SUPPLEMENTARY INFORMATION

# Oncolytic Virotherapy Induced CSDE1 Neo-antigenesis Restricts VSV Replication

But Can Be Targeted By Immunotherapy

## **Supplementary Figure 1**

Sanger sequencing shown begining at position 7-14 in CSDE1

A B16 Parental Exclusively GATCC (WT)



D B16-VSV-mIFNβ ESC Excusively GATTC (MUT)



G Hep3B Parental Exclusively GATCC (WT)



TTTGATCC

J Mel888 in vivo ESC VSV-hIFNβ tumor 1



B B16-VSV-GFP ESC Predominantly GATTC (MUT) Minority GATCC (WT)



E B16-shRNA mAPOBEC3-VSV-GFP ESC Exclusively GATCC (WT)



H Hep3B-VSV-GFP ESC Predominantly GATCC (WT) Minority GATTC (MUT)



TTTGATCC

Mel888 in vivo ESC VSV-hIFNβ tumor 3



C B16TK-GCV 21d ESC Exclusively GATCC (WT)



F B16-shRNA mAPOBEC3-VSV-mIFNβ ESC Exclusively GATCC (WT)



I Hep3B-VSV-hIFNβ ESC Excusively GATTC (MUT)



Supplementary Figure 1. Escape from VSV-IFN-ß selects for the CSDE1<sup>C-T</sup> **mutation.** B16 murine melanoma or Hep3B human HCC cells were infected (MOI 0.01) for 21d as in **Methods** and <sup>1</sup>. Surviving cells were pooled and genomic DNA prepared. Sanger sequencing of CSDE1 is shown for: A. Parental, uninfected B16 showing a homogenous population of wild type CSDE1 sequence; **B.** B16-VSV-GFP-ESC showing a mixed population of escape cells with either wild type ATCC or mutated ATTC sequence; **C.** B16 cells stably expressing the HSVtk suicide gene<sup>2</sup>, selected in GCV for 21d; D. B16-VSV-IFNß-ESC showing a nearly homogenous population of escape cells with mutated ATTC sequence; **E&F.** B16 cells stably expressing shRNA against APOBEC3 as in <sup>1</sup> selected for escape from VSV-GFP (**E**) or VSV-IFNß (**F**); **G.** Parental, uninfected Hep3B; H. Hep3B-VSV-GFP-ESC; and I. Hep3B-VSV-IFNß-ESC. J. Nude mice bearing s.c. Mel888 human melanoma tumors treated with i.t. VSV-hIFNß as described in <sup>1</sup> were excised upon escape from treatment, genomic DNA prepared and Sanger sequencing used to characterize the CSDE1 gene as shown. 4 of 4 excised, VSV-hIFNß-escaped tumors had mixed populations of CSDE1<sup>WT</sup> and CSDE1<sup>C-T</sup> tumor cells at ratios close to 1:1. Results are representative of seven separate clones (A-D); 3 separate clones (E,F) and (G-I); and 4 separate escape tumors (J).

### **Supplementary Figure 2**



Supplementary Figure 2. DC-CSDE1<sup>C-T</sup> vaccination. A. In vitro IL-4/GM-CSF matured murine DC were transfected with nothing, CSDE1<sup>WT</sup> or CSDE1<sup>C-T</sup> expression plasmids and 48 hrs later administered as vaccines with i.t. treatments with VSV-mIFN-ß and  $\alpha$ -PD1. **B.** On d40, spleens and LN were assayed for IFN- $\gamma$  following *in vitro* restimulation with live B16 cells or with live B16 cells stably expressing CSDE1<sup>C-T</sup>. WT: DC vaccines expressing wild type CSDE1; \* DC vaccine expressing mutated CSDE1<sup>C-T</sup>. C. C57BI/6 mice with 10d B16 tumors were injected i.t. with PBS or with VSV-mIFNß, (10<sup>7</sup>pfu/injection) and i.p. with unloaded DC (DC), DC vaccine expressing wild type CSDE1 (DC-CSDE1<sup>WT</sup>) or with DC vaccine expressing mutated CSDE1<sup>C-T</sup> (CSDE1<sup>C-T</sup>) followed by  $\alpha$ -PD-1 (n=8/grp) as shown in **A**. Kaplan-Meier survival for groups is shown. Representative of two separate experiments. P-values were determined using the Logrank Mantel Cox test. For multiple comparisons using the Bonferroni correction, overall statistical significance threshold was set at  $\alpha$  = 0.05 (3 comparisons at p<0.0167). **D.** Mice were injected as in A. (3 mice/grp). 24hrs after the last injection of virus (d22), or earlier if tumor size excised 1.0cm diameter in the PBS groups, tumors were excised and assayed for IL-12 or TNF- $\alpha$  by ELISA (normalized by protein concentration in whole tumor cell lysates and expressed as pg/ml protein). Representative of two separate experiments. Each symbol in B,D represents a mouse (n=2 or 3/group). Means  $\pm$  SD are shown. ND, not detected (below limit of detection). P-values were determined using a two-way ANOVA (B) with a Tukey multiple comparisons post-test. Statistical significance set at p < 0.05 for B. Source data are provided as a Source Data file.

## **Supplementary Figure 3**



**Supplementary Figure 3:** C57Bl/6 mice with 10d established s.c. B16 tumors (7 mice/grp) were treated **A.** i.v. with 2.5x10<sup>6</sup> OT-1 T cells activated *in vitro* for 5d with IL-2 (50 IU rhIL-2/ml) and SIINFEKL peptide (1µg/ml) (d10), and with anti-PD-1 antibody i.p. (300µg/injection) (d17,19,21); or with 2.5x10<sup>6</sup> CD8+ T cells recovered from the spleens of mice which had rejected B16 tumors treated with VSV-CSDE1<sup>P5S</sup> (**Fig.5F**) and activated *in vitro* for 5d with IL-2 and the CSDE1<sup>P5S</sup> mutated peptide MSFD<u>S</u>NLLH (1µg/ml) (d10) either with, or without, anti-PD-1 antibody (d17,19,21). **B.** 3 additional groups were treated intratumorally with VSV-IFN-ß (d10,12,14) (5x10<sup>7</sup> pfu/injection) and subsequently with either CD8+ T cells from naïve C57Bl/6 mice (d17); *in vitro* activated OT-I CD8+ T cells; or *in vitro* activated anti-CSDE1<sup>P5S</sup> CD8+ T cells. Survival with time is shown. Data (A, B) from a single experiment not repeated. P-values were determined using the Log-rank Mantel Cox test. Source data are provided as a Source Data file.

#### **Supplementary References**

- 1 Huff, A. L. *et al.* APOBEC3 Mediates Resistance to Oncolytic Viral Therapy. *Mol Ther Oncolytics* **11**, 1-13, doi:10.1016/j.omto.2018.003 (2018).
- 2 Evgin, L. *et al.* Suboptimal T-cell Therapy Drives a Tumor Cell Mutator Phenotype That Promotes Escape from First-Line Treatment. *Cancer Immunol Res* **7**, 828-840, doi:10.1158/2326-6066.CIR-18-0013 (2019).