Table S2. Primer and oligo sequences used in the study.

	Primer/Oligo sequences (5'----3')
A. Real-time qPCR for miRNAs/small nuclear RNA	
hsa-miR-765	
hsa-miR-765 qRT forward primer	TGGAGGAGAAGGAAGGTGATG
Universal qPCR reverse primer	N.A.
RNU6	
RNU6 qRT forward primer	CGCAAGGATGACACGCAAAT
Universal qPCR reverse primer	N.A.
B. Real-time RT-PCR for gene mRNA	
HMGA1 mRNA	
HMGA1-qRT-PCR-F	GAAAAGGACGGCACTGAGAA
HMGA1-qRT-PCR-R	CTTCCTGGAGTTGTGGTGGT
RPS3 mRNA	
RPS3-qRT-PCR-F	AGCCACCAGAACACAGAATG
RPS3-qRT-PCR-R	CTAGTGGCCACCTTTTCAGC
C. Promoter analysis	
5' upstream sequence of hsa-miR-765	
hsa-miR-765 promoter -F	GGAAGATCTCTGGCACAAACCAACTACCA
hsa-miR-765 promoter -R	CCCAAGCTTCTCCTCCAGAAACCCCAGTT
D. miR-765 target sites into pMIR-REPORT vector	
3'UTR HMGA1	
HMGA1 3 UTR-F	GGACTAGTCCTGCTCCTCACTGGAGGAG
HMGA1 3 UTR-R	CCCAAGCTTAGTAACTGCAAATAGGAAACC
Complementary hsa-miR-765 recognition site	
miR-765-Insert-sense oligo	CTAGTTTGCATCACCTTCCTTCTCCTCCACCCA
miR-765-Insert-antisense oligo	AGCTTGGGTGGAGGAGAAGGAAGGTGATGCAAA
E. Chromatin Immunoprecipitation assay	
ChIP-PCR-F	AAGTGATCCACCCACCTCAG
ChIP-PCR-R	CCAGTTTCTCGTAGGGCTTG