

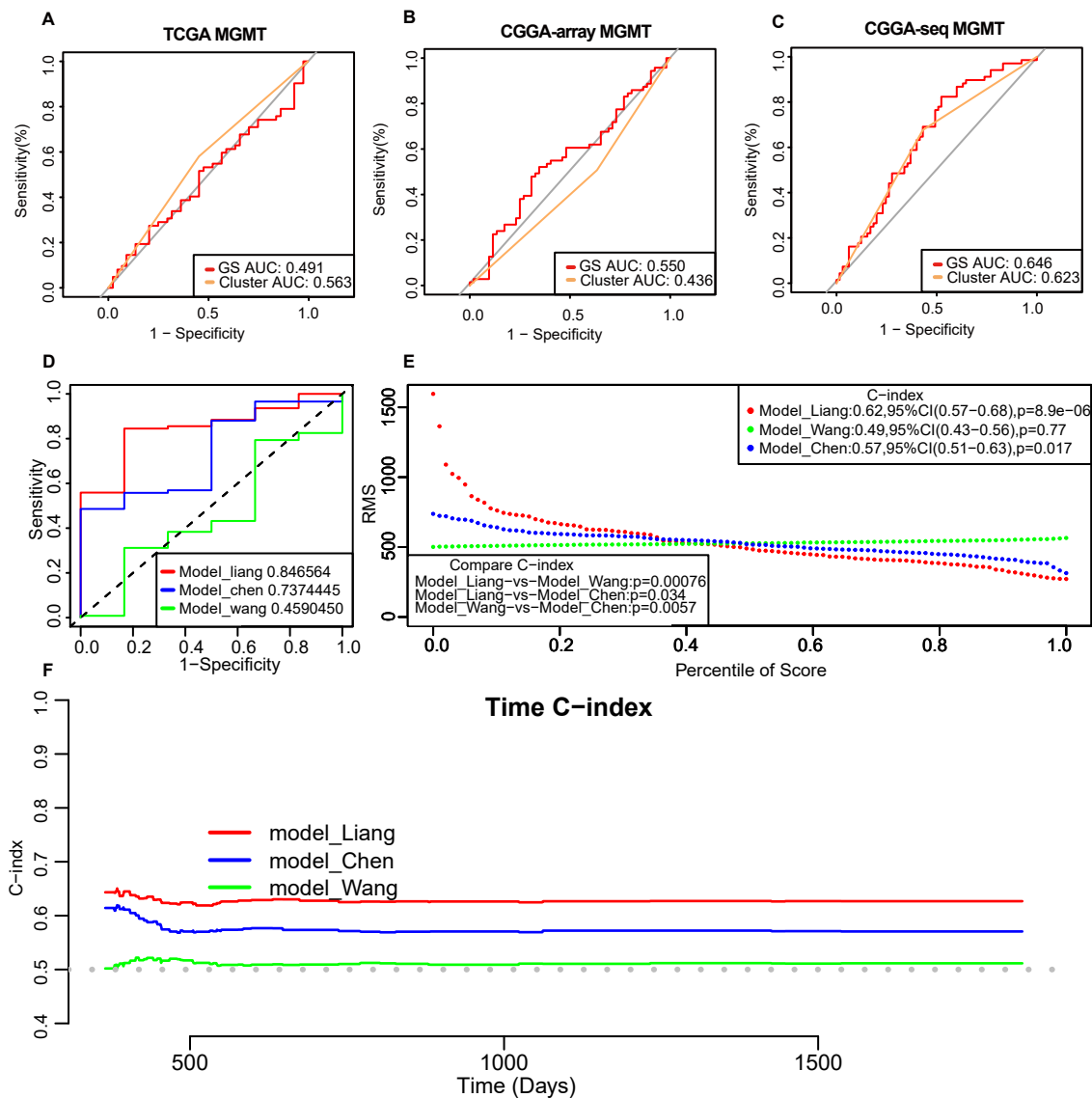


Figure S1: **Performance estimation and comparison of the Glycolytic Score.**

A-C: Receiver operating curves for MGMT methylation predictions by Glycolytic Score and PCA clusters in TCGA (A), CGGA-array (B), and CGGA-seq (C). D-F: Comparison of AUC (D), C-index (E), and time-sequencing C-index (F) for predicting glioblastoma patient OS among three established glycolysis-related prognostic model.