

Fig. S3. Clonality of 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). CThe total number of productive TCRB sequences are shown for each subset.