

Figure S1: The significantly enriched signaling pathways associated with the increased TMEM200A expression. (a) Cytokine-cytokine receptor interaction, (b) chemokine signaling pathway, (c) T cell receptor signaling pathway, (d) leukocyte transendothelial migration, (e) Toll-like receptor signaling pathway, (f) TGF-β signaling pathway, (g) JAK-STAT signaling pathway, (h) mTOR signaling pathway, (i) MAPK signaling pathway, (j) pathway in cancer.