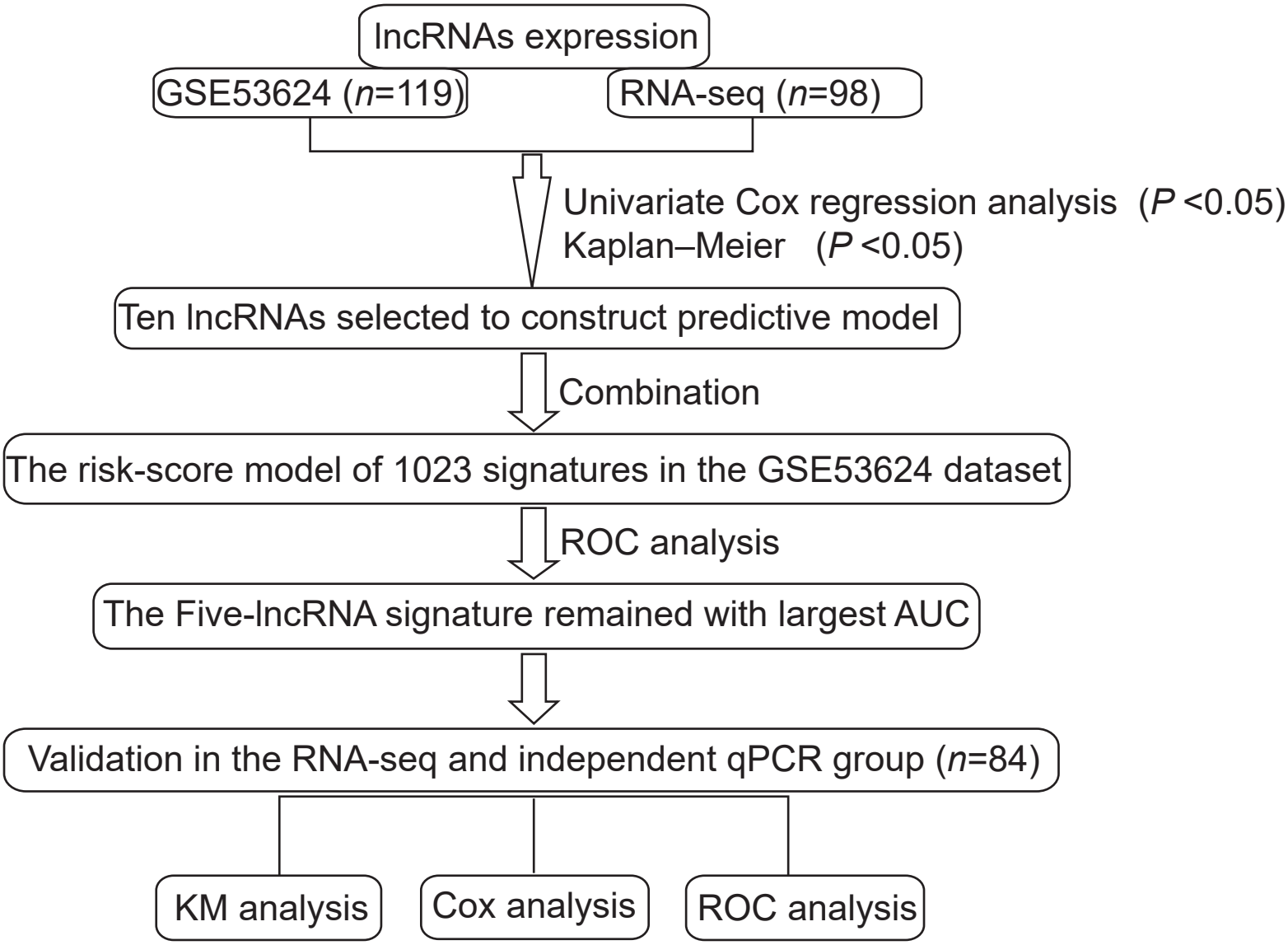
Figure S1


lncRNAs expression
GSE53624 (n=119)
RNA-seq (n=98)
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