

## Description of Additional Supplementary Files

File Name: Supplementary Data 1

Description: List of 801 genes considered to be cancer-related according to COSMIC and literature as described in the methods section.

File Name: Supplementary Data 2:

Description: List of published MYCN target genes.

File Name: Supplementary Data 3:

Description: List of differentially expressed genes of high-risk neuroblastoma.

File Name: Supplementary Data 4:

Description: List of the 100 most variable expressed genes assessed by RNA Seq in our 10 patient neuroblastoma cohort.

File Name: Supplementary Software 1

Description: Source code in *R* for running *sciClone* and for constructing the clonality trees provided as a zipped file.