## G9a orchestrates PCL3 and KDM7A to promote histone H3K27 methylation

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## **Supplementary Information**

**Supplementary figure legends** 

**Supplementary Figure 1-** Proliferation of gemcitabine-resistant PANC-1-R cells is similar to parental PANC-1 pancreatic cancer cells.

**Supplementary Figure 2-** Increase of G9a, H3K9 and H3K27 methylation in gemcitabine-resistant pancreatic cancer cells.

**Supplementary Figure 3-** Inhibition of G9a decreased cell migration and invasion without significantly affecting cellular proliferation of PANC-1-R cells.

**Supplementary Figure 4-** Inhibition of H3K9 and H3K27 methylation by overexpressing methyltransferase-dead G9a (DN-G9a).

**Supplementary Figure 5-** Inhibition of G9a by shRNA or UNC0638 reduced H3K27 methylation in lung and breast cancer cells.

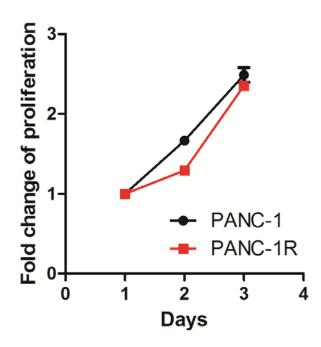
**Supplementary Figure 6-** UNC0638 treatment decreased the binding of PCL3 to E-cadherin promoter in PANC-1-R cells.

**Supplementary Figure 7-** UNC0638 treatment reduced PLC3 expression and decreased the binding of G9a to PCL3 gene prometer in PANC-1-R cells.

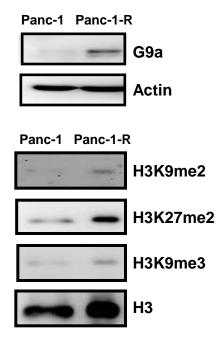
**Supplementary Figure 8-** Specific reduction of PCL3 by UNC0638.

**Supplementary Table 1-** Primer sequences used in this study.

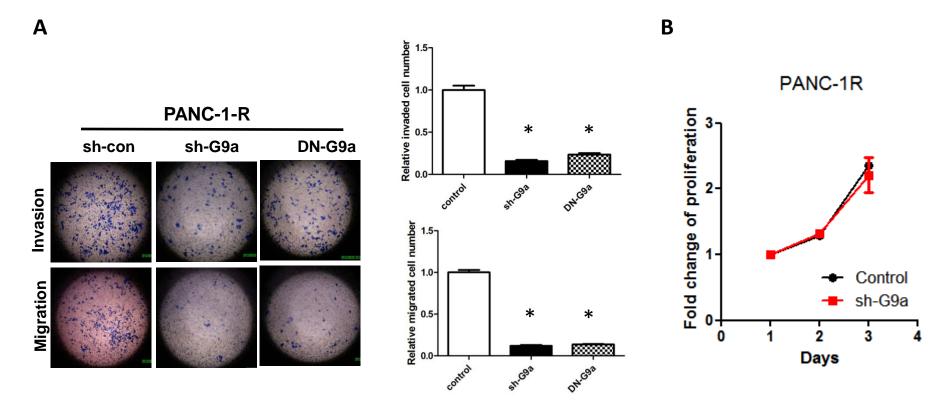
Supplementary Figure 1- Proliferation of gemcitabine-resistant PANC-1-R cells is similar to parental PANC-1 pancreatic cancer cells. Panc-1 and gemcitabine-resistant Panc-1-R cells were seeded at the density of 2000 cell/per well in 96-well plates. After different times, MTT assay was performed to investigate cellular proliferation. Results were the average of two independent assays.



**Supplementary Figure 2- Increase of G9a, H3K9 and H3K27 methylation in gemcitabine-resistant pancreatic cancer cells.** The protein level of G9a and the methylation status of H3K9 and H3K27 were compared in Panc-1 and gemcitabine-resistant Panc-1-R cells by western blotting.



Supplementary Figure 3- Inhibition of G9a decreased cell migration and invasion without significantly affecting cellular proliferation of PANC-1-R cells. (A) Knockdown of G9a (shRNA) or ectopic expression of methyltransferase-dead G9a (DN-G9a) reduced cell migration and invasion of Panc-1-R pancreatic cancer cells. (B) The proliferation rate of PANC-1-R cells transfected with control or G9a shRNA.



Supplementary Figure 4- Inhibition of H3K9 and H3K27 methylation by overexpressing methyltransferase-dead G9a (DN-G9a). Overexpression of methyltransferase-dead G9a decreased both H3K9 and H3K27 methylation in PANC-1-R cells.

PANC-1-R

C DN-G9a

G9a

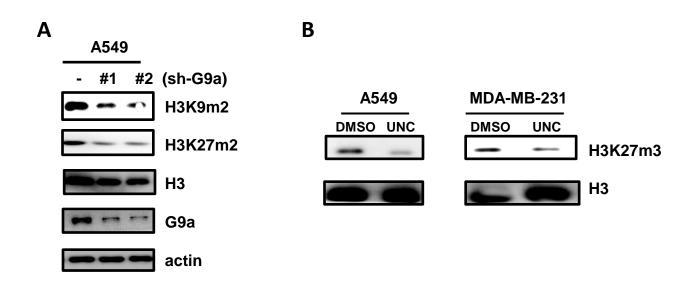
H3K9m2

H3K9m3

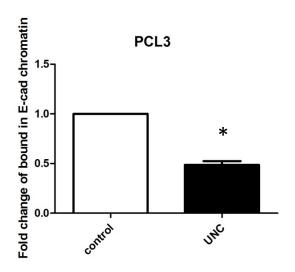
H3K27m2

H3

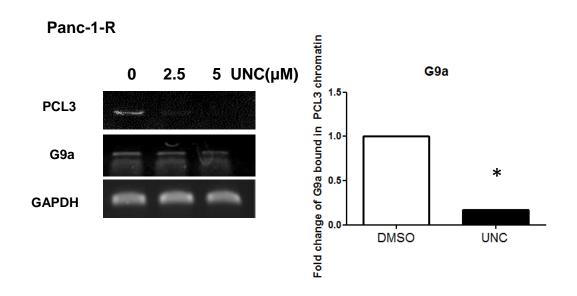
Supplementary Figure 5- Inhibition of G9a by shRNA or UNC0638 reduced H3K27 methylation in lung and breast cancer cells. (A) Knockdown of G9a decreased both H3K9 and H3K27 methylation in A549 lung cancer cells. (B) UNC0638 treatment decreased H3K27 methylation in A549 and MDA-MB-231 cells.



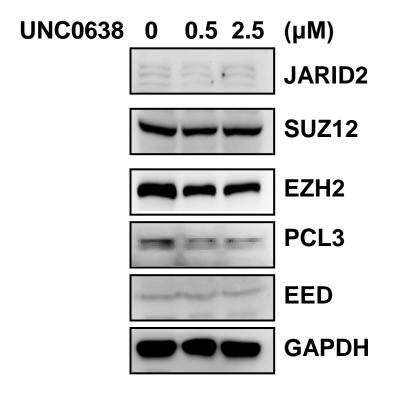
**Supplementary Figure 6-**UNC0638 treatment decreased the binding of PCL3 to E-cadherin promoter in PANC-1-R cells.



**Supplementary Figure 7-**UNC0638 treatment reduced PLC3 expression and decreased the binding of G9a to PCL3 gene prometer in PANC-1-R cells.



**Supplementary Figure 8- Specific reduction of PCL3 by UNC0638.** UNC0638 treatment reduced PLC3 protein in a dose-dependent manner. The protein level of EZH2 was decreased slightly. Other components of the PRC2 complex were not affected.



## Supplamantary Table 1. Primer sequences used in this study

Assay	Target	Sequence $(5' \rightarrow 3')$	Amplicon (bp)
shRNA	shG9a-C	GCTCCAGGAATTTAACAAGAT	
	sh-G9a-E	CTCCAGGAATTTAACAAGATT	
	sh-KDM7A-C	TGGATTTGATGTCCCTATTAT	
	sh-KDM7a-E	TTAGACCTGGACACCTTATTA	
	sh-PCL3-A	CCTCGTGACTTTCGAAGATAA	
	sh-PCL3-B	CCTGGCTAGCATATTTGACTT	
	sh-PCL3-C	CCCACCTCAAGTCATCTATCA	
	sh-PCL3-D	CAACGCTCTGAACAGTTATAA	
	sh-PCL3-E	ACCACCTGGCTAGCATATTTG	
Real-	E-cadherin	F:CCTGGGACTCCACCTACAGA	407
PCR		R: GGATGACACAGCGTGAGAGA	
	SUZ12	F:GAGCACGTCCAGGCTGACCAC	475
		R:TACTGGAAACTGCAAGGGACGGGA	
	JARID2	F:GGTCCGCGCTCAGGTGGAGA	594
		R:AGCTTGCTGCGGAAGCCGTT	
	PCL3	F:CCAGTATGTGCTGTGCCGGTGG	444
		R: CAGGTGTGGCCCTTCTGCTTGG	
	EED	F:GCCTGCGGCCAAGAAGCAGA	448
		R: AGCCAGCAGAGGATGGCTCGT	
	EZH2	F:GGGACTAGGGAGGTGGAAGA	359
		R: GCTGTGCCCTTATCTGGAAA	
	G9a	F: TGGGAAAGGTGACCTCAGAT	336
		R: TCCCTGACTCCTCATCTTCC	
	KDM7A	F: CGAGTGCGATATCTGCAAGG	154
		R: GTCATGTCTGTGCCAGTTCCT	
	GAPDH	F:GAGTCAACGGATTTGGTCGT	511
		R: TGTGGTCATGAGTCCTTCCA	
Chip	E-cadherin	F: CAGGTGAACCCTCAGCCAATCAGC	168
primer			
	promoter	R: GTGCGGTCGGGTCGGGCC	
	(-80~+88)		
	E-cadherin	F: CCCGCCCAGCTGTGTCATTTT	199
	promoter	R: AATGGTGCCCATCCACGTGG	
	(-10434~-10236)		
	PCL3 promoter	F:GGTCCCACTTGGAGTCTGG	238
	(-20~+218)	R:AATAATTCGCGGGAAAACG	
	KDM7A promoter	F:CTCCCTCCCTTTCCTT	237
	(-1584~-1343)	R: TGGTGAAACCCCGTCTCTAT	