Supplementary Table 1 - Primers for Pyrosequencing

	Forward	Reverse
1st/2nd CpG Island First Step	AGTTATGTATGTATGTGTGTTGTATATAGAGT	AAAAAATAATCCCCACTCTCCTATC
1st/2nd CpG Island Second Step	[Btn]TATATAGAGTAGATATATAGTTTATTAAG	CTATCTAATCCCTCCTCTC
1st/2nd CpG Island Sequencing A		СТТССТСТССАААААТСТАА
1st/2nd CpG Island Sequencing B		TTTCTTCTTCCCAATCTTTACC
1st/2nd CpG Island Sequencing C		CCTCTCCTCCCCCAATTCCA
1st/2nd CpG Island Sequencing D		CCAAATATCTTTTCTTCT
3rd CpG Island First Step	GAGGTGAGAGTGTTTTAGAT	CATAAACTTAATTTTCAATCTTAT
3rd CpG Island Second Step	GTTTATAATGATTAGTGTTTGTGGGAAAG	[Btn] AACCAAACCCCCTCAAACAACAAAACCTA
3rd CpG Island Sequencing A	AGGTTAGAGAGGAAATTTTG	
3rd CpG Island Sequencing B	GGTTGAGAGTTAGGAT	

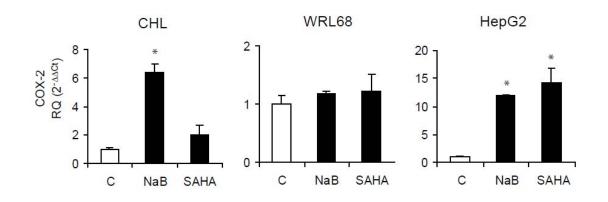
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Supplementary	Table 2 -	Primers 10	r methylation	-sensitive r	estriction as	SSAV (MISKA)

	Forward	Reverse
CpG -431	CTTAACCTTACTCGCCCCAGTCT	AGAAGGACACTTGGCTTCCTC
CpG -372	GGAAGCCAAGTGTCCTTCTGC	GGGCAGGGTTTTTTACCCAC
CpG -138	AGCTTCCTGGGTTTCCGATT	CCCCCACAAATTTTTCCCTC
CpG +75/84/98	GCCCTCAGACAGCAAAGCCTA	GGGAAAGCTGGAATATCCACG

	Forward	Reverse
Distal Promoter	GGAAGCCAAGTGTCCTTCTGC	TGATCGCCTTGGATGGGATA
Proximal Promoter	AGCTTCCTGGGTTTCCGATT	GGGCAGGGTTTTTTACCCAC
First Exon	CTCGGTTAGCGACCAATTGTC	CGTGCTCCTGACGCTCACT
First intron	AGCTTGGACCGCTAGAGTTCG	GGGAAAGCTGGAATATCCACG

Supplementary Table 3 - Primers for Chromatin immunoprecipitation

Figure. S1- Expression of COX-2 increases after treatment of HCC cells with HDAC inhibitors. Eight hours before the experiment HCC cells were change to 1% FBS and they were incubated without (white bars) or with 2.5 mM NaB or 2.5 μ M SAHA for 16 h. Total RNA was then isolated and mRNA levels for COX-2 and the internal control 18S were determined by qPCR. COX-2 mRNA amounts were calculated as RQ and normalized to the expression of ribosomal levels. Values represent fold change relative to each cell line without treatment. Data are reported as means \pm SD of three independent experiments. **p < 0.01 and *p < 0.05



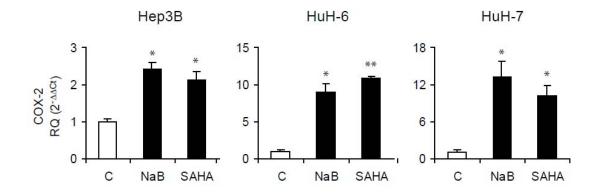
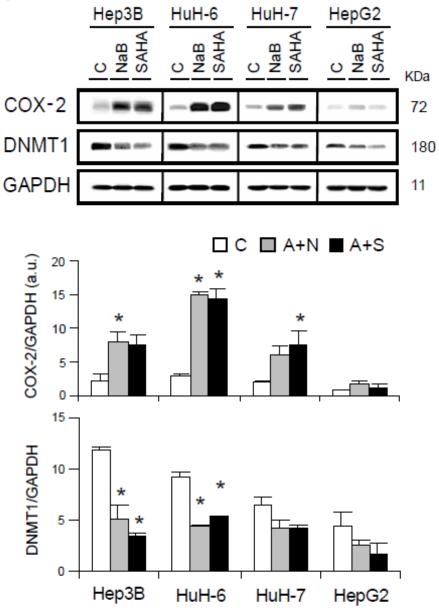
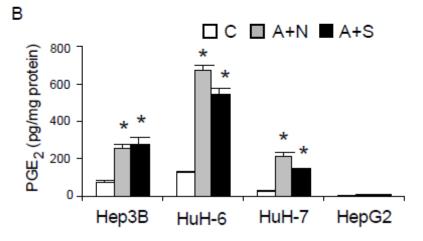


Figure. S2- Synergistic effect of HDAC inhibitors and AzadC on COX-2 expression.

(A) HCC cells in 1% FBS were incubated without (white) or with 5 μ M AzadC and 2.5 mM NaB (A+N, grey) or AzadC and 2.5 μ M SAHA (A+S, black) for 48 h. Cellular extracts were prepared and protein was analyzed by Western blot. A representative Western blot analyzing COX-2 and DNMT1 protein levels is shown. The expression of target proteins was normalized to GAPDH. Each panel has an optimal exposition time. Densitometric analysis of COX-2 and DNMT1 protein levels is shown. Results are expressed as arbitrary units. (B) PGE₂ concentration was determined by enzyme immunoassay in the supernatant of the cells. Data are expressed as means \pm SD of three independent experiments. *p < 0.05 vs. the corresponding cell line without treatment





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