Additional file 5. Previous literatures showing the implication of miR-15/107 family and cell cycle related biological events

miRNAs	Role in cell cycle related events	References
	Human miR-15a-5p inhibits pituitary tumor cell	(1)
	proliferation, invasion and migration by targeting Sox5.	
	miR-15a inhibits cell proliferation and epithelial to	(2)
	mesenchymal transition in pancreatic ductal	
	adenocarcinoma by down-regulating Bmi-1 expression.	
	Up-regulation of miR-15a-5p suppresses cell proliferation,	(3)
	migration and invasion of these ectopic stem cells by	
	targeting the 3' untranslated region of VEGFA.	
	Increased miR-15a suppresses the Cyclin E1, regulated	(4)
	CDK2 and promotes the $G_1/S$ transition.	
	Overexpression of VEGFA or inhibition of miR-15a-5p	(5)
	promotes cell proliferation in osteoarthritis chondrocytes.	
miR-15a-5p	miR-15a-5p suppresses the proliferation, motility,	(6)
mix-15a-5p	invasiveness and angiogenic potential and increased	
	apoptosis when combined with docetaxel in	
	chemotherapy-resistant ovarian cancer.	
	High levels of miR-15a-5p in the young poor ovarian	(7)
	response group represses granulosa cell proliferation by	
	regulating the PI3K-AKT-mTOR signaling pathway and	
	promoted apoptosis through BCL2 and BAD.	
	miR-15a-5p suppresses endometrial cancer cell growth via	(8)
	Wnt/ $\beta$ -catenin signaling pathway by inhibiting WNT3A.	
	Ectopic overexpression of miR-15a-5p suppresses cancer	(9)
	proliferation, induces cell cycle arrest in HepG2 or	
	SNU-182 cells <i>in vitro</i> , and inhibits HCC tumor growth <i>in</i>	
	vivo.	
miR-16-5p	MicroRNA-16-5p overexpression suppresses proliferation	(10)
	and invasion as well as triggers apoptosis by targeting	
	VEGFA in breast carcinoma.	
	Acute myeloid leukemia (AML) is associated with	(11)
	heterogeneous clonal proliferation, while numerous	
	microRNAs are involved in AML and associated with the	
'D 100 0	expression level of VEGF, including miR-16-5p.	(12)
miR-103a-3p	Overexpression of miR-103a-3p inhibits the proliferation	(12)

		,
	and osteogenic differentiation of hADSCs (human adipose tissue-derived stromal cells), downregulates protein and	
	mRNA levels of predicted target of miR-103a-3p (CDK6	
	and DICER1).	
	Over-expression of circTCF25 could down-regulate	(13)
	miR-103a-3p and miR-107, increase CDK6 expression, and	
	promote proliferation and migration <i>in vitro</i> and <i>in vivo</i> .	
	MicroRNA-107 promotes proliferation of gastriccancer cells	(14)
	by targeting cyclin dependent kinase 8.	
	miR-107 promotes hepatocellular carcinoma cell	(15)
	proliferation by targeting Axin2.	
	MicroRNA-107 inhibits tumor growth and metastasis by	(16)
	targeting the BDNF-mediated PI3K/AKT pathway in human	
	non-small lung cancer.	
	Up-regulation of MicroRNA-107 induces proliferation in	(17)
	human gastric cancer cells by targeting the transcription	
	factor FOXO1.	
	Up-regulation of miR-107 decreases proliferation rates,	(18)
	colony-forming abilities, tumorigenicity and increases	
	apoptosis rates by targeting DKK-1.	
	Treatment with miR-107 significantly blockes cell	(19)
miR-107	proliferation, DNA replication, colony formation, and	
	invasion in SCC25 and CAL27 cells by targeting protein	
	kinase Cɛ.	
	miR-107 directly targets MCL1 and activates ATR/Chk1	(20)
	pathway to inhibit proliferation, migration and invasiveness	
	of cervical cancer cells.	
	miR-107 overexpression inhibits cell proliferation in human	(21)
	neuroblastoma and ratpituitary adenoma cells by targeting	
	AIP.	
	Decreased miR-107 could inhibit tumor growth in a nude	(22)
	mouse model of triple-negative breast cancer.	(22)
	Overexpression of miR-107 in glioma cells leads to	(23)
	inhibition of HBMVEC proliferation, migration and tube	
	formation ability.	(24)
	MicroRNA-107 inhibits U87 glioma cell growth and invasion by target Notch?	(24)
	invasion by target Notch2.	(25)
	miR-107 overexpression suppresses cell proliferation in	(25)

	glioma cells, whereas miR-107 knockdown promotes cell	
	growth in MO59K cell line.	
	P53-induced microRNA-107 inhibits proliferation of glioma	(26)
	cells and down-regulates the expression of CDK6 and Notch-2.	
	MiR-107 suppresses proliferation of hepatoma cells through targeting HMGA2 mRNA 3'UTR	(27)
	miR-107 can simultaneously target large tumor suppressor 2	(28)
	(LATS2) and synchronically modulate its expression,	
	thereby promoting proliferation and invasion of GAC cells.	
	miR-107 improves the proliferative and migratory ability of HepG2 cells.	(29)
	miR-107 mimic leads to a significant increase in the $G_0/G_1$ population and a significant decrease in the $G_2/M$ population, and functions as a tumor suppressor in human esophageal squamous cell carcinoma.	(30)
	Overexpression of miR-195-5p significantly suppresses renal cell carcinoma growth in vitro and in vivo by targeting	(31)
	REGγ.	
	Overexpression of miR-195-5p inhibits cell proliferation,	(32)
	reduces cell colony formation, suppresses cell migration and	(- )
	causes an accumulation of cells in the $G_1$ phase of the cell	
	cycle by regulating CCNE1 in breast cancer.	
	miR-195-5p suppresses the proliferation, migration, and	(33)
	invasion of oral squamous cell carcinoma by targeting	
	TRIM14.	(2.4)
miR-195-5p	Overexpression of miR-195-5p in hepatoma cells reduces	(34)
	PHF19 expression, leads to suppression of hepatoma cell	
	invasion, migration and proliferation in <i>vitro</i> .	(25)
	MicroRNA-195-5p suppresses glucose uptake and	(35)
	proliferation of human bladder cancer T24 cells by	
	regulating GLUT3 expression.	(2.6)
	MicroRNA-195-5p suppresses osteosarcoma cell	(36)
	proliferation and invasion by suppressing naked cuticle	
	homolog 1.	(27)
	Anti-apoptosis endothelial cell-secreted microRNA-195-5p	(37)
	promotes pulmonary arterial smooth muscle cell	
	proliferation and migration in pulmonary arterial	
	hypertension.	

	miR-424-5p promotes proliferation of gastric cancer by	(38)
	targeting Smad3 through TGF-β signaling pathway.	
	Downregulated miR-424-5p causes the overexpression of	(39)
	MEK1 and cyclin E1 in the psoriatic keratinocytes, resulting	
	in the keratinocyte overgrowth and hyper-proliferation.	
	miR-424-5p mimics could decrease the invasion, metastasis	(40)
	and proliferation of esophageal squamous carcinom cells,	
	while SMAD7 is a specific target gene for miR-424-5p.	
miR-424-5p	Ectopic expression of miR-424 promotes proliferation and	(41)
	invasion of gastric cancer cells by targeting LATS1 gene.	
	miR-424-5p plays its role as an anti-onco gene in cervical	(42)
	cancer cell growth by targeting KDM5B via the Notch	
	signaling pathway.	
	miR-424-5p increases proliferation, migration and invasion	(43)
	of pancreatic cancer cells, while inhibits cell apoptosis.	
	miR-424-5p could prevent the proliferation, migration, and	(44)
	invasion in NSCLC cells.	
·D 407 5	miR-497-5p inhibits cell proliferation and invasion by	(45)
miR-497-5p	targeting KCa3.1 in angiosarcoma.	
·D 502 5	miR-503-3p (the passenger strand of miR-503-5p) inhibits	(46)
miR-503-5p	cell proliferation in lung cancer.	
	miR-646 negatively regulates NOB1 and inhibits the	(47)
miR-646	proliferation and migration of renal cancer cells.	
	miR-646 inhibits cell proliferation by targeting FOXK1 in	(48)
	gastric cancer.	
	Overexpression of miR-646 inhibits cell proliferation and	(49)
	metastasis in lung cancer.	
	Overexpression of miR-646 inhibits cell proliferation,	(50)
	migration, and invasion in osteosarcoma cells.	
miR-6838-5p	None	-
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