

Fig. S2

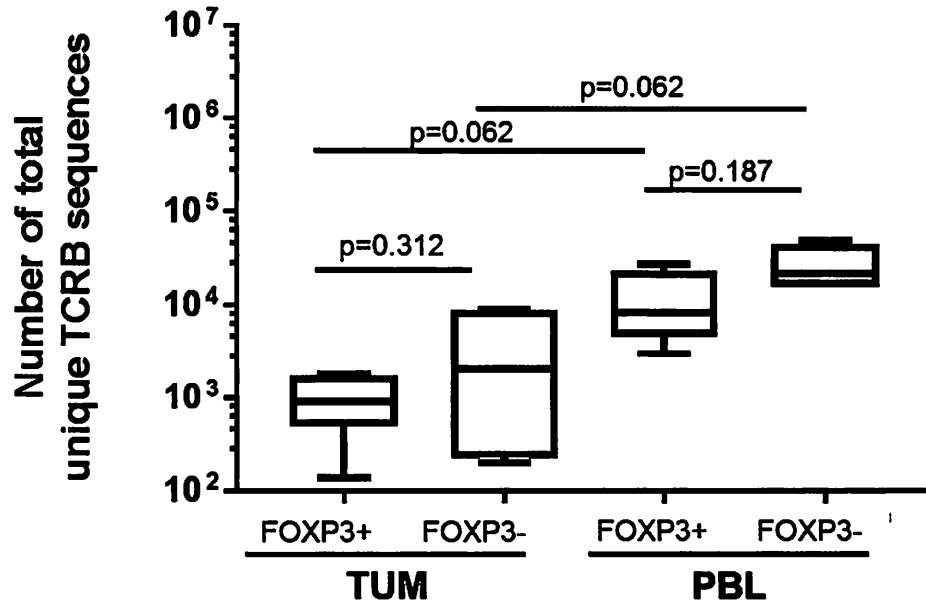


Fig. S2. Number of total unique TCRB reads for FOXP3⁺ and FOXP3⁻ subsets sorted from tumor or PBL.

TCRB immunosequencing was performed on each sorted FOXP3⁺ and FOXP3⁻ CD4 T cell subsets from tumor (n=6) and PBL (n=5). The total number of unique TCRB sequences are shown for each subset.