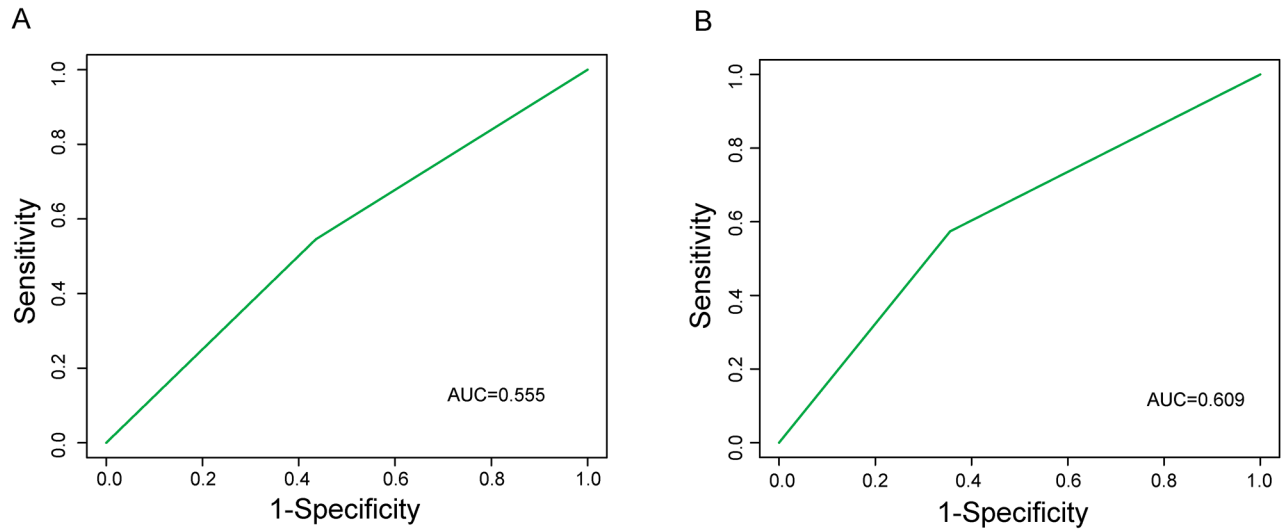
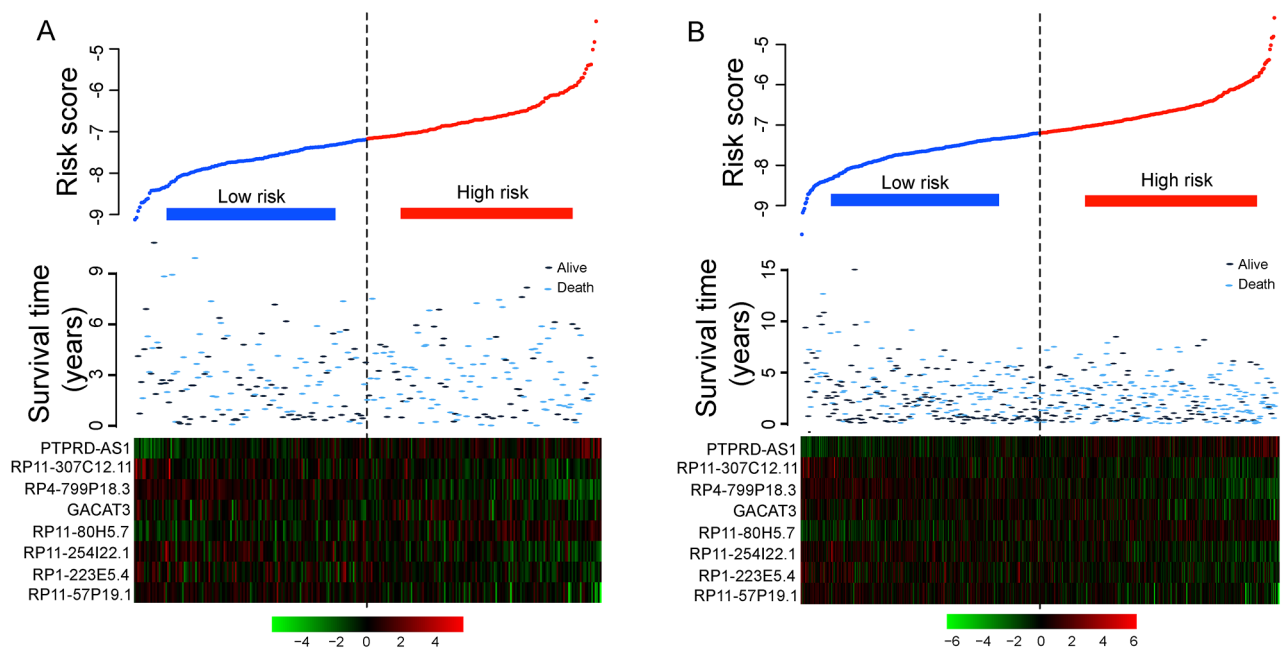


Comprehensive analysis of lncRNA expression profiles reveals a novel lncRNA signature to discriminate nonequivalent outcomes in patients with ovarian cancer

SUPPLEMENTARY FIGURES AND TABLE



Supplementary Figure S1: The time-dependent ROC curves within 5 years for the eight-lncRNA signature in the validation cohort A. and in the entire TCGA cohort B.



Supplementary Figure S2: The distribution of risk scores, survival status and lncRNA expression for patients in the validation cohort A. and in the entire TCGA cohort B.

Supplementary Table S1: Detailed clinical information of OvCa patients enrolled in this study.

See Supplementary File 1