

Supplementary Table S1. Primer sequences in this study

CircDOCK1	Forward	5'- CCAGAGGCACGTCCAGATTA -3'
	Reverse	5'- AGGAAACTCCGCGTCTAGG -3'
miR-186	Forward	5'- TTAATTCCGATAACGAACGAGA -3'
	Reverse	5'- CGCTGAGCCAGTCAGTG TAG -3'
DNMT3A	Forward	5'- TATGAACAGGCCGTTGGCATC -3'
	Reverse	5'- AAGAGGTGGCGGATGACTGG -3'
GAPDH	Forward	5'- CCTGCCGGTGACTAACCCTG -3'
	Reverse	5'- TCCACCACTGACACGTTGGC -3'