



Supplemental Figure 2: Single cell RNAseq analysis of CCR8⁺ and CCR8⁻ subpopulations of tumor CD4⁺FOXP3⁺ T cells

Single cell RNAseq analysis of CD4⁺FOXP3⁺ T cells examining effector and naïve T cell populations in relationship to CCR8 (*GZMA* vs *CCR8*, and *CCR7* vs *CCR8*) (A). (B) CCR8⁻ cells from CD4⁺FOXP3⁺ population showing mutual exclusivity between *GZMA* and *CCR7*, and the presence of naïve T_{regs} (CCR7⁺SELL⁺).