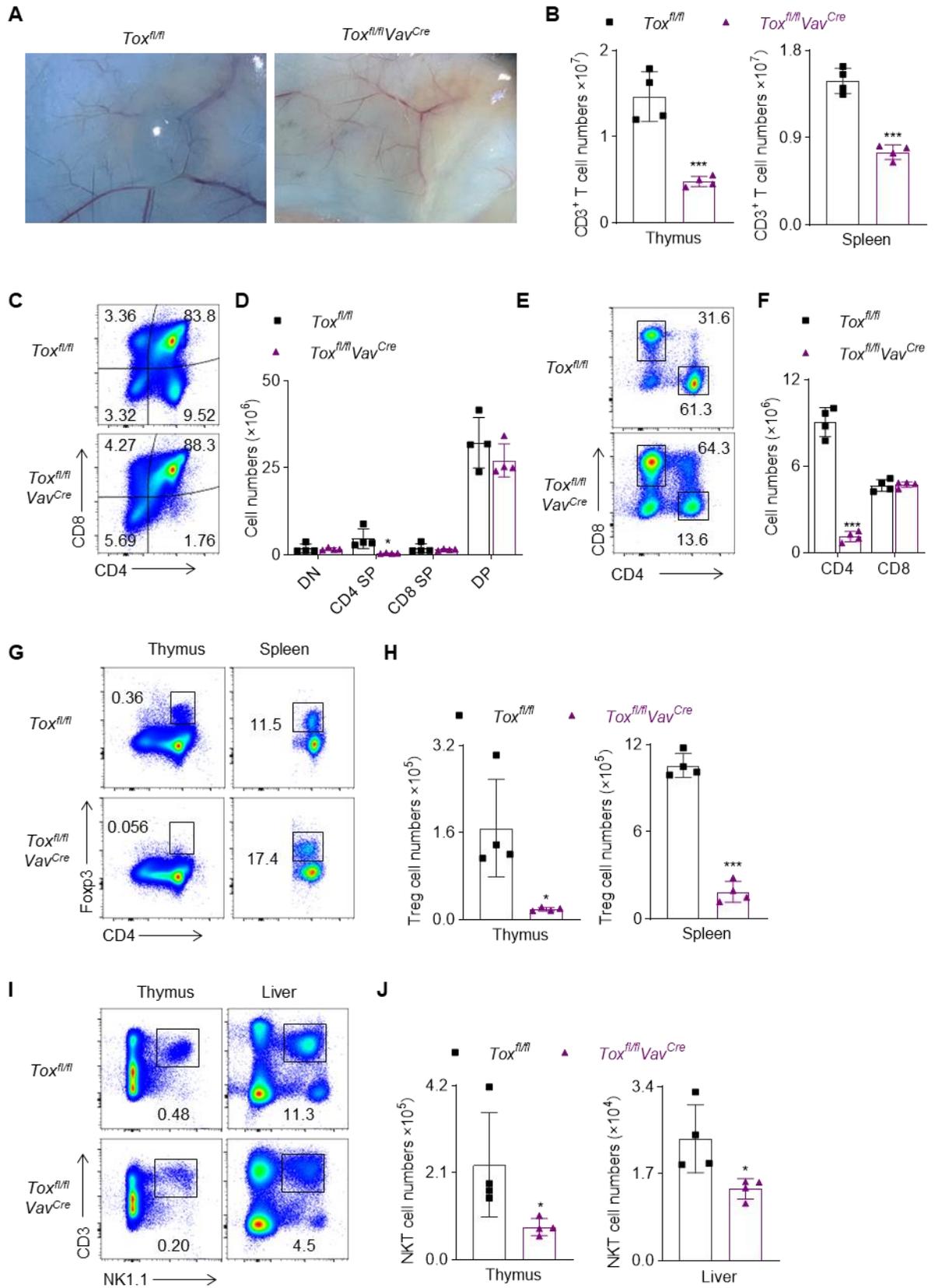
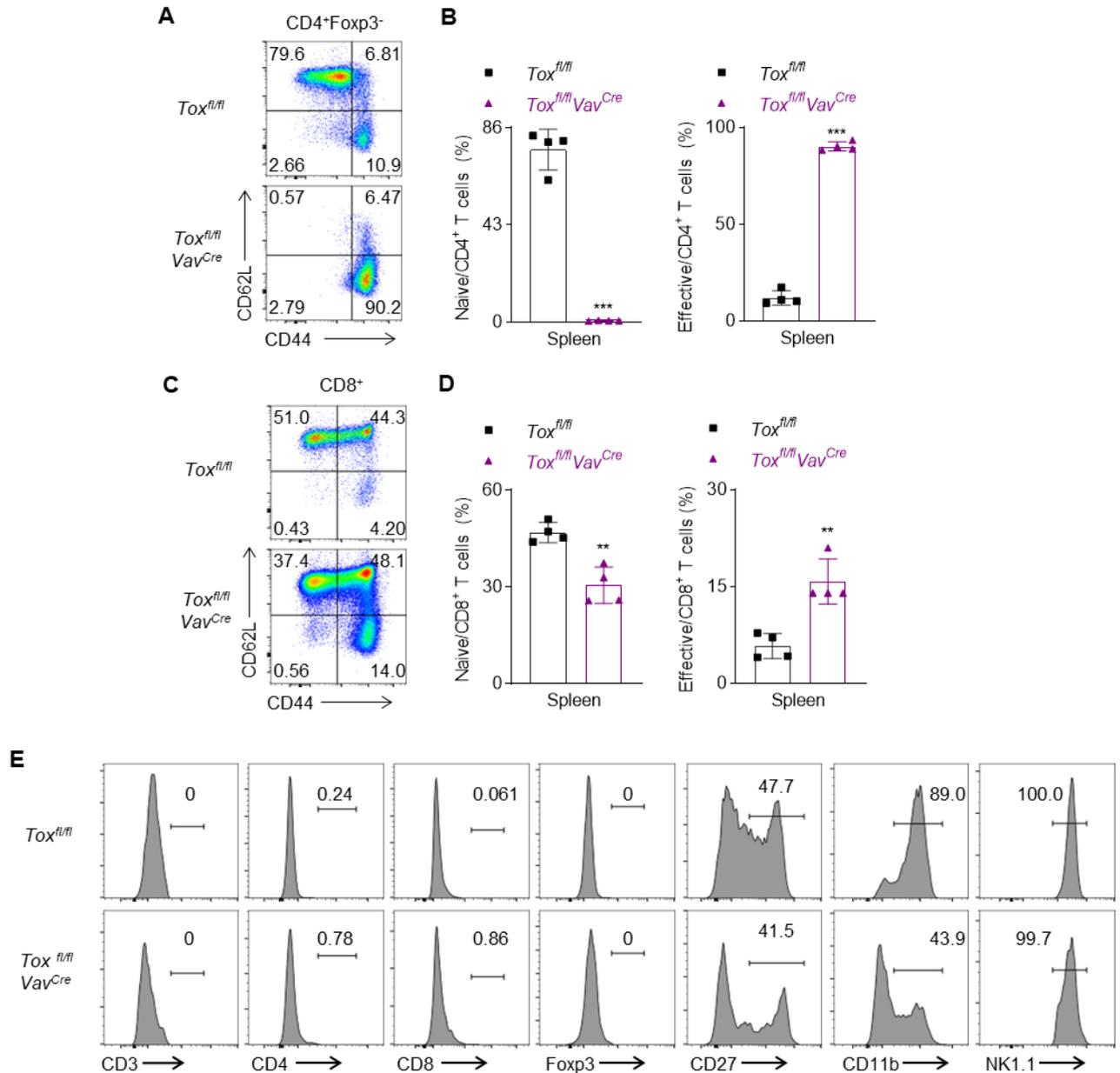


**Figure S1. Detecting TOX deletion efficiency in NK cells from the three conditional targeting mouse models. (A)** Schematic graphs showing the breeding strategies for the three conditional knockout mice. **(B)** Analysis of *Tox* mRNA expression in HSC, CLP and NK cells from BM of *Tox<sup>fl/fl</sup>* and *Tox<sup>fl/fl</sup>Vav<sup>Cre</sup>* mice (left), in NKp, iNK and mNK cells from spleen of *Tox<sup>fl/fl</sup>* and *Tox<sup>fl/fl</sup>CD122<sup>Cre</sup>* mice (middle), in NK cells from spleen and BM of the *Tox<sup>fl/fl</sup>* and *Tox<sup>fl/fl</sup>Ncr1<sup>Cre</sup>* mice (right) by quantitative PCR. All above results were normalized to *Gapdh* (n = 4).



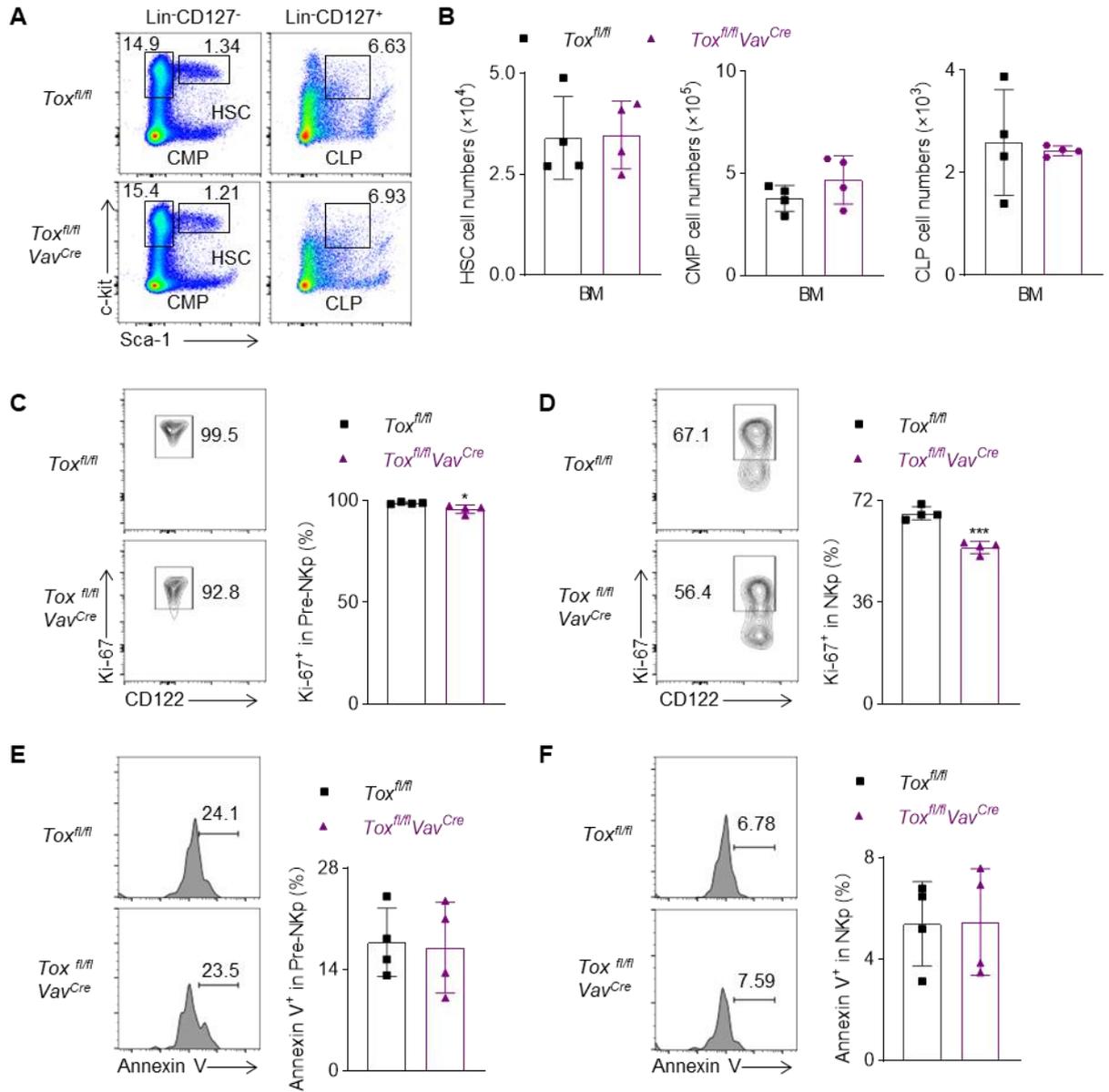
**Figure S2. The deletion of TOX at HSC stage impairs the development of CD4<sup>+</sup> T cells.** (A) Gross anatomy of lymph nodes visualized by the injection of Chicago sky

blue dye into  $Tox^{fl/fl}$  and  $Tox^{fl/fl}Vav^{Cre}$  mice. **(B)** The number of  $CD3^+$  T cells in thymus and spleen from  $Tox^{fl/fl}$  and  $Tox^{fl/fl}Vav^{Cre}$  mice (n = 4). **(C, D)** Representative flow cytometry plot **(C)** and the number **(D)** of different thymocyte T cell subsets in  $Tox^{fl/fl}$  and  $Tox^{fl/fl}Vav^{Cre}$  mice (n = 4). **(E, F)** Representative flow cytometry plot **(E)** and the number **(F)** of  $CD4^+$  T and  $CD8^+$  T cells in spleen from  $Tox^{fl/fl}$  and  $Tox^{fl/fl}Vav^{Cre}$  mice (n = 4). **(G, H)** Representative flow cytometry plot **(G)** and the absolute number **(H)** of Treg cells in thymus and spleen of  $Tox^{fl/fl}$  and  $Tox^{fl/fl}Vav^{Cre}$  mice (n = 4). **(I, J)** Representative flow cytometry plot **(I)** and the absolute number **(J)** of NKT cells in thymus and liver of  $Tox^{fl/fl}$  and  $Tox^{fl/fl}Vav^{Cre}$  mice (n = 4). Data are representative of three independent experiments.

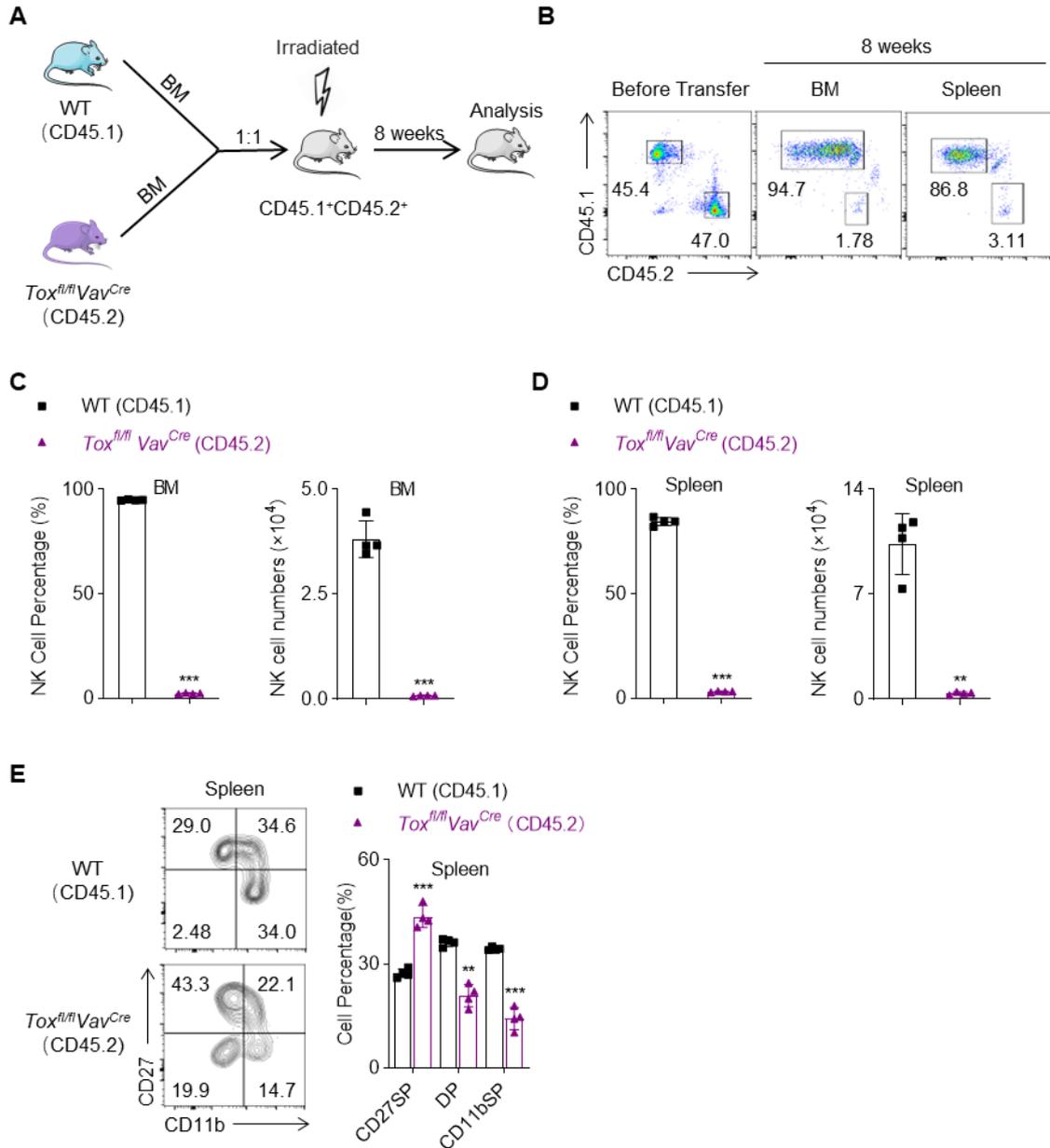


**Figure S3. Deletion of TOX at HSC stage influences the activated status of T cells.**

(A-D) Representative flow cytometry plot (A, C) and the percentage (B, D) of naïve T cells and effective CD4<sup>+</sup> (A, B) or CD8<sup>+</sup> (C, D) T cells in spleen of *Tox<sup>fl/fl</sup>* and *Tox<sup>fl/fl</sup> Vav<sup>Cre</sup>* mice (n = 4). (E) Comparison of the expression level of CD3, CD4, CD8, Foxp3, CD27, CD11b and NK1.1 in NK cells from spleen of *Tox<sup>fl/fl</sup>* and *Tox<sup>fl/fl</sup> Vav<sup>Cre</sup>* mice. Data are representative of at least two independent experiments.

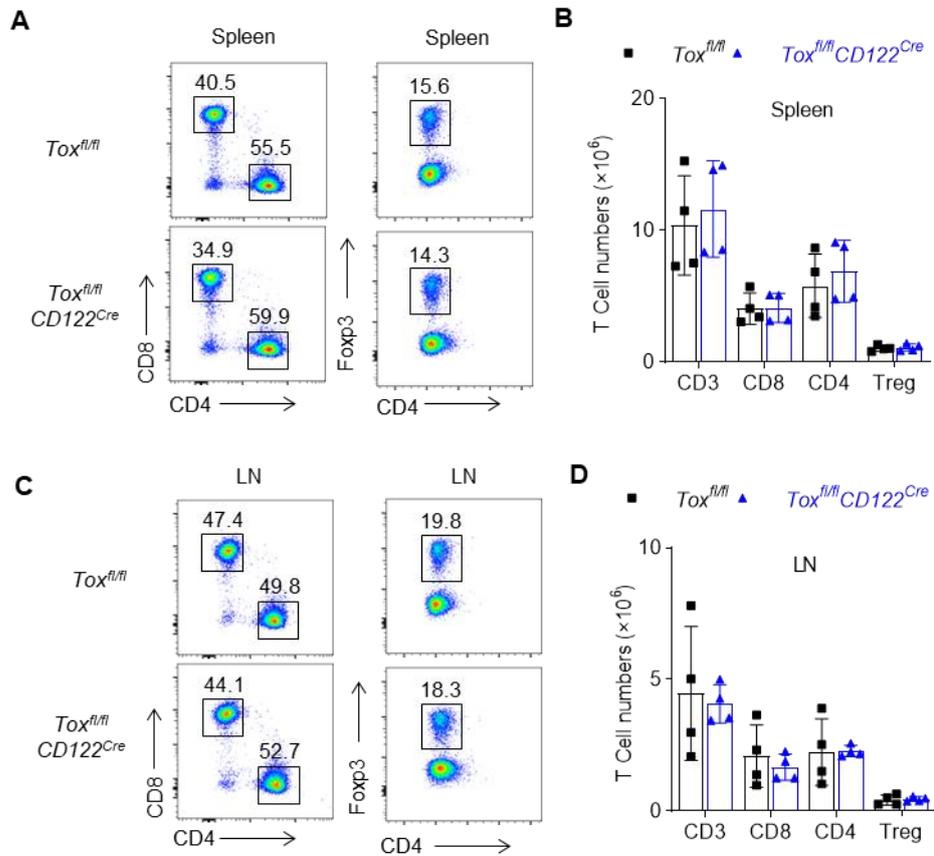


**Figure S4. TOX deletion at NKp stage impairs the proliferation capacities of Pre-NKp and NKp cells.** (A, B) Representative flow cytometry plot (A) and the absolute numbers (B) of HSC, CMP, and CLP in BM of *Tox<sup>fl/fl</sup>* and *Tox<sup>fl/fl</sup>Vav<sup>Cre</sup>* mice (n = 4). (C, D) Comparison the expression of Ki-67 in pre-NKp cells (C) and NKp cells (D) between *Tox<sup>fl/fl</sup>Vav<sup>Cre</sup>* mice and *Tox<sup>fl/fl</sup>* mice (n = 4). (E, F) Comparison the expression of Annexin V in pre-NKp cells (E) and NKp cells (F) between *Tox<sup>fl/fl</sup>Vav<sup>Cre</sup>* mice and *Tox<sup>fl/fl</sup>* mice (n = 4). Data are representative of two independent experiments.



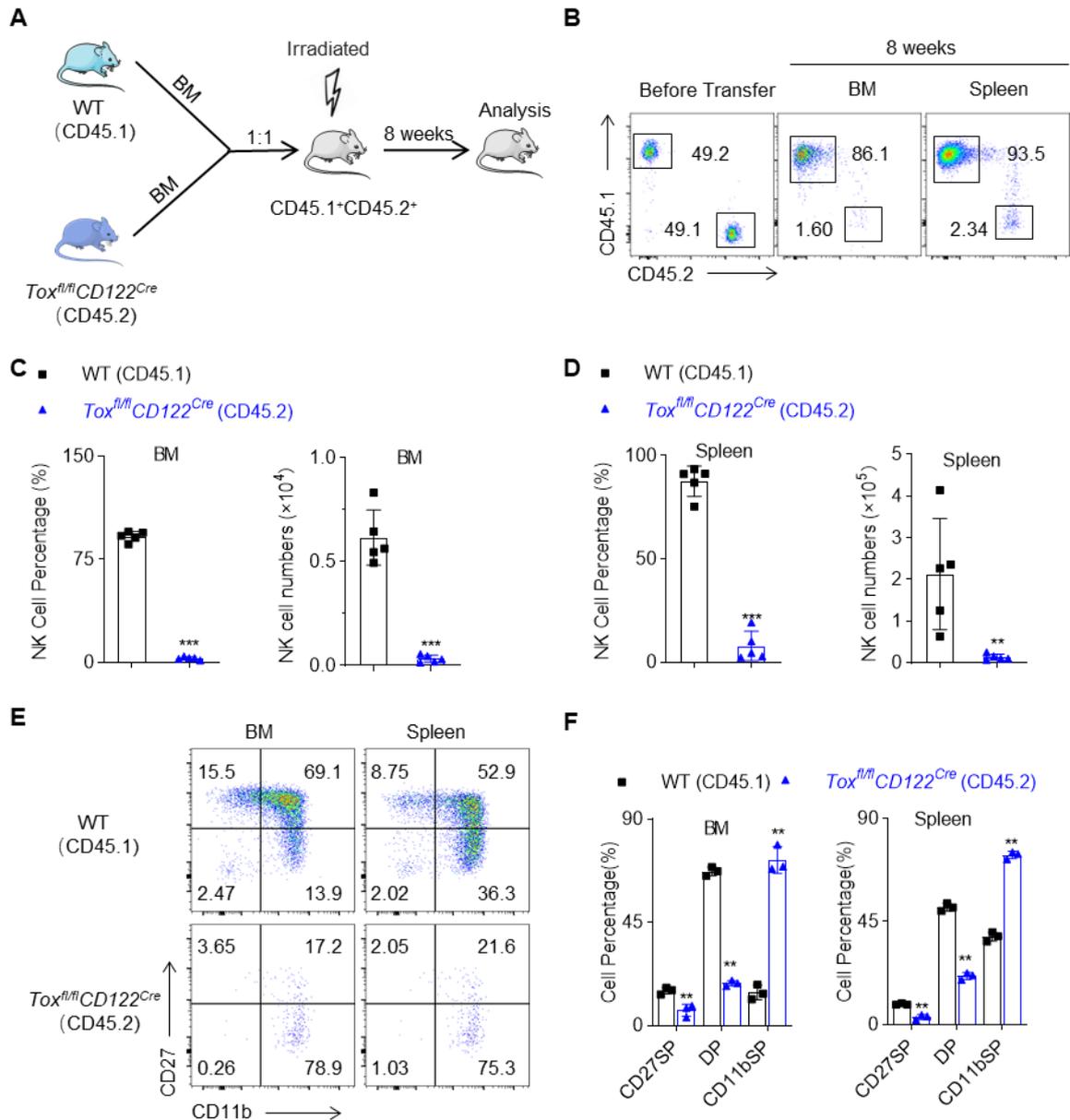
**Figure S5. TOX participates in NK cell commitment in a cell-intrinsic manner at HSC stage.** (A) Experimental design of congenic BM chimera assay. (B) Representative flow cytometry plots showing the percentage of NK cells from *Tox<sup>fl/fl</sup>Vav<sup>Cre</sup>* mice (CD45.2<sup>+</sup>) and WT mice (CD45.1<sup>+</sup>) in spleen and BM of recipient mice 8 weeks after congenic BM chimera assay. The injected cells are shown. (C, D) The percentage (left) and the number (right) of NK cells from *Tox<sup>fl/fl</sup>Vav<sup>Cre</sup>* mice (CD45.2<sup>+</sup>) and WT mice (CD45.1<sup>+</sup>) in BM (C) or spleen (D) of recipient mice 8 weeks after congenic BM chimera assay (n = 4). (E) Representative histogram and NK cell subset distribution from *Tox<sup>fl/fl</sup>Vav<sup>Cre</sup>* mice (CD45.2<sup>+</sup>) and WT mice (CD45.1<sup>+</sup>) in spleen

of recipient mice 8 weeks after congenic BM chimera assay (n = 4).



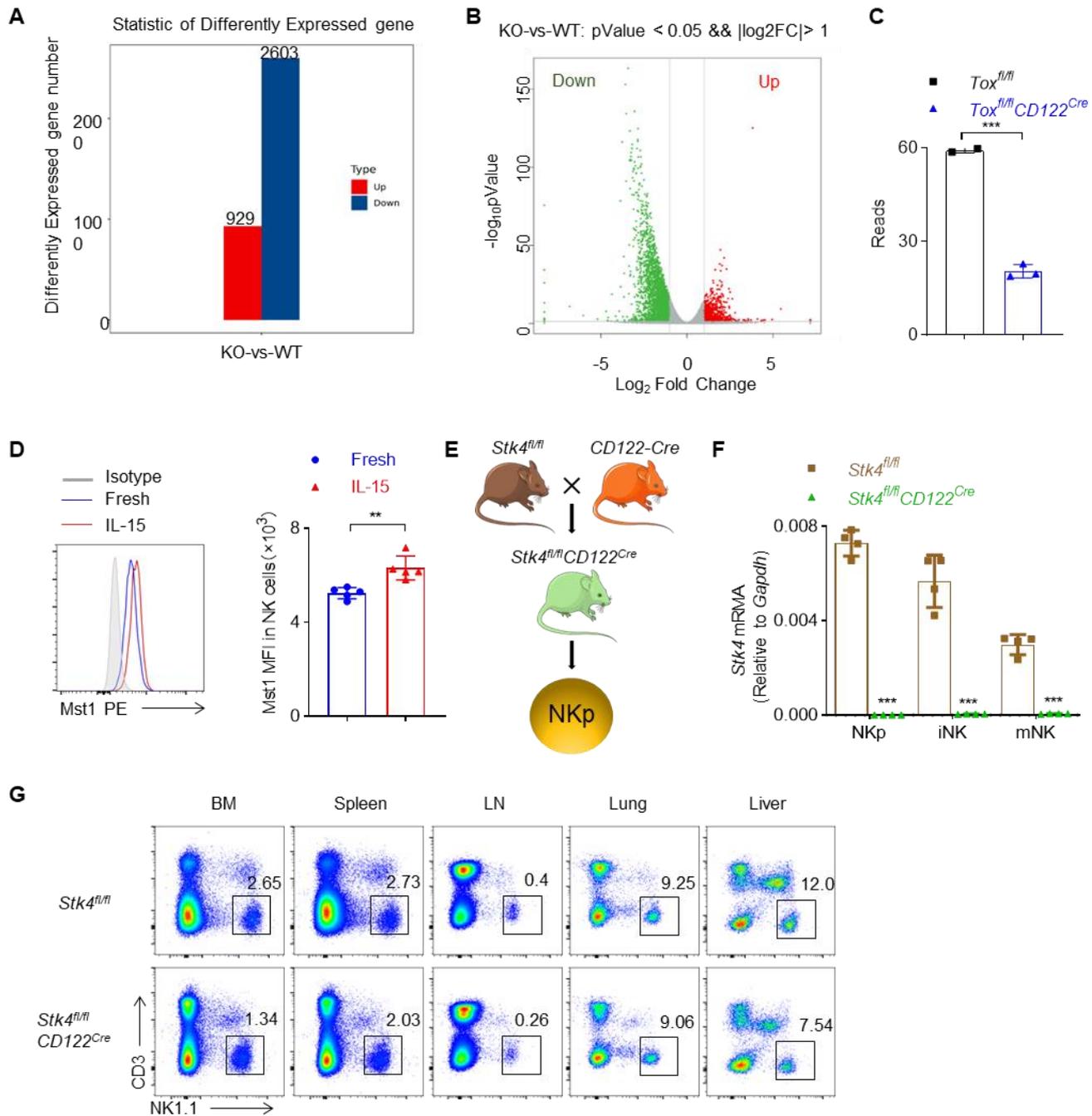
**Figure S6. The ablation of TOX at NKp stage doesn't affect T cell development.**

Representative flow cytometry plot (A, C) and the absolute numbers (B, D) of CD3<sup>+</sup> T cells, CD8<sup>+</sup> T cells, CD4<sup>+</sup> T cells and Treg cells in spleen (A, B) or LN (C, D) of *Tox<sup>fl/fl</sup>* and *Tox<sup>fl/fl</sup> CD122<sup>Cre</sup>* mice (n = 4). Data are representative of two independent experiments.



**Figure S7. Genetic ablation TOX at NKp stage leading to severe defects in NK cell development is cell-intrinsic. (A)** Experimental design of congenic BM chimera assay. **(B)** Representative flow cytometry plot showing the percentage of NK cells from *Tox<sup>fl/fl</sup>CD122<sup>Cre</sup>* mice (CD45.2<sup>+</sup>) and WT mice (CD45.1<sup>+</sup>) in spleen and BM of recipient mice 8 weeks after congenic BM chimera assay. The injected cells are shown. **(C, D)** The percentage (left) and the number (right) of NK cells from *Tox<sup>fl/fl</sup>CD122<sup>Cre</sup>* mice (CD45.2<sup>+</sup>) and WT mice (CD45.1<sup>+</sup>) in BM **(C)** or spleen **(D)** of recipient mice 8 weeks after congenic BM chimera assay (n = 5). **(E, F)** Representative flow cytometry plot **(E)** and the analysis of NK cell subset distribution **(F)** from *Tox<sup>fl/fl</sup>CD122<sup>Cre</sup>* mice (CD45.2<sup>+</sup>) and WT mice (CD45.1<sup>+</sup>) in BM and spleen of recipient mice 8 weeks after congenic

BM chimera assay (n = 3).



**Figure S8. TOX orchestrates NK cell homeostasis by controlling Mst1 expression.**

(A, B) The histogram (A) and the volcano plot (B) showing differentially expressed genes from RNA-seq. (C) The expression level of *Stk4* between WT NK cells and TOX-deficient NK cells in our RNA-seq dataset. (D) Comparison the expression level of Mst1 in NK cells from spleen of WT mice before and after stimulation with 10 ng/mL IL-15 for 16 h (n = 4). (E) Schematic graph showing the strategy for generating the *Stk4*<sup>fl/fl</sup> *CD122*<sup>Cre</sup> mice. (F) Analysis of *Stk4* mRNA expression in NKp, iNK and mNK

cells in spleen of *Stk4<sup>fl/fl</sup>* and *Stk4<sup>fl/fl</sup>CD122<sup>Cre</sup>* mice by quantitative PCR. **(G)** Representative flow cytometry plots showing the percentage of NK cells in BM, spleen, LN, lung and liver of *Stk4<sup>fl/fl</sup>* and *Stk4<sup>fl/fl</sup>CD122<sup>Cre</sup>* mice.