

Supplementary Figure 3. Single-cell RNA sequencing analysis of G1/S-phase cell cycle genes in breast cancer.

**(A)** Dimensionality reduction t-SNE maps of G1/S phase CDKs and cyclins expression in EpCAM+EGFR+ cells, and **(B)** analyses demonstrated CDK2 and cyclin E are upregulated in TNBCs (see Figure 2A and 2B). Although CDK4 is upregulated, its binding partners (CDK6 and cyclin D) are not in upregulated in TNBCs. *P*-values determined by two-tailed unpaired *t* test against TNBC subtype.