

Supplemental Methods

Generation of lentiviruses expressing *Cacna2d1* and *Gria3*

We constructed lentiviruses expressing the full-length rat *Cacna2d1* (Sequence ID: NM_012919) and rat *Gria3* (Sequence ID: NM_032990.2) coding sequences. Briefly, we performed PCR using CloneAMP HiFi PCR premix (#639298, TaKara) to obtain the full-length cDNA of *Cacna2d1* from the mRNA of rat spinal cord. The *Cacna2d1* cDNA was then cloned into a pEntry vector using the pEntry Directional Topo Cloning kit (#K2400-20, Invitrogen). The *Cacna2d1* in the pEntry vector was recombined into pLenti7.3 based on the instruction in the pLenti7.3/V5 Gateway Vector Kit (#V534-06, Invitrogen). The pLenti7.3/V5 vector containing *Cacna2d1* was then packaged into the lentivirus through co-transfecting with the packaging plasmids pCMV-VSV-G (#8454, Addgene) and psPAX2 (#12260, Addgene) into HEK293FT cells. The virus-containing culture medium was collected after 72 h of culture. The virus was concentrated 100 times using a Lenti-X concentrator reagent (#631231, Takara Bio), and the viral titer was determined using a titration reagent (#VPK-112, Cell Biolabs). The concentrated virus with a titer of 4×10^{11} viral particles/mL was stored at -80°C until use. The negative control virus was made using pLenti7.3/V5 vector (without *Cacna2d1*) with the same packaging method. Same methods are used for the construction of lentiviruses expressing *Gria3*. The titer of the concentrated *Gria3* lentivirus is 2.5×10^{11} viral particles/mL. The primers used for PCR are *Cacna2d1*-forward: 5'-CACCATggctgctgctgcctgctgg-3'; *Cacna2d1*-reverse: 5'-gtcaccatagatagtgtctgc-3'; *Gria3*-forward: 5'-CACCATggggcgaagcgtgctccgggcggtc-3'; *Gria3*-reverse: 5'-tcctagatcttaacactttctgtcc-3'. The CACC sequence in the forward primers provides overhang base pairs, allowing for insertion into the pEntry vector.

Cell culture and transfection

HEK293 cells were cultured in Dulbecco's Modified Eagle's Medium (DMEM; Gibco/Life Technologies) supplemented with 10% fetal bovine serum (FBS; Sigma-Aldrich) at 37°C in a 5% CO_2 incubator. For electrophysiological recordings, 1.2×10^4 cells were plated on poly-D-lysine-coated coverslips. DNA encoding GluA2(R) or GluA3 was transfected along with either empty vectors (pcDNA) or $\alpha 2\delta$ subunits. Empty vectors were used to keep the total DNA amount balance across different transfection groups. The transfection ratio of GluA2, GluA3, and $\alpha 2\delta$ -1 was maintained at 1:1:1 in most experiments using the PolyJet DNA In Vitro Transfection Reagent (SignaGen Laboratories). GFP coexpression was used to identify transfected cells. Electrophysiological recordings were conducted 48 h post-transfection. The cDNAs used for $\alpha 2\delta$, YFP-tagged WT $\alpha 2\delta$ -1, and $\alpha 2\delta$ chimeras have been previously described (13, 22). The GluA2(R) and GluA3 constructs were generously provided by Dr. Michael Hollmann (Ruhr University) and Dr. Li Niu (University at Albany), respectively.

Supplemental Figures

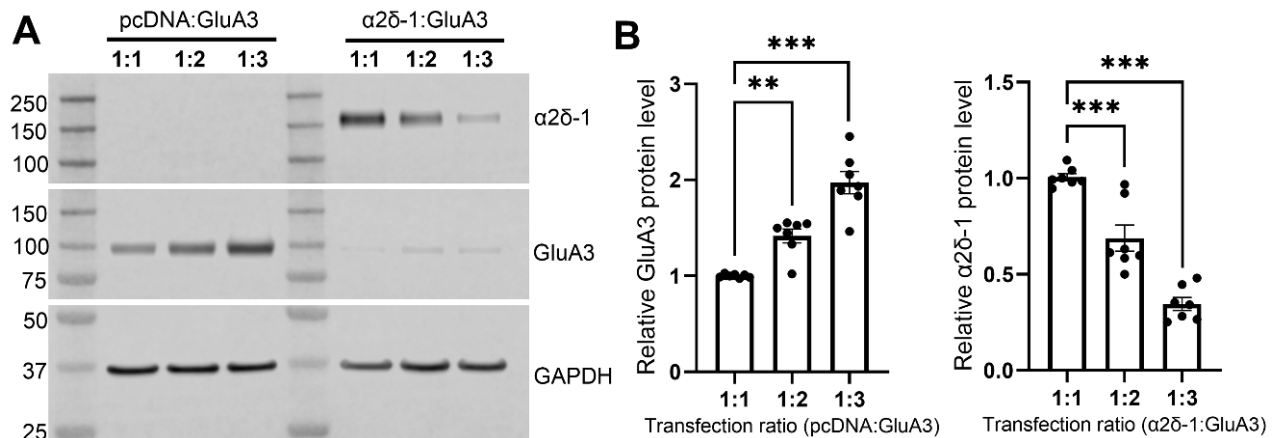


Figure S1. GluA3 overexpression reduces α2δ-1 protein levels. (A and B) Representative immunoblot images (A) and quantification (B) show the effects of coexpression with varying amounts of GluA3 on α2δ-1 protein levels in HEK293 cells. Empty vectors (pcDNA) were used to equalize the total DNA amount across different transfection groups (n = 7 independent experiments per group). **p < 0.01, ***p < 0.001; one-way ANOVA followed by Dunnett's *post hoc* test. Data are expressed as means ± SEM.

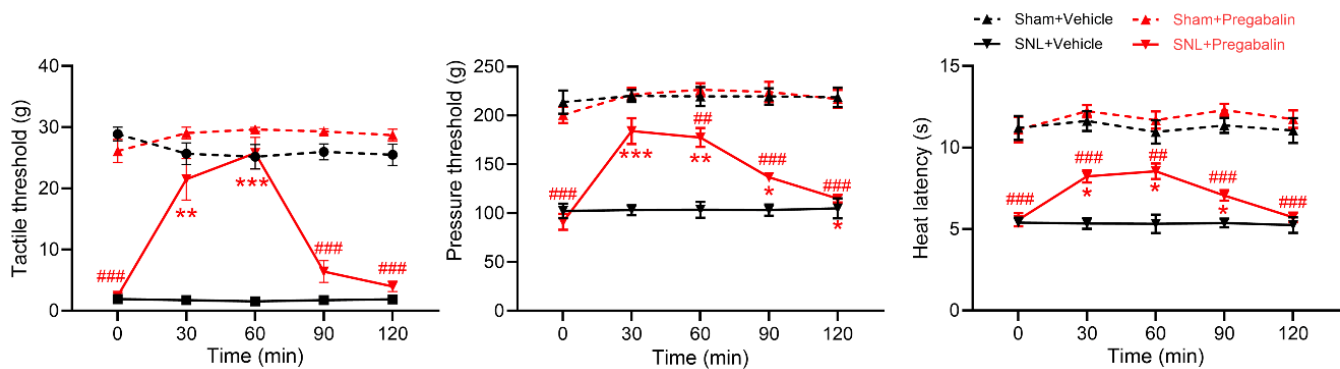


Figure S2. Pregabalin reverses nociceptive hypersensitivity induced by nerve injury. Time-course effects of intrathecal injection of pregabalin (10 μg) or vehicle on tactile, pressure, and heat withdrawal thresholds in sham and SNL rats 3 weeks after surgery (n = 7 rats per group). *p < 0.05, **p < 0.01, ***p < 0.001 vs. baseline (time 0); #p < 0.01, ###p < 0.001 vs. SNL+Vehicle group at the same time point (two-way ANOVA followed by Tukey's *post hoc* test). Data are expressed as means ± SEM.

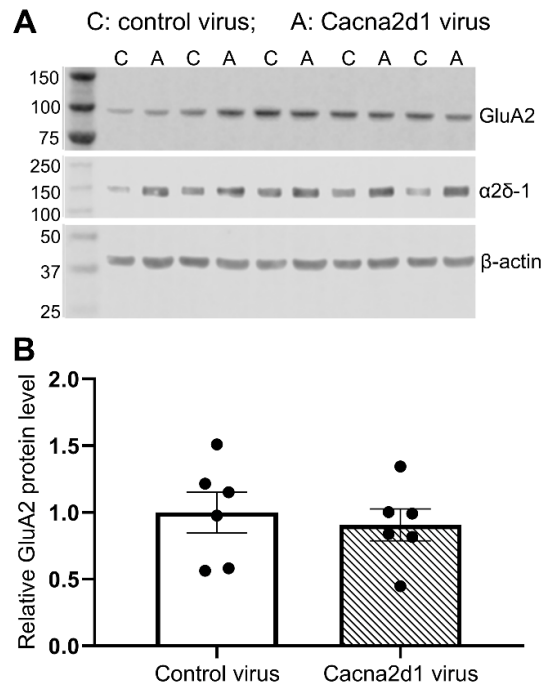


Figure S3. Effect of $\alpha 2\delta$ -1 overexpression on GluA2 protein levels in the spinal cord. (A and B) Representative immunoblot images (A) and quantification (B) show GluA2 protein levels in the dorsal spinal cord from naïve rats injected intrathecally with control lentiviruses or *Cacna2d1*-expressing lentiviruses (n = 6 rats per group). Two-tailed Student's t test. Data are expressed as means $\pm$ SEM.

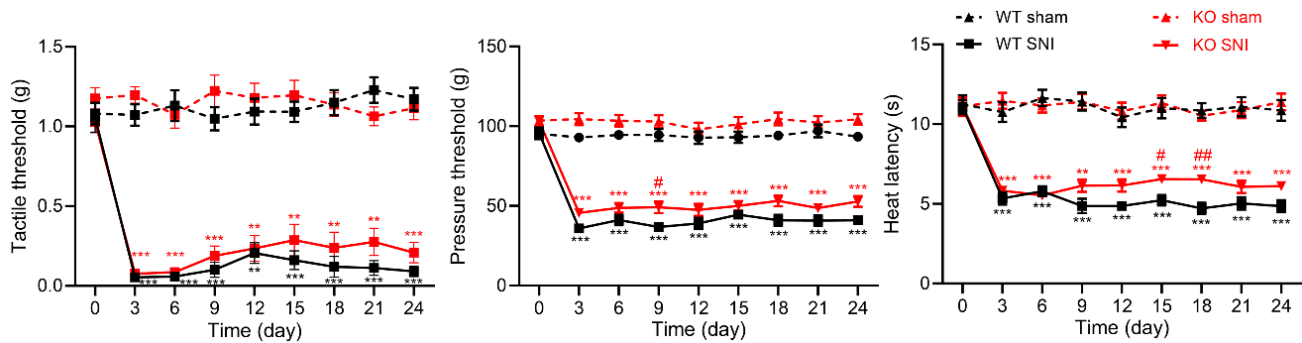


Figure S4. Development of nociceptive hypersensitivity after nerve injury in wild-type and *Cacna2d1* knockout mice. Time course of changes in tactile, pressure, and heat withdrawal thresholds in wild-type (WT) and *Cacna2d1* knockout (KO) mice after sham or SNI surgery (n = 10 rats per group). **p < 0.01, ***p < 0.001 vs. baseline (day 0). #p < 0.05, ##p < 0.01 vs. WT-SNI at the same time point (two-way ANOVA followed by Tukey's *post hoc* test). Data are expressed as means $\pm$ SEM.

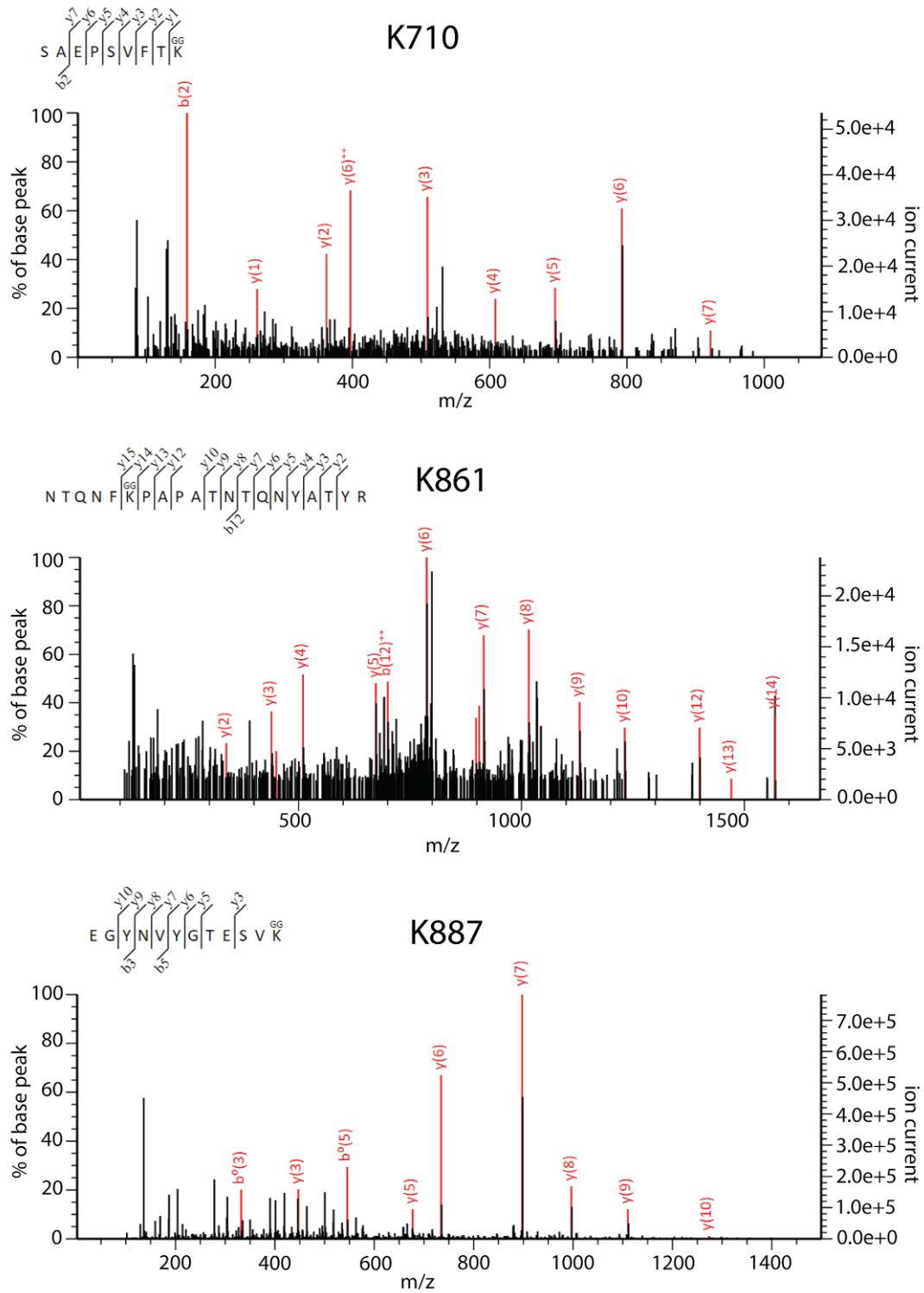


Figure S5. Annotated spectra of fragmented ions (tandem mass spectrometry) of putative GluA3 peptides including ubiquitination sites. The ubiquitination sites and peptide sequences are indicated on the top. Only b and y ions whose matches were qualified and used for scoring by Mascot version 2.7 are labeled. Annotation: b, B-ion; y, y-ion; number, position of fragmentation; m/z, mass-to-charge ratio; ^o loss of a water; ⁺⁺ double charge; GG, diglycine-modified peptides.

CLUSTAL O(1.2.4) multiple sequence alignment

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sp|P19490|GRIA1_RAT      ----MPYIFAFFCTGFLGAVGANFPNNIQIGGLFPNQSQEHAAFRFALSQLT----- 50
sp|P19491|GRIA2_RAT      -MQKIMHISVLLSPVLWGLI-FGVSSNSIQIGGLFPRGADQEYSAFRVGMVQFST----- 53
sp|P19492|GRIA3_RAT      MGQSVLRVAVFFLVLLGLLGH-SHGFPNTISIGGLFMRNTVQEHSARFAVQLYNTNQNTT 59
                          :      ::      *      .      *.*****      **::**..:      .

sp|P19490|GRIA1_RAT      -EPPKLLPQIDIVNISDSFEMTYRFSQFSKGVYAIFGFYERRTVNMLTSFCGALHVCFI 109
sp|P19491|GRIA2_RAT      -SEFRLTPHIDNLEVANSFAVTNAFCSQFSRGVYAIFGFYDKKSVNTITSFCGTLHVSFI 112
sp|P19492|GRIA3_RAT      EKPFLHLYHVDHLDSSNSFSVTNAFCSQFSRGVYAIFGFYDQMSMNTLSFCGALHTSFV 119
                          .      :*      ::*      ::**      :*      *****:*****::*      :*****:*.*:

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sp|P19492|GRIA3_RAT      TPSFPTADVQFVIQMRPALKGAILSLLSYYKWEKFVYLYDTERGFSVLQAIMEAAVQNN 179
*****.*      **::**      *      :      ::*      :      :      :      :      :      :      :      :      :      :

sp|P19490|GRIA1_RAT      WQVTAVNILT---TEEGYRMLFQDLEKKKERLVVDCESERLNAILGQIVKLEKNGIG 225
sp|P19491|GRIA2_RAT      WQVTAINVGNINNDKKDETYRSLFQDLEKKERRVILDCERDKVNDIVDQVITIGKHVK 232
sp|P19492|GRIA3_RAT      WQVTARSVGNIKD---VQEFRIIEEMDRRQEKRYLIDCEVERINTILEQVVILGKHSRG 236
*****      .:      .      :      :      :      :      :      :      :      :      :      :      :      :      :      :

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sp|P19491|GRIA2_RAT      YHYIIANLGFDTGDLKKIQFGGANVSGFQIVDYDSLVSKEIERWSTLEEKEYPGAHTAT 292
sp|P19492|GRIA3_RAT      YHYMLANLGFDTILLERVHGGANITGFQIVNNENPMVQFIQRWRLDEREFPEAKNAP 296
***::*****      *      *      .      .      :      :      :      :      :      :      :      :      :      :      :

sp|P19490|GRIA1_RAT      PKYTSALTYDGVKVMAEAFQSLRRQRIDISRRGNAGDCLANPAVPWGGQIDIQRALQQVR 345
sp|P19491|GRIA2_RAT      IKYTSALTYDAVQVMTEAFRNLRKQRIEISRRGNAGDCLANPAVPWGGQVEIERALKQVQ 352
sp|P19492|GRIA3_RAT      LKYTSALTHDAILVIAEAFRYLRRQRVDVSRGSGAGDCLANPAVPWSQGDIERALKMVQ 356
*****:*.:      :      :      :      :      :      :      :      :      :      :      :      :      :      :      :

sp|P19490|GRIA1_RAT      FEGLTGNVQFNEKGRRNTNYLTHVIEMKHDGIRKIGYWNEDDKFVPAATDAQAGDSSVQ 405
sp|P19491|GRIA2_RAT      VEGLSGNIKFDQNGKRINYINIMELKTNGPRKIGYWEVDKMMVTLTELPNGDTSGL 412
sp|P19492|GRIA3_RAT      VQGMTGNIQFDTYGRRNTNYIDVYEMKVSGRKAGYWNEYERFVP-FSDQQISNDSSSE 415
.:*::**::*      :      :      :      :      :      :      :      :      :      :      :      :      :      :      :

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sp|P19491|GRIA2_RAT      NKTVVTTILESPYVMMKKNHMELEGNERYEGYCVDLAAEIAKHGFKYKLTIVGDGKYG 472
sp|P19492|GRIA3_RAT      NRTIIVTTILESPYVMMKKNHMELEGNERYEGYCVDLAYEIAKHVRIKYKLSIVGDGKYG 475
*:      :      :      :      :      :      :      :      :      :      :      :      :      :      :      :

sp|P19490|GRIA1_RAT      ARDPDTKAWNGMVGLVYGRADVAAPLTITLVREEVIDFSKPFMSLGISIMIKKPQKSK 525
sp|P19491|GRIA2_RAT      ARDADTKIWNMGVGLVYGKADIAIAPLTITLVREEVIDFSKPFMSLGISIMIKKPQKSK 532
sp|P19492|GRIA3_RAT      ARDPETKIWNMGVGLVYGRADIAVAPLTITLVREEVIDFSKPFMSLGISIMIKKPQKSK 535
***      :      :      :      :      :      :      :      :      :      :      :      :      :      :      :

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sp|P19491|GRIA2_RAT      PGVFSFLDPLAYEIWMCIYFAYIGVSVVLFVSRFSPYEWHTTEFEDGRE--TQSSSESTN 590
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*****:*****:*****:*****:*****:*****:*****:*****:*****:*****:*****:*****:*****:*****:*****:*****:

sp|P19490|GRIA1_RAT      EFGIFNSLWFLSGLAFMQQGCDSIPRSLSGRIVGGVMWFFTLIISSYTANLAAFLTVERM 643
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sp|P19492|GRIA3_RAT      EFGIFNSLWFLSGLAFMQQGCDSIPRSLSGRIVGGVMWFFTLIISSYTANLAAFLTVERM 655
*****:*****:*****:*****:*****:*****:*****:*****:*****:*****:*****:*****:*****:*****:*****:

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sp|P19491|GRIA2_RAT      VSPIESAEDLSKQTEIAYGTLDGSGTKEFFRRSKIAVFDKMWYMRSAEPSVVFRTTAE 710
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sp|P19490|GRIA1_RAT      MIRVRKSKGYAYLLESTMNEYIEQRKPCDTMKVGGNLDKSGYGIATPKGSALRNPVLA 763

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: *****:*:*****:*****:*****:* * ****

sp|P19490|GRIA1_RAT  VLKLNEQGLLDKLKNKWYDKGECGSGGGDSKDKTSALSLSNVAGVFYILIGGLGLAMLV  823
sp|P19491|GRIA2_RAT  VLKLNEQGLLDKLKNKWYDKGECGSGGGDSKEKTSALSLSNVAGVFYILVGGGLGLAMLV  830
sp|P19492|GRIA3_RAT  VLKLNEQGLLDKLKNKWYDKGECGSGGGDSKDKTSALSLSNVAGVFYILVGGGLGLAMMV  835
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sp|P19491|GRIA2_RAT  ALIEFCYKSRAEAKRMKVAKNPQ--NINPSSSQN--SQNFATYKEGYNVVGIESVKI---  883
sp|P19492|GRIA3_RAT  ALIEFCYKSRAESKRMKLTKNQ--NFKPAPATN--TQNYATYREGYNVVGTESVKI---  888
*****:*:****          .:: : . :* .: .* . * :. :

sp|P19490|GRIA1_RAT  FPKSMQSI PCMSHSSGMPLGATGL          907
sp|P19491|GRIA2_RAT  ----- 883
sp|P19492|GRIA3_RAT  ----- 888

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Figure S6. Alignment of rat GluA1, GluA2, and GluA3 protein sequences. Putative ubiquitously modified peptides on GluA3, identified through a ubiquitinomic assay, are highlighted. Protein sequences and alignment were performed using UniProt databases.

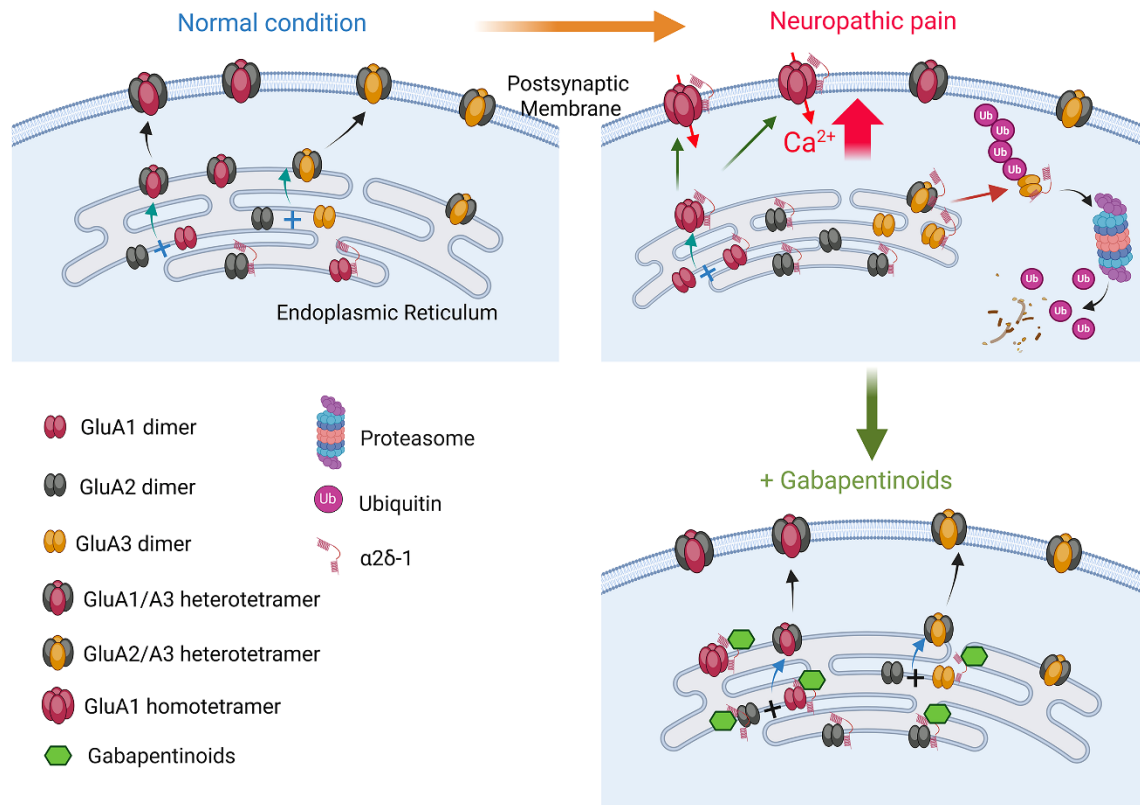


Figure S7. Graphical representations illustrating how $\alpha 2\delta -1$ facilitates the assembly and surface expression of CP-AMPA receptors at spinal cord synapses, along with associated gabapentinoid actions in neuropathic pain. Under normal conditions, $\alpha 2\delta -1$ has minimal interaction with GluA1, GluA2, or GluA3 in the spinal dorsal horn. GluA1/GluA2 and GluA2/GluA3 readily form Ca^{2+} -impermeable, heterotetrameric AMPARs for postsynaptic expression. In neuropathic pain conditions, $\alpha 2\delta -1$ expression increases and directly interacts with GluA1, GluA2, and GluA3. This interaction not only disrupts GluA1/GluA2 heteromeric assembly but also induces ubiquitination of GluA3, leading to its degradation in proteasomes. As a result of these two actions, $\alpha 2\delta -1$ promotes the assembly and synaptic expression of GluA1 homotetramers, resulting in elevated Ca^{2+} levels at spinal synapses. Gabapentinoids counteract these effects by inhibiting $\alpha 2\delta -1$'s actions, thereby restoring the heteromeric assembly and synaptic expression of GluA1/GluA2 and GluA2/GluA3.