

Supplementary data for Molecular Pharmacology**Mechanism of the anti-proliferative activity of some naphthalene diimide G-quadruplex ligands**

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name	nucleotide type	sequence	cation	Tm [°C]
F21T	G4-DNA	5'-FAM-GGG TTA GGG TTA GGG TTA GGG-TAMRA-3'	K ⁺	58
c-kit1	G4-DNA	5'-FAM-AGA GGG AGG GCG CTG GGA GGA GGG GCT-TAMRA-3'	K ⁺	52
c-kit2	G4-DNA	5'-FAM-CCC GGG CGG GCG CGA GGG AGG GGA GG-TAMRA-3'	K ⁺	55
bcl-2	G4-RNA	5'-FAM-GGG GGC CGU GGG GUG GGA GCU GGG G-TAMRA-3'	Na ⁺	68
hif-1α	G4-DNA	5'-FAM-GGG GAG GGG AGA GGG GGC GGG-TAMRA	K ⁺	78
T-loop	duplex-DNA	5'-FAM-TAT AGC TATA TTT TTT TATA GCT ATA-TAMRA-3'	K ⁺	54

Table S1: Nucleotide sequences used for FRET experiments.

<i>Correlatio n</i>	<i>Seed</i>	<i>target</i>	<i>mechanism of action</i>	<i>Common cell lines</i>	<i>Seed stand.</i>	<i>Target stand. Dev.</i>
0.82	BMSG-SH-3	NSC S342443	Inhibition of protein synthesis (Chan et al., 2004).	47	0.48	0.628
		NSC 753939	phyllanthoside			
0.79	BMSG-SH-3	NSC S707020	affects polyamine metabolism; derivatives interact with DNA (Kuska et al., 2002).	58	0.503	0.293
		NSC 753939	oxa-spermine homologue			
0.759	BMSG-SH-3	NSC S764111	Binds to GC-rich stretches in DNA minor groove and alters transcription and replication (Cheglakov et al., 2010).	47	0.511	0.783
		NSC 753939	olivomycin			
0.753	BMSG-SH-3	NSC S353076	DNA intercalating agent (Larne et al., 1987). Derivatives stabilise G-quadruplex DNA (Brooks et al., 2011).	47	0.48	0.517
		NSC 753939	ellipticinium derivative			
0.751	BMSG-SH-3	NSC S259969	Inhibits protein synthesis, decreases expression of cell cycle protein cyclin D (Wakita et al., 2001).	47	0.48	0.457
		NSC 753939	deoxybouvardin			
0.745	BMSG-SH-3	NSC S682066	Photochemical DNA cleavage (Saito et al., 1994). Analogs are G-quadruplex ligands (Izbicka et al., 1999).	59	0.499	0.53
		NSC 753939	A phosphonium-porphyrin			
0.74	BMSG-SH-3	NSC S259968	Inhibits protein synthesis (Zalacain et al., 1982).	47	0.48	0.683
		NSC 753939	bouvardin			
0.733	BMSG-SH-3	NSC S667645	n. a.	59	0.499	0.57
		NSC 753939				

			substituted acridine				
0.729	BMSG-SH-3	NSC S715654 NSC 753939	DNA binding agent (Spychala et al., 2008; Spychala et al., 2009).	56	0.51	0.371	
0.719	BMSG-SH-3	NSC S58514 NSC 753939	binds to GC rich DNA, inhibits RNA synthesis (NCI drug dictionary) chromomycin	47	0.48	0.413	
0.704	BMSG-SH-3	NSC 80467 NSC 753939	n. a.	46	0.473	0.716	
0.701	BMSG-SH-3	NSC S337766 NSC 753939	DNA intercalation, DNA single-strand breaks, DNA-protein crosslinking, inhibition of DNA replication (Elliott et al., 1989; NCI online drug dictionary). bisantrene hydrochloride	47	0.48	0.622	
0.7	BMSG-SH-3	NSC S35441 NSC 753939	DNA intercalator (Wakelin & Waring, 1974). a phenanthridinium derivative	59	0.501	0.513	
0.7	BMSG-SH-3	NSC S666489 NSC 753939	Inhibitor of histone methyltransferase Pr-Set7 (Kodama et al., 2011). Phosphonium aryl compound Analogs localise in mitochondria and decrease oxygen consumption (Millard et al., 2010).	59	0.499	0.522	
0.697	BMSG-SH-3	NSC S38270 NSC 753939	Binds to GC-rich stretches in DNA minor groove and alters transcription and replication (Cheglakov et al., 2010). olivomycin	59	0.499	0.45	
0.693	BMSG-SH-3	NSC S49842 NSC 753939	Mitotic block (Jordan et al., 1992). vinblastine sulfate	47	0.535	0.766	
0.693	BMSG-SH-3	NSC S682345	antiinflammatory C5a antagonist	59	0.499	0.265	

	NSC 753939	Aurantimycin	(Assem et al., 2008).				
0.692	BMSG-SH-3	NSC S219241	Analogs localise in mitochondria and	58	0.503	0.484	
	NSC 753939	a phosphonium aryl compound	decrease oxygen consumption (Millard et al., 2010).				
0.691	BMSG-SH-3	NSC S352671	analogs are DNA intercalators	56	0.51	0.572	
	NSC 753939	a phenanthridine	(Wakelin & Waring, 1974; De Stefano et al., 2009), and G-quadruplex ligands (Bai, 2008; Maiti & Kumar, 2010; Yang et al., 2010).				
0.688	BMSG-SH-3	NSC S673349	analogs are DNA intercalators	57	0.494	0.351	
	NSC 753939	anthraquinone derivative	(NSC724629)				
0.684	BMSG-SH-3	NSC S324646	Inhibitor of protein synthesis (Dawes, 1994).	53	0.506	0.454	
	NSC 753939	malformin a					
0.676	BMSG-SH-3	NSC S713157	blocks Ca ²⁺ channel (Berlinck et al., 1993).	54	0.514	0.205	
	NSC 753939	13,14,15-isocrambescidin 800					
0.672	BMSG-SH-3	NSC S246012	analogs are DNA intercalators	55	0.513	0.593	
	NSC 753939	Benzophenanthridinium	(Wakelin & Waring, 1974; De Stefano et al., 2009), and G-quadruplex ligands (Bai, 2008; Maiti & Kumar, 2010; Yang et al., 2010).				
0.669	BMSG-SH-3	NSC S674091	Topoisomerase-targeted anticancer drug, DNA binder (Spicer et al., 2002).	59	0.499	0.339	
	NSC 753939						
0.661	BMSG-SH-3	NSC S339570	Analogs localise in mitochondria and decrease oxygen consumption	51	0.518	0.334	
	NSC 753939						

		A phosphonium aryl derivative	(Millard et al., 2010).				
0.66	BMSG-SH-3	NSC S289922	Analogs localise in mitochondria and	59	0.499	0.668	
	NSC 753939	A phosphonium aryl derivative	decrease oxygen consumption (Millard et al., 2010).				
0.659	BMSG-SH-3	NSC S685703	Analog inhibits protein synthesis	59	0.499	0.314	
	NSC 753939	didemnin analog	(Ahuja et al., 2000).				
0.654	BMSG-SH-3	NSC S634791	DNA binder (Dimmock et al., 1992).	60	0.497	0.49	
	NSC 753939						
0.654	BMSG-SH-3	NSC S724629	a topoisomerase I inhibitor and DNA intercalator (Dezhenkova et al., 2008).	57	0.506	0.472	
	NSC 753939						
0.654	BMSG-SH-3	NSC S644614	n. a.	50	0.501	0.393	
	NSC 753939	a triarylethene					
0.652	BMSG-SH-3	NSC S632624	Derivatives stabilise G-quadruplex	47	0.479	0.389	
	NSC 753939	ellipticinium derivative	DNA (Brooks et al., 2011).				
0.651	BMSG-SH-3	NSC S368672	inhibits translation of hif-1α and c-myc through eIF4E phosphorylation pathway (Jin et al., 2008; Mata-Greenwood et al., 2002).	47	0.48	0.275	
	NSC 753939	quassinoïd					
0.648	BMSG-SH-3	NSC S176878	Analogs localise in mitochondria and	47	0.48	0.419	
	NSC 753939	A phosphonium aryl derivative	decrease oxygen consumption (Millard et al., 2010).				
0.648	BMSG-SH-3	NSC S179220	n. a.	59	0.496	0.544	
	NSC 753939	azothiazolium					

			derivative				
0.647	BMSG-SH-3	NSC S747162	mitochondria-targeted antioxidant,	60	0.497	0.346	
	NSC 753939	mitotempol	decreases reactive oxygen species and muscle differentiation (Lee et al., 2011)				
0.646	BMSG-SH-3	NSC S714616	nucleic acid binding (Kang et al., 2004; Spychala, 2008; Spychala, 2009).	57	0.507	0.207	
	NSC 753939						
0.641	BMSG-SH-3	NSC S710608	DNA binding (Spychala, 2008; Spychala, 2009).	59	0.499	0.303	
	NSC 753939						
0.638	BMSG-SH-3	NSC S665806	Analog inhibits protein synthesis	58	0.5	0.538	
	NSC 753939	didemnin analog	(Ahuja et al., 2000).				
0.638	BMSG-SH-3	NSC S226510	n. a.	47	0.483	0.192	
	NSC 753939						
0.637	BMSG-SH-3	NSC S694330	Analog inhibits protein synthesis	58	0.502	0.177	
	NSC 753939	didemnin analog	(Ahuja et al., 2000).				
0.637	BMSG-SH-3	NSC S711659	Derivatives stabilise G-quadruplex	57	0.505	0.326	
	NSC 753939	azaharman analog;	DNA (Brooks et al., 2011).				
		aza-ellipticine					
		derivative					
0.91	BMSG-SH-4	NSC:S753940	Analog of the seed, G-quadruplex	60	0.389	0.285	
	NSC 753938	Naphthalene diimide	ligand (Cuenca et al., 2008).				
		BMSG-SH-5					
0.584	BMSG-SH-4	NSC:S625331	DNA affinity (Nguyen et al., 1987).	46	0.382	0.336	
	NSC 753938	aza-ellipticine	Derivatives stabilise G-quadruplex DNA (Brooks et al., 2011).				

derivative							
0.577	BMSG-SH-4	NSC:S721046	n. a.	50	0.401	0.247	
		NSC 753938					
0.541	BMSG-SH-4	NSC:S656125	n. a.	49	0.375	0.213	
		NSC 753938					
0.531	BMSG-SH-4	NSC:S155694	Derivatives stabilise G-quadruplex	47	0.386	0.571	
		NSC 753938	DNA (Brooks et al., 2011). ellipticinium derivative				
0.525	BMSG-SH-4	NSC:S656575	n. a.	47	0.379	0.197	
		NSC 753938	Maleopimaric acid amide				
0.519	BMSG-SH-4	NSC:S268239	Analogs are G-quadruplex ligands	49	0.368	0.298	
		NSC 753938	(Manet et al., 2010; Clark et al., 2003). Inhibition of transcription by interacting with DNA in gene promoter regions (Banerjee & Mukhopadhyay, 2008; Sriram et al., 1991; Mansilla & Portugal, 2008).				
0.516	BMSG-SH-4	NSC:S298223	DNA alkylation (Gunz et al., 1996)	46	0.39	0.64	
		NSC 753938	CC-1065				
0.506	BMSG-SH-4	NSC:S671450	n. a.	54	0.36	0.168	
		NSC 753938					
0.505	BMSG-SH-4	NSC:S32938	antimalarial,	56	0.376	0.138	
		NSC 753938	a cresol derivative	n. a.			
0.91	BMSG-SH-5	NSC:S753938	Analog of the seed, G-quadruplex	60	0.285	0.389	
		NSC 753940	Naphthalene diimide				
			BMSG-SH-4				

0.593	BMSG-SH-5	NSC:S625331	DNA affinity (Nguyen et al., 1987).	46	0.263	0.336
	NSC 753940	aza-ellipticine derivative	Derivatives stabilise G-quadruplex DNA (Brooks et al., 2011).			
0.584	BMSG-SH-5	NSC:S58514	binds to GC rich DNA, inhibits RNA synthesis	47	0.275	0.413
	NSC 753940	Chromomycin A3	(http://www.cancer.gov/drugdictionary?cdrid=39184)			
0.577	BMSG-SH-5	NSC:S753939	Analog of the seed, G-quadruplex ligand (Hampel et al., 2010).	60	0.285	0.497
	NSC 753940	Naphthalene diimide				
		BMSG-SH-3				
0.572	BMSG-SH-5	NSC:S155694	Derivatives stabilise G-quadruplex DNA (Brooks et al., 2011).	47	0.275	0.571
	NSC 753940	ellipticinium derivative				
0.567	BMSG-SH-5	NSC:S325319	inhibits protein synthesis via binding to complexes between ribosomes and the elongation factor EF-1 α , which prevents binding to elongation factor EF-2 (Ahuja et al., 2000).	55	0.287	0.346
	NSC 753940	didemnin B				
0.547	BMSG-SH-5	NSC:S705956	Inhibitor of translation initiation, causes G2/M block (Cencic et al., 2009).	57	0.284	0.121
	NSC 753940	didesmethylrocaglamide				
0.537	BMSG-SH-5	NSC:S254681	Analogs are G-quadruplex ligands (Manet et al., 2010; Clark et al., 2003). Inhibition of transcription by interacting with DNA in gene promoter regions (Banerjee & Mukhopadhyay, 2008; Sriram et al., 1991; Mansilla & Portugal, 2008).	50	0.284	0.818
	NSC 753940	12-iminodaunorubicin (an anthracyclin)				

0.535	BMSG-SH-5 NSC 753940	NSC:S265450 nogalomycin c (an anthracyclin)	Analogs are G-quadruplex ligands (Manet et al., 2010; Clark et al., 2003). Inhibition of transcription by interacting with DNA in gene promoter regions (Banerjee & Mukhopadhyay, 2008; Sriram et al., 1991; Mansilla & Portugal, 2008).	47	0.275	0.43
0.53	BMSG-SH-5 NSC 753940	NSC:S259968 bouvardin	Inhibits protein synthesis (Zalacain et al., 1982).	51	0.281	0.563
0.526	BMSG-SH-5 NSC 753940	NSC:S259969 deoxybouvardin	Inhibits protein synthesis, decreases expression of cell cycle protein cyclin D (Wakita et al., 2001).	47	0.275	0.457
0.523	BMSG-SH-5 NSC 753940	NSC:S342443 Phyllantoside, S3'-desacetyl-	Analogs inhibit protein synthesis (Chan et al., 2004).	47	0.275	0.628
0.521	BMSG-SH-5 NSC 753940	NSC:S368672 quassinoïd	inhibits translation of <i>hif-1α</i> and <i>c-myc</i> through eIF4E phosphorylation pathway (Jin et al., 2008; Mata-Greenwood et al., 2002).	47	0.275	0.275
0.52	BMSG-SH-5 NSC 753940	NSC:S351710 2-methylellipticiniumiodide	Derivatives stabilise G-quadruplex DNA (Brooks et al., 2011).	47	0.275	0.62
0.517	BMSG-SH-5 NSC 753940	NSC:S131547 tubulosine	disrupts mitosis (Farjaudon et al., 1988).	59	0.286	0.458
0.516	BMSG-SH-5 NSC 753940	NSC:S672658	n. a.	54	0.283	0.236

			a peptide				
0.51	BMSG-SH-5	NSC:S83193	n. a.	60	0.285	0.193	
	NSC 753940	C.I. basic blue 1					
0.508	BMSG-SH-5	NSC:S671035	n. a.	58	0.274	0.141	
	NSC 753940						
0.502	BMSG-SH-5	NSC:S670163	Analogs localise in mitochondria and	55	0.28	0.478	
	NSC 753940	an aryl phosphonium compound	decrease oxygen consumption (Millard et al., 2010).				

Table S2: COMPARE study of compounds BMSG-SH-3 – 5 against NCI compound library. References for the mechanisms of action of correlated compounds are listed in the supplementary material section.

<i>Correlatio</i> <i>n</i>	<i>Seed</i>	<i>target vector</i>	<i>function of vector</i>	<i>Common</i> <i>cell lines</i>	<i>Seed</i> <i>stand.</i>	<i>Target</i> <i>stand.</i>
					<i>Dev.</i>	<i>Dev.</i>
0.627	BMSG-SH-3	MoltId: GC15718,	Encodes for RNA-binding protein	59	0.499	0.221
	NSC 753939	Genecard: HNRPLL	which regulates alternative splicing for multiple target mRNAs (Yamamoto et al., 2009).			
0.581	BMSG-SH-3	MoltId: GC17547,	Encodes for Acylphosphatase-1	59	0.499	0.189
	NSC 753939	Genecard: ACYP1	enzyme, which may be involved in alternative splicing and cell cycle regulation (Degl'innocenti et al., 2004; Stone et al., 2010).			
0.564	BMSG-SH-3	MoltId: GC60277,	chromosome 9 open reading frame,	59	0.499	88.127
	NSC 753939	Genecard: C9orf173	encoding for hypothetical protein LOC441476, function unknown (Ota et al., 2004).			
0.491	BMSG-SH-3	MoltId: GC67014,	Encodes for RNA-binding protein	59	0.499	165.655
	NSC 753939		which regulates alternative splicing			

		Genecard: HNRPLL	for multiple target mRNAs (Yamamoto et al., 2009).			
0.478	BMSG-SH-3	MoltId: GC91516,	Sorting nexin 31, involved in protein	59	0.499	22.043
	NSC 753939	Genecard: SNX31	transport and sorting, membrane trafficking, and alternative splicing (Ghaj et al., 2011).			
0.471	BMSG-SH-3	MoltId: GC66400,	unknown (Ota et al., 2004).	59	0.499	108.146
	NSC 753939	Genecard: FAM124A				
0.522	BMSG-SH-4	MoltId: GC160652,	Organic solute carrier partner 1, a	59	0.391	29,463
	NSC 753938	Genecard: OSCP1	transporter which may be involved in drug clearance in placenta (Kobayashi et al., 2005).			
0.494	BMSG-SH-4	MoltId: GC182826,	Sequestosome 1. Involved in	59	0.391	368.75
	NSC 753938	Genecard: SQSTM1	polyubiquitination, trafficking, inclusions, and adapter protein binding to ubiquitin. May regulate signaling cascades (e. g. apoptosis, K^+ channel regulation) (Young & Keiko, 2009).			
0.494	BMSG-SH-4	MoltId: GC84738,	Tumour protein 53 inducible protein	59	0.391	199.379
	NSC 753938	Genecard: TP53/13	13, may act as tumour suppressor (reference)			
0.486	BMSG-SH-4	MoltId: GC188357,	Encodes for proteins containing	59	0.391	26.648
	NSC 753938	Genecard: THAP10	domains similar to DNA binding zinc finger domains; regulates cytokine expression (Champagne et al., 2010 ; Clouaire et al., 2005).			
0.484	BMSG-SH-4	MoltId: GC169527,	Sequestosome 1. Involved in	59	0.391	68.33
	NSC 753938	Genecard: SQSTM1	polyubiquitination, trafficking,			

			inclusions, and adapter protein binding to ubiquitin. May regulate signaling cascades (e. g. apoptosis, K ⁺ channel regulation) (Young & Keiko, 2009).			
0.483	BMSG-SH-4 NSC 753938	MoltId: GC96959, Genecard: SQSTM1	Sequestosome 1. Involved in polyubiquitination, trafficking, inclusions, and adapter protein binding to ubiquitin. May regulate signaling cascades (e. g. apoptosis, K ⁺ channel regulation) (Young & Keiko, 2009).	59	0.391	476.204
0.48	BMSG-SH-5 NSC 753940	MoltId: GC152314, Genecard: ASB1	Encodes for ankyrin repeat and SOCs box protein 1 (suppressor of cytokine signaling) (Kohroki et al., 2005).	59	0.286	31.644
0.471	BMSG-SH-5 NSC 753940	MoltId: GC93247, Genecard: IDS	Encodes for Iduronate-2 sulfatase, which is required for lysosomal degradation of heparan and dermatan sulfate, and is related to the Hunter syndrome (Froissart et al., 1995).	59	0.286	92.605
0.469	BMSG-SH-5 NSC 753940	MoltId: GC187706, Genecard: ZHHC4	Encodes for zinc finger protein 374, a palmitoyltransferase (Ota et al., 2004).	59	0.286	89.778
0.467	BMSG-SH-5 NSC 753940	MoltId: GC36321, Genecard: NFIC	Encodes for protein binding to palindromic sequences in promoters and activates transcription and replication (Santoro et al., 1988).	59	0.286	53.55

0.464	BMSG-SH-5	MolId: GC47494, NSC 753940	Encodes for a subunit of V-ATPase, which acidifies intercellular compartments (Paunescu et al., 2007).	59	0.286	41.437
0.457	BMSG-SH-5	MolId: GC60277, NSC 753940	chromosome 9 open reading frame, encoding for hypothetical protein LOC441476, function unknown (Ota et al., 2004).	59	0.286	88.127

Table S3: COMPARE study of compounds BMSG-SH-3 – 5 against Brown/Botstein/Weinstein microarray dataset. References for the functions of vectors are listed in the supplementary material section.

Uptake study on MCF7 cells

Dimensions: x: 146.11 µm, Y: 146.11 µm

Objective: Plan-Apochromat 63x/1.4 Oil DIC M27

Pixel dwell: 1.60 µs

Average: line 4

Master gain: Ch1 : 775; ChD : 319

Digital gain: 1.00

Digital offset: Ch1 : -0.10; ChD : 0.08

Pinhole: 106 µm

Filter: LP 560

Beam Splitters: MBS : HFT 488/543; DBS1: Mirror; DBS2: Plate.

Laser: 543 nm : 100 %.

Manual corrections: Brightness 45 %; Contrast 55 %.

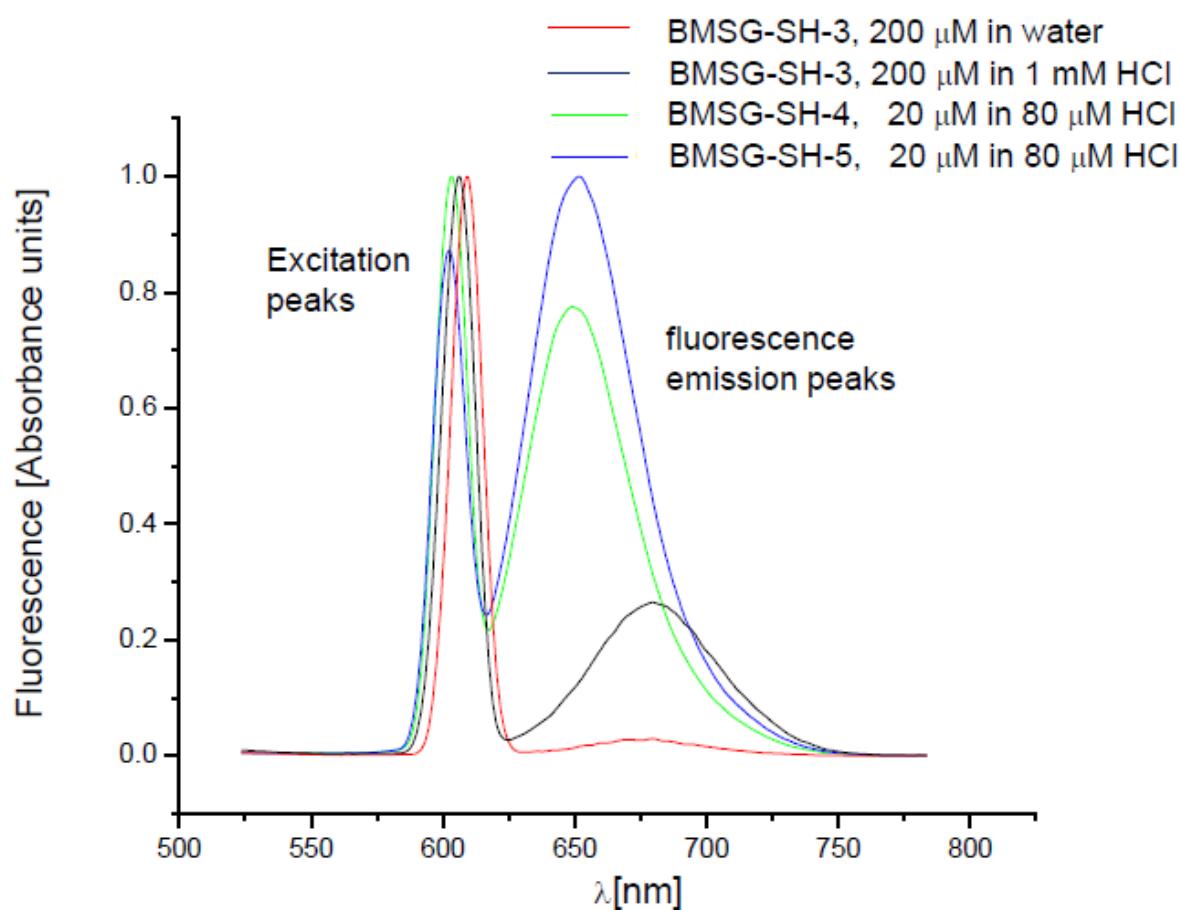
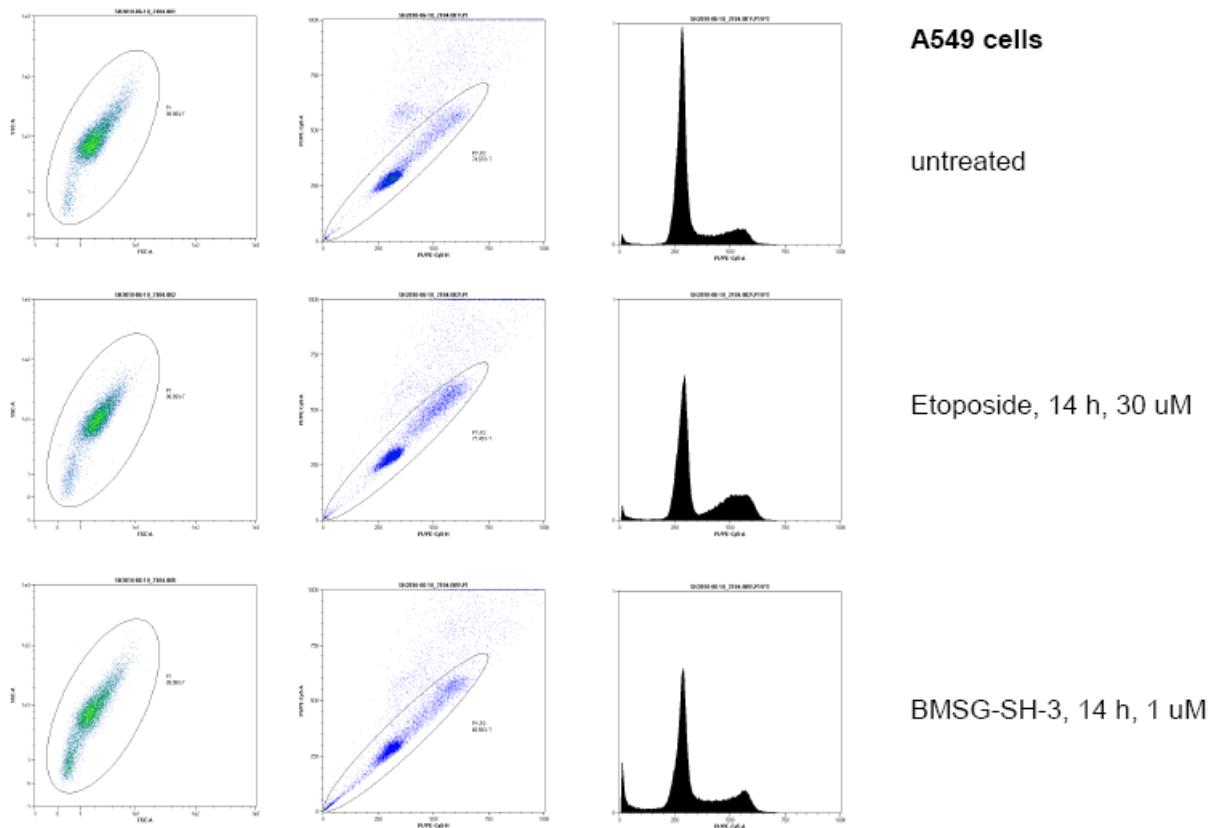


Figure S1: Fluorescence properties of compounds BMSG-SH-3 – 5 at 600 nm excitation. BMSG-SH-3 displays significantly lower fluorescence than the other derivatives, and it increases upon protonation. 4 equivalents of HCl were added to BMSG-SH-4 – 5 to form the soluble chloride salt.



FigureS2: Evaluation of cell cycle analysis of A549 cells. 10000 events each.

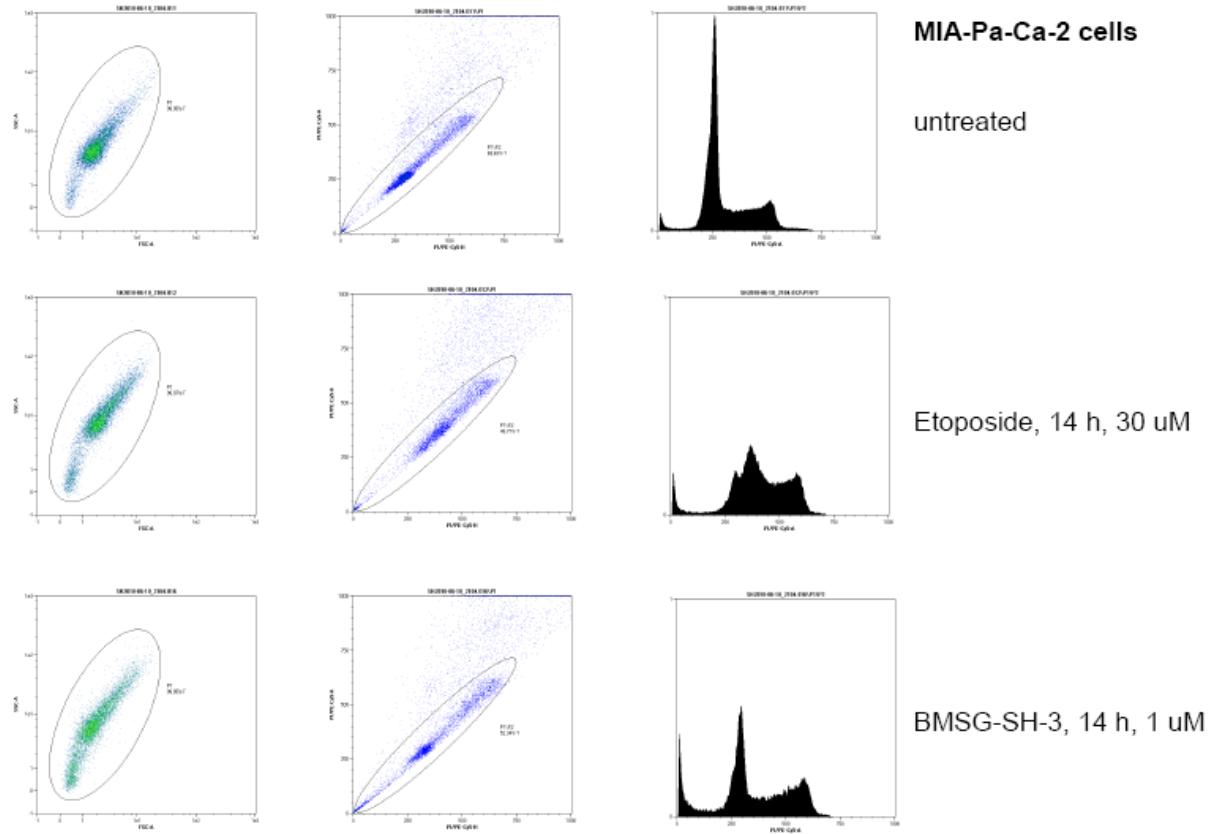
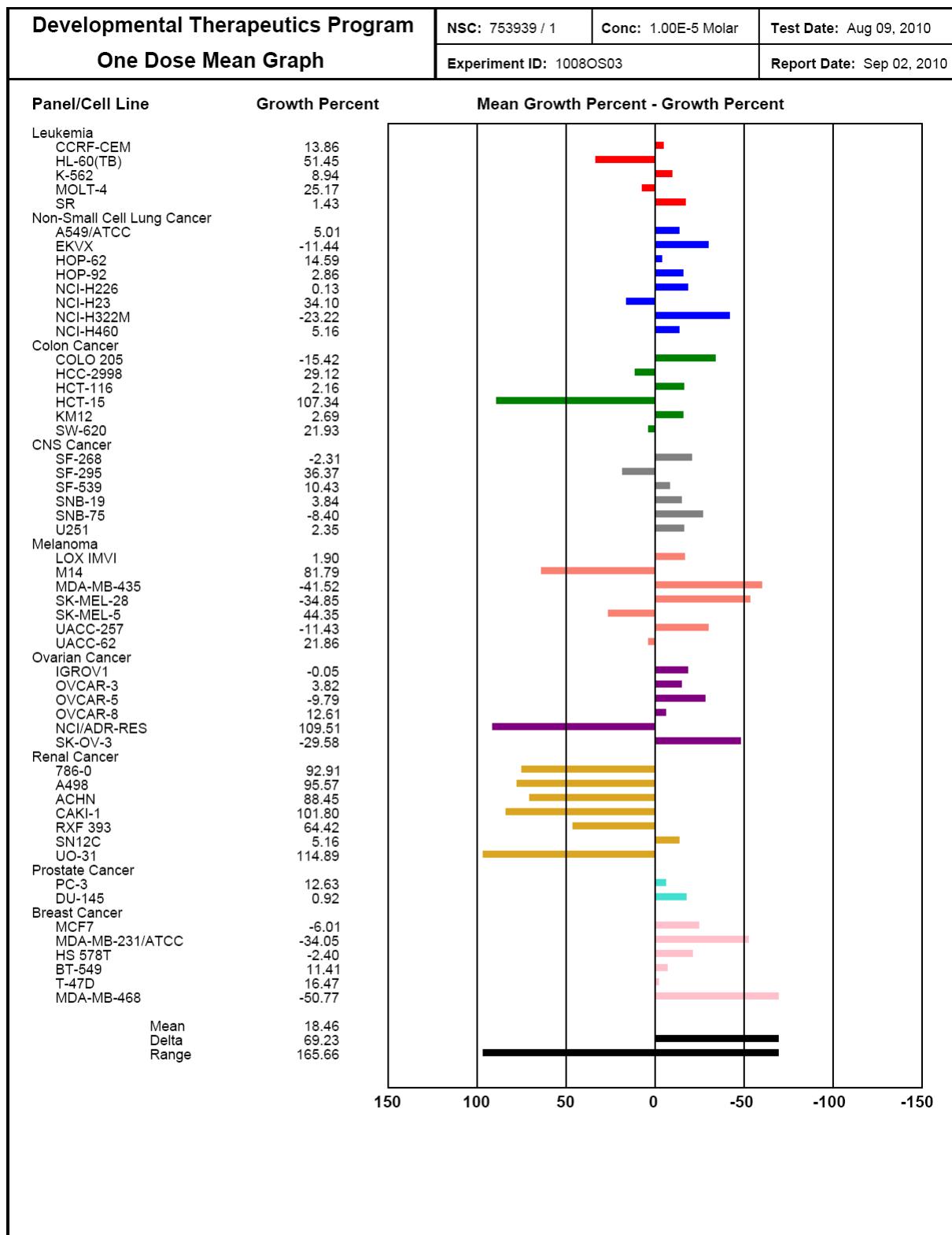


Figure S3: Evaluation of cell cycle analysis of MIA-Pa-Ca-2 cells. 10000 events each.



MOL #81075

Figure S4: NCI 60 cell line panel screen of BMSG-SH-3

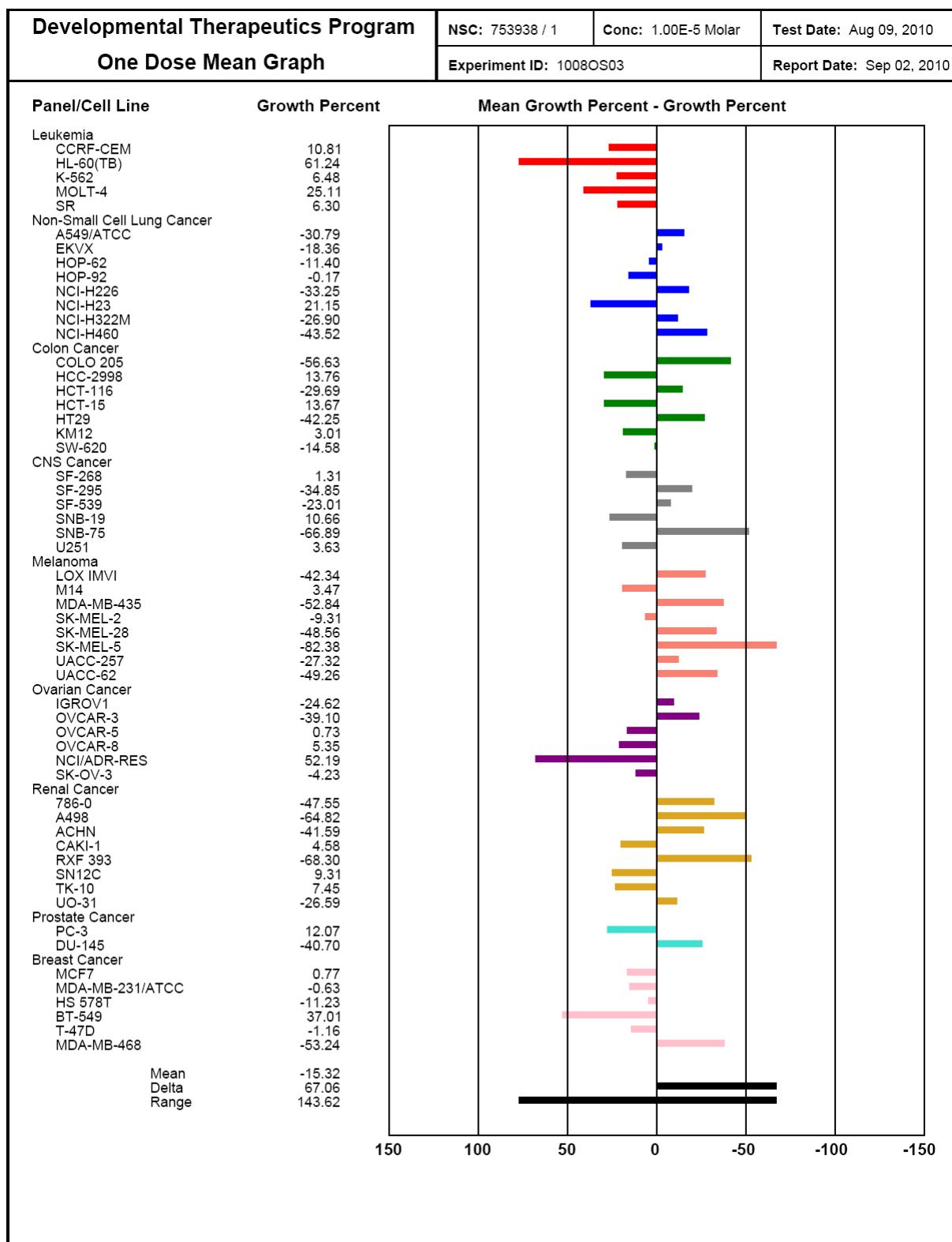


Figure S5: NCI 60 cell line panel screen of BMSG-SH-4

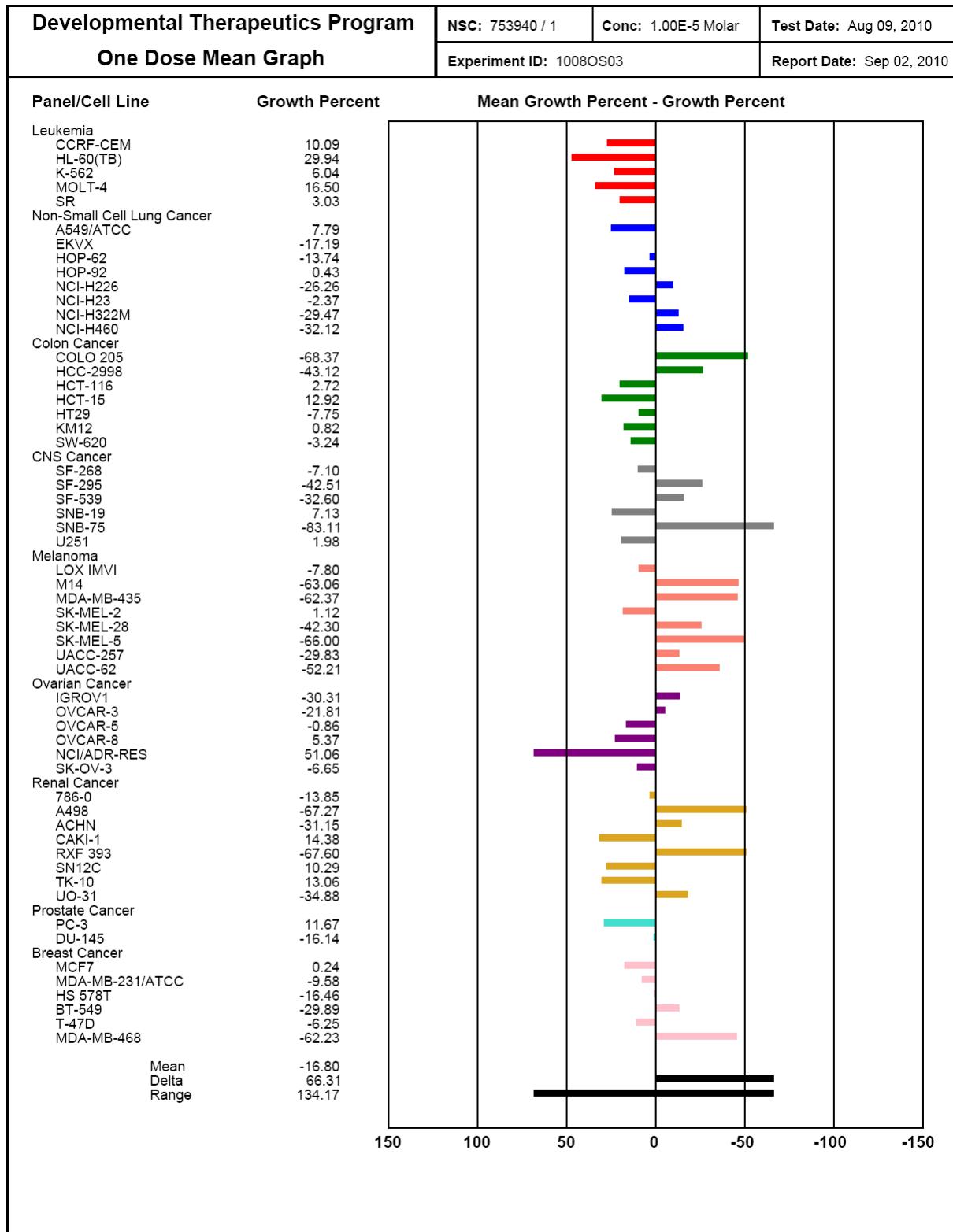


Figure S6: NCI 60 cell line panel screen of BMSG-SH-5

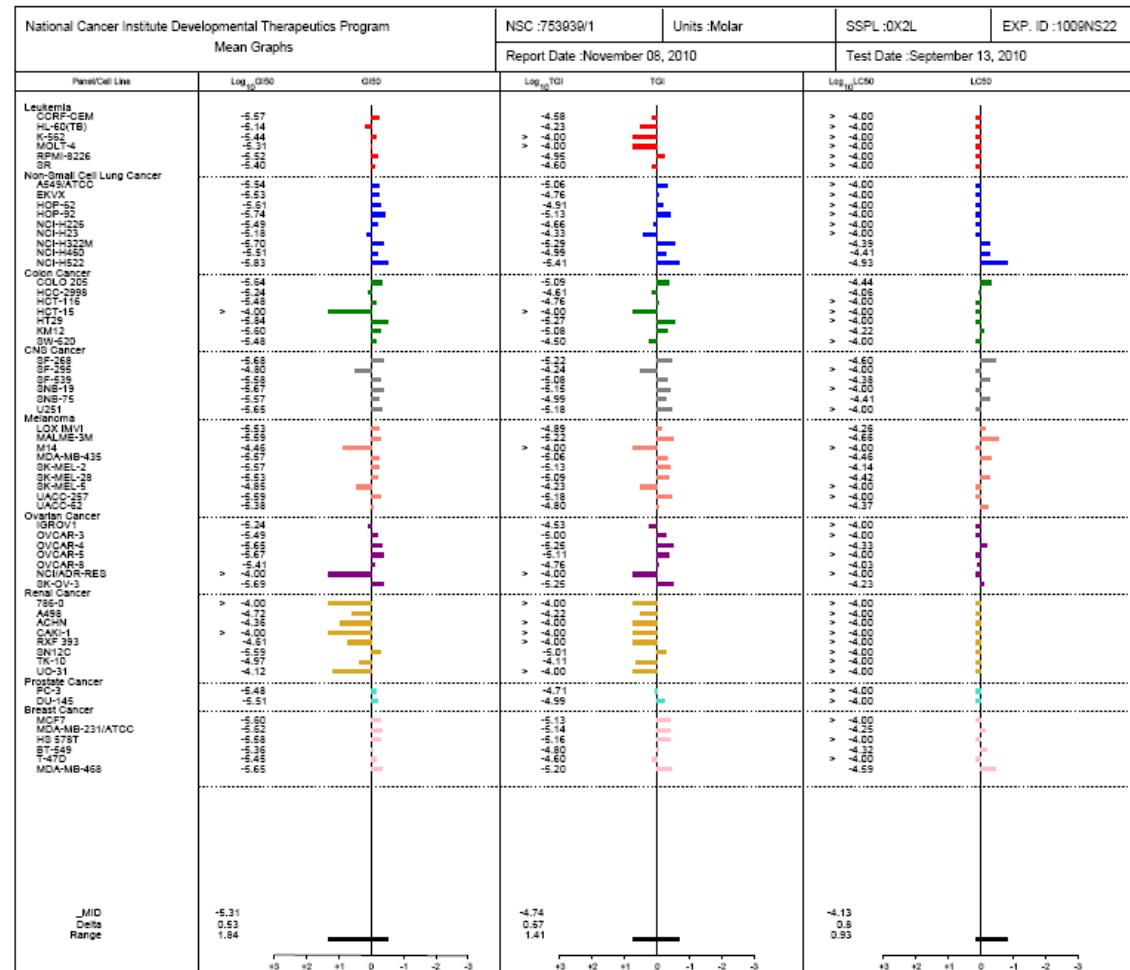


Figure S7: NCI 60 cell line panel screen of BMSG-SH-3, five dose range.

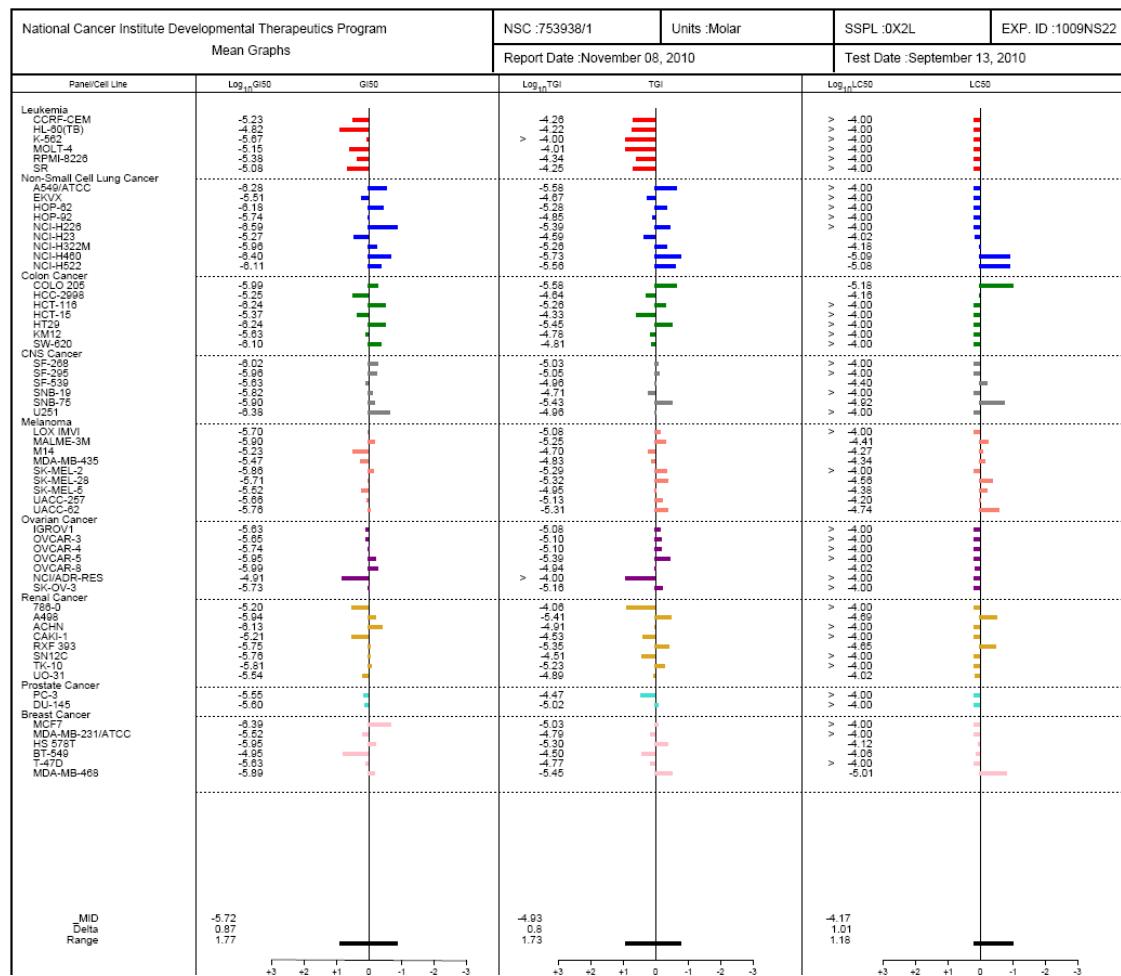


Figure S8: NCI 60 cell line panel screen of BMSG-SH-4, five dose range.

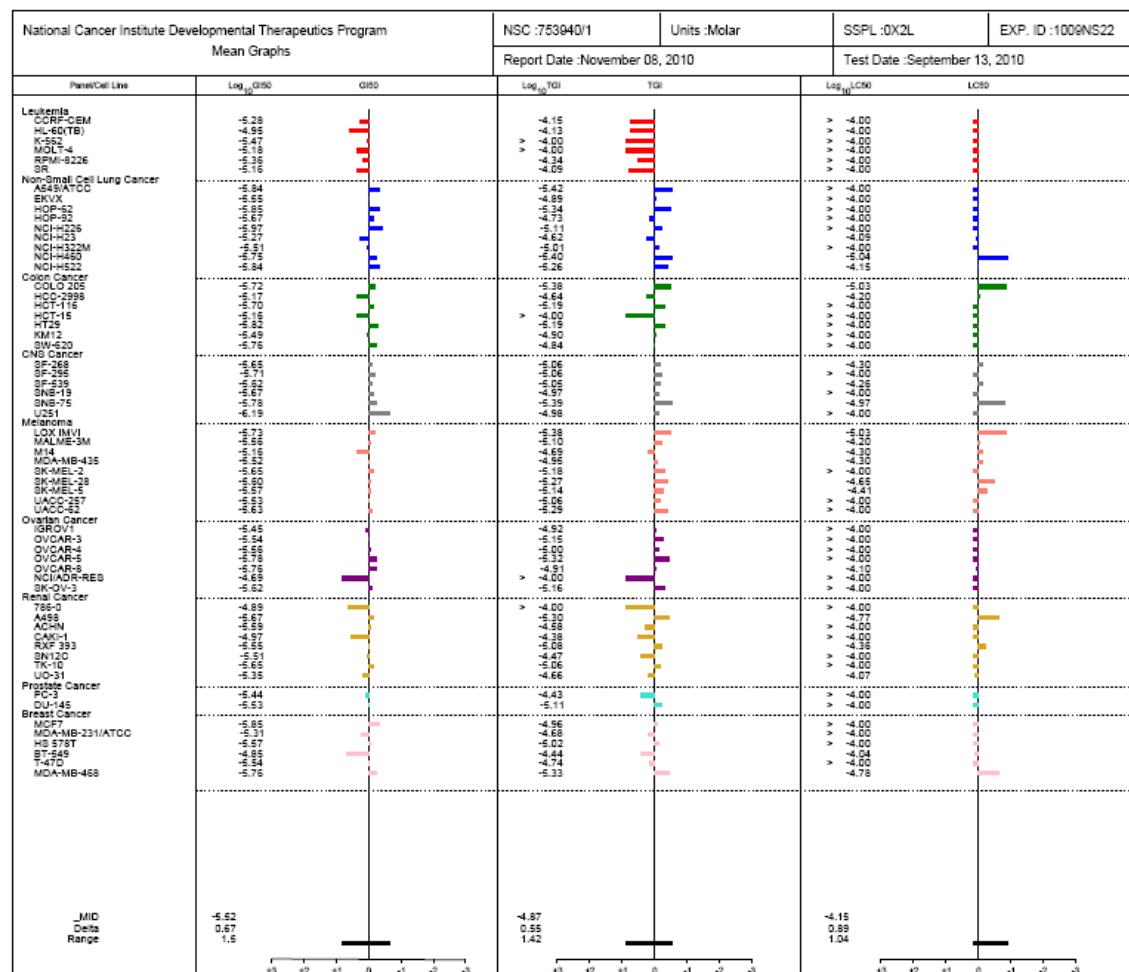


Figure S9: NCI 60 cell line panel screen of BMSG-SH-5, five dose range.

References for Tables S2 and S3

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