

## Expanded View Figures

**Figure EV1. Peripheral T cell compartment characterization of *CD4cre Ptpn6<sup>f/f</sup>* and *CD4cre Ptpn6<sup>f/f</sup>/Ptpn11<sup>f/f</sup>* mice.**

- A, B T cells from spleen (A) and inguinal lymph nodes (iLN) (B) of *CD4cre Ptpn6<sup>f/f</sup>* and control mice were analyzed. Cellularity, CD4<sup>+</sup> and CD8<sup>+</sup> T cell numbers (gated as TCR $\beta^+$  CD4<sup>+</sup> or CD8<sup>+</sup> respectively), percentages of CD44 high T cells are shown; regulatory T cell (Treg) numbers and frequency (gated as TCR $\beta^+$  CD4<sup>+</sup> Foxp3<sup>+</sup>) are illustrated for the spleen.
- C Cellularity from thymus and number of CD4 and CD8 double-negative, double-positive, CD4 or CD8 single-positive thymocytes (gated on lineage<sup>-</sup>; single-positives, an additional gate on TCR $\beta^+$  was performed) of *CD4cre Ptpn6<sup>f/f</sup>/Ptpn11<sup>f/fWbm</sup>* and control mice are depicted.
- D Treg number and frequency (gated as TCR $\beta^+$  CD4<sup>+</sup> Foxp3<sup>+</sup>) are illustrated for the spleen of *CD4cre Ptpn6<sup>f/f</sup>/Ptpn11<sup>f/fWbm</sup>* and control mice.
- E Cellularity, CD4<sup>+</sup> and CD8<sup>+</sup> T cell numbers, as well as percentages of CD44 high T cells from the iLN of *CD4cre Ptpn6<sup>f/f</sup>/Ptpn11<sup>f/fWbm</sup>* and control mice are shown.
- F Survival curves of *CD4cre Ptpn6<sup>f/f</sup>/Ptpn11<sup>f/fBgn</sup>* and control mice challenged with MC38.
- G–I Ten to twelve days following MC38 tumor inoculation, isotype-treated *CD4cre Ptpn6/11<sup>wt/wt</sup>* and *CD4cre Ptpn6<sup>f/f</sup>/Ptpn11<sup>f/fWbm</sup>* mice were sacrificed. Graph depicts the frequencies of PD-1<sup>+</sup> CD8<sup>+</sup> T cells (gated as CD45<sup>+</sup> TCR $\beta^+$  CD8<sup>+</sup>) in the tumor (G). Frequencies of CD4<sup>+</sup> T cells (gated as CD45<sup>+</sup> TCR $\beta^+$  CD4<sup>+</sup>) expressing IFN- $\gamma$  and TNF $\alpha$  upon re-stimulation (H) and percentages of CD4<sup>+</sup> T cells in the tumor are shown (I).

Data information: Results illustrate mean  $\pm$  SEM of  $n = 7$ –8 mice/group (A, B), of  $n = 10$ –12 mice/group (C–E), or of  $n = 9$ –14 mice/group (G–I). Student's t-test (unpaired, two-tailed) was used to compare differences between experimental groups (A–E, G–I). Results depict  $n = 11$  mice/group; statistical significance was calculated by log-rank (Mantel-Cox) test (F). \* $P \leq 0.05$ , \*\* $P \leq 0.01$ , \*\*\* $P \leq 0.001$ , \*\*\*\* $P \leq 0.0001$ .

Source data are available online for this figure.

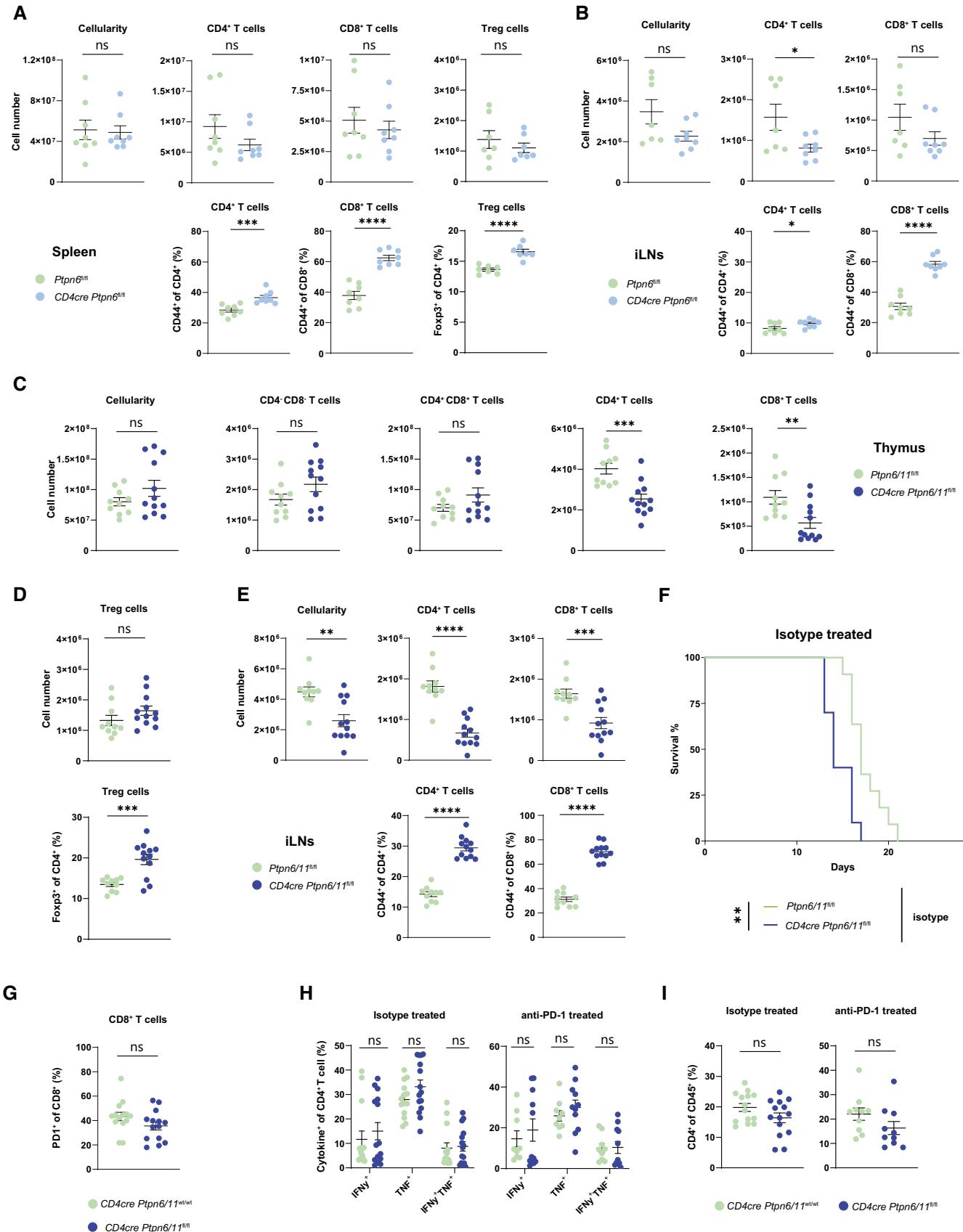
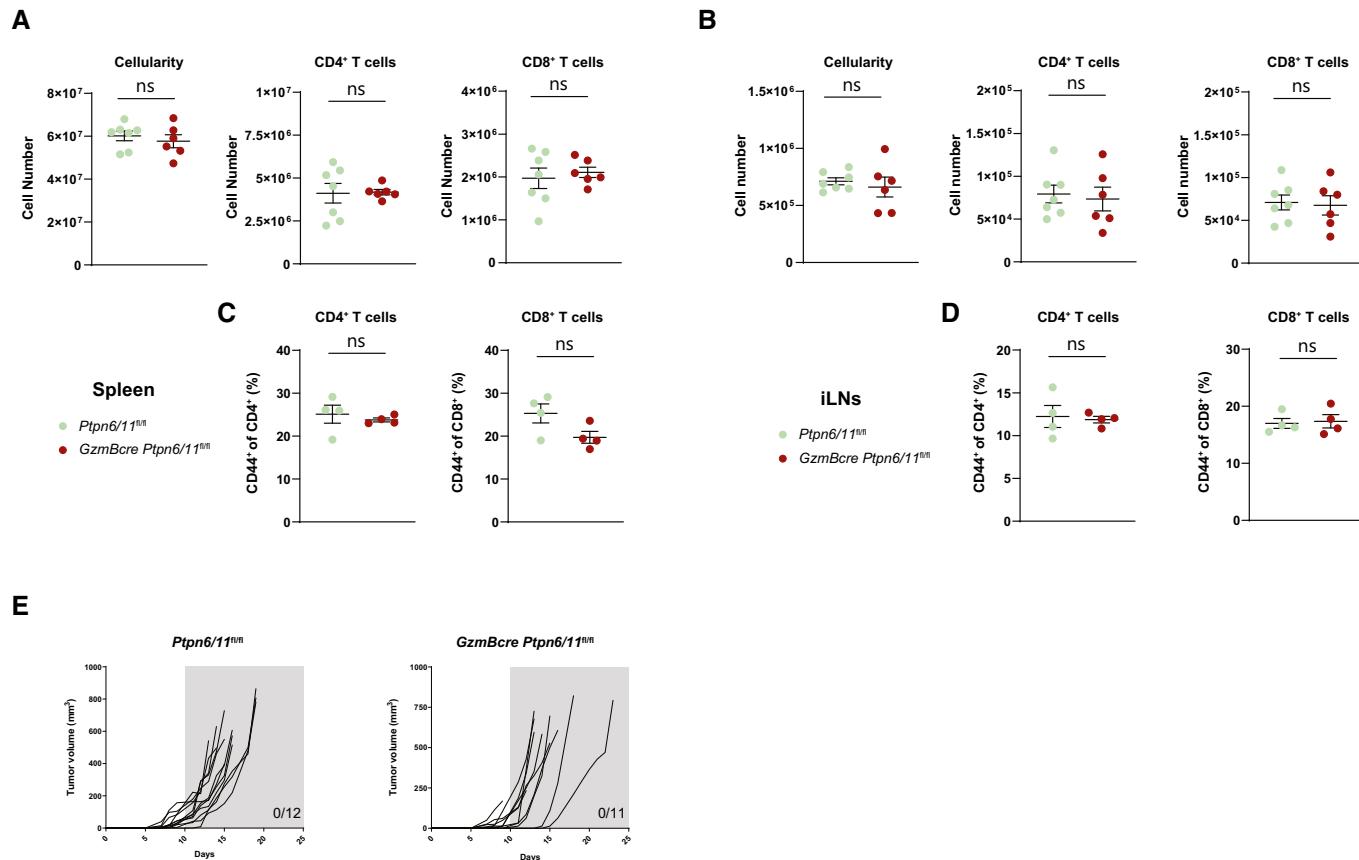


Figure EV1.

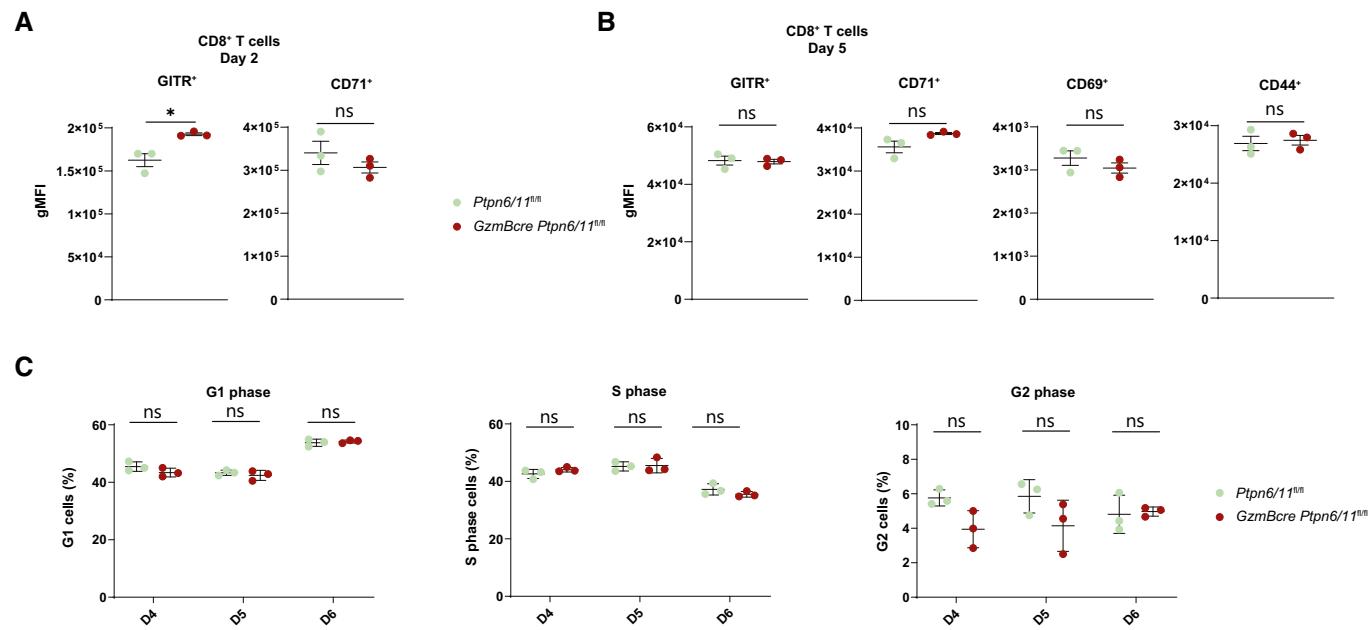


**Figure EV2. Peripheral T cell compartment characterization of *GzmBcre Ptprn6/11<sup>fl/fl</sup>* mice.**

**A–D** Spleen and iLN of *GzmBcre Ptprn6/11<sup>fl/fl</sup>* mice and control mice were analyzed. Cellularity, CD4<sup>+</sup> and CD8<sup>+</sup> T cell numbers are depicted for spleen (A) and iLN (B). Percentages of CD44 high of CD4<sup>+</sup> and CD8<sup>+</sup> T cells are shown for spleen (C) and iLN (D). Results depict  $n = 6\text{--}7$  mice/group (A, B) and  $n = 4$  mice/group for percentages of CD44 high cells (C, D). Student's t-test (unpaired, two-tailed) was used to compare differences between experimental groups.

**E** Tumor growth in individual mice challenged with MC38 is shown for the indicated genotypes; number of mice eradicating the tumor is shown within the graphs.

Source data are available online for this figure.



**Figure EV3. Characterization of CTLs from *Ptpn6/11<sup>fl/fl</sup>* and *GzmBcre Ptpn6/11<sup>fl/fl</sup>* mice.**

CD8<sup>+</sup> cytotoxic lymphocytes were generated from splenocytes of the indicated mice by anti-CD3, anti-CD28, and IL-12 stimulation and maintained in IL-2.

A, B CD8<sup>+</sup> T cells were analyzed by flow cytometry for surface expression of GITR and CD71 after 2 days (A) and for surface expression of GITR, CD71, CD69, and CD44 after 5 days (B); a quantification of these parameters is shown in the graphs (A and B).

C Graphs depict the percentages of cells in G1, S, and G2 phase as measured by flow cytometry-based cell cycle analysis of *Ptpn6/11<sup>fl/fl</sup>* and *GzmBcre Ptpn6/11<sup>fl/fl</sup>* CTLs (gated on CD8<sup>+</sup> T cells) at the indicated days.

Data information: Results depict mean  $\pm$  SD of  $n = 3$  biological replicates and Student's *t*-test (unpaired, two-tailed) was used to compare differences between experimental groups (A–C). \* $P \leq 0.05$ .

Source data are available online for this figure.