

Supplementary Materials

Protein Arginine Methyltransferase 3 Enhances Chemoresistance in Pancreatic Cancer by Methylating hnRNPA1 to Increase ABCG2 Expression

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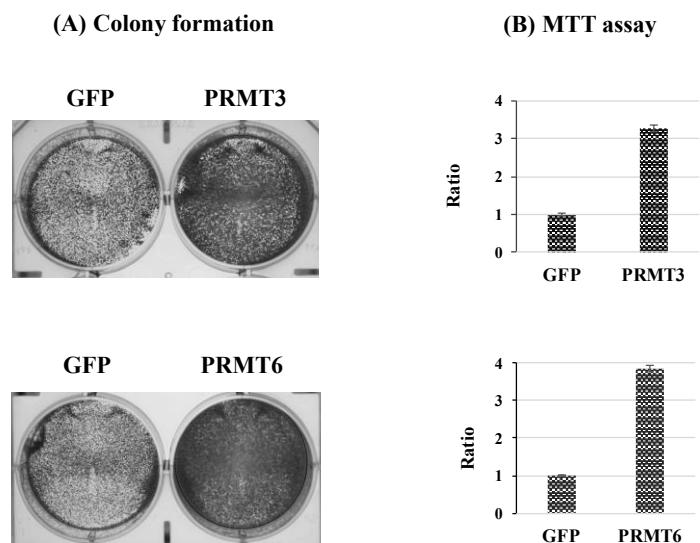


Figure S1. Overexpression of PRMT3 and PRMT6 facilitates cell proliferation. **(A)** GFP-, PRMT3- and PRMT6-overexpressing PANC-1 cells (1×10^4) were cultured for three days and the number of colonies was photographed. **(B)** GFP-, PRMT3- and PRMT6-overexpressing PANC-1 cells were cultured for three days and then the cell viability was measured by MTT assay. Error bars, SEM. $n = 3$.

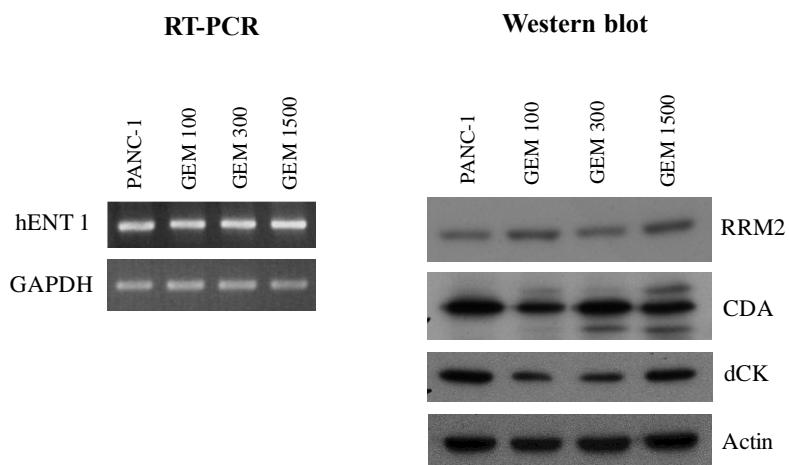
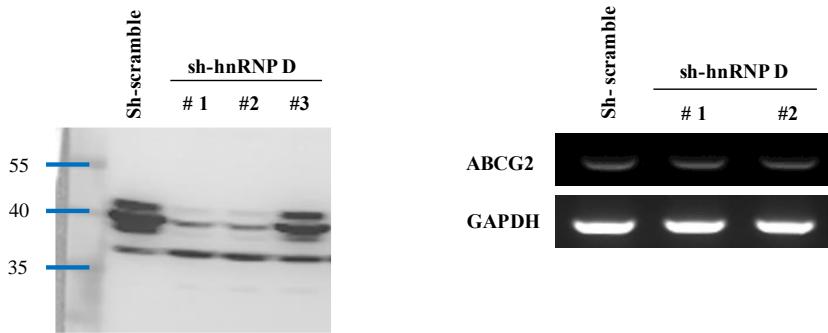


Figure S2. The expression of hENT1, RRM2, CDA and dCK was not altered in gemcitabine-resistant cell lines.



IB: hnRNP D

Figure S3. Depletion of hnRNP D had no effects on ABCG2 mRNA level. PRMT3-overexpressing PANC-1 cells were transfected with hnRNP D-targeting shRNAs. The protein level of hnRNP D was determined by western blot and mRNA levels of ABCG2 was measured by semi-quantitative RT-PCR.

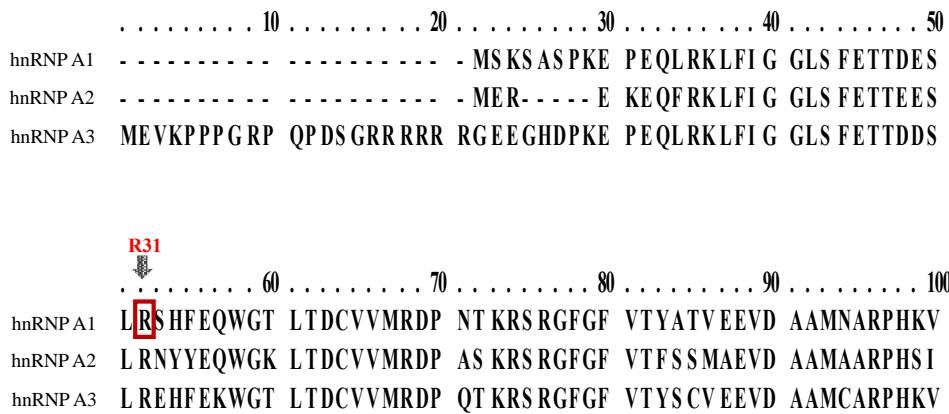


Figure S4. Sequence alignment of hnRNP A1, A2 and A3 proteins.

Table S1. PRMT3-associated heterogeneous nuclear ribonucleoproteins (hnRNPs).

HNRPK_HUMAN	Heterogeneous nuclear ribonucleoprotein K OS = Homo sapiens GN = HNRNPK PE = 1 SV = 1
HNRPD_HUMAN	Heterogeneous nuclear ribonucleoprotein D0 OS = Homo sapiens GN = HNRNPD PE = 1 SV = 1
HNRH1_HUMAN	Heterogeneous nuclear ribonucleoprotein H OS = Homo sapiens GN = HNRNPH1 PE = 1 SV = 4
ROA1_HUMAN	Heterogeneous nuclear ribonucleoprotein A1 OS = Homo sapiens GN = HNRNPA1 PE = 1 SV = 5
HNRPF_HUMAN	Heterogeneous nuclear ribonucleoprotein F OS = Homo sapiens GN = HNRNPF PE = 1 SV = 3
HNRDL_HUMAN	Heterogeneous nuclear ribonucleoprotein D-like OS = Homo sapiens GN = HNRNPDL PE = 1 SV = 3
ROA3_HUMAN	Heterogeneous nuclear ribonucleoprotein A3 OS = Homo sapiens GN = HNRNPA3 PE = 1 SV = 2

Table S2. Oligonucleotide sequences.

The Sequences of shRNA Plasmids	
sh-PRMT3#1	CGTGACCCTCACGGTGAATAA
sh-PRMT3#2	CCTTGGGAGAAAGAAGAGAT
sh-PRMT3#3	CCTTGTGGTATTAAGCATATA
sh-PRMT3#4	GCACTGAGTGATGTGAATAAA
sh-hnRNP A1#D1	AGATATTGTTGGTGGCATT
sh-hnRNP A1#G1	ACCCATGAAGGGAGGAAATT
sh-hnRNP A1#H1	CAACAATCAGTCTCAAATT
sh-hnRNP A1#F2	AGATATTGTTGGTGGCATT
The Sequences of Primers (5'-3')	
PRMT3-F	GCATGGTTCATAAACATGGACTT
PRMT3-R	CACCCAACATCCAAAACCTACC
PRMT6-F	CCTGGGTATCCTTCGGAACT
PRMT6-R	CTCCTTCAGCCACTTGGTTC
ABCG2-F	ACTGAGATTGAGAGACGC
ABCG2-R	TCTGGAGAGTTTTATCTTTCA
hnRNP A1-F	CCACGAAACCAAGGTGGCTA
hnRNP A1-R	CTGGCTCTCCTCTCCTGCTA
hnRNPA1-R31K-F	CAACTGATGAGAGCCTGAAGAGCCATTTGAGCAATGG
hnRNPA1-R31K-R	CCATTGCTAAAATGGCTCTCAGGCTCTCATCAGTTG
hnRNPA1-R31F-F	CAACTGATGAGAGCCTGTTAGCCATTTGAGCAATGG
hnRNPA1-R31F-R	CCATTGCTAAAATGGCTAACAGGCTCTCATCAGTTG
5'UTR (1-493)-F	(T7) GTCAGCGCTGCCTGAGCTCG
5'UTR (1-493)-R	GCCATTGGTGTTCCTGTGACACTGGG
3'UTR (2462-4445)-F	(T7) CACGTGGCCTGGCTGTATG
3'UTR (2462-4445)-R	CTCTCTGTGTAATAAGGGAAGGG
5'UTR-N.C. (1-295)-F	(T7) GTCAGCGCTGCCTGAGCTCG
5'UTR-N.C. (1-295)-R	CCTAAATCCTACCCAGTCCTCCAC
5'UTR-ARE1 (271-550)-F	(T7) GTGGAGGAACTGGTAGGATTAGG
5'UTR-ARE1 (271-550)-R	GCCATTGGTGTTCCTGTGACACTGGG
3'UTR-ARE1 (2381-2573)-F	(T7) CACGTGGCCTGGCTGTATG
3'UTR-ARE1 (2381-2573)-R	GCAACAGTGTGATGCCAAGGG
3'UTR-ARE2 (2553-2821)-F	(T7) CCCTTGCCATCACACTGTTGC
3'UTR-ARE2 (2553-2821)-R	GGGCTACTAACCTACCTATT
3'UTR-ARE3 (2801-3062)-F	(T7) GAATAGGTAGGTTAGTAGCCC
3'UTR-ARE3 (2801-3062)-R	CCAAACCCCTCAGCCAAGATTCC
3'UTR-ARE4 (3276-3585)-F	(T7) CTGACCAACTCCTGGATTGG
3'UTR-ARE4 (3276-3585)-R	GGAAGGAAGTAGTGAATGGAG
3'UTR-ARE5 (4027-4331)-F	(T7) GGGGCATCATGGTGTATAGACGCC
3'UTR-ARE5 (4027-4331)-R	CTCTCTGTGTAATAAGGGAAGGG



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