

**Figure S1** Differences in pyroptosis-related genes and immune cells between high and low risk groups. (A) Differences in the expression of pyroptosis-related genes between high- and low-risk groups. (B) Comparison of immune cells between high- and low-risk groups calculated using the single-sample gene set enrichment analysis (ssGSEA) method based on The Cancer Genome Atlas (TCGA) database. \*, P<0.05; \*\*, P<0.01; \*\*\*, P<0.001.