Figure	Panel	Cell line #1	Cell line #2	R	R^2	Adjusted R^2	Std. Error of the Estimate	Change Statistics				
								R^2 Change	F Change	df1	df2	Sig. F Change
1	E	*IDH1WT/WT* HCT116 P3	*IDH1WT/R132H* HCT116 P3	,894[b]	,800	,781	,22907	,295	31,706	2	43	,000
1	F	*IDH1WT/WT* HCT116 0 uM NAC	*IDH1WT/WT* HCT116 5 uM NAC	,915[b]	,836	,823	,71222	,009	1,418	2	49	,252
1	F	*IDH1WT/R132H* HCT116 0 uM NAC	*IDH1WT/R132H* HCT116 5 uM NAC	,914[b]	,836	,822	,62748	,142	21,265	2	49	,000
1	F	*IDH1WT/WT* HCT116 5 uM NAC	*IDH1WT/R132H* HCT116 5 uM NAC	,914[b]	,835	,822	,74231	,001	,167	2	49	,847
1	H	*IDH1WT/WT* HCT116 P30	*IDH1WT/R132H* HCT116 P30	,924[b]	,853	,840	,29245	,131	20,554	2	46	,000
5	A	*IDH1WT/WT* HCT116 0 nM AGI-5198	*IDH1WT/R132H* HCT116 0 nM AGI-5198	,898[b]	,806	,790	,62021	,096	12,150	2	49	,000
5	A	*IDH1WT/R132H* HCT116 0 nM AGI-5198	*IDH1WT/R132H* HCT116 10 nM AGI-5198 72 h	,956[b]	,913	,906	,30875	,005	1,407	2	48	,255
5	A	*IDH1WT/R132H* HCT116 0 nM AGI-5198	*IDH1WT/R132H* HCT116 25 nM AGI-5198 72 h	,914[b]	,835	,822	,47958	,022	3,298	2	49	,045
5	A	*IDH1WT/R132H* HCT116 0 nM AGI-5198	*IDH1WT/R132H* HCT116 100 nM AGI-5198 72 h	,950[b]	,903	,895	,35328	,032	8,099	2	49	,001
5	A	*IDH1WT/R132H* HCT116 0 nM AGI-5198	*IDH1WT/R132H* HCT116 400 nM AGI-5198 72 h	,941[b]	,886	,876	,42533	,070	15,110	2	49	,000
5	A	*IDH1WT/R132H* HCT116 0 nM AGI-5198	*IDH1WT/R132H* HCT116 800 nM AGI-5198 72 h	,868[b]	,753	,733	,88288	,125	12,419	2	49	,000
5	A	*IDH1WT/WT* HCT116 0 nM AGI-5198	*IDH1WT/R132H* HCT116 800 nM AGI-5198 72 h	,856[b]	,733	,712	1,01473	,009	,872	2	49	,424
5	A	*IDH1WT/WT* HCT116 0 nM AGI-5198	*IDH1WT/R132H* HCT116 400 nM AGI-5198 72 h	,907[b]	,823	,809	,65561	,005	,701	2	49	,501
5	A	*IDH1WT/WT* HCT116 0 nM AGI-5198	*IDH1WT/R132H* HCT116 100 nM AGI-5198 72 h	,910[b]	,828	,814	,61371	,027	3,864	2	49	,028
5	A	*IDH1WT/WT* HCT116 0 nM AGI-5198	*IDH1WT/R132H* HCT116 25 nM AGI-5198 72 h	,889[b]	,790	,772	,69059	,032	3,755	2	49	,030
5	A	*IDH1WT/WT* HCT116 0 nM AGI-5198	*IDH1WT/R132H* HCT116 10 nM AGI-5198 72 h	,910[b]	,828	,814	,59129	,067	9,310	2	48	,000
5	B	*IDH1WT/WT* HCT116 0 nM AGI-5198	*IDH1WT/WT* HCT116 25 nM AGI-5198 72 h	,869[b]	,755	,732	,31463	,002	,126	2	32	,882
5	B	*IDH1WT/WT* HCT116 0 nM AGI-5198	*IDH1WT/WT* HCT116 100 nM AGI-5198 72 h	,886[b]	,785	,765	,29018	,036	2,716	2	32	,081
5	B	*IDH1WT/WT* HCT116 0 nM AGI-5198	*IDH1WT/WT* HCT116 400 nM AGI-5198 72 h	,875[b]	,766	,744	,28977	,046	3,156	2	32	,056
5	C	U251 + IDH1R132H 0 nM AGI-5198	U251 + IDH1R132H 200 nM AGI-5198 72 h	,970[b]	,940	,935	,80890	,012	4,819	2	49	,012
5	C	U251 + IDH1R132H 0 nM AGI-5198	U251 + IDH1R132H 800 nM AGI-5198 72 h	,976[b]	,953	,949	,90083	,044	22,829	2	49	,000
5	C	U251 + IDH1R132H 200 nM AGI-5198 72 h	U251 + IDH1R132H 800 nM AGI-5198 72 h	,970[b]	,941	,937	1,06988	,015	6,289	2	49	,004
5	D	*IDH1WT/R132H* HCT116 0 nM AGI-5198	*IDH1WT/R132H* HCT116 100 nM AGI-5198 4 h	,933[b]	,870	,859	,35532	,000	,037	2	49	,964
5	D	*IDH1WT/R132H* HCT116 0 nM AGI-5198	*IDH1WT/R132H* HCT116 400 nM AGI-5198 4 h	,916[b]	,839	,826	,42895	,005	,713	2	49	,495

Supplementary Table S1. SPSS output statistics for comparisons between survival curves from colony-formation assays.

The model states the P value for the hypothesis that the data scatter was described best with one curve (for P < 0.05, the data scatter is best described with two curves
i.e., a statistically significant difference between the curve from cell line #1 and the curve from cell line #2)