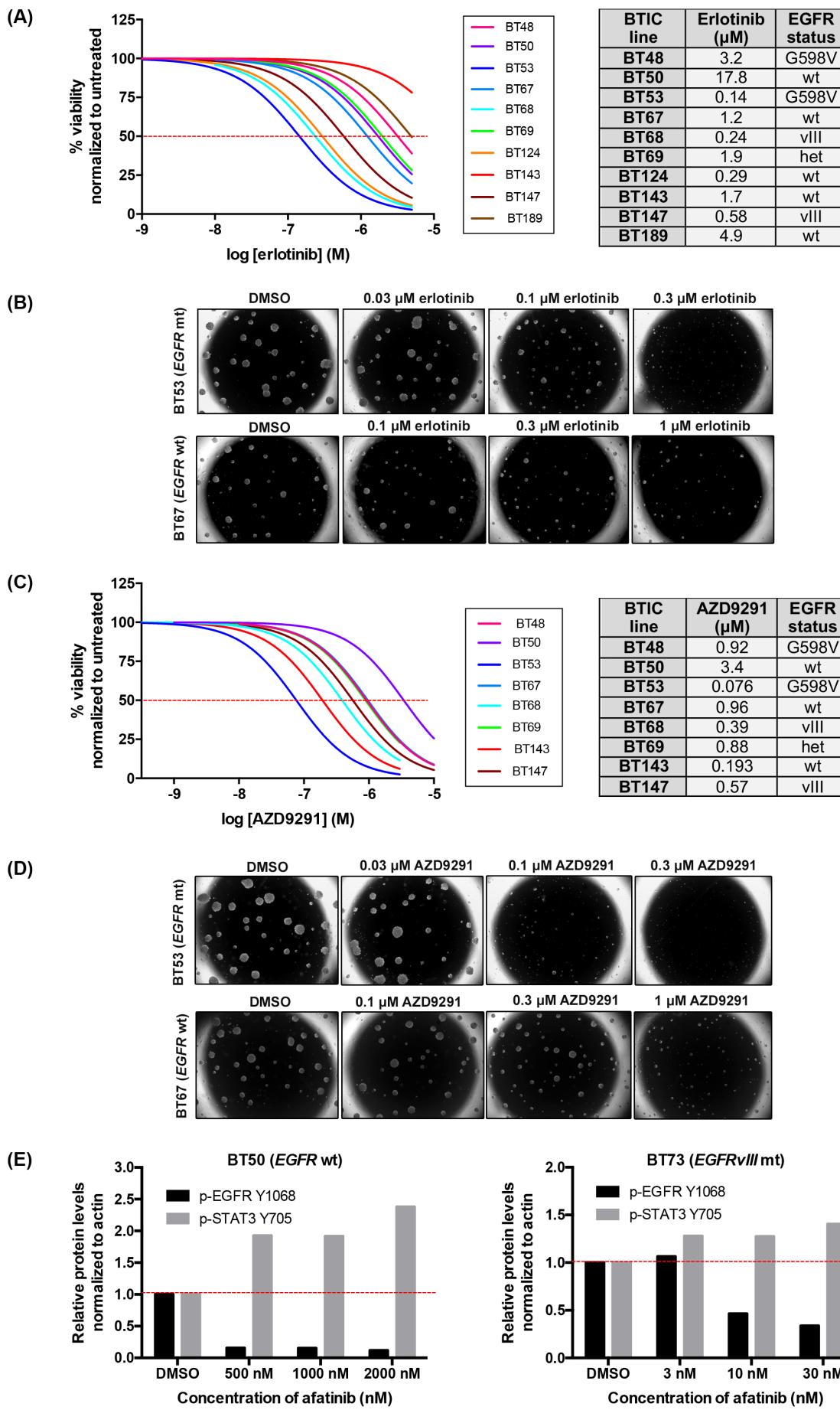


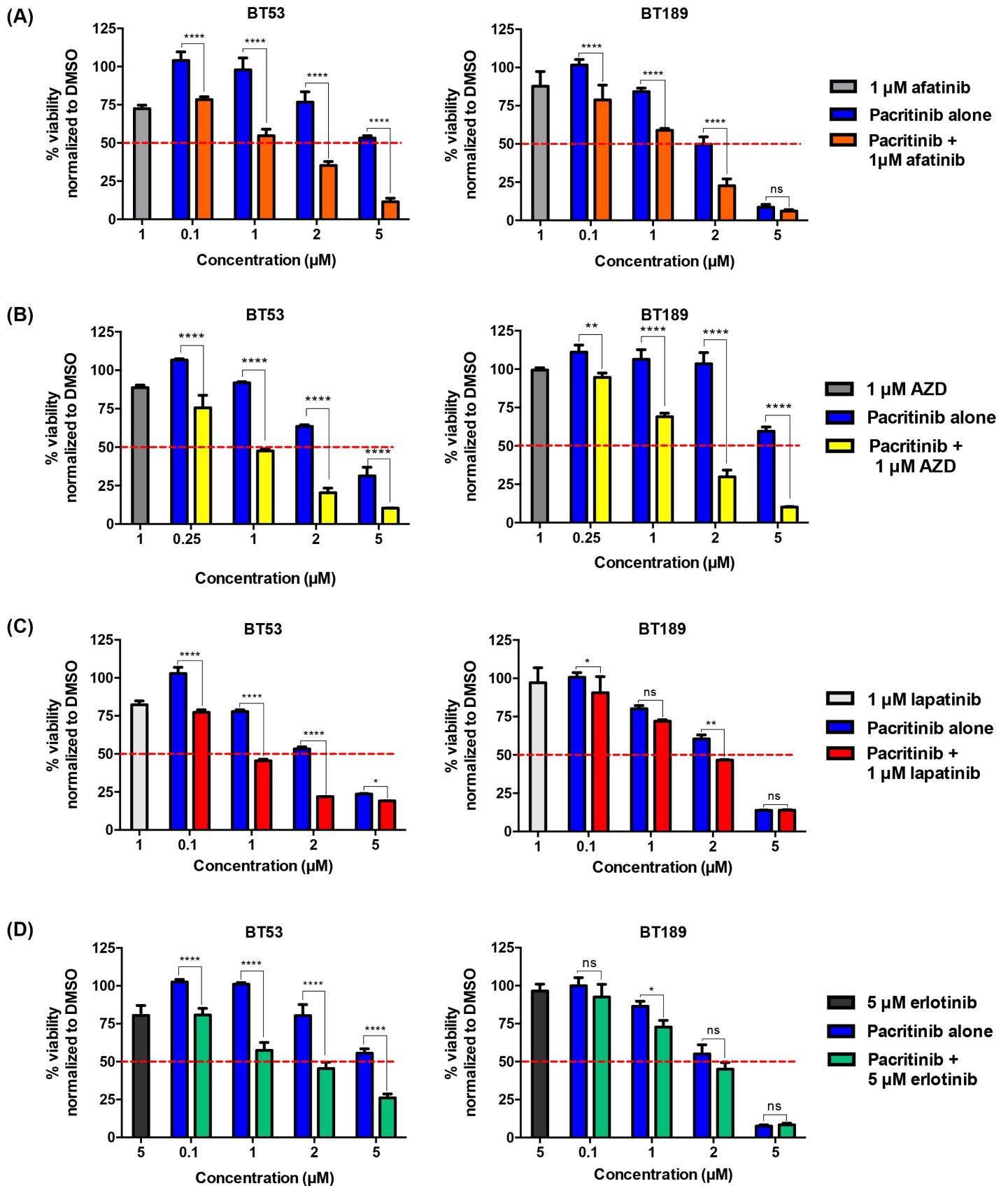
Supplemental Table 1: Identification of a diverse panel of GBM BTICs representative of the different mutational statuses of GBM patients.

BTIC Line	Diagnosis	Age	Sex	MGMT status	EGFR status	P53 status	PTEN status	IDH1 status	NF1 status	CDKN2A status
BT12	GBM-r	59	M	U	wt	wt	mut	wt	N/A	N/A
BT48	GBM	68	M	M	mut	wt	mut	wt	wt	homo del
BT50	GBM	61	M	M	wt	wt	mut	wt	wt	homo del
BT53	GBM	59	M	M	mut	mut	wt	wt	wt	homo del
BT67	GBM	44	M	M	wt	wt	mut	wt	mut/het del	homo del
BT68	GBM	59	M	U	vIII	wt	wt	wt	N/A	N/A
BT69	GBM	50	M	U/M	het	mut	mut	wt	N/A	homo del
BT73	GBM	52	M	M	vIII	mut	mut	wt	N/A	homo del
BT124	GBM	N/A	N/A	U	wt	mut	mut	wt	N/A	N/A
BT143	GBM-r	46	F	M	wt	mut	mut	wt	mut	homo del
BT147	GBM-r	56	M	U	vIII	mut	mut	wt	wt	homo del
BT189	GBM	55	F	U	wt	mut	mut	wt	wt	wt

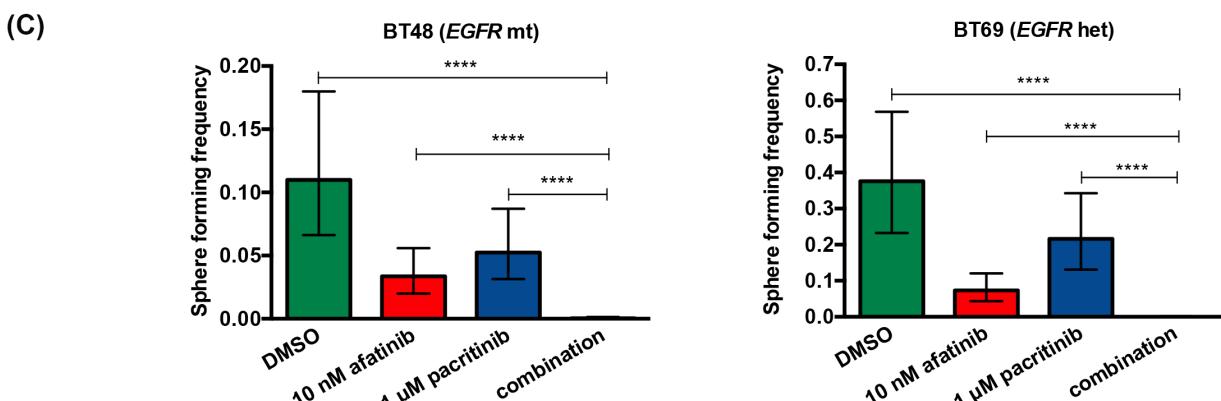
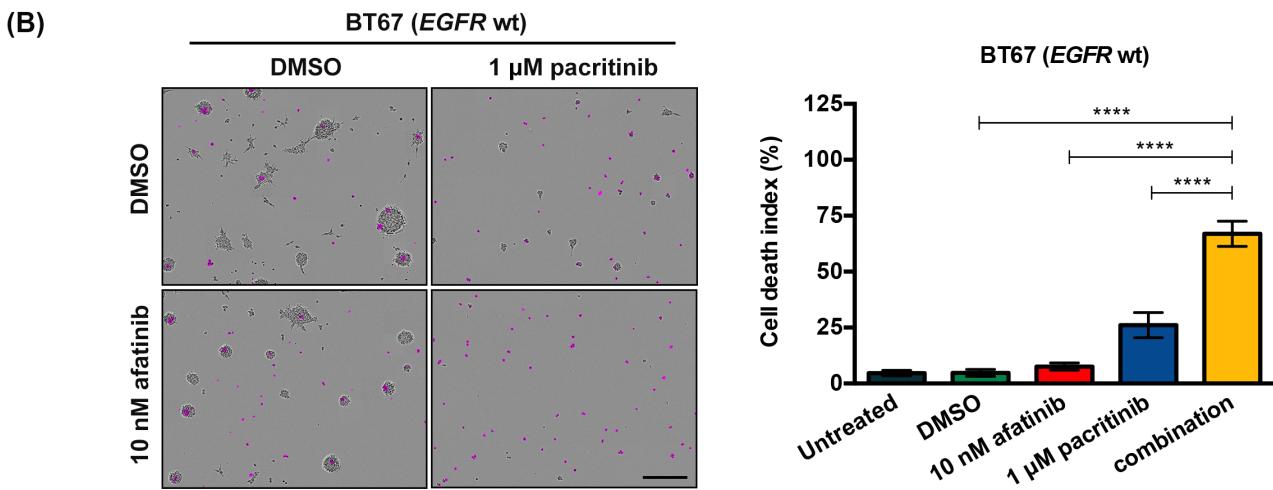
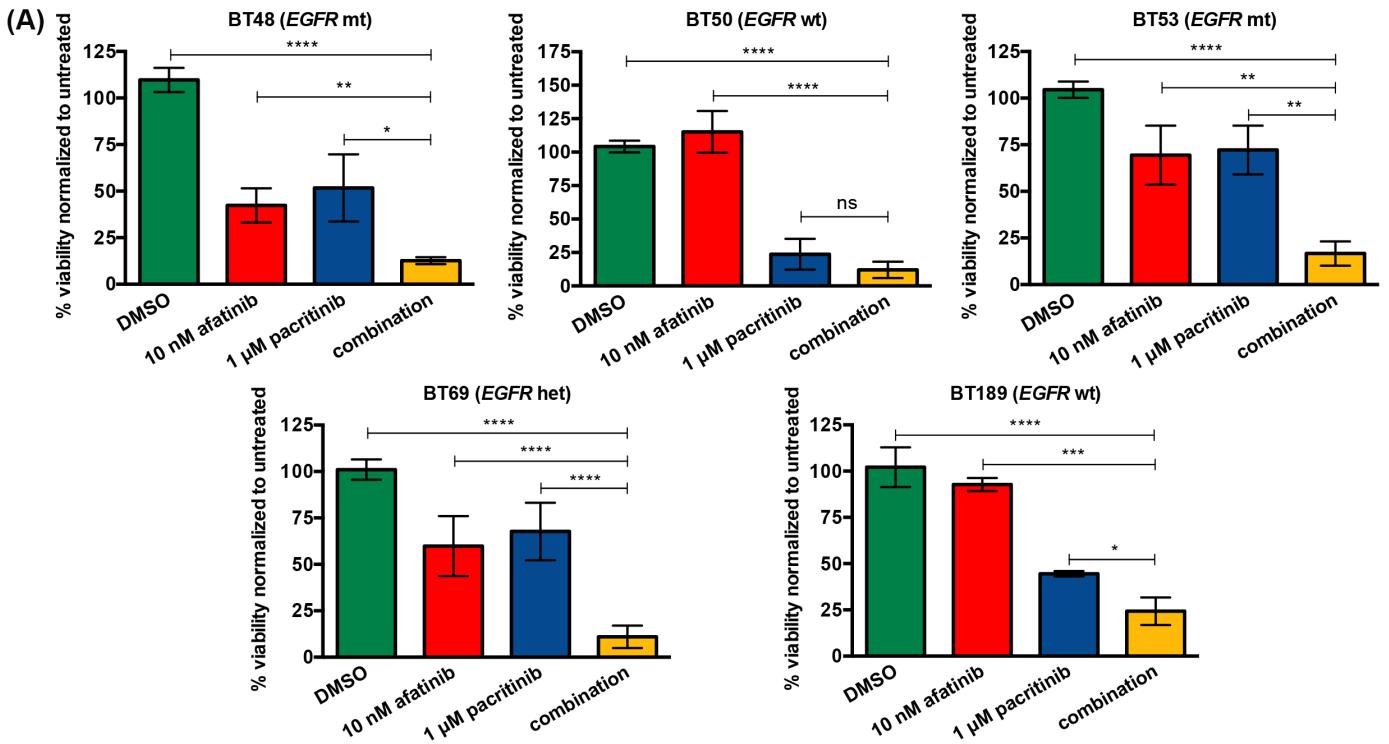
Common GBM molecular alterations including *MGMT* promoter methylation, *EGFR*, *PTEN*, *TP53*, *NF1*, *IDH1*, and *CDKN2A* mutation statuses for the 11 BTIC lines used in this study. Mut indicates mutant and wt indicates wild-type. U indicates unmethylated and M indicated methylated. U/M indicates hemi-methylation. vIII indicates *EGFR* variant III, an activating deletion characteristic of GBM, het del indicates a heterozygous deletion, homo del indicates a homozygous deletion, N/A indicates not available.



Supplemental Fig. 1

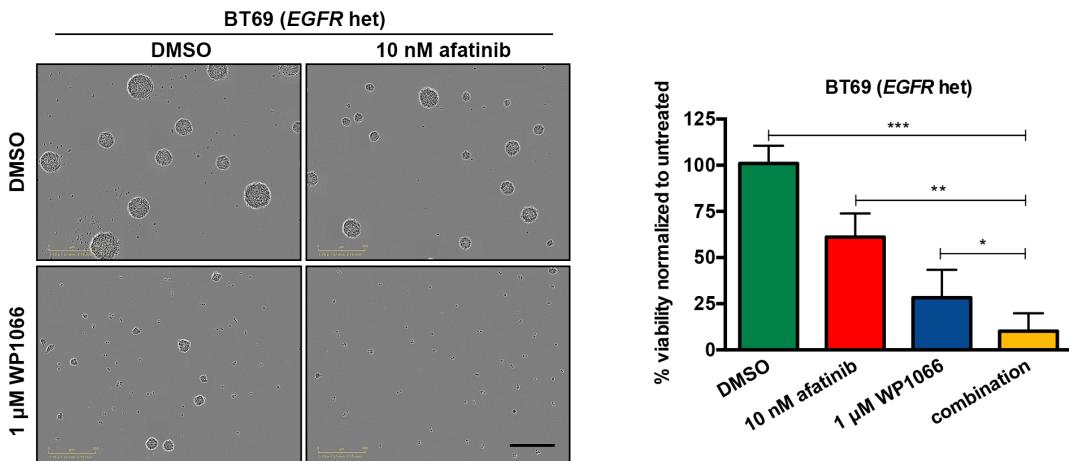


Supplemental Fig. 2

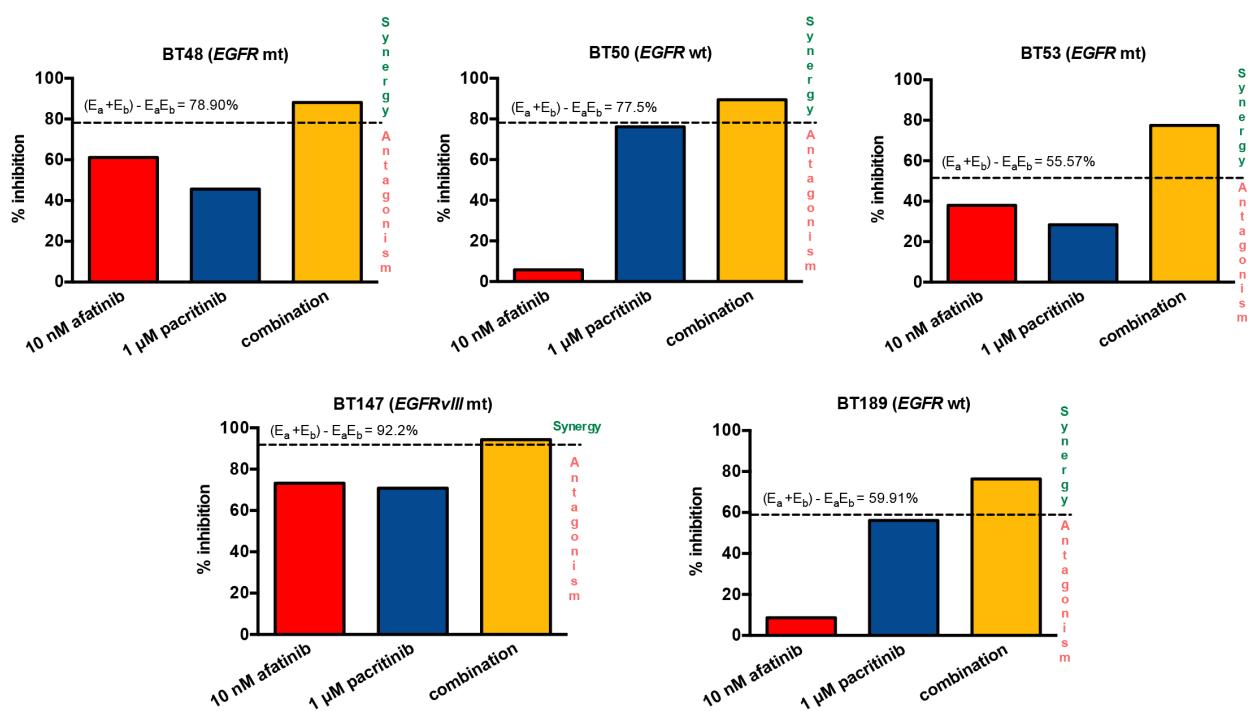


Supplemental Fig. 3

(A)



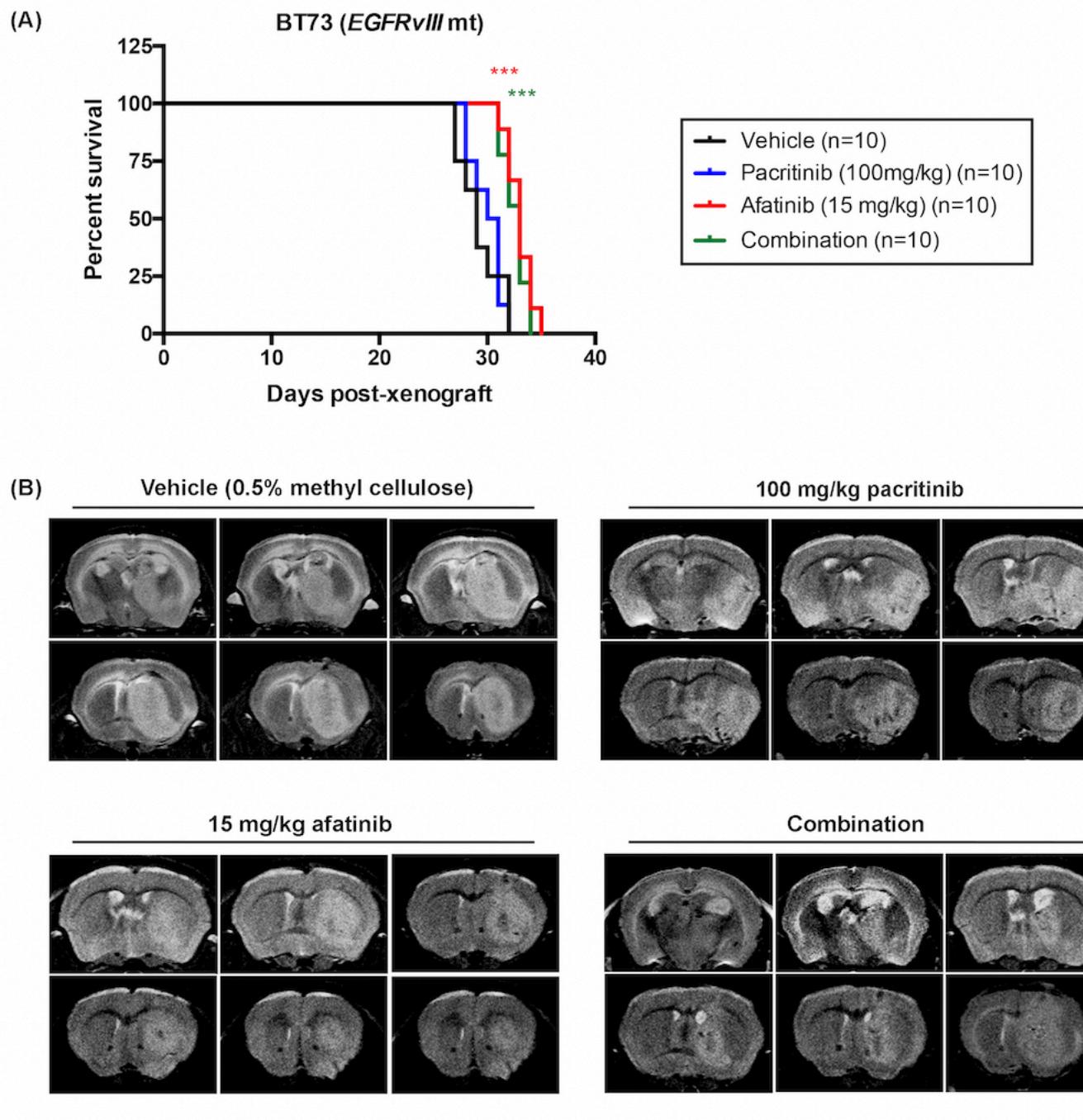
(B)



(C)

BT69 (EGFR het) erlotinib (μM)						BT147 (EGFRvIII mt) erlotinib (μM)							
		10	5	3	1	0			10	5	3	1	0
% inhibition pacritinib (μM)		85.0022841	80.3849223	78.2738674	70.8339955	-5.1306994			87.0287556	85.5860001	84.8352194	84.9468128	38.5645045
0.1		77.5243282	65.4335772	59.6033463	51.4351889	-14.877944			76.7758572	74.9831207	70.8651323	67.2372072	-11.39176
0.0		76.5056222	59.7428594	54.6965544	42.1872791	-8.7372337			76.7923093	71.3976478	67.3668418	61.2299209	-5.7182979
		68.7835636	58.7946963	50.6916304	38.5794293	0			76.0670927	75.2702882	66.7276967	64.7736307	0
Excess over bliss pacritinib (μM)		10	5	3	1	0	erlotinib (μM)		10	5	3	1	0
1		17.820342	23.7043463	30.1121012	35.4058712	0			1.73205579	0.77882109	5.27622379	6.58830734	0
0.3		13.3851286	12.7693831	16.2477875	21.9938779	0			3.43514385	2.52998203	7.92773668	6.4764801	0
0.1		10.4495115	4.54836681	8.3131114	8.9743086	0			2.09377154	-2.4585218	2.54175454	-1.529361	0
0		0	0	0	0	0			0	0	0	0	0

Supplemental Fig. 4



Supplemental Fig. 5