

Supplementary Table S1. List of predicted target genes of *TnP*-upregulated *Danio rerio* miR-21, miR-122, miR-731, and miR-26 by Diana microT-CDS (miTG score > 0.7) and TargetScanFish (score + context < -0.2).

miRNA	Ensembl Target Gene Id	Target gene name	Target Gene Synonyms	Description	DIANA microT-CDS	TargetScan
					miTG score	Total context + score
miR-21	ENSDARG00000008034	<i>skib</i>	<i>fc17g02, skiB, wu:fc17g02, wu:fk79g04</i>	v-ski avian sarcoma viral oncogene homolog b	0.99	-0.35
miR-21	ENSDARG000000021443	<i>zfp36l1b</i>	<i>fc06f10, wu:fc06f10, zfp36l1l, zgc:77882</i>	zinc finger protein 36, C3H type-like 1b	0.98	-0.32
miR-21	ENSDARG000000001898	<i>manea</i>	<i>fi29h09, wu:fi29h09, zgc:92825</i>	mannosidase, endo-alpha	0.96	-0.60
miR-21	ENSDARG000000076566	<i>kank3</i>	<i>NBP, Numb-Binding Protein, si:dkeyp-104h9.6</i>	KN motif and ankyrin repeat domains 3	0.96	-0.21
miR-21	ENSDARG000000038845	<i>ldhd</i>	<i>wu:fi36b04, zgc:55447</i>	lactate dehydrogenase D	0.96	-0.65
miR-21	ENSDARG000000074812	<i>fhdcl</i>	<i>si:ch211-62k15.3</i>	formin homology domain containing 1	0.95	-0.42
miR-21	ENSDARG000000008363	<i>mcl1b</i>	<i>zgc:109763</i>	MCL1 apoptosis regulator, BCL2 family member b	0.94	-0.25
miR-21	ENSDARG000000045160	<i>msrb1b</i>	<i>sepx1b, zgc:92822</i>	methionine sulfoxide reductase B1b	0.94	-0.38
miR-21	ENSDARG000000045516	<i>itih2</i>	<i>id:ibd5015, wu:fb34d10, wu:fb96f01, zgc:56119</i>	inter-alpha-trypsin inhibitor heavy chain 2	0.94	-0.39
miR-21	ENSDARG000000007369	<i>tcf7l1b</i>	<i>tcf3b</i>	transcription factor 7 like 1b	0.94	-0.37
miR-21	ENSDARG000000070781	<i>gjd2b</i>	<i>cx35, cx35.1, cx35b, gja9</i>	gap junction protein delta 2b	0.93	-0.54
miR-21	ENSDARG000000052594	<i>nkiras2</i>	<i>zgc:92870</i>	NFKB inhibitor interacting Ras-like 2	0.92	-0.42
miR-21	ENSDARG000000004218	<i>rnd1b</i>	<i>Rnd1a, rnd1l, wu:fk50c04, zgc:153089</i>	Rho family GTPase 1b	0.90	-0.45
miR-21	ENSDARG000000073883	<i>clstn3</i>	<i>si:dkey-197a20.4</i>	calsynenin 3	0.89	-0.38
miR-21	ENSDARG000000016835	<i>tcirg1a</i>	<i>si:dkey-9i23.9, tcirg1</i>	T cell immune regulator 1, ATPase H ⁺ transporting V0 subunit a3a	0.89	-0.63
miR-21	ENSDARG000000020228	<i>usf2</i>	<i>fb12g11, si:dkey-211g8.3, wu:fb12g11</i>	upstream transcription factor 2, c-fos interacting	0.89	-0.31
miR-21	ENSDARG000000075714	<i>hoatz</i>	<i>C5H11orf88, si:dkey-27p23.3</i>	HOATZ cilia and flagella associated protein	0.88	-0.42
miR-21	ENSDARG000000023058	<i>foxo3a</i>	<i>fb50g02, fi33f07, wu:fb50g02, wu:fi33f07, zgc:92176</i>	forkhead box O3A	0.88	-0.26
miR-21	ENSDARG000000011879	<i>foxn1</i>	<i>nude, whnb</i>	forkhead box N1	0.87	-0.52
miR-21	ENSDARG000000005343	<i>ccdc85a1</i>	<i>si:dkey-277e10.1, zgc:92780</i>	coiled-coil domain containing 85A, like	0.86	-0.38
miR-21	ENSDARG000000042029	<i>mbd3b</i>	<i>cb357, cb99, sb:cb357, sb:cb99, wu:fc41c05, zgc:111851</i>	methyl-CpG binding domain protein 3b	0.86	-0.34
miR-21	ENSDARG000000090072	<i>CU462878.2</i>	<i>KRTCAP3</i>	keratinocyte associated protein 3	0.85	-0.21
miR-21	ENSDARG000000075532	<i>slc4a11</i>	<i>si:dkey-12j14.4</i>	solute carrier family 4 member 11	0.85	-0.31
miR-21	ENSDARG0000000036457	<i>cacng6a</i>	<i>si:dkey-234p10.2</i>	calcium channel, voltage-dependent, gamma subunit 6a	0.85	-0.36
miR-21	ENSDARG000000022768	<i>gid8a</i>	<i>zgc:73100</i>	GID complex subunit 8 homolog a	0.85	-0.40
miR-21	ENSDARG000000043482	<i>b3gnt3.1</i>	<i>zgc:136488, zgc:86586</i>	UDP-GlcNAc:betaGal beta-1,3-N-acetylglucosaminyltransferase 3, tandem duplicate 1	0.85	-0.38
miR-21	ENSDARG000000042368	<i>kif26aa</i>	<i>si:dkey-31d8.1</i>	kinesin family member 26Aa	0.85	-0.30
miR-21	ENSDARG000000008904	<i>smarca2</i>	<i>wu:fa56c07, wu:fi27f11, zgc:66238</i>	SWI/SNF related, matrix associated, actin dependent regulator of chromatin, subfamily a, member 2	0.84	-0.31
miR-21	ENSDARG000000074262	<i>nck1a</i>	<i>si:dkey-4m11.4, nck1</i>	NCK adaptor protein 1a	0.83	-0.30
miR-21	ENSDARG000000002013	<i>grb10a</i>	<i>grb10, zgc:91987</i>	growth factor receptor-bound protein 10a	0.83	-0.31
miR-21	ENSDARG000000052957	<i>plcd3a</i>	<i>DrPLC-delta 3A, zgc:158396</i>	phospholipase C, delta 3a	0.83	-0.29
miR-21	ENSDARG000000090914	<i>si:ch211-117k10.3</i>	<i>klf15l, AL929286.1</i>	<i>si:ch211-117k10.3</i> ; orthologous to human KLF15 (Kruppel like factor 15)	0.83	-0.34
miR-21	ENSDARG000000076913	<i>emel</i>	<i>fc30c07, wu:fc30c07</i>	essential meiotic structure-specific endonuclease 1	0.82	-0.34
miR-21	ENSDARG000000052091	<i>rbpjb</i>	<i>Su(H)2</i>	recombination signal binding protein for immunoglobulin kappa J region b	0.82	-0.23
miR-21	ENSDARG0000000005943	<i>htra4</i>	<i>si:dkey-49o11.4</i>	HtrA serine peptidase 4	0.81	-0.31
miR-21	ENSDARG000000037421	<i>egr1</i>	<i>etID309970.14, krox-24, krox24, wu:ff64b05, wu:fq25f01</i>	early growth response 1	0.81	-0.22
miR-21	ENSDARG000000026236	<i>zgc:56585</i>	<i>zgc:77077</i>	<i>zgc:56585</i> ; orthologous to human HPGD (15-hydroxyprostaglandin dehydrogenase).	0.81	-0.37
miR-21	ENSDARG0000000088022	<i>mhc1zfa</i>	N/A	major histocompatibility complex class I ZFA	0.81	-0.24
miR-21	ENSDARG000000002006	<i>rxrb</i>	<i>NR2B2-B, rxrd, unp286</i>	retinoid x receptor, beta b	0.81	-0.28
miR-21	ENSDARG000000039423	<i>AL1</i>	<i>si:ch211-217g15.3</i>	<i>si:ch211-217g15.3</i>	0.81	-0.34
miR-21	ENSDARG000000012306	<i>syt13</i>	<i>zgc:112063</i>	synaptotagmin XIII	0.80	-0.39
miR-21	ENSDARG000000014828	<i>l3mbtl1a</i>	<i>l3mbtl1, si:ch211-262h21.1</i>	L3MBTL histone methyl-lysine binding protein 1a	0.80	-0.33

miR-21	ENSDARG00000017653	<i>rgs13</i>	<i>zgc:163081</i>	regulator of G protein signaling 13	0.80	-0.33
miR-21	ENSDARG00000045352	<i>cst14a.2</i>	<i>cst14a, zgc:56530</i>	cystatin 14a, tandem duplicate 2	0.79	-0.34
miR-21	ENSDARG00000016207	<i>zgc:112408</i>	N/A	<i>zgc:112408</i>	0.79	-0.38
miR-21	ENSDARG000000062164	<i>Atrn</i>	<i>si:dkey-259n11.1</i>	attractin	0.79	-0.29
miR-21	ENSDARG00000043128	<i>cldne</i>	<i>cb84, fb16e12, wu:fb16e12</i>	claudin e	0.79	-0.32
miR-21	ENSDARG00000039434	<i>oprm1</i>	ZFOR2, ZfMOR	opioid receptor, mu 1	0.78	-0.29
miR-21	ENSDARG00000088505	<i>fam217b</i>	<i>si:dkey-211h10.3</i>	family with sequence similarity 217 member B	0.78	-0.37
miR-21	ENSDARG00000040910	<i>ildr1b</i>	<i>fc95b03, ildr1, wu:fc95b03, zgc:101133</i>	immunoglobulin-like domain containing receptor 1b	0.78	-0.37
miR-21	ENSDARG00000076302	<i>DTX4a</i>	<i>dtx1, si:ch211-119e14.2, dtx4</i>	deltex 4, E3 ubiquitin ligase a	0.78	-0.32
miR-21	ENSDARG00000058365	<i>hspb8</i>	<i>fc09c11, wu:fc04b04, wu:fc09c11, zgc:64202</i>	heat shock protein b8	0.78	-0.34
miR-21	ENSDARG00000026444	<i>apob</i>	<i>apoo, mgc4825b, si:rp71-lf1.3, wu:fr42a11, zgc:103766</i>	apolipoprotein O, b	0.78	-0.30
miR-21	ENSDARG00000011459	<i>gsna</i>	<i>cb107, gsn, sb:cb107, u-gelsolin, wu:fi16f06</i>	gelsolin a	0.78	-0.24
miR-21	ENSDARG00000012381	<i>hsc70</i>	<i>zgc:63663</i>	heat shock cognate 70	0.78	-0.38
miR-21	ENSDARG00000079751	<i>megf8</i>	<i>fb77g03, si:dkey-108f8.4, wu:fb77g03, zgc:158741</i>	multiple EGF-like-domains 8	0.77	-0.29
miR-21	ENSDARG00000032010	<i>slc15a2</i>	<i>PEPT2, wu:fc84g07, wu:fi22a10, zgc:152684</i>	solute carrier family 15 member 2	0.77	-0.34
miR-21	ENSDARG00000027992	<i>hao2</i>	<i>zgc:63690</i>	hydroxyacid oxidase 2 (long chain)	0.77	-0.33
miR-21	ENSDARG00000079104	<i>mflas1</i>	<i>im:7154887, si:dkey-250j7.4</i>	malignant fibrous histiocytoma amplified sequence 1	0.77	-0.33
miR-21	ENSDARG00000034808	<i>kcnip1b</i>	<i>zgc:101531</i>	Kv channel interacting protein 1 b	0.76	-0.33
miR-21	ENSDARG00000015863	<i>socs6b</i>	<i>socs6l, wu:fb77f06, zgc:64107</i>	suppressor of cytokine signaling 6b	0.75	-0.25
miR-21	ENSDARG00000078676	<i>myrf</i>	<i>c11orf9, zgc:172183</i>	myelin regulatory factor	0.75	-0.21
miR-21	ENSDARG000000094894	<i>tyrobp</i>	<i>dap12</i>	transmembrane immune signaling adaptor TYROBP	0.75	-0.37
miR-21	ENSDARG00000070826	<i>bpgm</i>	<i>zgc:92230</i>	2,3-bisphosphoglycerate mutase	0.75	-0.31
miR-21	ENSDARG00000060092	<i>zgc:165481</i>	N/A	<i>zgc:165481</i>	0.75	-0.36
miR-21	ENSDARG00000056572	<i>nrsn11</i>	<i>si:dkey-261i16.5, NRSN2</i>	neurensin 1-like	0.75	-0.26
miR-21	ENSDARG00000095949	<i>si:dkey-22i16.9</i>	N/A	<i>si:dkey-22i16.9</i>	0.75	-0.30
miR-21	ENSDARG00000044924	<i>gdf11</i>	<i>bmp11</i>	growth differentiation factor 11	0.74	-0.40
miR-21	ENSDARG00000088140	<i>hsd17b7</i>	<i>zgc:153977</i>	hydroxysteroid (17-beta) dehydrogenase 7	0.74	-0.33
miR-21	ENSDARG00000009273	<i>ppm1da</i>	<i>ppm1d, wu:fi44c08, zgc:55836, zgc:77316</i>	protein phosphatase, Mg2+/Mn2+ dependent, 1Da	0.73	-0.37
miR-21	ENSDARG00000077442	<i>tubgcp5</i>	<i>fa03b04, fe01d03, wu:fa03b04, wu:fe01d03, zgc:110217</i>	tubulin, gamma complex associated protein 5	0.73	-0.31
miR-21	ENSDARG00000019335	<i>hes6</i>	<i>her13.2</i>	hes family bHLH transcription factor 6	0.73	-0.31
miR-21	ENSDARG00000046006	<i>med20</i>	<i>trfp, zgc:101043</i>	mediator complex subunit 20	0.73	-0.36
miR-21	ENSDARG00000056530	<i>cpamd8</i>	<i>si:ch211-87h14.4, si:dkey-94n12.1</i>	C3 and PZP like alpha-2-macroglobulin domain containing 8	0.73	-0.40
miR-21	ENSDARG00000073962	<i>ccdc32</i>	<i>si:ch211-210h14.3</i>	coiled-coil domain containing 32	0.72	-0.29
miR-21	ENSDARG00000024829	<i>tnn</i>	<i>etID309873.2, tenw, tmw, zgc:110729</i>	tenascin N	0.72	-0.28
miR-21	ENSDARG00000052470	<i>igfbp2a</i>	<i>IGFBP-2, igfbp2, igfbp2b</i>	insulin-like growth factor binding protein 2a	0.72	-0.21
miR-21	ENSDARG00000074866	<i>ptpn5</i>	N/A	protein tyrosine phosphatase non-receptor type 5	0.72	-0.23
miR-21	ENSDARG00000041022	<i>pdc4b</i>	<i>cb45, pdc4, sb:cb45, wu:fc84b08</i>	programmed cell death 4b	0.71	-0.25
miR-21	ENSDARG00000016527	<i>helz2</i>	<i>gb:eb955855</i>	helicase with zinc finger 2, transcriptional coactivator	0.71	-0.30
miR-21	ENSDARG00000053961	<i>slc2a11a</i>	<i>slc2a11, zglut11a, CR293534.1</i>	solute carrier family 2 member 11a	0.71	-0.22
miR-21	ENSDARG00000019529	<i>parp1</i>	<i>si:dkey-206f10.3, wu:fc60f12, zgc:110092</i>	poly (ADP-ribose) polymerase 1	0.71	-0.36
miR-21	ENSDARG00000037837	<i>ogfr</i>	<i>fb25b04, fj34c06, ogfr2, si:dkeyp-44b8.5, wu:fb25b04, wu:fj34c06</i>	opioid growth factor receptor	0.71	-0.25
miR-21	ENSDARG00000087867	<i>SLC2A13</i>	N/A	proton myo-inositol cotransporter-like	0.70	-0.22
miR-21	ENSDARG00000060021	<i>vcpkmt</i>	<i>im:7159042, mettl21d, zgc:153596</i>	valosin containing protein lysine (K) methyltransferase	0.70	-0.25
miR-21	ENSDARG00000031888	<i>mapk8a</i>	<i>jnk1a-2, zgc:172209</i>	mitogen-activated protein kinase 8a	0.70	-0.41
miR-21	ENSDARG00000038011	<i>cyth4a</i>	<i>cyth4</i>	cytohesin 4a	0.70	-0.26
miR-21	ENSDARG00000022659	<i>snx4</i>	<i>im:6971267, wu:fb92d09, zgc:113437</i>	sorting nexin 4	0.70	-0.32
miR-122	ENSDARG00000024443	<i>gnpda2</i>	<i>si:dkey-185e18.2</i>	glucosamine-6-phosphate deaminase 2	0.98	-0.86
miR-122	ENSDARG00000021374	<i>cab39l1</i>	<i>cab39, wu:fb72b01, wu:fi37c08, zgc:55451</i>	calcium binding protein 39, like 1	0.96	-0.60
miR-122	ENSDARG00000062954	<i>clk2a</i>	<i>fb20c02, si:ch211-81a5.7, wu:fb20c02, clk2</i>	CDC-like kinase 2a	0.96	-0.45
miR-122	ENSDARG00000031954	<i>rbck1</i>	<i>zgc:91964</i>	RanBP-type and C3HC4-type zinc finger containing 1	0.95	-0.49
miR-122	ENSDARG00000014047	<i>cldn7b</i>	<i>cb388, cldn7, fd19f08, wu:fd19f08</i>	claudin 7b	0.95	-0.43
miR-122	ENSDARG00000077364	<i>mapk9</i>	<i>jnk2, si:ch211-247j9.2</i>	mitogen-activated protein kinase 9	0.94	-0.42

miR-122	ENSDARG00000030095	<i>ccdc78</i>	N/A	coiled-coil domain containing 78	0.93	-1.18
miR-122	ENSDARG00000020326	<i>tyk2</i>	N/A	tyrosine kinase 2	0.93	-0.26
miR-122	ENSDARG00000038981	<i>zgc:153615</i>	N/A	<i>zgc:153615</i>	0.93	-0.38
miR-122	ENSDARG00000074041	<i>abca5</i>	<i>zgc:163009</i>	ATP-binding cassette, sub-family A (ABC1), member 5	0.92	-0.46
miR-122	ENSDARG00000009281	<i>dnm1b</i>	<i>si:ch73-310g9.3</i>	dynamins 1b	0.91	-0.76
miR-122	ENSDARG00000078887	<i>atp10d</i>	<i>si:dkey-20i10.6</i>	ATPase phospholipid transporting 10D	0.91	-0.59
miR-122	ENSDARG00000069507	<i>mettl27</i>	<i>Williams Beuren syndrome chromosome region 27, im:7139431, wbscr27, wu:fe50c11, zgc:136279</i>	methyltransferase like 27	0.91	-0.61
miR-122	ENSDARG00000061231	<i>tinagl1</i>	<i>LCN7, lipocalin-7, si:dkey-158b13.1</i>	tubulointerstitial nephritis antigen-like 1	0.90	-0.40
miR-122	ENSDARG00000044420	<i>dnajc19</i>	<i>zgc:73251</i>	DnaJ (Hsp40) homolog, subfamily C, member 19	0.90	-1.00
miR-122	ENSDARG00000021681	<i>asrgl1</i>	<i>zgc:103568</i>	asparaginase and isoaspartyl peptidase 1	0.90	-0.43
miR-122	ENSDARG00000060917	<i>Anln</i>	<i>si:ch211-197n10.2, wu:fb74d05, wu:fd10d09, wu:fi41f04</i>	anillin, actin binding protein	0.89	-0.41
miR-122	ENSDARG00000003776	<i>pip4k2aa</i>	<i>pip4k2a, zgc:194746, zgc:194777</i>	phosphatidylinositol-5-phosphate 4-kinase, type II, alpha a	0.89	-0.47
miR-122	ENSDARG00000075530	<i>Pigu</i>	N/A	phosphatidylinositol glycan anchor biosynthesis, class U	0.89	-0.42
miR-122	ENSDARG000000062081	<i>tbc1d1</i>	N/A	TBC1 (tre-2/USP6, BUB2, cdc16) domain family, member 1	0.89	-0.72
miR-122	ENSDARG00000068264	<i>grhpra</i>	<i>grhpr, si:ch211-240l14.3, zgc:109900</i>	glyoxylate reductase/hydroxypyruvate reductase a	0.89	-0.44
miR-122	ENSDARG00000071082	<i>p4ha1b</i>	<i>p4ha1, wu:fc69d06, wu:fi27h11, zgc:56058</i>	prolyl 4-hydroxylase, alpha polypeptide I b	0.88	-0.79
miR-122	ENSDARG00000091320	<i>nlrc6</i>	<i>NACHT, LRR and PYD domains-containing protein 12e, Nacht 1.01, nlrp12e, zgc:136791, zgc:136791</i>	NLR family CARD domain containing 6	0.88	-0.40
miR-122	ENSDARG00000017542	<i>fgf1a</i>	<i>fgf1, zgc:73249</i>	fibroblast growth factor 1a	0.88	-0.25
miR-122	ENSDARG00000061207	<i>ttc7b</i>	<i>wu:fi46e11, zgc:153460</i>	tetratricopeptide repeat domain 7B	0.87	-0.34
miR-122	ENSDARG00000000442	<i>slc39a13</i>	<i>ZIP13, si:by184l24.1, si:xx-184l24.1, wu:fa55a07, zgc:136440</i>	solute carrier family 39 member 13	0.87	-0.73
miR-122	ENSDARG00000003520	<i>tbc1d14</i>	<i>sb:eu1004, si:dkey-120m5.1, wu:fi35h01, wu:fi42a03</i>	TBC1 domain family, member 14	0.87	-0.43
miR-122	ENSDARG00000045465	<i>lancl2</i>	<i>zgc:158293</i>	LanC lantibiotic synthetase component C-like 2	0.86	-0.30
miR-122	ENSDARG00000056144	<i>zgc:123244</i>	<i>TMEM150B</i>	<i>zgc:123244; orthologous to human TMEM150B (transmembrane protein 150B)</i>	0.86	-0.47
miR-122	ENSDARG00000061436	<i>col6a2</i>	N/A	collagen, type VI, alpha 2	0.86	-0.40
miR-122	ENSDARG00000069264	<i>si:ch211-196h16.5</i>	<i>Gene1133</i>	<i>si:ch211-196h16.5</i>	0.86	-0.40
miR-122	ENSDARG00000034409	<i>pik3r3b</i>	<i>pik3r3, zgc:55564, zgc:85710</i>	phosphoinositide-3-kinase, regulatory subunit 3b (gamma)	0.85	-0.38
miR-122	ENSDARG00000038237	<i>tbc1d12a</i>	<i>wu:fc59a02, tbc1d12</i>	TBC1 domain family, member 12a	0.85	-0.52
miR-122	ENSDARG00000089444	<i>omgb</i>	<i>ogmb, omg</i>	oligodendrocyte myelin glycoprotein b	0.85	-0.44
miR-122	ENSDARG00000014181	<i>foxp1b</i>	<i>foxp1, im:7139058, wu:fc83a06</i>	forkhead box P1b	0.84	-0.30
miR-122	ENSDARG00000037837	<i>ogfr</i>	<i>fb25b04, fj34c06, ogfr2, si:dkeyp-44b8.5, wu:fb25b04, wu:fj34c06</i>	opioid growth factor receptor	0.84	-0.82
miR-122	ENSDARG00000056985	<i>Tpte</i>	<i>Dr-VSP, tpip, wu:fd20e11, wu:fi24b06</i>	transmembrane phosphatase with tensin homology	0.84	-0.40
miR-122	ENSDARG00000014674	<i>acsl3b</i>	<i>acsl3, fa07a08, im:7155129, si:dkey-20f20.5, wu:fa07a08, wu:fb34c03</i>	acyl-CoA synthetase long chain family member 3b	0.84	-0.73
miR-122	ENSDARG00000043934	<i>il17a/f2</i>	N/A	interleukin 17a/f2	0.83	-0.40
miR-122	ENSDARG00000078026	<i>zcchc24</i>	<i>fc45d05, wu:fc45d05, zgc:158323</i>	zinc finger, CCHC domain containing 24	0.83	-0.33
miR-122	ENSDARG00000076889	<i>ralgapa1</i>	<i>garnl1, ulip1</i>	Ral GTPase activating protein, alpha subunit 1 (catalytic)	0.83	-0.37
miR-122	ENSDARG00000052856	<i>khdrbs1a</i>	<i>fa18g12, fa56c01, fc91b01, fj90g10, khdrbs1, p62, wu:fa18g12, wu:fa56c01, wu:fc91b01, wu:fj90g10</i>	KH domain containing, RNA binding, signal transduction associated 1a	0.82	-0.25
miR-122	ENSDARG00000071017	<i>nt5e</i>	<i>zgc:63784</i>	5'-nucleotidase, ecto (CD73)	0.82	-0.41
miR-122	ENSDARG00000055945	<i>asph</i>	<i>cb971, fb69e10, fc06d04, fc95a08, junctin, wu:fb69e10, wu:fc06d04, wu:fc95a08</i>	aspartate beta-hydroxylase	0.82	-0.43
miR-122	ENSDARG00000073959	<i>pmch</i>	<i>pmch2, si:ch211-260p3.3</i>	pro-melanin-concentrating hormone	0.82	-0.40
miR-122	ENSDARG00000022668	<i>grapb</i>	<i>si:dkeyp-9d4.1, zgc:109892</i>	GRB2 related adaptor protein b	0.82	-0.26
miR-122	ENSDARG00000034643	<i>fhl3a</i>	<i>fd57e01, fhl3, wu:fd57e01, fhl3</i>	four and a half LIM domains 3a	0.81	-0.40
miR-122	ENSDARG00000061400	<i>csgalnact2</i>	<i>mp:zf637-2-001473, si:dkey-192p21.4, wu:fb98d03, zf637-2-001473</i>	chondroitin sulfate N-acetylglactosaminyltransferase 2	0.80	-0.39
miR-122	ENSDARG00000055502	<i>cicb</i>	<i>cic, wu:fb93c01</i>	capicua transcriptional repressor b	0.79	-0.32
miR-122	ENSDARG00000070132	<i>astela</i>	<i>astel, si:dkey-11p23.5</i>	asteroid homolog 1a	0.79	-0.48
miR-122	ENSDARG00000034240	<i>capza1a</i>	<i>capza1, wu:fa19e05, wu:fk22d04, zgc:56314, zgc:85654</i>	capping actin protein of muscle Z-line subunit alpha 1a	0.79	-0.38

miR-122	ENSDARG00000055118	<i>mylipb</i>	<i>zgc:153767</i>	myosin regulatory light chain interacting protein b	0.79	-0.42
miR-122	ENSDARG00000051814	<i>ptprz1a</i>	<i>ca16c, im:7144789, ptpz1, zCA16c</i>	protein tyrosine phosphatase receptor type Z1a	0.78	-0.40
miR-122	ENSDARG00000056762	<i>tm4sf21b</i>	<i>zgc:172079</i>	transmembrane 4 L six family member 21b	0.78	-0.45
miR-122	ENSDARG00000017244	<i>chd6</i>	<i>si:dkey-6e2.3, zmp:0000000556</i>	chromodomain helicase DNA binding protein 6	0.78	-0.25
miR-122	ENSDARG00000056184	<i>dvl2</i>	<i>sb:eu471, wu:fc05d12, wu:fo71e09, wu:fp54a02, zgc:55372</i>	dishevelled segment polarity protein 2	0.78	-0.26
miR-122	ENSDARG00000078502	<i>si:ch211-150o23.3</i>	<i>SSC5D</i>	<i>si:ch211-150o23.3</i>	0.78	-0.28
miR-122	ENSDARG00000087759	<i>cnih4</i>	<i>fb51e06, fc10d11, wu:fb51e06, wu:fc10d11, wu:fi84d07</i>	cornichon family AMPA receptor auxiliary protein 4	0.77	-0.40
miR-122	ENSDARG00000069476	<i>spint2</i>	<i>fb52h04, fe24a06, wu:fb52h04, wu:fe24a06, zgc:153795</i>	serine peptidase inhibitor, Kunitz type, 2	0.77	-0.43
miR-122	ENSDARG00000044295	<i>pip5k1ba</i>	<i>pip5k1b, zgc:101046</i>	phosphatidylinositol-4-phosphate 5-kinase, type I, beta a	0.77	-0.38
miR-122	ENSDARG00000015314	<i>zc4h2</i>	<i>un-named hi2729b, unim hi2729b, wu:fb99f08, zgc:63849</i>	zinc finger, C4H2 domain containing	0.77	-0.43
miR-122	ENSDARG00000006074	<i>uck2a</i>	<i>umpk, wu:fk91a01</i>	uridine-cytidine kinase 2a	0.77	-0.42
miR-122	ENSDARG00000062986	<i>pnpla7a</i>	<i>clica, zgc:158748</i>	patatin-like phospholipase domain containing 7a	0.77	-0.55
miR-122	ENSDARG00000056239	<i>tmem45b</i>	<i>zgc:77892</i>	transmembrane protein 45B	0.76	-0.37
miR-122	ENSDARG00000043843	<i>akap7</i>	<i>si:ch211-272f15.3</i>	A kinase (PRKA) anchor protein 7	0.76	-0.33
miR-122	ENSDARG00000033184	<i>prkar2aa</i>	<i>wu:fb55f12, wu:ff29e12, wu:ff33e06, zgc:153742</i>	protein kinase, cAMP-dependent, regulatory, type II, alpha A	0.76	-0.27
miR-122	ENSDARG00000016260	<i>fxr2</i>	<i>id:ibd5046, zgc:56215, zgc:77472</i>	fragile X mental retardation, autosomal homolog 2	0.76	-0.55
miR-122	ENSDARG00000095580	<i>si:ch211-67e16.11</i>	<i>wu:fk68g02</i>	<i>si:ch211-67e16.11</i>	0.75	-0.46
miR-122	ENSDARG00000052624	<i>spryd7a</i>	<i>fd05e05, si:busm1-79f24.1, si:dkey-7a20.6, si:dz79f24.1, wu:fd05e05, zgc:56247</i>	SPRY domain containing 7a	0.75	-0.31
miR-122	ENSDARG00000074535	<i>lingo2a</i>	<i>sc:d0348, lingo2</i>	leucine rich repeat and lg domain containing 2a	0.75	-0.36
miR-122	ENSDARG00000040565	<i>ckmb</i>	<i>ckm3, wu:fb55e09, zgc:64204</i>	creatine kinase, muscle b	0.75	-0.37
miR-122	ENSDARG00000086645	<i>hs3st3b1b</i>	<i>3-ost-3z, z3-ost-3z</i>	heparan sulfate (glucosamine) 3-O-sulfotransferase 3B1b	0.75	-0.33
miR-122	ENSDARG00000038822	<i>mrc1b</i>	<i>mrc1, si:ch211-106h4.2</i>	mannose receptor, C type 1b	0.75	-0.28
miR-122	ENSDARG00000069324	<i>vps53</i>	<i>wu:fb77d11, zgc:103741</i>	VPS53 subunit of GARP complex	0.75	-0.43
miR-122	ENSDARG00000079723	<i>si:dkey-46g23.1</i>	<i>TNRC6B</i>	<i>si:dkey-46g23.1</i> ; orthologous to human TNRC6B (trinucleotide repeat containing adaptor 6B)	0.75	-0.27
miR-122	ENSDARG00000077946	<i>smarcc2</i>	<i>baf170</i>	SWI/SNF related, matrix associated, actin dependent regulator of chromatin, subfamily c, member 2	0.75	-0.32
miR-122	ENSDARG00000042970	<i>gng8</i>	<i>zgc:110316</i>	guanine nucleotide binding protein (G protein), gamma 8	0.74	-0.36
miR-122	ENSDARG00000016263	<i>zdhhc5a</i>	<i>si:dkey-28b4.5</i>	zinc finger DHHC-type containing 5a	0.74	-0.30
miR-122	ENSDARG00000025309	<i>dpf3</i>	<i>fi43h12, si:ch211-268e23.1, wu:fi43h12</i>	D4, zinc and double PHD fingers, family 3	0.73	-0.43
miR-122	ENSDARG00000054848	<i>pdk4</i>	<i>si:rp71-57j15.4</i>	pyruvate dehydrogenase kinase, isozyme 4	0.73	-0.30
miR-122	ENSDARG00000056481	<i>vat1</i>	<i>si:dz163124.2</i>	vesicle amine transport 1	0.73	-0.50
miR-122	ENSDARG00000056108	<i>ndufa4</i>	<i>zgc:73405</i>	NADH:ubiquinone oxidoreductase subunit A4	0.73	-0.42
miR-122	ENSDARG000000061473	<i>tbkbp1</i>	<i>zmp:0000000698</i>	TBK1 binding protein 1	0.72	-0.38
miR-122	ENSDARG00000077506	<i>tjp1a</i>	<i>cb817, tjp1, tjp1.1, zo-1</i>	tight junction protein 1a	0.72	-0.21
miR-122	ENSDARG00000070579	<i>ggact.3</i>	<i>zgc:162208, A2LD1</i>	gamma-glutamylamine cyclotransferase, tandem duplicate 3	0.72	-0.44
miR-122	ENSDARG00000001733	<i>gulf1a</i>	<i>gulf1, si:by35d8.1, si:ch211-198b21.4, si:xx-35d8.1, wuj61e02</i>	GULP PTB domain containing engulfment adaptor 1a	0.71	-0.25
miR-122	ENSDARG00000058114	<i>impad1</i>	<i>zgc:123256</i>	inositol monophosphatase domain containing 1	0.71	-0.29
miR-122	ENSDARG0000004939	<i>midhb</i>	N/A	metadherin b	0.71	-0.63
miR-122	ENSDARG00000036028	<i>arrdc3b</i>	<i>zgc:92034</i>	arrestin domain containing 3b	0.71	-0.41
miR-122	ENSDARG00000006151	<i>pde11a1</i>	<i>pde11a, wu:fi39g12</i>	phosphodiesterase 11a, like	0.71	-0.21
miR-122	ENSDARG00000045360	<i>flj110111</i>	<i>zgc:136503</i>	hypothetical protein FLJ11011-like (H. sapiens)	0.70	-0.28
miR-122	ENSDARG00000019644	<i>ldhba</i>	<i>ldhb, wu:fa16d07, wu:fa20c10, wu:fa96a08</i>	lactate dehydrogenase Ba	0.70	-0.27
miR-122	ENSDARG00000036107	<i>txnipa</i>	<i>sb:cb368, txnip</i>	thioredoxin interacting protein a	0.70	-0.34
miR-122	ENSDARG00000020890	<i>tmod4</i>	<i>Sk-Tmod, fa12b03, skeletal tropomodulin, sktmod, wu:fa12b03</i>	tropomodulin 4 (muscle)	0.70	-0.42
miR-122	ENSDARG00000037773	<i>rybpa</i>	<i>cb337, rybp, wu:fi34f03</i>	RING1 and YY1 binding protein a	0.70	-0.34
miR-731	ENSDARG00000019516	<i>sp7</i>	<i>osterix, osx</i>	Sp7 transcription factor	0.99	-0.50
miR-731	ENSDARG00000004735	<i>hnrnpub</i>	<i>hnrnpu, hnrpu, wu:fa66f11, wu:fc18b08, wu:fi89c08, zgc:111835, zgc:136480</i>	heterogeneous nuclear ribonucleoprotein Ub	0.98	-0.32
miR-731	ENSDARG00000010186	<i>myo3a</i>	N/A	myosin IIIA	0.97	-0.48

miR-731	ENSDARG00000009031	<i>tnikb</i>	<i>im:6898857, zgc:123033, zgc:152783</i>	TRAF2 and NCK interacting kinase b	0.97	-0.52
miR-731	ENSDARG00000027750	<i>dpp7</i>	<i>wu:fb19g02, zgc:112274, zgc:113564</i>	dipeptidyl-peptidase 7	0.96	-0.31
miR-731	ENSDARG00000002840	<i>si:dkey-28b4.8</i>	N/A	<i>si:dkey-28b4.8</i>	0.96	-0.26
miR-731	ENSDARG000000018923	<i>fat2</i>	<i>fk09f07, wu:fk09f07</i>	FAT atypical cadherin 2	0.96	-0.30
miR-731	ENSDARG000000032565	<i>cacng2a</i>	<i>zgc:64200</i>	calcium channel, voltage-dependent, gamma subunit 2a	0.94	-0.30
miR-731	ENSDARG000000090764	<i>bcas3</i>	<i>zgc:63764</i>	BCAS3 microtubule associated cell migration factor	0.94	-0.61
miR-731	ENSDARG000000016854	<i>neurod2</i>	<i>NDF2, fj49a11, ndr2, wu:fj49a11, zNdr2</i>	neuronal differentiation 2	0.94	-0.30
miR-731	ENSDARG000000063572	<i>Perp</i>	<i>fk24g11, wu:fk24g11</i>	p53 apoptosis effector related to pmp22	0.93	-0.33
miR-731	ENSDARG000000062055	<i>rnf38</i>	<i>fa94e11, si:dkey-20n3.1, wu:fa94e11</i>	ring finger protein 38	0.92	-0.24
miR-731	ENSDARG000000044980	<i>kpna6</i>	<i>zmp:0000000623</i>	karyopherin alpha 6 (importin alpha 7)	0.91	-0.40
miR-731	ENSDARG000000043584	<i>ifit15</i>	<i>lfit17b, ifit2, wu:fb58d07</i>	interferon-induced protein with tetratricopeptide repeats 15	0.90	-0.21
miR-731	ENSDARG000000070337	<i>hoxc1a</i>	<i>z-42</i>	homeobox C1a	0.90	-0.29
miR-731	ENSDARG000000019752	<i>rom1a</i>	<i>zgc:73336, zgc:77401</i>	retinal outer segment membrane protein 1a	0.90	-0.38
miR-731	ENSDARG000000024092	<i>lmbr1</i>	<i>im:7156444</i>	limb development membrane protein 1	0.90	-0.37
miR-731	ENSDARG000000057586	<i>sall3b</i>	<i>im:7138196, sall3l, sallb, si:dkey-195c14.1, wu:fb01e09, sall3</i>	spalt-like transcription factor 3b	0.90	-0.36
miR-731	ENSDARG000000042526	<i>sebox</i>	<i>OG9 homeobox gene, mezzo, og9x</i>	SEBOX homeobox	0.89	-0.38
miR-731	ENSDARG000000059707	<i>znf423</i>	<i>zgc:158272, si:ch211-216123.1</i>	zinc finger protein 423	0.88	-0.23
miR-731	ENSDARG000000033957	<i>trmt44</i>	<i>mettl19, zgc:101657</i>	tRNA methyltransferase 44 homolog	0.88	-0.26
miR-731	ENSDARG0000000068369	<i>angptl2b</i>	<i>angptl2, wu:fc41g02</i>	angiopoietin-like 2b	0.88	-0.43
miR-731	ENSDARG000000029252	<i>Ssb</i>	<i>wu:fb17f11</i>	small RNA binding exonuclease protection factor La	0.88	-0.24
miR-731	ENSDARG000000043283	<i>prrg1</i>	<i>zgc:103657</i>	proline rich Gla (G-carboxyglutamic acid) 1	0.87	-0.39
miR-731	ENSDARG000000023999	<i>heatr5a</i>	<i>si:busm1-142b24.3, si:dz142b24.3, zgc:63665</i>	HEAT repeat containing 5a	0.87	-0.43
miR-731	ENSDARG000000043003	<i>pcyt2</i>	<i>im:7158585, wu:fb39h11, zgc:103434</i>	phosphate cytidyltransferase 2, ethanolamine	0.87	-0.53
miR-731	ENSDARG000000069922	<i>pla1a</i>	<i>wu:fj26h09, zgc:77160, popdc2</i>	popeye domain containing 2	0.87	-0.28
miR-731	ENSDARG0000000019326	<i>prpsap2</i>	<i>wu:fj14g04, zgc:55477, zgc:85610</i>	phosphoribosyl pyrophosphate synthetase-associated protein 2	0.86	-0.33
miR-731	ENSDARG000000010248	<i>wdr54</i>	<i>zgc:100930</i>	WD repeat domain 54	0.86	-0.23
miR-731	ENSDARG000000016141	<i>slc6a2</i>	<i>si:dkey-182h7.3</i>	solute carrier family 6 member 2	0.86	-0.49
miR-731	ENSDARG000000062837	<i>spata13</i>	<i>si:dkey-33c18.3</i>	spermatogenesis associated 13	0.86	-0.24
miR-731	ENSDARG000000074635	<i>abca1a</i>	<i>abca1, cb939, fa03d03, im:6795541, wu:fa03d03, wu:fj13h01</i>	ATP-binding cassette, sub-family A (ABC1), member 1A	0.86	-0.27
miR-731	ENSDARG000000057869	<i>cdc42l2</i>	<i>cdc42b, zgc:136923</i>	cell division cycle 42 like 2	0.85	-0.70
miR-731	ENSDARG000000070792	<i>lrrc15</i>	<i>cb894, zgc:158286</i>	leucine rich repeat containing 15	0.85	-0.45
miR-731	ENSDARG000000054683	<i>prdm8b</i>	<i>si:ch211-57h10.4</i>	PR domain containing 8b	0.85	-0.41
miR-731	ENSDARG000000030608	<i>rhogd</i>	<i>rhoga, zgc:66008</i>	ras homolog gene family, member Gd	0.85	-0.26
miR-731	ENSDARG000000015495	<i>klf3</i>	<i>klf12</i>	Kruppel-like factor 3 (basic)	0.85	-0.26
miR-731	ENSDARG000000023323	<i>ywhaqb</i>	<i>wu:fa91b01, wu:fb02e12, wu:fb36f06, wu:fk86h02, ywhaq1, ywhaz, zgc:55499</i>	tyrosine 3-monooxygenase/tryptophan 5-monooxygenase activation protein, theta polypeptide b	0.85	-0.46
miR-731	ENSDARG000000028173	<i>slc4a2a</i>	<i>ae2, ae2.1, slc4a2</i>	solute carrier family 4 member 2a	0.84	-0.27
miR-731	ENSDARG000000051916	<i>cdk7</i>	<i>zgc:85821</i>	cyclin-dependent kinase 7	0.84	-0.28
miR-731	ENSDARG000000044492	<i>ublcpl</i>	<i>fb33g09, wu:fb33g09, zgc:86634</i>	ubiquitin-like domain containing CTD phosphatase 1	0.84	-0.24
miR-731	ENSDARG000000035837	<i>mosmob</i>	<i>zgc:153595, C12H16orf52</i>	modulator of smoothened b	0.84	-0.47
miR-731	ENSDARG000000035660	<i>kiaa0319l</i>	<i>si:ch211-193k19.1</i>	KIAA0319 like; orthologous to human <i>KIAA0319L</i>	0.83	-0.29
miR-731	ENSDARG000000040253	<i>onecutl</i>	<i>id:ibd5005, ocl, onecut, onecut l, onecut1</i>	one cut domain, family member, like	0.83	-0.62
miR-731	ENSDARG000000035634	<i>rfc5</i>	<i>zgc:110313</i>	replication factor C (activator 1) 5	0.83	-0.32
miR-731	ENSDARG000000086173	<i>relb</i>	<i>fj85b02, wu:fj85b02</i>	bv-rel avian reticuloendotheliosis viral oncogene homolog B	0.83	-0.29
miR-731	ENSDARG0000000004160	<i>reep3b</i>	<i>im:6909180, reep3, sb:eu299, zgc:55529</i>	receptor accessory protein 3b	0.83	-0.24
miR-731	ENSDARG000000056292	<i>vsx1</i>	<i>etlD309864.17</i>	visual system homeobox 1 homolog, chx10-like	0.83	-0.53
miR-731	ENSDARG000000077721	<i>knop1</i>	<i>si:ch211-202h22.6, C1H16orf88</i>	lysine-rich nucleolar protein 1	0.83	-0.23
miR-731	ENSDARG000000043019	<i>exoc1</i>	<i>sec31l, wu:fb58e03, zgc:64145</i>	exocyst complex component 1	0.82	-0.24
miR-731	ENSDARG000000056347	<i>rab3aa</i>	<i>rab3a, zgc:92276</i>	RAB3A, member RAS oncogene family, a	0.82	-0.27
miR-731	ENSDARG000000016743	<i>angel2</i>	<i>f178e02, si:ch211-181h6.2, wu:f178e02, zgc:153829</i>	angel homolog 2 (Drosophila)	0.82	-0.27
miR-731	ENSDARG000000032496	<i>pon1</i>	<i>zgc:91887</i>	paraoxonase 1	0.82	-0.25
miR-731	ENSDARG000000008788	<i>camk1gb</i>	<i>fk61c03, wu:fk61c03, zgc:73155</i>	calcium/calmodulin-dependent protein kinase Igb	0.82	-0.29

miR-731	ENSDARG00000010454	<i>gucal1a</i>	<i>gcapl</i>	guanylate cyclase activator 1A	0.81	-0.35
miR-731	ENSDARG000000001676	<i>gpm6bb</i>	<i>DMgamma2, gpb6bb, zgc:110793</i>	glycoprotein M6Bb	0.81	-0.41
miR-731	ENSDARG000000059575	<i>uhmk1</i>	<i>zgc:153241</i>	U2AF homology motif (UHM) kinase 1	0.81	-0.28
miR-731	ENSDARG000000079740	<i>efna2b</i>	<i>si:dkey-174e10.1, EFNA2</i>	ephriin-A2b	0.81	-0.25
miR-731	ENSDARG000000013730	<i>slc4a4a</i>	<i>NBC1, NBCE1, id:ibd2520, si:dkey-256m11.1, slc4a4</i>	solute carrier family 4 member 4a	0.81	-0.59
miR-731	ENSDARG000000010791	<i>dla</i>	<i>cb335, delta A, fa04c10, wu:fa04c10</i>	deltaA	0.80	-0.26
miR-731	ENSDARG000000022372	<i>kng1</i>	<i>fb64g01, wu:fb64g01, zgc:103569</i>	kininogen 1	0.80	-0.27
miR-731	ENSDARG000000079345	<i>si:ch211-217k17.10</i>	<i>fk24c04, wu:fk24c04</i>	<i>si:ch211-217k17.10</i>	0.80	-0.57
miR-731	ENSDARG000000030871	<i>siah1</i>	<i>id:ibd2635, zgc:56026</i>	siah E3 ubiquitin protein ligase 1	0.80	-0.37
miR-731	ENSDARG000000043334	<i>ccdc6a</i>	<i>ccdc6, fj36e04, wu:fj36e04, zgc:77435</i>	coiled-coil domain containing 6a	0.80	-0.59
miR-731	ENSDARG000000056150	<i>rbms2b</i>	<i>zgc:100836</i>	RNA binding motif, single stranded interacting protein 2b	0.80	-0.22
miR-731	ENSDARG000000020607	<i>emc3</i>	<i>loc55831, pob, wu:fi32f05, wu:fj63h08, zgc:63727</i>	ER membrane protein complex subunit 3	0.79	-0.76
miR-731	ENSDARG000000070955	<i>hmx3a</i>	<i>Nkx5-1, hmx3, nkx5.1, zgc:109845</i>	H6 family homeobox 3a	0.79	-0.22
miR-731	ENSDARG000000035577	<i>cds2</i>	<i>zgc:66134</i>	CDP-diacylglycerol synthase (phosphatidate cytidyltransferase) 2	0.79	-0.27
miR-731	ENSDARG000000076710	<i>si:dkey-42123.7</i>	<i>zmp:0000000806, CR848836.6</i>	<i>si:dkey-42123.7</i> ; orthologous to human <i>GPR33</i> (G protein-coupled receptor 33)	0.79	-0.21
miR-731	ENSDARG000000009273	<i>ppm1da</i>	<i>ppm1d, wu:fj44c08, zgc:55836, zgc:77316</i>	protein phosphatase, Mg2+/Mn2+ dependent, 1Da	0.78	-0.50
miR-731	ENSDARG000000043608	<i>elf4ebp1</i>	<i>fc04h09, wu:fc04h09, zgc:64137</i>	eukaryotic translation initiation factor 4E binding protein 1	0.78	-0.23
miR-731	ENSDARG0000000091433	<i>abhd17aa</i>	<i>abhd17ca, fb50g01, wu:fb50g01, zgc:162293, FAM108A1</i>	abhydrolase domain containing 17Aa	0.78	-0.24
miR-731	ENSDARG000000044267	<i>sumo1</i>	<i>fd12c02, hm:zeh0670, wu:fb74c02, wu:fd12c02, zgc:65934, zgc:85634</i>	small ubiquitin like modifier 1	0.78	-0.52
miR-731	ENSDARG000000094377	<i>samd10b</i>	<i>samd10, si:ch211-278f9.1, samd10</i>	sterile alpha motif domain containing 10b	0.78	-0.24
miR-731	ENSDARG000000042961	<i>zgc:100920</i>	N/A	<i>zgc:100920</i> ; orthologous to human <i>NKAIN4</i> (sodium/potassium transporting ATPase interacting 4).	0.78	-0.40
miR-731	ENSDARG0000000053110	<i>pkib</i>	<i>im:7054531, si:dkey-150h13.1</i>	protein kinase (cAMP-dependent, catalytic) inhibitor beta	0.78	-0.27
miR-731	ENSDARG000000017740	<i>sec63</i>	<i>zgc:92718</i>	SEC63 homolog, protein translocation regulator	0.78	-0.39
miR-731	ENSDARG000000078095	<i>cipcb</i>	<i>si:ch73-387e10.3</i>	CLOCK-interacting pacemaker b	0.78	-0.22
miR-731	ENSDARG000000011611	<i>actr1</i>	<i>hm:zehn1110, wu:fc22g09, wu:fc56d01, wu:fj52e08, zehnl110, zgc:56317</i>	actin related protein 1, centractin	0.78	-0.21
miR-731	ENSDARG0000000053665	<i>gabrg2</i>	<i>si:ch211-145n14.1</i>	gamma-aminobutyric acid (GABA) A receptor, gamma 2	0.78	-0.29
miR-731	ENSDARG0000000058256	<i>draxin</i>	<i>neucrin, zgc:113312, C11H1orf187</i>	dorsal inhibitory axon guidance protein	0.78	-0.26
miR-731	ENSDARG000000070918	<i>si:ch211-284e20.8</i>	<i>FETUB</i>	<i>si:ch211-284e20.8</i> ; orthologous to human <i>FETUB</i> (fetuin B)	0.77	-0.22
miR-731	ENSDARG0000000020031	<i>cldn11a</i>	<i>zgc:92247</i>	claudin 11a	0.77	-0.22
miR-731	ENSDARG000000018621	<i>slc6a19a.1</i>	<i>slc6a19a, zgc:162095</i>	solute carrier family 6 member 19a, tandem duplicate 1	0.77	-0.25
miR-731	ENSDARG000000015860	<i>mtmr9</i>	<i>si:dkey-155d18.2</i>	myotubularin related protein 9	0.77	-0.26
miR-731	ENSDARG0000000058082	<i>birc7</i>	<i>zgc:165605</i>	baculoviral IAP repeat containing 7	0.77	-0.54
miR-731	ENSDARG0000000092501	<i>si:dkey-80c24.5</i>	N/A	<i>si:dkey-80c24.5</i>	0.77	-0.32
miR-731	ENSDARG000000012248	<i>rgma</i>	<i>id:ibd2030, wu:fb72h09, zgc:110534</i>	repulsive guidance molecule BMP co-receptor a	0.77	-0.49
miR-731	ENSDARG000000017886	<i>ZBTB11</i>	<i>man, marsanne, mne, si:dkey-98p3.1, si:dkeyp-116g9.2, wu:fc06h02, wu:fc36e08</i>	zinc finger and BTB domain containing 11	0.77	-0.44
miR-731	ENSDARG0000000029290	<i>stx11b.1</i>	<i>zgc:77855</i>	syntaxin 11b, tandem duplicate 1	0.76	-0.25
miR-731	ENSDARG0000000020134	<i>sipa1ll</i>	<i>cb600, id:ibd5092, sb:cb600, si:ch211-106p8.1, wu:fc43g06</i>	signal-induced proliferation-associated 1 like 1	0.76	-0.21
miR-731	ENSDARG0000000058535	<i>si:dkey-211g8.7</i>	<i>FFAR2</i>	<i>si:dkey-211g8.7</i> ; orthologous to human <i>FFAR2</i> (free fatty acid receptor 2)	0.76	-0.26
miR-731	ENSDARG0000000061738	<i>march2</i>	<i>si:ch211-197g15.3, zgc:158704</i>	membrane-associated ring finger (C3HC4) 2	0.76	-0.22
miR-731	ENSDARG0000000074363	<i>TTC9</i>	N/A	<i>si:ch211-259k16.3</i>	0.76	-0.31
miR-731	ENSDARG0000000021539	<i>ephb2b</i>	<i>fi18g12, wu:fi18g12, zgc:112296</i>	eph receptor B2b	0.76	-0.25
miR-731	ENSDARG0000000028379	<i>trub2</i>	<i>fc51e05, wu:fc51e05, zgc:91836</i>	TruB pseudouridine (psi) synthase family member 2	0.76	-0.26
miR-731	ENSDARG000000004954	<i>grna</i>	<i>grn, pgrn-a, progranulin, sb:eu337, wu:fb11g12, wu:fi18f05, zfPGRN-A</i>	granulin a	0.76	-0.28
miR-731	ENSDARG0000000039008	<i>SERP2</i>	<i>fc56c08, wu:fc56c08, zgc:85858</i>	<i>zgc:85858</i>	0.75	-0.24
miR-731	ENSDARG000000008904	<i>smarca2</i>	<i>wu:fa56c07, wu:fi27f11, zgc:66238</i>	SWI/SNF related, matrix associated, actin dependent regulator of chromatin, subfamily a, member 2	0.75	-0.32

miR-731	ENSDARG00000090889	<i>si:ch211-132p1.3</i>	<i>zmp:0000001007, BX005069.4, lect2</i>	<i>si:ch211-132p1.3</i> ; orthologous to human <i>LECT2</i> (leukocyte cell derived chemotaxin 2)	0.75	-0.25
miR-731	ENSDARG00000063544	<i>pip4k2ab</i>	<i>si:ch211-279l9.5</i>	phosphatidylinositol-5-phosphate 4-kinase, type II, alpha b	0.75	-0.33
miR-731	ENSDARG00000078492	<i>swi5</i>	<i>si:ch211-264f7.1</i>	SWI5 homologous recombination repair protein	0.75	-0.37
miR-731	ENSDARG00000074809	<i>zgc:194392</i>	<i>fd59g01, wu:fd59g01</i>	<i>zgc:194392</i> ; orthologous to human <i>C1orf115</i> (chromosome 1 open reading frame 115)	0.75	-0.26
miR-731	ENSDARG00000031809	<i>rbm24b</i>	<i>zgc:136803</i>	RNA binding motif protein 24b	0.75	-0.36
miR-731	ENSDARG00000037177	<i>zc3h13</i>	<i>zgc:66359</i>	zinc finger CCCH-type containing 13	0.74	-0.26
miR-731	ENSDARG00000059474	<i>mlxip</i>	<i>MLx, chrebp, si:ch211-110p13.4, wu:fb38e04</i>	MLX interacting protein	0.74	-0.37
miR-731	ENSDARG00000011885	<i>mrpl19</i>	<i>wu:fi69e01, zgc:100827</i>	mitochondrial ribosomal protein L19	0.74	-0.42
miR-731	ENSDARG00000027183	<i>namptb</i>	<i>im:7051384, wu:fc21g12, wu:fc27h03, wu:fd15d03, zgc:55764</i>	nicotinamide phosphoribosyltransferase b	0.74	-0.22
miR-731	ENSDARG00000074262	<i>nck1a</i>	<i>si:dkey-4m11.4, nck1</i>	NCK adaptor protein 1a	0.74	-0.23
miR-731	ENSDARG00000038737	<i>phf20b</i>	<i>zgc:91986</i>	PHD finger protein 20, b	0.74	-0.25
miR-731	ENSDARG00000094491	<i>si:ch211-202m22.1</i>	N/A	<i>si:ch211-202m22.1</i>	0.74	-0.31
miR-731	ENSDARG00000074367	<i>usp12b</i>	<i>CR381676.2</i>	ubiquitin specific peptidase 12b	0.74	-0.25
miR-731	ENSDARG00000007125	<i>asb6</i>	<i>zgc:110231</i>	ankyrin repeat and SOCS box containing 6	0.74	-0.36
miR-731	ENSDARG00000018643	<i>igf2a</i>	<i>IGF-II, etID31232.25, igf2, igf2b, wu:fc26e03, zgc:91781</i>	insulin-like growth factor 2a	0.74	-0.31
miR-731	ENSDARG00000028552	<i>plppr3b</i>	<i>PRG2A, lppr3, lppr3b, BX000430.1</i>	phospholipid phosphatase related 3b	0.74	-0.32
miR-731	ENSDARG00000074597	<i>lts2a</i>	<i>lts2, si:dkey-250a11.1</i>	leucine zipper, putative tumor suppressor 2a	0.74	-0.39
miR-731	ENSDARG00000054344	<i>gabpb2b</i>	<i>im:7142141, si:dkey-77n11.3, zgc:154003</i>	GA binding protein transcription factor subunit beta 2b	0.73	-0.40
miR-731	ENSDARG00000073813	<i>sclt</i>	<i>zgc:171514</i>	selenocysteine lyase	0.73	-0.40
miR-731	ENSDARG00000079784	<i>si:ch211-235o23.1</i>	<i>TMEM71</i>	<i>si:ch211-235o23.1</i> ; orthologous to human <i>TMEM71</i> (transmembrane protein 71)	0.73	-0.29
miR-731	ENSDARG000000008979	<i>golga1</i>	<i>zgc:63688</i>	golgin A1	0.73	-0.25
miR-731	ENSDARG00000035398	<i>enc1</i>	<i>klhl37, si:dkey-35i22.1</i>	ectodermal-neural cortex 1	0.73	-0.22
miR-731	ENSDARG00000055305	<i>ret</i>	<i>c-ret, cret, etID315074.13, ret1, wu:fd13h01</i>	ret proto-oncogene receptor tyrosine kinase	0.72	-0.36
miR-731	ENSDARG00000043640	<i>cenpn</i>	<i>id:ibd2033, si:busm1-241h12.3, si:dz241h12.3, un-named hi3634, unmi hi3634, zgc:92188</i>	centromere protein N	0.72	-0.28
miR-731	ENSDARG00000059244	<i>arfp2a</i>	<i>fj35g06, wu:fj35g06, zgc:92579</i>	ADP-ribosylation factor interacting protein 2a	0.72	-0.24
miR-731	ENSDARG00000061992	<i>dot1l</i>	<i>si:dkey-211f22.1</i>	DOT1-like histone H3K79 methyltransferase	0.72	-0.21
miR-731	ENSDARG00000069833	<i>rhbd12</i>	<i>zgc:55533</i>	rhomboid, veinlet-like 2 (Drosophila)	0.72	-0.31
miR-731	ENSDARG00000011553	<i>atp5pb</i>	<i>atp5f1, fb59g11, fj08b08, wu:fb59g11, wu:fj08b08, zgc:101887</i>	ATP synthase peripheral stalk-membrane subunit b	0.71	-0.42
miR-731	ENSDARG00000006010	<i>bmi1a</i>	<i>bmi1, pcg4, psc1, wu:fb17g03, wu:fd18f06</i>	bmi1 polycomb ring finger oncogene 1a	0.71	-0.22
miR-731	ENSDARG00000077265	<i>wdr89</i>	N/A	WD repeat domain 89	0.71	-0.23
miR-731	ENSDARG00000007216	<i>abce1</i>	<i>wu:fb34c09, wu:fe47b01, wu:fi09g07, zgc:111906, zgc:56045</i>	ATP-binding cassette, sub-family E (OABP), member 1	0.71	-0.34
miR-731	ENSDARG00000070567	<i>cadpsb</i>	N/A	Ca ²⁺ -dependent activator protein for secretion b	0.71	-0.31
miR-731	ENSDARG00000044827	<i>wnt7aa</i>	<i>wnt7a</i>	wingless-type MMTV integration site family, member 7Aa	0.71	-0.54
miR-731	ENSDARG000000076170	<i>pcsk1nl</i>	<i>proSAAS, uo:ion003, wu:fj32g06</i>	proprotein convertase subtilisin/kexin type 1 inhibitor, like	0.70	-0.31
miR-731	ENSDARG00000077096	<i>fncl7a</i>	<i>fncl7, si:dkeyp-51f11.6</i>	fibronectin type III domain containing 7a	0.70	-0.24
miR-731	ENSDARG00000079414	<i>sez6b</i>	N/A	seizure related 6 homolog b	0.70	-0.21
miR-731	ENSDARG00000078576	<i>setd5</i>	N/A	SET domain containing 5	0.70	-0.38
miR-731	ENSDARG00000091817	<i>CR391998.1</i>	N/A	putative protein TPRXL	0.70	-0.29
miR-731	ENSDARG00000053857	<i>ccdc187</i>	<i>zgc:162431</i>	coiled-coil domain containing 187	0.70	-0.24
miR-731	ENSDARG00000042115	<i>tmem198a</i>	<i>si:rp71-68g1.6</i>	transmembrane protein 198a	0.70	-0.31
miR-731	ENSDARG00000020278	<i>si:dkeyp-120h9.1</i>	N/A	<i>si:dkeyp-120h9.1</i> ; orthologous to human <i>SLC7A13</i> (solute carrier family 7 member 13)	0.70	-0.72
miR-26	ENSDARG00000059801	<i>pdzd2</i>	<i>si:dkey-3h3.1</i>	PDZ domain containing 2	0.96/0.97*	-0.58
miR-26	ENSDARG00000087176	<i>rskr</i>	<i>rskr, sgk494a, si:dkeyp-118e10.5, AL954372.2</i>	ribosomal protein S6 kinase related a	0.96/0.96	-0.51
miR-26	ENSDARG00000025766	<i>zgc:66440</i>	<i>fj25g12, wu:fj25g12, DTX3</i>	<i>zgc:66440</i>	0.72/-	-0.48
miR-26	ENSDARG00000055108	<i>gdel</i>	<i>zgc:56068, zgc:77135</i>	glycerophosphodiester phosphodiesterase 1	0.94/0.94	-0.47
miR-26	ENSDARG00000009901	<i>slc38a5a</i>	<i>fk81c02, si:dkey-48j7.5, wu:fk81c02</i>	solute carrier family 38 member 5a	0.93/0.91	-0.46

miR-26	ENSDARG00000040684	<i>plcd4b</i>	<i>DrPLC-delta 4B, plcd4</i>	phospholipase C, delta 4b	0.83/0.84	-0.46
miR-26	ENSDARG00000074301	<i>cth</i>	<i>fb58d08, wu:fb48e03, wu:fb58d08, zgc:66001, zgc:85785</i>	cystathionase (cystathionine gamma-lyase)	0.91/0.90	-0.44
miR-26	ENSDARG00000052438	<i>actr2a</i>	<i>actr2, hm:zeh1257, zgc:63719</i>	actin related protein 2a	0.78/0.80	-0.43
miR-26	ENSDARG00000078585	<i>mon1a</i>	<i>zgc:162873</i>	MON1 secretory trafficking family member A	0.91/0.90	-0.43
miR-26	ENSDARG00000044183	<i>prkab1a</i>	<i>zgc:92228</i>	protein kinase, AMP-activated, beta 1 non-catalytic subunit, a	0.73/0.71	-0.43
miR-26	ENSDARG00000001898	<i>manea</i>	<i>fi29h09, wu:fi29h09, zgc:92825</i>	mannosidase, endo-alpha	0.90/0.91	-0.42
miR-26	ENSDARG00000043026	<i>ttbk2b</i>	<i>ff62e10, sb:eu1102, si:dkey-12h9.11, si:dkey-12h9.14, wu:ff62e10</i>	tau tubulin kinase 2b	0.81/0.81	-0.41
miR-26	ENSDARG00000028027	<i>trim63a</i>	<i>MuRF1A, fc50c07, murf1, trim63, wu:fc50c07, zgc:86757</i>	tripartite motif containing 63a	0.91/0.93	-0.40
miR-26	ENSDARG00000036066	<i>si:dkey-78a14.4</i>	N/A	<i>si:dkey-78a14.4</i> ; orthologous to several human genes including <i>NAT1</i> (N-acetyltransferase 1)	0.70/-	-0.40
miR-26	ENSDARG00000006220	<i>ugt1a1</i>	<i>ugt1ab</i>	UDP glucuronosyltransferase 1 family, polypeptide A1	0.93/0.92	-0.39
miR-26	ENSDARG00000052386	<i>cldnd1b</i>	<i>cldnd1</i>	claudin domain containing 1b	0.82/0.81	-0.38
miR-26	ENSDARG00000017427	<i>phf20l1</i>	<i>fi20g04, si:dkey-97i18.2, wu:fi20g04</i>	PHD finger protein 20 like 1	0.95/0.95	-0.38
miR-26	ENSDARG00000003259	<i>lox</i>	<i>lox, zgc:77447</i>	lysyl oxidase a	0.94/0.94	-0.37
miR-26	ENSDARG00000039966	<i>prom1a</i>	<i>fb75c01, prom1, prom1l, wu:fb75c01</i>	prominin 1a	0.88/0.87	-0.37
miR-26	ENSDARG00000015184	<i>mpp3a</i>	<i>mpp3, si:ch73-368i2.1, zmp:0000000552</i>	membrane protein, palmitoylated 3a (MAGUK p55 subfamily member 3)	0.86/0.86	-0.37
miR-26	ENSDARG00000088377	<i>rp111b</i>	<i>si:dkey-33i22.3</i>	<i>rp1 like 1b</i> ; orthologous to human <i>RP111</i>	0.88/0.88	-0.36
miR-26	ENSDARG00000002696	<i>gnb3b</i>	<i>gnb3, wu:fk54b04, zgc:73058, zgc:77780</i>	guanine nucleotide binding protein (G protein), beta polypeptide 3b	0.86/0.86	-0.36
miR-26	ENSDARG00000038296	<i>tmem86b</i>	<i>zgc:153124</i>	transmembrane protein 86B	0.82/0.81	-0.36
miR-26	ENSDARG00000091548	<i>stard9</i>	<i>BX569781.2</i>	StAR-related lipid transfer (START) domain containing 9	0.91/0.90	-0.36
miR-26	ENSDARG00000045586	<i>zgc:172145</i>	N/A	<i>zgc:172145</i> ; orthologous to human <i>FTL</i> (ferritin light chain)	0.96/0.97	-0.36
miR-26	ENSDARG00000058486	<i>caps2</i>	<i>si:ch73-86n2.3</i>	calcyphosine 2	0.78/0.78	-0.36
miR-26	ENSDARG00000007485	<i>epc2</i>	<i>zgc:55784</i>	enhancer of polycomb homolog 2 (Drosophila)	0.99/0.99	-0.35
miR-26	ENSDARG00000028088	<i>galk1</i>	<i>zgc:101541</i>	galactokinase 1	0.79/0.81	-0.35
miR-26	ENSDARG00000016584	<i>rgs7a</i>	<i>rgs7, zgc:92793</i>	regulator of G protein signaling 7a	0.83/0.85	-0.35
miR-26	ENSDARG00000016141	<i>slc6a2</i>	<i>si:dkey-182h7.3</i>	solute carrier family 6 member 2	0.76/0.76	-0.35
miR-26	ENSDARG000000021255	<i>arhgap22</i>	<i>si:ch211-223a10.2</i>	Rho GTPase activating protein 22	-/0.71	-0.35
miR-26	ENSDARG00000076844	<i>plin6</i>	<i>zgc:162150</i>	perilipin 6	0.72/0.72	-0.34
miR-26	ENSDARG00000005236	<i>srcap</i>	<i>im:7137889, im:7138264</i>	Snf2-related CREBBP activator protein	0.90/0.90	-0.34
miR-26	ENSDARG00000058978	<i>zgc:113423</i>	<i>BEND2</i>	<i>zgc:113423</i> ; orthologous to human <i>BEND2</i> (BEN domain containing 2)	0.98/0.96	-0.34
miR-26	ENSDARG00000069045	<i>cht8</i>	<i>si:ch211-218c6.5, zgc:158641</i>	CTF8, chromosome transmission fidelity factor 8 homolog (S. cerevisiae)	0.78/0.79	-0.34
miR-26	ENSDARG00000063414	<i>rxylt1</i>	<i>tmem5, zgc:153239</i>	ribitol xylosyltransferase 1	0.91/0.93	-0.34
miR-26	ENSDARG00000079500	<i>kif3cb</i>	<i>kif3c, kif3c-like</i>	kinesin family member 3Cb	0.80/0.79	-0.34
miR-26	ENSDARG00000022550	<i>gxylt1b</i>	<i>gl18d3, gxylt1, si:ch211-155a11.6</i>	glucoside xylosyltransferase 1b	0.91/0.90	-0.34
miR-26	ENSDARG00000016022	<i>zic5</i>	N/A	zic family member 5 (odd-paired homolog, Drosophila)	0.99/0.99	-0.34
miR-26	ENSDARG00000060354	<i>samd7</i>	<i>si:dkey-52k20.12</i>	sterile alpha motif domain containing 7	0.87/0.87	-0.33
miR-26	ENSDARG00000060921	<i>pla2g6</i>	<i>wu:fc06g09, zgc:77476</i>	phospholipase A2, group VI (cytosolic, calcium-independent)	0.84/0.82	-0.33
miR-26	ENSDARG00000003167	<i>mlst8</i>	<i>fi37e04, gbl, wu:fi37e04, zgc:55455, zgc:85668</i>	MTOR associated protein, LST8 homolog (S. cerevisiae)	0.94/0.94	-0.33
miR-26	ENSDARG00000075612	<i>ercc6</i>	N/A	excision repair cross-complementation group 6	0.85/0.86	-0.33
miR-26	ENSDARG00000044513	<i>tpst1l</i>	N/A	tyrosylprotein sulfotransferase 1, like	0.77/0.75	-0.33
miR-26	ENSDARG00000058608	<i>wsb2</i>	<i>si:ch73-263f13.4</i>	WD repeat and SOCS box containing 2	0.79/0.82	-0.33
miR-26	ENSDARG00000052121	<i>rag2</i>	<i>zgc:136743</i>	recombination activating gene 2	0.82/0.83	-0.33
miR-26	ENSDARG00000002771	<i>slc4a5a</i>	<i>si:ch211-48m9.1</i>	<i>solute carrier family 4 member 5a</i>	0.89/0.88	-0.33
miR-26	ENSDARG00000058548	<i>bves</i>	<i>pop1, popdc1, zgc:86887</i>	blood vessel epicardial substance	0.76/0.76	-0.33
miR-26	ENSDARG00000039563	<i>lox13b</i>	<i>ff60c12, im:7157119, wu:ff60c12, zgc:162959</i>	lysyl oxidase-like 3b	0.75/0.74	-0.32
miR-26	ENSDARG00000045601	<i>cax1</i>	<i>si:dkey-180p18.2, wu:fc25f10, zgc:136271</i>	cation/H ⁺ exchanger protein 1	0.81/0.80	-0.32
miR-26	ENSDARG00000044325	<i>zgc:193690</i>	<i>sb:eu534, zgc:193696</i>	<i>zgc:193690</i> ; Orthologous to human iand <i>DDX19B</i> (DEAD-box helicase 19A and 19B)	0.83/0.83	-0.32
miR-26	ENSDARG00000069538	<i>gtf3c4</i>	<i>im:7138272, si:dkeyp-10e10.3</i>	general transcription factor IIIC, polypeptide 4	0.76/0.75	-0.32
miR-26	ENSDARG00000034427	<i>jpt2</i>	<i>fc04g11, hn11, wu:fb37f03, wu:fc04g11, zgc:55712</i>	Jupiter microtubule associated homolog 2	0.77/0.77	-0.32
miR-26	ENSDARG00000079712	<i>gal3st4</i>	<i>fb92g05, wu:fb92g05</i>	galactose-3-O-sulfotransferase 4	0.83/0.84	-0.32
miR-26	ENSDARG00000058357	<i>ankrd33ba</i>	<i>si:rp71-1f1.8</i>	ankyrin repeat domain 33ba	0.71/0.72	-0.32

miR-26	ENSDARG00000045911	<i>tulp4a</i>	<i>fb99g11, si:ch211-222h17.4, si:ch211-51e12.5, tulp4, wu:fb75h04, wu:fb99g11</i>	TUB like protein 4a	0.87/0.89	-0.31
miR-26	ENSDARG00000021664	<i>fzd3a</i>	<i>fz3a, fz9, fzd3, off-limits, olt, zg09</i>	frizzled class receptor 3a	0.90/0.89	-0.31
miR-26	ENSDARG000000041081	<i>kmt5b</i>	<i>suv420h1, wu:fb97g06, wu:fi57g03, zgc:103527</i>	lysine methyltransferase 5B	0.88/0.88	-0.31
miR-26	ENSDARG00000039390	<i>sdhaf4</i>	<i>si:busm1-265n4.4, si:busm1-46p4.2, si:ch211-103o12.3, C13H6orf57</i>	succinate dehydrogenase complex assembly factor 4	0.85/0.82	-0.31
miR-26	ENSDARG00000069540	<i>si:dkey-30c15.2</i>	N/A	<i>si:dkey-30c15.2</i>	0.85/0.87	-0.31
miR-26	ENSDARG00000054937	<i>badb</i>	<i>bad, fa01b12, proapoptotic BH3-only protein, wu:fa01b12, wu:fa96d04</i>	BCL2 associated agonist of cell death b	0.93/0.93	-0.31
miR-26	ENSDARG00000079730	<i>fuz</i>	N/A	fuzzy planar cell polarity protein	0.92/0.90	-0.31
miR-26	ENSDARG00000009021	<i>chrna1</i>	<i>AChR alpha 1, nic, nic-1, nic1, nicotinic receptor, zgc:86593</i>	cholinergic receptor, nicotinic, alpha 1 (muscle)	0.76/0.76	-0.31
miR-26	ENSDARG00000030512	<i>tsnaxip1</i>	<i>wu:fc26b06, zgc:162756</i>	translin-associated factor X interacting protein 1	0.89/0.89	-0.31
miR-26	ENSDARG00000017741	<i>g3bp1</i>	<i>fl17h05, wu:fl17h05, zgc:56034</i>	GTPase activating protein (SH3 domain) binding protein 1	0.94/0.94	-0.31
miR-26	ENSDARG00000078592	<i>nomo</i>	<i>loc283820</i>	nodal modulator	0.80/0.74	-0.31
miR-26	ENSDARG00000088290	<i>foxj1b</i>	<i>fl67h07, foxj1.2, wu:fl67h07, zgc:101713</i>	forkhead box J1b	0.74/0.75	-0.31
miR-26	ENSDARG00000074527	<i>chst15</i>	<i>si:dkey-59k9.3</i>	carbohydrate (N-acetylgalactosamine 4-sulfate 6-O) sulfotransferase 15	0.86/0.82	-0.30
miR-26	ENSDARG00000077950	<i>tor1</i>	<i>im:7165310, si:ch73-178d14.1</i>	torsin family 1	0.73/0.73	-0.30
miR-26	ENSDARG00000022183	<i>gstol</i>	<i>fd58h04, wu:fd58h04, zgc:92254</i>	glutathione S-transferase omega 1	0.81/0.84	-0.30
miR-26	ENSDARG00000011605	<i>dennd6b</i>	<i>fam116b, fl67e07, wu:fl67e07, zgc:55619</i>	DENN/MADD domain containing 6B	0.90/0.88	-0.30
miR-26	ENSDARG000000027199	<i>smad1</i>	<i>SMA- and MAD-related protein 1, fb39h09, madh1, sb:eu786, wu:fb39h09, wu:fb97e04</i>	SMAD family member 1	0.93/0.92	-0.30
miR-26	ENSDARG00000055070	<i>borcs7</i>	<i>zgc:114169, C13H10orf32</i>	BLOC-1 related complex subunit 7	0.70/0.70	-0.30
miR-26	ENSDARG000000092949	<i>si:ch73-366i20.1</i>	N/A	<i>si:ch73-366i20.1</i>	0.76/0.79	-0.30
miR-26	ENSDARG000000068551	<i>elovl8a</i>	<i>zgc:153394</i>	ELOVL fatty acid elongase 8a	0.76/0.76	-0.30
miR-26	ENSDARG00000020618	<i>gatd3a</i>	<i>zgc:112056, C9H21orf33</i>	glutamine amidotransferase like class 1 domain containing 3A	0.77/0.77	-0.30
miR-26	ENSDARG00000030616	<i>nfe2l1a</i>	<i>nfe2l1, nrfla</i>	nuclear factor, erythroid 2-like 1a	0.88/0.88	-0.30
miR-26	ENSDARG000000060383	<i>mmaa</i>	<i>zgc:171657</i>	metabolism of cobalamin associated A	0.88/0.87	-0.29
miR-26	ENSDARG000000002959	<i>gpr34b</i>	<i>zgc:110315</i>	G protein-coupled receptor 34b	0.84/0.83	-0.29
miR-26	ENSDARG000000012025	<i>hs6st2</i>	<i>fb80f12, hs6st, wu:fb80f12</i>	heparan sulfate 6-O-sulfotransferase 2	0.84/0.82	-0.29
miR-26	ENSDARG00000073948	<i>prok1</i>	<i>zgc:171925</i>	prokineticin 1	0.70/0.70	-0.29
miR-26	ENSDARG00000040607	<i>rtf2</i>	<i>rtfdcl, wu:fd99e08, zgc:55448, zgc:86650, C6H20orf43</i>	replication termination factor 2	0.90/0.91	-0.29
miR-26	ENSDARG00000075721	<i>zdhhc6</i>	<i>fa04d12, fc30b09, wu:fa04d12, wu:fc30b09</i>	zinc finger DHHC-type containing 6	0.90/0.90	-0.29
miR-26	ENSDARG00000070256	<i>suz12a</i>	<i>suz12, suz12l, zgc:100863</i>	SUZ12 polycomb repressive complex 2 subunit a	0.71/0.71	-0.29
miR-26	ENSDARG00000036864	<i>slc34a2b</i>	<i>NaPi-IIb2, wu:fi68d06</i>	solute carrier family 34 member 2b	0.76/0.75	-0.29
miR-26	ENSDARG00000012860	<i>tmprss4b</i>	<i>si:ch211-139a5.13, si:ch211-139a5.6, wu:fi21e09, wu:fk84e06</i>	transmembrane serine protease 4b	0.90/0.87	-0.29
miR-26	ENSDARG00000078891	<i>map11</i>	<i>C7orf43</i>	microtubule associated protein 1	0.81/0.81	-0.29
miR-26	ENSDARG00000012932	<i>mat2b</i>	<i>fc55d01, wu:fb48h02, wu:fc55d01, zgc:110308</i>	methionine adenosyltransferase II, beta	0.76/0.76	-0.29
miR-26	ENSDARG000000032103	<i>mapk6</i>	<i>ERK3, si:ch211-235f12.1</i>	mitogen-activated protein kinase 6	0.84/0.81	-0.29
miR-26	ENSDARG00000017803	<i>gsk3ba</i>	<i>GSK-3[b], fb68h05, fk80d11, gsk3b, wu:fb68h05, wu:fk80d11</i>	glycogen synthase kinase 3 beta, genome duplicate a	0.84/0.85	-0.28
miR-26	ENSDARG00000035986	<i>ptpn2b</i>	<i>cb806, ptpn2, ptpn2l, wu:fc10h07, zgc:55339, zgc:76973</i>	protein tyrosine phosphatase non-receptor type 2b	0.85/0.85	-0.28
miR-26	ENSDARG000000043857	<i>ubfm1</i>	<i>ubfm1, wu:fa10f05, wu:fb72g01, zgc:55335</i>	ubiquitin-fold modifier 1	0.72/0.71	-0.28
miR-26	ENSDARG00000010841	<i>fshb</i>	<i>FSH Beta, FSHB, Gth I, gthl</i>	follicle stimulating hormone subunit beta	0.88/0.88	-0.28
miR-26	ENSDARG00000054208	<i>phkg2</i>	<i>zgc:55863</i>	phosphorylase kinase, gamma 2 (testis)	0.86/0.86	-0.28
miR-26	ENSDARG00000040161	<i>zgc:92287</i>	N/A	<i>zgc:92287</i>	0.75/0.74	-0.28
miR-26	ENSDARG00000056196	<i>slc2a2</i>	<i>wu:fb62h10, zglut2</i>	solute carrier family 2 member 2	0.77/0.77	-0.28
miR-26	ENSDARG00000009681	<i>med27</i>	<i>crsp34, crsp8, trap37, zgc:66302</i>	mediator complex subunit 27	0.83/0.83	-0.28
miR-26	ENSDARG00000024503	<i>c6ast3</i>	<i>zgc:103606</i>	six-cysteine containing astacin protease 3	0.75/0.75	-0.28
miR-26	ENSDARG000000091678	<i>ints12</i>	<i>wu:fi17f05, zgc:86602</i>	integrator complex subunit 12	0.78/0.77	-0.28
miR-26	ENSDARG00000008986	<i>e2f7</i>	<i>fc43h03, si:dkey-217k21.2, wu:fc43h03</i>	E2F transcription factor 7	0.77/0.76	-0.28
miR-26	ENSDARG00000053569	<i>sox3</i>	<i>id:ibd2036, sb:cb493, wu:fd02a08, zgc:110279</i>	SRV-box transcription factor 3	0.92/0.92	-0.28
miR-26	ENSDARG00000093378	<i>si:ch211-235i11.5</i>	<i>CR382385.3</i>	<i>si:ch211-235i11.5</i>	0.82/0.82	-0.28

miR-26	ENSDARG00000018190	<i>get3</i>	<i>asna1, trc40, wu:fd05d03, zgc:56540, zgc:86799</i>	guided entry of tail-anchored proteins factor 3, ATPase	0.80/0.80	-0.28
miR-26	ENSDARG00000016837	<i>glipr2l</i>	<i>wu:fj53e10, zgc:103578</i>	GLI pathogenesis-related 2, like	0.80/0.80	-0.28
miR-26	ENSDARG00000018643	<i>igf2a</i>	<i>IGF-II, etID31232.25, igf2, igf2b, wu:fc26e03, zgc:91781</i>	insulin-like growth factor 2a	0.79/0.79	-0.28
miR-26	ENSDARG00000025679	<i>comtb</i>	<i>si:dkey-13a21.15, zgc:162236</i>	catechol-O-methyltransferase b	0.75/0.75	-0.28
miR-26	ENSDARG00000035719	<i>arl5c</i>	<i>zgc:77751</i>	ADP-ribosylation factor-like 5C	0.77/0.76	-0.28
miR-26	ENSDARG00000052527	<i>mul1b</i>	<i>im:7146383, zgc:165594</i>	mitochondrial E3 ubiquitin protein ligase 1b	0.79/0.77	-0.27
miR-26	ENSDARG00000034643	<i>fh13a</i>	<i>fd57e01, fh13, wu:fd57e01</i>	four and a half LIM domains 3a	0.70/-	-0.27
miR-26	ENSDARG00000092199	<i>si:ch211-250c4.4</i>	N/A	<i>si:ch211-250c4.4</i>	0.71/0.73	-0.27
miR-26	ENSDARG00000069980	<i>lman1</i>	<i>fc54c09, fi36e01, wu:fc54c09, wu:fi36e01</i>	lectin, mannose-binding, 1	0.71/0.72	-0.27
miR-26	ENSDARG00000036584	<i>st8sia5</i>	N/A	ST8 alpha-N-acetyl-neuraminide alpha-2,8-sialyltransferase 5	0.94/0.95	-0.27
miR-26	ENSDARG00000056381	<i>cfap97</i>	<i>fj68c08, wu:fj68c08, zgc:85910</i>	cilia and flagella associated protein 97	0.81/0.77	-0.27
miR-26	ENSDARG00000068883	<i>tspan13a</i>	<i>zgc:91858</i>	tetraspanin 13a	0.82/0.81	-0.27
miR-26	ENSDARG00000060054	<i>epc1b</i>	<i>zgc:158364</i>	enhancer of polycomb homolog 1 (Drosophila) b	0.99/0.99	-0.27
miR-26	ENSDARG00000079688	<i>tnrc6a</i>	<i>wu:fc10d10</i>	trinucleotide repeat containing adaptor 6A	0.99/0.99	-0.27
miR-26	ENSDARG00000078759	<i>dna2</i>	<i>si:ch211-1n9.7</i>	DNA replication helicase/nuclease 2	0.85/0.85	-0.27
miR-26	ENSDARG00000057736	<i>ano10</i>	<i>ano10, fc37b04, wu:fc37b04, zgc:114140</i>	anoctamin 10a	0.71/0.70	-0.27
miR-26	ENSDARG00000090646	<i>tnk2a</i>	<i>acka, si:dkey-150o13.3</i>	tyrosine kinase, non-receptor, 2a	0.71/0.71	-0.27
miR-26	ENSDARG00000029724	<i>pqbp1</i>	<i>pqbp1l, si:dz150f13.4, zgc:92646</i>	polyglutamine binding protein 1	0.74/0.75	-0.26
miR-26	ENSDARG00000027590	<i>sept2</i>	<i>nedd5, wu:fb52g01, wu:fb99a01</i>	septin 2	0.76/0.75	-0.26
miR-26	ENSDARG00000021345	<i>prph2lb</i>	<i>prph2l, si:dkey-85n7.2</i>	peripherin 2-like b	0.83/0.81	-0.26
miR-26	ENSDARG00000078430	<i>tiam1a</i>	<i>tiam1</i>	TIAM Rac1 associated GEF 1a	0.74/0.73	-0.26
miR-26	ENSDARG00000004082	<i>cx39.9</i>	<i>cx46l, zgc:110668</i>	connexin 39.9	0.82/0.82	-0.26
miR-26	ENSDARG00000031929	<i>stard14</i>	<i>stard10, wu:fc41g06, zgc:56270, zgc:77753</i>	START domain containing 14	0.73/0.73	-0.26
miR-26	ENSDARG00000010347	<i>acer1</i>	<i>zgc:110285</i>	alkaline ceramidase 1	0.74/0.74	-0.26
miR-26	ENSDARG00000055752	<i>npas4a</i>	<i>npas4, wu:fj52d04, zgc:136764</i>	neuronal PAS domain protein 4a	0.74/0.74	-0.26
miR-26	ENSDARG00000045360	<i>flj11011l</i>	<i>zgc:136503</i>	hypothetical protein FLJ11011-like (H. sapiens)	0.90/0.87	-0.26
miR-26	ENSDARG00000052304	<i>uqcrc1</i>	<i>zgc:73404, zgc:85750</i>	ubiquinol-cytochrome c reductase core protein 1	0.73/0.72	-0.26
miR-26	ENSDARG00000040284	<i>si:dkey-79d12.5</i>	<i>BX005256.1</i>	<i>si:dkey-79d12.5</i> ; orthologous to human <i>BTG3</i> (BTG anti-proliferation factor 3)	0.82/0.80	-0.26
miR-26	ENSDARG00000045910	<i>tmem18l</i>	<i>si:ch211-222h17.5, si:ch211-51e12.4</i>	transmembrane protein 181	0.87/0.87	-0.26
miR-26	ENSDARG00000055381	<i>bambia</i>	<i>Zfma, bambi, cb998, etID64300.12, fc44c12, id:ibd5020, nma, wu:fc44c12, wu:fe01e03</i>	BMP and activin membrane-bound inhibitor (Xenopus laevis) homolog a	0.93/0.91	-0.26
miR-26	ENSDARG00000070348	<i>hoxc10a</i>	<i>hoxc10, z-81</i>	homeobox C10a	0.94/0.94	-0.26
miR-26	ENSDARG000000091539	<i>ptprjb.1</i>	<i>density-enhanced phosphatase-1b, dep1b, ptprib</i>	protein tyrosine phosphatase receptor type Jb, tandem duplicate 1	0.77/0.78	-0.25
miR-26	ENSDARG00000075092	<i>si:dkey-52l18.4</i>	<i>sc:d808</i>	<i>si:dkey-52l18.4</i>	0.77/0.77	-0.25
miR-26	ENSDARG00000016093	<i>qtrt2</i>	<i>qtrtd1</i>	queuine tRNA-ribosyltransferase accessory subunit 2	0.73/0.75	-0.25
miR-26	ENSDARG00000062550	<i>rc3h1a</i>	<i>rc3h1, si:dkey-192m14.4</i>	ring finger and CCCH-type domains 1a	0.85/0.85	-0.25
miR-26	ENSDARG00000057699	<i>styx</i>	<i>si:ch211-270g19.2, zgc:110621</i>	serine/threonine/tyrosine interacting protein	0.74/0.76	-0.25
miR-26	ENSDARG00000040988	<i>tpi1b</i>	<i>DrTPI-B, cb179, wu:fb51c02</i>	triosephosphate isomerase 1b	0.89/0.89	-0.25
miR-26	ENSDARG00000063014	<i>dbpa</i>	<i>dbp1, si:dkey-109n11.1, zdbp1</i>	D site albumin promoter binding protein a	0.71/0.71	-0.25
miR-26	ENSDARG00000028213	<i>ttn.2</i>	<i>ttna, TTN2, pickwick, pik, si:busm1-258d18.1, si:dz258d18.1, ttn, ttna, wu:fi04b11, wu:fi20g08</i>	titin, tandem duplicate 2	0.92/0.92	-0.25
miR-26	ENSDARG00000010276	<i>ptgs2b</i>	<i>Cox-2b, si:dkey-97o5.6</i>	prostaglandin-endoperoxide synthase 2b	0.72/0.72	-0.25
miR-26	ENSDARG00000053362	<i>cilp</i>	<i>im:7137994</i>	cartilage intermediate layer protein, nucleotide pyrophosphohydrolase	0.77/0.76	-0.25
miR-26	ENSDARG00000020893	<i>slc25a55a</i>	<i>cb229, sb:cb229, slc25a22, wu:fa12h05, wu:fb72h03, zgc:56592</i>	solute carrier family 25 member 55a	0.72/0.72	-0.25
miR-26	ENSDARG00000075560	<i>kmt2cb</i>	<i>ml13b, si:dkey-193e13.2</i>	lysine (K)-specific methyltransferase 2Cb	0.77/0.76	-0.25
miR-26	ENSDARG00000073724	<i>nlr1</i>	<i>NLR-A5, Nod5, Nod9</i>	NLR family member X1	0.80/0.78	-0.25
miR-26	ENSDARG00000088505	<i>fam217b</i>	<i>si:dkey-211h10.3</i>	family with sequence similarity 217 member B	0.71/0.71	-0.24
miR-26	ENSDARG00000088318	<i>ubap2a</i>	<i>ubap2, zgc:158449</i>	ubiquitin associated protein 2a	0.71/0.71	-0.24
miR-26	ENSDARG0000008034	<i>skib</i>	<i>fc17g02, skiB, wu:fc17g02, wu:fk79g04</i>	v-ski avian sarcoma viral oncogene homolog b	0.78/0.79	-0.24
miR-26	ENSDARG00000071548	<i>jostl</i>	<i>zgc:153682</i>	Josephin domain containing 1	0.71/0.71	-0.24
miR-26	ENSDARG00000070408	<i>ccnd2b</i>	<i>si:ch211-240j22.2</i>	cyclin D2, b	0.72/0.71	-0.23
miR-26	ENSDARG0000002192	<i>aspn</i>	<i>aspn1, bg13, wu:fk08c11, zgc:109936</i>	asporin (LRR class 1)	0.76/0.76	-0.23

miR-26	ENSDARG00000039892	<i>gnpnat1</i>	<i>im:7138655, zgc:112267</i>	glucosamine-phosphate N-acetyltransferase 1	0.78/0.78	-0.23
miR-26	ENSDARG00000062323	<i>adam23a</i>	<i>wu:fj40f01, wu:fq14a08</i>	ADAM metalloproteinase domain 23a	0.95/0.94	-0.23
miR-26	ENSDARG00000070545	<i>top1l</i>	<i>wu:fc66a02, zgc:136349</i>	DNA topoisomerase I, like	0.87/0.86	-0.23
miR-26	ENSDARG00000025012	<i>tpi1a</i>	<i>DrTPI-A</i>	triosephosphate isomerase 1a	0.72/0.73	-0.23
miR-26	ENSDARG00000028396	<i>fkbp5</i>	<i>si:zc263a23.8, wu:fc31g11, wu:fl87b03, zgc:64082</i>	FKBP prolyl isomerase 5	0.82/0.83	-0.23
miR-26	ENSDARG00000011727	<i>b9d1</i>	<i>im:6905750, im:7069491, wu:fc09g07, zgc:110733</i>	B9 protein domain 1	0.87/0.89	-0.23
miR-26	ENSDARG00000038754	<i>plk3</i>	<i>cnk, wu:fi35f10</i>	polo-like kinase 3 (Drosophila)	0.72/0.71	-0.23
miR-26	ENSDARG00000042846	<i>disp2</i>	N/A	dispatched homolog 2 (Drosophila)	0.84/0.83	-0.23
miR-26	ENSDARG00000052975	<i>ubqln4</i>	<i>ataxin-1 ubiquitin-like interacting protein, ubin, wu:fb48e11, wu:fb54b10, wu:fi41e06, zgc:113928, zgc:66188</i>	ubiquilin 4	0.81/0.80	-0.23
miR-26	ENSDARG00000052950	<i>arhgap23a</i>	<i>sb:eu857</i>	Rho GTPase activating protein 23a	0.88/0.88	-0.22
miR-26	ENSDARG00000000142	<i>tmem267</i>	<i>si:busm1-79m10.4, si:dz79m10.4, zgc:101064, C10H5orf28</i>	transmembrane protein 267	0.77/0.77	-0.22
miR-26	ENSDARG00000003165	<i>nr2f6b</i>	<i>EAR2-B, fc72d04, nr2f6, wu:fc72d04, zgc:77260</i>	nuclear receptor subfamily 2, group F, member 6b	0.85/0.83	-0.22
miR-26	ENSDARG00000059280	<i>hoxd3a</i>	<i>Z-56, hoxd3, im:7144208</i>	homeobox D3a	0.90/0.90	-0.22
miR-26	ENSDARG00000044485	<i>sall4</i>	<i>cb372, cb614, ik:tdsubc_2c1, sb:cb372, sb:cb614, wu:fa01g03, xx:tdsubc_2c1</i>	spalt-like transcription factor 4	0.94/0.93	-0.22
miR-26	ENSDARG00000004218	<i>rnd1b</i>	<i>Rnd1a, rnd1l, wu:fk50c04, zgc:153089</i>	Rho family GTPase 1b	0.78/0.78	-0.22
miR-26	ENSDARG000000035578	<i>hs3st1l2</i>	<i>3-ost-5, z3-ost-5, zgc:136835</i>	heparan sulfate (glucosamine) 3-O-sulfotransferase 1-like 2	0.76/0.75	-0.22
miR-26	ENSDARG00000076163	<i>col19a1</i>	<i>si:dkey-49l3.2, stumpy, sty, zgc:153049</i>	collagen, type XIX, alpha 1	0.77/0.76	-0.22
miR-26	ENSDARG00000015734	<i>aicda</i>	<i>Aid</i>	activation-induced cytidine deaminase	0.72/0.73	-0.22
miR-26	ENSDARG00000025535	<i>clint1a</i>	<i>clint1, enth, enthoproten, epn4, kiaa0171, wu:fa05f05, wu:fb97f11, wu:fc12e09, zgc:103447</i>	clathrin interactor 1a	0.72/0.72	-0.22
miR-26	ENSDARG00000052535	<i>ubxn10</i>	<i>zgc:153648</i>	UBX domain protein 10	0.72/0.72	-0.22
miR-26	ENSDARG00000052011	<i>rrad</i>	<i>fi06a10, wu:fi06a10, zgc:63471</i>	Ras-related associated with diabetes	0.92/0.92	-0.22
miR-26	ENSDARG00000060893	<i>col8a2</i>	<i>si:ch211-197n10.4</i>	collagen, type VIII, alpha 2	0.75/0.74	-0.22
miR-26	ENSDARG00000059596	<i>clip2</i>	N/A	CAP-GLY domain containing linker protein 2	0.74/0.74	-0.22
miR-26	ENSDARG00000092692	<i>pimr133</i>	<i>si:dkey-197d18.1</i>	Pim proto-oncogene, serine/threonine kinase, related 133	0.73/0.75	-0.22
miR-26	ENSDARG00000042732	<i>tmem189</i>	<i>zgc:101801</i>	transmembrane protein 189	-0.70	-0.22
miR-26	ENSDARG00000039064	<i>st3gal7</i>	<i>SIAT5-r, ST3GalV-2, st3gal5-r2, st3gal5l, wu:fb60h07, zgc:153853</i>	ST3 beta-galactoside alpha-2,3-sialyltransferase 7	0.84/0.84	-0.22
miR-26	ENSDARG00000003109	<i>rcan1a</i>	<i>zgc:100963</i>	regulator of calcineurin 1a	0.82/0.82	-0.22
miR-26	ENSDARG00000057497	<i>stk35l</i>	<i>im:7138176, zgc:101750</i>	serine/threonine kinase 35, like	0.78/0.78	-0.21
miR-26	ENSDARG00000036171	<i>rnase13</i>	<i>Dr-RNase1, RNase ZF-3, im:6907218, zf-rnase-3</i>	ribonuclease like 3	0.79/0.79	-0.21
miR-26	ENSDARG00000076052	<i>sez6l2</i>	<i>si:dkey-282h22.6</i>	seizure related 6 homolog (mouse)-like 2	0.82/0.81	-0.21
miR-26	ENSDARG00000036791	<i>dnmt3bb.1</i>	<i>cb633, dnmt3b, dnmt3b1, dnmt4</i>	DNA (cytosine-5-)-methyltransferase 3 beta, duplicate b.1	0.86/0.86	-0.21
miR-26	ENSDARG00000004724	<i>tcea3</i>	<i>zgc:77732</i>	transcription elongation factor A (SII), 3	0.74/0.72	-0.21
miR-26	ENSDARG00000079783	<i>isg20</i>	<i>BX294110.1</i>	interferon stimulated exonuclease gene	0.77/0.78	-0.21
miR-26	ENSDARG00000014386	<i>galnt6</i>	<i>fc23d02, fc56f12, wu:fc23d02, wu:fc56f12, zgc:77836</i>	UDP-N-acetyl-alpha-D-galactosamine:polypeptide N-acetylgalactosaminyltransferase 6 (GalNAc-T6)	0.72/0.72	-0.21
miR-26	ENSDARG00000075757	<i>gig2e</i>	<i>DreE, BX510940.2</i>	grass carp reovirus (GCRV)-induced gene 2e	0.70/0.71	-0.21
miR-26	ENSDARG00000074365	<i>zgc:171901</i>	N/A	<i>zgc:171901</i>	0.75/0.76	-0.21
miR-26	ENSDARG00000057665	<i>sprn2</i>	<i>PrP-rel-1, Sho2, sprnb, wu:fi12d11, zgc:123052</i>	shadow of prion protein 2	0.70/-	-0.21
miR-26	ENSDARG00000073732	<i>myh14</i>	<i>si:ch211-120k19.2, si:ch211-120k19.3</i>	myosin, heavy chain 14, non-muscle	0.79/0.80	-0.21
Total	513					

*The miTG score to miR-26a and miR-26b is presented separately.