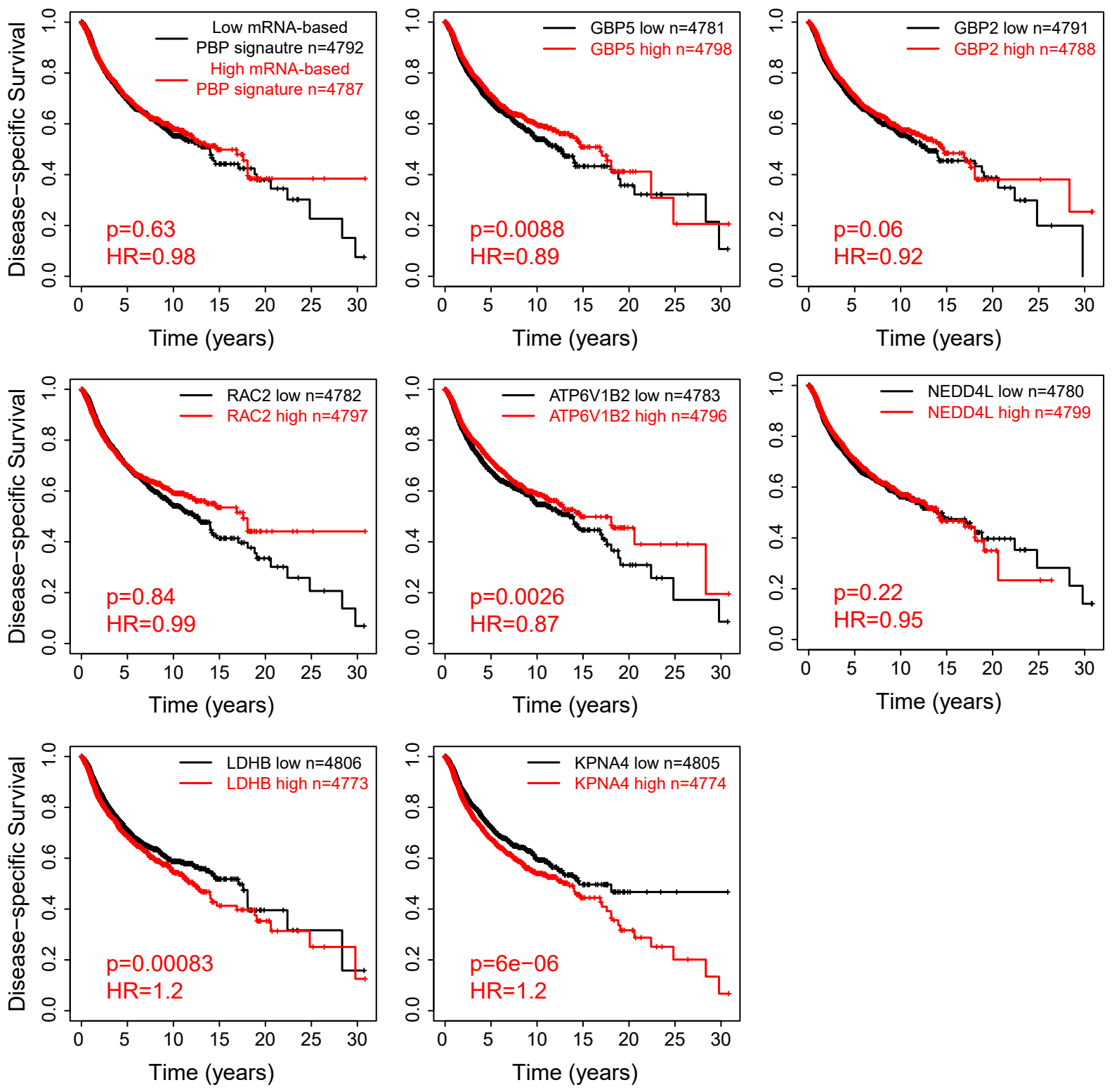# Supplementary Figure 7

TCGA pan-cancer analysis

Supplementary Figure 7. Kaplan-Meier (KM) curves show disease-specific survival of cancer samples from the TCGA pan-cancer dataset. Samples were categorized into high and low levels of the PBP signature score or individual signature gene (GBP5, GBP2, RAC2, ATP6V1B2, NEDD4L, LDHB, and KPNA4) based on the median mRNA expression cutoff.