

Table S4. Prediction of function in generation of iPS cells for 211 miRNA families.

miRNA family	Seed	B lymph.	MEF.a	MEF.b	NSC.a	NSC.b1	NSC.b2	BJ	MRC5	NFF	NHDF	SC	Mean
miR-124/506	AAGGCAC	-13.5	-22.2	-14.2	-26.5	-27.8	-28.2	-5.2	-8.5	-9.0	-9.3	6.3	-14.4
miR-291b-3p/519a/519b-3p/519c-3p	AAGUGCA	-6.2	-11.6	-10.7	-21.8	-26.9	-21.1	-6.4	-9.3	-14.1	-9.8	-5.5	-13.0
miR-17-5p/20/93.mr/106/519.d	AAAGUGC	-7.1	-11.9	-8.5	-21.4	-22.6	-17.5	-6.3	-6.1	-13.4	-12.1	-3.7	-11.9
miR-200bc/429	AAUACUG	4.7	-6.4	-6.2	-27.3	-22.9	-24.2	-2.6	-3.2	-4.6	-4.1	-6.3	-9.4
miR-30a/30a-5p/30b/30b-5p/30cde/384-5p	GUAAACA	-7.1	-9.7	-5.8	-26.2	-23.7	-19.9	-2.3	-4.2	-4.7	-3.1	3.6	-9.4
miR-19	GUGCAAA	-6.5	-9.9	-8.6	-20.9	-16.3	-22.4	-2.9	-3.7	-5.0	-9.6	3.3	-9.3
miR-130/301	AGUGCAA	-6.3	-7.9	-8.7	-21.7	-19.7	-18.9	-1.4	-4.4	-3.9	-5.6	-3.4	-9.2
miR-106/302	AAGUGCU	-6.0	-4.0	-5.4	-14.9	-18.8	-14.6	-4.2	-4.6	-11.5	-7.2	-3.7	-8.6
miR-340/340-5p	UAUAAAG	5.6	-10.2	-8.6	-25.2	-24.1	-28.8	3.2	5.7	4.0	-5.6	-7.4	-8.3
miR-218	UGUGCUU	-2.5	-8.7	-6.1	-18.7	-25.6	-28.0	2.1	-3.6	-2.2	-1.5	4.1	-8.3
miR-182	UUGGCAA	-6.0	-7.6	-6.7	-17.0	-19.2	-19.5	-5.6	-2.8	-7.7	-5.4	6.7	-8.2
miR-338-5p	ACAAUUAU	-4.4	-5.7	-10.4	-17.0	-20.5	-19.7	2.0	4.3	-4.7	-4.3	-8.3	-8.1
miR-96/1271	UUGGCAC	-6.1	-13.0	-10.0	-16.8	-18.8	-16.3	-4.8	3.4	-7.1	-4.2	7.1	-7.9
miR-590/590-3p	AAUUUUA	-5.1	-6.2	-6.5	-21.0	-15.7	-21.1	3.0	3.4	-4.0	-3.1	-8.7	-7.7
miR-181	ACAUUCA	-4.6	-5.1	-4.6	-19.2	-20.2	-20.5	3.9	3.4	-4.0	-4.7	-6.7	-7.5
miR-25/32/92/92ab/363/367	AUUGCAC	-4.8	-6.2	-7.4	-14.0	-14.4	-15.6	-2.3	-4.8	-5.9	-4.1	-1.2	-7.3
miR-302ac/520f	AGUGCUU	-4.1	-3.5	-3.1	-17.0	-14.9	-13.5	-2.0	-4.3	-5.5	-7.4	-2.9	-7.1
miR-495/1192	AACAAAC	-4.0	-4.4	-7.3	-22.7	-21.5	-22.1	6.0	6.1	6.1	1.7	-13.0	-6.8
miR-142-5p	AUAAAGU	-1.7	-6.0	-5.0	-14.7	-16.8	-12.8	-1.5	-4.4	-3.3	-4.9	-3.0	-6.7
miR-9	CUUUGGU	-5.8	-7.7	-11.2	-11.1	-10.1	-8.9	-6.1	-5.6	-9.6	-4.1	10.3	-6.3
miR-133	UUGGUCC	-3.9	-8.6	-7.4	-12.5	-15.9	-12.6	-2.5	-4.6	-3.5	-2.0	4.4	-6.3
miR-204/211	UCCCUUU	-3.9	-5.5	-8.6	-12.7	-14.1	-14.3	-2.4	-4.8	-3.7	-1.6	2.6	-6.3
miR-410	AUAUAAC	-4.8	-7.9	-4.3	-17.7	-12.1	-16.8	-2.5	4.5	-4.4	-4.1	1.9	-6.2
miR-145	UCCAGUU	-5.1	-4.3	-3.0	-13.8	-14.1	-18.5	-2.1	-2.8	-3.9	-3.3	3.1	-6.1
miR-141/200a	AACACUG	-2.9	-7.5	-4.8	-15.7	-17.7	-24.9	3.2	5.6	3.1	-1.6	-3.6	-6.1
miR-137	UAUUGCU	-4.2	-4.4	-4.7	-15.1	-18.7	-17.6	1.4	2.9	-5.3	-3.2	2.6	-6.0
miR-186	AAAGAAU	-5.4	-3.0	-4.1	-16.0	-12.9	-17.6	3.9	3.1	-2.6	-1.2	-8.1	-5.8
miR-144	ACAGUAU	-3.9	-5.3	-5.1	-17.1	-15.8	-18.6	4.4	3.9	-2.7	2.1	-4.7	-5.7
miR-323/323-3p	ACAUUAC	4.4	-4.6	-3.8	-21.1	-16.7	-14.3	2.8	-5.0	-2.2	2.6	-4.0	-5.6
miR-369/369-3p	AUAAUAC	-2.4	-3.3	-5.2	-14.0	-15.0	-15.1	-1.1	0.9	-2.3	-1.9	-2.3	-5.6
miR-203	UGAAAU	3.0	-5.1	-5.0	-18.7	-18.5	-20.5	5.0	3.6	3.6	-2.8	-3.7	-5.4
miR-512-3p/1186	AGUGCUG	-3.6	-2.5	-4.1	-11.8	-11.7	-12.6	-1.3	-1.3	-4.4	-2.8	-2.2	-5.3
miR-27ab	UCACAGU	-3.8	-8.6	-4.4	-15.0	-15.9	-17.3	3.9	5.0	-3.5	-1.5	3.8	-5.2
miR-300	AUACAAG	10.5	-3.7	-4.3	-22.7	-15.4	-21.0	6.2	9.3	-3.3	-2.5	-9.0	-5.1
miR-101	ACAGUAC	-4.7	-3.8	-4.1	-17.1	-15.2	-20.0	4.6	6.5	2.2	1.7	-5.9	-5.1

miR-543	AACAUUC	-3.9	-2.5	-4.2	-19.7	-16.2	-19.5	5.8	3.9	3.9	2.1	-3.8	-4.9
miR-183	AUGGCAC	-3.4	-5.6	-5.7	-11.4	-18.3	-14.4	2.3	3.4	-1.6	-1.1	2.0	-4.9
miR-448	UGCAUUAU	-3.3	-8.3	-7.0	-12.4	-11.4	-12.9	1.6	4.4	-2.6	-2.9	3.0	-4.7
miR-223	GUCAGUU	-4.6	-3.4	-2.6	-8.2	-12.4	-12.3	1.3	-1.5	-2.7	-1.6	-3.4	-4.7
miR-377	UCACACA	-4.3	-2.7	-3.2	-11.4	-9.5	-17.0	-1.2	2.0	-0.8	1.6	-4.3	-4.6
miR-1/206	GGAAUGU	3.7	-7.2	-4.9	-11.8	-11.1	-16.4	-2.3	2.8	-3.2	-2.1	2.8	-4.5
miR-135	AUGGCUU	3.3	-6.3	-4.0	-11.4	-12.7	-15.4	-1.4	5.7	-3.6	-1.8	-2.2	-4.5
miR-374/374ab	UAUAAUA	-2.4	-2.1	-3.2	-10.2	-9.1	-10.6	-1.6	-1.3	-3.0	-2.3	-3.9	-4.5
miR-23ab	UCACAUU	-4.4	-2.5	-2.2	-16.2	-14.7	-16.9	4.5	5.7	3.3	-1.7	-4.3	-4.5
miR-129/129-5p	UUUUUGC	-6.6	-4.1	-4.9	-11.7	-14.1	-15.3	2.3	4.3	3.2	3.5	-5.8	-4.5
miR-142-3p	GUAGUGU	-3.4	-6.7	-5.9	-8.6	-7.7	-10.9	-1.2	-2.5	-2.7	-1.9	2.4	-4.5
miR-153	UGCAUAG	-5.2	-5.7	-4.0	-11.8	-9.9	-12.8	-1.3	5.0	-2.7	-2.0	2.5	-4.4
miR-384/384-3p	UUCCUAG	-3.0	-4.3	-4.2	-8.7	-11.3	-8.7	1.2	-4.8	1.1	-0.8	-3.2	-4.3
miR-199/199-3p	CAGUAGU	-5.4	-1.8	-1.2	-13.2	-10.8	-11.6	-1.8	2.8	-4.0	-2.3	2.6	-4.2
miR-103/107	GCAGCAU	-2.2	-4.7	-2.5	-13.0	-10.5	-11.2	1.4	4.4	-3.1	-2.2	-2.9	-4.2
miR-568	UGUAUAA	-3.4	-3.6	-2.9	-9.8	-12.4	-9.2	-1.5	1.7	-3.8	-2.1	1.4	-4.1
miR-539	GAGAAAU	-4.2	-2.2	-1.3	-12.2	-11.8	-13.4	2.9	-2.8	-2.1	-1.1	2.6	-4.1
miR-26ab/1297	UCAAGUA	4.8	-4.0	-3.8	-16.1	-19.1	-21.2	4.9	7.6	2.7	2.8	-3.2	-4.1
miR-219/219-5p	GAUUGUC	2.6	-3.7	-3.4	-8.5	-9.5	-13.6	-1.6	-0.9	-1.5	-1.6	-1.7	-3.9
miR-494	GAAACAU	2.1	-2.9	-3.0	-11.9	-7.3	-10.1	2.9	-4.2	-4.0	-1.9	-2.8	-3.9
miR-128	CACAGUG	-3.5	-6.1	-4.2	-12.2	-12.5	-14.6	3.0	4.2	-2.8	2.0	4.0	-3.9
miR-15/16/195/424/497	AGCAGCA	4.3	-8.0	-8.8	-15.0	-15.2	-15.0	4.0	5.7	3.2	1.8	3.3	-3.6
miR-18ab	AAGGUGC	-2.0	-1.3	-2.0	-8.7	-6.9	-8.1	-3.2	-1.5	-4.0	-3.3	2.1	-3.5
miR-409-3p	AAUGUUG	-1.9	-4.4	-4.9	-9.0	-6.7	-11.5	1.0	3.3	3.1	-3.0	-4.2	-3.5
miR-376c	ACAUAGA	-3.1	-4.8	-5.6	-10.2	-6.6	-7.2	0.8	0.9	-1.4	-1.4	0.8	-3.4
miR-31	GGCAAGA	-3.4	-2.2	-3.5	-10.8	-10.5	-10.0	2.3	-1.1	2.8	-1.5	0.3	-3.4
miR-194	GUAACAG	3.2	-2.8	-2.6	-13.0	-13.3	-14.3	2.2	5.3	1.7	1.6	-4.3	-3.3
miR-138	GCUGGUG	2.7	-3.3	-5.6	-7.8	-11.3	-9.6	-1.3	5.5	-1.4	-1.3	-2.3	-3.3
miR-34a/34b-5p/34c/34c-5p/449/449abc/699	GGCAGUG	-2.3	-5.1	-5.7	-8.3	-10.5	-10.7	1.1	3.2	-1.2	2.1	1.9	-3.2
miR-409-5p	GGUUACC	-1.3	-1.3	-3.2	-8.5	-9.1	-11.9	-1.3	2.5	-1.7	-1.2	1.5	-3.2
miR-148/152	CAGUGCA	-1.8	-2.9	-3.1	-12.7	-9.6	-13.5	2.8	6.1	3.0	-1.2	-1.6	-3.1
miR-155	UAAUGCU	-2.2	-2.5	-3.5	-8.4	-6.3	-10.6	1.8	2.3	-1.0	-1.8	-2.0	-3.1
miR-140/140-5p/876-3p	AGUGGUU	-2.4	-2.8	-4.9	-7.7	-5.5	-6.0	-0.9	-1.0	-3.0	-2.6	3.3	-3.0
miR-375	UUGUUCG	-2.8	-3.8	-2.3	-6.5	-6.8	-5.8	1.7	-2.3	-2.0	-1.4	-1.5	-3.0
miR-653	UGUUGAA	-1.3	-1.9	-2.3	-9.6	-6.1	-8.8	-0.9	0.9	0.7	-2.5	-1.6	-3.0
miR-7/7ab	GGAAGAC	-2.2	-2.1	-3.8	-5.9	-5.5	-8.7	1.1	-2.0	-5.1	-1.7	2.9	-3.0
miR-221/222	GCUACAU	-1.6	-4.6	-4.8	-10.2	-12.5	-9.1	4.4	3.4	4.0	2.3	-4.4	-3.0
miR-365	AAUGCCC	-3.4	-1.2	-1.4	-5.5	-3.8	-8.3	-2.4	-1.9	-3.3	-3.5	2.3	-2.9
miR-592/599	UUGUGUC	-1.1	-0.9	-2.3	-7.2	-6.0	-10.4	-1.9	2.2	-1.6	-1.5	-1.4	-2.9
miR-129-3p	AGCCCUU	2.3	-1.8	-2.2	-10.6	-12.6	-11.2	1.3	-1.3	2.6	1.0	0.9	-2.9
miR-320/320abcd	AAAGCUG	8.1	-3.6	-4.2	-16.5	-15.4	-18.4	4.0	2.9	4.9	4.5	2.3	-2.9

miR-143	GAGAUGA	-4.1	-2.6	-2.1	-8.2	-10.4	-7.2	-0.6	2.3	-1.1	1.3	1.7	-2.8
miR-29abc	AGCACCA	7.7	-9.9	-6.2	-9.7	-10.3	-11.3	-4.3	3.1	3.1	-1.0	8.1	-2.8
miR-33/33ab	UGCAUUG	-1.2	-1.6	-1.1	-7.2	-5.7	-11.7	1.0	2.0	-1.9	-1.4	-1.7	-2.8
miR-344-5p/484	CAGGCUC	-2.5	-1.4	-5.3	-6.3	-8.7	-7.3	-1.2	2.2	-1.9	1.0	2.8	-2.6
miR-346	GUCUGCC	-1.7	-2.6	-1.8	-5.9	-4.4	-8.2	-1.3	1.0	-0.7	-1.7	-1.2	-2.6
miR-216/216b	AAUCUCU	-1.7	-4.9	-3.5	-4.7	-3.9	-4.6	-2.3	-1.6	-2.1	-1.0	1.9	-2.6
miR-335/335-5p	CAAGAGC	-0.8	-2.5	-4.3	-4.3	-4.0	-7.5	-1.2	2.5	-2.0	-3.0	-1.2	-2.6
miR-544	UUCUGCA	-2.5	-2.5	-3.3	-8.7	-5.2	-8.7	2.9	2.2	2.3	-1.0	-3.2	-2.5
miR-192/215	UGACCUA	0.9	-6.0	-5.4	-3.7	-3.1	-4.2	-1.1	-1.1	-1.1	-3.2	0.8	-2.5
miR-876-5p	GGAUUUC	-1.2	-1.5	-3.8	-4.7	-5.6	-8.5	1.4	-1.6	-1.3	-1.3	1.3	-2.4
miR-488	UGAAAGG	-3.4	-2.4	1.4	-6.0	-5.1	-7.6	1.6	1.3	-1.4	-2.4	-2.5	-2.4
miR-290-5p/292-5p/371-5p	CUCAAAC	-3.1	-2.1	-3.6	-8.4	-8.3	-11.0	4.0	3.9	4.3	2.2	-3.9	-2.4
miR-485/485-5p	GAGGCUG	-1.9	-1.8	-3.1	-3.5	-3.0	-5.0	-3.2	-2.7	-3.3	-1.0	2.5	-2.4
miR-139-5p	CUACAGU	4.0	1.6	-1.8	-11.8	-9.3	-15.8	4.1	-1.4	3.8	2.7	-1.7	-2.3
miR-205	CCUUCAU	3.2	-3.1	-4.2	-8.4	-9.2	-9.0	1.3	3.3	2.5	1.0	-2.8	-2.3
miR-338/338-3p	CCAGCAU	-2.0	-1.9	-1.6	-5.9	-7.6	-8.9	0.8	1.7	1.8	0.9	-2.0	-2.3
miR-214/761	CAGCAGG	-2.9	-1.9	-3.3	-5.2	-7.9	-11.2	0.5	3.3	2.0	0.8	2.2	-2.1
miR-216/216a	AAUCUCA	-3.2	-3.2	-2.6	-5.0	-4.8	-5.4	0.7	1.5	-1.3	0.7	-1.0	-2.1
miR-763/1207-3p	CAGCUGG	-1.4	-2.6	-5.4	-2.4	-3.6	-4.7	-2.3	-1.2	-2.3	-1.0	3.6	-2.1
miR-709/1827	GAGGCAG	1.0	-3.4	-4.2	-6.7	-6.2	-6.6	-1.9	1.2	-0.7	-1.0	5.6	-2.1
miR-425/489	AUGACAC	1.7	-1.3	-1.8	-4.5	-3.2	-7.1	-2.5	2.3	-3.6	-0.9	-1.9	-2.1
miR-150	CUCCCAA	-1.7	-4.6	-2.7	-5.0	-4.4	-5.8	-1.1	2.0	-1.9	-0.7	3.0	-2.1
miR-501/501-5p	AUCCUUU	-2.5	-1.4	-2.0	-6.2	-5.5	-6.4	1.9	-2.1	2.5	0.9	-1.8	-2.0
miR-190	GAUAUGU	-2.0	-2.7	-1.2	-9.6	-3.7	-4.9	0.5	1.6	2.6	1.8	-4.7	-2.0
miR-542/542-3p	GUGACAG	1.8	-2.6	-2.0	-4.7	-4.5	-7.2	1.4	-2.4	-2.3	-1.6	2.6	-2.0
miR-151-3p	UAGACUG	-2.6	-1.8	-1.3	-4.3	-6.1	-7.1	1.5	2.3	0.7	0.5	-2.6	-1.9
miR-361/361-5p	UAUCAGA	-2.4	-2.0	-1.5	-4.3	-3.8	-6.3	0.8	0.5	-0.7	-0.4	-0.9	-1.9
miR-370	CCUGCUG	-1.3	-1.8	-2.3	-2.5	-3.5	-5.8	-1.5	-3.6	1.4	-1.9	2.3	-1.9
miR-505.hm	GUCAACA	1.6	-2.2	2.7	-6.8	-3.9	-4.6	-1.2	-1.0	-2.2	-0.6	-1.7	-1.8
miR-431	GUCUUGC	-1.7	-1.3	-2.8	-5.0	-3.2	-5.0	-1.5	1.1	2.0	-1.0	-1.4	-1.8
miR-877	UAGAGGA	0.9	-1.2	-2.1	-4.1	-2.8	-4.7	-1.4	0.3	-2.3	-1.6	-0.8	-1.8
miR-217	ACUGCAU	4.2	0.9	-1.5	-8.9	-9.4	-12.4	3.8	3.6	1.9	0.9	-2.6	-1.8
miR-873	CAGGAAC	-2.5	-1.3	-1.4	-6.3	-7.6	-9.9	1.9	3.6	1.4	1.4	1.4	-1.7
miR-496	GAGUAUU	1.2	-2.5	-1.8	-5.2	-3.5	-4.9	-1.7	-0.5	-1.2	-0.8	1.7	-1.7
miR-379	GGUAGAC	-1.4	-3.5	-2.9	-2.0	-1.9	-2.6	0.6	-1.0	-0.8	-1.5	-1.9	-1.7
miR-671-5p	GGAAGCC	-0.9	-3.6	-1.1	-1.6	-2.9	-3.5	-4.1	1.4	-3.4	-2.9	3.9	-1.7
miR-329/362-3p	ACACACC	-1.5	1.9	-2.6	-6.5	-3.8	-8.0	1.6	1.8	-0.9	1.9	-2.2	-1.7
miR-499/499-5p	UAAGACU	-1.8	-2.0	-2.1	-5.7	-5.9	-8.0	3.1	2.5	1.7	2.8	-2.7	-1.7
miR-21/590-5p	AGCUUAU	3.3	1.1	-2.4	-3.7	-3.7	-6.0	-1.4	1.7	-2.0	-1.6	-3.4	-1.7
miR-122	GGAGUGU	-1.7	-3.8	-4.3	-2.7	-2.9	-3.6	-2.3	1.7	-2.1	0.4	3.3	-1.6
miR-376/376ab/376b-3p	UCAUAGA	2.4	-4.5	-5.9	-3.0	-2.4	-1.7	1.7	-1.6	-3.3	-1.1	1.4	-1.6
miR-149	CUGGCUC	-1.4	-4.1	-2.4	-3.3	-3.9	-7.4	1.0	1.9	-0.9	-0.9	3.5	-1.6

miR-508-5p	ACUCCAG	1.6	-2.6	-1.2	-5.3	-7.5	-8.7	1.8	1.0	0.9	1.1	1.3	-1.6
miR-500/501-3p/502/502-3p	AUGCACC	2.2	-1.3	-1.5	-6.9	-6.2	-10.8	2.5	1.8	2.5	0.7	-0.5	-1.6
miR-486/486-5p	CCUGUAC	0.7	-1.5	-1.5	-6.1	-6.1	-5.0	0.7	1.9	0.6	0.8	-1.9	-1.6
miR-199/199-5p	CCAGUGU	3.1	-2.0	-2.9	-7.1	-6.6	-6.1	1.3	3.5	0.9	0.9	-1.9	-1.5
miR-881/892a	ACUGUGU	1.1	-1.3	-1.4	-4.3	-3.7	-4.5	1.3	1.1	0.9	-1.7	-3.6	-1.5
miR-654-3p	AUGUCUG	-3.2	-0.9	0.9	-5.1	-6.8	-5.8	3.3	1.3	2.6	1.6	-4.2	-1.5
miR-581/669d	CUUGUGU	-1.3	-1.3	-3.8	-2.7	-2.2	-2.0	-1.1	-1.9	2.1	0.5	-2.2	-1.4
miR-326/330/330-5p	CUCUGGG	2.1	-3.6	-2.7	-2.7	-5.2	-5.9	-1.0	3.0	-1.1	-0.7	2.5	-1.4
miR-185/882	GGAGAGA	-1.8	-2.5	-3.8	-2.2	-1.9	-3.6	-0.5	2.5	-1.6	-1.3	1.4	-1.4
miR-503	AGCAGCG	2.6	-2.1	-4.9	-7.0	-6.7	-7.4	2.3	4.3	5.0	1.9	-2.5	-1.3
miR-1224/1224-5p	UGAGGAC	-2.0	-1.6	-2.2	-2.7	-3.2	-5.0	0.9	2.1	-2.6	-2.0	4.0	-1.3
miR-758	UUGUGAC	-0.9	-4.7	-0.8	-5.2	-3.0	-4.6	0.7	2.9	1.5	-2.3	2.0	-1.3
miR-532/532-5p	AUGCCUU	2.2	-1.9	-1.8	-4.6	-4.1	-4.5	-1.1	1.6	1.5	0.8	-2.3	-1.3
miR-297/297a/297b-5p/297c	UGUAUGU	-2.3	-1.9	-1.5	-4.6	-2.1	-6.3	1.5	2.8	1.7	0.5	-1.9	-1.3
miR-146	GAGAACU	-1.2	-1.4	-1.3	-4.2	-3.4	-3.2	-0.8	1.8	0.8	0.7	-1.4	-1.2
miR-668	GUCACUC	1.4	-2.6	-2.4	-2.8	-3.8	-2.9	1.1	-1.1	-0.8	-1.1	1.3	-1.2
miR-10	ACCCUGU	3.0	-2.4	-4.2	-4.6	-5.6	-4.4	1.8	1.7	1.3	3.3	-3.4	-1.2
miR-132/212	AACAGUC	2.3	-1.6	-1.2	-8.0	-6.5	-8.0	2.3	2.0	2.7	4.3	-1.8	-1.2
miR-328	UGGCCCCU	-1.2	-2.7	-4.6	-2.4	-3.0	-1.7	-0.7	0.4	-1.3	-0.9	4.9	-1.2
miR-154	AGGUUAU	2.1	-1.1	-2.9	-3.6	-2.8	-3.6	0.9	2.3	-1.5	0.7	-2.5	-1.1
miR-760/1842	GGCUCUG	2.8	-2.4	-2.1	-3.7	-4.8	-2.8	-1.0	1.6	-1.4	-0.4	2.4	-1.1
miR-22	AGCUGCC	1.6	-2.1	-2.4	-5.8	-6.2	-9.0	1.3	4.5	3.1	1.3	2.7	-1.0
miR-193ab	ACUGGCC	-3.1	1.4	1.0	-4.6	-6.7	-7.1	4.5	2.7	2.6	1.2	-2.8	-1.0
miR-324-5p	GCAUCCC	-1.0	-1.0	-2.9	-5.2	-4.4	-5.3	2.5	2.5	1.6	1.6	0.9	-1.0
miR-1197	AGGACAC	-1.4	-1.5	-1.0	-6.2	-4.6	-7.7	1.8	3.0	3.2	1.9	2.3	-0.9
miR-296/296-3p	AGGGUUG	-0.9	-1.8	-2.0	-3.2	-2.5	-2.9	1.4	1.5	-1.7	1.2	1.0	-0.9
miR-665	CCAGGAG	-1.1	-1.8	-2.1	-3.9	-2.9	-6.1	1.1	4.2	-1.3	1.3	3.1	-0.9
miR-202/202-3p	GAGGUAU	2.0	-4.9	-2.1	-6.8	-5.3	-5.7	1.7	5.0	2.5	2.5	1.8	-0.8
miR-455/455-3p	CAGUCCA	2.7	-1.4	-0.7	-5.4	-6.3	-6.5	2.7	4.2	2.6	2.2	-3.0	-0.8
miR-299/299-3p	AUGUGGG	-1.2	-1.8	-1.0	-2.6	-1.5	-2.9	0.8	1.0	0.7	-1.1	0.9	-0.8
miR-136	CUCCAUU	1.9	-2.9	-2.7	-2.1	-3.1	-7.7	1.8	0.7	1.6	1.2	2.8	-0.8
miR-433	UCAUGAU	-3.7	-0.6	-0.9	-5.8	-4.6	-2.9	3.7	2.3	3.3	-1.4	2.1	-0.8
miR-875-5p	AUACCUC	0.8	1.1	-0.8	-3.2	-2.2	-3.1	1.2	1.1	-0.6	-1.1	-1.6	-0.8
miR-362-5p	AUCCUUG	-1.3	-0.7	-1.7	-1.8	-2.3	-3.7	1.1	2.8	2.4	-0.6	-2.4	-0.7
miR-125/351	CCCUGAG	-3.7	-1.3	-2.1	-5.9	-7.6	-4.8	1.0	5.3	2.1	1.6	7.3	-0.7
miR-34b-3p/34c-3p	AUCACUA	1.9	0.9	2.0	-1.3	-3.3	-5.8	1.6	-2.6	2.9	-1.1	-3.2	-0.7
miR-191	AACGGAA	-0.5	-2.4	-1.2	-2.0	-1.1	-1.3	0.8	0.7	-0.5	-0.7	0.9	-0.7
miR-574-5p	GAGUGUG	-2.8	-1.1	-1.3	-3.3	-1.8	-2.6	1.4	2.8	1.6	1.1	-1.1	-0.7
miR-134	GUGACUG	-1.3	2.0	2.9	-1.5	-2.5	-1.6	-1.6	1.3	-1.1	-2.1	-1.5	-0.6
miR-208/208ab	UAAGACG	-1.6	1.4	-0.7	-3.7	-4.2	-5.7	3.9	2.0	1.6	3.2	-2.9	-0.6
miR-342/342-3p	CUCACAC	1.6	-1.5	-2.4	-3.3	-2.3	-6.5	1.3	4.1	1.0	2.9	-1.8	-0.6
miR-342-5p	GGGGUGC	-3.5	-1.4	-1.9	-1.5	-1.2	-2.2	-0.4	0.7	-1.3	0.5	5.4	-0.6

miR-127	CGGAUCC	-0.7	1.0	-0.7	-1.1	-1.1	-0.7	-0.7	-0.1	-1.6	-0.9	-0.1	-0.6
miR-296-5p	GGGCCCC	-0.7	-1.5	-2.9	-2.0	-0.8	-1.4	-1.4	0.5	-0.6	0.5	3.8	-0.6
miR-187	CGUGUCU	-1.4	-0.7	-0.2	-1.2	-1.6	-1.7	-0.4	1.3	-0.6	-0.6	0.7	-0.6
miR-378/422a	CUGGACU	2.0	-1.2	-1.6	-2.5	-4.1	-3.8	1.3	3.2	0.9	1.1	-1.5	-0.6
miR-339-5p	CCCUGUC	1.3	-2.1	-1.5	-4.5	-3.6	-4.5	1.7	3.7	2.8	1.8	-0.8	-0.5
let-7/98	GAGGUAG	3.4	-3.6	-3.3	-5.7	-4.0	-5.1	0.9	5.3	2.2	2.5	1.9	-0.5
miR-582-3p/1267	CUGUUGA	1.7	-1.1	-2.5	-2.0	-1.5	-4.9	1.7	0.4	0.8	0.7	1.5	-0.5
miR-224	AAGUCAC	-4.2	1.8	4.3	-3.5	-2.4	-5.1	2.6	1.7	1.2	1.1	-2.9	-0.5
miR-210	UGUGCGU	0.2	-0.2	-0.4	-0.8	-1.0	-0.6	-0.9	-0.8	-1.7	1.2	-0.2	-0.5
miR-331-5p	UAGGUAU	0.6	-1.4	-0.5	-2.3	-1.7	-1.4	0.7	0.5	0.5	0.6	-0.6	-0.5
miR-126/126-3p	CGUACCG	1.1	-0.6	0.4	-1.0	-2.3	-1.8	0.3	-0.4	-0.5	-0.6	0.5	-0.4
miR-504	GACCCUG	-0.8	-1.7	-2.4	-2.2	-2.2	-1.3	1.3	0.9	1.5	1.4	0.7	-0.4
miR-487/487b	AUCGUAC	-0.4	-0.8	-0.4	-0.6	-1.6	-1.8	-0.3	1.1	0.0	0.1	0.2	-0.4
miR-421	UCAACAG	1.7	3.4	2.7	-7.7	-5.3	-8.4	3.9	4.9	5.3	1.8	-6.8	-0.4
miR-675/675-5p	GGUGCGG	-0.6	-0.3	-1.7	-1.3	-1.1	-0.5	-0.4	0.6	-0.1	1.2	-0.1	-0.4
miR-411	AGUAGAC	1.3	-2.2	-0.3	-0.7	-3.2	-4.4	2.8	1.5	1.2	1.0	-1.1	-0.4
miR-652	AUGGCGC	-0.7	-0.5	-0.5	-0.4	-0.3	-0.7	-0.6	0.9	-0.6	0.6	-1.3	-0.4
miR-744	GCGGGGC	0.4	-0.3	-1.4	-0.6	-0.4	-0.7	-0.8	-0.3	-0.8	0.9	0.1	-0.4
miR-493	GAAGGUC	0.5	1.3	0.6	-1.4	-1.4	-4.1	0.8	0.2	-1.2	-0.5	1.3	-0.3
miR-874	UGCCCUG	-0.7	-0.6	0.9	-1.4	-2.4	-1.6	-0.9	1.7	-1.6	-1.1	4.5	-0.3
miR-196ab	AGGUAGU	5.6	2.4	-3.5	-4.1	-3.8	-7.8	3.2	2.6	1.4	2.5	-1.7	-0.3
miR-323-5p	GGUGGUC	1.5	-0.7	-1.5	-0.9	-0.7	-2.8	0.7	-1.6	1.5	0.8	0.7	-0.3
miR-382	AAGUUGU	0.8	1.2	1.9	-5.2	-2.9	-4.3	1.7	4.6	2.0	-1.2	-1.5	-0.3
miR-147/147b	UGUGCGG	0.2	-0.2	-0.2	-1.1	-1.1	-0.6	-0.6	-0.5	-0.3	1.1	0.4	-0.3
miR-450a/450a-5p	UUUGCGA	-0.2	-0.3	-0.4	-0.5	-1.3	-0.9	0.4	0.6	0.6	0.3	-0.7	-0.2
miR-615-5p	GGGGUCC	0.0	-0.3	-2.1	-0.2	-0.8	-1.1	-0.5	0.7	0.4	0.3	1.4	-0.2
miR-125a-3p	CAGGUGA	3.2	-1.0	0.8	-2.5	-2.6	-4.2	1.8	0.8	-1.3	1.3	1.6	-0.2
miR-188/188-5p	AUCCCUU	1.0	0.8	2.1	-3.4	-3.7	-4.8	3.6	3.0	1.9	-1.1	-1.1	-0.1
miR-151	CGAGGAG	-0.6	-0.1	-0.3	-0.3	-0.1	-0.5	0.3	-0.3	0.2	0.4	-0.1	-0.1
miR-184	GGACGGA	-0.6	0.7	0.7	-1.2	-1.4	-1.5	-1.0	0.5	1.1	0.4	1.1	-0.1
miR-28/28-5p/708	AGGAGCU	1.2	-1.2	-1.4	-1.4	-1.6	-2.2	0.7	1.7	1.1	0.6	1.3	-0.1
miR-383	GAUCAGA	1.0	0.8	-1.5	-2.9	-1.7	-3.4	-0.9	4.2	1.8	0.5	2.2	0.0
miR-369-5p	GAUCGAC	-1.4	0.3	-0.8	-0.3	0.5	0.2	-0.2	1.0	0.3	0.8	-0.3	0.0
miR-339-3p	GAGCGCC	0.6	0.5	0.4	1.1	0.7	0.3	-1.4	0.2	-2.1	-1.9	2.2	0.0
miR-197	UCACCAC	4.0	-1.8	-1.6	-3.6	-4.8	-5.9	3.3	2.2	3.5	3.0	2.0	0.0
miR-483/483-5p	AGACGGG	-0.1	2.4	0.7	-0.8	0.1	-0.3	0.6	-0.4	0.5	-0.5	-1.1	0.1
miR-532-3p	CUCCAC	1.5	-0.8	-1.2	-2.1	-1.4	-3.2	0.9	4.3	0.7	0.8	1.3	0.1
miR-491/491-5p	GUGGGGA	-0.5	-1.1	-1.5	1.0	-0.2	-0.3	-1.5	0.9	-0.6	1.1	3.7	0.1
miR-551ab	CGACCCA	-0.6	-0.4	-1.3	-0.2	0.4	0.1	0.4	1.6	0.3	1.2	-0.3	0.1
miR-331/331-3p	CCCCUGG	0.7	0.9	-1.5	-1.0	-0.4	-0.8	-1.0	2.8	-0.3	0.9	0.9	0.1
miR-574/574-3p	ACGCUCA	0.5	0.3	0.6	0.5	-0.1	0.2	0.4	-1.1	1.1	-0.4	-0.4	0.1
miR-615-3p	CCGAGCC	-1.0	1.1	1.5	0.8	0.7	-1.3	0.4	0.1	-0.5	0.3	-0.3	0.2
miR-188-3p	UCCCACA	4.3	0.7	-0.7	-3.8	-1.6	-3.1	2.0	3.9	1.2	1.8	-3.0	0.2
miR-598/598-3p	ACGUCAU	2.2	-0.8	0.9	-0.8	-0.7	-2.0	0.8	1.1	1.1	1.2	-0.5	0.2

miR-99ab/100	ACCCGUA	-0.9	-0.7	-1.2	-0.4	-0.6	-0.7	2.2	2.5	0.6	2.5	-0.8	0.2
miR-671/671-3p	CCGGUUC	-0.4	0.3	-0.2	0.4	0.7	0.7	0.3	0.7	0.6	0.3	-0.3	0.3
miR-423a/423-5p	GAGGGGC	1.2	1.1	-3.3	-0.6	-0.5	-1.1	-0.9	2.4	-0.8	1.5	4.1	0.3
miR-451	AACCGUU	0.3	0.8	0.5	-0.4	0.5	-0.2	0.8	1.5	0.2	1.3	-2.0	0.3
miR-423/423-3p	GCUCGGU	0.2	0.2	0.7	0.4	0.2	0.3	0.3	1.7	0.6	2.0	-0.7	0.5
miR-24	GGCUCAG	1.0	0.9	-1.3	-3.2	-2.9	-3.7	2.9	4.7	1.3	5.0	1.8	0.6
miR-490/490-3p	AACCUGG	3.4	1.4	2.3	-2.6	-3.1	-2.9	3.3	2.7	5.0	1.7	-1.9	0.8

Values in the table are significance scores (SS) described in Methods. miRNA families are sorted by the mean SS.