

Figure S1. Toca-1 transcript level profiling in breast cancer cell lines according to p53 mutation status. Toca-1 mRNA levels were measured for the same panel of cell lines by qRT-PCR ($2^{-\Delta\Delta}_{CT}$ values for Toca-1 were normalized to GAPDH for each cell line, and graph depicts transcript levels relative to HBL-100 cells (mean \pm SD; triplicate samples).