Description of Additional Supplementary Files

Supplementary Data 1: DEGs between TI-Treg cells and PB-Treg cells from ccRCC single cell RNAseq.

Wilcoxon rank sum test with p-values adjusted using the Bonferroni method.

Supplementary Data 2: DEGs between TI-Treg cells and PB-Treg cells from HCC single cell RNAseq

Wilcoxon rank sum test with p-values adjusted using the Bonferroni method.

Supplementary Data 3: All genes for pseudotime analysis in Figure 2B.

Supplementary Data 4: Treg gene signatures from ccRCC single cell RNAseq used for filtering in Figure 3.