Supplemental Methods

Survival curves

Kaplan Meier overall survival curves for individuals with high and low expression of \textit{KIF1B} and \textit{XAF1} were generated using the ‘Tumor Neuroblastoma public - Versteeg - 88 - MAS5.0 - u133p2’ dataset from R2: microarray analysis and visualization platform. The cutoff was determined using the Kaplan scanner tool within the web application. Samples are sorted according to the expression of the respective genes and divided into two groups based on a cutoff expression value. All cutoff expression levels and the resulting groups were analyzed for overall survival, with the criteria that each group includes at least eight samples. For each particular cutoff level and grouping, the raw \textit{p}-value significance of projected survival was calculated. The graph depicts the best \textit{p}-value corrected for multiple testing (Bonferroni correction). The numbers in parentheses indicate the number of individuals in each group.