

Supplementary Figure 8. (A) Scatter plot shows the correlation between the KIPA-based PBP signature score and protein-based or mRNA-based multigene proliferation scores (MGPS). (B) Single-cell RNA-seq data from a public dataset with 26 primary breast tumors. UMAP visualization shows the cell types annotated in the previous publication (30), generated from the Broad Institute Single Cell portal at https://singlecell.broadinstitute.org/single_cell/study/SCP1039. (C) Log-normalized expression of seven PBP signature genes in different clusters, generated from the Broad Institute Single Cell portal.

KPNA4

LDHB

NEDD4L