

Supplementary Material to “Coexpression network analysis of gastric adenocarcinoma identifies hub genes as biomarker candidates and their tumor microenvironment associations”

Table S2 - GS and MM. Correlation metrics between gene expression and clinical traits. For each gene, Gene Significance (GS) indicates the strength of association with a given trait, while Module Membership (MM) measures the correlation of the gene with its respective co-expression module. These values help identify hub genes with potential biological relevance.