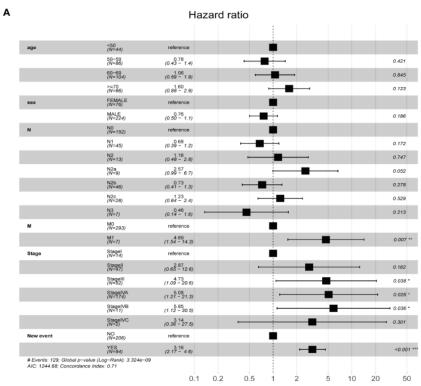
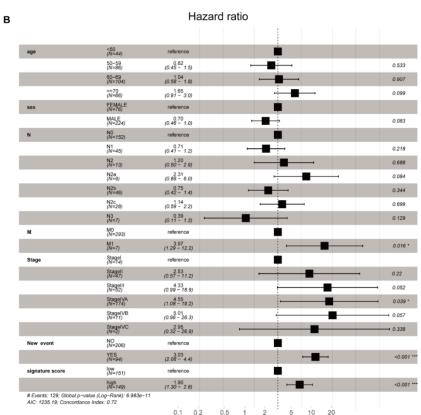
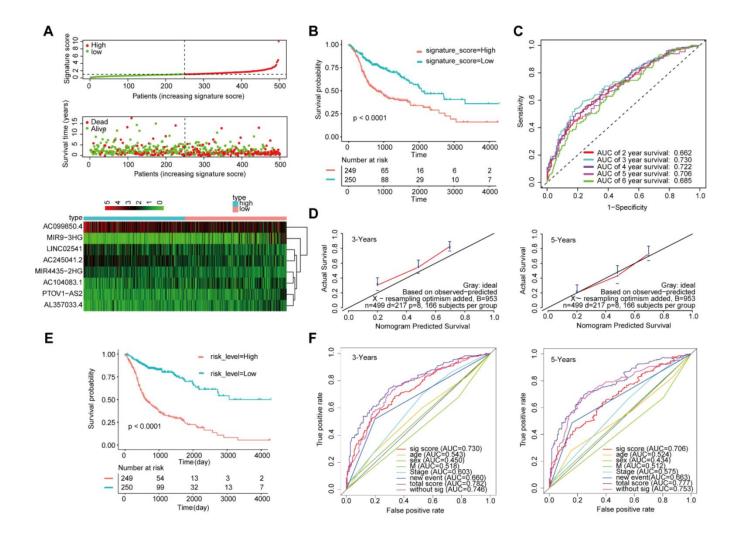
SUPPLEMENTARY FIGURES

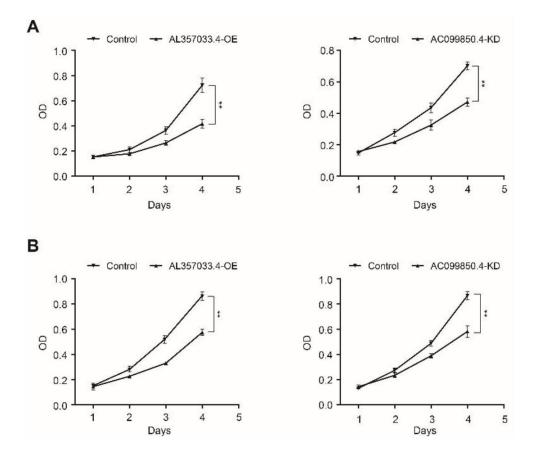




Supplementary Figure 1. Comparison of the multivariate Cox regression results of the two groups with and without the signature score. (A) The group without the signature score. (B) The group with the signature score.



Supplementary Figure 2. Validation of the model by the entire TCGA-HNSCC set (n=499). (A) Distribution of 8-IncRNA-based signature scores, IncRNA expression levels and patient survival durations in the entire set. (B) Kaplan-Meier curves of OS based on the 8-IncRNA signature. (C) ROC curve analyses based on the 8-IncRNA signature. (D) Calibration curves of the nomogram for the estimation of survival rates at 3 and 5 years. (E) Kaplan-Meier curves of OS according to the total risk score. (F) ROC curves according to the nomogram and IncRNA signature score.



Supplementary Figure 3. Proliferation of FaDu (A) and Hep-2 (B) cells was analyzed using CCK-8 assay following IncRNAs overexpression or knockdown.