

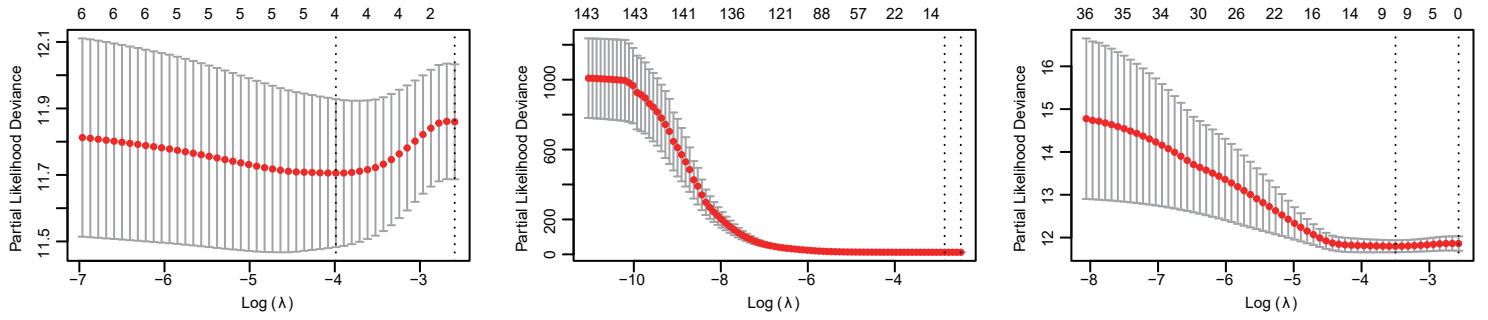
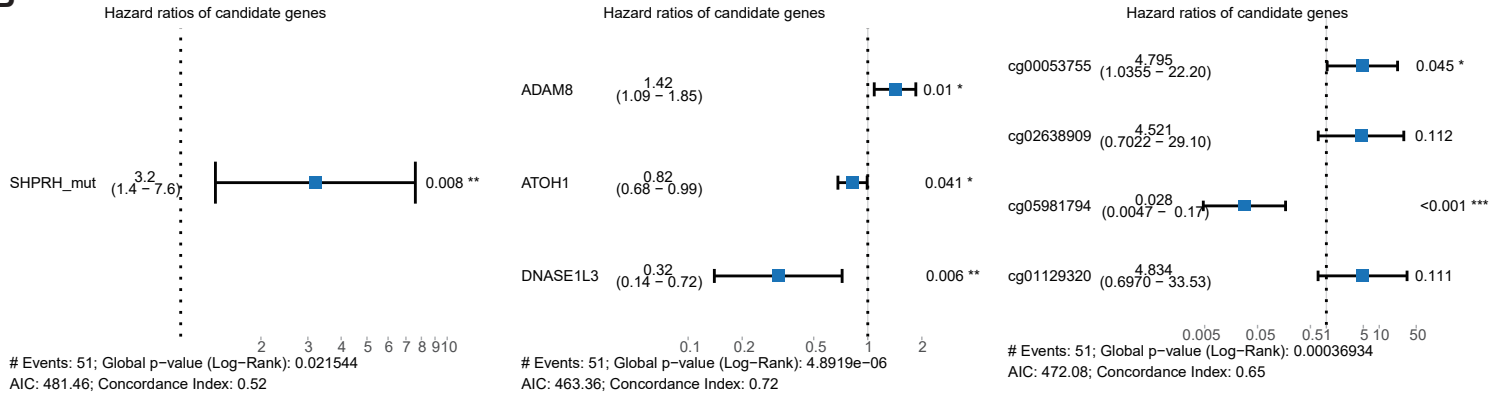
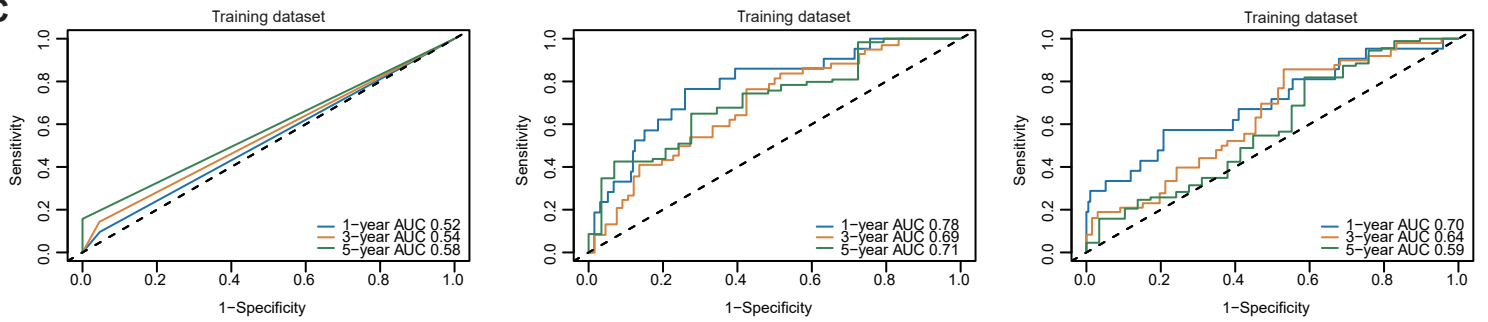
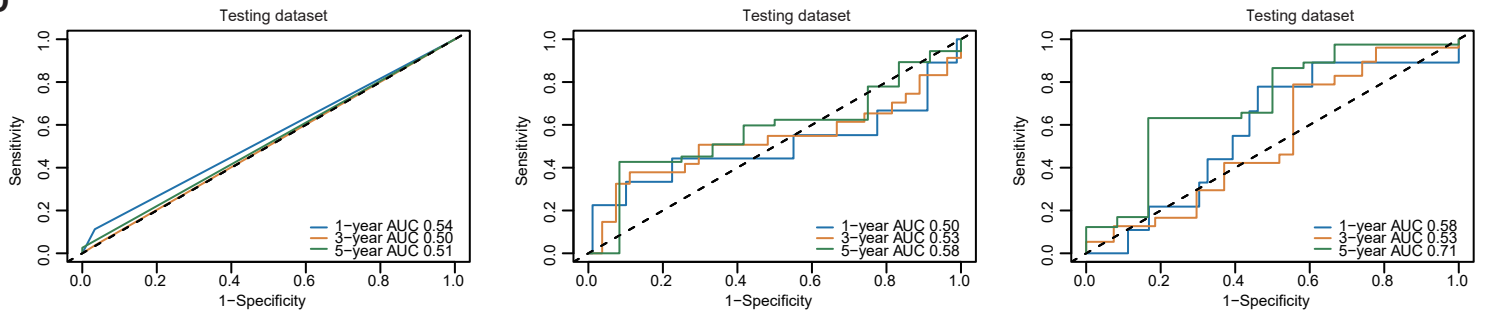
A**B****C****D**

Figure S4 The performance of the uni-omics model of RNA-seq (left), WES (middle), and methylation profiles (right). (A) Identification of the optimal penalization coefficient lambda in the Lasso regression model. (B) Forest plot of the prognostic impact of the candidate variables. (C-D) ROC curves of the risk score for predicting 1-year, 3-year, and 5-year survival in (C) training and (D) testing dataset.