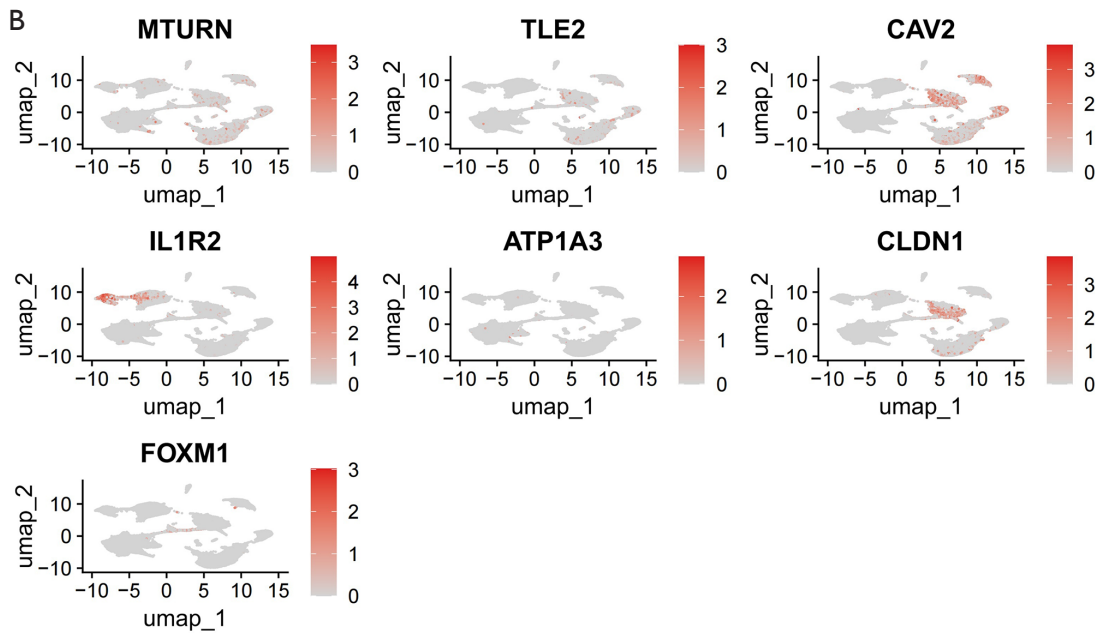
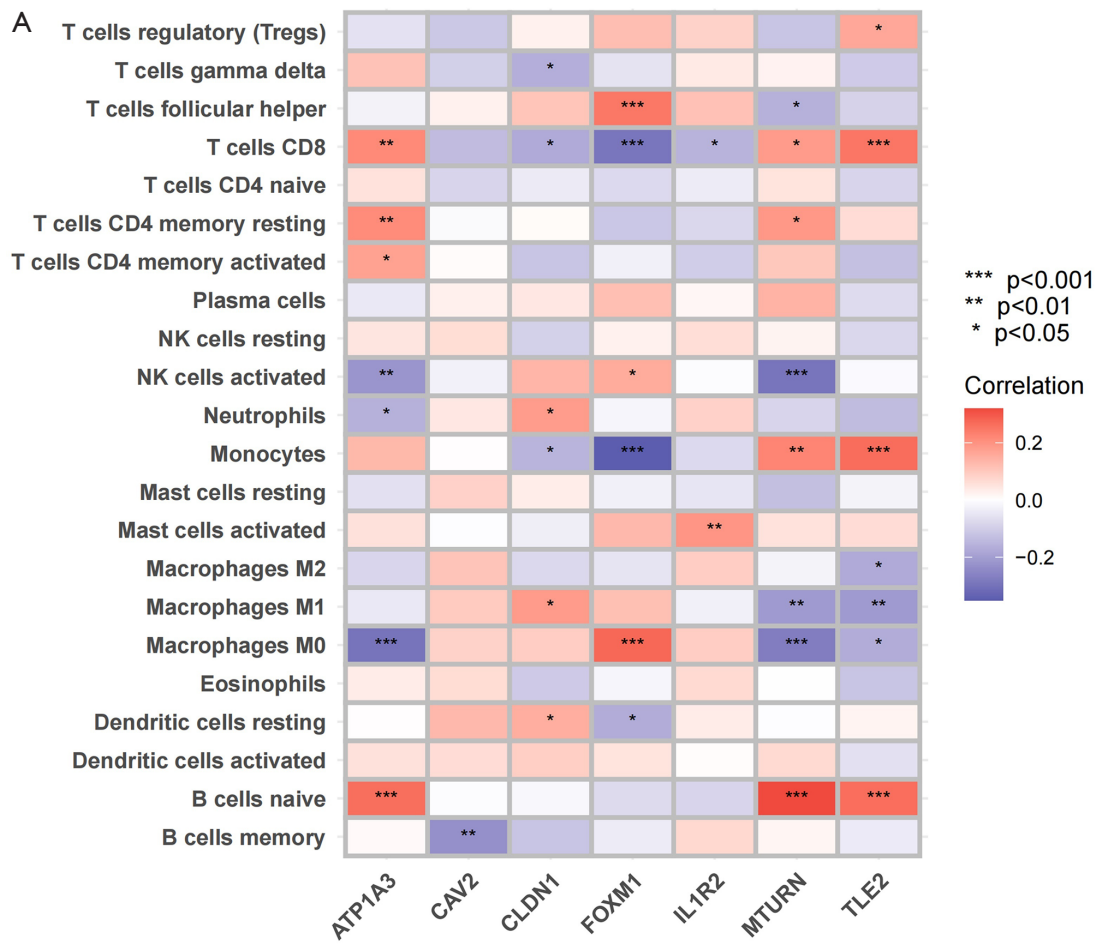
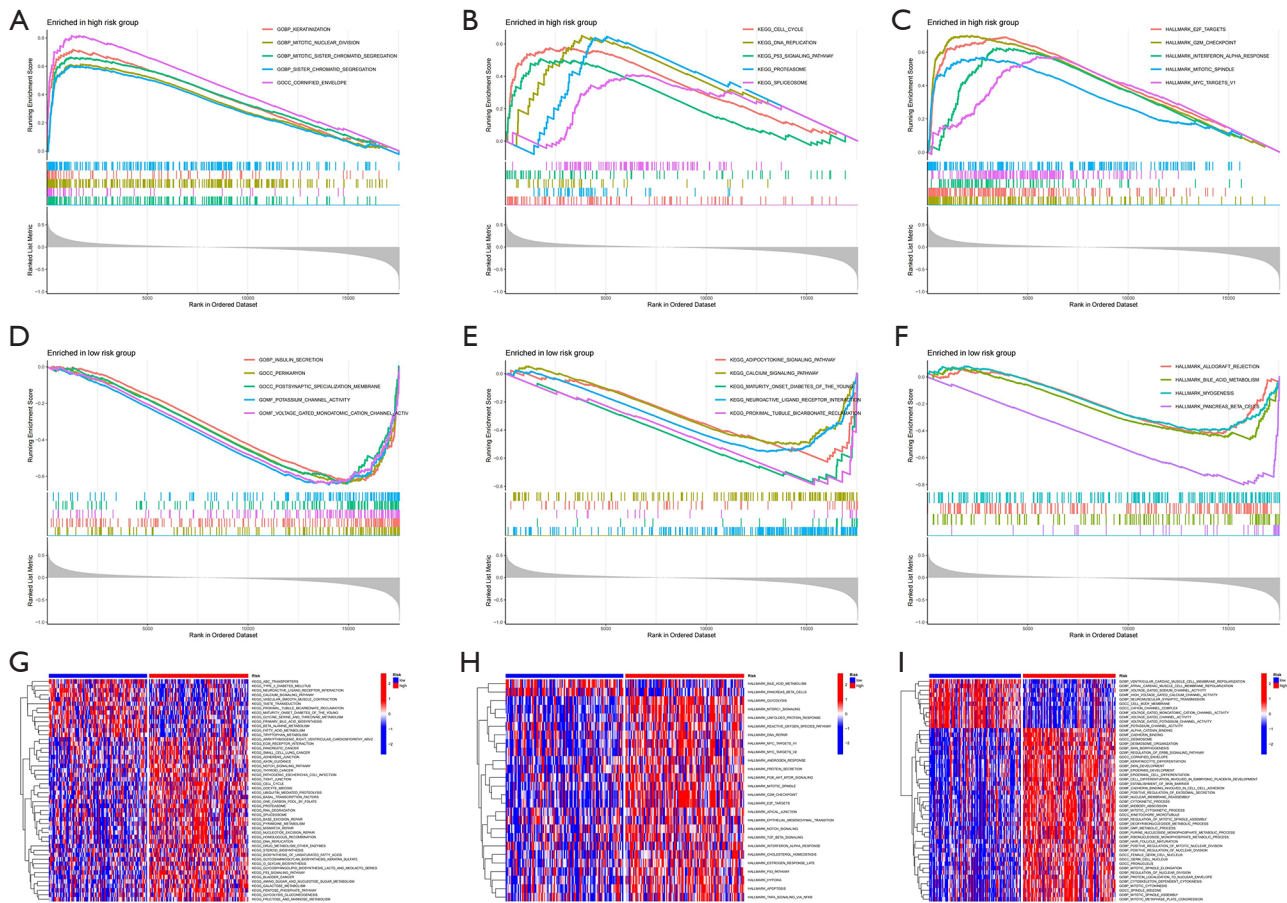


Figure S1 Unsupervised consensus clustering and identified distinct pancreatic cancer subtypes.



**Figure S2** Heatmap illustrating correlations between gene expression and immune cell infiltration in the TME.



**Figure S3** Enrichment analysis results of risk groups using the GO, KEGG, and HALLMARK datasets.