

Table SI. Sequences of the miRNA-93 mimics/inhibitors used for transfection.

Name	Sequence, 5'→3'
miRNA-93 inhibitor	CUACCUGCACGAACAGCACUUUG
miRNA-93 mimics	Sense: CAAAGUGCUGUUCGUGCAGGUAG Antisense: ACCUGCACGAACAGCACUUUGUU
miRNA, microRNA.	

Table SII. Sequence of mature miRs.

miR	miRBase accession no.	Sequence, 5'→3'
hsa-miR-93	MIMAT0000093	CAAAGUGCUGUUCGUGCAGGUAG
hsa-miR-106b	MIMAT0000680	UAAAGUGCUGACAGUGCAGAU
hsa-miR-210	MIMAT0026475	AGCCCCUGCCCACCGCACACUG
hsa-miR-374b	MIMAT0004955	AUAUAAUACAACCUGCUAAGUG
hsa-miR-125b	MIMAT0000423	UCCUGAGACCCUAACUUGUGA

miR, microRNA.

Table SIII. Reverse transcription-quantitative PCR primers.

Gene	Forward primer, 5' → 3'	Reverse primer, 5' → 3'
E2F1	CCGGGAATGAAGGTGAACA	AGAGCAAAGGGCCGAAAGT
E2F2	GAAGTCGGTGCAGTCGAGA	GCTTAGAGATCGCCGCTTGG
E2F3	ACCAACGTTCTGGAAGGCAT	TTGACAGGCCTTGACTG
CCND1	GATGCCAACCTCCTCAACGA	GGAAGCGGTCCAGGTAGTTC
CCNE1	CCTGTAAGTACTGAGCTGGGCAA	TTTTGGCTGCAGAAGAGGGT
GRB2	GTTCAACTCCCTCAACGAGC	TCCTGGGCTGAGAAGTCAAAC
MYC	AGAGAAGCTGGCCTCCTACC	AGAGAATCCGAGGACGGAGA
$\beta$ -actin	AAGGATTCCTATGTGGGCGAC	CGTACAGGGATAGCACAGCC

E2F, E2F transcription Factor; CCN, cyclin; GRB2, growth factor receptor bound protein 2.

Table SIV. Hub genes identified in the cytoHubba network.

A, Upregulated

Gene	Score
CDC16	$3.04 \times 10^{64}$
RPS27A	$3.04 \times 10^{64}$
ANAPC7	$3.04 \times 10^{64}$
UBC	$3.04 \times 10^{64}$
UBE2V2	$3.04 \times 10^{64}$
UBE2M	$3.04 \times 10^{64}$
ANAPC5	$3.04 \times 10^{64}$
ATG7	$3.04 \times 10^{64}$
TCEB1	$3.04 \times 10^{64}$
HUWE1	$3.04 \times 10^{64}$

B, Downregulated

Gene	Score
RPS3A	$1.12 \times 10^9$
RPS28	$1.12 \times 10^9$
RPL35A	$1.12 \times 10^9$
RPLP0	$1.12 \times 10^9$
RPL3	$1.12 \times 10^9$
RPS7	$1.12 \times 10^9$
RPS12	$1.12 \times 10^9$
RPL29	$1.08 \times 10^9$
ETF1	$1.08 \times 10^9$
PABPC1	$1.08 \times 10^9$