

Supplementary Material

Loss of tumor cell MHC Class II drives MAPK-inhibitor insensitivity of BRAF-mutant anaplastic thyroid cancers

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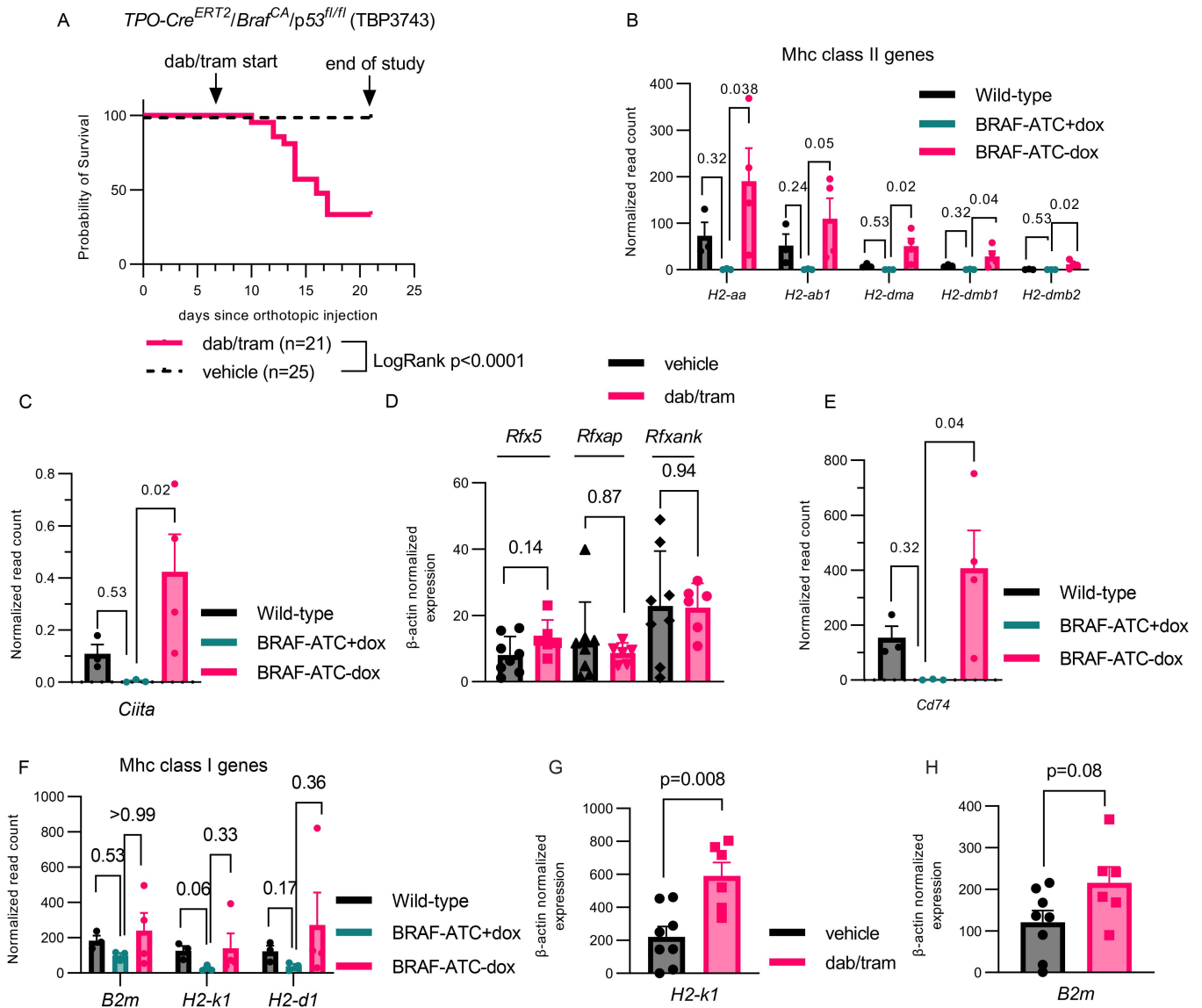
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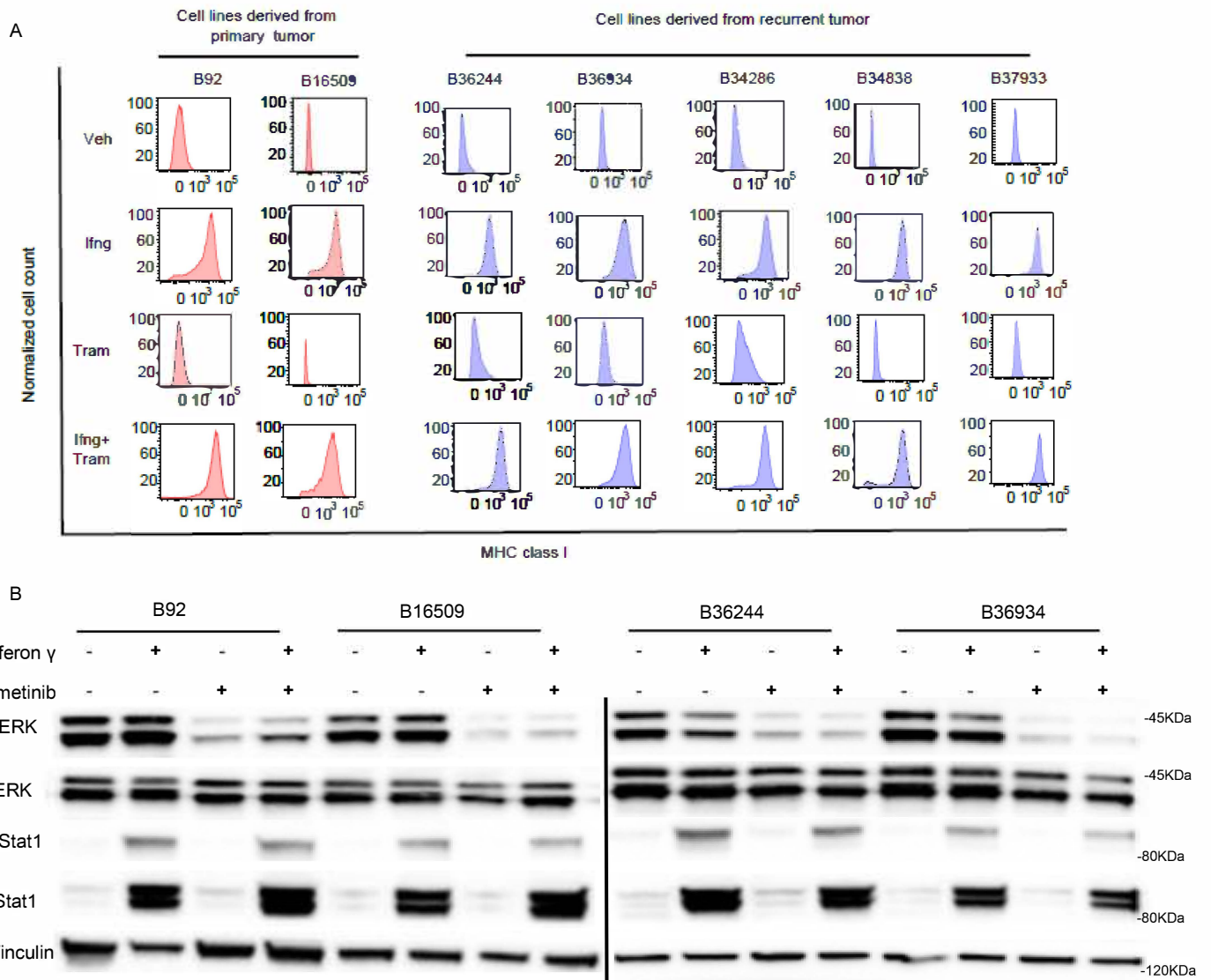
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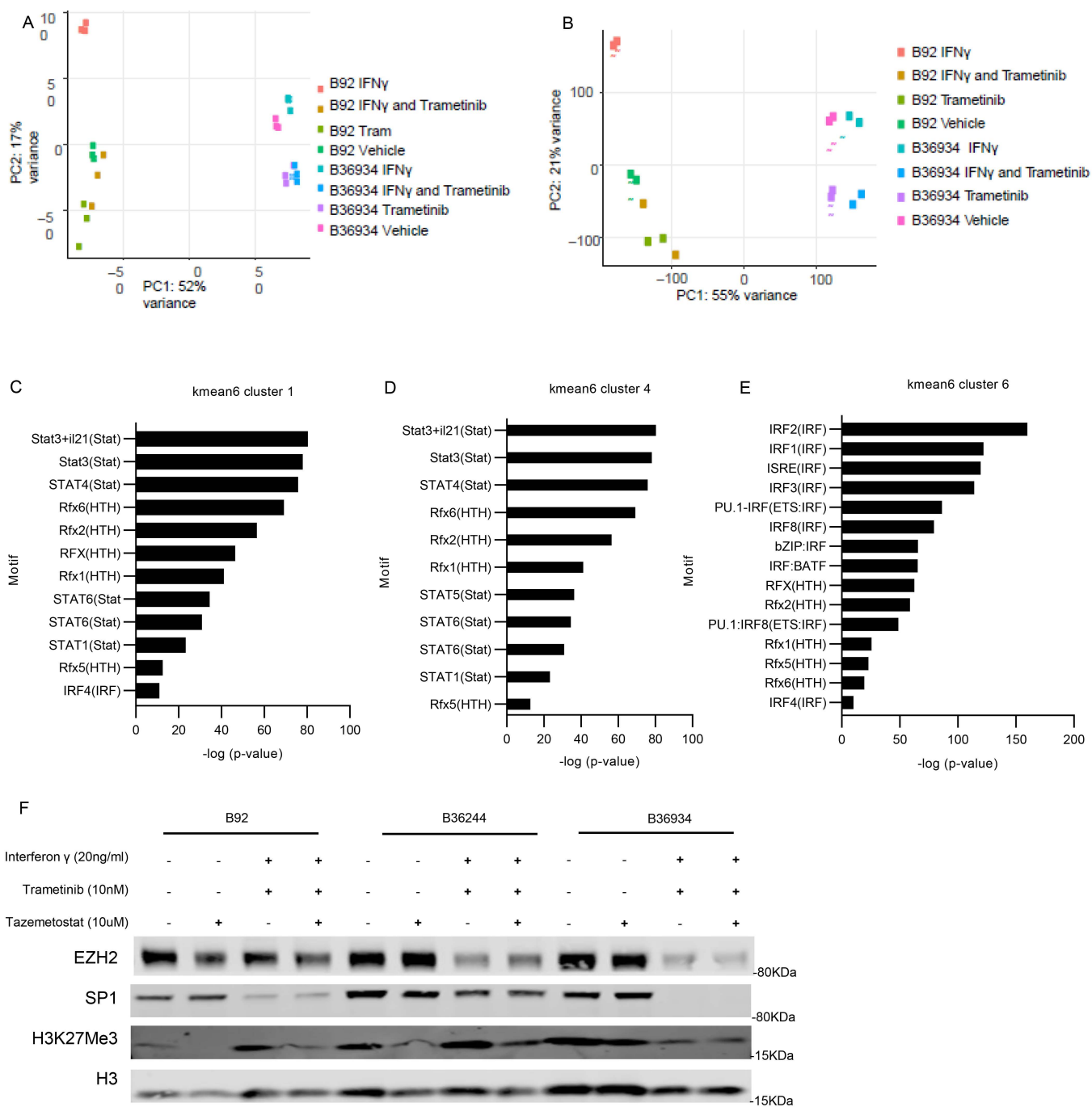
Conflict of interest statement: The authors declare no conflicts of interest with this work.



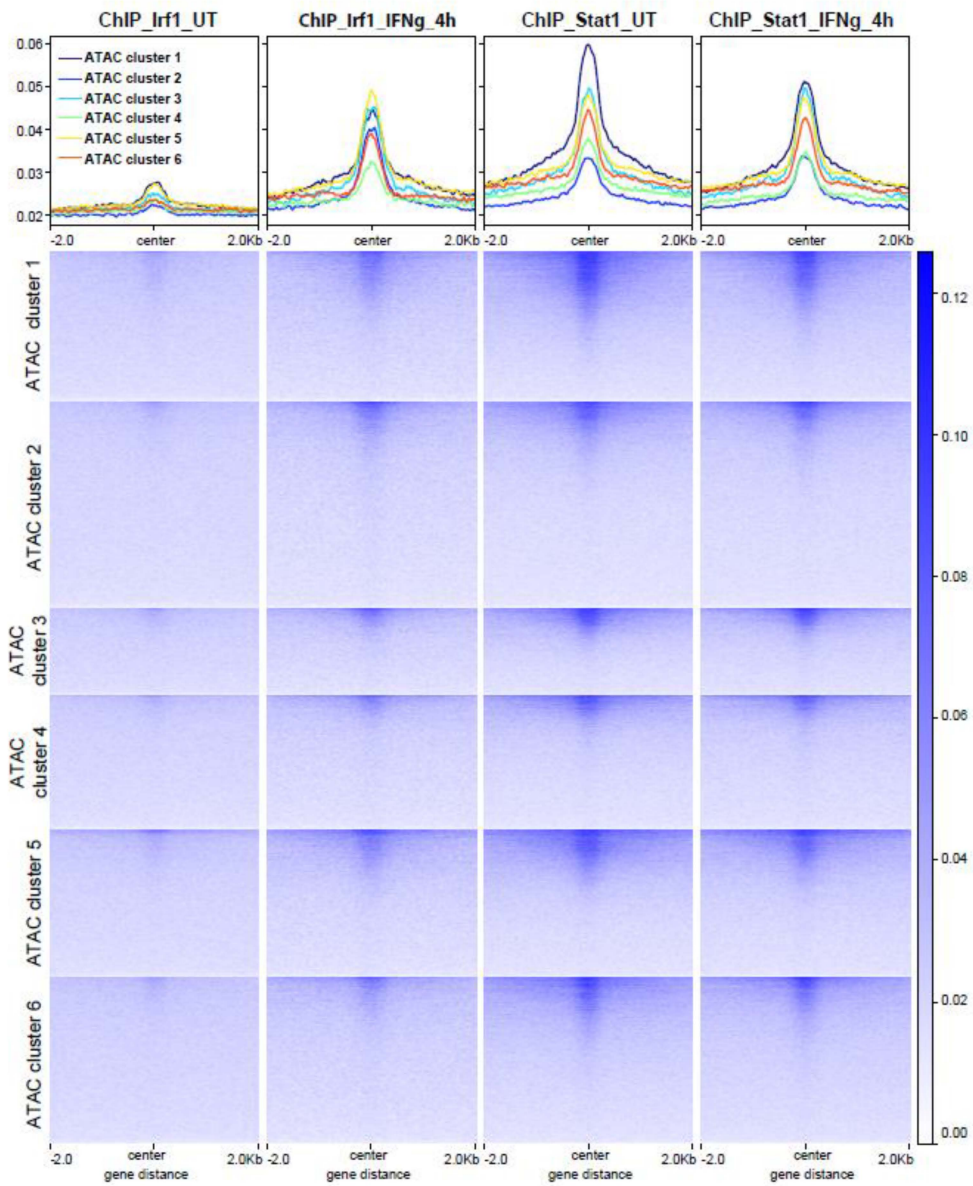
Supplementary Figure 1: A) Kaplan Meier analysis of survival of mice with orthotopic *Braf/p53* ATCs treated with vehicle or dab/tram 1 week after TBP3743 ATC cell injection. B) Expression of MhcII complex genes is low or absent in WT thyrocytes, low in BRAF/p53-ATC cells and induced by dox withdrawal. C) *Ciita* expression in WT thyrocytes and in ATC cells prior to or after dox withdrawal. D) Quantitative RT-PCR of the MhcII-related transcription factors *Rfx5*, *Rfxap* and *Rfxank* is not impacted by dab/tram treatment in the orthotopic *Braf/p53* model. E) *Cd74* expression in WT thyrocytes and in ATC cells prior to or after dox withdrawal. F) Expression of MhcI genes in WT thyrocytes or ATC cells prior to or after dox withdrawal. G and H) RT-PCR of *H2-k1* and *B2m* mRNAs in sorted thyrocytes from *Braf/p53* orthotopic ATCs treated with vehicle or dab/tram in vivo for 4 days. Log-rank (Mantel-Cox) test (A), Kruskal Wallis Test (B, C, E and F) Multiple Mann-Whitney tests (D, G and H). Bars represent SEM. ATC: Anaplastic thyroid cancer; IFN γ : Interferon γ ; GEMM: Genetic engineered mouse model; dab/tram: Dabrafenib and trametinib; *Ciita*: Class II major histocompatibility complex transactivator; WT: wild type; SEM: Standard error of the mean.



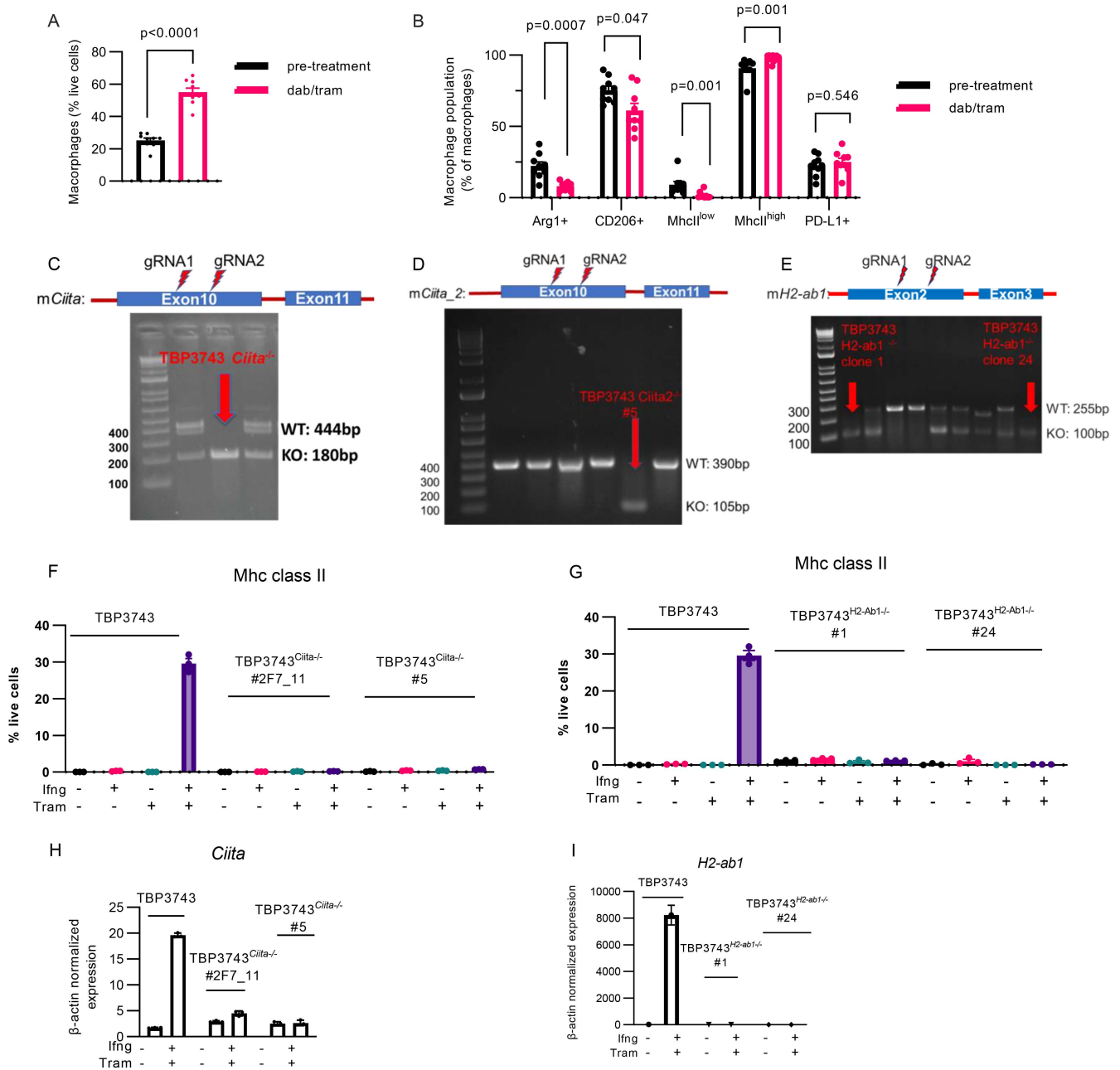
Supplementary Figure 2: A) Induction of Mhcl by IFN γ (20ng/ml) as determined by FACS in two primary cell lines derived from primary ATCs on dox diet (B92 and B16509) and five lines derived from recurrent tumors developing after dox withdrawal (B36244, B36934, B34286, B34838 and B37933). Addition of trametinib (10nM) does not significantly augment the IFN γ (20ng/ml) effect. B) Western blot of B92, B16509, B36244, B36934 cell lysates treated with trametinib alone or in combination with IFN γ for 96 h for the indicated proteins. IFN γ : Interferon γ ; dox: doxycycline.



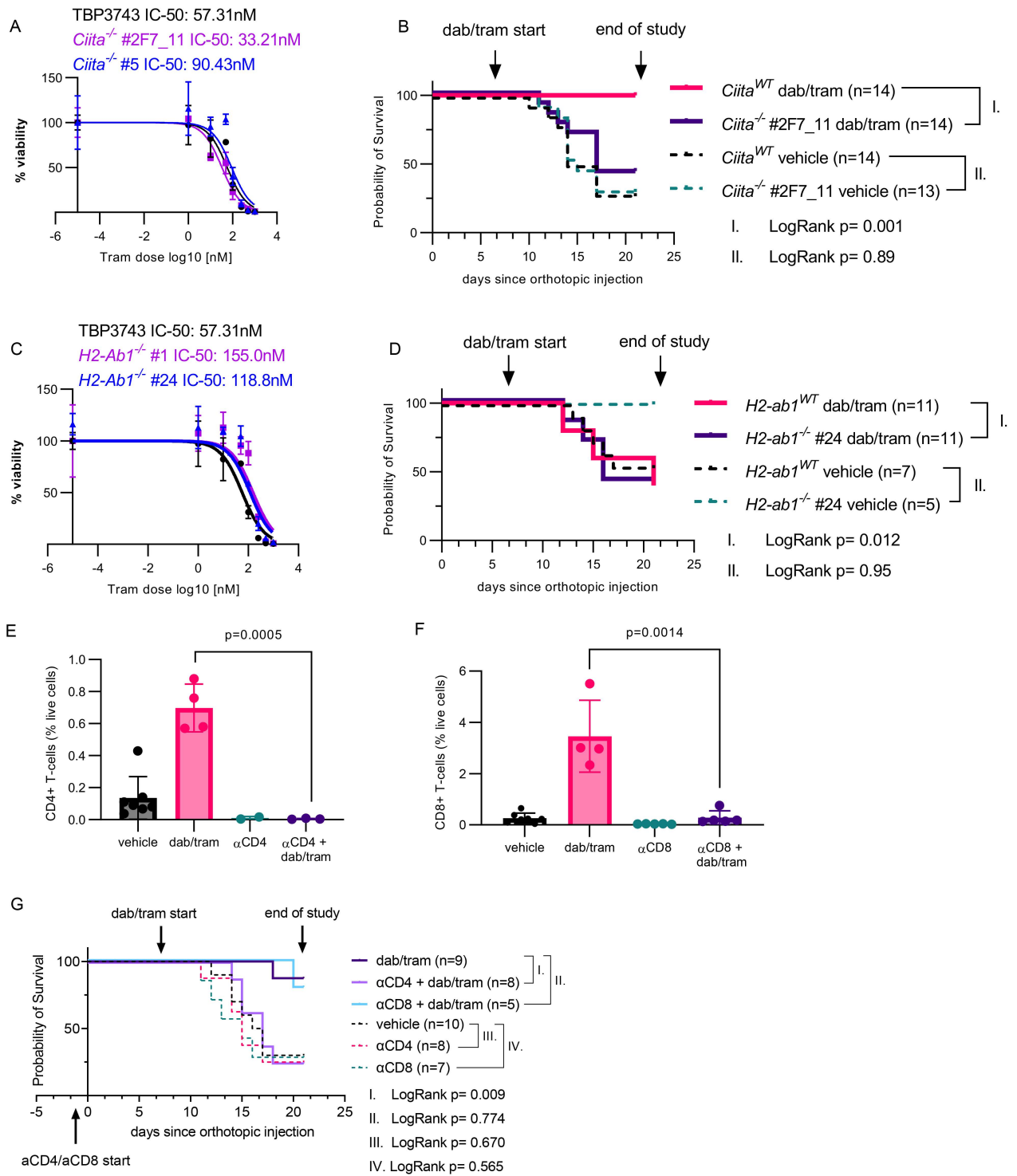
Supplementary Figure 3: A,B) Principal component analyses of RNA Seq and ATAC-Seq for the B92 primary and the B36934 recurrent ATC lines displaying triplicates and duplicates, respectively, for each treatment condition (DMSO, IFN γ (20ng/ml), trametinib (10nM) and the combination of IFN γ + trametinib (10nM) for 96 h. C–E) TF motifs of members of the Rfx, Stat and Irf families enriched in kmeans clusters 1, 4 and 6 identified using HOMER *de novo* motif discovery. F) Western blot for EZH2, Sp1, H3K27Me3 and H3 for the indicated cell lines and treatment conditions. IFN γ : Interferon γ ; tram: trametinib; TF: transcription factor.



Supplementary Figure 4: Tornado plots of averaged Stat1 and Irf1 transcription factor binding sites in the respective k-means clusters.



Supplementary Figure 5: A, B) Analysis of infiltrating macrophages by flow cytometry in orthotopic Braf/p53 ATC at baseline and 5 days after treatment with dab/tram. C, D) Top: Schema of two different gRNA pairs targeting Exon 10 for CRISPR KO of *Ciita*. Bottom: Image of gel electrophoresis of PCR products from the indicated CRISPR KO clones. E) Top: Schema for gRNA pair targeting exon2 of *H2-ab1* for generation of CRISPR KO clones. Bottom: Image of gel electrophoresis of PCR products of two homozygous *H2-ab1* KO clones. F, G) Loss of IFN γ and trametinib-induced MhclI in the *Ciita* CRISPR KO clones 2F7_11 and 5 and *H2-ab1* CRISPR KO clones 2 and 24, all derived from TBP3743 cells. MhclI was measured by FACS 96h after treatment with the indicated conditions. H, I) Quantitative RT-PCR of *Ciita* mRNA in parental and *Ciita*^{-/-} clones and *H2-ab1* mRNA in parental and *H2-ab1*^{-/-} clones. gRNA: guide RNA; IFN γ : Interferon γ ; *Ciita*: Class II major histocompatibility complex transactivator.



Supplementary Figure 6: A, C) IC50 for trametinib in parental, *Ciita*^{-/-} clones 2F7_11 and 5 and *H2-ab1*^{-/-} clones 1 and 24. B, D) Kaplan Meier analysis of survival of mice with orthotopic *Braf*/*p53* ATCs, treated with vehicle or dab/tram for 1 week after orthotopic injection of TBP3743 parental or *Ciita*^{-/-} or *H2-ab1*^{-/-} ATC cells. E) and F) CD4⁺ (E) and CD8⁺ T-cells (F) percentage of live cells in thyroids harvested from mice treated with in the indication conditions. G) Kaplan Meier analysis of survival of mice with orthotopic *Braf*/*p53* ATCs in the indicated treatment conditions. Tram: trametinib. dab/tram: dabrafenib and trametinib. *Ciita*: Class II major histocompatibility complex transactivator.

Supplementary Tables

Table 1 Tumor mutational burden in human and murine papillary and anaplastic thyroid cancer

Table 2 Whole exome sequencing results: High confidence Maf

Table 3 Antibodies for IHC, IF, FACS and WB

Table 4 CrisprKO gRNA Sequences

Table 5 Primers

Supplementary Table 1

	Human ATC	Murine ATC
Mutation per tumor	23.7 ¹	22 ³
	24 ²	

Reference

¹Kunstman et al 2015 (28)

²Ravi et al 2019 (29)

³Knauf et al 2018 (26)

Supplementary Table 3

Antibody	Clone	Source	Target Species
HLA-DR	TAL 1B5	Santa Cruz	human
Pan Cytokeratin	na	Thermo Fisher	human
CD8	4B11	Thermo Fisher	human
CD3	SP7	Abcam	human
CD163	10D6	Abcam	human
CD68	KP1	Abcam	human
CD15	MMA	Abcam	human
BUV395 Rat Anti-Mouse CD45	30-F11	BD Biosciences	mouse
BV510 Rat Anti-CD11b	M1/70	BD Biosciences	mouse
BV750 Rat Anti-Mouse Siglec-F	E50-2440	BD Biosciences	mouse
FITC Rat Anti-Mouse CD3	17A2	BD Biosciences	mouse
BUV805 Rat Anti-Mouse CD8a	53-6.7	BD Biosciences	mouse
BUV661 Mouse Anti-Mouse NK-1.1	PK136	BD Biosciences	mouse
PerCP anti-mouse Ly-6G Antibody	1A8	Biolegend	mouse
Brilliant Violet 711™ anti-mouse CD11c Antibody	N418	Biolegend	mouse
Spark Blue™ 550 anti-mouse I-A/I-E Antibody	M5/114.15.2	Biolegend	mouse
Brilliant Violet 650™ anti-mouse F4/80 Antibody	BM8	Biolegend	mouse
Brilliant Violet 570™ anti-mouse CD4 Antibody	RM4-5	Biolegend	mouse
Spark NIR™ 685 anti-mouse/human CD45R/B220 Antibody	RA3-6B2	Biolegend	mouse
APC-Cy™7 Rat Anti-Mouse Ly-6C	AL-21	BD Biosciences	mouse
Arginase 1 Monoclonal Antibody PerCP-eFluor™ 710 ¹	A1exF5	Thermo Fisher	mouse
FOXP3 Monoclonal Antibody eFluor™ 450 ¹	FJK-16s	Thermo Fisher	mouse
CD274 (PD-L1, B7-H1) Super Bright™ 436	MIH5	Thermo Fisher	mouse
BV605 Rat Anti-Mouse I-A/I-E	M5/114.15.2	BD Biosciences	mouse
LIVE/DEAD™ Fixable Blue Dead Cell Stain Kit, for UV excitation	na	Thermo Fisher	mouse
Alexa Fluor® 647 anti-mouse CD206 ¹	MMR	Biolegend	mouse
PE Mouse Anti-Mouse H-2Kb	AF6-88.5	BD Biosciences	mouse
Phospho-p44/42 MAPK (Erk1/2) (Thr202/Tyr204) (D13.14.4E) XP® Rabbit mAb		Cell Signaling	
p44/42 MAPK (Erk1/2) (L34F12) Mouse mAb		Cell Signaling	
Phospho-Stat1 (Ser727) (D3B7) Rabbit mAb		Cell Signaling	
Stat1 (D1K9Y) Rabbit mAb		Cell Signaling	
Vinculin (E1E9V) XP® Rabbit mAb		Cell Signaling	
SP1 (D4C3) Rabbit mAb		Cell Signaling	
Ezh2 (D2C9) XP® Rabbit mAb		Cell Signaling	
Histone H3 (D1H2) XP® Rabbit mAb		Cell Signaling	
Tri-Methyl-Histone H3 (Lys27) (C36B11) Rabbit mAb		Cell Signaling	

¹ Intracellular antibodies

Supplementary Table 4

Vector name	gRNA Sequence	Source	Identifier
mCiita_1	AGCAGGCCAAGACTTACATG, TAGTCGAGCTGGCCAAGCTG	Vectorbuilder	VB210324-1175xee
mCiita_2	CCCGGAGCCTTAGTCGAGCT, GAGACCCTATGACAACTGG	Vectorbuilder	VB230124-1305eyk
mH2-ab1	ACGGGACGCAGCGCATACGA, GGAGATCCTGGAGCGAACGC	Vectorbuilder	VB230124-1307xfy

Supplementary Table 5

Primer	Sequence
mCiita_ Ex3F	ATCTTCCAGCGGAAGCTACTGC
mCiita_ Ex3R	CCGGGTTTCTTGCAAGGTGC
mCiita_ Ex10F	GGACTCTATGTCAGCCTGCTAGG
mCiita_ Ex10R	TGGGCTCGAGGCTGGAAAAC
mH2-ab1_ Ex2F	CCGCAGGGCATTTCGTGTAC
mH2-ab1_ Ex2R	TCTCCGGCCCCTCGTAGTTGT
mCD74_ FW	AAGCAGTGGCTCTTGTTTGAG
mCD74_ RV	CTTCCATGTCCAGTGGCTCT
mCiita_ FW	AATCTACCACGGTGAGATGCCC
mCiita_ RV	TCGGGGAGACTGGGGATACTGA
mH2-aa_ FW	GAGCAGCTTCAGAGACCTCC
mH2-aa_ RV	CTACGTGGTCGGCCTCAAT
mH2-ab1_ FW	CACAGGAGTCAGAAAGGACCTC
mH2-ab1_ RV	TGGCAGTCAGGAATTCGGAG
mH2-dma_ FW	GAGATTGACCGCTACACGGCAA
mH2-dma_ RV	GAAGACAATGCCCATGATGGTGC
mH2-dmb1_ FW	AGAGCCTTCTCCAGCGTTTGCA
mH2-dmb1_ RV	TGTGGTTTGGGCTACTCGGACA
mH2-dmb2_ FW	ACCTTTCTGGGATGTGCTGACC
mH2-dmb2_ RV	GTGATGGTCACATCCGCTGGAT
mRfxap_ FW	ACGTCAAACCTGGAGGAAAGCAC
mRfxap_ RV	GAGTAGGTCTTGCAGGGCG
mRfx5_ FW	GGAAGACCTTGGTATCCATGCC
mRfx5_ RV	GGCTGCTTCTACCAGTTCATCC
Rfxank_ FW	GCACATGCCTGTCTGGAAAC
Rfxank_ RV	AGCAGGAAGCGAACTGTCTC
Irf1_ FW	CAAAGCCACCATGCCAATCACTCG
Irf1_ RV	GGCCCAGCTCCGGAACAGACAG
mH2-k1_ FW	GGAGCAGGAGGGGCCCGAGTATTG
mH2-k1_ RV	CGCCGTCCACGTTTTTCAGGTCTTC
mB2m_ FW	TCACTGACCGGCTGTATGCTATC
mB2m_ RV	AATGTGAGGCGGGTGGAACTGT