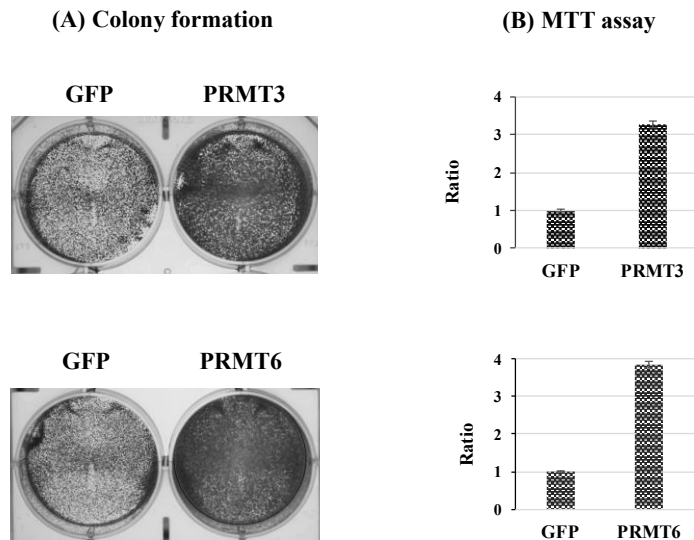
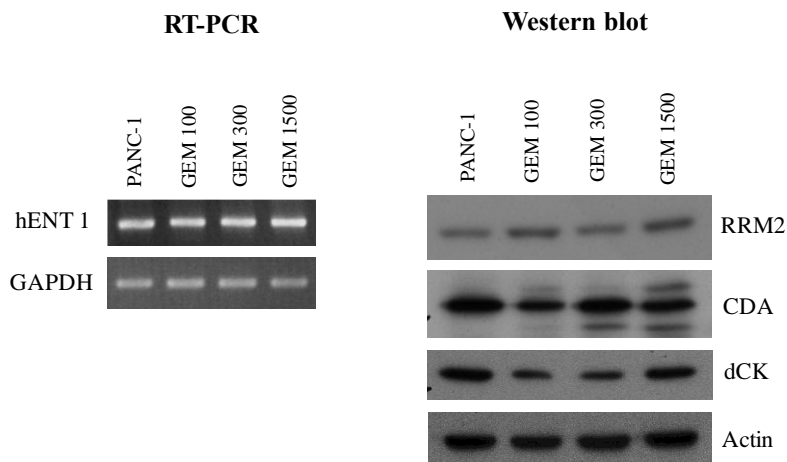


# Protein Arginine Methyltransferase 3 Enhances Chemoresistance in Pancreatic Cancer by Methylating hnRNPA1 to Increase ABCG2 Expression

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**Figure S1.** Overexpression of PRMT3 and PRMT6 facilitates cell proliferation. (A) GFP-, PRMT3- and PRMT6-overexpressing PANC-1 cells ( $1 \times 10^4$ ) 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.



**Table S2.** Oligonucleotide sequences.

<b>The Sequences of shRNA Plasmids</b>	
sh-PRMT3#1	CGTGACCCTCACGTTGAATAA
sh-PRMT3#2	CCTGGGAGAAAGAAGAGAT
sh-PRMT3#3	CCTGTGGTATTAAGCATATA
sh-PRMT3#4	GCAGTGAGTGATGTGAATAAA
sh-hnRNP A1#D1	AGATATTTGTTGGTGGCATT
sh-hnRNP A1#G1	ACCCATGAAGGGAGGAAATTT
sh-hnRNP A1#H1	CAACAATCAGTCTTCAAATTT
sh-hnRNP A1#F2	AGATATTTGTTGGTGGCATT
<b>The Sequences of Primers (5'–3')</b>	
PRMT3-F	GCATGGTTCATAAACATGGACTT
PRMT3-R	CACCCAACATCCAAAACCTACC
PRMT6-F	CCTGGGTATCCTTCGGAAC
PRMT6-R	CTCCTTCAGCCACTTGGTTC
ABCG2-F	ACTGAGATTGAGAGACGCGG
ABCG2-R	TCTGGAGAGTTTTATCTTTTCAGC
hnRNP A1-F	CCACGAAACCAAGGTGGCTA
hnRNP A1-R	CTGGCTCTCCTCTCCTGCTA
hnRNPA1-R31K-F	CAACTGATGAGAGCCTGAAGAGCCATTTGAGCAATGG
hnRNPA1-R31K-R	CCATTGCTCAAAATGGCTCTTCAGGCTCTCATCAGTTG
hnRNPA1-R31F-F	CAACTGATGAGAGCCTGTTTAGCCATTTGAGCAATGG
hnRNPA1-R31F-R	CCATTGCTCAAAATGGCTAAACAGGCTCTCATCAGTTG
5'UTR (1-493)-F	(T7) GTCAGCGCTGCCTGAGCTCG
5'UTR (1-493)-R	GCCATTGGTGTTCCTTGTGACACTGGG
3'UTR (2462-4445)-F	(T7) CACGTGGCCTTGGCTTGTATG
3'UTR (2462-4445)-R	CTCTCTGTGTAATAAGGGAAGGG
5'UTR-N.C. (1-295)-F	(T7) GTCAGCGCTGCCTGAGCTCG
5'UTR-N.C. (1-295)-R	CCTAAATCCTACCCAGTTCCTCCAC
5'UTR-ARE1 (271-550)-F	(T7) GTGGAGGAACTGGGTAGGATTTAGG
5'UTR-ARE1 (271-550)-R	GCCATTGGTGTTCCTTGTGACACTGGG
3'UTR-ARE1 (2381-2573)-F	(T7) CACGTGGCCTTGGCTTGTATG
3'UTR-ARE1 (2381-2573)-R	GCAACAGTGTGATGGCAAGGG
3'UTR-ARE2 (2553-2821)-F	(T7) CCCTTGCCATCACACTGTTGC
3'UTR-ARE2 (2553-2821)-R	GGGCTACTAACCTACCTATTC
3'UTR-ARE3 (2801-3062)-F	(T7) GAATAGGTAGGTTAGTAGCCC
3'UTR-ARE3 (2801-3062)-R	CCAAACCCTCAGCCAAGATTCC
3'UTR-ARE4 (3276-3585)-F	(T7) CTGACCACTCCTGGATTTGG
3'UTR-ARE4 (3276-3585)-R	GGAAGGAAGTAGTACTGGGAG
3'UTR-ARE5 (4027-4331)-F	(T7) GGGGCATCATGGTGTATAGACGCC
3'UTR-ARE5 (4027-4331)-R	CTCTCTGTGTAATAAGGGAAGGG



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