

Supplementary Material figures

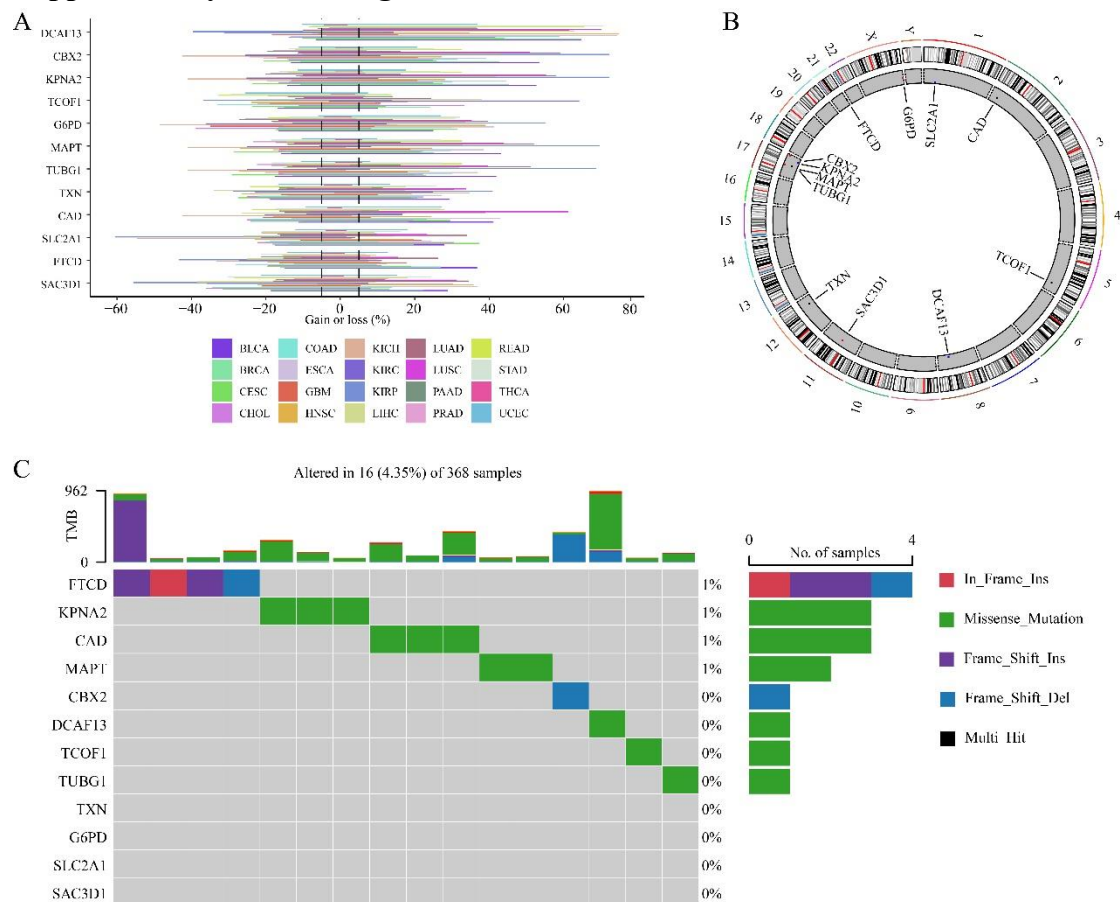


Figure S1. Variation and Chromosomal Distribution of LLPS Prognosis-Related Genes in HCC. (A) Copy number variation (CNV) values of LLPS prognosis-related genes across various cancers as per TCGA database. (B) Chromosomal distribution of LLPS prognosis-related genes. (C) Somatic mutations in LLPS prognosis-related genes according to TCGA data.

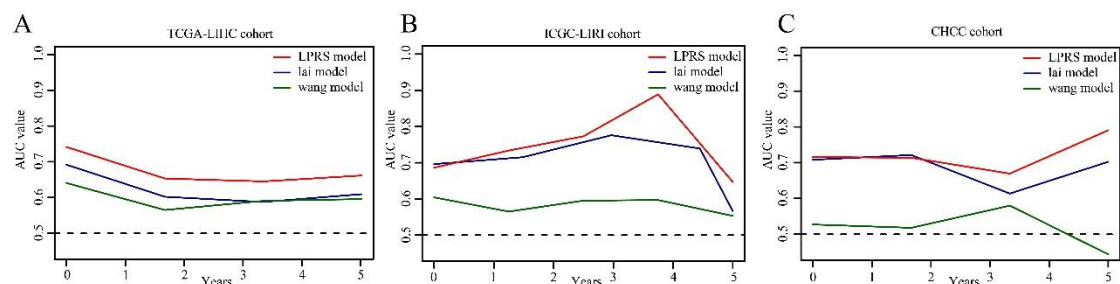


Figure S2. The time-dependent AUCs of the signature scores in three cohorts. (A) The time-dependent AUCs of the LPRS, lai and wang model in TCGA-LIHC cohort. (B) The time-dependent AUCs of three model in ICGC-LIRI cohort. (C) The time-dependent AUCs of three model in CHCC cohort.

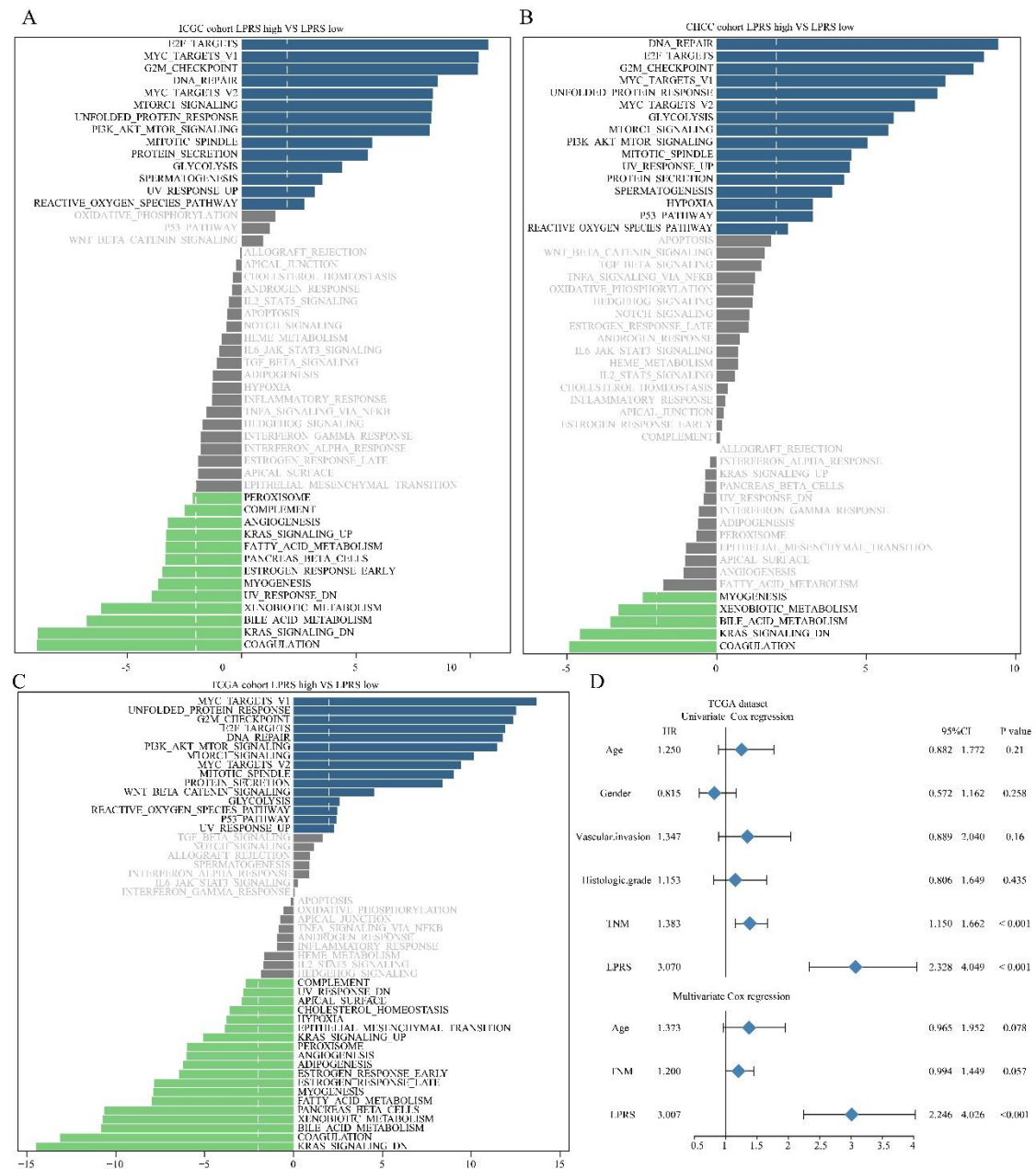


Figure S3. Prognostic and Pathway Characteristics of LPRS. (A-C) Gene Set Variation Analysis (GSEA) comparing high- and low-LPRS patients. (D) Univariate and multivariate Cox analyses of clinicopathological characteristics and LPRS in the TCGA cohort.