

Description of Additional Supplementary Files

Supplementary Data 1. Alternative splicing events (exon skipping) associated with *Tnpo3*-deficiency at a False Discovery Rate (FDR) threshold of 1% (Sheet 1, CD4⁺ single-positive T cells; sheet 2, CD4⁺CD8⁺ double-positive T cells). Chromosome, strand, and skipped exon start and end denote the coordinates of the spliced-out exon. FDR denotes the false discovery rate of the differential alternative splicing event. Inclusion level denotes the mean exon inclusion level for either wild-type or mutant. Δ Inclusion level denotes the difference in inclusion level between wild-type and mutant.