Figure S1. Expression levels of miR-330-5p, miR-193a-5p and miR-326. Cells were transfected with NC or miR-330-5p/miR-193a-5p/miR-326 mimics. RT-qPCR results demonstrated that the relative expression levels of (A) miR-330-5p, (B) miR-193a-5p and (C) miR-326 were significantly upregulated by corresponding mimics. Cells were transfected with NC or miR-330-5p/miR-193a-5p/miR-326 inhibitor. RT-qPCR results demonstrated that the relative expression levels of (D) miR-330-5p, (E) miR-193a-5p and (F) miR-326 were significantly downregulated by corresponding inhibitor. $^{**}P<0.01$. miR, microRNA; NC, negative control; RT-q, reverse transcription-quantitative.

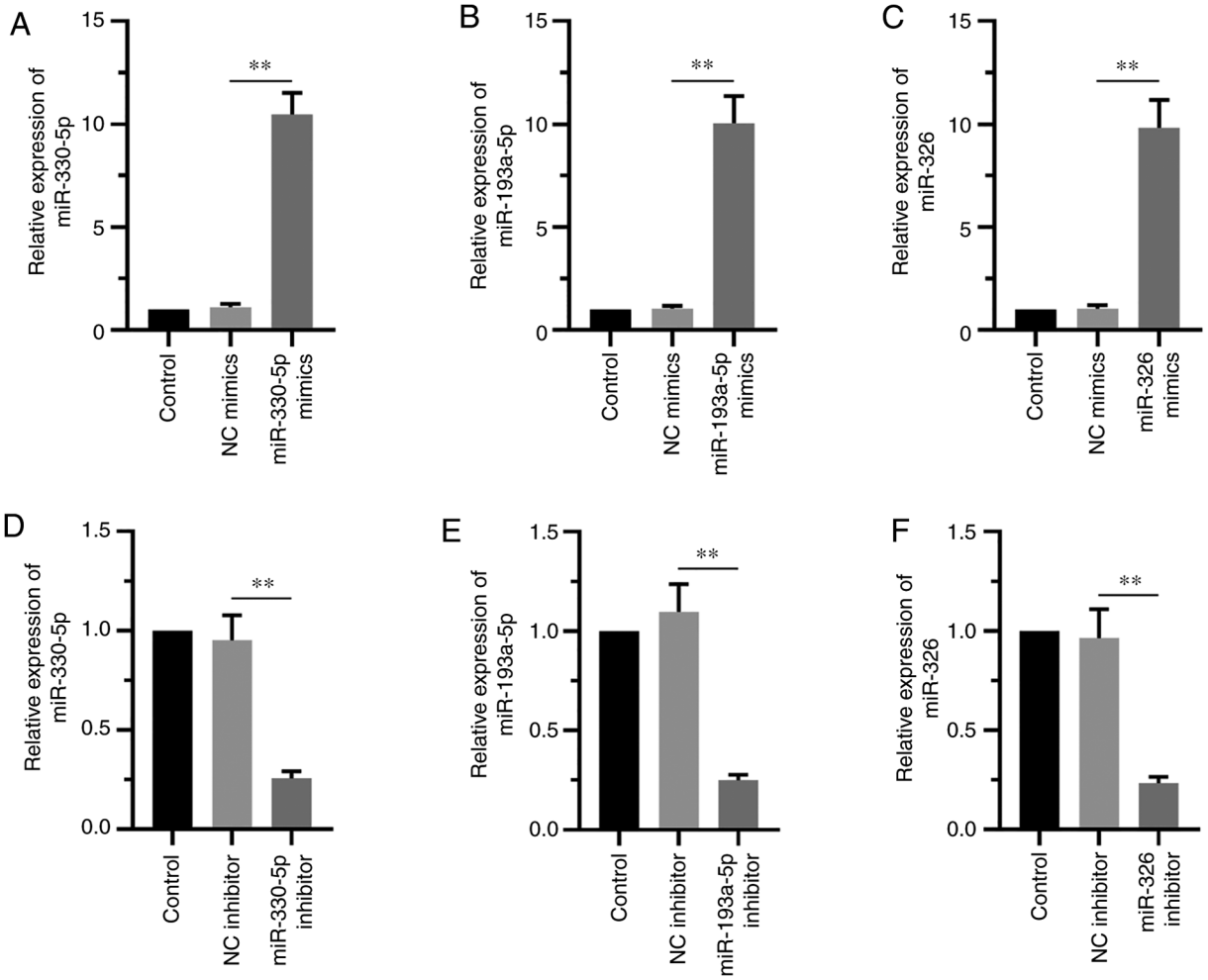

Table SI. Sequences of transfected oligonucleotides.

Oligonucleotide	Sequence (5'→3')
NC siRNA	
Sense	UUCUCCGAACGUGUCACGUTT
Antisense	ACGUGACACGUUCGGAGAATT
hsa_circ-0001666 siRNA1	
Sense	GAUGACCAUUCCAGAUCCUUUTT
Antisense	AAAGGAUCUGGAAUGGUCAUCTT
hsa_circ-0001666 siRNA2	
Sense	AGAUGACCAUUCCAGAUCCUUTT
Antisense	AAGGAUCUGGAAUGGUCAUCUTT
hsa_circ-0001666 siRNA3	
Sense	CAGAUGACCAUUCCAGAUCCUTT
Antisense	AGGAUCUGGAAUGGUCAUCUGTT
hsa_circ-0001666 siRNA4	
Sense	UCCAGAUCCUUUCCCGGAGUUTT
Antisense	AACUCCGGGAAAGGAUCUGGATT
hsa_circ-0001666 siRNA5	
Sense	CAUUCCAGAUCCUUUCCCGGATT
Antisense	UCCGGGAAAGGAUCUGGAAUGTT
NC shRNA	TTCTCCGAACGTGTCACGTttcaagaga
	ACGTGACACGTTCGGAGAAttttt
hsa_circ_0001666 shRNA	GATGACCATTCCAGATCCTTTttcaagaga
	AAAGGATCTGGAATGGTCATCttttt
NC mimics	UUCUCCGAACGUGUCACGUTT
miR-330-5p mimics	UCUCUGGGCCUGUGUCUUAGGC
miR-193a-5p mimics	UGGGUCUUUGCGGGCGAGAUGA
miR-326 mimics	CCUCUGGGCCCUUCCUCCAG
NC inhibitor	UUGUACUACACAAAAGUACUG
miR-330-5p inhibitor	GCCUAAGACACAGGCCCAGAGA
miR-193a-5p inhibitor	UCAUCUCGCCCGCAAAGACCCA
miR-326 inhibitor	CUGGAGGAAGGGCCCAGAGG

NC, negative control; si, small interfering; sh, short hairpin; miR, microRNA.

Table SII. Primer sequences.

Primer	Sequence (5'→3')
hsa_circ_0001666 convergent	
Forward	CAAGGATTTGTGGGAAGT
Reverse	CTAGGCAGGTGCTGGTGA
hsa_circ_0001666 divergent	
Forward	CGACAGCGGGTCTACTCA
Reverse	GGTGCTCTGCCAGTTCTT
FAM120B	
Forward	ACCAAGAGCCAGAAATACAG
Reverse	ATCAAGAGGCAGCACAAA
ETV4	
Forward	GGGCATCCAGAAGAACCG
Reverse	AACGCTCACCAGCCACCT
β-actin	
Forward	GGCACCCAGCACAATGAA
Reverse	TAGAAGCATTTGCGGTGG
miR-330-5p	
Forward	TCTCTGGGCCTGTGTCTTAGGC
Reverse	GCAGGGTCCGAGGTATTC
miR-193a-5p	
Forward	TGGGTCTTTGCGGGCGAGATGA
Reverse	GCAGGGTCCGAGGTATTC
miR-326	
Forward	CCTCTGGGCCCTTCCTCCAG
Reverse	GCAGGGTCCGAGGTATTC
U6	
Forward	GCTTCGGCAGCACATATACT
Reverse	GCAGGGTCCGAGGTATTC

FAM, family with sequence similarity; ETV, ETS variant transcription factor; miR, microRNA.