

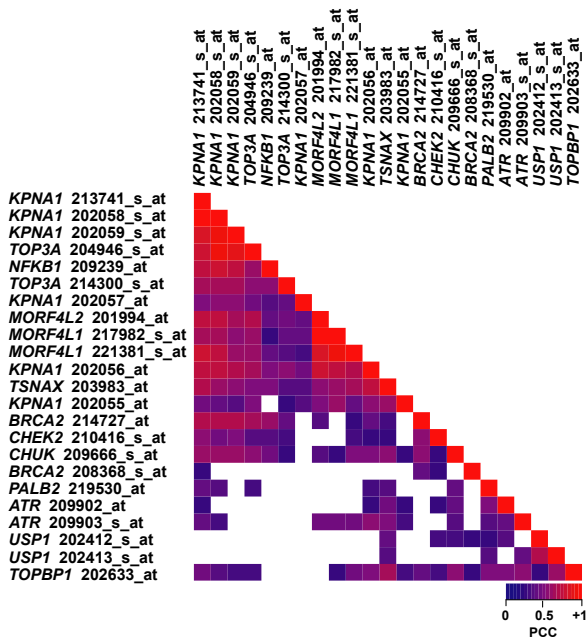


Figure S1. Matrix of gene co-expression values across human samples (cell lines, tissues and organs; GSE1133; $n = 158$) using Pearson correlation coefficients (PCCs, colored scale values are shown at the bottom right) and all microarray probes for known and potential FA/BrCa pathway components. Blanks represent non-significant correlations.