

miR-99a-5p wild-type vector

gatattttccaggccttacgaccctgggctc**acgggt**acctatttatatg

miR-99a-5p deletion-type vector

gatattttccaggccttacgaccctgggctc-----acctatttatatgctcagtg

miR-99a-3p wild-type vector

ttctgctttcctaggggactctt**gagctt**agaaactcatcgtaacttg

miR-99a-3p deletion-type vector

ttctgctttcctaggggactctt-----agaaactcatcgtaacttgacct

Figure S1. Vector inserted sequences.

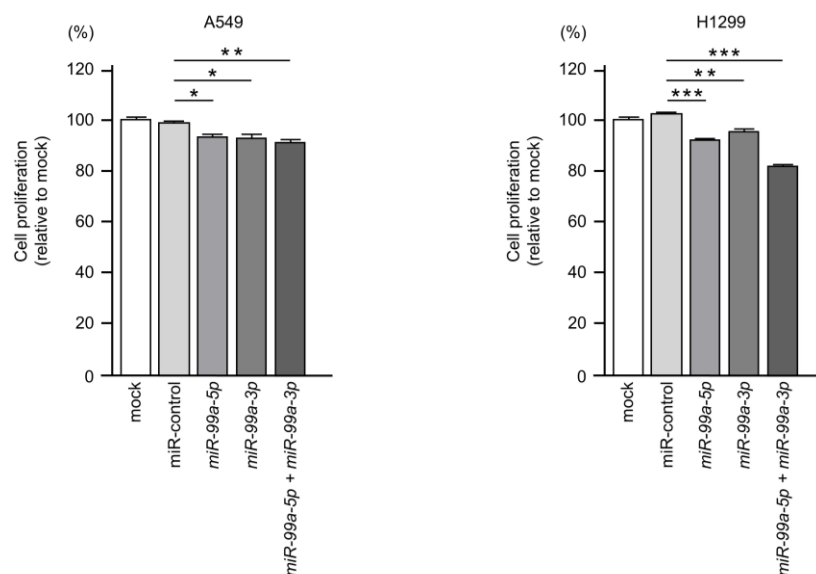


Figure 2. Effects of ectopic expression of *miR-99a-5p* and *miR-99a-3p* on LUAD cells. Cell proliferation was determined by XTT assays 72 h following transfection with *miR-99a-5p* (5nM), *miR-99a-3p* (5nM) or *miR-99a-5p* (5nM) + *miR-99a-3p* (5nM) (* $p < 0.05$, ** $p < 0.01$, *** $p < 0.0001$).

Table S1. Reagent used in this study.

Antibody	Dilution	Catalog number	Company
FAM64A	IHC 1:100	ab251896	Abcam, Cambridge, UK

miRNA species	Concentration	Assay ID	Company
	n		
<i>miR-99a-5p</i>	10 nM	PM 10719	Applied Biosystems, Foster City, CA, USA
<i>miR-99a-3p</i>	10 nM	PM 12983	Applied Biosystems, Foster City, CA, USA
anti-miR Negative Control #1	10 nM	AM 17010	Applied Biosystems, Foster City, CA, USA

siRNA	Concentration	Catalog number	Company
	n		
siFAM64A	10 nM	HSS147670 and HSS147671	Invitrogen, Carlsbad, CA, USA

primer and probe	Assay ID	Company
<i>FAM64A</i>	Hs00760463_m1	Applied Biosystems, Foster City, CA, USA
<i>GUSB</i>	Hs00939627_m1	Applied Biosystems, Foster City, CA, USA

Plasmid vector	Catalog number	Company
psiCHECK-2 vector	C8021	Promega, Madison, WI, USA

Table S2. Characteristics of the patients used in immunostaining.

Patients	Age	Sex	T	N	M	Pathological stage	Type
A	47	M	2	0	0	IB	LUAD
B	56	F	2	1	0	IIA	LUAD
C	67	M	2	0	0	IB	LUAD
D	16	F	-	-	-	-	Normal lung tissue

LUAD: lung adenocarcinoma

Table S3. 23 target genes analyzed by Benjamini-Hochberg method.

Entrez GeneID	Gene symbol	<i>p</i> -value	i	$q_i = p_i * N / i$	FDR threshold	Judgement
983	<i>CDK1</i>	0.0003	3	0.0023	0.05	Significant
4288	<i>MKI67</i>	0.0005	4	0.0029	0.05	Significant
57082	<i>CASC5</i>	0.0003	2	0.0035	0.05	Significant
4751	<i>NEK2</i>	0.0002	1	0.0046	0.05	Significant
6241	<i>RRM2</i>	0.002	6	0.0077	0.05	Significant
1033	<i>CDKN3</i>	0.0017	5	0.0078	0.05	Significant
3832	<i>KIF11</i>	0.0034	8	0.0098	0.05	Significant
4173	<i>MCM4</i>	0.0032	7	0.0105	0.05	Significant
1063	<i>CENPF</i>	0.0059	9	0.0151	0.05	Significant
1163	<i>CKS1B</i>	0.0073	10	0.0168	0.05	Significant
11130	<i>ZWINT</i>	0.0094	12	0.0180	0.05	Significant
84908	<i>FAM136A</i>	0.0087	11	0.0182	0.05	Significant
55215	<i>FANCI</i>	0.0108	13	0.0191	0.05	Significant
56888	<i>KCMF1</i>	0.0125	14	0.0205	0.05	Significant
417685	<i>FAM64A</i>	0.0175	15	0.0268	0.05	Significant
64151	<i>NCAPG</i>	0.0208	16	0.0299	0.05	Significant
151648	<i>SGOL1</i>	0.0235	18	0.0300	0.05	Significant
157570	<i>ESCO2</i>	0.0235	17	0.0318	0.05	Significant
10797	<i>MTHFD2</i>	0.0321	19	0.0389	0.05	Significant
11260	<i>XPOT</i>	0.0669	20	0.0769	0.05	NS
55854	<i>ZC3H15</i>	0.0942	22	0.0985	0.05	NS
51290	<i>ERGIC2</i>	0.0929	21	0.1017	0.05	NS
993	<i>CDC25A</i>	0.1585	23	0.1585	0.05	NS

NS: not significant

Table S4A. Significantly enriched annotations regulated by *miR-99a-5p* in LUAD cells.

Genes	<i>p</i> -value	Annotations
<i>CKS1B</i> <i>CENPF</i> <i>CDC25</i> <i>A</i> <i>FAM64</i> <i>A</i>	4.14×10^{-5}	(GO) 0051301: cell division
<i>CKS1B</i> <i>CENPF</i> <i>CDC25</i> <i>A</i> <i>FAM64</i> <i>A</i>	0.000137009	(GO) 0007049: cell cycle
<i>CKS1B</i> <i>CDC25</i> <i>A</i>	0.00127197	(GO) 0000079: regulation of cyclin-dependent protein serine/threonine kinase activity
<i>CDC25</i> <i>A</i>	0.00529465	(GO) 0110032: positive regulation of G2/M1 transition of meiotic cell cycle
<i>CKS1B</i> <i>CDC25</i> <i>A</i>	0.00547439	(GO) 0008283: cell population proliferation

Table S4B. Significantly enriched annotations regulated by *miR-99a-3p* in LUAD cells.

Genes	<i>p</i> -value	Annotations
ZWINT	9.15×10^{-14}	(GO) 0007049: cell cycle
CDKN3		
CDK1		
NEK2		
MKI67		
MCM4		
KIF11		
ESCO2		
SGOL1		
NCAPG		
CASC5		
FANCI		
FAM64A		
ZWINT	1.10×10^{-7}	(GO) 0051301: cell division
CDK1		
NEK2		
KIF11		
SGOL1		
NCAPG		
CASC5		
FAM64A		
NEK2	7.25×10^{-5}	(GO) 0007059: chromosome segregation
ESCO2		
SGOL1		
CASC5		
ZWINT	0.000165787	(GO) 0000070: mitotic sister chromatid segregation
NEK2		
SGOL1		
NEK2	0.000243324	(GO) 0046602: regulation of mitotic centrosome separation
KIF11		

Table S4C. Pathways regulated by *miR-99a-5p* in LUAD cells.

Cell division pathway	
Gene symbol	Gene name
<i>CKS1B</i>	CDC28 protein kinase regulatory subunit 1B
<i>CENPF</i>	Centromere protein F
<i>CDC25A</i>	Cell division cycle 25A
<i>FAM64A</i>	Family with sequence similarity 64 member A
Cell cycle pathway	
Gene symbol	Gene name
<i>CKS1B</i>	CDC28 protein kinase regulatory subunit 1B
<i>CENPF</i>	Centromere protein F
<i>CDC25A</i>	Cell division cycle 25A
<i>FAM64A</i>	Family with sequence similarity 64 member A
Regulation of cyclin-dependent protein serine/threonine kinase activity pathway	
Gene symbol	Gene name
<i>CKS1B</i>	CDC28 protein kinase regulatory subunit 1B
<i>CDC25A</i>	Cell division cycle 25A
Positive regulation of G2/M1 transition of meiotic cell cycle pathway	
Gene symbol	Gene name
<i>CDC25A</i>	Cell division cycle 25A
Cell population proliferation pathway	
Gene symbol	Gene name
<i>CKS1B</i>	CDC28 protein kinase regulatory subunit 1B
<i>CDC25A</i>	Cell division cycle 25A

Table S4D. Pathways regulated by *miR-99a-3p* in LUAD cells.

Cell cycle pathway	
Gene symbol	Gene name
<i>ZWINT</i>	ZW10 interacting kinetochore protein
<i>CDKN3</i>	Cyclin dependent kinase inhibitor 3
<i>CDK1</i>	Cyclin dependent kinase 1
<i>NEK2</i>	NIMA related kinase 2
<i>MKI67</i>	Marker of proliferation Ki-67
<i>MCM4</i>	Minichromosome maintenance complex component 4
<i>KIF11</i>	Kinesin family member 11
<i>ESCO2</i>	Establishment of sister chromatid cohesion N-acetyltransferase 2
<i>SGOL1</i>	Shugoshin 1
<i>NCAPG</i>	Non-SMC condensin I complex subunit G
<i>CASC5</i>	Cancer susceptibility candidate 5
<i>FANCI</i>	FA complementation group I
<i>FAM64A</i>	Family with sequence similarity 64 member A
Cell division pathway	
Gene symbol	Gene name
<i>ZWINT</i>	ZW10 interacting kinetochore protein
<i>CDK1</i>	Cyclin dependent kinase 1
<i>NEK2</i>	NIMA related kinase 2
<i>KIF11</i>	Kinesin family member 11
<i>SGOL1</i>	Shugoshin 1
<i>NCAPG</i>	Non-SMC condensin I complex subunit G
<i>CASC5</i>	Cancer susceptibility candidate 5
<i>FAM64A</i>	Family with sequence similarity 64 member A
Chromosome segregation pathway	
Gene symbol	Gene name
<i>NEK2</i>	NIMA related kinase 2
<i>ESCO2</i>	Establishment of sister chromatid cohesion N-acetyltransferase 2
<i>SGOL1</i>	Shugoshin 1
<i>CASC5</i>	Cancer susceptibility candidate 5
Mitotic sister chromatid segregation pathway	
Gene symbol	Gene name
<i>ZWINT</i>	ZW10 interacting kinetochore protein
<i>NEK2</i>	NIMA related kinase 2
<i>SGOL1</i>	Shugoshin 1
Regulation of mitotic centrosome separation pathway	

Gene symbol	Gene name
<i>NEK2</i>	NIMA related kinase 2
<i>KIF11</i>	Kinesin family member 11