

Supplementary materials

Molecular mechanisms of anticancer activity of N-glycosides of indolocarbazoles LSC-1208 and LSC-1269

Roman G. Zenkov^{*1}, Lidiya V. Ektova¹, Vera A. Eremina¹, Olga A. Vlasova¹, Varvara P. Maksimova¹, Timur I. Fetisov¹, Natalia Y. Karpechenko¹, Valeriia G. Popova^{1,4}, Olga G. Usalka^{1,3}, Ekaterina A. Lesovaya^{1,5}, Gennady A. Belitskiy¹, Marianna G. Yakubovskaya¹, Kirill I. Kirsanov^{1,2}

¹ N. N. Blokhin Russian Cancer Research Center, 24 Kashirskoe shosse, Moscow 115478, Russia;

² RUDN University, 6 Miklukho-Maklaya St., Moscow 117198, Russia

³ I.M. Sechenov First Moscow State Medical University, 8-2 Trubetskaya St., Moscow 119991, Russia

⁴ Mendeleev University of Chemical Technology of Russia, 9 Miusskaya ploshchad, Moscow 125047, Russia

⁵ I.P. Pavlov Ryazan State Medical University, 9 Vysokovolt'naya st., Ryazan 390026, Russia

Table S1. Relative expression of the genes in HT29 cell line after treatment with LCS-1208 and LCS-1269 for 24 hours. Expression was upregulated or downregulated by 1.5 times or more in triplicate.

Signal pathway	Genes	Compounds		Signal pathway	Genes	Compounds	
		LCS-1208	LCS-1269			LCS-1208	LCS-1269
TGFβ Pathway	ATF4			Angiogenesis	ANGPT1		
	CDKN1B		1.59		ANGPT2		
	EMP1				CCL2		
	GADD45B	1.93			FGF2		
	HERPUD1				FLT	1.82	
	IFRD1				KDR		0.47
	TNFSF10				PGF		
WNT Pathway	AXIN2	0.41	0.53		SERPINF1		
	CCND2				TEK		
	DAB2				VEGFC		0.41
	FOSL1			APAF1			
	MMP7	4.20		BCL2L11	1.87		
	PPARD			BIRC3	2.38	2.79	
	WISP1			CASP2			
NFκB Pathway	BCL2A1			Apoptosis	CASP7	2.05	
	BIRC3				CASP9		
	CCL5	3.29	1.73		CFLAR		
	CSF1	2.85			FASLG		0.57
	ICAM1	2.61			NOL3		
	IFNG				XIAP		
	STAT1				Cell Cycle	AURKA	

	TNF	1.79				CCND2		
JAK/STAT Pathway	IRF1					CCND3		
	CEBPD					CDC20		
	LRG1					E2F4		
	MCL1	2.06				MCM2		
	SOCS3					MKI67		
	FCER2					SKP2		0.66
	GATA3					STMN1		
		BAX					WEE1	
p53 Pathway	BBC3				Cell Senescence	BMI1		
	BTG2					ETS2		
	CDKN1A					IGFBP3	0.16	0.54
	EGFR					IGFBP5		
	FAS					IGFBP7	0.41	
	GADD45 A					MAP2K1		
	PCNA					MAP2K3		
	RB1					MAPK14		
		HES1					SERPINB 2	
Notch Pathway	HES5				SOD1			
	HEY1				TBX2			
	HEY2				DNA Damage & Repair	DDB2		
	HEYL					DDIT3		
	ID1					ERCC3		
	JAG1	1.54				ERCC5		
	LFNG		0.43			GADD45 G		0.34
	NOTCH1					LIG4		
		BCL2					POLB	
Hedgehog Pathway	BMP2					PPP1R15A		
	BMP4	1.59	1.87			EMT	CDH2	
	PTCH1				DSP			
	WNT1				FOXC2			
	WNT2B				GSC			
	WNT3A				KRT14			0.57
	WNT5A				OCLN			
	WNT6				SNAI1			
		ACSL3					SNAI2	
PPAR Pathway	ACSL4				SNAI3			
	ACSL5				SOX10			
	CPT2				Hypoxia	ADM		
	FABP1					ARNT		
	OLR1	3.34				CA9		
	SLC27A4					EPO		0.48

	SORBS1	0.31	0.39		HMOX1		
Oxidative Stress	FTH1			Metabolism	LDHA		
	GCLC				SLC2A1		
	GCLM				ACLY		
	GSR				ACSL4		
	NQO1				ATP5A1		
	SQSTM1				COX5A		
	TXN				CPT2		
	TXNRD1				G6PD		
						GPD2	
Hypoxia	ADM			Telomeres & Telomerase	LPL		
	ARNT				PFKL		
	CA9	1.91			UQCRFS1		
	EPO				DKC1		
	LDHA	0.58	0.55		PINX1		
	SERPINE1				TEP1		
	SLC2A1				TERF1		
	VEGFA				TERF2IP		
TGFβ Pathway and WNT Pathway	MYC			TINF2			
WNT Pathway and JAK/STAT Pathway	CCND1	0.37	0.47	TNKS			
NFκB Pathway and JAK/STAT Pathway	BCL2L1			TNKS2			
Oxidative Stress and Hypoxia	HMOX1						

Table S2. The data on cytotoxicity of gold standard treatment compounds and LCSs in 11 tumor cell lines.

Cell line	Anticancer agent (Gold standard treatment)	Cytotoxicity (IC ₅₀)	Cytotoxicity of LCSs	
			LCS-1208	LCS-1269
MCF-7	Doxorubicin	0.267 μM [1]	5.5 μM	31 μM
	Cisplatin	9.7 μM [2]		
HepG2	Doxorubicin	0.478 μM [1]	1.7 μM	2.5 μM
U251	Temozolomide	176.5 μM [3]	0.36 μM	1.2 μM
A549	Cisplatin	6.59 μM [4]	1.0 μM	3.2 μM
	Doxorubicin	0.32 μM [5]		
PC-3	Docetaxel	0.0005 μM [6]	0.97 μM	24 μM
	Cisplatin	3.3 μM [7]		
HT29	Doxorubicin	750 nM [8]	0.13 μM	1.4 μM
	Cisplatin	75,7 μM [9]		
HeLa	Cisplatin	5.0 μM [2]	28,1	26,6
	Doxorubicin	0.374 μM [10]		
CCRF CEM	Cisplatin	0.7 μM [11]	2.0 μM	6.8 μM
Granta-519	Doxorubicin	212.7 nM (48 h) [12]	0.071 μM	0.60 μM
K562	Doxorubicin	About 1 μM [13]	6.0 μM	>500 μM
KG-1	Doxorubicin	About 9 μM (48 h) [14]	0.6 μM	7.1 μM

Figure S1. Analysis of the LCS-1269 effects on histone modifications, Western blot results.

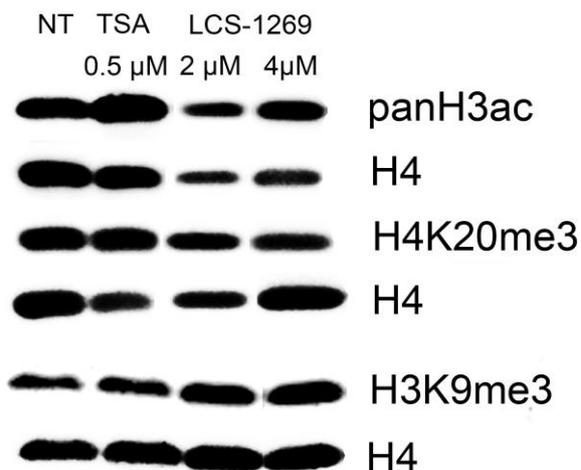


Figure S2. Analysis of the LCS-1269 effects on HDACs expression, Western blot results.

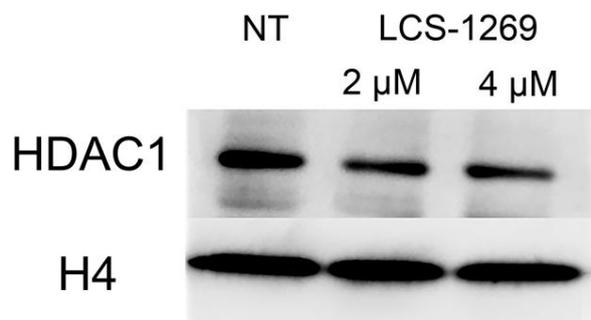
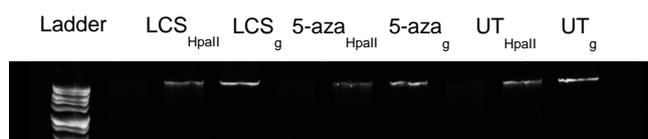


Figure S3. Analysis of the LCS-1269 effects on DNA methylation MSRE assay results.



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