

CG200745, an HDAC inhibitor, induces anti-tumour effects in cholangiocarcinoma cell lines via miRNAs targeting the Hippo pathway

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*Equal contribution.

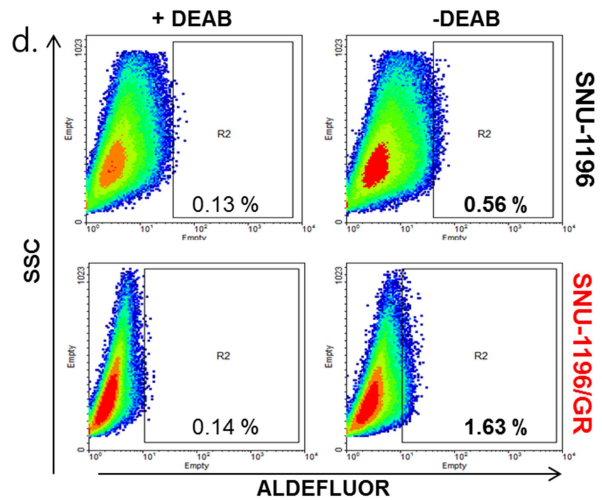
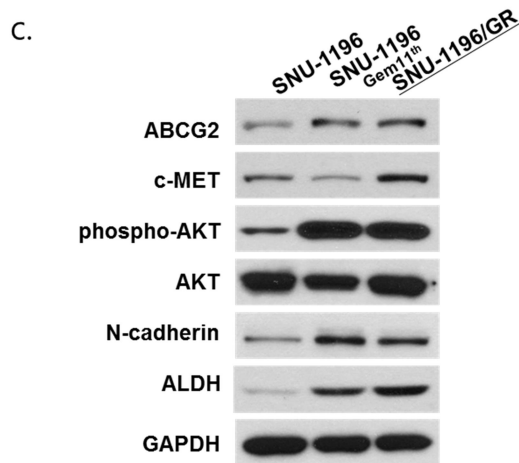
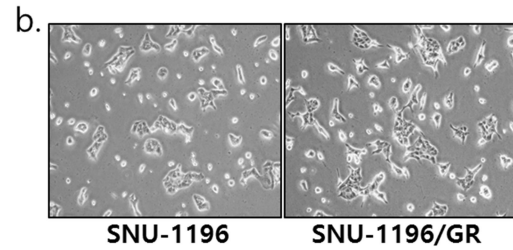
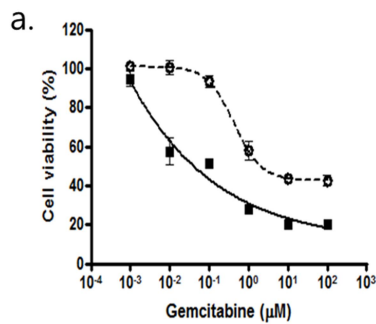
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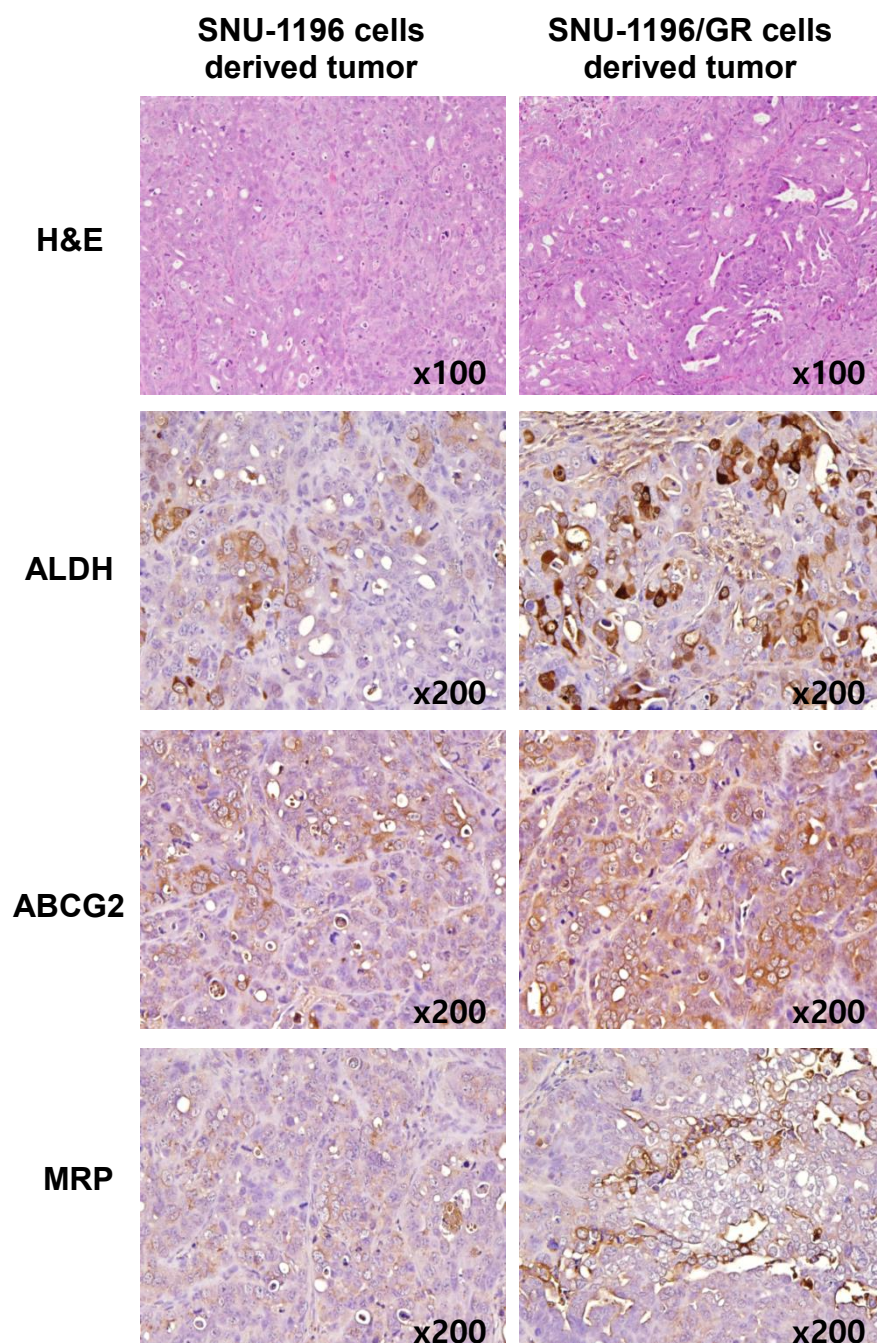
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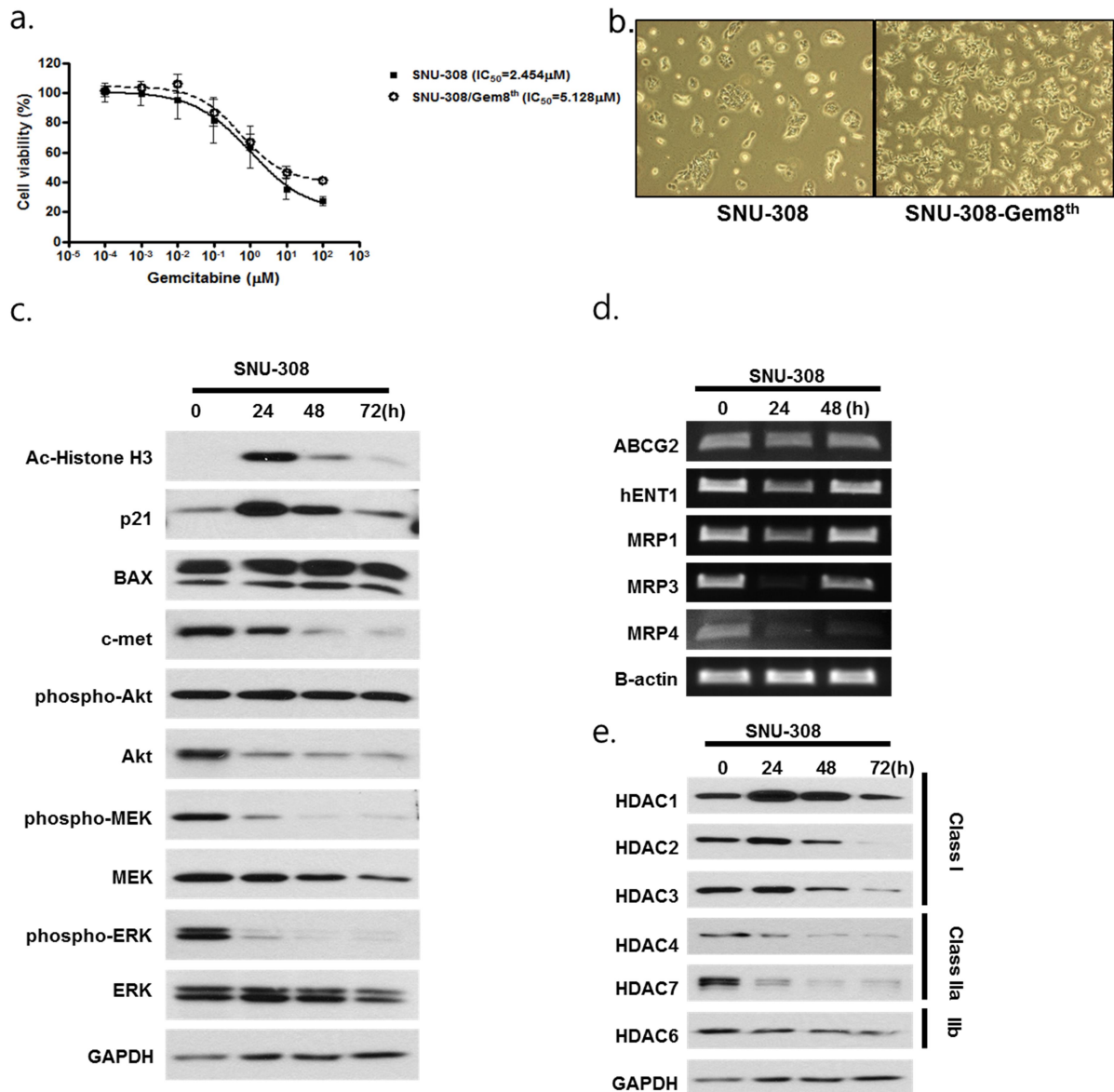
Supplementary Figure 1. SNU-1196 cells were treated with increasing doses of gemcitabine to establish a GR cell line (SNU-1196/GR). (a, b) Cell viability was assessed with the MTT assay; gemcitabine IC₅₀ values were 0.038 and 2.291 μM, respectively. (c) Elevated protein levels of c-MET, phospho-AKT, N-cadherin, and ALDH were analysed by western blotting. (d) ALDH enzymatic activity was measured with the ALDEFLUOR kit; cells treated with the specific ALDH inhibitor diethyliminobenzaldehyde served as a negative control. ALDH activity was also measured by flow cytometry.



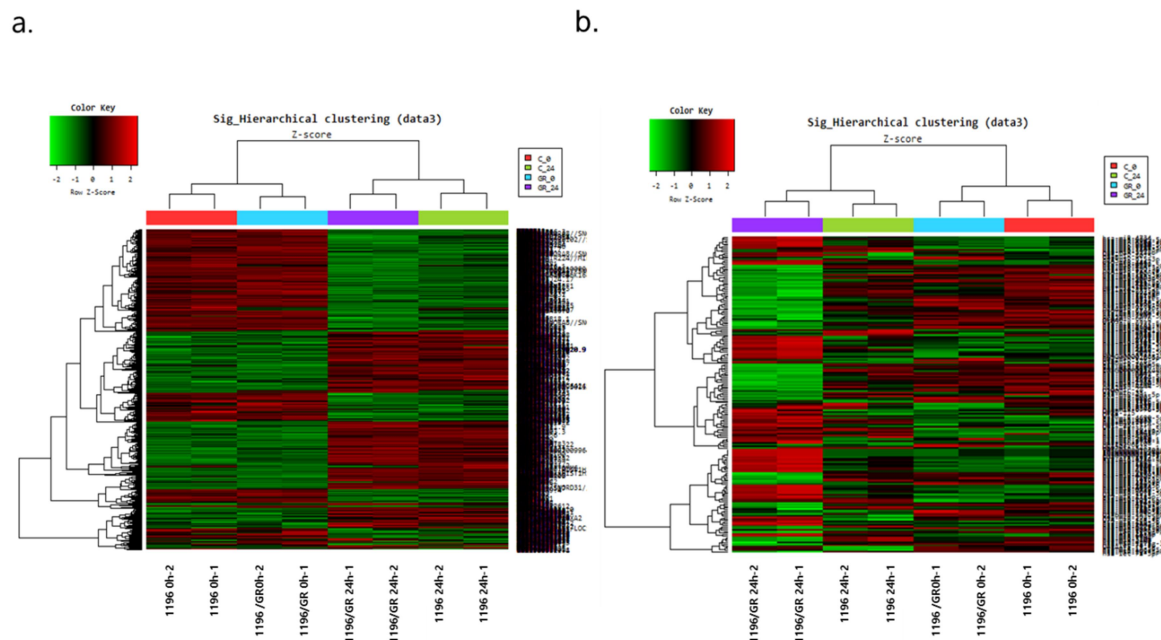
Supplementary Figure 2. Detection of proteins differentially expressed in tumours derived from SNU-1196 and SNU-1196/GR cells, as detected by immunohistochemistry using antibodies against ALDH, ABCG2, and MRP4.



Supplementary Figure 3. Establishment of a GR cell line. (a, b) SNU-308 was treated with increasing doses of gemcitabine to establish a GR cell line (SNU-308/GR). Cell viability was assessed with the MTT assay; gemcitabine IC₅₀ values were 2.454 and 5.128 μM, respectively. (c–e) CG200745 (IC₅₀) induced the expression of apoptosis-related protein p21 and inhibited that of c-MET, AKT, phospho-MEK, and phospho-ERK, as determined by western blotting (c); inhibited the expression of multidrug resistance genes, as determined by PCR (d); and altered the expression level of HDAC isozyme, as determined by western blotting (e).



Supplementary Figure 5. (a) Gene expression profiles were analysed using Affymetrix Human Genechip 2.0 and differentially expressed genes (cut-off, fold change ≥ 2 , $P < 0.05$) were identified by cluster analysis-hierarchical clustering (Euclidean method, complete linkage). Based on clustering results, overall gene expression patterns in SNU-1196 and SNU-1196/GR either pre- (0 h) or post- (24 h) treatment with CG200745 were hierarchically organised. (b) MiRNA expression profiles were analysed with a human Affymetrix GeneChip miRNA 4.0 Array and differentially expressed miRNAs (cut-off, fold change ≥ 1.5 , $P < 0.05$) were identified by cluster analysis-hierarchical clustering (Euclidean method, complete linkage). Based on clustering results, overall miRNA expression patterns in SNU-1196 and SNU-1196/GR either pre- (0 h) or post- (24 h) treatment with CG200745 were hierarchically organised.



Supplementary Table 1. Combination index (CI) and dose reduction index (DRI) of CG200745 in SNU-1196 cells

a.

SNU-1196	Gemcitabine (μM)		Cisplatin (μM)		5-FU (μM)		Oxaliplatin (μM)	
Single (IC_{50})	0.038		4.90		104.713		3.236	
CG200745 (μM)	0.25	0.50	0.25	0.50	0.25	0.50	0.25	0.50
Combi (IC_{50})	0.002	<0.0001	0.56	0.11	1.86	0.01	1.59	0.25
Dose reduction index	19	>100	8.75	44.55	56.29	>100	2.04	12.94

b. Gemcitabine and CG200745

Combination Index	0.489	0.281	0.273	0.281	0.341
Gemcitabine(μM)	0.001	0.010	0.100	1.000	10.000
CG200745(μM)	0.250	0.250	0.250	0.250	0.250
Combination Index	0.570	0.518	0.444	0.508	0.594
Gemcitabine(μM)	0.001	0.010	0.100	1.000	10.000
CG200745(μM)	0.500	0.500	0.500	0.500	0.500

Cisplatin and CG200745

Combination Index	0.567	0.447	0.635
Cisplatin(μM)	0.400	2.000	10.000
CG200745(μM)	0.250	0.250	0.250
Combination Index	1.039	0.591	0.946
Cisplatin(μM)	0.400	2.000	10.000
CG200745(μM)	0.500	0.500	0.500

5-FU and CG200745

Combination Index	0.627	0.454	0.342	0.432	0.770
5-FU(μM)	0.100	1.000	10.000	100.000	1000.000
CG200745(μM)	0.250	0.250	0.250	0.250	0.250
Combination Index	0.665	0.494	0.449	0.424	0.718
5-FU(μM)	0.100	1.000	10.000	100.000	1000.000
CG200745(μM)	0.500	0.500	0.500	0.500	0.500

Oxaliplatin and CG200745

Combination Index	0.84091	0.92858	0.78218	0.77119
Oxaliplatin(μM)	0.080	0.400	2.000	10.000
CG200745(μM)	0.250	0.250	0.250	0.250
Combination Index	0.72283	0.62079	0.63581	0.77325
Oxaliplatin(μM)	0.080	0.400	2.000	10.000
CG200745(μM)	0.500	0.500	0.500	0.500

c. Gemcitabine plus Cisplatin and CG200745

Combination Index	0.763	0.583	0.494	0.468	0.545	0.654
Gemcitabine(μM)	0.00313	0.00625	0.01250	0.02500	0.05000	0.10000
Cisplatin(μM)	0.12813	0.25625	0.51240	1.02500	2.05000	4.10000
CG200745(μM)	0.50000	0.50000	0.50000	0.50000	0.50000	0.50000

CI>1 indicates antagonism; CI=1 indicates an additive effect; and CI<1 indicates synergism. DRI>1 is favorable dose reduction that lead to toxicity reduction.

Supplementary Table 2. Combination index (CI) and dose reduction index (DRI) of CG200745 in SNU-308 cells

a.

SNU-308	Gemcitabine (μM)		Cisplatin (μM)		5-FU (μM)		Oxaliplatin (μM)	
Single (IC_{50})	1.413		3.80		74.131		6.918	
CG200745 (μM)	0.50	1.00	0.50	1.00	0.50	1.00	0.50	1.00
Combi (IC_{50})	0.32	0.13	2.34	1.82	44.67	17.78	4.57	2.82
Dose reduction index	4.47	11.21	1.62	2.09	1.66	4.17	1.51	2.45

b. Gemcitabine and CG200745

Combination Index	0.559	0.369	0.244	0.640
Gemcitabine(μM)	0.001	0.010	0.100	1.000
CG200745(μM)	0.500	0.500	0.500	0.500
Combination Index	0.723	0.507	0.312	0.614
Gemcitabine(μM)	0.001	0.010	0.100	1.000
CG200745(μM)	1.000	1.000	1.000	1.000

Cisplatin and CG200745

Combination Index	1.474	1.005	1.001	0.484	0.475
Cisplatin(μM)	0.001	0.010	0.100	1.000	10.000
CG200745(μM)	0.500	0.500	0.500	0.500	0.500
Combination Index	1.016	1.381	0.772	0.366	0.484
Cisplatin(μM)	0.001	0.010	0.100	1.000	10.000
CG200745(μM)	1.000	1.000	1.000	1.000	1.000

5-FU and CG200745

Combination Index	0.892	0.772	1.096	1.313	0.589
5-FU(μM)	0.001	0.010	0.100	1.000	10.000
CG200745(μM)	1.000	1.000	1.000	1.000	1.000

Oxaliplatin and CG200745

Combination Index	0.907	0.914	0.825	0.887	0.233
Oxaliplatin(μM)	0.010	0.100	1.000	10.000	10.000
CG200745(μM)	1.000	1.000	1.000	1.000	1.000

c. Gemcitabine plus Cisplatin and CG200745

Combination Index	0.818	0.696	0.478	0.408	0.301
Gemcitabine(μM)	0.063	0.125	0.250	0.500	1.000
Cisplatin(μM)	0.144	0.288	0.575	1.150	2.300
CG200745(μM)	1.000	1.000	1.000	1.000	1.000

CI>1 indicates antagonism; CI=1 indicates an additive effect; and CI<1 indicates synergism. DRI>1 is favorable dose reduction that lead to toxicity reduction.

Supplementary Table 3. Combination index (CI) and dose reduction index (DRI) of CG200745 in SNU-1196/GR

a.

SNU-1196/GR	Gemcitabine (μM)		5-FU (μM)		
Single (IC_{50})	2.291		229.0		
CG200745 (μM)	0.25	0.50	0.25	0.50	1.00
Combi (IC_{50})	0.21	0.05	35.48	6.76	0.42
Dose reduction index	10.91	45.82	6.45	33.88	>100

b. Gemcitabine and CG200745

Combination Index	0.637	0.589	0.279	0.083	0.149	1.140
Gemcitabine(μM)	0.001	0.010	0.100	1.000	10.000	100.000
CG200745(μM)	0.250	0.250	0.250	0.250	0.250	0.250
Combination Index	0.613	0.618	0.206	0.114	0.151	0.937
Gemcitabine(μM)	0.001	0.010	0.100	1.000	10.000	100.000
CG200745(μM)	0.500	0.500	0.500	0.500	0.500	0.500

5-FU and CG200745

Combination Index	0.803	0.620	0.478	0.356	0.606	0.785
5-FU(μM)	0.010	0.100	1.000	10.000	100.000	1000.000
CG200745(μM)	0.250	0.250	0.250	0.250	0.250	0.250
Combination Index	1.062	1.037	0.652	0.336	0.234	0.552
5-FU(μM)	0.010	0.100	1.000	10.000	100.000	1000.000
CG200745(μM)	0.500	0.500	0.500	0.500	0.500	0.500
Combination Index	0.856	0.757	0.522	0.221	0.177	0.427
5-FU(μM)	0.010	0.100	1.000	10.000	100.000	1000.000
CG200745(μM)	1.000	1.000	1.000	1.000	1.000	1.000

CI>1 indicates antagonism; CI=1 indicates an additive effect; and CI<1 indicates synergism. DRI>1 is favorable dose reduction that lead to toxicity reduction.

Supplementary Table 4. List of primers used in PCR analysis.

Gene	Sense	Antisense
<i>ABCG2</i>	TATGAGTGGCTTATCCTGCT	CACTGATCCTTCCATCTTGT
<i>hENT1</i>	GACAACCAGTCACCAGCCTCAG	AGAGCATCCAGCTGCACCTTCA
<i>MRP1</i>	CTGACAAGCTAGACCATGAATGT	TCACACCAAGCCGGCGTCTTT
<i>MRP3</i>	GGACCCTGCGCATGAACCTG	AGGCAAGTCCAGCATCTCTGG
<i>MRP4</i>	GGATCCAAGAACTGATGAGTTAAT	TCACAGTGCTGTCTCGAAAATAG
<i>MRP5</i>	GCTGTTCAAGTGGCACTGTCAG	TCAGCCCTTGACAGCGACCTT
<i>β-Actin</i>	GGCATCCTCACCTGAAGTA	GGGGTGTGAAGGTCTCAA

Supplementary Table 5. List of antibodies used in western blotting and immunohistochemistry

Primary antibodies	Supplier	Species	Type
Acetylated histone h3	Santa Cruz Biotechnology (Santa Cruz, CA)	Rabbit	polyclonal
C-Met	Santa Cruz Biotechnology	Rabbit	polyclonal
Caspase-3	Santa Cruz Biotechnology	Mouse	monoclonal
p21	Santa Cruz Biotechnology	Rabbit	polyclonal
BAX	Santa Cruz Biotechnology	Rabbit	polyclonal
AKT	Santa Cruz Biotechnology	Rabbit	polyclonal
ERK	Santa Cruz Biotechnology	Rabbit	polyclonal
MRP4	Santa Cruz Biotechnology	Mouse	monoclonal
HDAC7	Santa Cruz Biotechnology	Mouse	monoclonal
GAPDH	Santa Cruz Biotechnology	Mouse	monoclonal
SMAD3	Santa Cruz Biotechnology	Mouse	monoclonal
NOTCH3	Santa Cruz Biotechnology	Rabbit	polyclonal
HES5	Santa Cruz Biotechnology	Goat	polyclonal
ABCG2	Santa Cruz Biotechnology	Rabbit	polyclonal
Phospho-AKT	Cell Signaling Technology (Danvers, MA)	Rabbit	polyclonal
Phospho-MEK	Cell Signaling Technology	Rabbit	polyclonal
MEK	Cell Signaling Technology	Rabbit	polyclonal
Phospho-ERK	Cell Signaling Technology	Rabbit	polyclonal
HDAC1	Cell Signaling Technology	Mouse	monoclonal
HDAC2	Cell Signaling Technology	Mouse	monoclonal
HDAC3	Cell Signaling Technology	Mouse	monoclonal
HDAC4	Cell Signaling Technology	Rabbit	polyclonal
HDAC6	Cell Signaling Technology	Rabbit	polyclonal
YAP	Cell Signaling Technology	Rabbit	polyclonal
AXL	Cell Signaling Technology	Rabbit	polyclonal
PARP	Cell Signaling Technology	Rabbit	polyclonal
TEAD4	Abcam (Cambridge, UK)	Rabbit	polyclonal
N-cadherin	Abcam	Mouse	monoclonal
ALDH	BD Transduction Laboratories (San Jose, CA)	Mouse	monoclonal
TGF- β 2	Proteintech (Rosemont, IL)	Rabbit	polyclonal
Cleaved-Caspase 3	Cell Signaling Technology	Rabbit	polyclonal