# **Supplemental Figures and Table for:**

# STING Activation Reprograms the Microenvironment to Sensitize NF1-related Malignant Peripheral Nerve Sheath Tumors for Immunotherapy

Bandarigoda N. Somatilaka<sup>1</sup>, Laasya Madana<sup>1</sup>, Ali Sadek<sup>1</sup>, Zhiguo Chen<sup>1</sup>, Sanjay Chandrasekaran<sup>2,3</sup>, Renee M. McKay<sup>1</sup>, and Lu Q. Le<sup>1,2,4,5,6,7</sup>

<sup>1</sup>Department of Dermatology, <sup>2</sup> Simmons Comprehensive Cancer Center, <sup>3</sup>Department of Internal Medicine, Division of Hematology/Oncology, <sup>4</sup>UTSW Comprehensive Neurofibromatosis Clinic, <sup>5</sup>Hamon Center for Regenerative Science and Medicine, <sup>6</sup>O'Donnell Brain Institute, University of Texas Southwestern Medical Center at Dallas, Dallas, Texas, 75390-9069, USA, <sup>7</sup>Department of Dermatology, University of Virginia School of Medicine, Charlottesville, VA, 22903, USA

#### Author for correspondence:

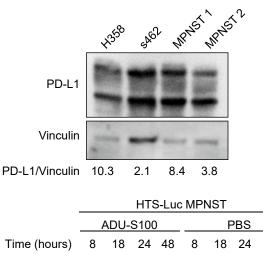
Lu Q. Le, M.D., Ph.D. Professor and Chair Department of Dermatology University of Virginia School of Medicine Phone: (434) 982-5974 Fax: (434) 244-4504 E-mail: bkn6qd@uvahealth.org

**Short title:** Reprogramming MPNST for immune checkpoint blockade **Conflicts of interest:** The authors have declared that they have no competing interests.

### **Keywords**

Neurofibromatosis Type 1, NF1, neurofibroma, plexiform neurofibroma, malignant peripheral nerve sheath tumor, MPNST, STING, immune checkpoint blockade, ICB, ADU-S100

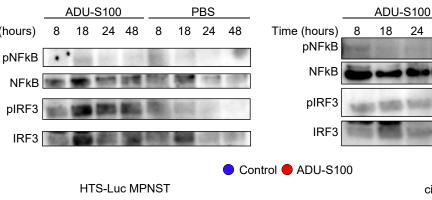
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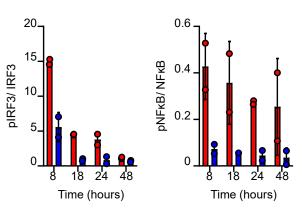


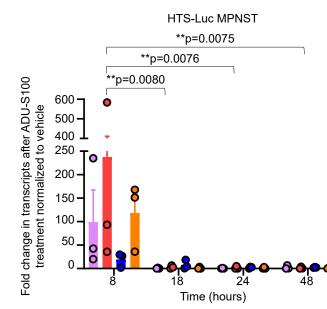


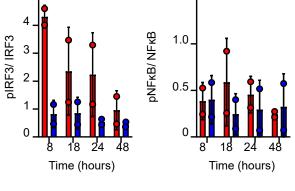
С

D









cisMPNST

1.5

cisMPNST

8

24 48

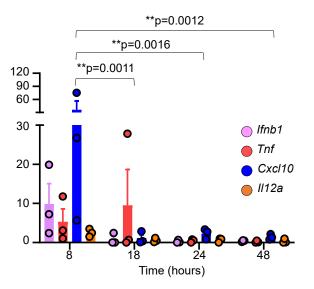
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PBS

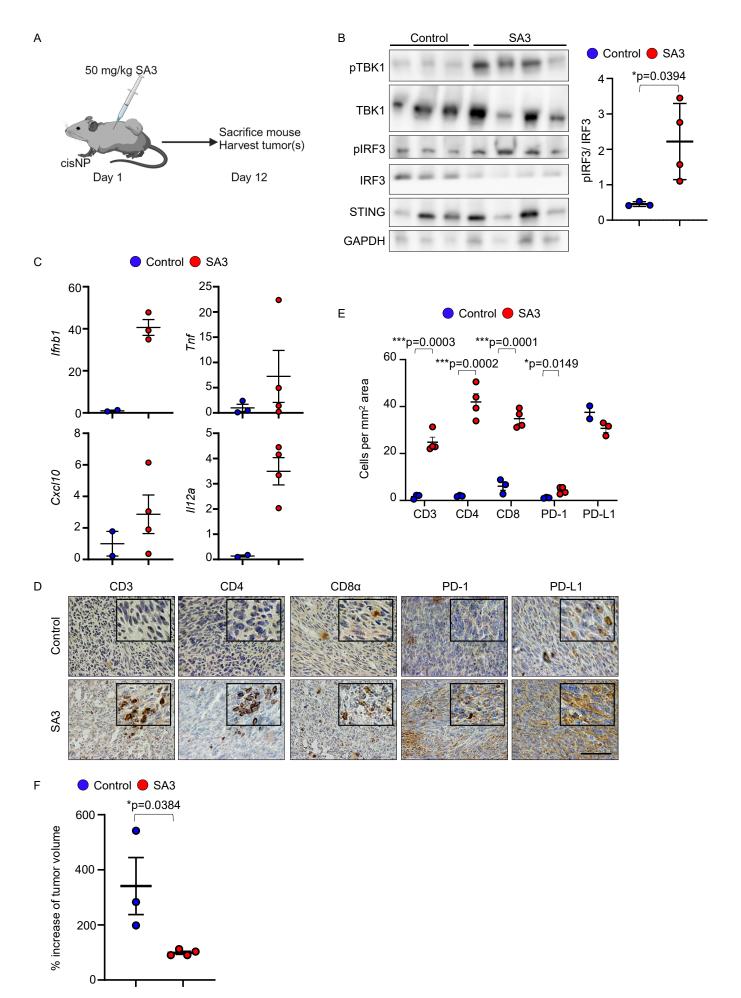
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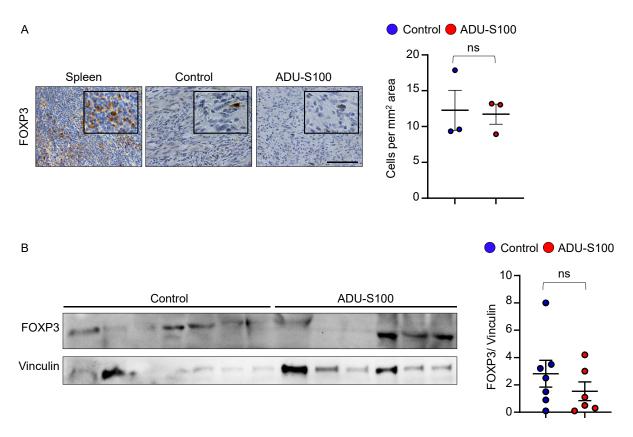
cisMPNST



Supplemental Figure 1. STING agonist ADU-S100 activates the STING pathway in MPNST cell lines. (A) Western blot of the indicated cell lines and 2 cisMPNST tumors for PD-L1. The total PD-L1 band intensity normalized to that of vinculin is shown below each lane. (B) Western blot analysis for expression of the indicated proteins in mouse MPNST cell lines derived from *Nf1* and *p53* null skin progenitor cells (HTS-Luc MPNST) or harvested from *cisNP* mice (*cis*MPNST) treated with vehicle control (PBS) or ADU-S100. (C) Quantified protein band intensities from (B) are shown graphically. Data shown are as mean  $\pm$  SD. (D) PCR analysis of fold change in cytokine gene expression (*lfnb1*, *Tnf*, *Cxcl10*, and *ll12a*) in MPNST cells as in (B) treated with PBS (n=3/ time point) or ADU-S100 (n=3/ time point). Data are represented as mean  $\pm$  SEM and p values are determined by Tukey's multiple comparisons test as indicated (D). \*\**P* < 0.01.

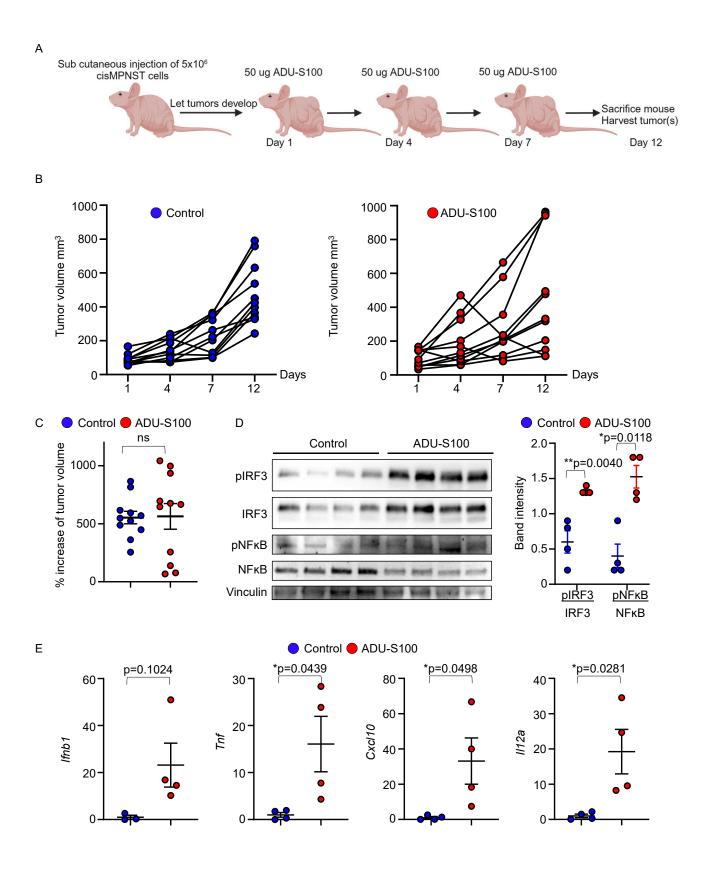


Supplemental Figure 2. SA3 treatment of *cisNP* mice activates the STING pathway in tumors. (A) Schema of SA3 treatment protocol. (B) Western blot analysis for expression of the indicated proteins in MPNSTs harvested from *cisNP* mice treated with vehicle control or SA3. Quantified protein band intensities for pIRF3/ IRF3 are shown graphically on the right. (C) PCR analysis of fold change in cytokine gene expression (*lfnb1*, *Tnf*, *Cxcl10*, and *ll12a*) in *cis*MPNSTs harvested from control-treated (n=3) and SA3-treated (n=4) mice. (D) Paraffin sections of MPNSTs harvested from vehicle-treated or SA3 treated *cisNP* mice were stained with antibodies against CD3, CD4, CD8 $\alpha$ , PD-1, and PD-L1 and quantified (E). (F) Percent increase in tumor volume was measured in control-treated or SA3-treated *cisNP* mice. Data are represented as mean ± SEM and p values are determined by two-tailed t test as indicated. \**P* < 0.05. Scale bar: 50 um.

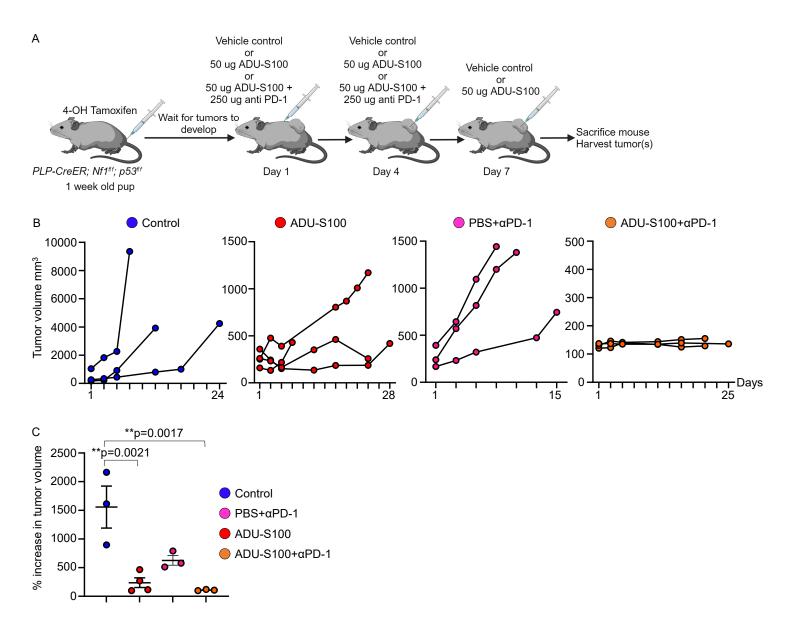


### Supplemental Figure 3. Foxp3 expression is unaltered upon ADU-S100 treatment. (A)

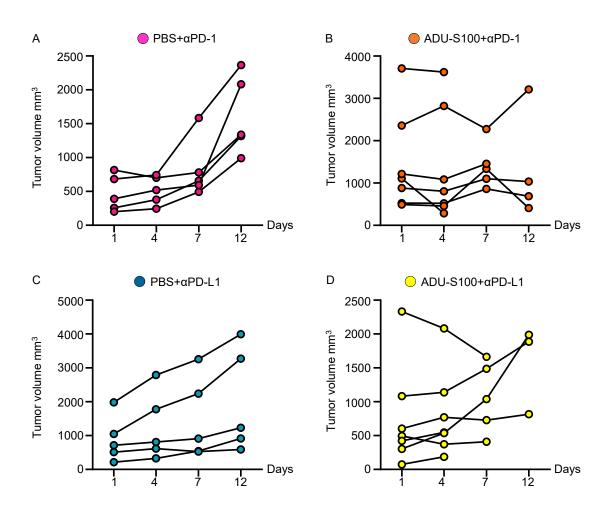
Paraffin sections of murine spleen and control-treated (n=3) or ADU-S100-treated (n=3) *cis*MPNST are stained for FOXP3. Quantification of FOXP3-positive cells is shown on the right. (**B**) Western blot analysis for expression of FOXP3 in MPNSTs harvested from *cisNP* mice treated with vehicle control (n=7) or ADU-S100 (n=6). Quantified protein band intensities are shown graphically on the right. Data are represented as mean  $\pm$  SEM and p values are determined by unpaired t test as indicated. ns = not significant.



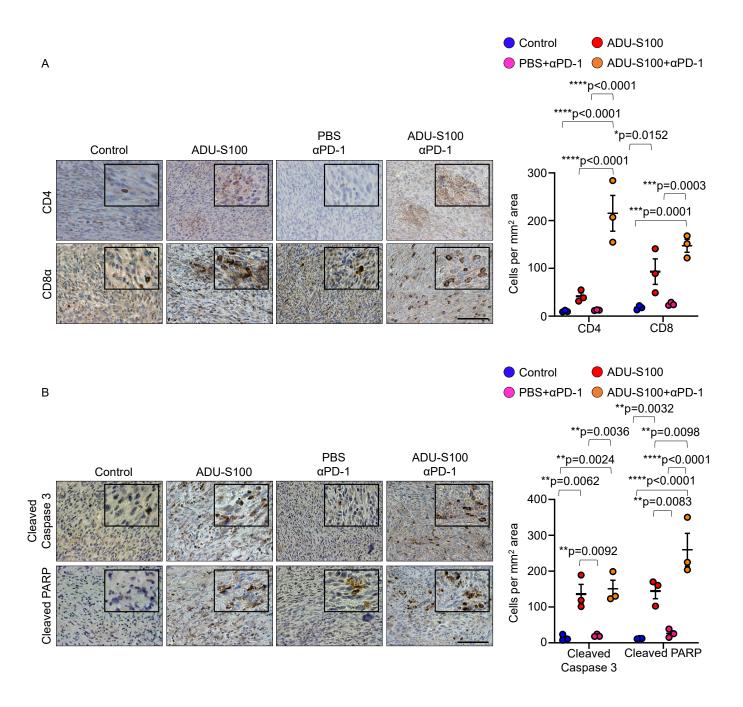
Supplemental Figure 4. STING agonist does not decrease MPNST volume in allograft MPNST in nude mice. (A) Schema of ADU-S100 treatment protocol in athymic nude mice bearing *cis*MPNST. (B) Tumor volume change over time in response to indicated treatments. (C) Tumor volume increase in athymic mice treated with PBS control (n=11) or ADU-S100 (n=11) as shown in (A). (D) Western blot analysis for expression of the indicated proteins in MPNSTs harvested from athymic mice treated with vehicle control (n=4) or ADU-S100 (n=4) for 24 hours. Quantified protein band intensities are shown graphically on the right. (E) PCR analysis of fold change in cytokine gene expression (*Ifnb1*, *Tnf*, *Cxcl10*, and *Il12a*) in *cis*MPNSTs harvested from control-treated (n=4) and ADU-S100-treated (n=4) athymic mice 24 hours after treatment. Data are represented as mean  $\pm$  SEM and p values are determined by two-tailed t test with respect to vehicle control. \**P* < 0.05, \*\**P* < 0.01, ns = not significant.



Supplemental Figure 5. Combination treatment of STING agonist and ICB decreases MPNST growth in *PLP-Cre; Nf1<sup>ff</sup>; p53<sup>ff</sup>* mice. (A) Schema of STING activation and ICB combination treatment protocol in *PLP-Cre; Nf1<sup>ff</sup>; p53<sup>ff</sup>* mice. (B) Tumor volume change of MPNST in *PLP-Cre; Nf1<sup>ff</sup>; p53<sup>ff</sup>* mice in response to the indicated treatments. (C) Percent increase in tumor volume in *PLP-Cre; Nf1<sup>ff</sup>; p53<sup>ff</sup>* mice upon indicated treatments. Control, n=3; ADU-S100, n=4; PBS +  $\alpha$ PD-1, n=3; ADU-S100 +  $\alpha$ PD-1, n=3. Data are represented as mean  $\pm$  SEM and p values are determined by Tukey's multiple comparisons test as indicated. \**P* < 0.05, \*\**P* < 0.01.

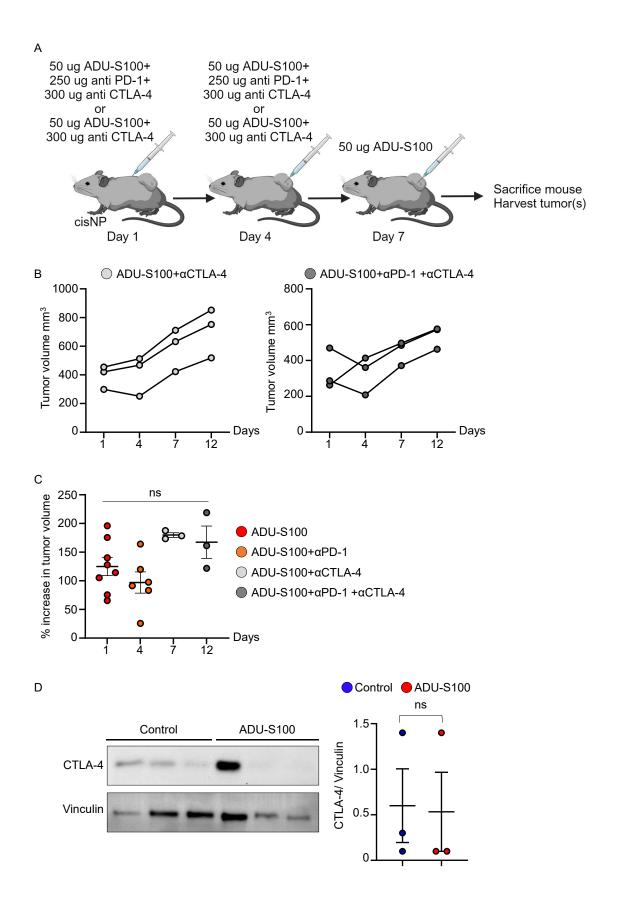


Supplemental Figure 6. Combination treatment of *cis*MPNSTs with STING activation plus ICB slows tumor growth. Tumor volume change over time in *cisNP* mice treated with (A) PBS +  $\alpha$ PD-1, (B) ADU-S100 +  $\alpha$ PD-1, (C) PBS +  $\alpha$ PD-L1, or (D) ADU-S100 +  $\alpha$ PD-L1.



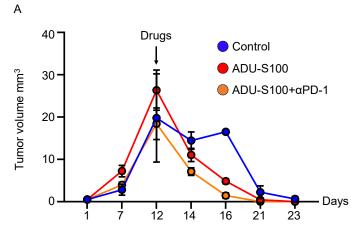
Supplemental Figure 7. Combination treatment of STING agonist plus ICB increases T cell infiltration and expression of apoptotic markers in MPNST in *PLP-Cre; Nf1<sup>ff</sup>; p53<sup>ff</sup>* mice.

(A) Paraffin sections from MPNSTs harvested from *PLP-Cre; Nf1<sup>ff</sup>; p53<sup>ff</sup>* mice treated as indicated were stained for T cell markers. Quantification of images are shown on the right. (B) Paraffin sections from MPNSTs in (A) stained for Cleaved Caspase 3 and Cleaved PARP. Quantification of images are shown on the right. Control, n=3; ADU-S100, n=3; PBS +  $\alpha$ PD-1, n=3; ADU-S100 +  $\alpha$ PD-1, n=3. Data are represented as mean ± SEM and p values are determined by Tukey's multiple comparisons test as indicated. \**P* < 0.05, \*\**P* < 0.01, \*\*\**P* < 0.001. Scale bar: 50 µm.

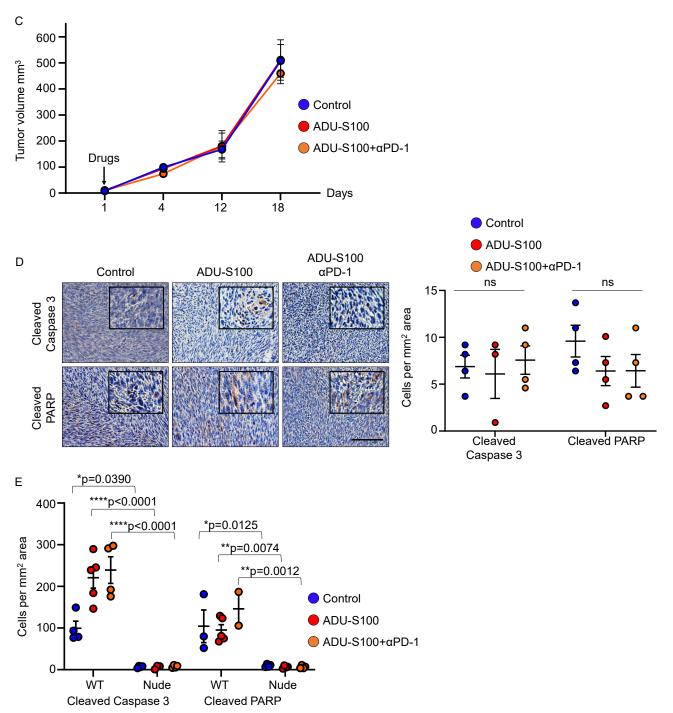


Supplemental Figure 8. Combination treatment of STING agonist plus αPD-1 and αCTLA-4 antibodies does not further decrease MPNST volume compared to STING agonist plus αPD-1 treatment.

(A) Schema of ADU-S100 plus  $\alpha$ PD-1 and  $\alpha$ CTLA-4 treatment protocol in *cisNP* mice. (B) Tumor volume changes in response to the indicated treatments. (C) Percentage increase in tumor volume upon indicated treatments. Data sets for ADU-S100 and ADU-S100 plus  $\alpha$ PD-1 used in this graph are borrowed from Figure 4D for clarity and ease of comparison. ADU-S100 +  $\alpha$ CTLA-4, n=3; ADU-S100 +  $\alpha$ PD-1 +  $\alpha$ CTLA-4, n=3. (D) Western blot analysis for expression of CTLA-4 in MPNSTs harvested from *cisNP* mice treated with vehicle control (n=3) or ADU-S100 (n=3). Quantified protein band intensities are shown graphically on the right. Data are represented as mean ± SEM and p values are determined by Tukey's multiple comparisons test in (C) and two-tailed t test in (D) as indicated. ns = not significant.



В		Control vs ADU-S100	Control vs ADU-S100 + αPD-1	ADU-S100 vs ADU-S100 + αPD-1
	Day 14	ns	p=0.0169	ns
	Day 16	p<0.0001	p<0.0001	p=0.0081



Supplemental Figure 9. Combination treatment of STING activation plus ICB accelerates tumor regression in a human MPNST xenograft mouse model. (A) Difference in tumor volume in human MPNST xenograft mice upon treatment with PBS (n=15), ADU-S100 (n=15), or ADU-S100 +  $\alpha$ PD-1 (n=15). (B) Statistical significance of the volume differences in (A) determined by multiple unpaired t tests. (C) Difference in tumor volume in human MPNST xenograft in nude mice upon treatment with PBS (n=4), ADU-S100 (n=4) or ADU-S100 +  $\alpha$ PD-1 (n=4). (D) Paraffin sections from nude mouse xenograft MPNST treated as indicated were stained for Cleaved Caspase 3 and Cleaved PARP. (E) Comparison of Cleaved Caspase 3 and Cleaved PARP levels in wild-type mouse (repeated from data in Figure 6D) and nude mouse xenograft MPNST. Data are represented as mean ± SEM and p values are determined by Tukey's multiple comparisons test as indicated. Scale bar: 50 µm. \**P* < 0.05, \*\**P* < 0.01, \*\*\**P* < 0.001, ns = not significant.

Ifnb1	mIfnb1-F1	5' CCC TAT GGA GAT GAC GGA GA 3'
	mIfnb1-R1	5' ACC CAG TGC TGG AGA AAT TG 3'
Tnf	mTnf-F1	5' ACG GCA TGG ATC TCA AAG AC 3'
	mTnf-R1	5' GTG GGT GAG GAG CAC GTA GT 3'
Cxcl10	mCxcl10-F1	5' GCT GCA ACT GCA TCC ATA TC 3'
	mCxcl10-R1	5' GTG GCA ATG ATC TCA ACA CG 3'
Il12a	mIl12a-F1	5' CTC CTG TGG GAG AAG CAG AC 3'
	mIl12a-R1	5' CAG ATA GCC CAT CAC CCT GT 3'

Supplemental Table 1: Primer sequences for qRT-PCR