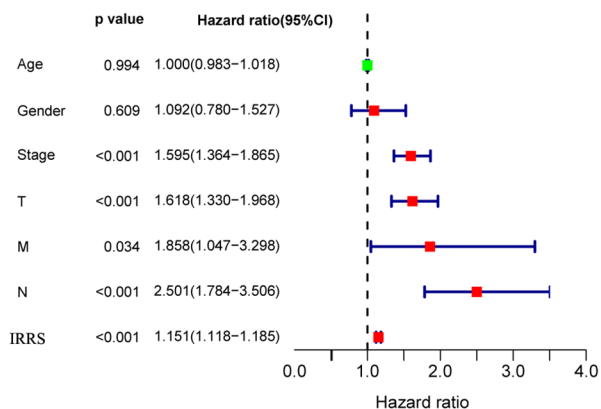
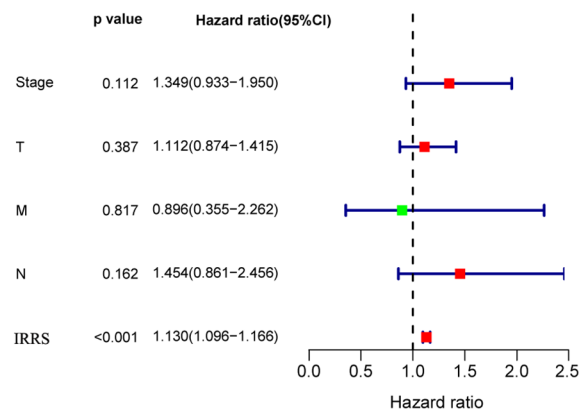


**Figure S1** Prognostic-related genes incorporated into the IRRS formula. Forest plot illustrating the 34 genes selected and incorporated into the risk score formula after the LASSO regression analysis. \*, P<0.05; \*\*, P<0.01.

A



B



**Figure S2** The IRRS independently predicted OS in patients with LUAD. (A) Univariate Cox regression analysis revealed that clinical stage, T stage, N stage, M stage, and IRRS were independent predictors for OS in LUAD. (B) Multivariate regression analysis revealed that the IRRS was an independent predictor for overall OS in patients with LUAD.