

Description of Additional Supplementary Files

File Name: Supplementary Data 1

Description: Genes differentially expressed between four MYCN gene-amplified [BE(2)-C, Kelly, CHP134 and SK-N-DZ] and two MYCN-non-amplified (SY5Y and SK-N-AS) neuroblastoma cell lines, as identified by RNA sequencing. FC and CPM represented fold change and counts per million reads, and FDR represented false discovery rate. Transcripts with an absolute log₂ fold change of >1 and a false discovery rate of <0.05 were listed.