

Suppl. Table 1

Cell line	TMM	MYCN/TERT status	TERT expression	ALK status
BE(2)C	Telomerase pos.	<i>MYCN</i> amplified	10.31	WT
CLB-GA	Telomerase pos.	<i>TERT</i> rearr.	13.67	R1275Q
GI-MEN	Telomerase pos.	<i>TERT</i> rearr.	13.88	WT
IMR-32	Telomerase pos.	<i>MYCN</i> amplified	10.95	WT
KELLY	Telomerase pos.	<i>MYCN</i> amplified	11.37	F1174L
LS	Telomerase pos.	<i>MYCN</i> amplified	10.49	WT
SH-SY5Y	Telomerase pos.	WT	12.38	F1174L
SK-NFI	ALT	WT	8.81	D1529E
TR-14	Telomerase pos.	<i>MYCN</i> amplified	8.92	amplified

Suppl. Table 1: Molecular characteristics of cell lines used in this study; TMM, telomere maintenance mechanism; WT, wild-type; ALT, alternative lengthening of telomeres. *TERT* expression is given as log₂ intensity according to microarray analysis.