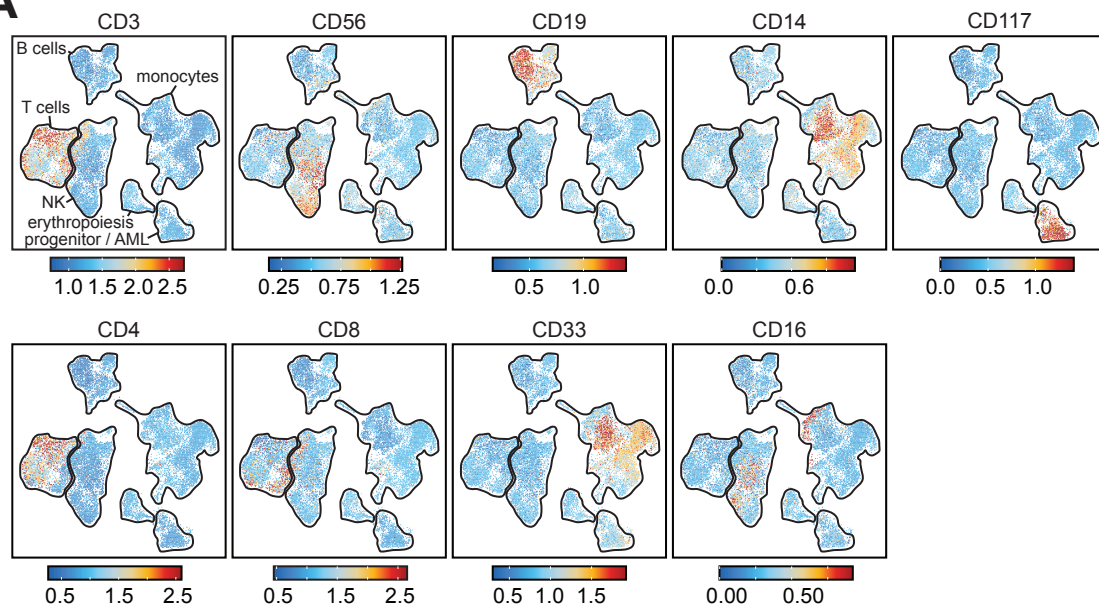
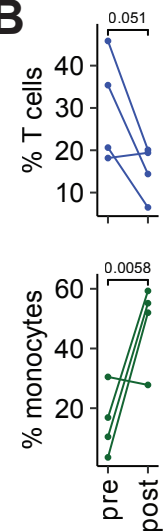
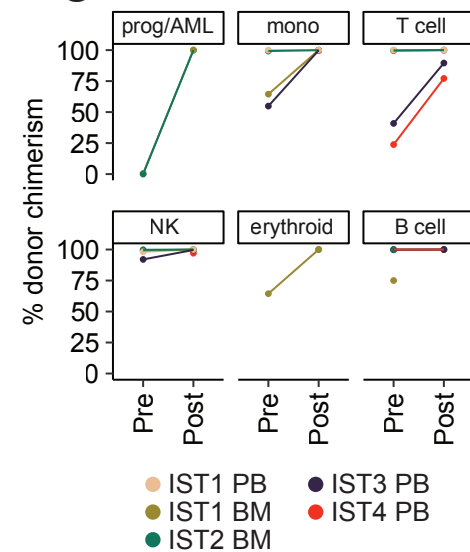
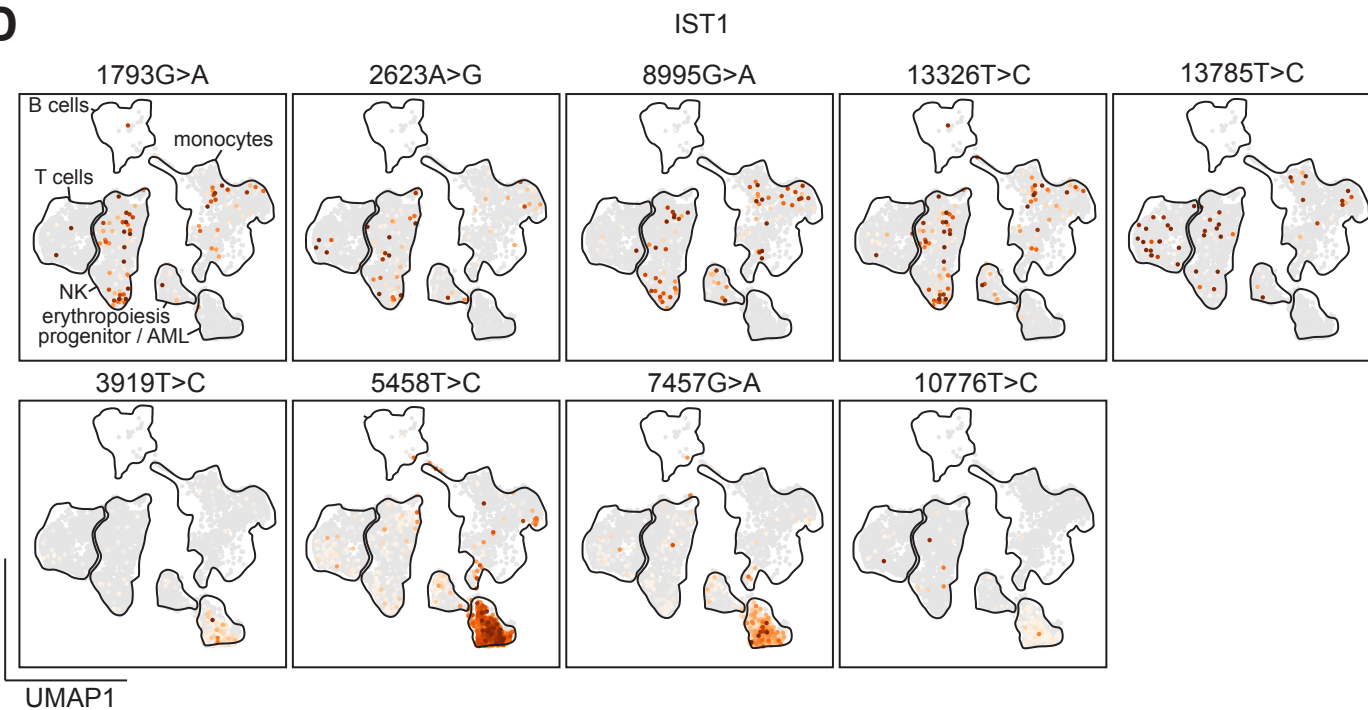
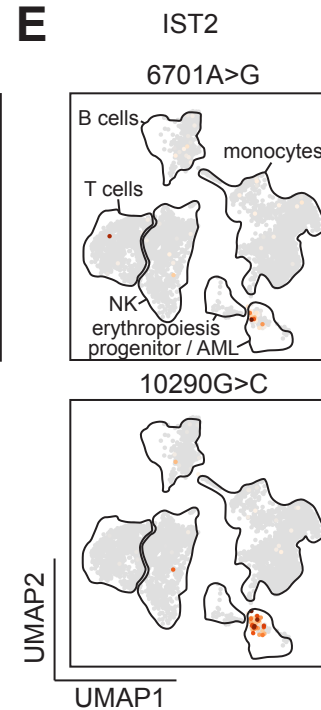


**A****B****C****D****E**

Suppl. Fig. 14. Identification of cell types and tracking of mitochondrial DNA mutations in immunosuppression tapering (IST) cohort using ASAP-seq. A UMAP projection of single cell chromatin accessibility profiles annotated by surface marker expression of CD3, CD4 and CD8 (T cells), CD56 (NK cells), and CD14, CD16, CD33 and CD117 (myeloid populations) measured using ASAP-seq and Total-seq B antibodies. B Percentage of T cells and monocytes in patients IST1-4 before (pre) and after (post) IST. C Donor chimerism across progenitor/AML, monocytic (mono), T cell, NK, erythroid and B cell populations before (Pre) and after (Post) IST. D-E UMAP projection of single cell chromatin accessibility profiles annotated by detection of mitochondrial DNA (mtDNA) mutations in IST1 (D) and IST2 (E).