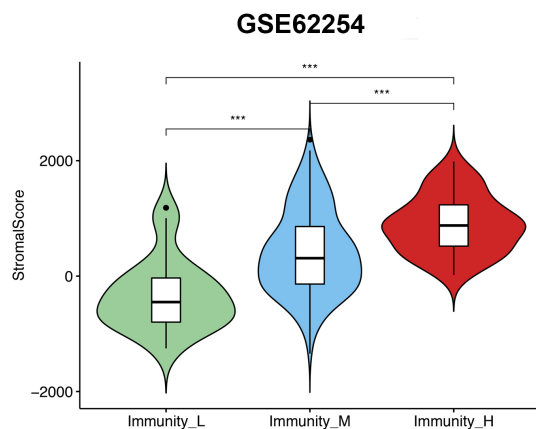
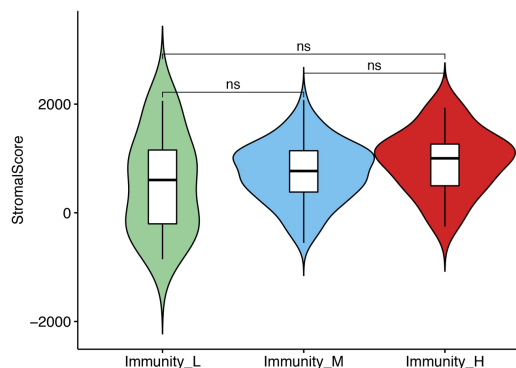
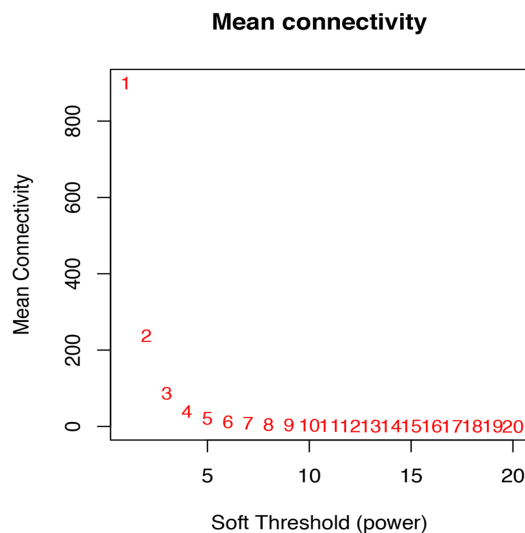
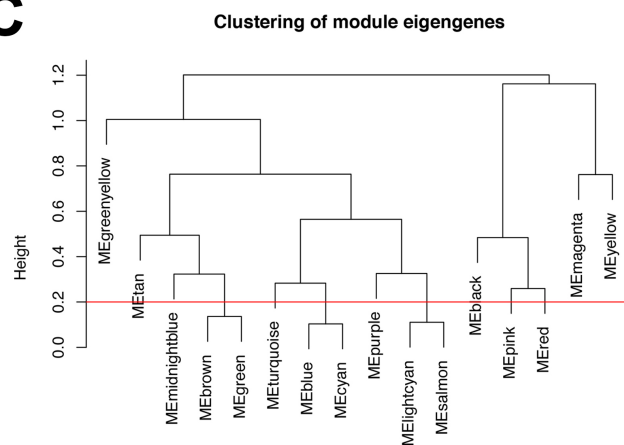
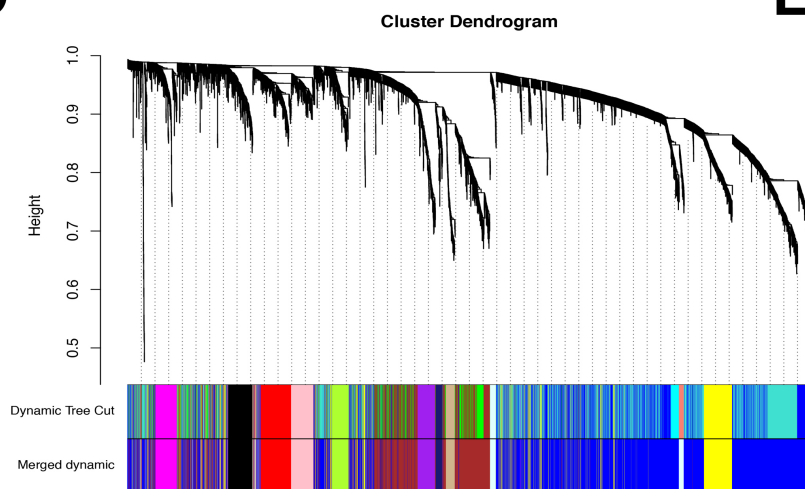
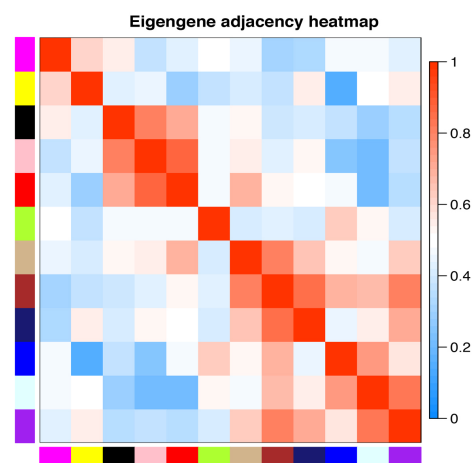
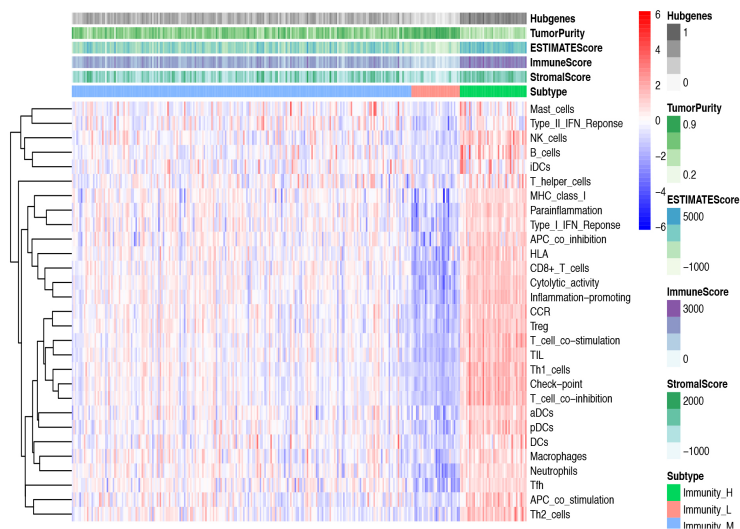
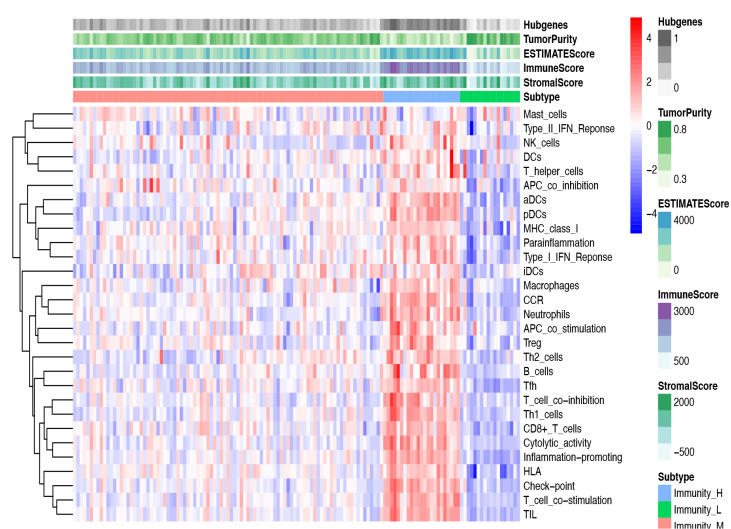
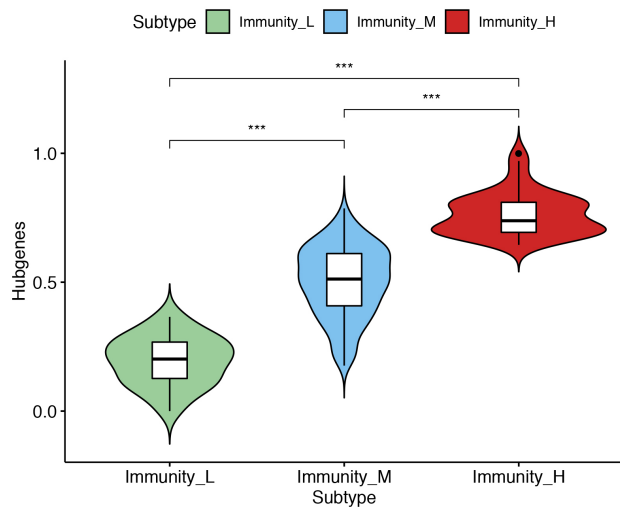
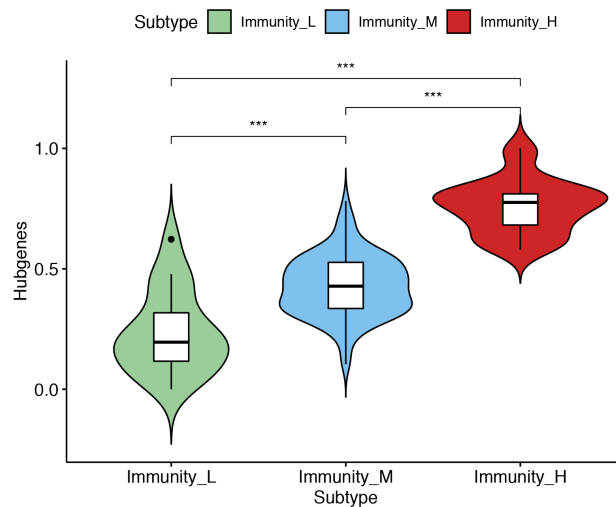


A**GSE29272****B****C****D****E**

Supplementary Figure 1. (A) The level of Stromal score in different immune subtypes (MannWhitney U test). (B) Identification of the best soft-thresholding power value. (C) 4 pairs of gene modules were merged due to their similarity according to the threshold (Red line). (D) The gene dendrogram was constructed by hierarchical clustering base on dissTOM of genes. (E) Heatmap plot of the adjacencies of modules.

A**GSE62254****GSE29272****B****GSE62254****GSE29272**

Supplementary Figure 2. (A, B) High expression level of 275 hub-genes was correlated with Immunity_H subtype and high immune cell infiltration. ***P<0.001.