Supplementary Figure 1. Expression of PLK2, PLK3 and PLK4 in triple-negative breast cancer. (A,C,E) The expressions of PLK2, PLK3 and PLK4 were studied at the mRNA level in the different tumor subgroups of our cohort (TNBC, HER2, luminal A and luminal B). The expression was also analyzed in healthy breast tissues. (B,D,F) the PLK2, PLK3 and PLK4 genes were analyzed at the DNA level. The smoothed segmented copy number (CN) signal is presented in boxplots, with dashed lines indicating the thresholds retained to call copy number gains and losses.