Supplemental Figure S1. Technical validation of array data. RT-qPCR data for the 16 genes using RNA of T47D cells that was utilized for the microarrays. 13 of the 16 genes are significantly upregulated in the Vehicle vs. R5020 cohort, *p< 0.05. KIF11, PKMYT1 and TOP2A were not able to be validated. The same 13 genes were downregulated # p<0.05 for R5020 vs. R5020 plus TPA.