## Supplementary material

Table 1: Downstream molecules list of miR-100-5p

Target gene	Representative transcript	Gene name	Number of 3P-seq tags supporting UTR+5	Link to	(	Conserved sites				Poorly conserved site				
				Link to sites in UTRs	total	8mer	7mer- m8	7mer- A1	total	8mer	7mer- m8	7mer- A1		Representativ miRNA
PDR1	ENST00000199448.4	ependymin related 1	84	Sites in UTR	1	1	0	0	1	0	0	1	0	hsa-miR-100-5
P1AR	ENST00000274000.5	adaptor-related protein complex 1 associated regulatory protein	89	Sites in UTR	1	1	0	0	0	0	0	0	0	hsa-miR-99b-5
T6GALNAC4	ENST00000335791.5	ST6 (alpha-N-acetyl-neuraminyl-2,3-beta-galactosyl-1,3)-N-acetylgalactosaminide alpha-2,6-sialyltransferase 4	178	Sites in UTR	1	1	0	0	0	0	0	0	0	hsa-miR-99a-5
TC39A	ENST00000530004.1	tetratricopeptide repeat domain 39A	174	Sites in UTR	1	1	0	0	0	0	0	0	1	hsa-miR-99a-5
S3ST2	ENST00000261374.3	heparan sulfate (glucosamine) 3-O-sulfotransferase 2	5	Sites in UTR	1	1	0	0	0	0	0	0	0	hsa-miR-100-5
AZ2A	ENST00000379441.3	bromodomain adjacent to zinc finger domain, 2A	48	Sites in UTR	2	1	1	0	0	0	0	0	0	hsa-miR-99a-
BTBD8	ENST00000295568.4	kelch repeat and BTB (POZ) domain containing 8	22	Sites in UTR	1	1	0	0	0	0	0	0	0	hsa-miR-100-
KBP5	ENST00000536438.1	FK506 binding protein 5	32	Sites in UTR	1	1	0	0	0	0	0	0	0	hsa-miR-100-
ES7	ENST00000541682.2	hairy and enhancer of split 7 (Drosophila)	416	Sites in UTR	1	1	0	0	0	0	0	0	0	hsa-miR-99b-5
S3ST3B1	ENST00000360954.2	heparan sulfate (glucosamine) 3-O-sulfotransferase 3B1	1489	Sites in UTR	1	1	0	0	0	0	0	0	0	hsa-miR-99a-5
GFR3	ENST00000340107.4	fibroblast growth factor receptor 3	1085	Sites in UTR	1	1	0	0	0	0	0	0	0	hsa-miR-99a-
TDSPL	ENST00000443503.2	CTD (carboxy-terminal domain, RNA polymerase II, polypeptide A) small phosphatase-like	372	Sites in UTR	1	1	0	0	0	0	0	0	0	hsa-miR-99b-
AVER2	ENST00000371072.4	ribonucleoprotein, PTB-binding 2	218	Sites in UTR	1	0	1:	0	0	0	0	0	0	hsa-miR-99a-
MARCA5	ENST00000283131.3	SWI/SNF related, matrix associated, actin dependent regulator of chromatin, subfamily a, member 5	478	Sites in UTR	1	0	1	0	0	0	0	0	0	hsa-miR-100-
RRC8B	ENST00000330947.2	leucine rich repeat containing 8 family, member B	326	Sites in UTR	1	1	0	0	0	0	0	0	0	hsa-miR-99a-
OXA1	ENST00000355633.5	homeobox A1	157	Sites in UTR	1	0	1	0	0	0	0	0	0	hsa-miR-99a-
MPS	ENST00000496455.2	guanine monphosphate synthase	867	Sites in UTR	1	1	0	0	0	0	0	0	0	hsa-miR-99a-
MTOR	ENST00000376838.	1 mechanistic target of rapamycin (serine/threonine kinase)	653	Sites in UTF	3 1	0	4	0	0	0	0	0	0	hsa-miR-100
NR6A1	ENST00000376838.		1082	Sites in UTF			0	0	0		0	0	0	hsa-miR-998
FZD8	ENST00000374694.		791	Sites in UTF			1	0	0	The state of	0	0	1	hsa-miR-10
			10.500											
NTRK3 TMEM30A		2 neurotrophic tyrosine kinase, receptor, type 3	7	Sites in UTF			0	0	0	1122574	0	0	0	hsa-miR-99a
RASGRP3	ENST00000230461.	6 transmembrane protein 30A 3 RAS guanyl releasing protein 3 (calcium and DAG-regulated)	24	Sites in UTI				0	0		0	0	0	hsa-miR-100
CLDN11	ENST00000402338.		1116	Sites in UTI			1	0	0		0	0	0	hsa-miR-100
TRIB2		4 tribbles pseudokinase 2	68	Sites in UTF			1	0	0	1000	0	0	0	hsa-miR-10
TRIB1	ENST00000311922.	3 tribbles pseudokinase 1	313	Sites in UTF	₹ 1	0	0	1	0	0	0	0	0	hsa-miR-99
PPP3CA	ENST00000512215.	1 protein phosphatase 3, catalytic subunit, alpha isozyme	259	Sites in UTF	₹ 1	0	1	0	0	0	0	0	0	hsa-miR-99
ZBTB7A	ENST00000322357.	4 zinc finger and BTB domain containing 7A	37	Sites in UTF	₹ 1	0	0	1	0	0	0	0	1	hsa-miR-10
GRHL1	ENST00000405379.	2 grainyhead-like 1 (Drosophila)	147	Sites in UTI	₹ 1	0	0	1	0	0	0	0	0	hsa-miR-99
TMEM135	ENST00000340353.	7 transmembrane protein 135	963	Sites in UTF	₹ 1	0	0	- 1	0	0	0	0	0	hsa-miR-100
ST5	ENST00000526757.		139	Sites in UTF	₹ 1	100	1	0	0		0	0	0	hsa-miR-100
PPP1CB	ENST00000395366.		5795	Sites in UTF	(A)	100		1	0	1500	0	0	0	hsa-miR-10
NIPBL	ENST00000448238.	2 Nipped-B homolog (Drosophila)	250	Sites in UTF	₹ 1	0	0	1	0	0	0	0	0	hsa-miR-99
MTMR3	ENST00000333027.	3 myotubularin related protein 3	123	Sites in UTI	٦ 1	0	1	0	0	0	0	0	0	hsa-miR-99
PI15		2 peptidase inhibitor 15	12	Sites in UTI			0	1	0	1,000	0	0	0	hsa-miR-99
DESI2	ENST00000302550.	11 desumoylating isopeptidase 2	147	Sites in UTF	₹ 1	0	0	1	0	0	0	0	0	hsa-miR-99
INSM1	ENST00000310227.	1 insulinoma-associated 1	20	Sites in UTF	₹ 1	0	1	0	0	0	0	0	0	hsa-miR-99a
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CEP85	ENST00000252992.	4 centrosomal protein 85kDa	471	Sites in UTI	₹ 1	0	0	1	0	0	0	0	0	hsa-miR-99

Table 1: Continued

SMARCD1	ENST00000394963.4	SWI/SNF related, matrix associated, actin dependent regulator of chromatin, subfamily d, member 1	952	Sites in UTR	1	0	0	1	0	0	0	0	0	hsa-miR-99b-5p
NXF1	ENST00000531709.2	nuclear RNA export factor 1	1017	Sites in UTR	1	0	0	1	0	0	0	0	0	hsa-miR-100-5p
SLC44A1	ENST00000374720.3	solute carrier family 44 (choline transporter), member 1	407	Sites in UTR	1	0	1	0	0	0	0	0	0	hsa-miR-100-5p
ETV3	ENST00000368192.4	ets variant 3	37	Sites in UTR	1	0	1	0	0	0	0	0	0	hsa-miR-99a-5p
KDM6B	ENST00000254846.5	lysine (K)-specific demethylase 6B	8	Sites in UTR	1	0	1	0	0	0	0	0	0	hsa-miR-99b-5p
RRAGD	ENST00000369415.4	Ras-related GTP binding D	644	Sites in UTR	1	0	1	0	1	0	1	0	0	hsa-miR-100-5p
ZZEF1	ENST00000381638.2	zinc finger, ZZ-type with EF-hand domain 1	83	Sites in UTR	1	0	1	0	0	0	0	0	0	hsa-miR-100-5p
TRIM71	ENST00000383763.5	tripartite motif containing 71, E3 ubiquitin protein ligase	67	Sites in UTR	1	0	1	0	0	0	0	0	1	hsa-miR-99a-5p
CYP26B1	ENST00000001146.2	cytochrome P450, family 26, subfamily B, polypeptide 1	1315	Sites in UTR	1	0	1	0	0	0	0	0	0	hsa-miR-100-5p
CDYL2	ENST00000570137.2	chromodomain protein, Y-like 2	239	Sites in UTR	1	0	1	0	0	0	0	0	0	hsa-miR-99b-5p
IGF1R	ENST00000268035.6	insulin-like growth factor 1 receptor	561	Sites in UTR	1	1	0	0	0	0	0	0	0	hsa-miR-99b-5p
TAOK1	ENST00000261716.3	TAO kinase 1	164	Sites in UTR	1	1	0	0	0	0	0	0	0	hsa-miR-100-5p
THAP2	ENST00000308086.2	THAP domain containing, apoptosis associated protein 2	109	Sites in UTR	1	1	0	0	0	0	0	0	0	hsa-miR-100-5p
SATB1	ENST00000338745.6	SATB homeobox 1	312	Sites in UTR	1	1	0	0	0	0	0	0	0	hsa-miR-99b-5p
AGO2	ENST00000220592.5	argonaute RISC catalytic component 2	350	Sites in UTR	1	0	1	0	0	0	0	0	0	hsa-miR-100-5p
BMPR2	ENST00000374574.2	bone morphogenetic protein receptor, type II (serine/threonine kinase)	287	Sites in UTR	1	0	0	1	0	0	0	0	0	hsa-miR-100-5p
ZNRF2	ENST00000323037.4	zinc and ring finger 2	403	Sites in UTR	1	0	0	1	0	0	0	0	0	hsa-miR-100-5p
ICMT	ENST00000343813.5	isoprenylcysteine carboxyl methyltransferase	558	Sites in UTR	1	0	1	0	0	0	0	0	0	hsa-miR-99a-5p
RMND5A	ENST00000283632.4	required for meiotic nuclear division 5 homolog A (S. cerevisiae)	261	Sites in UTR	1	0	0	1	0	0	0	0	0	hsa-miR-99a-5p
FZD5	ENST00000295417.3	frizzled family receptor 5	122	Sites in UTR	1	0	1	0	0	0	0	0	0	hsa-miR-100-5p