



Supplemental Figure 2. Comparison of the Relative Expression of CD24 across all screened neuroblastoma cells and tumors. qRT-PCR was performed comparing IMR-32, SK-N-AS, SMS-KAN, and SMS-KANR cells (*in vitro*, prior to introduction into the mouse host) to tumors (*in vivo*, averaged from post-treatment vehicle treated control tumors) for each neuroblastoma. Expression was normalized to GAPDH. All qPCR data shown are the composite of triplicate wells acquired from triplicate experiments. Error bars represent standard deviation. * $p > 0.05$ from SK-N-AS Cells, unpaired t-test.