The Five-lncRNA signature remained with largest AUC validation in the RNA-seq and independent qPCR group (n=84)

Univariate Cox regression analysis (P < 0.05)
Kaplan–Meier (P < 0.05)

Ten lncRNAs selected to construct predictive model

Combination

The risk-score model of 1023 signatures in the GSE53624 dataset

ROC analysis

The Five-lncRNA signature remained with largest AUC validation in the RNA-seq and independent qPCR group (n=84)

KM analysis  Cox analysis  ROC analysis