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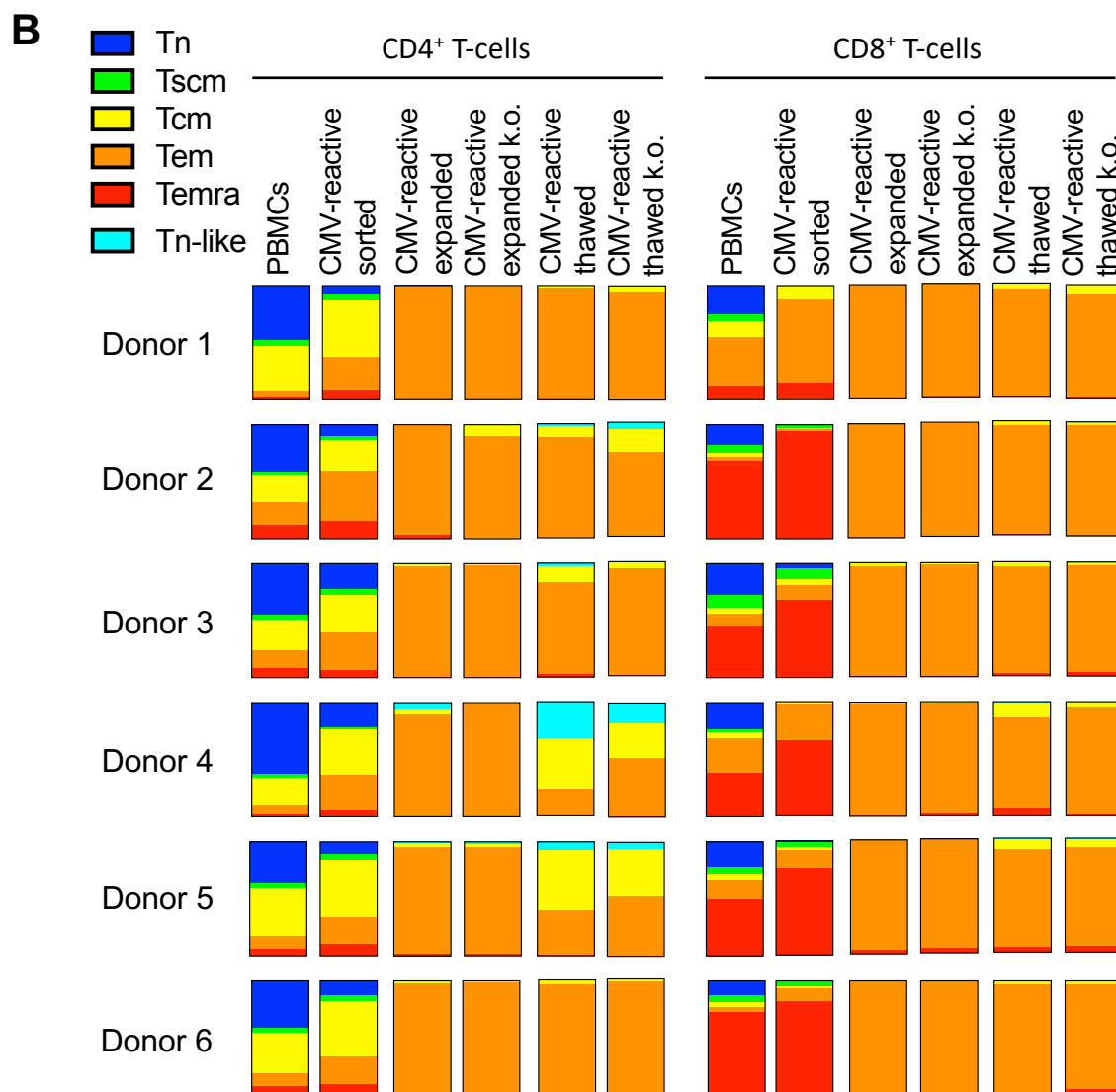
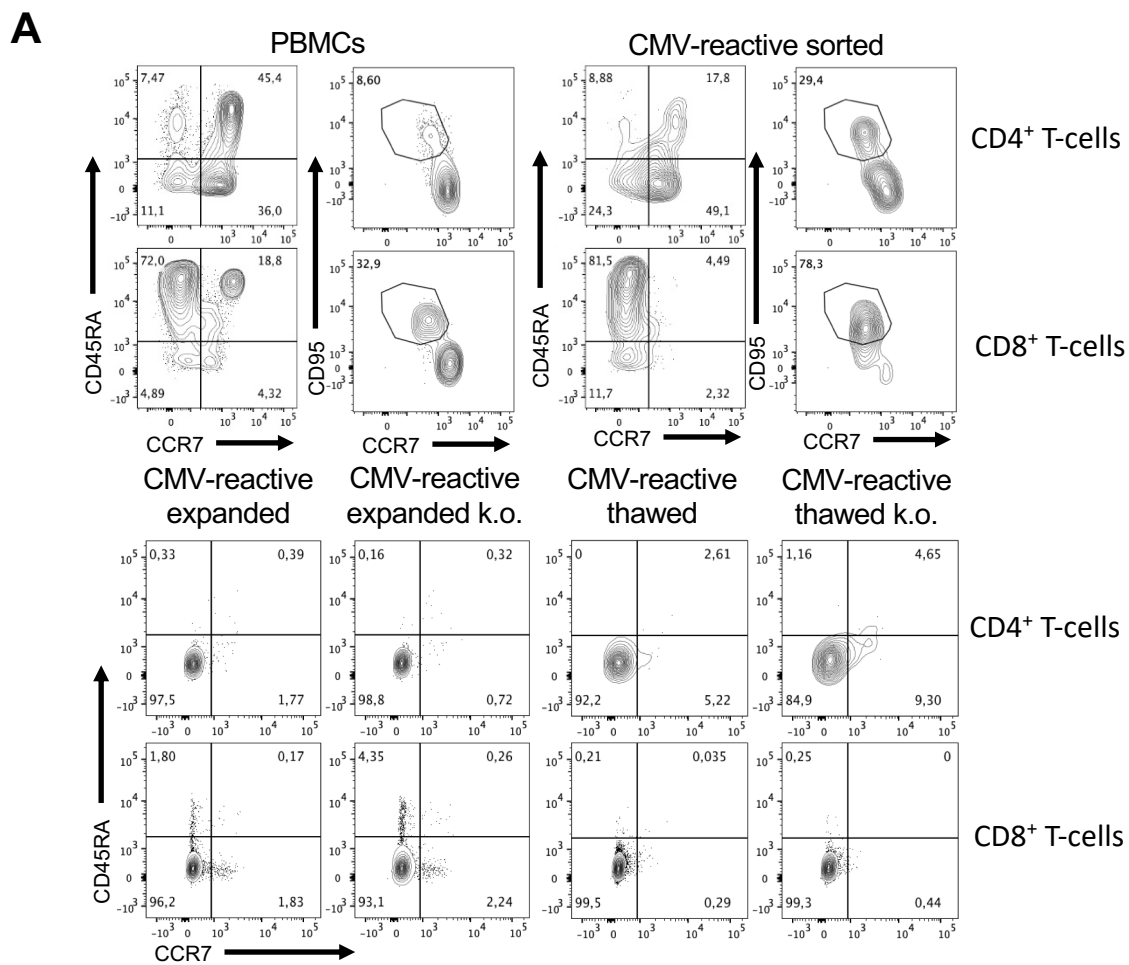
Supplemental Information

CRISPR-Cas9-Edited Tacrolimus-Resistant

Antiviral T Cells for Advanced Adoptive

Immunotherapy in Transplant Recipients

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Supplementary Figure 1: T-cell memory subsets in the starting material, sorted CMV-specific T-cells and the expanded and thawed T-cell products.

(A) Representative flow cytometric dot plots illustrating T-cell memory subsets in the different populations indicated above the plots.

(B) Distribution of CD4⁺ (left panel) and CD8⁺ (right panel) T-cell memory subsets in the different populations indicated for each donor: T_N = naïve T-cells (CCR7⁺, CD45RA⁺, CD95⁻), T_{SCM} = stem cell memory T-cells (CCR7⁺, CD45RA⁺, CD95⁺), T_{CM} = central memory T-cells (CCR7⁺ CD45RA⁻), T_{EM} = effector memory T-cells (CCR7⁻ CD45RA⁻), T_{EMRA} = CD45RA-expressing effector memory T-cells (CCR7⁻ CD45RA⁺), T_{N-like} = expanded naïve-like T-cells (CCR7⁺ CD45RA⁺).