

Figure S1 The abundance and expression score of 26 tumor-infiltrating cells between high and low TICS group in the TCGA training cohort. ns, not significant; *, $P < 0.05$; **, $P < 0.01$; ****, $P < 0.0001$. NK, natural killer; TICS, tumor-infiltrating cell score; TCGA, The Cancer Genome Atlas.

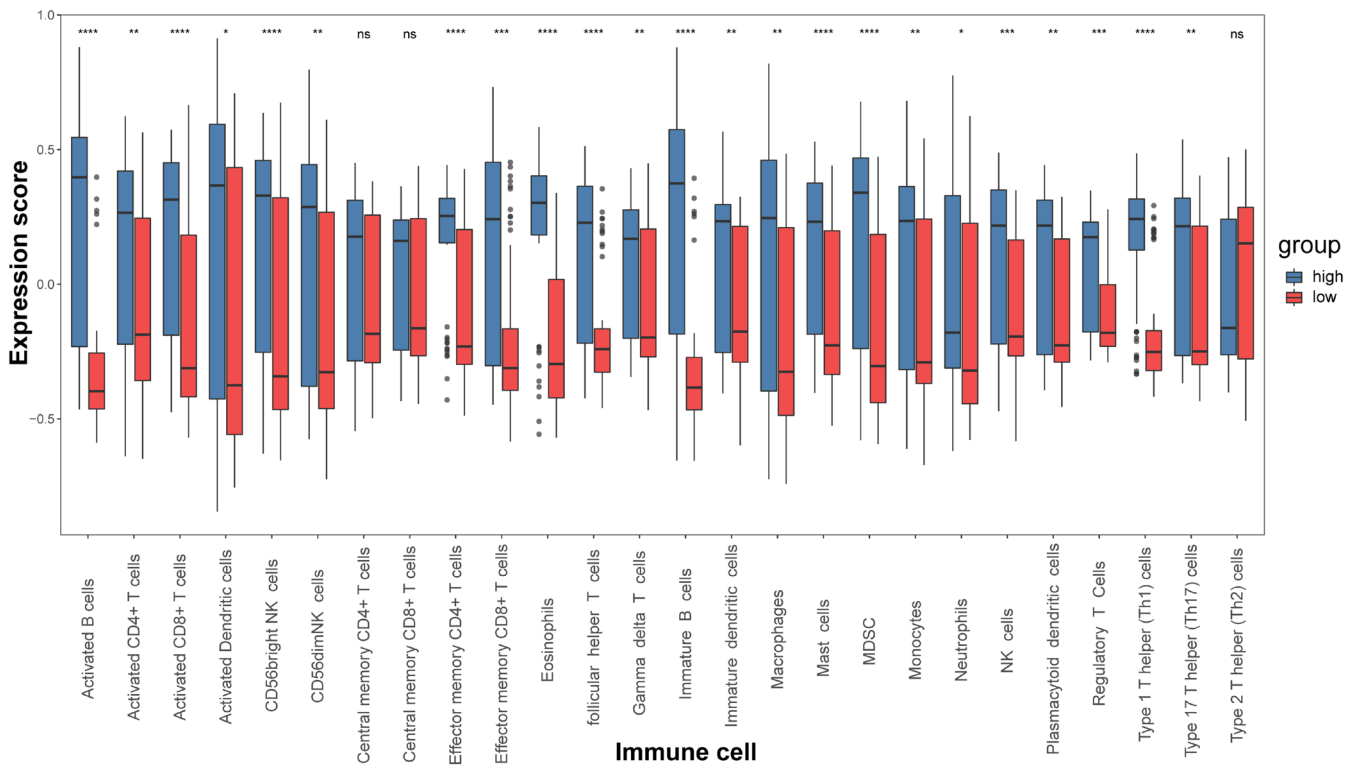


Figure S2 The abundance and expression score of 26 tumor infiltrating cells between high and low TICS group in the validation cohort. ns, not significant; *, $P < 0.05$; **, $P < 0.01$; ***, $P < 0.001$; ****, $P < 0.0001$. MDSC, myeloid-derived suppressor cells; NK, natural killer; TICS, tumor-infiltrating cell score.

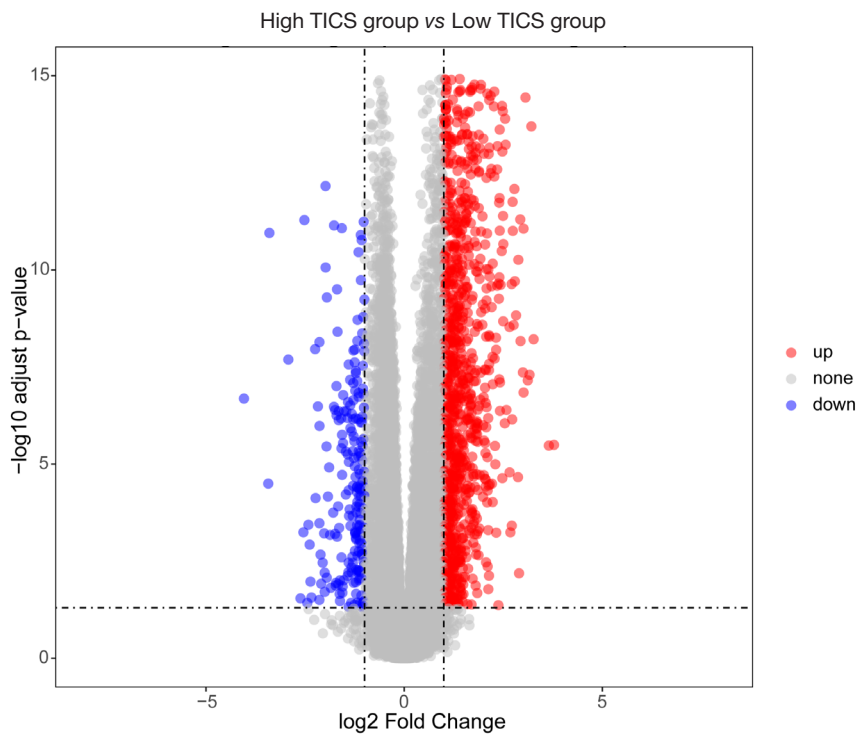


Figure S3 The volcano plot of the DEGs analysis between TICS subgroup in the TCGA training cohort. TICS, tumor-infiltrating cell score; DEGs, differentially expressed genes; TCGA, The Cancer Genome Atlas.