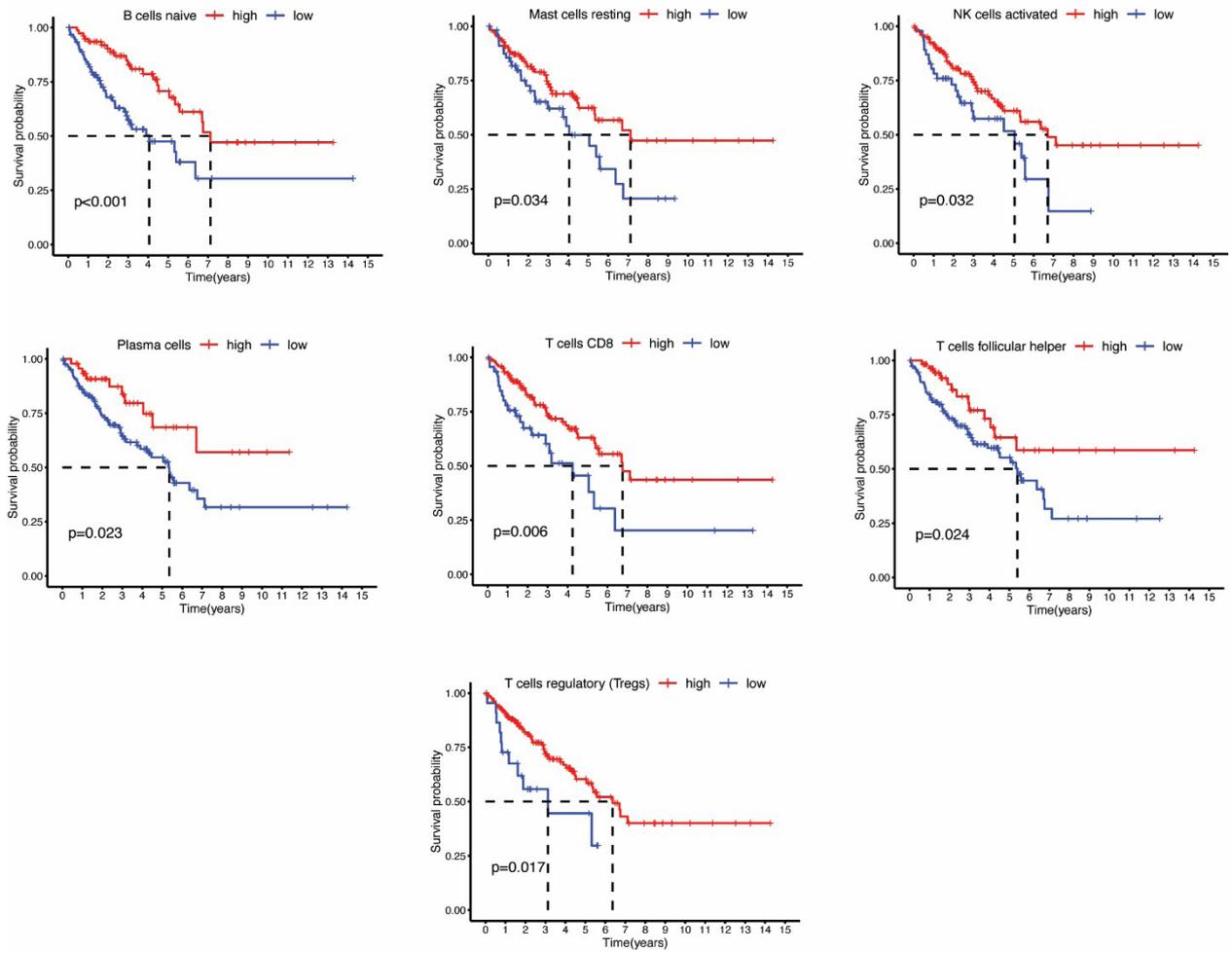
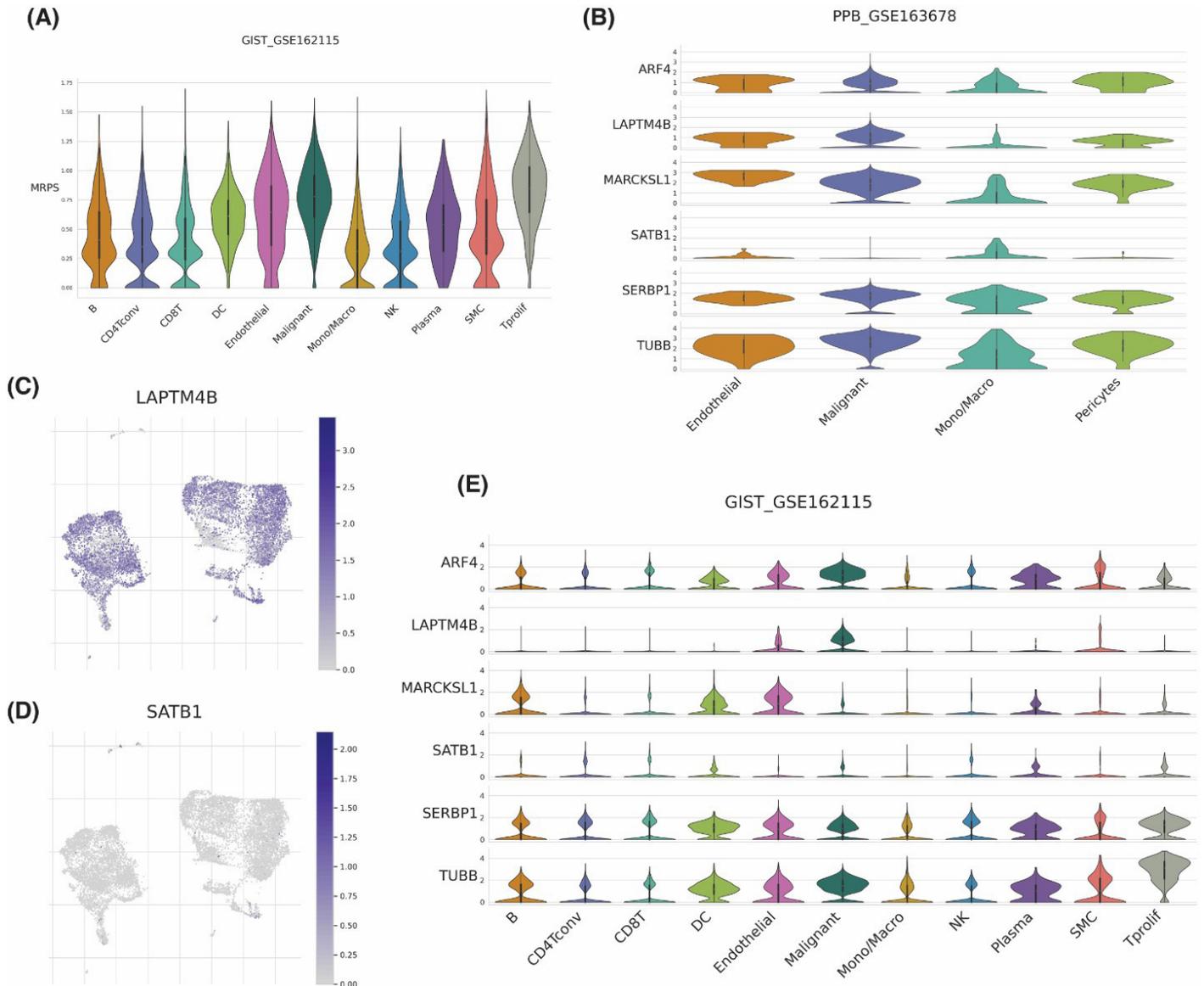


Risk stratification of soft tissue sarcoma based on activity of prognostic molecules associated with unpolarized macrophages



Supplementary Figure S1. Kaplan-Meier survival analysis of various immune cells deconvoluted based on CIBERSORT algorithm in TCGA SARC cohort



Supplementary Figure S2. **A** – Violin plot showing enrichment of MRPS score in each cell type in single-cell gastrointestinal stromal tumor dataset (GIST_GSE162115). **B** – Violin plot of showing enrichment of individual MRPS genes in each cell type in single-cell in pleuropulmonary blastoma dataset (PPB_GSE163678). **C, D** – UMAP (uniform manifold approximation and projection) plots showing enrichment of **(C)** LAPTM4B and **(D)** SATB1 in single cell in pleuropulmonary blastoma dataset (PPB_GSE163678). **E** – Violin plot showing enrichment of individual MRPS genes in each cell type in single-cell gastrointestinal stromal tumor dataset (GIST_GSE162115)