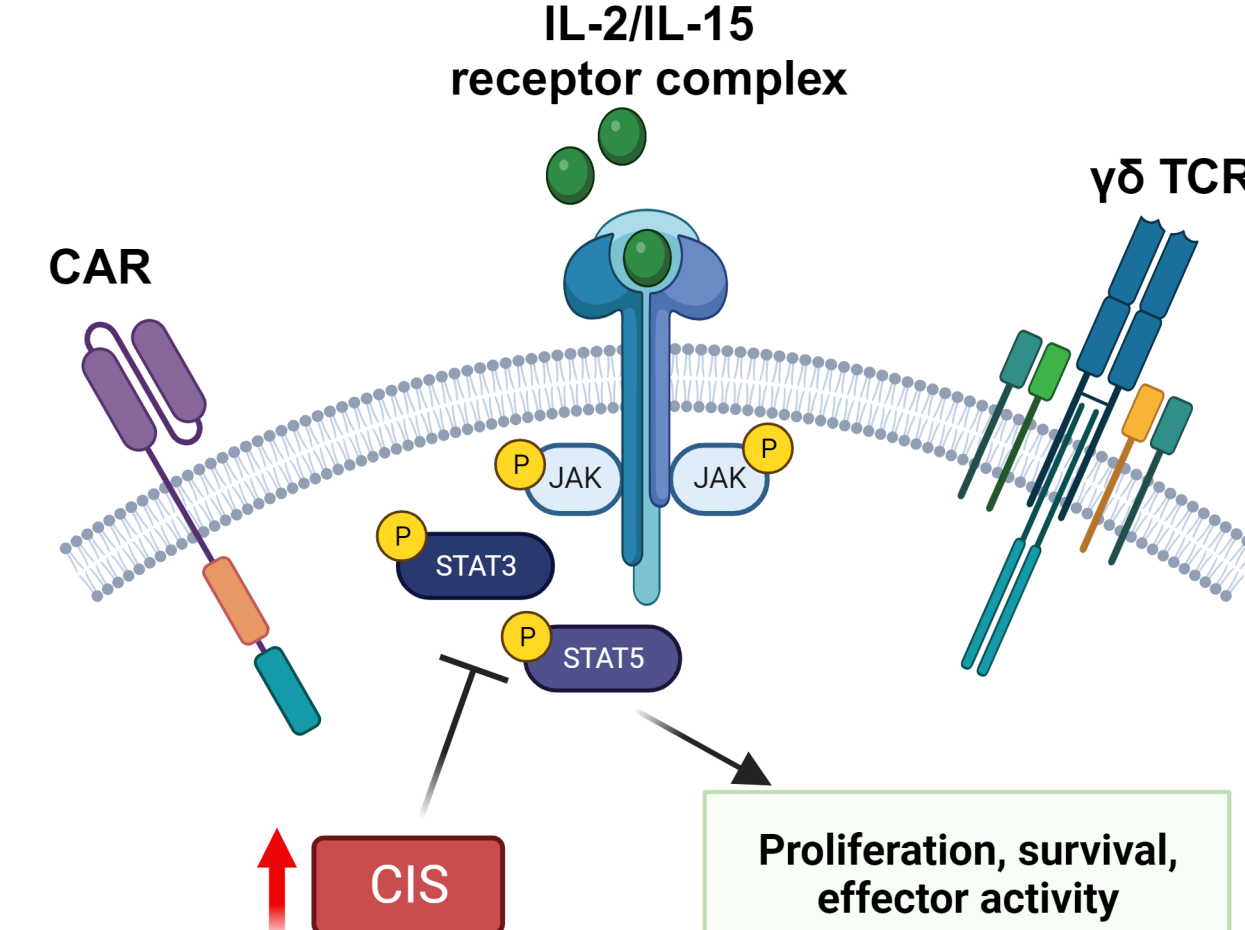


INTRODUCTION

$\gamma\delta$ T cells have emerged as a promising allogeneic cell-based immunotherapy platform for the treatment of cancer. Armed with a unique combination of innate and adaptive immunity to target malignant cells, the infiltration and presence of $\gamma\delta$ T cells in various solid tumors is significantly correlated with survival benefit. Additional modifications to engineer $\gamma\delta$ T cells to express CARs have shown enhanced tumoricidal activity and encouraging clinical efficacy^{1,2}. Opportunities to understand the mechanisms that drive these potent anti-tumor responses in $\gamma\delta$ T cells are integral to further augmentation of their effector functions. Cytokine inducible SH2-containing protein (CIS) is a key negative regulator of IL-2/15 and TCR signaling. CIS disruption in $\alpha\beta$ T cells and NK cells promotes increased persistence and anti-tumor activity^{3,4}, but the role of CIS in $\gamma\delta$ T cells is not well understood. Here, we report, for the first time, a strategy that couples targeted disruption of the CIS loci within allogeneic CAR engineered $\gamma\delta$ T cells for enhanced anti-tumor potential.



CIS up-regulation by cytokine/TCR/CAR stimulation

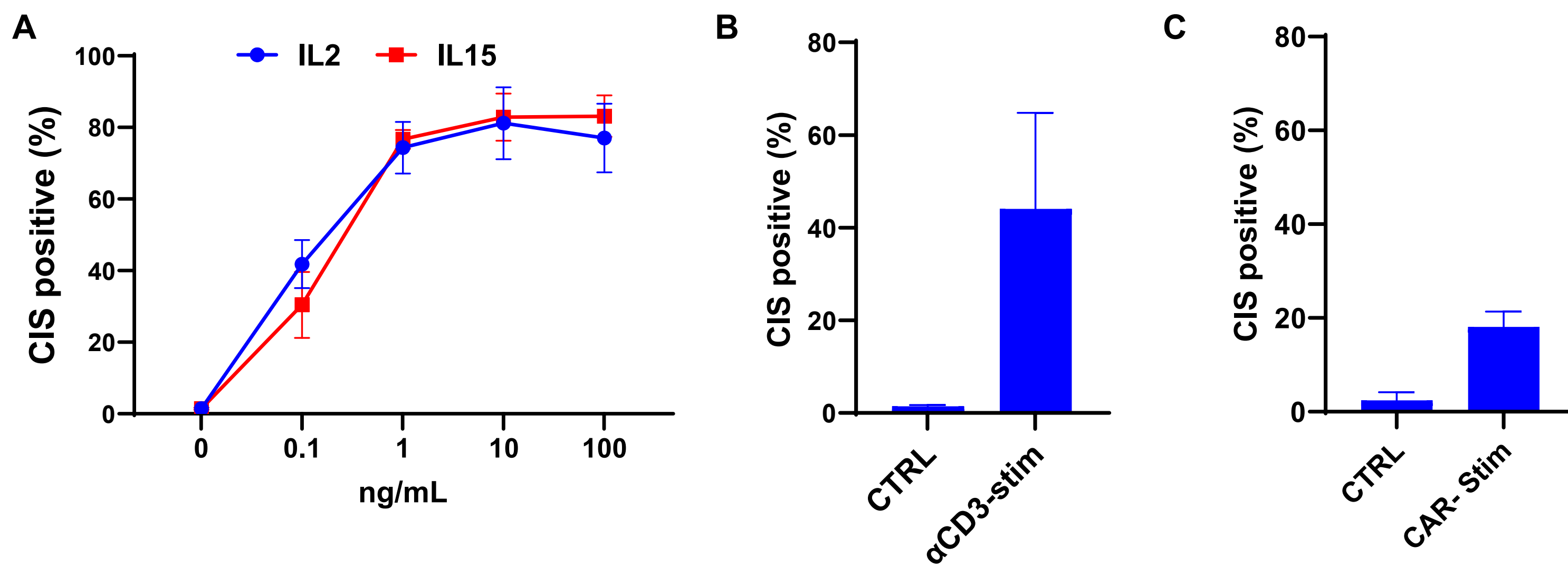


Figure 1. (A) CIS up-regulation by IL2/IL15 stimulation. V δ 1 T cells were cultured overnight in cytokine free medium, treated with IL2 or IL15 at the indicated concentrations for five hours, then CIS expression was analyzed using flow cytometry. (B) CIS up-regulation through TCR stimulation. Cytokine starved V δ 1 cells were stimulated with plate bound CD3 antibody (OKT3) for five hours, then CIS expression was analyzed using flow cytometry. (C) Cytokine starved CAR V δ 1 cells were stimulated by tumor antigen for five hours and CIS expression was analyzed using flow cytometry.

Efficient Generation of *CISH* gene knockout CAR V δ 1 T cells

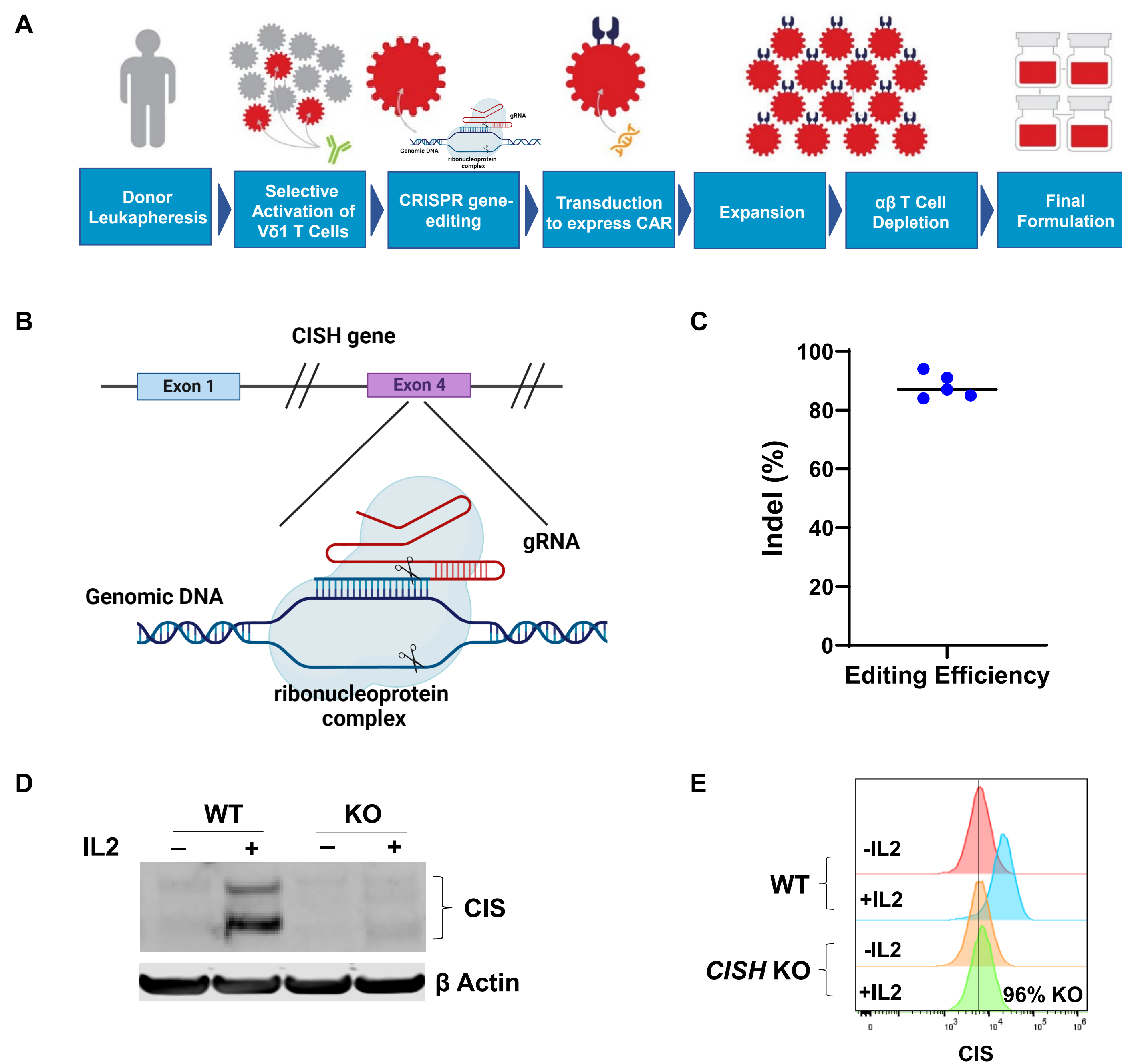


Figure 2. (A) Flow chart highlighting the key steps in generation of *CISH* gene knockout CAR V δ 1 T cells. (B) Scheme of CRISPR mediated *CISH* KO using guide RNAs (gRNA) located in exon 4 of the *CISH* gene. (C) Indel frequencies of the *CISH* gene knockout in V δ 1 cells were analyzed by Inference of CRISPR Edits (ICE) analysis tool. (D) WT or *CISH* gene edited CAR transduced V δ 1 T cells were incubated overnight in cytokine free medium and then stimulated with or without IL2 for five hours, and CIS expression was analyzed by immunoblotting. β actin was used as loading control. (E) Representative flow cytometry plot of CIS expression on WT or *CISH* knockout CAR V δ 1 T cells stimulated with or without IL2.

Robust CAR expression and a less differentiated T cell memory phenotype with minimal activation/exhaustion-associated markers in *CISH* KO V δ 1 T cells

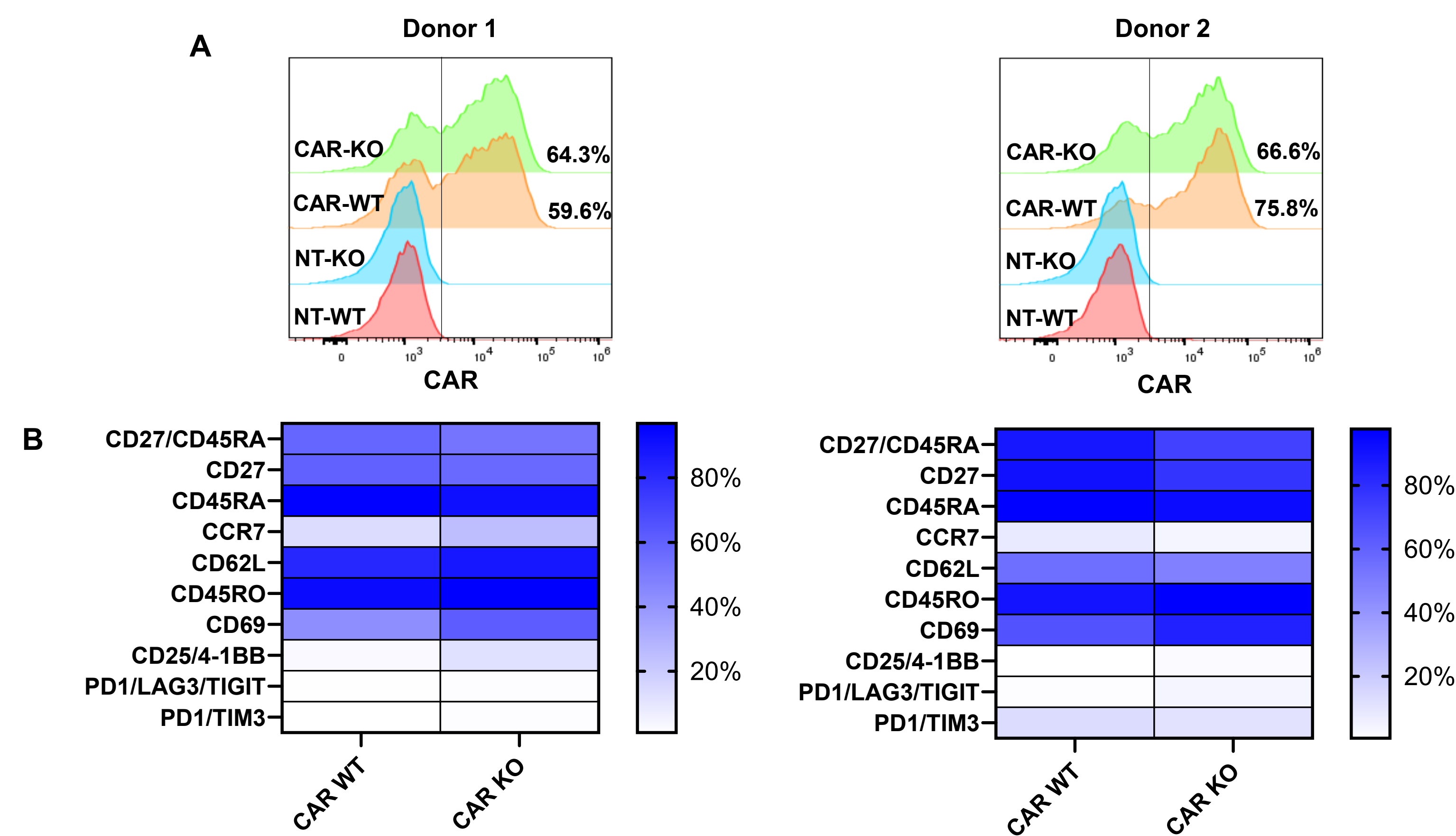


Figure 3. (A) Representative flow cytometry plot of anti-B7-H6 CAR expression on WT or *CISH* knockout CAR and non-transduced (NT) V δ 1 T cells from two different donors. (B) Heatmap showing percentages of T cell memory, activation, and exhaustion associated markers expressed by WT or *CISH* KO CAR V δ 1 T cells.

CISH KO in V δ 1 T cells promoted increased expression of genes and gene-enrichment sets associated with T cell effector function

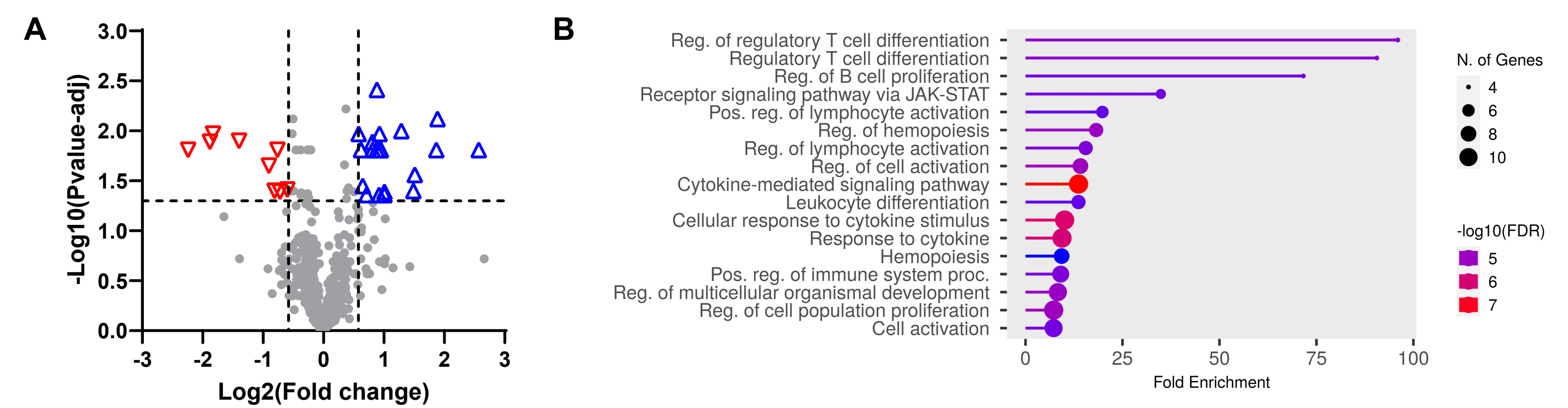


Figure 4. (A) Volcano plot of differentially expressed genes (DEGs) between WT and *CISH* KO CAR V δ 1 T cells. Blue triangles Δ represent DEGs upregulated and red triangles ∇ represent DEGs downregulated in *CISH* KO vs WT conditions. Gene expression was quantitated using the Nanostring nCounter[®] CAR T Cell Characterization panel. (B) Gene Ontology analysis was performed using ShinyGO 0.77 (<http://bioinformatics.sdstate.edu/go/>) to identify the biological pathways associated with DEGs that were upregulated in the *CISH* KO CAR V δ 1 T cells.

Enhanced phosphorylation of STAT3/5 and proliferation in *CISH* KO V δ 1 cells

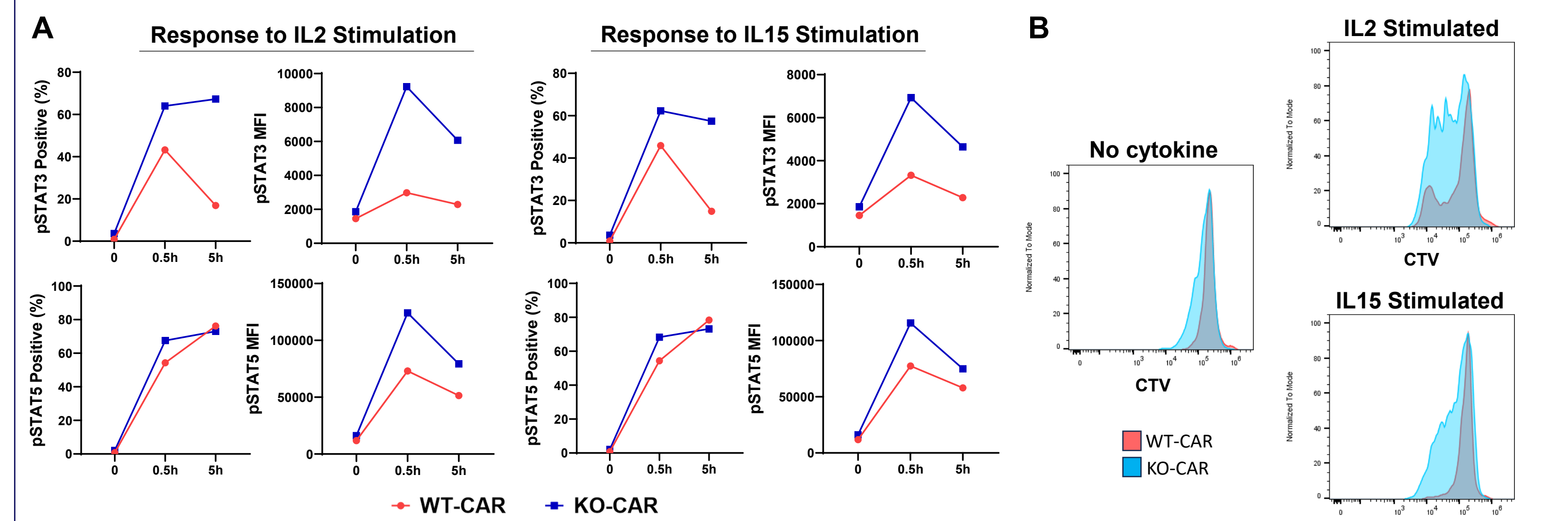


Figure 5. (A) Enhanced phosphorylation of STAT3/5 signaling in *CISH* KO V δ 1 T cells. WT or *CISH* KO CAR V δ 1 T cells were cultured overnight in cytokine free medium, then treated with IL2 or IL15 at the indicated times, and percentage of pSTAT3 and pSTAT5 expression (left) or mean fluorescence intensity (MFI, right) was analyzed using flow cytometry. (B) Improved proliferation of *CISH* KO compared to WT CAR V δ 1 T cells. WT or *CISH* KO CAR V δ 1 T cells were labeled with cell trace violet (CTV) and cultured for seven days in cytokine free or low levels of IL2/IL15. Representative FACS plots are shown.

CISH KO CAR V δ 1 T cells demonstrated enhanced serial killing and persistence against tumor cell lines with diverse antigen expression levels

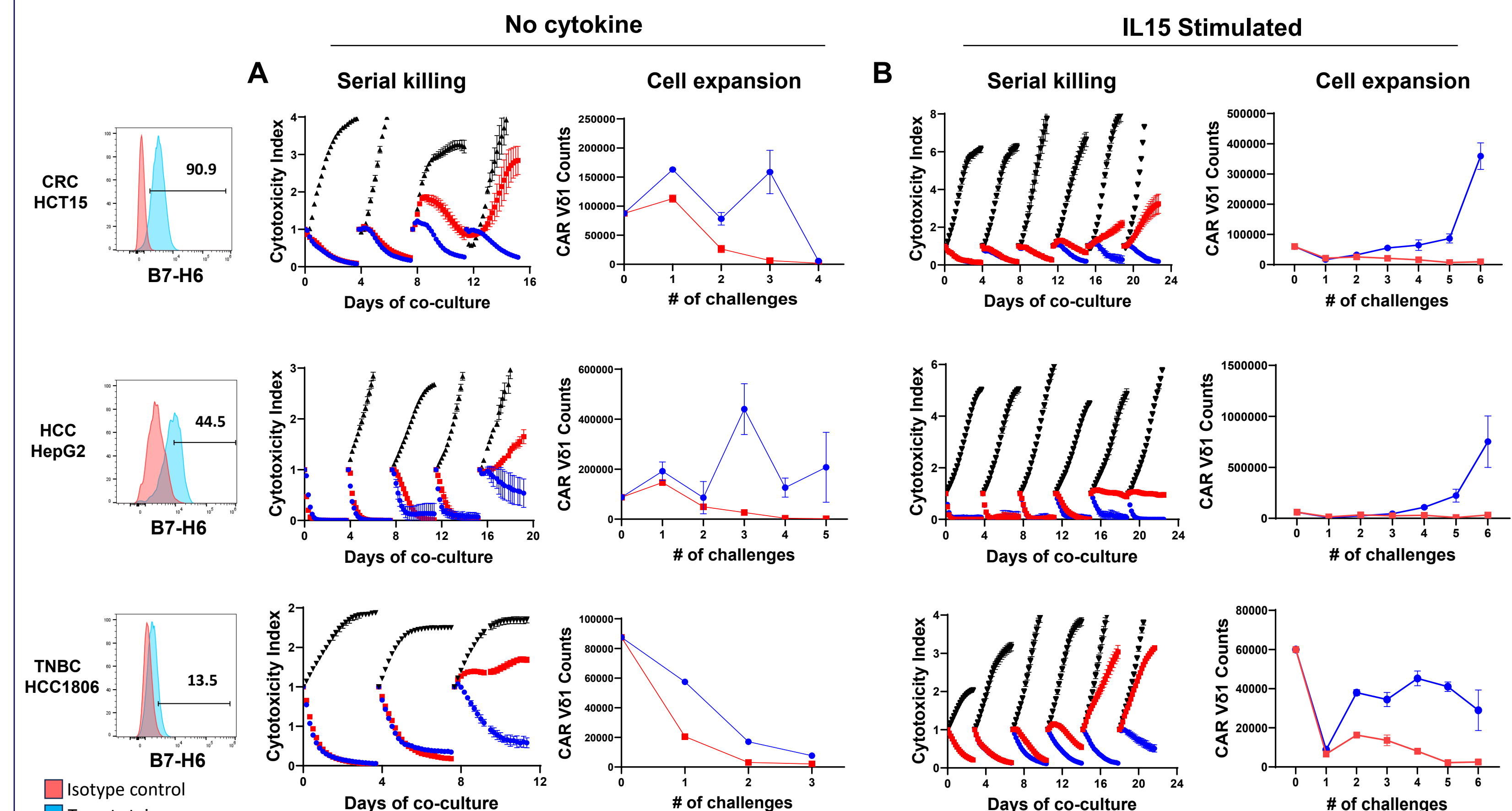


Figure 6. Cytotoxic potentials of WT CAR V δ 1 T cells (red square) and *CISH* KO CAR V δ 1 T cells (blue circle) were evaluated against tumor cell lines (black triangle) with diverse antigen expression levels: HCT15 (high), HepG2 (medium) and HCC1806 (low) in the repetitive stimulation Incubate Immune Cell Killing Assay in cytokine free medium (A) or IL15 (B). CAR V δ 1 T cells were cocultured with target cell line for three to four days, then half of the cells were transferred into a new vessel with freshly plated target cells for an additional three to four days. The Cytotoxicity Index was calculated by dividing the total NIR object area ($\mu\text{m}^2/\text{well}$) of all time points by the value at time of tumor challenge. Leftover cells in the previous plates were collected and analyzed by flow cytometry to monitor the CAR positive V δ 1 T cell expansion.

Demonstration of One Step Knock-out Knock-in (KOKI) of CAR through *CISH* locus in V δ 1 T cells

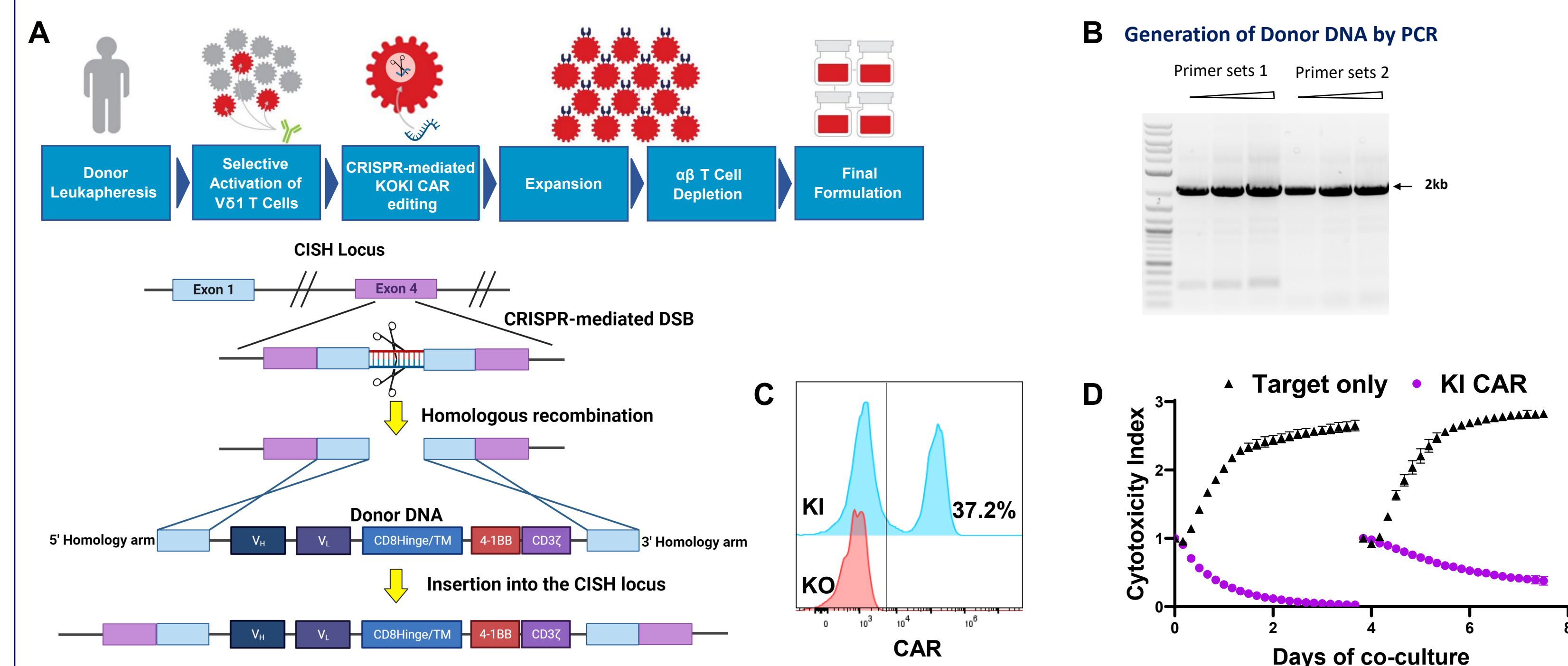


Figure 7. (A) Flow chart describing the process of generation of knockout knock-in CAR through the *CISH* locus in V δ 1 T cells. (B) Use of PCR primer sets on the homologous arms to generate donor DNA for electroporation. (C) Representative flow cytometry plot of CAR expression on *CISH* knockout CAR knock-in V δ 1 T cells. (D) Cytotoxic potential of *CISH* knockout knock-in CAR V δ 1 T cells (purple circles) were evaluated against B7-H6 positive tumor cell line HCT-15 in a repetitive stimulation Incubate Immune Cell Killing Assay. The Cytotoxicity Index was calculated by dividing the total NIR object area ($\mu\text{m}^2/\text{well}$) of all time points by the value at time of tumor challenge.

Summary & Conclusions

- CRISPR-gene editing can be successfully applied to CAR V δ 1 T cells with a high KO efficiency
- CISH* KO in CAR V δ 1 T cells enhanced pSTAT3/5 signaling, increased proliferation, and promoted robust *in vitro* anti-tumor activity
- Transcripts associated with T cell effector activity were upregulated in *CISH* KO CAR in V δ 1 T cells
- One step knock-out knock-in experiments supports PoC for non-viral vector CAR insertion and expression in V δ 1 T cells
- These approaches provide opportunities to explore novel gene-editing strategies to further enhance allogeneic CAR $\gamma\delta$ T cells as a promising platform for cancer immunotherapy.

References

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