



**Supplementary Figure S5. Mutations identified in RNA-Seq analysis.** The data obtained in RNA-Seq studies were graphed using the Oviz-Bio tool (<https://academic.oup.com/nar/article/48/W1/W415/5835823>). Only driver breast cancer genes (99) were selected from the Intogene data base (<https://www.intogen.org/download>) and we found mutations in 19 driver genes after read depth filtering (DP >50). Of note is the missense mutation found in ESR1 in the M070 patient. Left, Percentage of samples with mutations. Color defines the type of mutations.