Supplementary Figure S1: Distribution of normalized TOP2A expression values among breast cancers

The distribution of normalized TOP2A expression values in the combined dataset of 1681 samples is given in the upper panels stratified according to positive (left panel) and negative (right panel) ER status, respectively. For comparison, the distribution among the cohort of ErbB2 negative tumors only is given in the two lower panels, demonstrating that the subset of ErbB2 positive tumors have only a marginal effect on the overall distribution of TOP2A in the total cohort.

All samples (n=1681)

ER positive (n=1200)  ER negative (n=481)

normalized TOP2A expression

ErbB2 negative only (n=1432)

ER positive (n=1085)  ER negative (n=347)