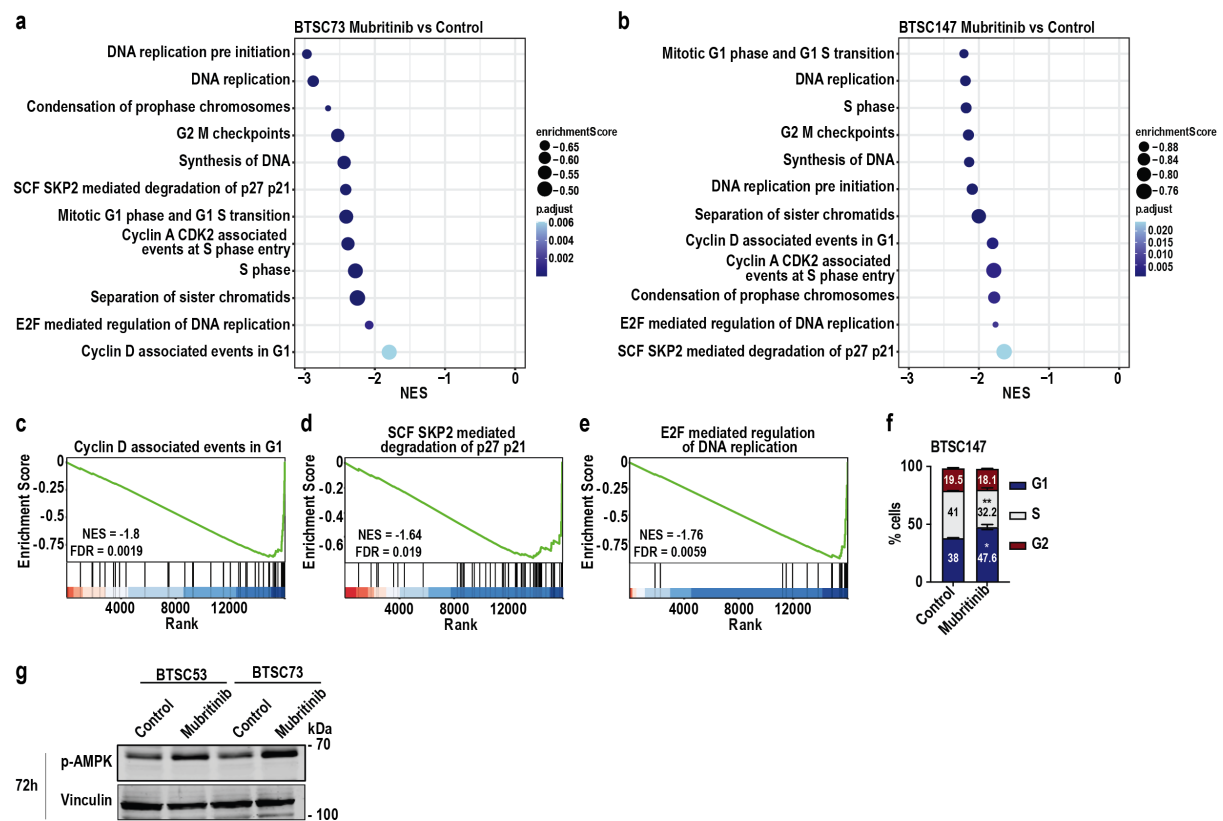


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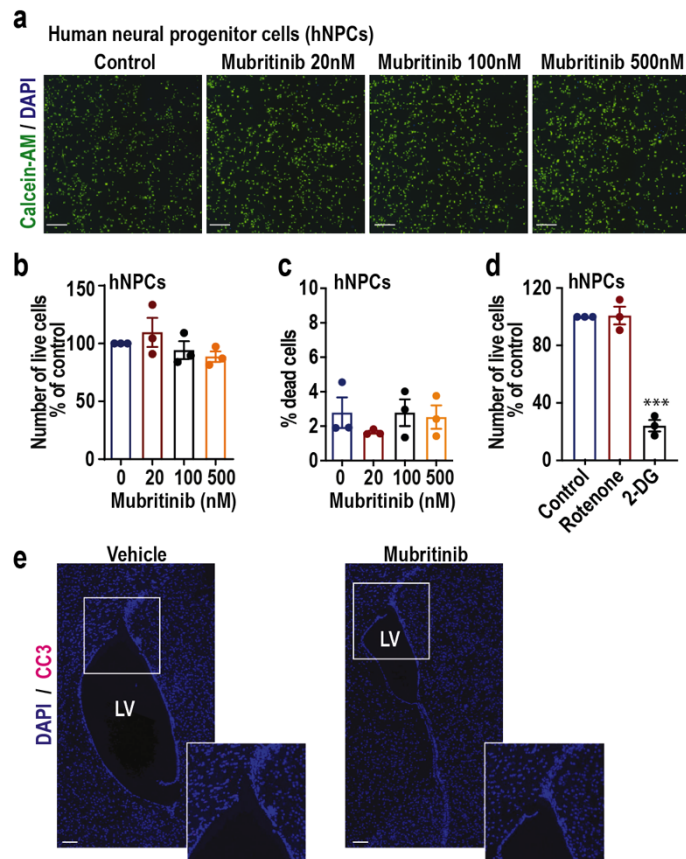
Appendix Fig S1. Mubritinib downregulates cell-cycle related pathways in BTSCs.

(a-b) Gene set enrichment analysis of deregulated genes in BTSC73 (a) and BTSC147 (b) treated with 500 nM of mubritinib for 24 h demonstrates enrichment for gene sets corresponding to cell-cycle related pathways.

(c-e) Gene set enrichment analysis of deregulated genes in BTSC147 treated with 500 nM mubritinib for 24 h demonstrates enrichment of gene sets corresponding to cyclin D1 associated events in G1 (c), SCF SKP2 mediated degradation of p27/p21 (d) and E2F mediated regulation of DNA replication (e).

(f) Cell cycle distribution was assessed by flow cytometry after PI staining in BTSC147 following 24 h of treatment with 500 nM of mubritinib. Data are presented as the means $\pm$ SEM, $n = 3$ independent biological experiments. Unpaired two-tailed t test. $*p_{G1} = 0.0128$, $**p_S = 0.0074$.

(g) BTSC53 and BTSC73 were treated for 72 h with 500 nM of mubritinib or vehicle control and subjected to immunoblotting using antibody against p-AMPK. Tubulin was used as loading control.



Appendix fig S2. Mubritinib does not impact normal non-oncogenic cells and has a safe profile *in vivo*.

(a-c) Human neural progenitor cells (hNPCs) were cultured in 2D monolayer on matrigel-coated plate and treated with increasing concentrations of mubritinib (0 to 500 nM) for 4 days and subjected to calcein-AM/DAPI double staining. Representative images are shown (a). Scale bars = 100 μ m. The number of live cells (calcein-AM positive) (b) and dead cells (DAPI positive) (c) were quantified by Fiji software.

(d) hNPCs were treated with 100 nM rotenone or 1mM 2-Deoxy-D-glucose (2-DG) and subjected to live cell counting by PI staining followed by flow cytometry after 7 days of treatment. Data are presented as the mean $\pm$ SEM. One-way ANOVA followed by Dunnett's test vs control. *** $p_{2-DG} = 2.6e-5$.

(e) Representative immunofluorescence images of cleaved caspase 3 (CC3) in the SVZ of mouse brains subjected to the indicated treatments are shown. Scale bars = 100 μ m.

BTSCs	Diagnosis	Age, y	Sex	IDH1	EGFR status	TP53	PTEN	NF1	CDKN2A	MGMT	References
12	GB-rec	59	M	wt	wt	mut	mut	NA	NA	U	(Cusulin <i>et al.</i> , 2015; Kelly <i>et al.</i> , 2009)
25	GB	57	M	wt	wt	mut	mut	NA	NA	U/M	(Cusulin <i>et al.</i> , 2015; Kelly <i>et al.</i> , 2009)
50	GB	61	M	wt	amplified	wt	mut	wt	homo del	M	(Cusulin <i>et al.</i> , 2015; Luchman <i>et al.</i> , 2014)
53	GB	59	M	wt	amplified / mut G598V	mut	wt	wt	homo del	M	(Cusulin <i>et al.</i> , 2015; Kelly <i>et al.</i> , 2009; Stechishin <i>et al.</i> , 2013)
73	GB	52	M	wt	amplified / VIII	mut	mut	NA	homo del	M	(Cusulin <i>et al.</i> , 2015; Kelly <i>et al.</i> , 2009; Stechishin <i>et al.</i> , 2013)
75	GB	74	M	wt	wt	wt	wt	wt	wt	U	(Cusulin <i>et al.</i> , 2015; Kelly <i>et al.</i> , 2009; Shen <i>et al.</i> , 2019; Stechishin <i>et al.</i> , 2013)
100	GB	63	M	wt	wt	wt	wt	NA	NA	U	(Cusulin <i>et al.</i> , 2015; Shen <i>et al.</i> , 2019)
119	GB-rec	69	F	wt	amplified / mut Y620C	mut	mut	mut	homo del	U/M	(Nguyen <i>et al.</i> , 2014; Shen <i>et al.</i> , 2019)
147	GB-rec	56	M	wt	VIII	mut	mut	wt	homo del	U	(Cusulin <i>et al.</i> , 2015; Stechishin <i>et al.</i> , 2013)
198	GB	52	F	wt	amplified	mut	mut	wt	NA	U/M	(Grinshtein <i>et al.</i> , 2016; Shen <i>et al.</i> , 2019)
P3	GB	NA	NA	wt	wt	wt	mut	wt	homo del	U	(Eskilsson <i>et al.</i> , 2016; Joseph <i>et al.</i> , 2022; Keunen <i>et al.</i> , 2011)

Appendix Table S1: Characterization of patient-derived BTSCs.

Mutant (mut) or wild-type (WT) status of genes frequently mutated in GB for the BTSC lines used in this study isolated from primary glioblastoma tumour (GB) or recurrent glioblastoma (GB-rec). VIII indicates EGFR variant III, homo del indicates a homozygous deletion, NA – not available; U indicates unmethylated; M indicates methylated.

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Analyte	Parent ion (m/z)	Transition ion (m/z)	Confirmation ion (m/z)	Cone voltage (V)	Collision energy (eV)
Mubritinib	469.2	252.3	224.2	30	25 / 20
[¹³ C, ² H ₃]-Sorafenib	469.3	274.2	256.2	50	30 / 25

Appendix Table S2. Mass spectrometer settings for the quantification of mubritinib by UPLC-MS/MS.