

Figure S1. Comparison of the expression of 14 key genes in endometrial tumor tissue and normal endometrial tissue in Genotype-Tissue Expression project-The Cancer Genome Atlas database. Comparison of (A) ZAP70; (B) LCK; (C) FOXP3; (D) TIGIT; (E) CTLA4; (F) ICOS; (G) CD5; (H) IL2RG; (I) PDCD1; (J) TNFRSF4; (K) CD27; (L) CCR7; (M) GZMB; (N) CXCL9 expression levels in endometrial carcinoma samples and normal endometrial samples in TCGA. TPM, transcripts per million; ZAP70, ζ -chain-associated protein kinase 70. * $P < 0.05$, ** $P < 0.01$, *** $P < 0.001$.

