

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

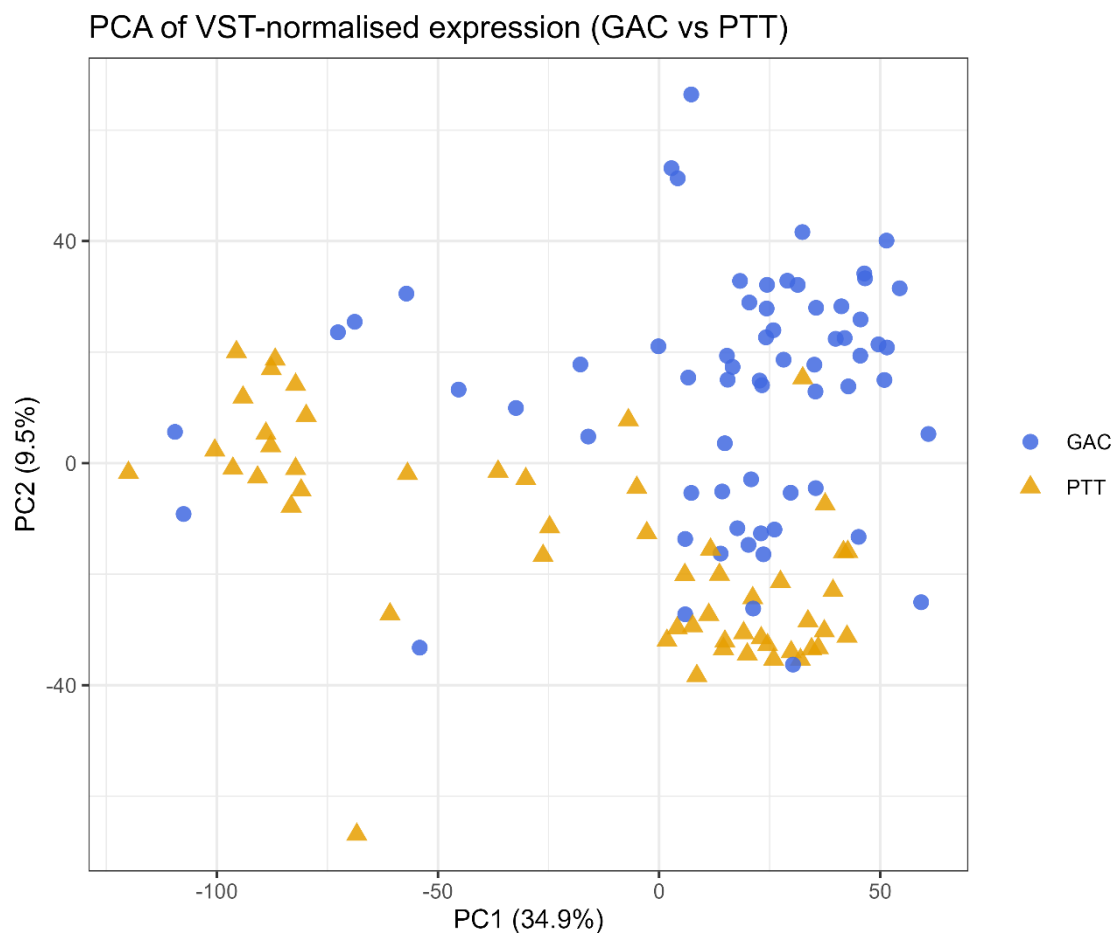


Figure S1 - Principal Component Analysis (PCA) differentiating GAC and PTT samples. Each point represents an individual sample, plotted based on the first two principal components (PC1 and PC2). Samples are color-coded according to their respective tumor groups. The distinct clustering of GAC and PTT samples indicates substantial differences in their molecular characteristics.