gene 4930412F12Rik
1458265_at 0.98795 Kinesin 2 Kns2
1458266_at 1.00113 Mitotic arrest deficient 1-like 1 Mad111
1458267_at 1.01956 zinc finger protein 248 Zfp248
1458268_s_at 0.71803 insulin-like growth factor binding protein 3 Igfbp3
1458269_at 0.82782 protocadherin 9 /// similar to protocadherin 9 isoform 1 precursor Pcdh9 ///
LOC638275
1458270_at 0.99358 --- ---
1458271_at 1.13750 "12 days embryo spinal cord cDNA, RIKEN full-length enriched library, clone:
C530003P15 product:unclassifiable, full insert sequence" ---
1458272_at 1.10576 Transcribed locus ---
1458273_at 0.96186 Transcribed locus ---
1458274_at 0.92665 zinc finger protein 69 Zfp69
1458275_at 1.06117 RIKEN cDNA 2310045A20 gene 2310045A20Rik
1458276_x_at 1.12924 citron Cit
1458277_at 1.15299 Chemokine (C-C motif) ligand 25 Ccl25
1458278_at 0.84724 --- ---
1458279_at 1.23871 Forkhead box O3a Foxo3a
1458280_at 1.40912 Activating signal cointegrator 1 complex subunit 2 Ascc2
1458281_at 0.91748 Arylsulfatase B Arsb
1458282_at 1.45790 --- ---
1458283_at 0.95215 RIKEN cDNA 4931406P16 gene 4931406P16Rik
1458284_at 0.90645 polypyrimidine tract binding protein 1 Ptbp1
1458285_at 1.02966 "Glutamate receptor, ionotropic, AMPA1 (alpha 1)" Gria1
1458286_at 0.93694 RIKEN cDNA 0610007P14 gene 0610007P14Rik
1458287_at 0.83405 Transcribed locus ---
1458288_at 1.00227 RIKEN cDNA 4732475C15 gene 4732475C15Rik
1458289_at 1.06160 UDP-N-acetyl-alpha-D-galactosamine:polypeptide N-
acetylgalactosaminyltransferase-like 4 Galntl4
1458290_at 1.90576 R3H domain 1 (binds single-stranded nucleic acids) R3hdm1
1458291_at 0.99874 Inhibin beta-A Inhba
1458292_at 1.18433 "proteasome (prosome, macropain) subunit, alpha type 1" Psma1
1458293_at 1.09617 RIKEN cDNA 5730403B10 gene 5730403B10Rik
1458294_at 1.14991 Transcribed locus ---
1458295_at 0.91375 cDNA sequence BC038331 BC038331
1458296_at 2.19604 Exostoses (multiple) 1 Ext1
1458297_s_at 1.01709 macrophage receptor with collagenous structure Marco
1458298_at 1.01502 Ca<2+>dependent activator protein for secretion Cadps
1458299_s_at 1.30663 "nuclear factor of kappa light polypeptide gene enhancer in B-cells inhibitor,
epsilon" Nfkbie
1458300_at 0.98914 "aldo-keto reductase family 7, member A5 (aflatoxin aldehyde reductase)"
Akr7a5

1458301_x_at 1.03780 Zinc finger protein 292 Zfp292
1458302_at 1.01447 Transcribed locus ---
1458303_at 1.08218 "gene model 96, (NCBI)" Gm96
1458304_at 0.98009 MAD homolog 2 (Drosophila) Smad2
1458305_at 0.94834 --- ---
1458306_at 0.99479 "phosphatidylinositol glycan anchor biosynthesis, class Z" Pigz
1458307_at 1.03320 RIKEN cDNA B230334C09 gene B230334C09Rik
1458308_at 2.16084 strawberry notch homolog (Drosophila) Stno
1458309_at 2.08241 Transcribed locus ---
1458310_at 0.88459 RIKEN cDNA A230056J06 gene A230056J06Rik
1458311_at 1.09351 ubiquitin specific peptidase 36 Usp36
1458312_at 0.94076 "Calcium channel, voltage-dependent, P/Q type, alpha 1A subunit" Cacna1a
1458313_at 0.87423 Cell division cycle 2-like 5 (cholinesterase-related cell division controller)
Cdc215
1458314_at 0.98760 "Solute carrier family 1 (glial high affinity glutamate transporter), member 2"
Slc1a2
1458315_at 1.00392 enhancer of polycomb homolog 1 (Drosophila) Epc1
1458316_at 1.05153 Transcribed locus ---
1458317_at 1.07703 Echinoderm microtubule associated protein like 1 Eml1
1458318_at 1.44423 RIKEN cDNA 3110047M12 gene 3110047M12Rik
1458319_at 0.89674 G protein-coupled receptor 177 Gpr177
1458320_at 1.07143 RIKEN cDNA 4932431H17 gene /// similar to RIKEN cDNA 4932431H17
gene 4932431H17Rik /// LOC635623
1458321_at 0.99388 phosphatidylinositol 3-kinase catalytic delta polypeptide Pik3cd
1458322_x_at 0.99233 RIKEN cDNA E230008N13 gene E230008N13Rik
1458323_at 0.96704 RIKEN cDNA 4930444A19 gene 4930444A19Rik
1458324_x_at 1.07021 "16 days embryo head cDNA, RIKEN full-length enriched library, clone:
C130051K14 product:unclassifiable, full insert sequence" ---
1458325_x_at 1.15505 Bcl2 modifying factor Bmf
1458326_at 1.04051 polymerase (RNA) II (DNA directed) polypeptide E Polr2e
1458327_x_at 0.99048 "solute carrier family 26 (sulfate transporter), member 1" Slc26a1
1458328_x_at 0.97730 SH3-domain GRB2-like (endophilin) interacting protein 1 Sgip1
1458329_x_at 1.05699 RIKEN cDNA 1110051B16 gene 1110051B16Rik
1458330_x_at 0.87663 Cyclin M3 Cnm3
1458331_x_at 0.95951 carboxylesterase 7 Ces7
1458332_x_at 0.93917 SRY-box containing gene 4 Sox4
1458333_x_at 0.93954 undifferentiated embryonic cell transcription factor 1 Utf1
1458334_at 1.00942 wingless-related MMTV integration site 7A Wnt7a
1458335_x_at 1.52583 RIKEN cDNA 2900073H19 gene 2900073H19Rik
1458336_at 0.94998 RIKEN cDNA 2310035C23 gene 2310035C23Rik
1458337_at 1.15472 Transcription factor 12 Tcf12
1458338_x_at 0.76140 "nuclear receptor subfamily 2, group C, member 1" Nr2c1
1458339_at 1.10253 cytidine and dCMP deaminase domain containing 1 Cdadc1

1458340_at 0.93059 Transcribed locus ---
 1458341_x_at 0.78689 --- ---
 1458342_at 1.03826 transmembrane protein 90a Tmem90a
 1458343_x_at 1.02895 "mediator of RNA polymerase II transcription, subunit 6 homolog (yeast)"
 Med6
 1458344_at 0.97953 N-acetylglucosamine-1-phosphodiester alpha-N-acetylglucosaminidase
 Nagpa
 1458345_s_at 0.99355 collectin sub-family member 11 Colec11
 1458346_x_at 1.11874 "ATPase, Ca⁺⁺ transporting, plasma membrane 1" Atp2b1
 1458347_s_at 0.98275 "transmembrane protease, serine 2" Tmprss2
 1458348_at 1.00012 Transcribed locus ---
 1458349_s_at 1.02152 "10 days neonate skin cDNA, RIKEN full-length enriched library,
 clone:4732488M12 product:unclassifiable, full insert sequence" ---
 1458350_at 0.91260 Optic atrophy 3 (human) Opa3
 1458351_s_at 0.96620 "kelch-like 2, Mayven (Drosophila)" Klh2
 1458352_at 1.11212 RIKEN cDNA A830039H10 gene A830039H10Rik
 1458353_at 0.97063 RIKEN cDNA A230063L24 gene A230063L24Rik
 1458354_x_at 1.04281 keratin 28 Krt28
 1458355_x_at 0.95464 expressed sequence AI553587 AI553587
 1458356_at 0.92093 --- ---
 1458357_x_at 0.92542 Prostate tumor over expressed gene 1 Ptov1
 1458358_at 1.10982 pantothenate kinase 2 (Hallervorden-Spatz syndrome) Pank2
 1458359_at 0.95244 Transcribed locus ---
 1458360_at 0.95608 "Sparc/osteonectin, cwcv and kazal-like domains proteoglycan 2" Spock2
 1458361_at 0.91617 "DNA cross-link repair 1C, PSO2 homolog (S. cerevisiae)" Dclre1c
 1458362_at 0.98876 RIKEN cDNA D730005E14 gene D730005E14Rik
 1458363_at 1.08666 "zinc finger, DHHC domain containing 17" Zdhhc17
 1458364_s_at 0.96522 interferon alpha responsive gene Ifrg15
 1458365_at 0.86900 SCY1-like 1 binding protein 1 Scyl1bp1
 1458366_at 0.88309 Additional sex combs like 2 (Drosophila) Asxl2
 1458367_at 1.01535 "gene model 94, (NCBI)" Gm94
 1458368_at 0.99191 "myosin, heavy polypeptide 4, skeletal muscle" Myh4
 1458369_at 1.10729 RIKEN cDNA 2310050B05 gene 2310050B05Rik
 1458370_at 0.99359 BMP2 inducible kinase Bmp2k
 1458371_at 1.08503 RIKEN cDNA E130006N16 gene E130006N16Rik
 1458372_at 0.98326 transmembrane channel-like 8 Tmc8
 1458373_at 1.11948 RIKEN cDNA 5830483C08 gene 5830483C08Rik
 1458374_at 1.01295 expressed sequence C79407 C79407
 1458375_at 1.07757 RIKEN cDNA 6430524H05 gene 6430524H05Rik
 1458376_at 2.83702 RIKEN cDNA B930025B16 gene B930025B16Rik
 1458377_at 1.01678 "cofactor required for Sp1 transcriptional activation, subunit 7" Crsp7
 1458378_at 0.99234 "glutamate receptor ionotropic, NMDA3A" Grin3a
 1458379_at 0.97575 RIKEN cDNA 9330159F19 gene 9330159F19Rik

1458380_at	0.85568	additional sex combs like 1 (Drosophila)	Asxl1
1458381_at	0.77899	chloride intracellular channel 5	Clic5
1458382_a_at	1.26951	zinc fingers and homeoboxes protein 1	Zhx1
1458383_at	1.08420	"Isocitrate dehydrogenase 2 (NADP+), mitochondrial"	Idh2
1458384_at	1.08472	Transcribed locus	---
1458385_at	1.38038	heat shock protein 4 like	Hspa4l
1458386_at	1.02747	RIKEN cDNA C030009J22 gene	C030009J22Rik
1458387_at	0.97519	phosphatidylserine synthase 2	Ptdss2
1458388_at	0.99760	"gene model 172, (NCBI)"	Gm172
1458389_at	1.03226	Kruppel-like factor 7 (ubiquitous)	Klf7
1458390_at	1.01510	anaphase promoting complex subunit 10	Anapc10
1458391_at	1.03592	---	---
1458392_at	0.89729	Expressed sequence C87414	C87414
1458393_at	1.03626	serine racemase	Srr
1458394_at	1.02875	"solute carrier family 6 (neurotransmitter transporter), member 17"	Slc6a17
1458395_at	0.85683	hypothetical protein B930054O08	B930054O08
1458396_at	0.91231	"purinergic receptor P2X, ligand-gated ion channel, 3"	P2rx3
1458397_at	0.97198	hypothetical LOC622208	LOC622208
1458398_at	0.98593	RIKEN cDNA 1700027F06 gene	1700027F06Rik
1458399_at	0.98930	leucine rich repeat containing 3	Lrrc3
1458400_at	0.95852	RIKEN cDNA 9630050P21 gene	9630050P21Rik
1458401_at	1.05365	RIKEN cDNA 4932438A13 gene	/// similar to CG15133-PA
CG15133-PA	4932438A13Rik	/// LOC671298	/// LOC676099
1458402_at	0.91817	6-pyruvoyl-tetrahydropterin synthase	Pts
1458403_at	0.98598	TRAF2 and NCK interacting kinase	/// similar to [Segment 1 of 2] Traf2 and NCK-interacting kinase Tnik
1458404_at	0.95803	NADH dehydrogenase (ubiquinone) 1 beta subcomplex 8	Ndufb8
1458405_at	1.10646	hypothetical protein B430201F14	B430201F14
1458406_at	0.94479	Expressed sequence AI429294	AI429294
1458407_s_at	1.10497	Expressed sequence AI429294	AI429294
1458408_at	0.96813	sterile alpha motif domain containing 8	Samd8
1458409_at	0.83616	expressed sequence C86595	C86595
1458410_at	1.05719	GTPase activating RANGAP domain-like 1	Garnl1
1458411_at	1.03266	kinase non-catalytic C-lobe domain (KIND) containing 1	Kndc1
1458412_at	0.83349	zinc finger protein 78	Zfp78
1458413_at	1.02312	F-box and WD-40 domain protein 8	Fbxw8
1458414_at	1.38916	"DNA segment, Chr 2, ERATO Doi 93, expressed"	D2Ertd93e
1458415_at	0.90581	"C-type lectin domain family 2, member e"	Clec2e
1458416_at	0.91553	RIKEN cDNA A330048O09 gene	A330048O09Rik
1458417_at	0.95233	jumonji domain containing 1B	Jmjd1b
1458418_at	1.14303	"Mus musculus, clone IMAGE:5368497, mRNA"	---
1458419_at	1.00991	RIKEN cDNA E130215H24 gene	E130215H24Rik
1458420_at	0.75695	RIKEN cDNA 9430057O19 gene	9430057O19Rik

1458421_at 0.99643 "potassium voltage-gated channel, subfamily Q, member 3" Kcnq3
1458422_at 0.98581 RIKEN cDNA D730002M21 gene D730002M21Rik
1458423_at 1.05940 LUC7-like 2 (*S. cerevisiae*) Luc712
1458424_at 0.91709 Fatty acid 2-hydroxylase Fa2h
1458425_at 0.88320 "16 days neonate thymus cDNA, RIKEN full-length enriched library, clone:
A130088J11 product:unclassifiable, full insert sequence" ---
1458426_at 1.07484 Kinesin family member 1B Kif1b
1458427_at 1.10634 BRCA1 interacting protein C-terminal helicase 1 Brip1
1458428_at 0.95035 "potassium channel, subfamily T, member 1" Kcnt1
1458429_at 0.97950 Rho GTPase activating protein 30 Arhgap30
1458430_at 1.23790 expressed sequence C87436 C87436
1458431_at 0.89383 "eukaryotic translation initiation factor 4, gamma 2" Eif4g2
1458432_at 1.66061 Non-catalytic region of tyrosine kinase adaptor protein 2 Nck2
1458433_at 0.89825 "RASD family, member 2" Rasd2
1458434_at 1.04448 RIKEN cDNA A730009L09 gene A730009L09Rik
1458435_at 0.99776 RIKEN cDNA 2010000I03 gene 2010000I03Rik
1458436_at 1.01695 AU RNA binding protein/enoyl-coenzyme A hydratase Auh
1458437_at 0.95418 Transcribed locus ---
1458438_at 0.83756 RIKEN cDNA 4933415L06 gene 4933415L06Rik
1458439_a_at 1.04689 RIKEN cDNA 2310047C04 gene 2310047C04Rik
1458440_at 1.14222 spectrin domain with coiled-coils 1 Specc1
1458441_at 1.12268 "solute carrier family 6 (neurotransmitter transporter), member 17" Slc6a17
1458442_at 1.00826 expressed sequence AI132709 AI132709
1458443_at 0.97387 CREB regulated transcription coactivator 1 Crtc1
1458444_at 0.92987 Expressed sequence AI195381 AI195381
1458445_at 0.93217 RIKEN cDNA 6530409C15 gene 6530409C15Rik
1458446_at 0.94291 RIKEN cDNA 3222401L13 gene 3222401L13Rik
1458447_at 0.86248 centromere protein F Cenpf
1458448_at 0.92426 expressed sequence AA408296 AA408296
1458449_at 1.12184 F-box and leucine-rich repeat protein 11 Fbxl11
1458450_at 2.07191 zinc finger RNA binding protein Zfr
1458451_at 1.07449 "Sno, strawberry notch homolog 1 (*Drosophila*)" Sbnol
1458452_at 1.87046 ankyrin repeat domain 11 Ankrd11
1458453_at 1.11506 LIM domain only 7 Lmo7
1458454_at 0.88369 Transcribed locus ---
1458455_at 0.91848 actin-binding Rho activating protein Abra
1458456_x_at 1.06163 RIKEN cDNA 6430571L13 gene 6430571L13Rik
1458457_at 0.91113 --- ---
1458458_at 0.98760 schlafen 5 Slfn5
1458459_a_at 0.89309 RIKEN cDNA E230029C05 gene /// expressed sequence AI452102 ///
hypothetical protein LOC671286 E230029C05Rik /// AI452102 /// LOC671286
1458460_at 1.19852 Stromal interaction molecule 1 Stim1
1458461_at 1.03959 RIKEN cDNA E330021D16 gene E330021D16Rik

1458462_at 1.03393 Rho GTPase activating protein 30 Arhgap30
1458463_at 0.96734 --- ---
1458464_at 1.07443 "HECT, C2 and WW domain containing E3 ubiquitin protein ligase 2"
Hecw2
1458465_at 1.10774 PAN3 polyA specific ribonuclease subunit homolog (S. cerevisiae) Pan3
1458466_at 0.98977 protective protein for beta-galactosidase Ppgb
1458467_at 0.88556 RIKEN cDNA 1110032D16 gene 1110032D16Rik
1458468_at 1.11884 Coenzyme Q10 homolog A (yeast) Coq10a
1458469_at 1.08441 Casitas B-lineage lymphoma b Cblb
1458470_at 0.95797 --- ---
1458471_at 1.05471 WW domain containing adaptor with coiled-coil Wac
1458472_at 1.01330 RIKEN cDNA 4932417H02 gene 4932417H02Rik
1458473_at 1.05304 "POU domain, class 6, transcription factor 2" Pou6f2
1458474_at 1.20620 --- ---
1458475_at 1.13177 "Bruno-like 4, RNA binding protein (Drosophila)" Brunol4
1458476_at 1.10565 "eukaryotic translation initiation factor 2C, 3" Eif2c3
1458477_at 1.08877 "3 days neonate thymus cDNA, RIKEN full-length enriched library, clone:
A630098N09 product:unclassifiable, full insert sequence" ---
1458478_at 0.85900 RIKEN cDNA 9230110F11 gene 9230110F11Rik
1458479_at 1.16296 SMC2 structural maintenance of chromosomes 2-like 1 (yeast) Smc2l1
1458480_at 1.17388 topoisomerase (DNA) II beta binding protein Topbp1
1458481_at 0.90567 "gene model 614, (NCBI)" Gm614
1458482_at 0.98681 TNNI3 interacting kinase Tnni3k
1458483_at 1.06105 RIKEN cDNA A330102I10 gene A330102I10Rik
1458484_at 1.05223 RIKEN cDNA A730020M07 gene A730020M07Rik
1458485_at 0.93468 F-box and leucine-rich repeat protein 7 Fbxl7
1458486_at 1.27819 --- ---
1458487_at 0.92565 Kruppel-like factor 3 (basic) /// RIKEN cDNA C230096K16 gene Klf3 ///
C230096K16Rik
1458488_at 0.91611 doublecortin and CaM kinase-like 2 Dcamkl2
1458489_at 1.38911 Transcribed locus ---
1458490_at 0.98673 RIKEN cDNA 9230116N13 gene 9230116N13Rik
1458491_at 0.84696 RIKEN cDNA 4930422I07 gene 4930422I07Rik
1458492_x_at 0.93761 neurotrimin Hnt
1458493_a_at 1.08550 --- ---
1458494_at 0.91849 Transcribed locus ---
1458495_at 0.93242 "Protein phosphatase 3, catalytic subunit, alpha isoform" Ppp3ca
1458496_at 1.10458 Cytochrome b-561 Cyb561
1458497_at 2.47012 Cysteine rich transmembrane BMP regulator 1 (chordin like) Crim1
1458498_at 1.01257 --- ---
1458499_at 0.97679 phosphodiesterase 10A Pde10a
1458500_at 1.01288 expressed sequence AU021034 AU021034
1458501_at 0.98817 "vesicle-associated membrane protein, associated protein B and C" Vapb

1458502_at	0.91525	Kinectin 1	Ktn1	
1458503_at	1.02131	---	---	
1458504_at	0.94044	zinc finger CCCH type containing 12D	Zc3h12d	
1458505_at	0.86927	hypothetical LOC552901	LOC552901	
1458506_at	1.05159	"gene model 1582, (NCBI)"	Gm1582	
1458507_at	1.26080	RIKEN cDNA 2810055G22 gene	2810055G22Rik	
1458508_at	2.75334	matrin 3	Matr3	
1458509_at	1.26042	Meiotic recombination 11 homolog A (S. cerevisiae)	Mre11a	
1458510_at	0.90517	expressed sequence AU019823	AU019823	
1458511_at	0.97957	RIKEN cDNA 2410025L10 gene	2410025L10Rik	
1458512_at	1.28423	"transducin-like enhancer of split 3, homolog of Drosophila E(spl)"	Tle3	
1458513_at	1.02949	mitogen activated protein kinase 10	Mapk10	
1458514_at	0.98998	expressed sequence AA536887	AA536887	
1458515_at	0.92968	zinc finger protein 128	Zfp128	
1458516_at	1.18033	NCK associated protein 1 like	Nckap11	
1458517_at	0.94003	"Aldo-keto reductase family 1, member E1"	Akr1e1	
1458518_at	0.94508	cytoplasmic polyadenylation element binding protein 2	Cpeb2	
1458519_at	0.98077	"0 day neonate head cDNA, RIKEN full-length enriched library, clone:4833418G05 product:unclassifiable, full insert sequence"	---	
1458520_at	1.04665	Homeodomain interacting protein kinase 2	Hipk2	
1458521_at	0.93170	Jumonji domain containing 1C	Jmjd1c	
1458522_at	0.98246	---	---	
1458523_at	1.04118	"zinc finger, matrin-like"	Zfml	
1458524_at	1.19476	fibronectin type III domain containing 3a	Fndc3a	
1458525_at	1.98164	Amyloid beta (A4) precursor protein	App	
1458526_at	1.42116	---	---	
1458527_at	0.92723	RIKEN cDNA 2610005L07 gene	2610005L07Rik	
1458528_at	0.99132	RIKEN cDNA C330002I19 gene	C330002I19Rik	
1458529_at	0.99684	hypothetical protein A930010C09	LOC215714	
1458530_at	1.06991	Zinc finger protein 532	Zfp532	
1458531_at	0.90991	"DNA segment, Chr 6, ERATO Doi 588, expressed"	D6Ertd588e	
1458532_at	1.05469	5-methyltetrahydrofolate-homocysteine methyltransferase	Mtr	
1458533_at	0.97299	RIKEN cDNA 9330169L03 gene	9330169L03Rik	
1458534_at	0.93269	"DNA segment, Chr 13, Brigham & Women's Genetics 1146 expressed"	D13Bwg1146e	
1458535_at	1.39309	"Adult male spinal cord cDNA, RIKEN full-length enriched library, clone: A330040D20 product:unclassifiable, full insert sequence"	---	
1458536_at	0.93321	Cyclin I	Ccni	
1458537_at	0.99282	---	---	
1458538_at	1.13656	"Sirtuin 1 ((silent mating type information regulation 2, homolog) 1 (S. cerevisiae)"	Sirt1	
1458539_at	0.92713	R3H domain 1 (binds single-stranded nucleic acids)	R3hdm1	
1458540_at	1.08098	Transcribed locus	---	

1458541_at	0.89894	dynactin 4	Dctn4	
1458542_at	1.22186	Musashi homolog 2 (Drosophila)	Msi2	
1458543_at	0.85323	---	---	
1458544_at	1.05281	cDNA sequence BC059841	BC059841	
1458545_at	1.11457	Transmembrane protein 57	Tmem57	
1458546_at	1.01154	"DNA segment, Chr 18, ERATO Doi 653, expressed"	D18Ertd653e	
1458547_at	0.89416	Transcribed locus	---	
1458548_at	1.02301	Transcribed locus	---	
1458549_at	1.06847	LOC436015	LOC436015	
1458550_at	1.50981	myosin ID	Myo1d	
1458551_at	1.05927	Myocyte enhancer factor 2C	Mef2c	
1458552_at	0.81505	RIKEN cDNA D430050E20 gene	D430050E20Rik	
1458553_at	0.94372	"plasminogen activator, tissue"	Plat	
1458554_at	1.04515	RIKEN cDNA 9330169B04 gene	9330169B04Rik	
1458555_at	1.02059	---	---	
1458556_at	1.04419	Trinucleotide repeat containing 6C	Tnrc6c	
1458557_at	1.02440	breast carcinoma amplified sequence 3	Bcas3	
1458558_at	0.98108	---	---	
1458559_at	0.92566	RIKEN cDNA A730028G07 gene	A730028G07Rik	
1458560_at	1.02105	"asp (abnormal spindle)-like, microcephaly associated (Drosophila)"	Aspm	
1458561_at	1.00468	neuralized-like homolog (Drosophila)	Neurl	
1458562_at	1.17108	---	---	
1458563_at	0.96849	zinc finger protein 668	Zfp668	
1458564_at	0.97483	---	---	
1458565_at	1.10754	---	---	
1458566_at	1.17394	G patch domain containing 2	Gpatc2	
1458567_at	1.11830	RIKEN cDNA D130017N08 gene	D130017N08Rik	
1458568_at	0.85132	Tumor suppressor candidate 3	Tusc3	
1458569_at	1.02620	elongation factor RNA polymerase II 2	Ell2	
1458570_at	0.93584	"Alveolar soft part sarcoma chromosome region, candidate 1 (human)"	Aspscr1	
1458571_at	0.91072	RIKEN cDNA D430047D06 gene	D430047D06Rik	
1458572_at	0.85064	chromatin modifying protein 5	Chmp5	
1458573_at	0.94761	Transcribed locus	---	
1458574_at	1.23257	E2F transcription factor 2	E2f2	
1458575_at	1.27937	SET binding protein 1	Setbp1	
1458576_at	1.26953	N-acetyltransferase 11	Nat11	
1458577_at	0.88305	---	---	
1458578_at	3.10036	"RNA binding motif, single stranded interacting protein 1"	Rbms1	
1458579_at	1.02610	---	---	
1458580_at	1.05468	Transcribed locus	---	
1458581_at	1.09612	hypothetical LOC553093	LOC553093	
1458582_at	1.17448	expressed sequence AI461788	AI461788	

1458583_at	1.08540	RIKEN cDNA 1700123O12 gene	1700123O12Rik
1458584_at	1.02247	RIKEN cDNA 4832406H04 gene	4832406H04Rik
1458585_at	0.69560	Transcribed locus	---
1458586_at	2.01328	---	---
1458587_at	0.96237	RIKEN cDNA 2310047D07 gene	2310047D07Rik
1458588_at	1.93066	Schwannomin interacting protein 1	Schip1
1458589_at	1.11649	---	---
1458590_at	1.13336	Integrator complex subunit 7	Ints7
1458591_at	0.92860	RAS and EF hand domain containing	Rasef
1458592_at	0.97853	Autism susceptibility candidate 2	Auts2
1458593_at	0.94969	Fibrillin 1	Fbn1
1458594_at	0.85731	SNF2 histone linker PHD RING helicase	Shprh
1458595_at	1.06298	PDZ domain containing 6	Pdzd6
1458596_at	0.85733	dynactin 4	Dctn4
1458597_at	1.10775	Transcribed locus	---
1458598_at	1.08738	paternally expressed 13	Peg13
1458599_at	1.01673	---	---
1458600_at	1.02566	Transcribed locus	---
1458601_at	1.23742	RIKEN cDNA 8030447M02 gene	8030447M02Rik
1458602_at	0.78825	bobby sox homolog (Drosophila)	Bbx
1458603_at	1.38103	---	---
1458604_at	1.57003	Rho/rac guanine nucleotide exchange factor (GEF) 18	Arhgef18
1458605_at	1.09434	ADP-ribosylation factor-like 15	Arl15
1458606_at	0.96330	---	---
1458607_at	0.97042	"Transforming growth factor, beta receptor II"	Tgfbr2
1458608_at	0.96174	Protein-tyrosine sulfotransferase 2	Tpst2
1458609_at	0.98071	"Adult male bone cDNA, RIKEN full-length enriched library, clone:9830162N06 product:unclassifiable, full insert sequence"	---
1458610_at	0.94181	---	---
1458611_at	1.01798	Transcribed locus	---
1458612_at	0.94619	RIKEN cDNA A930006A01 gene	A930006A01Rik
1458613_at	1.10313	arrestin domain containing 1	Arrdc1
1458614_at	0.90758	DCP1 decapping enzyme homolog b (S. cerevisiae)	Dcp1b
1458615_at	1.07780	DEP domain containing 5	Depdc5
1458616_at	1.25174	"UDP-Gal:betaGlcNAc beta 1,4- galactosyltransferase, polypeptide 1"	B4galt1
1458617_at	1.48970	protein kinase C binding protein 1	Prkcbp1
1458618_at	1.45981	Iron responsive element binding protein 2	Ireb2
1458619_at	0.88046	"Transcribed locus, moderately similar to XP_576460.1 PREDICTED: similar to hypothetical protein PB402898.00.0 [Rattus norvegicus]"	---
1458620_at	0.97358	similar to ubiquitin A-52 residue ribosomal protein fusion product 1	LOC332993
1458621_at	0.93271	WD repeat domain 78	Wdr78

1458622_at 0.99133 "neurotrophic tyrosine kinase, receptor, type 2" Ntrk2
1458623_at 1.06414 Transcribed locus ---
1458624_at 1.05037 similar to RNA binding motif protein 24 /// similar to RNA binding motif
protein 24 LOC380843 /// LOC666794
1458625_at 1.00471 "10 days neonate cortex cDNA, RIKEN full-length enriched library, clone:
A830014C22 product:unclassifiable, full insert sequence" ---
1458626_at 0.95482 "nitric oxide synthase 1, neuronal" Nos1
1458627_at 1.08202 "9 days embryo whole body cDNA, RIKEN full-length enriched library, clone:
D030053I05 product:unclassifiable, full insert sequence" ---
1458628_at 1.04812 LIM motif-containing protein kinase 2 Limk2
1458629_at 0.97254 --- ---
1458630_at 1.00787 Vav 3 oncogene Vav3
1458631_at 0.92612 RIKEN cDNA 4921533L14 gene 4921533L14Rik
1458632_at 1.00577 Dihydropyrimidinase-like 2 Dpysl2
1458633_at 1.06582 --- ---
1458634_at 0.96801 "CD47 antigen (Rh-related antigen, integrin-associated signal transducer)"
Cd47
1458635_at 0.88906 RIKEN cDNA 4832428D23 gene 4832428D23Rik
1458636_at 1.12013 RIKEN cDNA 2610206B13 gene 2610206B13Rik
1458637_x_at 1.23377 Ubiquitin protein ligase E3C Ube3c
1458638_at 0.90557 Metadherin Mtdh
1458639_at 1.14868 Formin binding protein 1 Fnbp1
1458640_at 0.95656 "6 days neonate skin cDNA, RIKEN full-length enriched library, clone:
A030007I19 product:hypothetical protein, full insert sequence" ---
1458641_at 1.07371 Braf transforming gene Braf
1458642_at 1.01254 killer cell lectin-like receptor family E member 1 Klre1
1458643_at 0.91690 deltex 1 homolog (Drosophila) Dtx1
1458644_at 0.88848 "expressed sequence AI661708 /// gene model 150, (NCBI)" AI661708 ///
Gm150
1458645_at 1.01457 Transcribed locus ---
1458646_at 1.14964 MMS19 (MET18 S. cerevisiae)-like Mms19l
1458647_at 1.01400 RIKEN cDNA 6430553K19 gene 6430553K19Rik
1458648_at 3.08241 expressed sequence AU042950 AU042950
1458649_at 0.96101 expressed sequence AI662245 AI662245
1458650_at 0.96817 --- ---
1458651_at 1.05929 Mitogen activated protein kinase kinase kinase 10 Map3k10
1458652_at 0.96550 "CD8 antigen, beta chain 1" Cd8b1
1458653_at 1.04428 Pannexin 1 Panx1
1458654_at 0.97723 Transcribed locus ---
1458655_at 1.35520 WW domain containing transcription regulator 1 Wwtr1
1458656_at 0.95345 expressed sequence AW050198 AW050198
1458657_at 1.08830 Anthrax toxin receptor 1 Antxr1
1458658_at 0.92083 --- ---

1458659_at 1.06727 placenta specific 9 Plac9
1458660_at 0.89604 expressed sequence AI414330 AI414330
1458661_at 1.63594 Mortality factor 4 like 1 Morf411
1458662_at 0.77229 dishevelled associated activator of morphogenesis 1 Daam1
1458663_at 1.99685 Like-glycosyltransferase Large
1458664_at 1.19251 Transcribed locus ---
1458665_at 1.15369 Chloride intracellular channel 1 Clic1
1458666_at 0.99120 "DNA segment, Chr 2, ERATO Doi 391, expressed" D2Ertd391e
1458667_at 0.82453 RIKEN cDNA 4930519N13 gene 4930519N13Rik
1458668_at 0.99408 Tumor protein D52 Tpd52
1458669_at 0.97119 Transcribed locus ---
1458670_at 0.84048 "Adenosine deaminase, RNA-specific, B2" Adarb2
1458671_at 0.94799 hypothetical protein LOC669460 LOC669460
1458672_at 0.97024 T-complex-associated testis expressed 2 Tcte2
1458673_at 0.91123 RIKEN cDNA 1700093J21 gene 1700093J21Rik
1458674_at 1.01794 Protein kinase N1 Pkn1
1458675_at 0.85198 --- ---
1458676_at 0.99834 natural killer tumor recognition sequence Nktr
1458677_at 0.87157 ectonucleoside triphosphate diphosphohydrolase 5 Entpd5
1458678_at 0.97253 "NADH dehydrogenase (ubiquinone) 1, alpha/beta subcomplex, 1" Ndufab1
1458679_a_at 1.04322 TatD DNase domain containing 1 Tatdn1
1458680_at 1.02853 Hephaestin-like 1 Heph11
1458681_at 0.96852 Transmembrane 9 superfamily member 3 Tm9sf3
1458682_at 0.97440 RIKEN cDNA D130059P03 gene D130059P03Rik
1458683_at 0.92617 signal-regulatory protein beta 1 /// similar to SIRP beta 1 isoform 1 Sirpb1 ///
LOC673172
1458684_at 1.18841 "synovial sarcoma translocation, Chromosome 18" Ss18
1458685_at 0.91316 GTPase activating RANGAP domain-like 1 Garn11
1458686_at 1.05864 Gametogenetin binding protein 2 Ggnbp2
1458687_at 0.88926 hypothetical protein LOC668172 /// hypothetical protein LOC672498 ///
hypothetical protein LOC677134 LOC668172 /// LOC672498 /// LOC677134
1458688_at 1.17326 Rho GTPase activating protein 28 Arhgap28
1458689_at 1.15283 Autophagy-related 10 (yeast) Atg10
1458690_at 1.64455 membrane-associated ring finger (C3HC4) 7 7-Mar
1458691_at 1.01430 Synaptopodin 2-like Synpo2l
1458692_at 1.09302 Translocase of outer mitochondrial membrane 20 homolog (yeast) Tomm20
1458693_at 1.04054 "Sodium channel, voltage-gated, type VIII, alpha" Scn8a
1458694_at 1.00485 expressed sequence AI789011 AI789011
1458695_at 0.94147 Zinc finger and BTB domain containing 10 Zbtb10
1458696_at 0.87831 ARP1 actin-related protein 1 homolog A (yeast) Actr1a
1458697_at 0.97687 Transcribed locus ---
1458698_at 0.96799 CDNA sequence AB041803 AB041803
1458699_at 1.19345 CCR4 carbon catabolite repression 4-like (*S. cerevisiae*) Ccrn4l

1458700_at	1.69493	"Leucine rich repeat containing 8 family, member C"	Lrrc8c
1458701_at	1.04491	preimplantation protein 4	Prei4
1458702_at	1.07929	RIKEN cDNA 2610316D01 gene	2610316D01Rik
1458703_at	2.15432	---	---
1458704_at	0.96854	"Transcribed locus, moderately similar to NP_663461.2 membrane-associated ring finger (C3HC4) 2 [Mus musculus]"	---
1458705_at	1.03325	RIKEN cDNA 1700031C06 gene	1700031C06Rik
1458706_at	1.16719	RIKEN cDNA 2610035D17 gene	2610035D17Rik
1458707_at	0.99044	RIKEN cDNA E430016F16 gene	E430016F16Rik
1458708_at	1.75059	Bromodomain containing 4	Brd4
1458709_a_at	1.06218	ataxin 7-like 1	Atxn7l1
1458710_at	1.82056	Polymerase (RNA) II (DNA directed) polypeptide A	Polr2a
1458711_at	1.00578	---	---
1458712_at	0.99892	---	---
1458713_at	0.87773	---	---
1458714_at	0.94093	RIKEN cDNA 1700093C20 gene	1700093C20Rik
1458715_at	1.13058	Caspase 6	Casp6
1458716_at	1.00027	dual specificity phosphatase 27 (putative)	Dusp27
1458717_at	1.02621	RIKEN cDNA C030010L15 gene	C030010L15Rik
1458718_at	0.97284	"PREDICTED: Mus musculus hypothetical LOC243025 (LOC243025), mRNA"	---
1458719_at	1.04165	Transcribed locus	---
1458720_at	0.95920	"DNA segment, Chr 9, ERATO Doi 356, expressed"	D9Ertd356e
1458721_at	0.91954	"Protocadherin gamma subfamily C, 3"	Pcdhgc3
1458722_at	1.07283	RIKEN cDNA 1200015F23 gene	1200015F23Rik
1458723_at	1.06315	"Similar to Poly(A) binding protein, cytoplasmic 4, isoform 1"	LOC544782
1458724_at	1.33331	RIKEN cDNA E230008O15 gene	E230008O15Rik
1458725_at	1.01362	---	---
1458726_at	1.09165	expressed sequence C87033	C87033
1458727_at	1.03131	"Protein tyrosine phosphatase-like (proline instead of catalytic arginine), member b"	Ptplb
1458728_at	0.95339	---	---
1458729_at	1.10725	FK506 binding protein 4	Fkbp4
1458730_at	1.13205	BTB and CNC homology 2	Bach2
1458731_at	1.03680	Transcribed locus	---
1458732_at	1.05826	---	---
1458733_at	0.86823	G protein-coupled receptor 149	Gpr149
1458734_at	0.86714	expressed sequence C78128	C78128
1458735_at	1.00490	Transcribed locus	---
1458736_at	0.96614	---	---
1458737_at	0.94542	expressed sequence C77097	C77097
1458738_at	0.93678	"Guanine nucleotide binding protein, alpha 14"	Gna14
1458739_at	1.08425	---	---

1458740_at	0.87063 --- ---
1458741_at	0.95924 Phosphoglycerate dehydrogenase like 1 Phgdh1
1458742_at	0.89976 latrophilin 3 Lphn3
1458743_at	1.07079 --- ---
1458744_at	0.90584 "TAF1 RNA polymerase II, TATA box binding protein (TBP)-associated factor" Taf1
1458745_at	0.87690 "DNA segment, Chr 18, ERATO Doi 201, expressed" D18Ertd201e
1458746_at	0.88075 RIKEN cDNA A430107O13 gene A430107O13Rik
1458747_at	0.92882 F-box protein 45 Fbxo45
1458748_at	1.03505 RNA binding motif protein 27 Rbm27
1458749_at	0.96741 Single-stranded DNA binding protein 3 Ssbp3
1458750_at	0.99095 "15 days embryo head cDNA, RIKEN full-length enriched library, clone: D930028N23 product:unclassifiable, full insert sequence" ---
1458751_at	0.87664 --- ---
1458752_at	1.06732 "PREDICTED: Mus musculus similar to otogelin (LOC627748), mRNA" ---
1458753_at	0.94669 "gene model 941, (NCBI)" Gm941
1458754_at	0.95138 --- ---
1458755_at	1.09731 Transcribed locus ---
1458756_at	1.00734 "Glutamate receptor, ionotropic, delta 2" Grid2
1458757_at	0.94518 Polypyrimidine tract binding protein 2 Ptbp2
1458758_at	1.15016 Vav2 oncogene Vav2
1458759_at	0.95881 Microtubule-associated protein 4 Mtap4
1458760_at	1.01702 --- ---
1458761_at	0.83727 U box domain containing 5 Ubox5
1458762_at	0.83048 expressed sequence AU015230 AU015230
1458763_at	0.96206 expressed sequence C78024 C78024
1458764_at	1.19801 --- ---
1458765_at	1.13516 RNA binding motif protein 10 Rbm10
1458766_at	0.96270 P21 (CDKN1A)-activated kinase 7 Pak7
1458767_at	0.90525 "Glucan (1,4-alpha-), branching enzyme 1" Gbe1
1458768_at	1.58241 Erythrocyte protein band 4.1-like 5 Epb4.115
1458769_at	0.95477 "DNA segment, Chr 7, ERATO Doi 558, expressed" D7Ertd558e
1458770_at	1.14076 "DNA segment, Chr 11, ERATO Doi 4, expressed" D11Ertd4e
1458771_at	0.91385 --- ---
1458772_at	1.01851 Latrophilin 3 Lphn3
1458773_at	0.93313 "UDP-Gal:betaGlcNAc beta 1,4- galactosyltransferase, polypeptide 1" B4galt1
1458774_at	0.94046 "DNA segment, Chr 10, ERATO Doi 584, expressed" D10Ertd584e
1458775_at	1.09458 RIKEN cDNA 1700041E20 gene 1700041E20Rik
1458776_at	1.09124 --- ---
1458777_at	1.07269 "solute carrier family 18 (vesicular monoamine), member 1" Slc18a1
1458778_at	1.14424 WD repeats and SOF domain containing 1 Wdsof1
1458779_at	0.95684 RIKEN cDNA 8030445P17 gene 8030445P17Rik

1458780_at	0.99695	"DNA segment, Chr 8, ERATO Doi 620, expressed"	D8Ertd620e
1458781_at	1.07157	"Potassium channel, subfamily K, member 13"	Kcnk13
1458782_at	0.99980	Transcribed locus	---
1458783_at	0.89689	---	---
1458784_at	0.95011	expressed sequence	C78148 C78148
1458785_at	0.99324	Synovial sarcoma translocation gene on chromosome 18-like 1	Ss1811
1458786_at	0.89904	expressed sequence	C79250 C79250
1458787_at	1.00361	Mitogen activated protein kinase kinase kinase 7	Map3k7
1458788_at	2.50576	Ecotropic viral integration site 2a	Evi2a
1458789_at	1.12654	---	---
1458790_at	1.04307	---	---
1458791_at	0.88656	expressed sequence	AU015849 AU015849
1458792_at	1.11238	"DNA segment, Chr 12, ERATO Doi 208, expressed"	D12Ertd208e
1458793_at	1.23858	Tensin 3	Tns3
1458794_at	0.97800	---	---
1458795_at	0.89680	Transcribed locus	---
1458796_at	0.99192	---	---
1458797_at	1.05532	---	---
1458798_at	0.96550	Transcribed locus	---
1458799_at	0.93954	RIKEN cDNA A330033J07 gene	A330033J07Rik
1458800_at	0.84613	Trinucleotide repeat containing 6C	Tnrc6c
1458801_at	1.05055	"fusion, derived from t(12;16) malignant liposarcoma (human)"	Fus
1458802_at	0.94931	human immunodeficiency virus type I enhancer binding protein 3	Hivep3
1458803_at	0.88684	schlafen 9 /// similar to schlafen 9	Slfn9 /// LOC677619
1458804_at	0.86319	"Solute carrier organic anion transporter family, member 4C1"	Slco4c1
1458805_at	0.88682	---	---
1458806_at	0.91666	RIKEN cDNA D230005D02 gene	D230005D02Rik
1458807_at	1.26631	erythrocyte protein band 4.1	Epb4.1
1458808_at	1.03934	"Sushi, nidogen and EGF-like domains 1"	Sned1
1458809_at	1.35752	---	---
1458810_at	0.99949	---	---
1458811_at	0.95479	RIKEN cDNA 9430047L24 gene	9430047L24Rik
1458812_at	1.02731	Adenylate kinase 3	Ak3
1458813_at	1.01275	"sodium channel, voltage-gated, type V, alpha"	Scn5a
1458814_at	1.07409	expressed sequence	AU022855 AU022855
1458815_at	0.94394	expressed sequence	AU021884 AU021884
1458816_at	1.03378	---	---
1458817_at	0.93978	---	---
1458818_at	0.91365	"DNA segment, Chr 3, ERATO Doi 162, expressed"	D3Ertd162e
1458819_at	0.96958	GTPase activating RANGAP domain-like 3	Garnl3
1458820_at	0.96607	zinc finger protein 560	Zfp560
1458821_at	0.90052	"SEC24 related gene family, member D (S. cerevisiae)"	Sec24d
1458822_at	1.00731	---	---

1458823_at	1.36858	Leucine-rich repeats and immunoglobulin-like domains 1	Lrig1
1458824_at	1.04724	unc-45 homolog A (C. elegans)	Unc45a
1458825_at	1.10664	"Mesoderm induction early response 1, family member 3"	Mier3
1458826_at	1.04787	expressed sequence C78809	C78809
1458827_at	0.97219	RAB11 family interacting protein 4 (class II)	Rab11fip4
1458828_at	1.15906	Expressed sequence AW146242	AW146242
1458829_at	1.03887	---	---
1458830_at	0.96657	Fibroblast growth factor 14	Fgf14
1458831_at	0.98168	NAD(P) dependent steroid dehydrogenase-like	Nsdhl
1458832_at	0.99243	Growth hormone receptor	Ghr
1458833_at	0.87514	RIKEN cDNA C130076O07 gene	C130076O07Rik
1458834_at	0.99093	Mitochondrial tumor suppressor 1	Mtus1
1458835_at	0.95720	---	---
1458836_at	1.02899	calbindin-28K	Calb1
1458837_at	0.82525	RIKEN cDNA 2810006K23 gene	2810006K23Rik
1458838_at	1.01594	RIKEN cDNA 4732468M13 gene	4732468M13Rik
1458839_at	1.04635	exocyst complex component 8	Exoc8
1458840_at	0.95545	Transcribed locus	---
1458841_at	0.87875	NudC domain containing 1	Nudcd1
1458842_at	1.07311	---	---
1458843_at	0.95228	DEP domain containing 5	Depdc5
1458844_at	0.97895	RIKEN cDNA 4933415L06 gene	4933415L06Rik
1458845_at	0.98242	"DNA segment, Chr 4, ERATO Doi 264, expressed"	D4Ertd264e
1458846_at	1.09792	---	---
1458847_at	1.01342	---	---
1458848_at	0.94287	Hypothetical protein LOC624549	LOC624549
1458849_at	1.03609	N-acylsphingosine amidohydrolase 2	Asah2
1458850_at	0.95681	"Myosin, heavy polypeptide 6, cardiac muscle, alpha"	Myh6
1458851_at	0.96991	Transcribed locus	---
1458852_at	0.98867	---	---
1458853_at	0.89436	Pre B-cell leukemia transcription factor 1	Pbx1
1458854_at	1.12853	expressed sequence C78891	C78891
1458855_at	0.92914	"DNA segment, Chr 11, ERATO Doi 49, expressed"	D11Ertd49e
1458856_at	1.01393	---	---
1458857_at	1.01391	Transcribed locus	---
1458858_at	1.01045	---	---
1458859_at	0.93774	---	---
1458860_at	0.81034	---	---
1458861_at	0.95961	---	---
1458862_at	0.96856	expressed sequence C78704	C78704
1458863_at	0.96613	RIKEN cDNA 6330415G19 gene	6330415G19Rik
1458864_at	1.10328	---	---
1458865_at	1.08251	---	---

1458866_at	0.90673	RIKEN cDNA A930001M12 gene	A930001M12Rik
1458867_at	0.99444	EGF-like repeats and discoidin I-like domains 3	Edil3
1458868_at	1.08957	"ATPase, Na ⁺ /K ⁺ transporting, alpha 1 polypeptide"	Atp1a1
1458869_at	0.90789	RIKEN cDNA 2900076A13 gene	2900076A13Rik
1458870_x_at	0.98514	"pam, highwire, rpm 1"	Phr1
1458871_at	1.01455	"transmembrane protease, serine 11e"	Tmprss11e
1458872_at	0.95740	---	---
1458873_at	0.94728	"asp (abnormal spindle)-like, microcephaly associated (Drosophila)"	Aspm
1458874_at	0.96053	FAT tumor suppressor homolog 3 (Drosophila)	Fat3
1458875_at	0.95053	"SEC24 related gene family, member B (S. cerevisiae)"	Sec24b
1458876_at	0.99319	Enabled homolog (Drosophila)	Enah
1458877_at	1.02524	Transcribed locus	---
1458878_at	1.04631	Yamaguchi sarcoma viral (v-yes) oncogene homolog 1	Yes1
1458879_at	1.07548	expressed sequence C76798	C76798
1458880_at	0.98518	---	---
1458881_at	1.05372	RIKEN cDNA 9530001P21 gene	9530001P21Rik
1458882_at	1.06159	"serine (or cysteine) peptidase inhibitor, clade B, member 8"	Serpinb8
1458883_at	1.11160	---	---
1458884_at	1.38309	---	---
1458885_at	0.86809	---	---
1458886_at	1.00184	Speckle-type POZ protein	Spop
1458887_at	0.92836	transcription factor 21	Tcf21
1458888_at	1.02839	CDNA clone IMAGE:5250818	---
1458889_at	1.10038	Ena-vasodilator stimulated phosphoprotein	Evl
1458890_at	0.96871	RIKEN cDNA D030055H07 gene	D030055H07Rik
1458891_at	1.09839	GTPase activating RANGAP domain-like 3	Garnl3
1458892_at	0.85546	RIKEN cDNA 9430047G12 gene	9430047G12Rik
1458893_at	0.93119	Transcribed locus	---
1458894_at	1.05043	---	---
1458895_at	1.09902	expressed sequence C80008	C80008
1458896_at	1.01636	glucocorticoid modulatory element binding protein 1	Gmeb1
1458897_at	1.01051	uronyl-2-sulfotransferase	Ust
1458898_at	0.99751	---	---
1458899_at	1.29222	ubiquitin specific peptidase 53	Usp53
1458900_at	0.92780	"Solute carrier family 24 (sodium/potassium/calcium exchanger), member 3"	Slc24a3
1458901_at	0.98453	Myocyte enhancer factor 2D	Mef2d
1458902_at	0.99616	---	---
1458903_at	1.00836	RIKEN cDNA A230057G18 gene	A230057G18Rik
1458904_at	0.85899	Frizzled homolog 6 (Drosophila)	Fzd6
1458905_x_at	0.88029	RIKEN cDNA 2610002D18 gene	2610002D18Rik
1458906_at	1.11279	---	---
1458907_at	0.85658	"Solute carrier organic anion transporter family, member 6b1"	Slco6b1

1458908_at	0.99913	Dystrobrevin alpha	Dtna
1458909_at	1.04167	---	---
1458910_at	1.02217	---	---
1458911_at	0.87285	RIKEN cDNA C230086A09 gene	C230086A09Rik
1458912_at	1.18906	RNA binding motif protein 6	Rbm6
1458913_at	0.97366	RIKEN cDNA 4831440E17 gene	4831440E17Rik
1458914_at	1.09082	---	---
1458915_at	1.04202	expressed sequence C77949	C77949
1458916_at	1.33830	"Solute carrier family 12, member 6"	Slc12a6
1458917_at	1.17224	DEP domain containing 5 Depdc5	
1458918_at	0.88991	"solute carrier family 12 (potassium/chloride transporters), member 8"	Slc12a8
1458919_at	1.36315	"muskelin 1, intracellular mediator containing kelch motifs"	Mkln1
1458920_at	1.16710	Syntaxin 6	Stx6
1458921_at	1.36173	hypothetical gene supported by AK043371	LOC434348
1458922_at	0.92291	mitogen-activated protein kinase kinase kinase 9	Map3k9
1458923_at	0.93538	Methionine sulfoxide reductase A	Msra
1458924_at	1.38895	RIKEN cDNA D430013B06 gene	D430013B06Rik
1458925_at	1.05063	zinc finger protein 82	Zfp82
1458926_at	1.26766	RIKEN cDNA 5830415L20 gene	5830415L20Rik
1458927_at	0.95447	"ATPase, Ca ⁺⁺ transporting, cardiac muscle, slow twitch 2"	Atp2a2
1458928_at	0.96791	---	---
1458929_at	0.92821	ATP-binding cassette transporter sub-family A member 9	Abca9
1458930_at	0.89545	"alpha-1,4-N-acetylglucosaminyltransferase"	A4gnt
1458931_at	1.13635	RIKEN cDNA 6030427F01 gene	6030427F01Rik
1458932_at	1.05699	peroxin 2	Pex2
1458933_at	0.92910	RIKEN cDNA A530052I06 gene	A530052I06Rik
1458934_at	1.08743	"DNA segment, Chr 5, ERATO Doi 505 , expressed"	D5ErtD505e
1458935_at	0.99909	expressed sequence C79744	C79744
1458936_at	1.05180	---	---
1458937_at	0.97026	5'-nucleotidase domain containing 1	Nt5dc1
1458938_at	0.97339	---	---
1458939_at	0.99812	Synaptojanin 2 binding protein	Synj2bp
1458940_at	0.96114	RIKEN cDNA 5730507A11 gene	5730507A11Rik
1458941_at	1.00024	RIKEN cDNA D130016B08 gene	D130016B08Rik
1458942_at	1.03426	RIKEN cDNA C230037E05 gene	C230037E05Rik
1458943_at	0.99319	---	---
1458944_at	1.06060	"Adult male testis cDNA, RIKEN full-length enriched library, clone:4930542D16 product:unclassifiable, full insert sequence"	---
1458945_at	0.94353	expressed sequence AU015148	AU015148
1458946_at	1.02177	---	---
1458947_at	1.80266	RIKEN cDNA 2010111I01 gene	2010111I01Rik
1458948_at	0.99333	RIKEN cDNA D630013N20 gene	D630013N20Rik
1458949_at	1.08827	RIKEN cDNA 5330434G04 gene	5330434G04Rik

1458950_at	0.87212	Transcribed locus	---	
1458951_at	0.96271	vaccinia related kinase 1	Vrk1	
1458952_at	1.00115	WD repeat and FYVE domain containing 3	Wdfy3	
1458953_at	0.98841	"DNA segment, Chr 5, ERATO Doi 215, expressed"	D5Ertd215e	
1458954_at	0.90827	---	---	
1458955_at	1.05842	Eukaryotic translation elongation factor 1 gamma	Eef1g	
1458956_at	0.96676	RIKEN cDNA A330068P14 gene	A330068P14Rik	
1458957_at	0.93977	"Similar to pyridoxal (pyridoxine, vitamin B6) kinase"	LOC623548	
1458958_at	0.86241	neighbor of Punc E11	Nope	
1458959_at	1.00212	"AF4/FMR2 family, member 4"	Aff4	
1458960_at	1.21776	Receptor-like tyrosine kinase	Ryk	
1458961_at	1.04350	eyes absent 3 homolog (Drosophila)	Eya3	
1458962_at	1.08079	transmembrane emp24 protein transport domain containing 5	Tmed5	
1458963_at	1.10099	RIKEN cDNA 1810054D07 gene	1810054D07Rik	
1458964_at	1.08842	Interleukin 1 receptor accessory protein-like 1	Il1rapl1	
1458965_at	0.96546	---	---	
1458966_at	0.87954	expressed sequence C80278	C80278	
1458967_at	0.97344	Transcribed locus	---	
1458968_at	0.98693	expressed sequence C80283	C80283	
1458969_at	1.17448	expressed sequence AU019559	AU019559	
1458970_at	0.96418	RAD50 homolog (<i>S. cerevisiae</i>)	Rad50	
1458971_at	0.87377	---	---	
1458972_at	1.28960	RIKEN cDNA 9330112F22 gene	9330112F22Rik	
1458973_at	1.02984	"PREDICTED: Mus musculus similar to RAB5A, member RAS oncogene family (LOC629071), mRNA"	---	
1458974_at	0.93485	Thymocyte selection-associated HMG box gene	Tox	
1458975_at	0.93428	Transcribed locus	---	
1458976_at	1.02598	RIKEN cDNA C230035I16 gene	C230035I16Rik	
1458977_at	0.98766	hypothetical protein A530021J07	A530021J07	
1458978_at	0.93718	---	---	
1458979_at	0.97173	---	---	
1458980_at	2.36464	---	---	
1458981_at	1.28453	Transcribed locus	---	
1458982_at	1.02628	expressed sequence AU014965	AU014965	
1458983_at	1.14123	expressed sequence AU015559	AU015559	
1458984_at	0.92097	RIKEN cDNA E130308A19 gene	E130308A19Rik	
1458985_at	1.30805	furry homolog (Drosophila)	Fry	
1458986_at	1.00353	RIKEN cDNA 4930534H18 gene	4930534H18Rik	
1458987_at	0.82554	Cyclin-dependent kinase 7 (homolog of Xenopus MO15 cdk-activating kinase)	Cdk7	
1458988_at	1.08344	---	---	
1458989_at	1.05793	"LIM-domain containing, protein kinase"	Limk1	
1458990_at	0.88752	SET domain containing 5	Setd5	

1458991_at 1.17468 "Solute carrier family 37 (glycerol-3-phosphate transporter), member 1" Slc37a1
 1458992_at 0.99184 SUMO/sentrin specific peptidase 5 Senp5
 1458993_at 0.94919 expressed sequence AU022882 AU022882
 1458994_at 1.36527 "Casein kinase 1, gamma 3" Csnk1g3
 1458995_at 0.89725 Expressed sequence AA408296 AA408296
 1458996_at 1.14318 integrin alpha 5 (fibronectin receptor alpha) Itga5
 1458997_at 0.99532 RIKEN cDNA A230106M20 gene A230106M20Rik
 1458998_at 1.02210 oligonucleotide/oligosaccharide-binding fold containing 1 Obfc1
 1458999_at 1.17838 Mitochondrial ribosomal protein S10 Mrps10
 1459000_at 1.11216 --- ---
 1459001_at 1.13856 Vacuolar protein sorting 33A (yeast) Vps33a
 1459002_at 1.01035 ribosomal protein L21 /// similar to 60S ribosomal protein L21 /// similar to
 ribosomal protein L21 /// similar to ribosomal protein L21 /// similar to ribosomal protein L21 /// similar
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 hypothetical protein LOC665245 /// similar to ribosomal protein L21 /// similar to ribosomal protein
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 ribosomal protein L21 /// hypothetical protein LOC671626 /// similar to 60S ribosomal protein L21 ///
 similar to ribosomal protein L21 /// hypothetical protein LOC673532 /// similar to 60S ribosomal protein
 L21 /// similar to ribosomal protein L21 /// similar to 60S ribosomal protein L21 /// similar to ribosomal
 protein L21 /// similar to 60S ribosomal protein L21 /// hypothetical protein LOC675371 /// similar to
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 protein L21 /// similar to 60S ribosomal protein L21 /// similar to ribosomal protein L21 Rpl21 ///
 LOC268676 /// LOC382843 /// LOC432665 /// LOC432725 /// LOC432801 /// LOC433125 ///
 LOC433168 /// LOC433319 /// LOC433387 /// LOC433630 /// LOC434067 /// LOC434253 ///

LOC545172 /// LOC545226 /// LOC545423 /// LOC545572 /// LOC545877 /// LOC620520 ///
 LOC620771 /// LOC625038 /// LOC627939 /// LOC628449 /// LOC628822 /// LOC631359 ///
 LOC632448 /// LOC633404 /// LOC634339 /// LOC634904 /// LOC635258 /// LOC636187 ///
 LOC636855 /// LOC637748 /// LOC637883 /// LOC637900 /// LOC637907 /// LOC637993 ///
 LOC638534 /// LOC639424 /// LOC639991 /// LOC640096 /// LOC665245 /// LOC665251 ///
 LOC665821 /// LOC666340 /// LOC666977 /// LOC667610 /// LOC668319 /// LOC668785 ///
 LOC668919 /// LOC669703 /// LOC669967 /// LOC670607 /// LOC670636 /// LOC670852 ///
 LOC670966 /// LOC671028 /// LOC671397 /// LOC671626 /// LOC672341 /// LOC672798 ///
 LOC673532 /// LOC674403 /// LOC674827 /// LOC675035 /// LOC675278 /// LOC675313 ///
 LOC675371 /// LOC675601 /// LOC675818 /// LOC675935 /// LOC676253 /// LOC676590 ///
 LOC676825 /// LOC676831 /// LOC676903 /// LOC677032 /// LOC677133 /// LOC677463 ///
 LOC677495 /// LOC677530 /// LOC677659

1459003_at 1.01405 four and a half LIM domains 1 Fhl1
 1459004_at 0.93974 --- ---
 1459005_at 2.02258 RIKEN cDNA D030051N19 gene D030051N19Rik
 1459006_a_at 0.91483 Hypothetical gene supported by AK083927 LOC432938
 1459007_at 0.99545 Transcribed locus ---
 1459008_at 2.20071 --- ---
 1459009_at 1.36637 Utrophin Utrn
 1459010_at 1.08818 RIKEN cDNA 5430437P03 gene 5430437P03Rik
 1459011_at 1.19987 Ubiquitin carboxyl-terminal esterase L5 Uchl5
 1459012_at 1.01830 RIKEN cDNA A730041O05 gene A730041O05Rik
 1459013_at 0.92729 --- ---
 1459014_at 1.02267 RIKEN cDNA 2900019G14 gene 2900019G14Rik
 1459015_at 1.02662 "Potassium voltage-gated channel, subfamily H (eag-related), member 8" Kcnh8
 1459016_at 1.05105 "Insulin-like growth factor 2, binding protein 3" Igf2bp3
 1459017_at 1.03210 --- ---
 1459018_at 0.91206 "DNA segment, Chr 10, ERATO Doi 447, expressed" D10Ertd447e
 1459019_at 1.07573 shadow of prion protein Sprn
 1459020_at 0.94080 adhesion molecule with Ig like domain 1 Amigo1
 1459021_at 0.98617 Transcribed locus ---
 1459022_at 1.11735 --- ---
 1459023_at 0.91802 --- ---
 1459024_at 0.90227 Transcribed locus ---
 1459025_at 1.02204 RIKEN cDNA 1700001J04 gene 1700001J04Rik
 1459026_at 1.44282 SNW domain containing 1 Snw1
 1459027_at 0.85976 Transcribed locus ---
 1459028_at 1.15565 "UTP20, small subunit (SSU) processome component, homolog (yeast)"
 Utp20
 1459029_at 0.91496 P21 (CDKN1A)-activated kinase 1 Pak1
 1459030_at 1.00610 --- ---
 1459031_at 1.03916 "tubulin tyrosine ligase-like family, member 4" Ttl14
 1459032_at 1.09515 Transcribed locus ---

1459033_at	0.99735	Plexin A2	Plxna2	
1459034_at	0.69583	WD repeat domain 19	Wdr19	
1459035_at	0.98698	---	---	
1459036_at	0.94188	RIKEN cDNA 4930402E16 gene	4930402E16Rik	
1459037_at	0.90028	"gene model 1614, (NCBI)"	Gm1614	
1459038_at	0.96026	Transcribed locus	---	
1459039_at	1.05433	HEAT repeat containing 1	Heatr1	
1459040_at	0.88244	High mobility group nucleosomal binding domain 3	Hmgn3	
1459041_at	1.06269	Sorting nexin 12	Snx12	
1459042_at	0.98579	regulating synaptic membrane exocytosis 3	Rims3	
1459043_at	0.97153	"Farnesyltransferase, CAAX box, beta"	Fntb	
1459044_at	1.00499	expressed sequence AU015558	AU015558	
1459045_at	1.00605	golgi phosphoprotein 4	Golph4	
1459046_at	1.08761	RIKEN cDNA 4930473A06 gene	4930473A06Rik	
1459047_x_at	1.01003	RIKEN cDNA 5430405G24 gene	5430405G24Rik	
1459048_s_at	0.95377	zinc finger protein 142	Zfp142	
1459049_at	1.01890	RAS protein-specific guanine nucleotide-releasing factor 1	Rasgrf1	
1459050_at	0.97669	"gene model 996, (NCBI)"	Gm996	
1459051_at	0.95186	RIKEN cDNA 6530418L21 gene	6530418L21Rik	
1459052_at	0.95712	"Hypoxia inducible factor 3, alpha subunit"	Hif3a	
1459053_at	0.97750	"centaurin, gamma 2"	Centg2	
1459054_at	1.00244	cDNA sequence BC035954	BC035954	
1459055_at	1.17707	forkhead box J3	Foxj3	
1459056_at	0.95399	RAS protein-specific guanine nucleotide-releasing factor 1	Rasgrf1	
1459057_at	0.92261	Tudor domain containing 3	Tdrd3	
1459058_at	0.92577	sialic acid acetyltransferase	Siae	
1459059_at	1.00901	RIKEN cDNA 2010308F09 gene	2010308F09Rik	
1459060_at	1.03461	Shc SH2-domain binding protein 1	Shcbp1	
1459061_at	0.93255	Transferrin receptor 2	Trfr2	
1459062_x_at	1.10495	Expressed sequence AU041133	AU041133	
1459063_at	0.93242	Transcribed locus	---	
1459064_at	0.96701	"Discs, large homolog 2 (Drosophila)"	Dlgh2	
1459065_at	1.15496	"Kalirin, RhoGEF kinase"	Kalrn	
1459066_at	0.98395	zinc finger protein 612	Zfp612	
1459067_at	0.91726	RIKEN cDNA 1190002N15 gene	1190002N15Rik	
1459068_at	1.37904	Transmembrane protein 109	Tmem109	
1459069_at	1.04871	"Protein phosphatase 1D magnesium-dependent, delta isoform"	Ppm1d	
1459070_at	0.88103	Transcribed locus	---	
1459071_at	0.87587	Transcribed locus	---	
1459072_at	1.03929	Transcribed locus	---	
1459073_x_at	1.02460	Fibroblast growth factor 14	Fgf14	
1459074_at	1.02873	---	---	
1459075_at	0.92197	expressed sequence AU022479	AU022479	

1459076_at 0.95491 CDNA sequence BC052040 BC052040
1459077_at 0.90919 "Cytochrome P450, family 2, subfamily d, polypeptide 26" Cyp2d26
1459078_at 1.26389 RNA binding protein gene with multiple splicing Rbpms
1459079_at 1.03008 --- ---
1459080_at 0.88478 --- ---
1459081_at 0.94061 RIKEN cDNA 1700007B14 gene 1700007B14Rik
1459082_at 1.05618 Transcribed locus ---
1459083_at 0.97514 RIKEN cDNA A330084C13 gene A330084C13Rik
1459084_at 0.94552 "PREDICTED: Mus musculus similar to cyclin-dependent kinase 10 isoform 2
(LOC624138), mRNA" ---
1459085_at 1.00792 "DNA segment, Chr 1, ERATO Doi 259, expressed" D1Ert259e
1459086_at 0.97023 RIKEN cDNA A430107P09 gene A430107P09Rik
1459087_at 0.89497 --- ---
1459088_at 0.95416 expressed sequence C79557 C79557
1459089_at 0.92945 hypothetical LOC545637 /// hypothetical LOC623272 /// hypothetical
LOC623281 /// hypothetical protein LOC636750 /// hypothetical protein LOC667780 LOC545637 ///
LOC623272 /// LOC623281 /// LOC636750 /// LOC667780
1459090_at 1.11781 UBX domain containing 4 Ubx4
1459091_at 1.12922 --- ---
1459092_at 1.01685 "solute carrier organic anion transporter family, member 6c1" Slco6c1
1459093_at 0.96916 Fraser syndrome 1 homolog (human) Fras1
1459094_at 0.97534 Synaptosomal-associated protein 91 Snap91
1459095_at 0.97854 --- ---
1459096_at 0.98933 Activating transcription factor 7 interacting protein Atf7ip
1459097_at 0.96641 Interleukin 17 receptor D Il17rd
1459098_at 1.61196 Dystonin Dst
1459099_at 0.91794 transmembrane channel-like gene family 7 Tmc7
1459100_at 0.97395 expressed sequence AU015253 AU015253
1459101_at 0.96810 expressed sequence C78760 C78760
1459102_at 0.90197 Myosin ID Myo1d
1459103_at 0.97913 Histone cell cycle regulation defective homolog A (S. cerevisiae) Hira
1459104_at 1.13816 budding uninhibited by benzimidazoles 3 homolog (S. cerevisiae) Bub3
1459105_at 0.96131 --- ---
1459106_at 1.00523 "PREDICTED: Mus musculus hypothetical protein LOC622817 (LOC622817),
mRNA" ---
1459107_at 0.98269 "potassium voltage-gated channel, subfamily H (eag-related), member 3" Kcnh3
1459108_a_at 1.07657 YEATS domain containing 2 Yeats2
1459109_at 0.97697 "sema domain, seven thrombospondin repeats (type 1 and type 1-like),
transmembrane domain (TM) and short cytoplasmic domain, (semaphorin) 5A" Sema5a
1459110_at 1.00604 Myosin IXb Myo9b
1459111_at 0.97000 RIKEN cDNA 4831440E17 gene 4831440E17Rik
1459112_at 1.06541 "Adult male testis cDNA, RIKEN full-length enriched library,
clone:4930558M01 product:unclassifiable, full insert sequence" ---

1459113_at	0.92559	Neurofascin	Nfasc	
1459114_at	0.96929	Ankyrin repeat and SAM domain containing 1	Anks1	
1459115_at	1.06728	Transcribed locus	---	
1459116_at	1.02006	Neural cell adhesion molecule 2	Ncam2	
1459117_at	0.94658	Transcribed locus	---	
1459118_at	1.25019	hypothetical protein D230038C21	D230038C21	
1459119_at	1.03891	centromere protein O	Cenpo	
1459120_at	1.03216	erythrocyte protein band 4.1-like 5	Epb4.115	
1459121_at	0.96568	"9 days embryo whole body cDNA, RIKEN full-length enriched library, clone: D030030M01 product:unclassifiable, full insert sequence"	---	
1459122_at	1.03049	RIKEN cDNA C230086A09 gene	C230086A09Rik	
1459123_at	1.07337	---	---	
1459124_at	0.87856	Proprotein convertase subtilisin/kexin type 5	Pcsk5	
1459125_at	1.06486	RIKEN cDNA D030059C06 gene	D030059C06Rik	
1459126_at	0.93567	Neuronal growth regulator 1	Negr1	
1459127_at	0.92950	Transcribed locus	---	
1459128_at	1.04270	Nicotinamide nucleotide adenylyltransferase 3	Nmnat3	
1459129_at	0.97289	"Adult male testis cDNA, RIKEN full-length enriched library, clone:4930548A09 product:unclassifiable, full insert sequence"	---	
1459130_at	1.07354	"HECT domain and ankyrin repeat containing, E3 ubiquitin protein ligase 1"	Hace1	
1459131_at	1.07914	Transformation related protein 53 binding protein 1	Trp53bp1	
1459132_at	1.00625	basonuclin 2	Bnc2	
1459133_at	1.10353	"ER degradation enhancer, mannosidase alpha-like 3"	Edem3	
1459134_at	1.19192	CDC91 cell division cycle 91-like 1 (S. cerevisiae)	Cdc911	
1459135_at	1.09653	"Decapping enzyme, scavenger"	Dcps	
1459136_at	0.97886	BTB and CNC homology 2	Bach2	
1459137_at	1.04471	promyelocytic leukemia	Pml	
1459138_at	0.97071	---	---	
1459139_at	0.73355	---	---	
1459140_at	1.01587	---	---	
1459141_at	1.12546	RIKEN cDNA 1810008I18 gene	1810008I18Rik	
1459142_at	1.03731	sperm associated antigen 17	Spag17	
1459143_at	1.11919	coiled-coil-helix-coiled-coil-helix domain containing 3	Chchd3	
1459144_at	1.35113	Fibronectin type III domain containing 3a	Fndc3a	
1459145_at	1.08433	RIKEN cDNA A930033H14 gene	A930033H14Rik	
1459146_at	0.97731	mitogen-activated protein kinase 6	Mapk6	
1459147_at	1.13498	---	---	
1459148_at	1.08271	Paired box gene 3	Pax3	
1459149_at	0.92833	---	---	
1459150_at	1.07030	Leucine-rich repeats and calponin homology (CH) domain containing 1	Lrch1	
1459151_x_at	1.05737	interferon-induced protein 35	Ifi35	
1459152_at	1.02068	expressed sequence C78515	C78515	

1459153_at	1.04897	stonin 2	Ston2	
1459154_at	1.65937	Transcribed locus	---	
1459155_at	1.00569	expressed sequence	AU015696	AU015696
1459156_at	1.03472	---	---	
1459157_at	1.17226	Storkhead box 2	Stox2	
1459158_at	0.87778	---	---	
1459159_a_at	1.01522	coiled-coil domain containing 6	Ccdc6	
1459160_at	1.08314	"Gene model 879, (NCBI)"	Gm879	
1459161_at	0.97618	expressed sequence	AU015777	AU015777
1459162_at	0.91479	Zinc finger protein 644	Zfp644	
1459163_at	0.90557	---	---	
1459164_at	1.05270	expressed sequence	AU014678	AU014678
1459165_at	0.90962	"DNA segment, Chr 4, ERATO Doi 179, expressed"	D4Erttd179e	
1459166_at	0.98665	---	---	
1459167_at	1.13866	Cdc42 binding protein kinase beta	Cdc42bpb	
1459168_at	1.01110	Zinc finger protein 521	Zfp521	
1459169_at	0.90976	"Coronin, actin binding protein 1C"	Coro1c	
1459170_at	1.06504	Forkhead box O1	Foxo1	
1459171_at	1.03416	Epithelial membrane protein 1	Emp1	
1459172_at	0.97249	---	---	
1459173_at	0.97729	"protein tyrosine phosphatase, receptor type, D"	Ptprd	
1459174_at	1.02184	---	---	
1459175_at	1.09530	Transcribed locus	---	
1459176_at	1.06795	---	---	
1459177_at	1.02086	expressed sequence	AU015741	AU015741
1459178_at	1.02164	"DNA segment, Chr 8, ERATO Doi 503, expressed"	D8Erttd503e	
1459179_at	1.01926	"DNA segment, Chr 8, ERATO Doi 268, expressed"	D8Erttd268e	
1459180_at	1.02724	Transcribed locus	---	
1459181_at	1.02787	"PREDICTED: Mus musculus similar to Tubulin alpha-1 chain (LOC629601), mRNA"	---	
1459182_at	1.00131	Transcribed locus	---	
1459183_at	1.01720	"PREDICTED: Mus musculus similar to C05G5.5 (LOC230628), mRNA"	---	
1459184_at	0.92606	---	---	
1459185_at	1.56710	Nuclear protein in the AT region	Npat	
1459186_at	1.00528	expressed sequence	C80165	C80165
1459187_at	1.15949	"ST3 beta-galactoside alpha-2,3-sialyltransferase 4"	St3gal4	
1459188_at	0.95081	Receptor-like tyrosine kinase	Ryk	
1459189_at	0.95377	RIKEN cDNA 2610037D02 gene	2610037D02Rik	
1459190_at	1.16916	"DNA segment, Chr 3, ERATO Doi 258, expressed"	D3Erttd258e	
1459191_at	0.99249	RIKEN cDNA B130020M22 gene	B130020M22Rik	
1459192_at	0.97448	Transcribed locus	---	
1459193_at	0.96234	RAD51-like 1 (S. cerevisiae)	Rad5111	
1459194_at	0.99762	RIKEN cDNA 2210408I21 gene	2210408I21Rik	

1459195_at	1.02892	---	---	
1459196_at	1.13398	Dynamin binding protein	Dnmbp	
1459197_at	1.01256	---	---	
1459198_at	1.02095	"DNA segment, Chr 4, ERATO Doi 199, expressed"	D4Ertd199e	
1459199_at	1.00496	---	---	
1459200_at	0.97024	---	---	
1459201_at	1.01949	---	---	
1459202_at	0.95796	---	---	
1459203_at	1.14462	importin 8	Ipo8	
1459204_at	0.89720	expressed sequence AU015449	AU015449	
1459205_at	1.01911	Cytoplasmic polyadenylation element binding protein 1	Cpeb1	
1459206_at	0.89324	---	---	
1459207_at	1.08453	Transcribed locus	---	
1459208_at	0.95346	expressed sequence AU021889	AU021889	
1459209_at	0.97209	Ring finger protein 10	Rnf10	
1459210_at	0.97183	transmembrane protein 108	Tmem108	
1459211_at	0.67321	GLI-Kruppel family member GLI2	Gli2	
1459212_at	1.00920	anaphase promoting complex subunit 11 homolog (yeast)	Anapc11	
1459213_at	1.00991	"Adult male aorta and vein cDNA, RIKEN full-length enriched library, clone: A530096K17 product:unclassifiable, full insert sequence"	---	
1459214_at	1.07939	Odd Oz/ten-m homolog 2 (Drosophila)	Odz2	
1459215_at	0.99375	Lysosomal-associated protein transmembrane 4B	Laptm4b	
1459216_at	1.35666	Tudor domain containing 7	Tdrd7	
1459217_at	1.00225	oncostatin M receptor	Osmr	
1459218_at	1.51850	RIKEN cDNA D330001F17 gene	D330001F17Rik	
1459219_at	1.62915	---	---	
1459220_at	0.93218	expressed sequence C78651	C78651	
1459221_at	1.03193	---	---	
1459222_x_at	0.98922	---	---	
1459223_at	0.98006	RIKEN cDNA B930095G15 gene	B930095G15Rik	
1459224_at	1.08501	"Protein phosphatase 1, regulatory (inhibitor) subunit 13B"	Ppp1r13b	
1459225_at	1.10935	---	---	
1459226_at	1.07162	Transcribed locus	---	
1459227_at	0.88628	Transcribed locus	---	
1459228_at	1.03941	mirror-image polydactyly gene 1 homolog (human)	Mipol1	
1459229_at	0.94868	"7 days neonate cerebellum cDNA, RIKEN full-length enriched library, clone: A730008E08 product:unclassifiable, full insert sequence"	---	
1459230_at	0.93243	RIKEN cDNA B430319G15 gene	D530025C14Rik	
1459231_at	1.16261	protein kinase N1	Pkn1	
1459232_at	0.89656	---	---	
1459233_at	1.16895	CLIP associating protein 1	Clasp1	
1459234_at	0.77422	Arachidonate 5-lipoxygenase activating protein	Alox5ap	
1459235_at	0.98163	---	---	

1459236_at	0.95414	expressed sequence AU022436	AU022436
1459237_at	1.74938	Activating transcription factor 2	Atf2
1459238_at	1.22760	---	---
1459239_at	1.02809	Activating signal cointegrator 1 complex subunit 3	Ascc3
1459240_at	1.19254	Transcription factor 25 (basic helix-loop-helix)	Tcf25
1459241_at	0.94639	Transcribed locus	---
1459242_at	1.07128	"ELOVL family member 5, elongation of long chain fatty acids (yeast)"	Elov15
1459243_at	1.11211	"PREDICTED: Mus musculus hypothetical protein LOC626097 (LOC626097), mRNA"	---
1459244_at	0.96738	expressed sequence AU014756	AU014756
1459245_s_at	1.17401	"glutamate receptor, ionotropic, delta 2"	Grid2
1459246_at	0.99150	RIKEN cDNA D230021J17 gene	D230021J17Rik
1459247_at	0.99728	ATP/GTP binding protein-like 3	Agbl3
1459248_at	1.00997	"Protein tyrosine phosphatase, non-receptor type 9"	Ptpn9
1459249_at	1.00406	Tudor domain containing 3	Tdrd3
1459250_at	0.93827	teashirt zinc finger family member 2	Tshz2
1459251_at	1.14814	RIKEN cDNA A430088I17 gene	A430088I17Rik
1459252_at	0.95335	"similar to double homeobox, 4"	LOC245263
1459253_at	0.93956	---	---
1459254_at	0.97765	cell division cycle associated 2	Cdca2
1459255_at	0.99591	RIKEN cDNA 1700034P13 gene	1700034P13Rik
1459256_at	0.98501	Neuropilin 2	Nrp2
1459257_at	1.01775	RIKEN cDNA A230048G03 gene	A230048G03Rik
1459258_at	1.34570	RIKEN cDNA 2610021K21 gene	2610021K21Rik
1459259_at	1.06671	hypothetical protein A030007L22	A030007L22
1459260_at	0.99970	---	---
1459261_at	0.92641	SRY-box containing gene 5	Sox5
1459262_at	1.04795	MTERF domain containing 1	Mterfd1
1459263_at	0.96913	---	---
1459264_at	0.99614	tripartite motif protein 33 /// RIKEN cDNA 8030451N04 gene	Trim33 /// 8030451N04Rik
1459265_at	0.94679	myosin head domain containing 1	Myohd1
1459266_at	1.24182	---	---
1459267_at	0.85121	RIKEN cDNA 2310033P09 gene	2310033P09Rik
1459268_at	1.03154	RIKEN cDNA 5330439C02 gene	5330439C02Rik
1459269_at	0.89790	RIKEN cDNA 2310014G06 gene	2310014G06Rik
1459270_at	1.07265	"ATP-binding cassette, sub-family C (CFTR/MRP), member 1"	Abcc1
1459271_at	1.09528	Transcribed locus	---
1459272_at	1.01423	Pellino 1	Peli1
1459273_at	1.01504	---	---
1459274_at	0.99876	G protein-coupled receptor 135	Gpr135
1459275_at	0.87965	ring finger protein 17	Rnf17
1459276_at	1.05007	RIKEN cDNA A530088E08 gene	A530088E08Rik

1459277_at	0.98512 Nuclear receptor coactivator 1	Ncoa1
1459278_at	1.11178 "FYVE, RhoGEF and PH domain containing 1"	Fgd1
1459279_at	0.92536 Dedicator of cytokinesis 4	Dock4
1459280_at	0.96971 "DNA segment, Chr 1, ERATO Doi 185, expressed"	D1Erttd185e
1459281_at	0.98679 Neuropeptide Y receptor Y2	Npy2r
1459282_at	0.82825 --- ---	
1459283_at	1.51370 RIKEN cDNA 6430510B20 gene	6430510B20Rik
1459284_at	0.89897 Transcribed locus	---
1459285_at	0.96217 --- ---	
1459286_at	1.08835 Transcribed locus	---
1459287_at	0.86541 WD repeat and FYVE domain containing 3	Wdfy3
1459288_at	0.89620 "Potassium voltage-gated channel, Shal-related family, member 2"	Kcnd2
1459289_at	1.16446 "Phosphodiesterase 4D, cAMP specific"	Pde4d
1459290_at	1.12610 RIKEN cDNA C030015D19 gene	C030015D19Rik
1459291_at	0.94991 "zinc binding alcohol dehydrogenase, domain containing 1"	Zadh1
1459292_at	1.09601 Mitogen activated protein kinase kinase 6	Map2k6
1459293_at	0.99298 RIKEN cDNA C030038J10 gene	C030038J10Rik
1459294_at	1.05173 mitochondrial intermediate peptidase	Mipep
1459295_at	1.05809 Latrophilin 3	Lphn3
1459296_at	1.08578 ubiquitin-conjugating enzyme E2Z (putative)	Ube2z
1459297_at	1.09266 CDC91 cell division cycle 91-like 1 (S. cerevisiae)	Cdc9111
1459298_at	1.06796 --- ---	
1459299_at	0.90070 myosin IIIB	Myo3b
1459300_at	1.11737 Transcribed locus	---
1459301_at	1.07457 Myeloid ecotropic viral integration site-related gene 1	Mrg1
1459302_at	1.22551 RIKEN cDNA A030007N12 gene	A030007N12Rik
1459303_at	0.85176 Glycoprotein m6a	Gpm6a
1459304_at	1.01368 RIKEN cDNA A730054J21 gene	A730054J21Rik
1459305_at	1.21258 Retinoic acid induced 17	Rai17
1459306_at	1.01668 "Beaded filament structural protein 2, phakinin"	Bfsp2
1459307_at	1.57939 "SEC24 related gene family, member B (S. cerevisiae)"	Sec24b
1459308_at	0.87827 "Potassium intermediate/small conductance calcium-activated channel, subfamily N, member 3"	Kcnn3
1459309_at	0.97262 RUN and FYVE domain containing 3	Rufy3
1459310_at	0.98398 RIKEN cDNA 5330434G04 gene	5330434G04Rik
1459311_at	0.95702 "phosphodiesterase 4D, cAMP specific"	Pde4d
1459312_at	1.02229 Transcribed locus	---
1459313_at	0.96950 Transcribed locus	---
1459314_at	1.07128 CDK5 regulatory subunit associated protein 1-like 1	Cdkal1
1459315_at	1.27196 --- ---	
1459316_at	1.82864 RIKEN cDNA 2900026A02 gene	2900026A02Rik
1459317_at	0.98655 "ankyrin 2, brain"	Ank2
1459318_at	1.15733 "sema domain, transmembrane domain (TM), and cytoplasmic domain,	

(semaphorin) 6D"	Sema6d		
1459319_at	0.91417	"Polymerase (DNA directed), alpha 1"	Pola1
1459320_at	1.08215	Transcribed locus	---
1459321_at	0.98664	---	---
1459322_at	1.75352	Synaptic nuclear envelope 2	Syne2
1459323_at	0.87143	RIKEN cDNA 1700106N22 gene	1700106N22Rik
1459324_at	1.02885	---	---
1459325_at	0.92082	"7 days neonate cerebellum cDNA, RIKEN full-length enriched library, clone: A730031J22 product:unclassifiable, full insert sequence"	---
1459326_at	1.05172	"DNA segment, Chr 9, ERATO Doi 256, expressed"	D9Ertd256e
1459327_at	1.05670	RIKEN cDNA 4931406I20 gene	4931406I20Rik
1459328_at	1.29964	RIKEN cDNA 4930429A22 gene	4930429A22Rik
1459329_at	1.04351	tetratricopeptide repeat domain 7	Ttc7
1459330_at	0.99186	Ribosomal protein L31	Rpl31
1459331_at	0.94462	surfeit gene 6	Surf6
1459332_at	1.05055	itchy	Itch
1459333_at	1.06890	Transcribed locus	---
1459334_at	0.98613	---	---
1459335_at	0.97924	---	---
1459336_at	1.02594	"Guanylate cyclase 1, soluble, alpha 2"	Gucy1a2
1459337_at	0.98160	"asparagine-linked glycosylation 6 homolog (yeast, alpha-1,3,-glucosyltransferase)"	Alg6
1459338_at	1.15025	Autism susceptibility candidate 2	Auts2
1459339_at	0.80954	RIKEN cDNA 1700049G17 gene	1700049G17Rik
1459340_at	0.95052	---	---
1459341_at	1.27169	RIKEN cDNA 8030448K23 gene	8030448K23Rik
1459342_at	0.96516	hypothetical protein C230029M16	C230029M16
1459343_at	0.99682	RIKEN cDNA 2700008B19 gene	2700008B19Rik
1459344_at	1.13076	RIKEN cDNA 9630019E01 gene	9630019E01Rik
1459345_at	0.91289	---	---
1459346_at	1.09660	"tRNA splicing endonuclease 2 homolog (SEN2, S. cerevisiae)"	Tsen2
1459347_at	0.92142	---	---
1459348_at	1.07016	Early B-cell factor 1	Ebf1
1459349_at	0.75863	RIKEN cDNA A930011G23 gene	A930011G23Rik
1459350_at	1.08882	Suppressor of cytokine signaling 6	Socs6
1459351_at	1.04326	"adrenergic receptor, alpha 1a"	Adra1a
1459352_at	0.94275	RIKEN cDNA 2410003P15 gene	2410003P15Rik
1459353_at	1.20125	potassium channel tetramerisation domain containing 3	Kctd3
1459354_at	1.17910	"ST3 beta-galactoside alpha-2,3-sialyltransferase 6"	St3gal6
1459355_at	0.94532	"Transforming growth factor, beta receptor III"	Tgfbr3
1459356_at	1.05927	Dystonin	Dst
1459357_at	0.92705	Neurofascin	Nfasc
1459358_at	1.58981	Plakophilin 2	Pkp2

1459359_at 0.97744 "0 day neonate cerebellum cDNA, RIKEN full-length enriched library, clone:
C230091N23 product:unclassifiable, full insert sequence" ---
1459360_at 2.67508 Trans-acting transcription factor 3 Sp3
1459361_at 1.08335 --- ---
1459362_at 1.08749 Transcribed locus ---
1459363_at 1.26279 Ataxin 2 Atxn2
1459364_at 1.01817 RIKEN cDNA C230094A16 gene C230094A16Rik
1459365_at 1.00316 RIKEN cDNA 4930429A22 gene 4930429A22Rik
1459366_at 0.96146 "RGM domain family, member B" Rgmb
1459367_at 1.00459 "Transient receptor potential cation channel, subfamily C, member 5" Trpc5
1459368_at 1.02521 --- ---
1459369_at 1.04300 Eph receptor A6 Epha6
1459370_at 0.93046 "Calcium channel, voltage-dependent, gamma subunit 5" Cacng5
1459371_at 1.28877 Eukaryotic translation initiation factor 4E Eif4e
1459372_at 0.98767 neuronal PAS domain protein 4 Npas4
1459373_at 1.14351 Erythrocyte protein band 4.1 Epb4.1
1459374_at 1.24087 Diaphanous homolog 3 (Drosophila) Diap3
1459375_at 0.91222 Pregnancy-associated plasma protein A Pappa
1459376_at 1.05340 Human immunodeficiency virus type I enhancer binding protein 2 Hivep2
1459377_at 0.88984 paralemmin 2 Palm2
1459378_at 0.98685 "Protein phosphatase 3, catalytic subunit, beta isoform" Ppp3cb
1459379_at 1.03248 --- ---
1459380_at 1.09709 "transcriptional adaptor 1 (HFI1 homolog, yeast) like" Tada11
1459381_at 0.84826 Dachshund 1 (Drosophila) Dach1
1459382_at 0.86907 Dedicator of cyto-kinesis 2 Dock2
1459383_at 0.99231 Zinc finger protein 52 Zfp52
1459384_at 0.96802 DNA methyltransferase 3B Dnmt3b
1459385_at 0.95652 calmodulin regulated spectrin-associated protein 1-like 1 Camsap111
1459386_at 0.94757 zinc finger protein 709 Zfp709
1459387_at 0.87366 microtubule associated serine/threonine kinase family member 4 Mast4
1459388_at 1.05847 Deleted in colorectal carcinoma Dcc
1459389_at 1.03061 "Centaurin, gamma 2" Centg2
1459390_at 1.03709 Transcribed locus ---
1459391_at 0.96083 "ATP-binding cassette, sub-family A (ABC1), member 5" Abca5
1459392_at 1.07284 PHD finger protein 8 Phf8
1459393_at 0.96590 Transcription factor AP-2 beta Tcfap2b
1459394_at 1.28542 RIKEN cDNA D130052B06 gene D130052B06Rik
1459395_at 0.90466 Transcribed locus ---
1459396_at 1.02929 RIKEN cDNA 6030446N20 gene 6030446N20Rik
1459397_at 1.06322 ariadne ubiquitin-conjugating enzyme E2 binding protein homolog 1
(Drosophila) Arih1
1459398_at 1.08977 --- ---
1459399_at 1.10624 "Solute carrier family 31, member 2" Slc31a2

1459400_at	1.00265	RIKEN cDNA A630038E17 gene	A630038E17Rik
1459401_at	1.08914	Similar to zinc finger protein 616	LOC545191
1459402_at	1.02192	"opioid receptor, mu 1" Oprm1	
1459403_at	1.61514	low density lipoprotein receptor	Ldlr
1459404_at	1.12506	RIKEN cDNA 4930583H14 gene	4930583H14Rik
1459405_at	1.26005	IQ motif containing GTPase activating protein 3	Iqgap3
1459406_at	1.00593	Choline/ethanolaminephosphotransferase 1	Cept1
1459407_at	1.05380	Neurobeachin like 1	Nbeal1
1459408_at	0.90617	SEC22 vesicle trafficking protein-like A (S. cerevisiae)	Sec22a
1459409_at	1.99844	RIKEN cDNA D130073L02 gene	D130073L02Rik
1459410_at	0.97709	hypothetical LOC493582	LOC493582
1459411_at	1.16237	Serologically defined colon cancer antigen 10	Sdccag10
1459412_at	0.88696	Malignant fibrous histiocyoma amplified sequence 1	Mfhas1
1459413_at	1.00577	Trf (TATA binding protein-related factor)-proximal protein homolog	
(Drosophila)		Trfp	
1459414_at	1.07504	WD repeat and FYVE domain containing 3	Wdfy3
1459415_at	1.07715	---	---
1459416_at	1.11412	B-cell CLL/lymphoma 7A	Bcl7a
1459417_at	1.14972	Zinc finger protein 609	Zfp609
1459418_at	0.88813	RIKEN cDNA 2810457I06 gene	2810457I06Rik
1459419_at	1.13675	---	---
1459420_at	1.21410	---	---
1459421_at	1.06954	hexokinase domain containing 1	Hkdc1
1459422_at	0.97358	---	---
1459423_at	0.99883	myeloid ecotropic viral integration site 1	Meis1
1459424_at	0.79218	"DNA segment, Chr 3, ERATO Doi 108, expressed"	D3Ertd108e
1459425_at	0.89434	---	---
1459426_at	1.01728	Sciellin	Scel
1459427_at	1.01984	signal transducing adaptor molecule (SH3 domain and ITAM motif) 1	Stam
1459428_at	0.97339	expressed sequence C76662	C76662
1459429_at	1.01652	kinectin 1	Ktn1
1459430_at	0.97440	G protein-coupled receptor 158	Gpr158
1459431_at	1.01510	Nuclear factor I/A	Nfia
1459432_at	1.00421	Transcribed locus	---
1459433_at	2.05859	RIKEN cDNA C130051F05 gene	C130051F05Rik
1459434_at	1.21608	"Pyruvate dehydrogenase complex, component X"	Pdhx
1459435_at	0.99198	Transcribed locus	---
1459436_at	0.96074	"Zinc finger, RAN-binding domain containing 3"	Zranb3
1459437_at	1.09566	similar to kinesin-like motor protein C20orf23	LOC640536
protein C20orf23	LOC668880	LOC670644	LOC670644
1459438_at	0.92502	---	---
1459439_at	1.00550	mitochondrial ribosomal protein S5	Mrps5

1459440_at	0.95034	Neuron navigator 3	Nav3
1459441_at	1.09907	Odd Oz/ten-m homolog 3 (Drosophila)	Odz3
1459442_at	1.45452	"Guanine nucleotide binding protein, beta 1"	Gnb1
1459443_at	0.81350	"Potassium voltage-gated channel, subfamily Q, member 1"	Kcnq1
1459444_at	1.17949	zinc finger protein 91	Zfp91
1459445_at	0.94255	---	---
1459446_at	1.04965	leucine rich repeat containing 41	Lrrc41
1459447_at	0.98591	Transcribed locus	---
1459448_at	1.06361	cytochrome b5 reductase 2	Cyb5r2
1459449_at	1.17944	"CCR4-NOT transcription complex, subunit 4"	Cnot4
1459450_at	1.09510	Chromodomain helicase DNA binding protein 9	Chd9
1459451_at	1.09673	---	---
1459452_at	1.02397	Transcribed locus	---
1459453_at	0.96872	Roundabout homolog 1 (Drosophila)	Robo1
1459454_at	1.05718	SAP30 binding protein	Sap30bp
1459455_at	0.90803	"Arginine-rich, mutated in early stage tumors"	Armet
1459456_at	0.99511	RIKEN cDNA 0610011L14 gene	0610011L14Rik
1459457_at	1.22509	"calcium/calmodulin-dependent protein kinase II, delta"	Camk2d
1459458_at	1.01100	"DNA segment, Chr 9, ERATO Doi 338, expressed"	D9Ertd338e
1459459_at	1.01845	"Protein tyrosine phosphatase, non-receptor type 12"	Ptpn12
1459460_at	1.07091	---	---
1459461_at	1.85283	---	---
1459462_at	1.22686	Glucocorticoid induced gene 1	Gig1
1459463_at	1.07123	"Adult male spinal cord cDNA, RIKEN full-length enriched library, clone: A330073N09 product:unclassifiable, full insert sequence"	---
1459464_at	1.15562	Transcribed locus	---
1459465_at	1.04767	RIKEN cDNA 9630020C08 gene	9630020C08Rik
1459466_at	1.05244	---	---
1459467_at	1.04193	expressed sequence AA986715	AA986715
1459468_at	1.02603	expressed sequence C79743	C79743
1459469_at	1.06846	expressed sequence C78516	C78516
1459470_at	1.57238	Runt related transcription factor 1	Runx1
1459471_at	0.96273	---	---
1459472_at	0.95076	---	---
1459473_at	0.97032	"Prenyl (solanesyl) diphosphate synthase, subunit 2"	Pdss2
1459474_at	0.96553	"DNA segment, Chr 1, ERATO Doi 86, expressed"	D1Ertd86e
1459475_at	1.08217	"DNA segment, Chr 10, ERATO Doi 761, expressed"	D10Ertd761e
1459476_s_at	1.10286	cysteine and glycine-rich protein 2 binding protein	Csrp2bp
1459477_at	1.04986	Gamma-aminobutyric acid (GABA-A) receptor-associated protein-like 2	Gabarapl2
1459478_at	0.98384	expressed sequence AA408396	AA408396
1459479_at	1.02372	"PREDICTED: Mus musculus similar to Uridine 5-monophosphate synthase (UMP synthase) (LOC628594), mRNA"	---

1459480_at	0.97794	"DNA segment, Chr 6, ERATO Doi 90, expressed"	D6Ertd90e
1459481_at	1.06776	Protocadherin 11 X-linked	Pcdh11x
1459482_at	1.07259	"transient receptor potential cation channel, subfamily M, member 2"	Trpm2
1459483_at	0.93533	Transmembrane protein 28	Tmem28
1459484_at	1.01579	Transmembrane protein 2	Tmem2
1459485_at	1.54807	Neogenin	Neo1
1459486_at	1.06537	expressed sequence AU022240	AU022240
1459487_at	1.27641	RIKEN cDNA 4930554I06 gene	4930554I06Rik
1459488_at	1.51797	RIKEN cDNA 3110050K21 gene	3110050K21Rik
1459489_at	0.92033	---	---
1459490_at	0.99616	RIKEN cDNA 1110032E23 gene	1110032E23Rik
1459491_at	1.02609	"DNA segment, Chr 12, ERATO Doi 673, expressed"	D12Ertd673e
1459492_at	0.96726	---	---
1459493_at	0.96128	SET binding protein 1	Setbp1
1459494_at	0.98520	"Rabaptin, RAB GTPase binding effector protein 1"	Rabep1
1459495_at	1.13929	Transmembrane protein 16A	Tmem16a
1459496_at	0.87951	expressed sequence C77267	C77267
1459497_at	1.71704	Farnesyl diphosphate farnesyl transferase 1	Fdft1
1459498_at	1.04922	---	---
1459499_at	1.00160	Transcribed locus	---
1459500_at	1.08962	expressed sequence C85351	C85351
1459501_at	0.91432	expressed sequence AU022194	AU022194
1459502_at	1.05508	"DNA segment, Chr 17, ERATO Doi 589, expressed"	D17Ertd589e
1459503_at	0.95806	"10 days neonate cerebellum cDNA, RIKEN full-length enriched library, clone: B930063G06 product:unclassifiable, full insert sequence"	---
1459504_at	1.10110	expressed sequence AU022121	AU022121
1459505_at	0.85424	---	---
1459506_at	0.96206	---	---
1459507_at	0.90809	---	---
1459508_at	0.99166	expressed sequence C85600	C85600
1459509_at	1.13212	Mortality factor 4 like 1	Morf4l1
1459510_at	0.97987	"transmembrane protease, serine 2"	Tmprss2
1459511_at	1.01798	RIKEN cDNA 3110004L20 gene	3110004L20Rik
1459512_at	1.06601	Myeloid ecotropic viral integration site 1	Meis1
1459513_at	0.96722	---	---
1459514_at	1.03924	---	---
1459515_at	1.01596	RIKEN cDNA 2610037D02 gene	2610037D02Rik
1459516_at	1.19268	Cullin 1	Cul1
1459517_at	1.08821	CDC14 cell division cycle 14 homolog A (S. cerevisiae)	Cdc14a
1459518_at	0.85329	"Gene model 489, (NCBI)"	Gm489
1459519_at	0.81563	---	---
1459520_at	0.98265	---	---
1459521_at	0.91200	---	---

1459522_s_at 0.92642 glycogenin Gyg
1459523_at 0.97465 expressed sequence C85403 C85403
1459524_at 0.91256 mutated in colorectal cancers Mcc
1459525_at 1.06350 Calcium binding atopy-related autoantigen 1 Cbara1
1459526_at 0.99695 Trinucleotide repeat containing 6a Tnrc6a
1459527_at 0.84763 --- ---
1459528_at 1.07085 --- ---
1459529_at 1.16147 RIKEN cDNA E230016K23 gene E230016K23Rik
1459530_at 1.61713 Hydroxysteroid (17-beta) dehydrogenase 12 Hsd17b12
1459531_at 1.02731 Lemur tyrosine kinase 2 Lmtk2
1459532_at 0.91817 "glutamate receptor, metabotropic 7" Grm7
1459533_at 0.95615 "DNA segment, Chr 19, ERATO Doi 79, expressed" D19Ert79e
1459534_at 0.92539 "Adult retina cDNA, RIKEN full-length enriched library, clone:A930028F22
product:unclassifiable, full insert sequence" ---
1459535_at 0.92986 Cadherin 13 Cdh13
1459536_at 1.24436 Calcitonin receptor-like Calcr1
1459537_at 1.13267 RIKEN cDNA 4631427C17 gene 4631427C17Rik
1459538_at 1.03787 CDNA sequence BC050092 BC050092
1459539_at 1.02213 --- ---
1459540_at 0.98871 Cell division cycle 26 Cdc26
1459541_at 0.91289 --- ---
1459542_at 1.01191 --- ---
1459543_at 1.13556 expressed sequence AA407787 AA407787
1459544_at 0.94929 --- ---
1459545_at 1.03923 --- ---
1459546_s_at 0.77177 ectonucleotide pyrophosphatase/phosphodiesterase 1 Enpp1
1459547_at 1.08999 Transcribed locus ---
1459548_at 0.98239 spire homolog 1 (Drosophila) Spire1
1459549_at 0.95417 DPH5 homolog (S. cerevisiae) Dph5
1459550_at 1.01666 Synapsin III Syn3
1459551_at 0.95960 --- ---
1459552_at 0.99903 Transcribed locus ---
1459553_at 1.01229 --- ---
1459554_at 0.99040 --- ---
1459555_at 0.82217 --- ---
1459556_at 0.97983 CDNA sequence BC013481 BC013481
1459557_at 1.08983 Zinc finger and BTB domain containing 16 Zbtb16
1459558_at 1.05872 Microtubule associated serine/threonine kinase 2 Mast2
1459559_at 1.09043 --- ---
1459560_at 2.16733 RIKEN cDNA 4921517N04 gene 4921517N04Rik
1459561_at 1.08140 --- ---
1459562_at 0.91369 --- ---
1459563_x_at 0.89919 Transcribed locus ---

1459564_at	1.03511	expressed sequence C87580	C87580
1459565_at	1.04320	"ubiquitously transcribed tetratricopeptide repeat gene, Y chromosome"	Uty
1459566_at	1.07287	Regulator of G-protein signaling 12	Rgs12
1459567_at	0.85381	expressed sequence C77905	C77905
1459568_at	1.27981	Transmembrane and tetratricopeptide repeat containing 4	Tmtc4
1459569_at	0.91876	Transcribed locus	---
1459570_at	0.91625	G protein-coupled receptor 143	Gpr143
1459571_at	1.21597	SH3-binding domain glutamic acid-rich protein like	Sh3bgrl
1459572_at	1.04148	expressed sequence C86371	C86371
1459573_at	1.07713	brix domain containing 5	Bxdc5
1459574_at	1.07374	expressed sequence C86865	C86865
1459575_at	1.21548	"DNA segment, Chr 1, ERATO Doi 704, expressed"	D1ErtD704e
1459576_at	0.99744	---	---
1459577_at	1.05586	---	---
1459578_at	0.89700	expressed sequence AA407175	AA407175
1459579_at	0.82722	---	---
1459580_at	0.99204	RIKEN cDNA G630055G22 gene	G630055G22Rik
1459581_at	1.08581	transformation related protein 63	Trp63
1459582_at	1.01033	---	---
1459583_at	0.98242	Glutamate receptor interacting protein 1	Grip1
1459584_at	0.96154	vacuolar protein sorting 24 (yeast)	Vps24
1459585_at	1.16029	growth arrest specific 6	Gas6
1459586_at	1.00946	expressed sequence C78452	C78452
1459587_at	1.12685	RIKEN cDNA 2410015B03 gene	2410015B03Rik
1459588_at	0.94506	Kit oncogene	Kit
1459589_at	0.88896	"crystallin, lamda 1"	Cryl1
1459590_at	1.00781	Transcribed locus	---
1459591_at	0.96493	---	---
1459592_a_at	0.87941	similar to NFkB interacting protein 1	LOC669005
1459593_x_at	0.98752	similar to NFkB interacting protein 1	LOC669005
1459594_at	0.97888	zinc finger protein 294	Zfp294
1459595_at	1.21665	Expressed sequence AA516738	AA516738
1459596_at	0.82313	---	---
1459597_at	0.96279	myotrophin	Mtpn
1459598_at	1.11507	Hermansky-Pudlak syndrome 1 homolog (human)	Hps1
1459599_at	1.06327	---	---
1459600_at	0.97265	"Calcium/calmodulin-dependent protein kinase II, delta"	Camk2d
1459601_at	1.06181	SNF1-like kinase	Snf1lk
1459602_at	0.91824	cyclin I	Ccni
1459603_at	1.09909	RIKEN cDNA C330008K14 gene	C330008K14Rik
1459604_at	1.00578	SNRPN upstream reading frame	Snurf
1459605_at	0.94212	"amyloid beta (A4) precursor protein binding, family A, member 1"	Apba1
1459606_at	0.99502	expressed sequence C79130	C79130

1459607_at 0.93235 RIKEN cDNA 2900001O04 gene 2900001O04Rik
1459608_at 1.07888 Tripartite motif protein 24 Trim24
1459609_at 1.12904 Rho GTPase activating protein 10 Arhgap10
1459610_at 0.98422 expressed sequence C87490 C87490
1459611_at 0.94581 "membrane protein, palmitoylated 3 (MAGUK p55 subfamily member 3)"
Mpp3
1459612_at 1.05152 PH domain and leucine rich repeat protein phosphatase Phlpp
1459613_at 1.07522 Transcribed locus ---
1459614_at 1.13241 "laminin, alpha 5" Lama5
1459615_at 1.06294 RIKEN cDNA B130052P14 gene B130052P14Rik
1459616_at 1.02544 polo-like kinase 1 (Drosophila) Plk1
1459617_at 1.14602 Mitogen activated protein kinase 14 Mapk14
1459618_at 1.04750 Three prime repair exonuclease 1 Trex1
1459619_at 1.51450 erythrocyte protein band 4.1-like 2 Epb4.112
1459620_at 1.04719 RIKEN cDNA 9130004C02 gene 9130004C02Rik
1459621_at 0.93598 intersectin 1 (SH3 domain protein 1A) Itsn1
1459622_at 0.86708 "gene model 22, (NCBI)" Gm22
1459623_at 0.96050 --- ---
1459624_at 1.01206 --- ---
1459625_at 1.03998 FK506 binding protein 12-rapamycin associated protein 1 Frap1
1459626_at 0.88184 Transcribed locus ---
1459627_at 1.97076 Sterol-C4-methyl oxidase-like Sc4mol
1459628_at 1.01448 "Zinc finger, matrin type 4" Zmat4
1459629_at 0.98981 --- ---
1459630_at 0.97111 Transcribed locus ---
1459631_at 0.95513 ribosomal protein S12 /// similar to ribosomal protein S12 /// similar to
ribosomal protein S12 /// similar to ribosomal protein S12 /// similar to ribosomal protein S12 /// similar
to ribosomal protein S12 /// similar to ribosomal protein S12 /// similar to ribosomal protein S12 ///
similar to ribosomal protein S12 /// similar to ribosomal protein S12 /// similar to ribosomal protein
S12 /// similar to ribosomal protein S12 /// similar to ribosomal protein S12 /// similar to ribosomal
protein S12 /// similar to ribosomal protein S12 /// similar to 40S ribosomal protein S12 /// similar to
ribosomal protein S12 /// similar to ribosomal protein S12 /// similar to ribosomal protein S12 /// similar
to ribosomal protein S12 Rps12 /// LOC432865 /// LOC434843 /// LOC546371 /// LOC626048 ///
LOC627835 /// LOC665463 /// LOC665501 /// LOC665522 /// LOC665555 /// LOC665579 ///
LOC665611 /// LOC668354 /// LOC670751 /// LOC670832 /// LOC671641 /// LOC672008 ///
LOC672415 /// LOC673522 /// LOC676277
1459632_at 3.56592 Cysteine rich transmembrane BMP regulator 1 (chordin like) Crim1
1459633_at 1.06510 --- ---
1459634_at 0.88737 RIKEN cDNA A230048G03 gene A230048G03Rik
1459635_at 1.29393 "Discs, large homolog 1 (Drosophila)" Dlgh1
1459636_at 1.14354 Angiopoietin 2 Angpt2
1459637_at 1.00563 --- ---
1459638_at 0.94202 "Protein tyrosine phosphatase, receptor type, E" Ptpre

1459639_at 0.95546 BR serine/threonine kinase 2 Brsk2
1459640_at 1.02779 --- ---
1459641_at 1.00489 "Proteasome (prosome, macropain) subunit, beta type 2" Psmb2
1459642_at 1.09781 A disintegrin and metallopeptidase domain 23 Adam23
1459643_at 0.91686 Transcribed locus ---
1459644_at 0.92947 expressed sequence AU020772 AU020772
1459645_at 0.99915 Adenosine kinase Adk
1459646_at 0.93266 heparan sulfate (glucosamine) 3-O-sulfotransferase 6 Hs3st6
1459647_at 1.12181 "Polymerase (DNA-directed), delta 3, accessory subunit" Pold3
1459648_at 1.16726 RUN and TBC1 domain containing 3 Rutbc3
1459649_at 1.14286 Transcribed locus ---
1459650_at 0.89606 telomeric repeat binding factor 2 Terf2
1459651_s_at 1.04196 nucleoporin 54 Nup54
1459652_at 0.93705 expressed sequence C79356 C79356
1459653_at 1.02504 --- ---
1459654_at 1.01848 --- ---
1459655_at 0.98784 --- ---
1459656_at 0.90286 "PREDICTED: Mus musculus similar to potassium voltage-gated channel, subfamily G, member 1 isoform 2, transcript variant 2 (LOC241794), mRNA" ---
1459657_s_at 1.05965 RNA polymerase 1-3 Rpo1-3
1459658_at 0.88930 --- ---
1459659_at 1.02151 "PREDICTED: Mus musculus hypothetical protein LOC623278 (LOC623278), mRNA" ---
1459660_at 0.94555 --- ---
1459661_at 1.04561 expressed sequence AW492955 AW492955
1459662_at 0.98373 "Adult male diencephalon cDNA, RIKEN full-length enriched library, clone:9330197M14 product:unclassifiable, full insert sequence" ---
1459663_at 0.98511 --- ---
1459664_at 1.10935 WD repeat domain 31 Wdr31
1459665_s_at 1.04938 MRV integration site 1 Mrvi1
1459666_at 0.87818 "trafficking protein, kinesin binding 1" Trak1
1459667_at 1.34232 Rho GTPase activating protein 17 Arhgap17
1459668_at 1.12737 TRAF family member-associated Nf-kappa B activator Tank
1459669_at 0.88236 Transcribed locus ---
1459670_at 1.37231 Heterogeneous nuclear ribonucleoprotein L-like Hnrpll
1459671_at 0.91377 Transcribed locus ---
1459672_at 1.10186 Topoisomerase (DNA) I Top1
1459673_at 0.97680 general transcription factor III C 1 Gtf3c1
1459674_at 1.05622 "Protein kinase C, beta 1" Prkcb1
1459675_at 0.90281 --- ---
1459676_at 0.98952 tetratricopeptide repeat domain 15 Ttc15
1459677_at 1.43208 --- ---
1459678_at 1.30164 hypothetical LOC497255 LOC497255

1459679_s_at 1.02355 myosin IB Myo1b
1459680_at 1.27479 Translocase of outer mitochondrial membrane 40 homolog (yeast) Tomm40
1459681_at 0.90692 --- ---
1459682_at 0.94996 hypothetical LOC434249 LOC434249
1459683_at 1.04506 similar to spermatogenic cell-specific gene 2 LOC218695
1459684_at 0.97846 RIKEN cDNA 2810449C10 gene 2810449C10Rik
1459685_at 0.99833 expressed sequence C77683 C77683
1459686_at 0.89723 --- ---
1459687_x_at 1.06500 Cleavage and polyadenylation specific factor 6 Cpsf6
1459688_at 1.20722 Transcribed locus ---
1459689_at 1.13710 "Glycerol phosphate dehydrogenase 2, mitochondrial" Gpd2
1459690_at 1.14806 --- ---
1459691_at 1.10743 --- ---
1459692_at 0.92309 expressed sequence AI661323 AI661323
1459693_x_at 0.97039 "3-monooxygenase/tryptophan 5-monooxygenase activation protein, gamma polypeptide" Ywhag
1459694_at 0.91741 expressed sequence AI428898 AI428898
1459695_at 1.33599 --- ---
1459696_at 0.90623 Furry homolog (Drosophila) Fry
1459697_at 1.09108 Thyroid adenoma associated Thada
1459698_at 1.04442 "Similar to retinoic acid, EGF, and NGF upregulated" LOC622320
1459699_at 0.88543 Transcribed locus ---
1459700_at 0.96958 LEM domain containing 3 Lemd3
1459701_x_at 1.07713 --- ---
1459702_at 1.44487 --- ---
1459703_at 1.07487 "DNA segment, Chr 7, ERATO Doi 443, expressed" D7Ertd443e
1459704_at 1.06910 "DnaJ (Hsp40) homolog, subfamily C, member 7" Dnajc7
1459705_at 0.94982 Olfactomedin 3 Olfm3
1459706_at 1.01347 Exocyst complex component 4 Exoc4
1459707_at 1.56121 Phosphofurin acidic cluster sorting protein 1 Pacs1
1459708_at 0.92135 "Ubiquitination factor E4A, UFD2 homolog (S. cerevisiae)" Ube4a
1459709_at 0.93478 --- ---
1459710_at 0.98395 Transcribed locus ---
1459711_at 0.84914 sidekick homolog 1 (chicken) Sdk1
1459712_at 1.02894 zinc finger protein 182 Zfp182
1459713_s_at 0.74383 transmembrane protein 16A Tmem16a
1459714_at 0.87741 RIKEN cDNA 6330545A04 gene 6330545A04Rik
1459715_at 1.36147 RIKEN cDNA 1500011J06 gene 1500011J06Rik
1459716_at 1.01615 expressed sequence AI835735 AI835735
1459717_at 1.01259 Olfactory receptor 1344 Olfr1344
1459718_x_at 0.96810 Kruppel-like factor 6 Klf6
1459719_at 1.00196 DEP domain containing 6 Depdc6
1459720_x_at 0.91870 --- ---

1459721_at 1.03297 CDNA clone IMAGE:6406267 ---
 1459722_at 2.65200 "Zinc finger, SWIM domain containing 6" Zswim6
 1459723_at 0.89876 "zinc finger, DHHC-type containing 22" Zdhhc22
 1459724_at 0.82991 "zinc finger, DHHC domain containing 20" Zdhhc20
 1459725_s_at 1.14268 demilune cell and parotid protein /// similar to common salivary protein 1
 Dcpp /// LOC630537
 1459726_at 0.98308 "Protein tyrosine phosphatase, receptor type, N polypeptide 2" Ptprn2
 1459727_at 0.92120 expressed sequence AW125646 AW125646
 1459728_at 0.97606 RIKEN cDNA 5830446M03 gene 5830446M03Rik
 1459729_at 0.98688 "solute carrier family 13 (sodium-dependent citrate transporter), member 5"
 Slc13a5
 1459730_at 1.00823 Transcribed locus ---
 1459731_at 0.90999 --- ---
 1459732_at 1.01160 Transcribed locus ---
 1459733_at 2.24130 Phosphoprotein enriched in astrocytes 15 Pea15
 1459734_at 1.65595 "Proteasome (prosome, macropain) 26S subunit, non-ATPase, 14" Psm14
 1459735_at 1.09783 RIKEN cDNA C430014M02 gene C430014M02Rik
 1459736_at 0.88880 Serine/threonine kinase 10 Stk10
 1459737_s_at 0.87765 transthyretin Ttr
 1459738_x_at 1.06732 --- ---
 1459739_at 1.09977 RIKEN cDNA E130308A19 gene E130308A19Rik
 1459740_s_at 0.92718 "uncoupling protein 2 (mitochondrial, proton carrier)" Ucp2
 1459741_x_at 0.94129 "uncoupling protein 2 (mitochondrial, proton carrier)" Ucp2
 1459742_at 0.96188 "15 days embryo brain cDNA, RIKEN full-length enriched library, clone:
 G630032A08 product:unclassifiable, full insert sequence" ---
 1459743_at 0.95589 Transcribed locus ---
 1459744_at 0.97749 CD53 antigen Cd53
 1459745_at 1.10236 Ubiquitin-associated protein 2 Ubap2
 1459746_at 2.19075 R3H domain 1 (binds single-stranded nucleic acids) R3hdm1
 1459747_at 0.74599 --- ---
 1459748_at 0.96039 --- ---
 1459749_s_at 0.74811 FAT tumor suppressor homolog 4 (Drosophila) Fat4
 1459750_s_at 1.02299 G protein-coupled receptor 123 Gpr123
 1459751_s_at 0.94982 "protein phosphatase 1, regulatory (inhibitor) subunit 16A" Ppp1r16a
 1459752_at 1.07077 mutated in colorectal cancers Mcc
 1459753_x_at 0.96816 transformer 2 alpha homolog (Drosophila) Tra2a
 1459754_x_at 0.78436 eukaryotic translation elongation factor 1 delta (guanine nucleotide exchange
 protein) Eef1d
 1459755_x_at 0.95711 RIKEN cDNA 1700082C02 gene 1700082C02Rik
 1459756_at 0.92955 "CCR4-NOT transcription complex, subunit 10" Cnot10
 1459757_x_at 0.90618 "CCR4-NOT transcription complex, subunit 10" Cnot10
 1459758_at 0.94289 upstream transcription factor 1 Usf1
 1459759_s_at 0.98336 RIKEN cDNA 1700065I17 gene 1700065I17Rik

1459760_at 0.93813 NADH dehydrogenase (ubiquinone) Fe-S protein 4 Ndufs4
1459761_x_at 0.89220 "immunoglobulin superfamily, member 4" Igsf4d
1459762_x_at 1.05988 WD repeat domain 12 Wdr12
1459763_at 1.10154 "Adiponectin, C1Q and collagen domain containing" Adipoq
1459764_x_at 1.44549 Sorting nexin 10 Snx10
1459765_s_at 1.00968 Splicing factor 1 Sf1
1459766_x_at 0.84430 --- ---
1459767_x_at 1.00289 chondroitin sulfate proteoglycan 2 Cspg2
1459768_x_at 1.11647 "RAP1, GTP-GDP dissociation stimulator 1" Rap1gds1
1459769_at 1.09252 Zinc finger like protein 1 Zfp11
1459770_at 0.90949 TAO kinase 1 Taok1
1459771_x_at 1.04266 "phytoceramidase, alkaline" Phca
1459772_at 0.94512 Hermansky-Pudlak syndrome 3 homolog (human) Hps3
1459773_x_at 0.99154 Smad nuclear interacting protein 1 Snip1
1459774_at 0.92207 Tetraspanin 5 Tspan5
1459775_at 0.91698 3-phosphoinositide dependent protein kinase-1 Pdpk1
1459776_x_at 1.04310 3-phosphoinositide dependent protein kinase-1 Pdpk1
1459777_at 0.91744 "interleukin 1 receptor, type I" Il1r1
1459778_at 1.04276 Ribophorin I Rpn1
1459779_s_at 1.19488 SET domain containing (lysine methyltransferase) 8 Setd8
1459780_at 0.90233 transformation related protein 53 Trp53
1459781_x_at 0.89951 transformation related protein 53 Trp53
1459782_x_at 0.82873 "ATP-binding cassette, sub-family C (CFTR/MRP), member 5" Abcc5
1459783_s_at 0.95470 cappuccino Cno
1459784_x_at 0.94284 RIKEN cDNA 1300002A08 gene 1300002A08Rik
1459785_at 0.92911 Zinc finger protein 383 Zfp383
1459786_at 0.99279 Transcribed locus ---
1459787_at 0.94600 --- ---
1459788_at 1.10705 expressed sequence AI790205 AI790205
1459789_at 0.95069 --- ---
1459790_x_at 1.12492 aristaless 3 Alx3
1459791_at 1.00966 "DnaJ (Hsp40) homolog, subfamily C, member 1" Dnajc1
1459792_at 1.00525 --- ---
1459793_s_at 0.79037 --- ---
1459794_at 1.07197 BCL2-like 11 (apoptosis facilitator) Bcl2l11
1459795_at 0.92994 G protein-coupled receptor 85 Gpr85
1459796_at 1.04753 RIKEN cDNA 1700061E18 gene 1700061E18Rik
1459797_at 0.97033 Erythrocyte protein band 4.1-like 2 Epb4.112
1459798_x_at 1.00437 RIKEN cDNA 4732479N06 gene 4732479N06Rik
1459799_at 0.92638 "FERM, RhoGEF (Arhgef) and pleckstrin domain protein 1 (chondrocyte-derived)" Farp1
1459800_s_at 0.91131 mitogen activated protein kinase kinase kinase 4 Map3k4
1459801_at 0.78876 "UDP-Gal:betaGlcNAc beta 1,3-galactosyltransferase, polypeptide 5"

B3galt5

1459802_at 0.91800 --- ---
 1459803_x_at 1.02793 RIKEN cDNA 2810408A11 gene 2810408A11Rik
 1459804_at 0.75978 CREB binding protein Crebbp
 1459805_x_at 1.01899 dihydrouridine synthase 3-like (*S. cerevisiae*) Dus3l
 1459806_x_at 0.97121 mitochondrial ribosomal protein S23 Mrps23
 1459807_x_at 1.06972 RIKEN cDNA 4933406E20 gene 4933406E20Rik
 1459808_at 0.97531 FK506 binding protein 4 Fkbp4
 1459809_x_at 0.93054 RIKEN cDNA 1700063D05 gene 1700063D05Rik
 1459810_at 1.03063 RIKEN cDNA 1110008F13 gene 1110008F13Rik
 1459811_at 1.05464 metal response element binding transcription factor 2 Mtf2
 1459812_x_at 0.91592 tetraspanin 7 Tspan7
 1459813_at 0.89938 RIKEN cDNA 1700012D01 gene 1700012D01Rik
 1459814_at 0.96598 transmembrane protein 109 Tmem109
 1459815_at 0.89211 "zinc finger, MYND domain containing 12" Zmynd12
 1459816_x_at 1.03292 --- ---
 1459817_at 1.03052 --- ---
 1459818_x_at 0.97560 "zinc finger, MYM-type 3" Zmym3
 1459819_at 0.94551 RIKEN cDNA 5330439K02 gene 5330439K02Rik
 1459820_at 0.93816 RIKEN cDNA A630082K20 gene A630082K20Rik
 1459821_x_at 1.02150 RIKEN cDNA A630082K20 gene A630082K20Rik
 1459822_at 0.88321 RAE1 RNA export 1 homolog (*S. pombe*) Rae1
 1459823_at 0.97605 EH-domain containing 2 /// similar to EH-domain containing 2 Ehd2 ///
 LOC673251
 1459824_at 0.94172 "SWI/SNF related, matrix associated, actin dependent regulator of chromatin, subfamily c, member 1" Smarcc1
 1459825_x_at 0.93463 insulin-like growth factor 2 mRNA binding protein 1 Igf2bp1
 1459826_at 1.00873 "Potassium voltage-gated channel, subfamily Q, member 2" Kcnq2
 1459827_x_at 1.00229 Hermansky-Pudlak syndrome 1 homolog (human) Hps1
 1459828_at 1.07650 Serine/arginine repetitive matrix 1 Srrm1
 1459829_x_at 1.04329 CDNA sequence BC031781 BC031781
 1459830_at 1.02741 Transcribed locus ---
 1459831_s_at 1.04785 transmembrane and coiled-coil domains 5 Tmco5
 1459832_s_at 0.98703 "adaptor-related protein complex AP-1, mu subunit 1" Ap1m1
 1459833_x_at 0.94108 calcium and integrin binding family member 2 Cib2
 1459834_x_at 1.08641 --- ---
 1459835_s_at 0.82054 "DnaJ (Hsp40) homolog, subfamily A, member 1" Dnaj1
 1459836_x_at 1.06330 "splicing factor 3b, subunit 2" Sf3b2
 1459837_at 1.08247 Cytochrome b-561 domain containing 1 Cyb561d1
 1459838_s_at 0.64007 BTB (POZ) domain containing 11 Btbd11
 1459839_x_at 0.97246 BTB (POZ) domain containing 11 Btbd11
 1459840_s_at 1.03980 coiled coil domain containing 28B Ccdc28b
 1459841_x_at 0.92322 lysosomal-associated protein transmembrane 5 Laptm5

1459842_x_at 0.81952 nucleotide binding protein 2 Nubp2
1459843_s_at 0.84529 MAD homolog 1 (Drosophila) Smad1
1459844_at 0.93889 TNFAIP3 interacting protein 2 Tnip2
1459845_at 1.04063 Chromobox homolog 5 (Drosophila HP1a) Cbx5
1459846_x_at 0.99965 transmembrane protein 4 Tmem4
1459847_x_at 0.88483 glial cell line derived neurotrophic factor family receptor alpha 2 Gfra2
1459848_x_at 0.98121 dopamine beta hydroxylase Dbh
1459849_x_at 1.11893 valosin containing protein (p97)/p47 complex interacting protein 1 Vcpip1
1459850_x_at 0.88297 "glycine receptor, beta subunit" Glrb
1459851_x_at 0.86060 RIO kinase 1 (yeast) Riok1
1459852_x_at 0.98463 tribbles homolog 2 (Drosophila) Trib2
1459853_x_at 0.99134 --- ---
1459854_s_at 0.86232 dynein light chain Tctex-type 3 Dynlt3
1459855_x_at 0.99050 ARP1 actin-related protein 1 homolog A (yeast) Actr1a
1459856_at 0.97924 Mitochondrial ribosomal protein L13 Mrpl13
1459857_at 1.47634 ubiquitin specific peptidase 32 Usp32
1459858_x_at 0.92723 cartilage acidic protein 1 Crtac1
1459859_x_at 1.08017 chromatin accessibility complex 1 Chrac1
1459860_x_at 0.78962 tripartite motif protein 2 Trim2
1459861_s_at 1.00873 F-box and leucine-rich repeat protein 10 Fbxl10
1459862_at 0.94739 Transcribed locus ---
1459863_x_at 0.88000 "golgi associated, gamma adaptin ear containing, ARF binding protein 1" Ggal
1459864_at 1.11982 G protein-coupled receptor 146 Gpr146
1459865_x_at 1.04403 carboxylesterase 7 Ces7
1459866_x_at 0.97616 non imprinted in Prader-Willi/Angelman syndrome 2 homolog (human)
Nipa2
1459867_x_at 0.91666 unc-45 homolog A (C. elegans) Unc45a
1459868_x_at 0.85235 "interleukin 11 receptor, alpha chain 1 /// interleukin 11 receptor, alpha chain
2 /// similar to interleukin 11 receptor, alpha chain 2" Il11ra1 /// Il11ra2 /// LOC621887
1459869_x_at 1.08146 RIKEN cDNA 4930402E16 gene 4930402E16Rik
1459870_x_at 0.92813 "Interleukin 2 receptor, beta chain" Il2rb
1459871_x_at 0.97713 membrane-associated ring finger (C3HC4) 2 2-Mar
1459872_x_at 0.78165 "histocompatibility 2, class II, locus DMa" H2-DMa
1459873_x_at 0.86556 "Melanoma antigen, family D, 2" Maged2
1459874_s_at 0.91693 myotubularin related protein 4 Mtmr4
1459875_x_at 0.98455 RIKEN cDNA 5730494M16 gene 5730494M16Rik
1459876_at 0.93389 Natriuretic peptide receptor 1 Npr1
1459877_x_at 0.91306 Ring finger protein 185 Rnf185
1459878_a_at 0.94205 RIKEN cDNA A430107O13 gene A430107O13Rik
1459879_at 1.67445 RIKEN cDNA 4921513D23 gene 4921513D23Rik
1459880_at 1.36479 heterogeneous nuclear ribonucleoprotein A/B Hnrpab
1459881_at 1.02157 expressed sequence AI595406 AI595406
1459882_at 1.09190 --- ---

1459883_at 0.94738 "Adult male epididymis cDNA, RIKEN full-length enriched library, clone:9230001K01 product:unclassifiable, full insert sequence" ---

1459884_at 0.90709 "cytochrome c oxidase, subunit VIIc" Cox7c

1459885_s_at 0.84279 "cytochrome c oxidase, subunit VIIc" Cox7c

1459886_at 0.92327 Transcribed locus ---

1459887_at 1.19338 Transcribed locus ---

1459888_at 0.95084 hypothetical protein LOC545261 LOC545261

1459889_at 0.96837 chloride channel calcium activated 3 Clca3

1459890_s_at 1.15532 RIKEN cDNA 1110008P14 gene 1110008P14Rik

1459891_at 0.97423 expressed sequence C78444 C78444

1459892_at 0.89291 expressed sequence AA408156 AA408156

1459893_at 0.90508 expressed sequence C76472 C76472

1459894_at 0.94988 IQ motif containing GTPase activating protein 2 Iqgap2

1459895_at 0.86789 RIKEN cDNA C030010B13 gene C030010B13Rik

1459896_at 0.87265 pogo transposable element with KRAB domain Pogk

1459897_a_at 0.89559 suprabasin Sbsn

1459898_at 1.10439 suprabasin Sbsn

1459900_at 1.19128 expressed sequence C79468 C79468

1459901_at 0.98180 hypothetical protein 6820429M01 6820429M01

1459902_at 1.00928 RIKEN cDNA 2700007P21 gene 2700007P21Rik

1459903_at 0.97561 "sema domain, immunoglobulin domain (Ig), and GPI membrane anchor, (semaphorin) 7A" Sema7a

1459904_at 0.89177 cDNA sequence BC030870 BC030870

1459905_at 1.12459 "10 days neonate cerebellum cDNA, RIKEN full-length enriched library, clone: B930063G06 product:unclassifiable, full insert sequence" ---

1459906_at 0.93899 "Diacylglycerol kinase, eta" Dgkh

1459907_a_at 1.18629 Wolf-Hirschhorn syndrome candidate 1-like 1 (human) Whsc111

1459908_at 1.14687 "CD47 antigen (Rh-related antigen, integrin-associated signal transducer)" Cd47

1459909_at 0.98729 nuclear factor I/X Nfix

1459910_at 0.78267 "tankyrase, TRF1-interacting ankyrin-related ADP-ribose polymerase" Tnks

1459911_at 0.92671 cerebellar degeneration-related protein 2-like Cdr2l

1459912_at 0.92815 mitogen-activated protein kinase kinase kinase 4 Map4k4

1459913_at 0.99688 "tumor necrosis factor (ligand) superfamily, member 10" Tnfsf10

1459914_at 0.95618 RIKEN cDNA 2810402A17 gene 2810402A17Rik

1459916_at 1.08917 Paired box gene 5 Pax5

1459917_at 2.42144 expressed sequence AI451896 AI451896

1459918_at 1.00087 budding uninhibited by benzimidazoles 3 homolog (S. cerevisiae) Bub3

1459919_a_at 0.92265 "MOB1, Mps One Binder kinase activator-like 2C (yeast)" Mobk12c

1459920_at 1.01697 RIKEN cDNA 4933415E08 gene 4933415E08Rik

1459921_at 0.94034 neurexophilin 1 Nxph1

1459922_at 1.01671 expressed sequence AI646383 AI646383

1459923_at 0.92587 brain expressed gene 6 Bex6

1459924_at	0.88299	"ATPase, H ⁺ transporting, lysosomal V0 subunit A1"	Atp6v0a1
1459925_at	1.11722	expressed sequence C78282	C78282
1459926_at	1.04758	expressed sequence C77068	C77068
1459927_at	1.07072	RIKEN cDNA 4833445I07 gene	4833445I07Rik
1459928_at	1.21121	ligand dependent nuclear receptor corepressor	Lcor
1459929_at	1.02358	zinc finger protein 568	Zfp568
1459930_at	0.86830	Ubiquitin-conjugating enzyme E2H	Ube2h
1459931_a_at	0.93944	Similar to zinc finger protein 709	LOC626877
1459932_at	0.89848	"DNA segment, Chr 8, ERATO Doi 28, expressed"	D8Ertd28e
1459933_at	0.99478	---	---
1459934_at	1.11104	ubiquitin specific peptidase 42	Usp42
1459935_at	1.08483	EST AA517562	AA517562
1459936_at	0.93432	Transcribed locus	---
1459937_at	0.98392	Argininosuccinate synthetase 1	Ass1
1459938_at	1.05160	RIKEN cDNA C330024D21 gene	C330024D21Rik
1459939_at	1.01757	Transcribed locus	---
1459940_at	1.09491	---	---
1459941_at	1.02849	RIKEN cDNA 4933402J24 gene	4933402J24Rik
1459942_at	1.03955	Open reading frame 34	ORF34
1459943_at	0.86687	RIKEN cDNA A030012G06 gene	A030012G06Rik
1459944_at	1.16032	"UTP11-like, U3 small nucleolar ribonucleoprotein, (yeast)"	Utp11
1459945_at	1.10406	"Adult male corpora quadrigemina cDNA, RIKEN full-length enriched library, clone:B230215A15 product:hypothetical protein, full insert sequence"	---
1459946_at	1.00680	"Adult male diencephalon cDNA, RIKEN full-length enriched library, clone:9330150G21 product:unclassifiable, full insert sequence"	---
1459947_at	0.93758	Bone morphogenetic protein 6	Bmp6
1459948_at	1.00657	Growth hormone receptor	Ghr
1459949_at	1.23187	"ATP synthase, H ⁺ transporting, mitochondrial F0 complex, subunit s"	Atp5s
1459950_at	0.95330	Doublecortin and CaM kinase-like 2	Dcamk12
1459951_at	1.34503	"ribosomal protein S6 kinase, polypeptide 1"	Rps6kb1
1459952_at	1.01505	Ankyrin repeat domain 17	Ankrd17
1459953_at	1.12210	Expressed sequence AW011752	AW011752
1459954_at	1.06153	---	---
1459955_at	1.03499	RIKEN cDNA 5830484A20 gene	5830484A20Rik
1459956_at	0.93029	---	---
1459957_at	1.53651	Trinucleotide repeat containing 6a	Tnrc6a
1459958_at	1.75394	arginine/serine-rich coiled-coil 1	Rsrc1
1459959_at	1.08755	Glycine decarboxylase	Glhc
1459960_at	1.00418	Transcribed locus	---
1459961_a_at	1.12543	Signal transducer and activator of transcription 3	Stat3
1459962_at	1.09214	RIKEN cDNA 4930523C07 gene	4930523C07Rik
1459963_at	1.04733	"Polymerase (DNA directed), alpha 1"	Pola1
1459964_at	0.98609	Jun proto-oncogene related gene d1	Jund1

1459965_at 0.93912 Dyslexia susceptibility 1 candidate 1 homolog (human) Dyx1c1
1459966_at 1.23713 Oxysterol binding protein-like 7 Osbpl7
1459967_at 1.02021 Transcribed locus ---
1459968_at 1.27632 "RNA binding motif, single stranded interacting protein 1" Rbms1
1459969_x_at 1.06934 Transcribed locus ---
1459970_at 1.05698 Uronyl-2-sulfotransferase Ust
1459971_at 0.96263 "Potassium channel, subfamily T, member 2" Kcnt2
1459972_x_at 0.78876 seizure related gene 6 Sez6
1459973_x_at 0.90328 dipeptidylpeptidase 4 Dpp4
1459974_x_at 1.02408 RIKEN cDNA 2700008N14 gene 2700008N14Rik
1459975_at 1.04752 Transcribed locus ---
1459976_s_at 1.14694 "superoxide dismutase 1, soluble" Sod1
1459977_x_at 0.98506 "COX10 homolog, cytochrome c oxidase assembly protein, heme A:
farnesyltransferase (yeast)" Cox10
1459978_x_at 1.06265 cyclin J-like Ccnj1
1459979_x_at 0.91544 zinc finger protein 68 Zfp68
1459980_x_at 0.98503 "RAB3A, member RAS oncogene family" Rab3a
1459981_s_at 1.08005 "rosbin, round spermatid basic protein 1" Rsb1
1459982_a_at 0.89403 --- ---
1459983_at 0.97022 inositol (myo)-1(or 4)-monophosphatase 2 Impa2
1459984_at 0.98512 melanoma inhibitory activity 3 Mia3
1459985_at 1.10772 WD repeat domain 61 Wdr61
1459986_a_at 1.03782 ribosomal protein S17 Rps17
1459987_s_at 1.05413 chaperonin subunit 3 (gamma) Cct3
1459988_at 0.89699 methyl-CpG binding domain protein 3-like 2 Mbd3l2
1459989_at 0.97035 Transcribed locus ---
1459990_at 0.93605 "discoidin domain receptor family, member 1" Ddr1
1459991_at 1.09930 myosin IXa Myo9a
1459992_x_at 0.93281 "ceroid-lipofuscinosis, neuronal 8" Cln8
1459993_at 0.95393 "0 day neonate lung cDNA, RIKEN full-length enriched library, clone:
E030043J12 product:twist homolog 2 (Drosophila), full insert sequence" ---
1459994_x_at 1.01868 transferrin receptor 2 Trfr2
1459995_at 0.94196 RIKEN cDNA 1700015G11 gene 1700015G11Rik
1459996_at 0.85176 "Calcium channel, voltage-dependent, P/Q type, alpha 1A subunit" Cacna1a
1459997_s_at 0.83232 transmembrane protein 17 Tmem17
1459998_at 0.93392 zinc finger protein 407 Zfp407
1459999_at 0.97627 "splicing factor, arginine/serine-rich 15 /// similar to splicing factor, arginine/
serine-rich 15" Sfrs15 /// LOC670368
1460000_at 0.97520 RIKEN cDNA D830007B15 gene D830007B15Rik
1460001_at 0.89145 pyroglutamyl-peptidase I Pgpep1
1460002_at 1.71619 tousled-like kinase 1 Tlk1
1460003_at 1.04720 expressed sequence AI956758 AI956758
1460004_x_at 0.99013 syntaxin 6 Stx6

1460005_at 0.99545 "expressed sequence AF013969 /// similar to CG5514-PB, isoform B /// region containing expressed sequence AF013969; RIKEN cDNA A230054D04 gene" AF013969 /// LOC665775 /// LOC675766

1460006_at 0.94500 AT motif binding factor 1 Atbf1

1460007_at 0.91688 thymocyte nuclear protein 1 Thyn1

1460008_x_at 1.01144 ERO1-like (S. cerevisiae) /// ribosomal protein L31 /// similar to ribosomal protein L31 /// similar to ribosomal protein L31 /// similar to ribosomal protein L31 Ero11 /// Rpl31 /// LOC667682 /// LOC672957 /// LOC676592

1460009_at 0.89074 Transcribed locus ---

1460010_a_at 0.90665 phosphatidylserine synthase 2 Ptdss2

1460011_at 0.78438 "cytochrome P450, family 26, subfamily b, polypeptide 1" Cyp26b1

1460012_at 0.98749 WAP four-disulfide core domain 3 Wfdc3

1460013_at 1.02621 neuroligin 2 Nlgn2

1460014_at 0.86283 triggering receptor expressed on myeloid cells-like 4 Trem14

1460015_at 1.01524 spermatogenesis and oogenesis specific basic helix-loop-helix 1 Sohlh1

1460016_at 1.29915 transmembrane protein 164 Tmem164

1460017_at 1.03255 hypothetical LOC434179 LOC434179

1460018_at 0.90110 "DNA segment, Chr 11, ERATO Doi 759, expressed" D11Ert759e

1460019_at 0.94862 similar to leucine rich repeat containing 45 LOC627607

1460020_at 1.03341 hypothetical protein LOC629623 LOC629623

1460021_at 0.99137 hypothetical protein LOC626231 LOC626231

1460022_at 0.92488 "ATP-binding cassette, sub-family B (MDR/TAP), member 8" Abcb8

1460023_at 0.97851 --- ---

1460024_at 0.90002 trinucleotide repeat containing 6b Tnrc6b

1460025_at 1.00690 leucine-rich repeats and immunoglobulin-like domains 2 Lrig2

1460026_s_at 0.95418 "zinc finger, FYVE domain containing 27" Zfyve27

1460027_at 0.95282 cDNA sequence BC061194 BC061194

1460028_at 0.88000 glutamate receptor interacting protein 2 Grip2

1460029_at 1.03746 GTP binding protein 5 Gtpbp5

1460030_at 1.05973 "HECT, C2 and WW domain containing E3 ubiquitin protein ligase 1" Hecw1

1460031_at 1.00368 G protein-coupled receptor 45 Gpr45

1460032_at 2.49940 RIKEN cDNA 4933407C03 gene 4933407C03Rik

1460033_at 0.95166 RIKEN cDNA C030002C11 gene C030002C11Rik

1460034_at 0.93685 sterile alpha motif domain containing 4B Samd4b

1460035_at 1.27894 prohibitin 2 Phb2

1460036_at 1.14184 "adaptor-related protein complex 1, sigma 2 subunit" Ap1s2

1460037_at 1.14775 RIKEN cDNA 2610510H03 gene 2610510H03Rik

1460038_at 0.91706 "POU domain, class 3, transcription factor 1" Pou3f1

1460039_at 0.84439 "C-type lectin domain family 1, member a" Clec1a

1460040_at 0.95991 RIKEN cDNA 4933407H18 gene 4933407H18Rik

1460041_at 0.87131 RIKEN cDNA D630040I23 gene D630040I23Rik

1460042_at 0.95576 "solute carrier family 23 (nucleobase transporters), member 3" Slc23a3

1460043_at 0.99689 --- ---
1460044_at 0.91037 "0 day neonate cerebellum cDNA, RIKEN full-length enriched library, clone: C230062K19 product:unclassifiable, full insert sequence" ---
1460045_at 1.00509 "cadherin 7, type 2" Cdh7
1460046_at 0.95197 "pleckstrin homology domain containing, family G (with RhoGef domain) member 1" Plekhg1
1460047_at 1.33918 Formin binding protein 1 Fnbp1
1460048_at 1.07012 "Zinc finger, DHHC domain containing 13" Zdhhc13
1460049_s_at 0.51616 RIKEN cDNA 1500015O10 gene 1500015O10Rik
1460050_x_at 1.01857 RIKEN cDNA 5630401D24 gene 5630401D24Rik
1460051_at 0.89967 cDNA sequence BC038613 BC038613
1460052_at 0.94222 Acyl-CoA synthetase medium-chain family member 1 Acsm1
1460053_at 0.99007 SET and MYND domain containing 4 Smyd4
1460054_at 1.07557 trafficking protein particle complex 6A Trappc6a
1460055_at 1.04812 Ecotropic viral integration site 2a Evi2a
1460056_at 0.99645 RIKEN cDNA 1700109F18 gene 1700109F18Rik
1460057_at 0.95845 growth differentiation factor 3 Gdf3
1460058_at 0.80803 "10 days neonate cortex cDNA, RIKEN full-length enriched library, clone: A830028J12 product:unclassifiable, full insert sequence" ---
1460059_at 1.01161 uridine phosphorylase 2 Upp2
1460060_at 0.98681 --- ---
1460061_at 1.15382 "Adult male olfactory brain cDNA, RIKEN full-length enriched library, clone:6430530M09 product:unclassifiable, full insert sequence" ---
1460062_at 0.94432 "pleckstrin homology domain containing, family H (with MyTH4 domain) member 2" Plekhh2
1460063_at 1.47774 "DNA segment, Chr 5, ERATO Doi 798, expressed" D5Ertd798e
1460064_at 1.03111 cDNA sequence BC028789 BC028789
1460065_at 0.97828 expressed sequence AA763515 AA763515
1460066_at 1.06077 expressed sequence C80889 C80889
1460067_at 0.84733 chemokine (C-C motif) receptor 2 Ccr2
1460068_at 1.16409 RIKEN cDNA 1110034C04 gene 1110034C04Rik
1460069_at 1.29269 structural maintenance of chromosomes 6 Smc6
1460070_at 0.84698 Carbohydrate sulfotransferase 2 Chst2
1460071_at 0.97607 SAP30-like Sap30l
1460072_at 0.82480 Transcribed locus ---
1460073_at 1.04268 "Protein kinase, cAMP dependent regulatory, type I beta" Prkar1b
1460074_x_at 0.80334 ependymin related protein 2 (zebrafish) Epdr2
1460075_x_at 0.95815 Small nuclear ribonucleoprotein N Snrpn
1460076_x_at 0.91186 --- ---
1460077_at 1.58734 tetratricopeptide repeat domain 3 Ttc3
1460078_at 1.00544 RIKEN cDNA G630024C07 gene G630024C07Rik
1460079_at 0.90144 Tetratricopeptide repeat domain 5 Ttc5
1460080_at 1.04227 expressed sequence AI645535 AI645535

1460081_at	0.93451	Synaptotagmin VII	Syt7	
1460082_at	0.91662	"inhibitor of growth family, member 3"	Ing3	
1460083_at	0.99670	A disintegrin and metallopeptidase domain 10	Adam10	
1460084_at	1.14192	---	---	
1460085_at	0.90991	Zinc finger protein 281	Zfp281	
1460086_at	1.11971	---	---	
1460087_at	0.96185	cDNA sequence BC017643	BC017643	
1460088_at	0.90280	RIKEN cDNA 6430531B16 gene	6430531B16Rik	
1460089_at	0.93931	Period homolog 3 (Drosophila)	Per3	
1460090_at	0.98475	Ariadne ubiquitin-conjugating enzyme E2 binding protein homolog 1 (Drosophila)	Arih1	
1460091_at	1.14903	"protein phosphatase 1, regulatory (inhibitor) subunit 12B"	Ppp1r12b	
1460092_at	0.96643	Limbic system-associated membrane protein	Lsamp	
1460093_at	0.89800	---	---	
1460094_at	1.14282	---	---	
1460095_at	1.07092	RIKEN cDNA 1110030E23 gene	1110030E23Rik	
1460096_at	1.73288	Lysosomal membrane glycoprotein 1	Lamp1	
1460097_at	1.00412	"Glutamate receptor, ionotropic, AMPA4 (alpha 4)"	Gria4	
1460098_at	0.99872	---	---	
1460099_at	2.01795	"ATPas, class II, type 9B"	Atp9b	
1460100_at	1.05464	Ladybird homeobox 1 homolog (Drosophila) corepressor 1	Lbxcor1	
1460101_at	0.90255	Neurexin III	Nrxn3	
1460102_at	1.35937	CLIP associating protein 1	Clasp1	
1460103_at	0.97433	---	---	
1460104_at	1.08268	vacuolar protein sorting 4b (yeast)	Vps4b	
1460105_at	0.87104	"UDP-GalNAc:betaGlcNAc beta 1,3-galactosaminyltransferase, polypeptide 2"	B3galnt2	
1460106_at	0.90045	hydrocephalus inducing	Hydin	
1460107_at	0.99588	RIKEN cDNA 1700129I04 gene	1700129I04Rik	
1460108_at	0.92417	TRNA isopentenyltransferase 1	Trit1	
1460109_at	1.28617	Transcription factor 25 (basic helix-loop-helix)	Tcf25	
1460110_at	1.13900	rotatin	Rtnn	
1460111_at	0.94104	myelin transcription factor 1-like	Myt1l	
1460112_at	1.11088	Transcribed locus	---	
1460113_at	1.41676	RIKEN cDNA B930093H17 gene	B930093H17Rik	
1460114_at	1.01790	RIKEN cDNA 1700123M08 gene	1700123M08Rik	
1460115_at	0.93786	"Solute carrier family 26, member 11"	Slc26a11	
1460116_s_at	1.05425	"sprouty protein with EVH-1 domain 1, related sequence"	Spred1	
1460117_at	0.99862	---	---	
1460118_at	1.08915	Transcribed locus	---	
1460119_at	1.02534	"16 days embryo lung cDNA, RIKEN full-length enriched library, clone:8430415C15 product:unclassifiable, full insert sequence"	---	
1460120_at	1.10597	actin-binding LIM protein 1	Ablim1	

1460121_at	1.97386	RIKEN cDNA 9630010G10 gene	9630010G10Rik
1460122_at	0.82866	transmembrane protein 41B	Tmem41b
1460123_at	1.01820	G protein-coupled receptor 1	Gpr1
1460124_at	0.94912	RIKEN cDNA 1190020J12 gene	1190020J12Rik
1460125_at	1.00718	RIKEN cDNA 2610301F02 gene	2610301F02Rik
1460126_at	2.48988	Ubiquitin specific peptidase 34	Usp34
1460127_at	1.01304	"hepatocyte nuclear factor 4, gamma"	Hnf4g
1460128_at	0.94205	Transmembrane emp24 domain trafficking protein 2	Tmed2
1460129_at	0.97230	"solute carrier family 6 (neurotransmitter transporter, noradrenalin), member 2"	Slc6a2
1460130_at	1.01799	hypothetical LOC433632	LOC433632
1460131_at	0.98006	Ring finger protein 170	Rnf170
1460132_at	1.10443	"Ubiquitin-conjugating enzyme E2B, RAD6 homology (S. cerevisiae)"	Ube2b
1460133_at	2.95039	---	---
1460134_at	0.91410	RIKEN cDNA 9330175E14 gene	9330175E14Rik
1460135_at	1.03207	RIKEN cDNA A930005H10 gene	A930005H10Rik
1460136_at	1.00401	expressed sequence AW047481	AW047481
1460137_at	0.89189	Fibroblast growth factor 12	Fgf12
1460138_at	1.40957	RIKEN cDNA 5730446C15 gene	5730446C15Rik
1460139_at	1.01513	expressed sequence C79256	C79256
1460140_at	1.14234	CD2-associated protein	Cd2ap
1460141_at	0.98601	Transcribed locus	---
1460142_at	0.87958	"DNA segment, Chr 13, ERATO Doi 324, expressed"	D13Ertd324e
1460143_at	0.99041	---	---
1460144_at	1.40276	CDNA sequence BC052040	BC052040
1460145_at	0.94924	---	---
1460146_at	0.99581	---	---
1460147_at	0.97329	RIKEN cDNA A730009E18 gene	A730009E18Rik
1460148_at	1.03319	Transcribed locus	---
1460149_at	0.99342	ATPase type 13A5	Atp13a5
1460150_at	0.95791	Transcribed locus	---
1460151_at	1.31355	---	---
1460152_at	1.02792	---	---
1460153_at	1.00038	Abl-interactor 2	Abi2
1460154_at	0.98147	expressed sequence AI194348	AI194348
1460155_at	0.94751	RIKEN cDNA 2410005O16 gene	2410005O16Rik
1460156_at	1.10265	Acyloxyacyl hydrolase	Aoah
1460157_at	1.00295	---	---
1460158_at	1.02447	---	---
1460159_at	1.11891	"myb-like, SWIRM and MPN domains 1"	Mysm1
1460160_at	0.97449	"Adult male testis cDNA, RIKEN full-length enriched library, clone:4930520J05 product:unclassifiable, full insert sequence"	---

1460161_at 0.90196 "DNA segment, Chr 5, ERATO Doi 422, expressed" D5Ert422e
 1460162_at 1.19057 similar to Serine/threonine-protein kinase Nek1 (NimA-related protein kinase 1)
 LOC640559
 1460163_at 1.03967 C1q and tumor necrosis factor related protein 7 C1qtnf7
 1460164_at 0.96332 spindlin Spin
 1460165_at 1.00338 "protein phosphatase 1, catalytic subunit, alpha isoform" Ppp1ca
 1460166_at 1.03208 Ras-like without CAAX 2 Rit2
 1460167_at 0.82878 "aldehyde dehydrogenase family 7, member A1" Aldh7a1
 1460168_at 0.88528 stem-loop binding protein Slbp
 1460169_a_at 1.01965 PCTAIRE-motif protein kinase 1 Pctk1
 1460170_at 0.88143 exostoses (multiple) 2 Ext2
 1460171_at 1.07379 "COP9 (constitutive photomorphogenic) homolog, subunit 5 (Arabidopsis thaliana)" Cops5
 1460172_at 0.94332 RIKEN cDNA 5730403B10 gene 5730403B10Rik
 1460173_at 1.05332 LIM and SH3 protein 1 Laspl
 1460174_at 0.86700 dexamethasone-induced transcript Dexi
 1460175_at 1.03642 ribosomal protein S23 Rps23
 1460176_at 0.87777 v-crk sarcoma virus CT10 oncogene homolog (avian) Crk
 1460177_at 0.97697 CNDP dipeptidase 2 (metallopeptidase M20 family) Cndp2
 1460178_at 0.86608 RIKEN cDNA 1300002A08 gene 1300002A08Rik
 1460179_at 0.90743 "DnaJ (Hsp40) homolog, subfamily A, member 1" Dnaja1
 1460180_at 0.88821 hexosaminidase B Hexb
 1460181_at 0.94681 stathmin-like 3 Stmn3
 1460182_at 1.08452 sorting nexin 4 Snx4
 1460183_at 1.02232 ADP-ribosylarginine hydrolase Adprh
 1460184_at 0.91847 "L-3-hydroxyacyl-Coenzyme A dehydrogenase, short chain" Hadhsc
 1460185_at 0.98415 keratin 85 Krt85
 1460186_at 1.03347 transmembrane 9 superfamily member 3 Tm9sf3
 1460187_at 0.97529 secreted frizzled-related sequence protein 1 Sfrp1
 1460188_at 1.07669 "protein tyrosine phosphatase, non-receptor type 6" Ptpn6
 1460189_at 0.92741 WD repeat domain 23 Wdr23
 1460190_at 0.97663 "adaptor protein complex AP-1, mu 2 subunit" Ap1m2
 1460191_at 0.92884 YKT6 homolog (S. Cerevisiae) Ykt6
 1460192_at 0.74837 oxysterol binding protein-like 1A Osbp1a
 1460193_at 1.04234 suppression of tumorigenicity 13 St13
 1460194_at 0.78533 phytanoyl-CoA hydroxylase Phyh
 1460195_at 0.97974 mitochondrial ribosomal protein S11 Mrps11
 1460196_at 0.88295 carbonyl reductase 1 Cbr1
 1460197_a_at 0.88554 STEAP family member 4 Steap4
 1460198_a_at 1.09468 "proteasome (prosome, macropain) subunit, beta type 3 /// similar to proteasome beta 3 subunit /// similar to proteasome beta 3 subunit /// similar to proteasome beta 3 subunit" Psm3 /// LOC240289 /// LOC636660 /// LOC672201
 1460199_a_at 1.06811 "platelet-activating factor acetylhydrolase, isoform 1b, beta1 subunit" Pafah1b1

1460200_s_at 1.00773 leucine zipper transcription factor-like 1 Lztf1
 1460201_a_at 1.06186 ribosomal protein L24 /// similar to ribosomal protein L24 /// similar to ribosomal protein L24 Rpl24 /// LOC668829 /// LOC675789
 1460202_at 1.02227 myozenin 1 Myoz1
 1460203_at 0.75109 "inositol 1,4,5-triphosphate receptor 1" Itpr1
 1460204_at 0.91811 "cytoplasmic tyrosine kinase, Dscr28C related (Drosophila)" Tec
 1460205_at 0.96850 dephospho-CoA kinase domain containing Dcakd
 1460206_at 0.89096 GRP1 (general receptor for phosphoinositides 1)-associated scaffold protein Grasp
 1460207_s_at 0.98140 E2F transcription factor 5 E2f5
 1460208_at 1.03037 fibrillin 1 Fbn1
 1460209_at 1.01149 ubiquitin specific peptidase 39 Usp39
 1460210_at 0.92331 polycystic kidney disease 1 homolog Pkd1
 1460211_a_at 0.83267 KDEL (Lys-Asp-Glu-Leu) endoplasmic reticulum protein retention receptor 1 Kdelr1
 1460212_at 0.91077 "guanine nucleotide binding protein, alpha transducing 1" Gnat1
 1460213_at 0.98963 "golgi autoantigen, golgin subfamily a, 4" Golga4
 1460214_at 1.04820 Purkinje cell protein 4 Pcp4
 1460215_at 1.08286 RNA polymerase 1-4 Rpo1-4
 1460216_at 0.91719 "acyl-Coenzyme A dehydrogenase, short chain" Acads
 1460217_at 1.00181 proliferin related protein Plfr
 1460218_at 0.83089 CD52 antigen Cd52
 1460219_at 0.98002 myelin-associated glycoprotein Mag
 1460220_a_at 1.27256 colony stimulating factor 1 (macrophage) Csf1
 1460221_at 1.05487 prostaglandin E synthase 3 (cytosolic) Ptges3
 1460222_at 0.91120 SH3-domain binding protein 1 Sh3bp1
 1460223_a_at 0.90008 erythrocyte protein band 4.9 Epb4.9
 1460224_at 1.00565 sorting nexin 2 Snx2
 1460225_at 0.88261 mitochondrial ribosomal protein 63 Mrp63
 1460226_at 0.94666 tumor rejection antigen P1A Trap1a
 1460227_at 1.19118 tissue inhibitor of metalloproteinase 1 Timp1
 1460228_at 0.96914 upstream transcription factor 2 Usf2
 1460229_at 1.04263 stromal antigen 3 Stag3
 1460230_at 1.07343 synapsin II Syn2
 1460231_at 1.06310 interferon regulatory factor 5 Irf5
 1460232_s_at 0.97653 "hydroxy-delta-5-steroid dehydrogenase, 3 beta- and steroid delta-isomerase 2 /// hydroxy-delta-5-steroid dehydrogenase, 3 beta- and steroid delta-isomerase 3 /// hydroxy-delta-5-steroid dehydrogenase, 3 beta- and steroid delta-isomerase 6" Hsd3b2 /// Hsd3b3 /// Hsd3b6
 1460233_at 0.83527 guanylate cyclase activator 2b (retina) Guca2b
 1460234_at 1.14335 WD repeat domain 81 Wdr81
 1460235_at 0.84948 "scavenger receptor class B, member 2" Scarb2
 1460236_at 1.07028 kallikrein 10 Klk10
 1460237_at 0.94652 tripartite motif protein 8 Trim8

1460238_at	1.36634	mesothelin	Msln	
1460239_at	0.94612	tetraspanin 13	Tspan13	
1460240_a_at	0.91997	heterogeneous nuclear ribonucleoprotein C	Hnrpc	
1460241_a_at	0.87513	"ST3 beta-galactoside alpha-2,3-sialyltransferase 5"	St3gal5	
1460242_at	1.05377	CD55 antigen	Cd55	
1460243_at	1.04795	"serine palmitoyltransferase, long chain base subunit 2"	Sptlc2	
1460244_at	0.99044	"ureidopropionase, beta"	Upb1	
1460245_at	0.92964	"killer cell lectin-like receptor, subfamily D, member 1"	Klrd1	
1460246_at	0.86694	methyl CpG binding protein 2	Mecp2	
1460247_a_at	1.12254	S-phase kinase-associated protein 2 (p45)	Skp2	
1460248_at	1.02211	carboxypeptidase X 2 (M14 family)	Cpxm2	
1460249_at	0.92204	ligand of numb-protein X 2	Lnx2	
1460250_at	1.03086	sclerostin domain containing 1	Sostdc1	
1460251_at	1.87646	Fas (TNF receptor superfamily member)	Fas	
1460252_s_at	0.98523	zinc finger protein 105	Zfp105	
1460253_at	1.13771	CKLF-like MARVEL transmembrane domain containing 7	Cmtm7	
1460254_at	0.95299	RIKEN cDNA 1810049H13 gene	1810049H13Rik	
1460255_at	0.93286	"tumor necrosis factor (ligand) superfamily, member 13b"	Tnfsf13b	
1460256_at	1.06203	carbonic anhydrase 3	Car3	
1460257_a_at	1.01440	"5, 10-methenyltetrahydrofolate synthetase"	Mthfs	
1460258_at	1.03556	leukocyte cell derived chemotaxin 1	Lect1	
1460259_s_at	2.30582	chloride channel calcium activated 1 /// chloride channel calcium activated 2	Clca1 /// Clca2	
1460260_s_at	1.05647	karyopherin (importin) alpha 1	Kpna1	
1460261_at	1.03831	"discs, large homolog 4 (Drosophila)"	Dlgh4	
1460262_a_at	1.05568	RIKEN cDNA 1700037H04 gene	1700037H04Rik	
1460263_at	0.84900	Programmed cell death 6 interacting protein	Pdcd6ip	
1460264_at	0.94277	large tumor suppressor	Lats1	
1460265_at	0.73835	Proliferation-associated 2G4	Pa2g4	
1460266_at	1.02696	RIKEN cDNA 1700094J05 gene	1700094J05Rik	
1460267_at	0.95279	doublesex and mab-3 related transcription factor 3	Dmrt3	
1460268_at	0.93559	---	---	
1460269_at	1.00071	Phenylethanolamine-N-methyltransferase	Pnmt	
1460271_at	1.11365	triggering receptor expressed on myeloid cells 3	Trem3	
1460272_at	0.98981	ubiquitin-like 4B	Ubl4b	
1460273_a_at	1.19057	baculoviral IAP repeat-containing 1b	Birc1b	
1460274_at	0.97818	RIKEN cDNA 4921530G04 gene	4921530G04Rik	
1460275_at	0.99735	G-protein coupled receptor 3	Gpr3	
1460276_a_at	0.98593	G protein-coupled receptor 175	Gpr175	
1460277_at	0.92866	diencephalon/mesencephalon homeobox 1	Dmbx1	
1460278_a_at	0.98844	"DNA segment, Chr 15, Wayne State University 75, expressed"	D15Wsu75e	
1460279_a_at	0.87938	general transcription factor II I /// similar to General transcription factor II-I (GTFII-I) (TFII-I) (Bruton tyrosine kinase-associated protein 135) (BTK-associated protein 135) (BAP-		

135) Gtf2i /// LOC669007

1460280_at 0.92250 GRB2-related adaptor protein 2 /// similar to GRB2-related adaptor protein 2 (GADS protein) (Growth factor receptor binding protein) (GRBLG) (GRB-2-like protein) (GRB2L) (Hematopoietic cell-associated adaptor protein GrpL) (GRB-2-related monocytic adapter protein) (Monocytic adapter... Grap2 /// LOC674050

1460281_at 1.04567 ankyrin repeat and SOCS box-containing protein 15 Asb15

1460282_at 1.09693 triggering receptor expressed on myeloid cells 1 Trem1

1460283_at 1.03256 Mediterranean fever Mefv

1460284_at 0.79413 leucine zipper-EF-hand containing transmembrane protein 1 Letm1

1460285_at 1.12086 integrin alpha 9 Itga9

1460286_at 0.97945 septin 6 6-Sep

1460287_at 0.91558 tissue inhibitor of metalloproteinase 2 Timp2

1460288_a_at 0.97413 "protein phosphatase 4, catalytic subunit" Ppp4c

1460289_at 1.02892 neuregulin 3 Nrg3

1460290_at 1.25882 lipin 2 Lpin2

1460291_at 1.32932 cyclin-dependent kinase 6 Cdk6

1460292_a_at 0.71418 "SWI/SNF related, matrix associated, actin dependent regulator of chromatin, subfamily a, member 1" Smarca1

1460293_at 0.98553 frequenin homolog (Drosophila) Freq

1460294_at 1.36026 "ATPase, aminophospholipid transporter-like, class I, type 8A, member 2" Atp8a2

1460295_s_at 1.00687 interleukin 6 signal transducer Il6st

1460296_a_at 0.96027 fibroblast growth factor 22 Fgf22

1460297_at 1.06001 "nephrosis 2 homolog, podocin (human)" Nphs2

1460298_at 0.96445 RIKEN cDNA 4930483J18 gene 4930483J18Rik

1460299_at 0.90906 homeobox gene HB9 Hlxb9

1460300_a_at 0.86812 leukocyte tyrosine kinase Ltk

1460301_at 0.92333 "vomeronasal 1 receptor, D2" V1rd2

1460302_at 0.77422 thrombospondin 1 /// similar to thrombospondin 1 Thbs1 /// LOC640441

1460303_at 0.91612 "nuclear receptor subfamily 3, group C, member 1" Nr3c1

1460304_a_at 0.92001 "upstream binding transcription factor, RNA polymerase I" Ubtf

1460305_at 0.91897 integrin alpha 3 Itga3

1460306_at 0.95632 myocilin Myoc

1460307_at 0.90915 thymoma viral proto-oncogene 3 Akt3

1460308_a_at 1.06299 immature colon carcinoma transcript 1 Ict1

1460309_at 1.08424 T-cell acute lymphocytic leukemia 2 Tal2

1460310_a_at 1.03689 growth hormone Gh

1460311_at 0.97978 simple repeat sequence-containing transcript Srst

1460312_at 0.95216 olfactory receptor 157 Olfr157

1460313_at 0.96859 olfactory receptor 2 Olfr2

1460314_s_at 1.20938 "histone 2, H3c1 /// histone 2, H3c2 /// histone 1, H3g /// histone 1, H3f /// histone 1, H3c /// histone 1, H3d /// histone 1, H3b /// histone 1, H3e /// histone 1, H3h /// histone 1, H3i /// histone 2, H3b /// histone 1, H3a" Hist2h3c1 /// Hist2h3c2 /// Hist1h3g /// Hist1h3f ///

Hist1h3c /// Hist1h3d /// Hist1h3b /// Hist1h3e /// Hist1h3h /// Hist1h3i /// Hist2h3b /// Hist1h3a
 1460315_s_at 1.03235 TANK-binding kinase 1 Tbk1
 1460316_at 0.92993 acyl-CoA synthetase long-chain family member 1 Acs11
 1460317_s_at 1.09684 "guanine nucleotide binding protein, alpha 13" Gna13
 1460318_at 1.19402 cysteine and glycine-rich protein 3 Csrp3
 1460319_at 0.95129 fucosyltransferase 8 Fut8
 1460320_at 0.99398 "beclin 1 (coiled-coil, myosin-like BCL2-interacting protein)" Becn1
 1460321_at 0.90725 contactin 4 Cntn4
 1460322_at 1.01444 carbohydrate (chondroitin 6/keratan) sulfotransferase 3 Chst3
 1460323_at 1.28005 threonyl-tRNA synthetase Tars
 1460324_at 0.61142 DNA methyltransferase 3A Dnmt3a
 1460325_at 1.04832 pumilio 1 (Drosophila) Pum1
 1460326_at 0.96978 "phosphatidylinositol 3-kinase, catalytic, alpha polypeptide" Pik3ca
 1460327_at 0.90115 G-protein coupled receptor 88 Gpr88
 1460328_at 0.99678 bromodomain containing 3 Brd3
 1460329_at 1.14043 "similar to Beta-1,4-galactosyltransferase 6 (Beta-1,4-GalTase 6) (Beta4Gal-T6) (b4Gal-T6) (UDP-galactose:beta-N-acetylglucosamine beta-1,4-galactosyltransferase 6) (UDP-Gal:beta-GlcNAc beta-1,4-galactosyltransferase 6)" LOC675709
 1460330_at 1.04584 annexin A3 Anxa3
 1460331_at 0.94138 transmembrane 9 superfamily member 2 Tm9sf2
 1460332_at 1.07785 phospholamban Pln
 1460333_at 1.04538 DEAD (Asp-Glu-Ala-Asp) box polypeptide 59 Ddx59
 1460334_at 1.11177 drebrin-like Dbnl
 1460335_at 1.02483 "LysM, putative peptidoglycan-binding, domain containing 3" Lysmd3
 1460336_at 0.73310 "peroxisome proliferative activated receptor, gamma, coactivator 1 alpha" Ppargc1a
 1460337_at 0.73244 SH3-domain kinase binding protein 1 Sh3kbp1
 1460338_a_at 1.12708 cytokine receptor-like factor 3 Crlf3
 1460339_at 1.09118 "proteasome (prosome, macropain) subunit, alpha type 4" Psma4
 1460340_at 0.95967 piwi-like homolog 1 (Drosophila) Piwil1
 1460341_at 0.94006 "pleckstrin homology domain containing, family B (evectins) member 2" Plekhh2
 1460342_s_at 0.89619 expressed sequence AA536749 AA536749
 1460343_at 0.89296 neuralized-like homolog (Drosophila) Neurl
 1460344_at 0.88171 RIKEN cDNA 2310033F14 gene 2310033F14Rik
 1460345_at 1.06616 RIKEN cDNA 2610208M17 gene 2610208M17Rik
 1460346_at 1.01665 arylsulfatase A Arsa
 1460347_at 1.15832 keratin 14 Krt14
 1460348_at 0.92847 MAD2 mitotic arrest deficient-like 2 (yeast) Mad2l2
 1460349_at 0.89190 proline rich 14 Prr14
 1460350_at 0.99937 oxysterol binding protein Osbp
 1460351_at 0.99833 S100 calcium binding protein A11 (calizzarin) S100a11
 1460352_s_at 0.92968 "phosphatidylinositol 3 kinase, regulatory subunit, polypeptide 4, p150"

Pik3r4

1460353_at 0.85358 transmembrane protein 48 Tmem48
 1460354_a_at 0.98967 mitochondrial ribosomal protein L13 Mrp13
 1460355_at 0.97052 BUD13 homolog (yeast) Bud13
 1460356_at 0.79958 endothelial cell-specific adhesion molecule Esam1
 1460357_at 0.97787 YTH domain family 2 Ythdf2
 1460358_s_at 0.93619 nudix (nucleoside diphosphate linked moiety X)-type motif 22 Nudt22
 1460359_at 1.01145 "armadillo repeat containing, X-linked 3" Armcx3
 1460360_at 0.90853 asparaginase like 1 Asrg11
 1460361_at 0.94326 RIKEN cDNA 5033414D02 gene 5033414D02Rik
 1460362_at 0.99564 RIKEN cDNA 2410001C21 gene 2410001C21Rik
 1460363_at 0.94525 trinucleotide repeat containing 6C Tnrc6c
 1460364_at 1.09496 general transcription factor II I repeat domain-containing 1 Gtf2ird1
 1460365_a_at 0.99465 dynamin 1 Dnm1
 1460366_at 0.99672 echinoderm microtubule associated protein like 3 Eml3
 1460367_at 0.86053 high mobility group box transcription factor 1 Hbp1
 1460368_at 1.00004 "membrane protein, palmitoylated 4 (MAGUK p55 subfamily member 4)"

Mpp4

1460369_at 0.95949 cDNA sequence BC003267 BC003267
 1460370_at 1.06715 "DNA topoisomerase 1, mitochondrial" Top1mt
 1460371_at 1.05336 heat shock protein 12B Hspa12b
 1460372_at 0.92951 dual oxidase maturation factor 1 Duoxa1
 1460373_a_at 0.96986 SET domain containing 4 Setd4
 1460374_at 0.94584 a disintegrin and metallopeptidase domain 2 Adam2
 1460375_at 1.11086 RIKEN cDNA 0610038D11 gene 0610038D11Rik
 1460376_a_at 0.98965 "COX15 homolog, cytochrome c oxidase assembly protein (yeast)" Cox15
 1460377_a_at 1.00366 transmembrane protein 8 (five membrane-spanning domains) Tmem8
 1460378_a_at 1.01608 testis derived transcript Tes
 1460379_at 1.07103 homeo box B4 Hoxb4
 1460380_at 0.86081 desmoglein 2 Dsg2
 1460381_at 1.00983 cDNA sequence BC023179 BC023179
 1460382_at 0.95119 cDNA sequence BC020535 BC020535
 1460383_at 0.95726 "guanine nucleotide binding protein, alpha o" Gnao1
 1460384_a_at 0.95872 AT rich interactive domain 4B (Rbp1 like) Arid4b
 1460385_a_at 0.96598 zinc finger protein 179 Zfp179
 1460386_a_at 1.08138 "solute carrier family 1 (neuronal/epithelial high affinity glutamate transporter, system Xag), member 1" Slc1a1
 1460387_a_at 0.87287 sialic acid acetyltransferase Siae
 1460388_at 0.96045 RIKEN cDNA 4632408A20 gene 4632408A20Rik
 1460389_at 1.05828 cyclin-dependent kinase 8 Cdk8
 1460390_at 0.92465 "sortilin-related receptor, LDLR class A repeats-containing" Sorl1
 1460391_at 1.06567 RIKEN cDNA 2810409H07 gene 2810409H07Rik
 1460392_a_at 1.08752 enhancer of yellow 2 homolog (Drosophila) Eny2

1460393_a_at 0.98366 dual specificity phosphatase 7 /// similar to dual specificity phosphatase 7
 Dusp7 /// LOC674944
 1460394_a_at 0.97771 inositol polyphosphate phosphatase-like 1 Inpp1
 1460395_at 1.07992 NudC domain containing 3 Nudcd3
 1460396_at 1.09154 DEAD (Asp-Glu-Ala-Asp) box polypeptide 54 Ddx54
 1460397_at 0.99395 armadillo repeat containing 6 Armc6
 1460398_at 0.95021 PHD finger protein 8 Phf8
 1460399_at 0.99594 coiled-coil domain containing 117 Ccdc117
 1460400_at 1.01640 spastin Spast
 1460401_at 0.92443 "ER degradation enhancer, mannosidase alpha-like 3" Edem3
 1460402_at 0.97523 "bromodomain and PHD finger containing, 1" Brpf1
 1460403_at 0.96302 PC4 and SFRS1 interacting protein 1 Psip1
 1460404_at 1.08522 stretch responsive protein 278 Sr278
 1460405_at 1.01650 Rho guanine nucleotide exchange factor (GEF) 10-like Arhgef10
 1460406_at 1.00913 expressed sequence AI427122 AI427122
 1460407_at 1.21206 Spi-B transcription factor (Spi-1/PU.1 related) Spib
 1460408_at 0.99186 "gamma-aminobutyric acid (GABA-A) receptor, subunit gamma 1" Gabrg1
 1460409_at 0.79520 "carnitine palmitoyltransferase 1a, liver" Cpt1a
 1460410_at 1.01181 RIKEN cDNA 2310079F23 gene 2310079F23Rik
 1460411_s_at 1.03061 expressed sequence AW548124 AW548124
 1460412_at 0.91978 RIKEN cDNA 1600015H20 gene 1600015H20Rik
 1460413_s_at 0.83118 "zinc finger, DHHC domain containing 1" Zdhhc1
 1460414_at 0.91891 expressed sequence AW822216 AW822216
 1460415_a_at 0.99901 CD40 antigen Cd40
 1460416_s_at 0.96854 component of Sp100-rs /// similar to component of Sp100-rs /// similar to
 component of Sp100-rs /// similar to component of Sp100-rs /// similar to component of Sp100-
 rs /// similar to component of Sp100-rs /// similar to component of Sp100-rs /// similar to component of Sp100-
 rs /// similar to component of Sp100-rs /// similar to component of Sp100-rs Csprs /// Sp100-rs ///
 LOC665338 /// LOC665378 /// LOC673105 /// LOC673206 /// LOC673671 /// LOC673849 ///
 LOC673925 /// LOC673997
 1460417_at 1.16825 cDNA sequence AB041803 AB041803
 1460418_x_at 0.95749 "histocompatibility 2, T region locus 18" H2-T18
 1460419_a_at 0.95339 "protein kinase C, beta 1" Prkcb1
 1460420_a_at 1.02629 epidermal growth factor receptor Egfr
 1460421_at 0.98945 zinc finger protein 133 Zfp133
 1460422_at 0.96120 "DNA segment, KIST 5" D0Kist5
 1460423_x_at 0.99834 immunoglobulin heavy chain (V7183 family) /// immunoglobulin kappa chain
 variable 1 (V1) /// immunoglobulin kappa chain variable 1-117 /// Ig kappa chain Igh-V7183 /// Igk-
 V1 /// Igkv1-117 /// Cr1
 1460424_at 1.14540 transmembrane protein 160 Tmem160
 1460425_at 0.88627 RIKEN cDNA 1700001C19 gene 1700001C19Rik
 1460426_at 0.79822 phosphodiesterase 4D interacting protein (myomegalin) Pde4dip
 1460427_a_at 1.07547 a disintegrin and metallopeptidase domain 28 Adam28

1460428_at 0.96581 ankyrin repeat domain 13a Ankrd13a
1460429_at 1.00906 cell division cycle 5-like (*S. pombe*) Cdc5l
1460430_at 1.00130 "RAP2C, member of RAS oncogene family" Rap2c
1460431_at 0.91440 "glucosaminyl (N-acetyl) transferase 1, core 2" Gcnt1
1460432_a_at 1.02345 "eukaryotic translation initiation factor 3, subunit 6" Eif3s6
1460433_at 0.99564 ectonucleoside triphosphate diphosphohydrolase 6 Entpd6
1460434_at 1.09495 FUN14 domain containing 2 Fundc2
1460435_at 0.98296 RIKEN cDNA 1500002O20 gene 1500002O20Rik
1460436_at 0.95209 N-deacetylase/N-sulfotransferase (heparan glucosaminyl) 1 Ndst1
1460437_at 0.99334 "pleckstrin homology, Sec7 and coiled/coil domains 4" Pscd4
1460438_at 0.92826 "LysM, putative peptidoglycan-binding, domain containing 1" Lysmd1
1460439_at 1.09480 cDNA sequence BC033915 BC033915
1460440_at 0.79115 latrophilin 3 Lphn3
1460441_at 0.97056 similar to Zinc finger X-linked protein ZXDB LOC668166
1460442_at 1.08476 ribosomal protein S19 Rps19
1460443_at 1.05100 breast cancer metastasis-suppressor 1-like Brms1l
1460444_at 0.79032 "arrestin, beta 1" Arrb1
1460445_at 0.98938 "splicing factor, arginine/serine-rich 2, interacting protein" Sfrs2ip
1460446_at 1.03449 tryptophan rich basic protein Wrb
1460447_at 1.10477 pseudouridylate synthase 7 homolog (*S. cerevisiae*)-like Pus7l
1460448_s_at 1.01414 tetratricopeptide repeat domain 14 Ttc14
1460449_at 1.05490 ankyrin repeat and sterile alpha motif domain containing 1B /// similar to AIDA-1b Anks1b /// LOC544718
1460450_at 0.95494 RIKEN cDNA E230022H04 gene E230022H04Rik
1460451_at 1.01684 transmembrane protein 52 Tmem52
1460452_at 0.95746 motile sperm domain containing 3 Mospd3
1460453_at 1.07797 T-cell activation Rho GTPase-activating protein Tagap
1460454_at 1.04531 RIKEN cDNA 2010001H14 gene 2010001H14Rik
1460455_at 0.91961 RIKEN cDNA 1110059H15 gene 1110059H15Rik
1460456_at 1.02313 RIKEN cDNA 2010316F05 gene 2010316F05Rik
1460457_at 0.98888 RIKEN cDNA 2810405F17 gene 2810405F17Rik
1460458_at 0.91988 cysteine-rich secretory protein LCCL domain containing 2 Crisp1d2
1460459_at 0.90767 progesterin and adipoQ receptor family member V Paqr5
1460460_a_at 0.91403 golgi reassembly stacking protein 2 Gorasp2
1460461_at 0.97979 RIKEN cDNA 4921521K07 gene 4921521K07Rik
1460462_at 0.93399 "mediator of RNA polymerase II transcription, subunit 18 homolog (yeast)" Med18
1460463_at 1.02898 RIKEN cDNA 4632413I24 gene 4632413I24Rik
1460464_at 0.89325 RIKEN cDNA 2700089E24 gene 2700089E24Rik
1460465_at 0.68287 RIKEN cDNA A930038C07 gene A930038C07Rik
1460466_at 1.20570 RIKEN cDNA 1700047I17 gene /// similar to signal recognition particle 54 1700047I17Rik /// LOC665155
1460467_at 0.97974 IQ motif containing G Iqcg

1460468_s_at 0.91042 RIKEN cDNA 2810451A06 gene 2810451A06Rik
1460469_at 1.50378 "tumor necrosis factor receptor superfamily, member 9" Tnfrsf9
1460470_at 1.15755 acyl-Coenzyme A oxidase-like Acox1
1460471_at 0.97245 RIKEN cDNA 2410146L05 gene 2410146L05Rik
1460472_at 0.92696 cyclin-dependent kinase 3 Cdk3
1460473_at 0.86266 RIKEN cDNA 6230400D17 gene 6230400D17Rik
1460474_at 1.46753 RIKEN cDNA 2610028L16 gene 2610028L16Rik
1460475_at 0.91977 RIKEN cDNA 3110056J03 gene 3110056J03Rik
1460476_s_at 0.94371 RIKEN cDNA 1200015N20 gene 1200015N20Rik
1460477_at 1.01286 RIKEN cDNA 1700010C24 gene 1700010C24Rik
1460478_at 0.97540 RIKEN cDNA 2200002J24 gene 2200002J24Rik
1460479_at 0.84992 RIKEN cDNA A330094K24 gene A330094K24Rik
1460480_at 1.00547 RIKEN cDNA 1600014E20 gene 1600014E20Rik
1460481_at 1.10449 --- ---
1460482_at 1.08106 RIKEN cDNA 3110047P20 gene 3110047P20Rik
1460483_at 1.14228 RIKEN cDNA 2610034E01 gene 2610034E01Rik
1460484_at 0.99070 tubulin tyrosine ligase Ttl
1460485_at 0.90827 RIKEN cDNA 8430434A19 gene 8430434A19Rik
1460486_at 0.91462 RAB GTPase activating protein 1 Rabgap1
1460487_at 1.00457 RIKEN cDNA 1110014L15 gene 1110014L15Rik
1460488_at 0.84001 zinc finger and BTB domain containing 4 Zbtb4
1460489_at 0.88696 RIKEN cDNA 1700097M23 gene 1700097M23Rik
1460490_at 1.00469 mitochondrial ribosomal protein L15 Mrpl15
1460491_at 1.09240 inversin Invs
1460492_at 0.99540 RIKEN cDNA 8030498B09 gene 8030498B09Rik
1460493_at 0.96206 RIKEN cDNA 4921506J03 gene 4921506J03Rik
1460494_at 0.94721 "tubulin tyrosine ligase-like family, member 10" Ttl10
1460495_s_at 1.24135 HtrA serine peptidase 2 Htra2
1460496_at 1.02477 similar to RIKEN cDNA 5430413K10 LOC433492
1460497_a_at 0.92250 RIKEN cDNA 4933413B09 gene 4933413B09Rik
1460498_a_at 1.22825 "DnaJ (Hsp40) homolog, subfamily C, member 5" Dnajc5
1460499_at 1.17478 RIKEN cDNA 9230110I02 gene 9230110I02Rik
1460500_at 1.19585 RIKEN cDNA 5033421C21 gene 5033421C21Rik
1460501_at 1.00860 RIKEN cDNA 4930521G14 gene 4930521G14Rik
1460502_at 0.95916 RIKEN cDNA 2010007E15 gene 2010007E15Rik
1460503_at 0.92581 zona pellucida binding protein 2 Zpbp2
1460504_at 0.94637 --- ---
1460505_at 0.92959 RIKEN cDNA 8030442B05 gene 8030442B05Rik
1460506_s_at 1.03213 "NADH dehydrogenase (ubiquinone) 1, subcomplex unknown, 2 /// similar to NADH dehydrogenase (ubiquinone) 1, subcomplex unknown, 2" Ndufc2 /// LOC675851
1460507_at 0.92406 RIKEN cDNA 4930552P12 gene 4930552P12Rik
1460508_at 0.97851 mitochondrial ribosomal protein L1 Mrpl1
1460509_at 0.97748 "UDP-Gal:betaGlcNAc beta 1,3-galactosyltransferase, polypeptide 1"

B3galt1

1460510_a_at 0.94978 coenzyme Q10 homolog B (*S. cerevisiae*) Coq10b

1460511_at 1.47650 plakophilin 2 Pkp2

1460512_a_at 0.77555 RIKEN cDNA 4930558O21 gene 4930558O21Rik

1460513_a_at 1.09032 endothelin receptor type A Ednra

1460514_s_at 0.97266 achaete-scute complex homolog-like 2 (*Drosophila*) Ascl2

1460515_at 0.85219 RIKEN cDNA 8430419K02 gene 8430419K02Rik

1460516_at 1.23989 RIKEN cDNA 3110006I21 gene 3110006I21Rik

1460517_at 1.00440 RIKEN cDNA 9130409J20 gene 9130409J20Rik

1460518_at 0.88767 RIKEN cDNA E130119H09 gene E130119H09Rik

1460519_a_at 1.03509 methyltransferase like 5 Mettl5

1460520_at 0.94311 molybdenum cofactor synthesis 1 Mocs1

1460521_a_at 1.26020 RIKEN cDNA 5830411E10 gene 5830411E10Rik

1460522_at 0.91459 RIKEN cDNA 6030458E02 gene 6030458E02Rik

1460523_at 0.91949 --- ---

1460524_at 0.97392 RIKEN cDNA 6430711C07 gene 6430711C07Rik

1460525_at 1.01196 RIKEN cDNA 2810047J09 gene 2810047J09Rik

1460526_at 1.27799 RIKEN cDNA 4833420L08 gene 4833420L08Rik

1460527_at 0.89633 RIKEN cDNA 4933407C09 gene 4933407C09Rik

1460528_at 0.98637 RIKEN cDNA 4930413G21 gene /// hypothetical protein LOC668574 ///
 hypothetical protein LOC670667 /// hypothetical protein LOC676722 4930413G21Rik ///
 LOC668574 /// LOC670667 /// LOC676722

1460529_at 1.02925 RIKEN cDNA 4930455M05 gene 4930455M05Rik

1460530_at 1.05059 RIKEN cDNA 4933416A02 gene 4933416A02Rik

1460531_at 0.96290 RIKEN cDNA 2900069M18 gene 2900069M18Rik

1460532_at 0.91771 RIKEN cDNA 2900011F02 gene 2900011F02Rik

1460533_at 0.91865 RIKEN cDNA 4933440K10 gene 4933440K10Rik

1460534_at 0.97011 RIKEN cDNA 5530401A10 gene 5530401A10Rik

1460535_at 1.22960 "golgi autoantigen, golgin subfamily b, macrogolgin 1" Golgb1

1460536_at 1.30107 RIKEN cDNA 2310079F09 gene 2310079F09Rik

1460537_at 0.99891 RIKEN cDNA 9430085L16 gene 9430085L16Rik

1460538_at 1.05971 cadherin 10 Cdh10

1460539_at 1.03597 RIKEN cDNA 4933404K13 gene 4933404K13Rik

1460540_at 0.94446 neurexin I Nrxn1

1460541_at 1.07779 "solute carrier family 7 (cationic amino acid transporter, y+ system), member 6" Slc7a6

1460542_s_at 1.01398 transformed mouse 3T3 cell double minute 4 Mdm4

1460543_x_at 0.96248 ribosomal protein L37a Rpl37a

1460544_at 0.95570 "MAK10 homolog, amino-acid N-acetyltransferase subunit, (*S. cerevisiae*)"
 Mak10

1460545_at 0.95518 thyroid hormone receptor associated protein 3 Thrap3

1460546_at 1.01153 "leucine-rich repeat LGI family, member 3" Lgi3

1460547_a_at 0.98830 heterogeneous nuclear ribonucleoprotein K Hnrpk

1460548_a_at 0.96939 Era (G-protein)-like 1 (E. coli) Eral1
1460549_a_at 0.99383 "CDC23 (cell division cycle 23, yeast, homolog)" Cdc23
1460550_at 0.93750 myotubularin related protein 11 Mtmr11
1460551_at 1.02815 "RAN, member RAS oncogene family" Ran
1460552_at 0.95595 activating signal cointegrator 1 complex subunit 3-like 1 Ascc311
1460553_at 0.87311 RIKEN cDNA 1700025K23 gene 1700025K23Rik
1460554_s_at 0.88877 golgi apparatus protein 1 Glg1
1460555_at 0.89495 RIKEN cDNA 6330500D04 gene 6330500D04Rik
1460556_at 0.89889 --- ---
1460557_at 1.06664 "suppressor of var1, 3-like 1 (S. cerevisiae)" Supv311
1460558_at 0.85183 coiled-coil domain containing 32 Ccdc32
1460559_at 1.04535 ankyrin repeat domain 25 Ankrd25
1460560_at 0.98043 BAH domain and coiled-coil containing 1 Bahcc1
1460561_x_at 0.75482 "selenoprotein W, muscle 1" Sepw1
1460562_at 1.00565 elongation factor Tu GTP binding domain containing 1 Eftud1
1460563_at 1.06781 "DNA segment, Chr 10, ERATO Doi 438, expressed" D10Ert438e
1460564_at 1.05315 suppressor of hairy wing homolog 2 (Drosophila) Suhw2
1460565_at 0.94319 "solute carrier family 41, member 1" Slc41a1
1460566_at 0.88166 microtubule-associated protein 1 A Mtap1a
1460567_at 1.27225 regulatory factor X domain containing 2 homolog (human) Rfxdc2
1460568_at 1.16902 tripartite motif protein 46 Trim46
1460569_x_at 0.96220 claudin 3 Cldn3
1460570_at 1.04653 piggyBac transposable element derived 5 Pgbd5
1460571_at 0.80597 "Dicer1, Dcr-1 homolog (Drosophila)" Dicer1
1460572_a_at 1.00353 zinc finger protein 511 Zfp511
1460573_at 0.97136 expressed sequence AI848100 AI848100
1460574_at 0.62271 FAT tumor suppressor homolog 4 (Drosophila) Fat4
1460575_at 1.00892 eukaryotic translation initiation factor 2a Eif2a
1460576_at 0.97158 exocyst complex component 6 Exoc6
1460577_at 1.03466 Expressed sequence AA591059 AA591059
1460578_at 1.03916 "FYVE, RhoGEF and PH domain containing 5" Fgd5
1460579_at 1.05809 aspartyl aminopeptidase Dnpep
1460580_at 0.80015 pecanex homolog (Drosophila) Pcnx
1460581_a_at 0.99101 ribosomal protein L13 /// similar to 60S ribosomal protein L13 /// similar to 60S ribosomal protein L13 /// similar to 60S ribosomal protein L13 /// similar to 60S ribosomal protein L13 /// similar to 60S ribosomal protein L13 Rpl13 /// LOC280047 /// LOC547376 /// LOC668182 /// LOC672281 /// LOC674921 /// LOC675192
1460582_x_at 0.92281 Nonagouti a
1460583_at 0.92601 golgi transport 1 homolog B (S. cerevisiae) Golt1b
1460584_at 1.02955 histidine acid phosphatase domain containing 2A Hisppd2a
1460585_x_at 0.93512 phosphatidylserine decarboxylase Pisd
1460586_at 1.07532 multiple EGF-like-domains 8 Megf8
1460587_at 1.04041 RIKEN cDNA B230215L15 gene B230215L15Rik

1460588_at 1.07068 "10 days lactation, adult female mammary gland cDNA, RIKEN full-length enriched library, clone:D730013G10 product:unclassifiable, full insert sequence" ---
1460589_at 0.84544 zinc finger protein 597 Zfp597
1460590_s_at 0.97146 "tyrosine 3-monooxygenase/tryptophan 5-monooxygenase activation protein, theta polypeptide" Ywhaq
1460591_at 0.85469 estrogen receptor 1 (alpha) Esr1
1460592_at 0.91632 erythrocyte protein band 4.1-like 1 Epb4.111
1460593_at 0.94342 sushi domain containing 4 Susd4
1460594_a_at 0.97290 GDP-mannose pyrophosphorylase A Gmppa
1460595_at 1.03360 "16 days embryo head cDNA, RIKEN full-length enriched library, clone: C130021L12 product:unclassifiable, full insert sequence" ---
1460596_at 0.92534 "angiotensin II, type I receptor-associated protein" Agrap
1460597_at 1.19321 Additional sex combs like 2 (Drosophila) Asxl2
1460598_at 1.03351 interleukin 28 receptor alpha Il28ra
1460599_at 0.86199 "DNA segment, Chr 19, Wayne State University 12, expressed" D19Wsu12e
1460600_at 0.94063 similar to ubiquitin-conjugating enzyme E2Q (putative) 2 LOC245350
1460601_at 1.08968 myosin VIIA and Rab interacting protein Myrip
1460602_at 0.98318 deleted in liver cancer 1 Dlc1
1460603_at 1.10221 sterile alpha motif domain containing 9-like Samd9l
1460604_at 0.97236 cytochrome b reductase 1 Cybrd1
1460605_at 1.02936 Crx opposite strand transcript 1 Crxos1
1460606_at 1.05609 hydroxysteroid (17-beta) dehydrogenase 13 Hsd17b13
1460607_at 0.94444 "immunoglobulin superfamily, member 11" Igsf11
1460608_at 0.88799 "calcium channel, voltage-dependent, N type, alpha 1B subunit" Cacna1b
1460609_at 0.94046 Integrator complex subunit 8 Ints8
1460610_at 0.78945 RIKEN cDNA 9430057O19 gene 9430057O19Rik
1460611_at 0.98078 expressed sequence C87499 C87499
1460612_at 1.10759 RAN binding protein 3 Ranbp3
1460613_x_at 0.98693 growth hormone Gh
1460614_at 0.83269 "mesoderm induction early response 1, family member 3" Mier3
1460615_at 1.02671 5'-nucleotidase domain containing 1 Nt5dc1
1460616_at 0.95959 "solute carrier organic anion transporter family, member 4C1" Slco4c1
1460617_s_at 1.15179 "RAB6B, member RAS oncogene family" Rab6b
1460618_x_at 0.97985 RIKEN cDNA 1700001J11 gene 1700001J11Rik
1460619_at 1.09426 RIKEN cDNA 4931419K03 gene 4931419K03Rik
1460620_at 0.80428 zinc finger protein 592 Zfp592
1460621_x_at 0.89304 "tyrosine 3-monooxygenase/tryptophan 5-monooxygenase activation protein, theta polypeptide" Ywhaq
1460622_x_at 1.21497 WD repeat domain 18 Wdr18
1460623_at 0.86727 src family associated phosphoprotein 2 Scap2
1460624_at 1.02257 RIKEN cDNA 6330564D18 gene 6330564D18Rik
1460625_at 1.01850 "gene model 1568, (NCBI)" Gm1568
1460626_at 1.05666 septin 11 11-Sep

1460627_at 1.03234 RIKEN cDNA D130067I03 gene D130067I03Rik
1460628_at 1.03882 Mitogen-activated protein kinase 8 interacting protein 3 Mapk8ip3
1460629_at 0.97246 tripartite motif protein 16 Trim16
1460630_at 1.05130 "calcium/calmodulin-dependent protein kinase II, delta" Camk2d
1460631_at 1.03087 O-linked N-acetylglucosamine (GlcNAc) transferase (UDP-N-acetylglucosamine:polypeptide-N-acetylglucosaminyl transferase) Ogt
1460632_at 0.77896 --- ---
1460633_at 1.02588 PRP19/PSO4 pre-mRNA processing factor 19 homolog (S. cerevisiae) Prpf19
1460634_at 0.85779 ral guanine nucleotide dissociation stimulator Ralgds
1460635_at 0.88882 Fas-activated serine/threonine kinase Fastk
1460636_at 0.96192 mitogen activated protein kinase kinase 2 Map2k2
1460637_s_at 0.93696 prefoldin 5 Pfdn5
1460638_at 1.07150 tyrosyl-tRNA synthetase Yars
1460639_a_at 0.98614 ATX1 (antioxidant protein 1) homolog 1 (yeast) Atox1
1460640_at 0.99345 ATX1 (antioxidant protein 1) homolog 1 (yeast) Atox1
1460641_a_at 0.85584 RIKEN cDNA 3100004P22 gene /// RIKEN cDNA 2310022K01 gene 3100004P22Rik /// 2310022K01Rik
1460642_at 0.94384 Tnf receptor associated factor 4 Traf4
1460643_at 0.96117 elongation factor RNA polymerase II Ell
1460644_at 0.98910 branched chain ketoacid dehydrogenase kinase Bckdk
1460645_at 0.94039 "cysteine and histidine-rich domain (CHORD)-containing, zinc-binding protein 1" Chordc1
1460646_at 0.98421 "casein kinase 2, alpha prime polypeptide" Csnk2a2
1460647_a_at 0.83678 "nuclear receptor subfamily 2, group F, member 6" Nr2f6
1460648_at 0.89847 "nuclear receptor subfamily 2, group F, member 6" Nr2f6
1460649_at 1.12708 interleukin-1 receptor-associated kinase 1 Irak1
1460650_at 0.89474 "ATPase, H+ transporting, lysosomal V0 subunit A1" Atp6v0a1
1460651_at 1.04011 linker for activation of T cells Lat
1460652_at 0.94866 "estrogen related receptor, alpha" Esrra
1460653_at 1.01072 ataxin 2 Atxn2
1460654_at 1.02850 "solute carrier family 30 (zinc transporter), member 3" Slc30a3
1460655_a_at 1.07179 upstream binding protein 1 Ubp1
1460656_a_at 1.08095 RIKEN cDNA 5630401J11 gene 5630401J11Rik
1460657_at 1.12760 wingless related MMTV integration site 10a Wnt10a
1460658_at 0.97641 "Adaptor protein complex AP-1, gamma 1 subunit" Ap1g1
1460659_at 0.91126 "CDNA, clone:Y0G0114H22, strand:minus, reference:ENSEMBL:Mouse-Transcript-ENST:ENSMUST00000058552, based on BLAT search" ---
1460660_x_at 0.94643 RER1 retention in endoplasmic reticulum 1 homolog (S. cerevisiae) Rer1
1460661_at 1.05698 "endothelial differentiation, sphingolipid G-protein-coupled receptor, 3" Edg3
1460662_at 0.93092 period homolog 3 (Drosophila) Per3
1460663_at 0.90689 cholecystokinin B receptor Cckbr

1460664_at 0.91066 "zinc finger protein, subfamily 1A, 4 (Eos)" Zfpn1a4
1460665_a_at 0.97932 "CCR4-NOT transcription complex, subunit 7" Cnot7
1460666_a_at 1.03207 early B-cell factor 3 Ebf3
1460667_at 0.89971 cDNA sequence U90926 U90926
1460668_at 0.83932 galanin Gal
1460669_at 1.03154 interleukin enhancer binding factor 3 Ilf3
1460670_at 0.97828 RIO kinase 3 (yeast) Riok3
1460671_at 0.87787 glutathione peroxidase 1 Gpx1
1460672_at 1.03722 RIKEN cDNA 2410002F23 gene 2410002F23Rik
1460673_at 0.96345 foylypolyglutamyl synthetase Fpgs
1460674_at 0.83126 progesterin and adipoQ receptor family member VII Paqr7
1460675_at 0.84846 "immunoglobulin superfamily, member 8" Igsf8
1460676_at 0.97317 Josephin domain containing 1 Josd1
1460677_at 0.94553 --- ---
1460678_at 0.81706 kelch domain containing 2 Klhdc2
1460679_at 1.03553 exosome component 4 Exosc4
1460680_a_at 1.04476 ribosomal protein L23 Rpl23
1460681_at 0.92310 CEA-related cell adhesion molecule 1 Ceacam1
1460682_s_at 0.99327 CEA-related cell adhesion molecule 1 /// CEA-related cell adhesion molecule
2 Ceacam1 /// Ceacam2
1460683_at 0.96662 TEA domain family member 4 Tead4
1460684_at 1.44284 transmembrane 7 superfamily member 2 Tm7sf2
1460685_at 1.13553 translocase of inner mitochondrial membrane 17b Timm17b
1460686_at 0.89193 cyclin N-terminal domain containing 1 Cntd1
1460687_at 1.03113 SET domain containing (lysine methyltransferase) 8 Setd8
1460688_s_at 0.95679 expressed sequence AA407659 AA407659
1460689_at 1.00999 "DNA segment, Chr 15, Wayne State University 75, expressed" D15Wsu75e
1460690_at 0.92271 cDNA sequence BC003940 BC003940
1460691_at 1.10101 zinc finger protein 598 Zfp598
1460692_at 1.01269 euchromatic histone lysine N-methyltransferase 2 Ehmt2
1460693_a_at 1.02342 "procollagen, type IX, alpha 3" Col9a3
1460694_s_at 0.84188 supervillin Svil
1460695_a_at 0.88930 RIKEN cDNA 2010111I01 gene 2010111I01Rik
1460696_at 0.98187 cDNA sequence BC026585 BC026585
1460697_s_at 1.06143 RIKEN cDNA 2610209M04 gene 2610209M04Rik
1460698_a_at 1.06961 Sec11-like 3 (S. cerevisiae) Sec113
1460699_at 1.05824 ribosomal protein S27 Rps27
1460700_at 1.05282 signal transducer and activator of transcription 3 Stat3
1460701_a_at 0.98051 mitochondrial ribosomal protein L52 Mrpl52
1460702_at 1.05274 TP53 regulated inhibitor of apoptosis 1 Triap1
1460703_at 0.90448 activating signal cointegrator 1 complex subunit 1 Ascc1
1460704_at 1.03894 radical fringe gene homolog (Drosophila) Rfng
1460705_at 1.12530 "ribosomal protein S6 kinase, polypeptide 1" Rps6kb1

1460706_s_at 0.91702 RER1 retention in endoplasmic reticulum 1 homolog (S. cerevisiae) Rer1
1460707_at 1.07920 protein tyrosine phosphatase 4a2 Ptp4a2
1460708_s_at 0.94856 cell division cycle 42 homolog (S. cerevisiae) Cdc42
1460709_a_at 0.87084 HLA-B associated transcript 5 Bat5
1460710_at 0.94074 adenosine A2a receptor Adora2a
1460711_at 1.20865 RIKEN cDNA 4930461P20 gene 4930461P20Rik
1460712_s_at 0.99701 "adaptor protein complex AP-1, gamma 1 subunit" Ap1g1
1460713_at 1.10430 cDNA sequence BC048355 BC048355
1460714_at 1.05416 Transcription factor CP2-like 1 Tcfcp211
1460715_x_at 1.00043 Transcription factor CP2-like 1 Tcfcp211
1460716_a_at 0.93578 core binding factor beta Cbfb
1460717_at 0.77780 "testis-specific protein, Y-encoded-like 1" Tspyl1
1460718_s_at 0.93960 mitochondrial carrier homolog 1 (C. elegans) Mtch1
1460719_a_at 1.19189 "purinergic receptor P2X, ligand-gated ion channel, 1" P2rx1
1460720_at 1.01451 "transient receptor potential cation channel, subfamily C, member 4 associated protein" Trpc4ap
1460722_at 0.98314 sterol O-acyltransferase 2 Soat2
1460723_at 1.18770 melanocortin 5 receptor Mc5r
1460724_at 1.04478 "adaptor protein complex AP-2, alpha 1 subunit" Ap2a1
1460725_at 1.03133 "xeroderma pigmentosum, complementation group A" Xpa
1460726_at 1.04044 "adenylosuccinate synthetase, non muscle" Adss
1460727_at 1.11886 "excision repair cross-complementing rodent repair deficiency, complementation group 2" Ercc2
1460728_s_at 0.88043 "inhibitor of growth family, member 4" Ing4
1460729_at 0.92807 Rho-associated coiled-coil containing protein kinase 1 Rock1
1460730_at 1.02535 "eukaryotic translation initiation factor 2B, subunit 1 (alpha)" Eif2b1
1460731_at 1.04217 "solute carrier family 35, member C2" Slc35c2
1460732_a_at 1.21989 periplakin Ppl
1460733_at 0.91260 expressed sequence AA407659 AA407659
1460734_at 0.94802 "procollagen, type IX, alpha 3" Col9a3
1460735_at 0.91450 supervillin Svl
1460736_at 0.96891 "similar to CG5514-PB, isoform B /// region containing expressed sequence AF013969; RIKEN cDNA A230054D04 gene" LOC665775 /// LOC675766
1460737_at 1.05426 immunoglobulin (CD79A) binding protein 1 Igbp1
1460738_at 0.82223 LIM domain containing 2 Limd2
1460739_at 0.86070 "DNA segment, Chr 11, Brigham & Women's Genetics 0280e expressed" D11Bwg0280e
1460740_at 0.93999 "clathrin, light polypeptide (Lcb)" Cltb
1460741_x_at 1.01052 "DNA segment, Chr 17, Wayne State University 92, expressed" D17Wsu92e
1460742_at 1.04346 "asparagine-linked glycosylation 3 homolog (yeast, alpha-1,3-mannosyltransferase)" Alg3
1460743_at 0.95243 tigger transposable element derived 5 Tigd5
1460744_at 0.87689 RIKEN cDNA 2810002I04 gene 2810002I04Rik

1460745_at	0.95347	RIKEN cDNA A630098A13 gene	A630098A13Rik
1460746_at	0.91760	fidgetin-like 1 Fignl1	