



**Supplemental Figure 10: Transcriptional profiling of CD8+ FR-B TALs: A)** Normalized expression of key effector and checkpoint genes measured by RNAseq for CD8+ FR-B 2/7 and 2/15 TALs (n=2-3/group) **B)** Hierarchical clustering of differentially expressed genes between CD8+ FR-B 2/7 and 2/15 TALs (n=2-3/group).