Figure S1. Gene Ontology-enriched functional analysis of DEGs and PPI network of diffuse-type GC DEGs. (A) Diffuse-type DEGs. (B) Intestinal-type DEGs. (C) PPI network of diffuse type DEGs. Node colors represent the logFC of genes, with darker shades of blue indicating higher $\log_{10}$FC values. Node size is proportional to gene connectedness. DEG, differentially expressed gene; PPI, protein-protein interaction; FC, fold change.