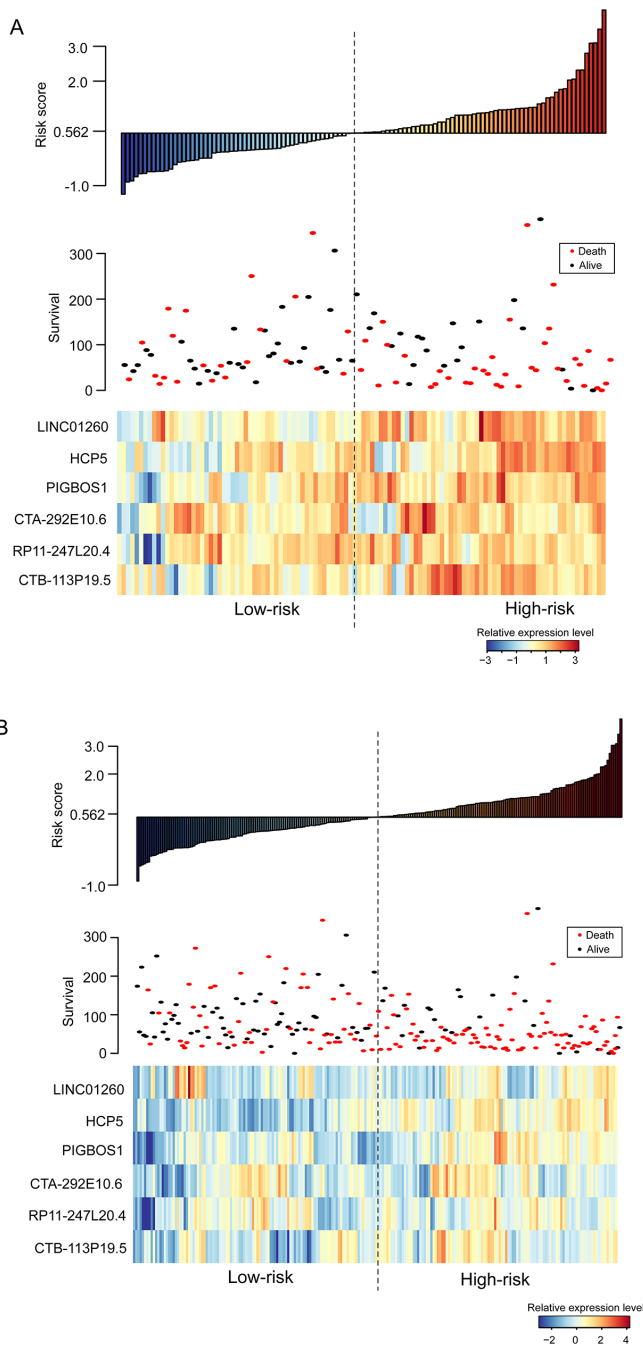


Expression profiles analysis of long non-coding RNAs identified novel lncRNA biomarkers with predictive value in outcome of cutaneous melanoma

SUPPLEMENTARY MATERIALS



Supplementary Figure 1: Distribution of risk scores, patient survival status and lncRNA expression map in the validation cohort (A) and the entire TCGA cohort (B).