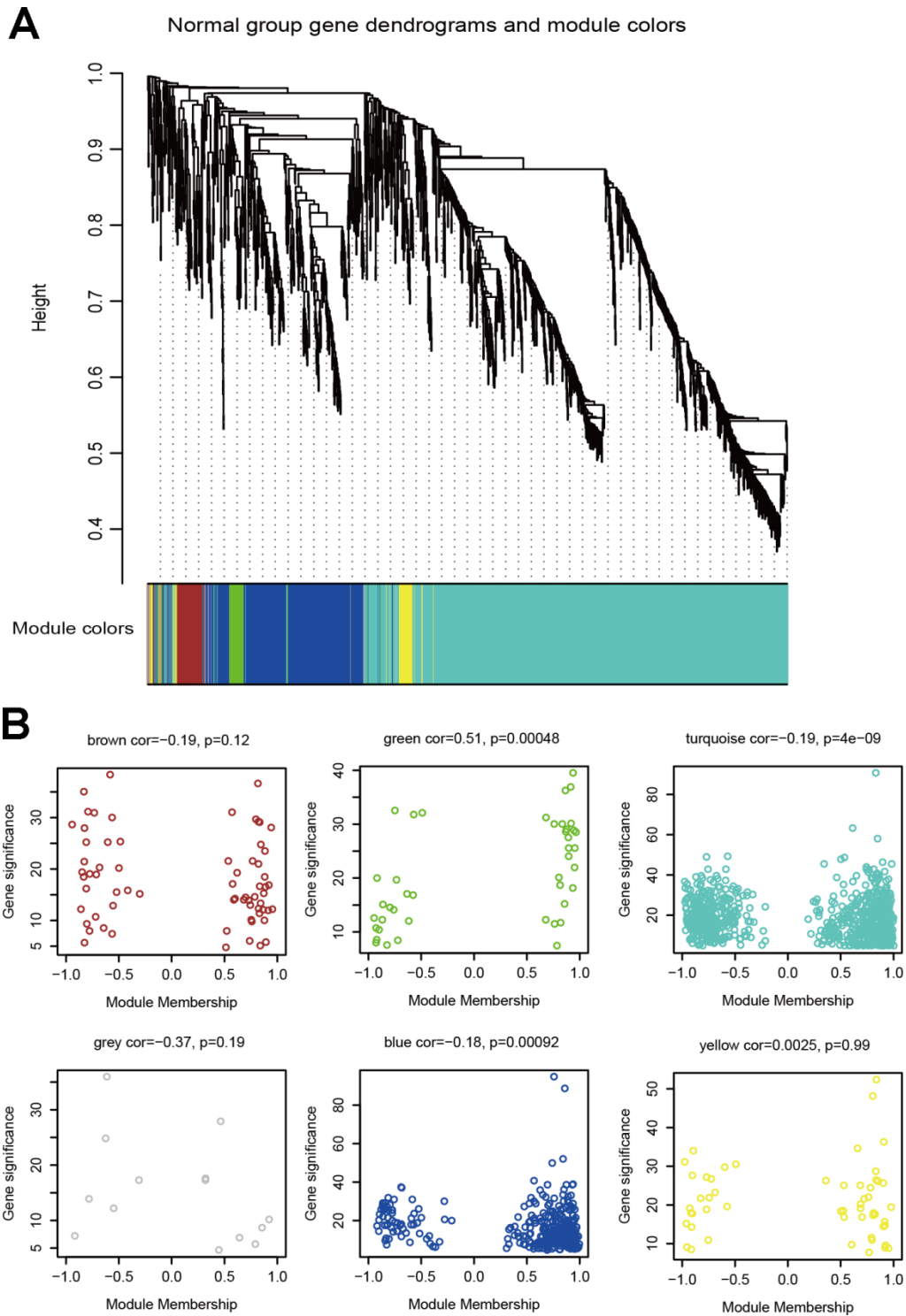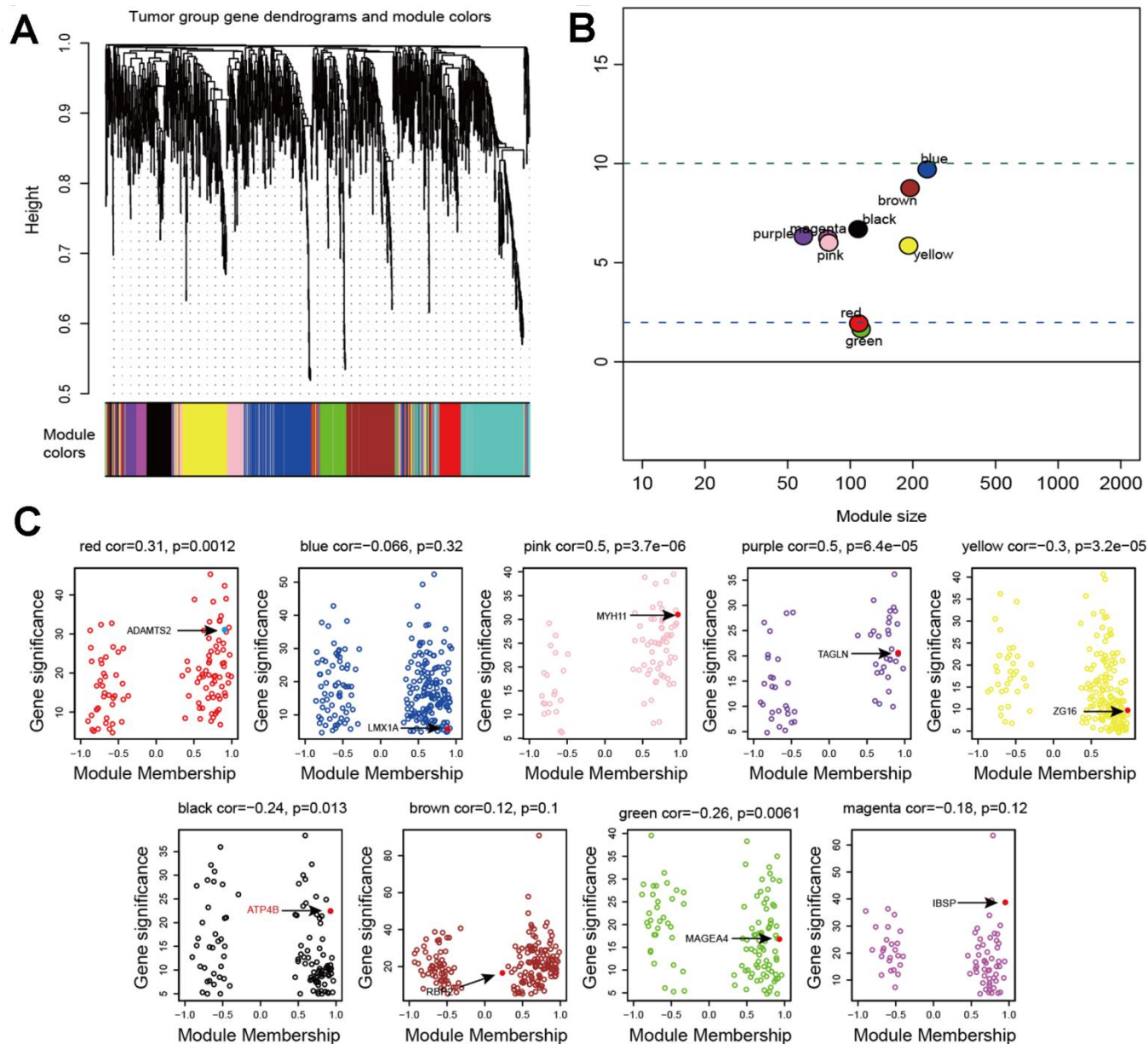# SUPPLEMENTARY FIGURES

Supplementary Figure 1. WGCNA analysis for normal tissues. (**A**) Hierarchical cluster analysis showed coexpression clusters for normal tissues. (**B**) The details of each module for the normal group.

Supplementary Figure 2. WGCNA analysis for tumor tissues. (**A**) Hierarchical cluster analysis showed coexpression clusters for tumor tissues. (**B**) The size of each module for GC tissues. (**C**) The details of each module for the tumor group and marked hub gene for each module.