

Plate Name(s)	Well Position	Sequence Name	Sequence
Human-miRNome_S15-Plate #1			
Plate 1	A1	hsa-let-7a	TGAGGTAGTAGTTGTATAGTT
Plate 1	A2	hsa-let-7b	TGAGGTAGTAGTTGTGTGGTT
Plate 1	A3	hsa-let-7c	TGAGGTAGTAGTTGTATGGTT
Plate 1	A4	hsa-let-7d	AGAGGTAGTAGTTGCATAGTT
Plate 1	A5	hsa-let-7e	TGAGGTAGTAGTTGTATAGTT
Plate 1	A6	hsa-let-7f	TGAGGTAGTAGTTGTATAGTT
Plate 1	A7	hsa-let-7g	TGAGGTAGTAGTTGTACAGTT
Plate 1	A8	hsa-let-7i	TGAGGTAGTAGTTGTGCTGTT
Plate 1	A9	hsa-miR-1	TGGAATGTAAGAAGTATGTAT
Plate 1	A10	hsa-miR-7	TGGAAGACTAGTGATTTTGTGTT
Plate 1	A11	hsa-miR-9	TCTTTGGTTATCTAGCTGTATGA
Plate 1	A12	hsa-miR-10a	TACCCTGTAGATCCGAATTTGTG
Plate 1	A13	hsa-miR-10b	TACCCTGTAGAACCGAATTTGTG
Plate 1	A14	hsa-miR-15a	TAGCAGCACATAATGGTTTGTG
Plate 1	A15	hsa-miR-15b	TAGCAGCACATCATGGTTTACA
Plate 1	A16	hsa-miR-16	TAGCAGCACGTAATATTGGCG
Plate 1	A17	hsa-miR-17	CAAAGTGCTTACAGTGCAGGTAG
Plate 1	A18	hsa-miR-18a	TAAGGTGCATCTAGTGCAGATAG
Plate 1	A19	hsa-miR-18b	TAAGGTGCATCTAGTGCAGTTAG
Plate 1	A20	hsa-miR-19a	TGTGCAAATCTATGCAAACTGA
Plate 1	A21	hsa-miR-19b	TGTGCAAATCCATGCAAACTGA
Plate 1	A22	hsa-miR-20a	TAAAGTGCTTATAGTGCAGGTAG
Plate 1	A23	hsa-miR-20b	CAAAGTGCTCATAGTGCAGGTAG
Plate 1	A24	hsa-miR-21	TAGCTTATCAGACTGATGTTGA
Plate 1	B1	hsa-miR-22	AAGCTGCCAGTTGAAGAACTGT
Plate 1	B2	hsa-miR-23a	ATCACATTGCCAGGGATTCC
Plate 1	B3	hsa-miR-23b	ATCACATTGCCAGGGATTACC
Plate 1	B4	hsa-miR-24	TGGCTCAGTTCAGCAGGAACAG
Plate 1	B5	hsa-miR-25	CATTGCACCTGTCTCGGTCTGA
Plate 1	B6	hsa-miR-26a	TTCAAGTAATCCAGGATAGGCT
Plate 1	B7	hsa-miR-26b	TTCAAGTAATCCAGGATAGGT
Plate 1	B8	hsa-miR-27a	TTACAGTGGCTAAGTTCCGC
Plate 1	B9	hsa-miR-27b	TTACAGTGGCTAAGTTCTGC
Plate 1	B10	hsa-miR-28-3p	CACTAGATTGTGAGCTCCTGGA
Plate 1	B11	hsa-miR-28-5p	AAGGAGCTCACAGTCTATTGAG
Plate 1	B12	hsa-miR-29a	TAGCACCATCTGAAATCGGTTA
Plate 1	B13	hsa-miR-29b	TAGCACCATTTGAAATCAGTGTT
Plate 1	B14	hsa-miR-29c	TAGCACCATTTGAAATCGGTTA
Plate 1	B15	hsa-miR-30a	TGTAACATCCTCGACTGGAAG
Plate 1	B16	hsa-miR-30b	TGTAACATCCTACACTCAGCT
Plate 1	B17	hsa-miR-30c	TGTAACATCCTACACTCTCAGC
Plate 1	B18	hsa-miR-30d	TGTAACATCCTCGACTGGAAG
Plate 1	B19	hsa-miR-30e	TGTAACATCCTGACTGGAAG
Plate 1	B20	hsa-miR-31	AGGCAAGATGCTGGCATAGCT
Plate 1	B21	hsa-miR-32	TATTGCACATTACTAAGTTGCA
Plate 1	B22	hsa-miR-33a	GTGCATTGTAGTTGCATTGCA
Plate 1	B23	hsa-miR-33b	GTGCATTGCTGTTGCATTGC
Plate 1	B24	hsa-miR-34a	TGGCAGTGTCTTAGCTGGTTGT
Plate 1	C1	hsa-miR-34b	CAATCACTAACTCCACTGCCAT
Plate 1	C2	hsa-miR-34c-3p	AATCACTAACCACACGGCCAGG
Plate 1	C3	hsa-miR-34c-5p	AGGCAGTGTAGTTAGCTGATTGC
Plate 1	C4	hsa-miR-92a	TATTGCACCTGTCCCGCCTGT
Plate 1	C5	hsa-miR-92b	TATTGCACCTGTCCCGCCTCC
Plate 1	C6	hsa-miR-93	CAAAGTGCTGTTCTGCAGGTAG
Plate 1	C7	hsa-miR-95	TTCAACGGGTATTTATTGAGCA
Plate 1	C8	hsa-miR-96	TTTGGCATTAGCACATTTTGTCT
Plate 1	C9	hsa-miR-98	TGAGGTAGTAAGTTGTATTGTT
Plate 1	C10	hsa-miR-99a	AACCCGTAGATCCGATCTTGTG
Plate 1	C11	hsa-miR-99b	CACCCGTAGAACCGACCTTGCG
Plate 1	C12	hsa-miR-100	AACCCGTAGATCCGAACCTTGTG
Plate 1	C13	hsa-miR-101	TACAGTACTGTGATAACTGAA
Plate 1	C14	hsa-miR-103	AGCAGCATTGTACAGGGCTATGA

Plate 1	C15	hsa-miR-103-as	TCATAGCCCTGTACAATGCTGCT
Plate 1	C16	hsa-miR-105	TCAAATGCTCAGACTCCTGTGGT
Plate 1	C17	hsa-miR-106a	AAAAGTGCTTACAGTGCAGGTAG
Plate 1	C18	hsa-miR-106b	TAAAGTGCTGACAGTGCAGAT
Plate 1	C19	hsa-miR-107	AGCAGCATTGTACAGGGCTATCA
Plate 1	C20	hsa-miR-122	TGGAGTGTGACAATGGTGTTTG
Plate 1	C21	hsa-miR-124	TAAGGCACGGGTGAATGCC
Plate 1	C22	hsa-miR-125a-3p	ACAGGTGAGGTTCTTGGGAGCC
Plate 1	C23	hsa-miR-125a-5p	TCCCTGAGACCCTTTAACCTGTGA
Plate 1	C24	hsa-miR-125b	TCCCTGAGACCCTAACCTGTGA
Plate 1	D1	hsa-miR-126	TCGTACCGTGAGTAATAATGCG
Plate 1	D2	hsa-miR-127-3p	TCGGATCCGTCTGAGCTTGGCT
Plate 1	D3	hsa-miR-127-5p	CTGAAGCTCAGAGGGCTCTGAT
Plate 1	D4	hsa-miR-128	TCACAGTGAACCGGTCTCTTT
Plate 1	D5	hsa-miR-129-3p	AAGCCCTTACCCCAAAAGCAT
Plate 1	D6	hsa-miR-129-5p	CTTTTTGCGGTCTGGGCTTGC
Plate 1	D7	hsa-miR-130a	CAGTGCATGTTAAAAGGGCAT
Plate 1	D8	hsa-miR-130b	CAGTGCATGATGAAAGGGCAT
Plate 1	D9	hsa-miR-132	TAACAGTCTACAGCCATGGTCG
Plate 1	D10	hsa-miR-133a	TTTGGTCCCCTTCAACCAGCTG
Plate 1	D11	hsa-miR-133b	TTTGGTCCCCTTCAACCAGCTA
Plate 1	D12	hsa-miR-134	TGTGACTGGTTGACCAGAGGGG
Plate 1	D13	hsa-miR-135a	TATGGCTTTTTATTCTATGTGA
Plate 1	D14	hsa-miR-135b	TATGGCTTTTCATTCTATGTGA
Plate 1	D15	hsa-miR-136	ACTCCATTTGTTTTGATGATGGA
Plate 1	D16	hsa-miR-137	TTATTGCTTAAGAATACGCGTAG
Plate 1	D17	hsa-miR-138	AGCTGGTGTGTGAATCAGGCCG
Plate 1	D18	hsa-miR-139-3p	GGAGACGCGGCCCTGTTGGAGT
Plate 1	D19	hsa-miR-139-5p	TCTACAGTGCACGTGTCTCCAG
Plate 1	D20	hsa-miR-140-3p	TACCACAGGGTAGAACCACGG
Plate 1	D21	hsa-miR-140-5p	CAGTGGTTTTACCCTATGGTAG
Plate 1	D22	hsa-miR-141	TAACACTGTCTGGTAAAGATGG
Plate 1	D23	hsa-miR-142-3p	TGTAGTGTTCCTACTTTATGGA
Plate 1	D24	hsa-miR-142-5p	CATAAAGTAGAAAGCACTACT
Plate 1	E1	hsa-miR-143	TGAGATGAAGCACTGTAGCTC
Plate 1	E2	hsa-miR-144	TACAGTATAGATGATGTACT
Plate 1	E3	hsa-miR-145	GTCCAGTTTTCCAGGAATCCCT
Plate 1	E4	hsa-miR-146a	TGAGAACTGAATTCATGGGTT
Plate 1	E5	hsa-miR-146b-3p	TGCCCTGTGGACTCAGTTCTGG
Plate 1	E6	hsa-miR-146b-5p	TGAGAACTGAATTCATAGGCT
Plate 1	E7	hsa-miR-147	GTGTGTGAAATGCTTCTGC
Plate 1	E8	hsa-miR-147b	GTGTGCGGAAATGCTTCTGCTA
Plate 1	E9	hsa-miR-148a	TCAGTGCACACAGAACTTTGT
Plate 1	E10	hsa-miR-148b	TCAGTGCATCACAGAACTTTGT
Plate 1	E11	hsa-miR-149	TCTGGCTCCGTGTCTTCACTCCC
Plate 1	E12	hsa-miR-150	TCTCCCAACCCTTGTAACAGTG
Plate 1	E13	hsa-miR-151-3p	CTAGACTGAAGCTCCTTGAGG
Plate 1	E14	hsa-miR-151-5p	TCGAGGAGCTCACAGTCTAGT
Plate 1	E15	hsa-miR-152	TCAGTGCATGACAGAACTTGG
Plate 1	E16	hsa-miR-153	TTGCATAGTCACAAAAGTGATC
Plate 1	E17	hsa-miR-154	TAGGTTATCCGTGTTGCCTTCG
Plate 1	E18	hsa-miR-155	TTAATGCTAATCGTGATAGGGGT
Plate 1	E19	hsa-miR-181a	AACATTCACGCTGTCGGTGAGT
Plate 1	E20	hsa-miR-181b	AACATTCATTGCTGTCGGTGGGT
Plate 1	E21	hsa-miR-181c	AACATTCACCTGTCGGTGAGT
Plate 1	E22	hsa-miR-181d	AACATTCATTGTTGTCGGTGGGT
Plate 1	E23	hsa-miR-182	TTTGCAATGGTAGAACTCACACT
Plate 1	E24	hsa-miR-183	TATGGCACTGGTAGAATTCACT
Plate 1	F1	hsa-miR-184	TGGACGGAGAAGTATAAGGGT
Plate 1	F2	hsa-miR-185	TGGAGAGAAAGGCAGTTCCTGA
Plate 1	F3	hsa-miR-186	CAAAGAATTCTCCTTTGGGCT
Plate 1	F4	hsa-miR-187	TCGTGTCTTGTGTTGCAGCCGG
Plate 1	F5	hsa-miR-188-3p	CTCCCACATGCAGGGTTTGCA
Plate 1	F6	hsa-miR-188-5p	CATCCCTTGCATGGTGGAGGG
Plate 1	F7	hsa-miR-190	TGATATGTTTGATATATTAGGT
Plate 1	F8	hsa-miR-190b	TGATATGTTTGATATTGGGTT

Plate 1	F9	hsa-miR-191	CAACGGAATCCCAAAAGCAGCTG
Plate 1	F10	hsa-miR-192	CTGACCTATGAATTGACAGCC
Plate 1	F11	hsa-miR-193a-3p	AACTGGCCTACAAAGTCCCAGT
Plate 1	F12	hsa-miR-193a-5p	TGGGTCTTTGGGGCGAGATGA
Plate 1	F13	hsa-miR-193b	AACTGGCCCTCAAAGTCCCGCT
Plate 1	F14	hsa-miR-194	TGTAACAGCAACTCCATGTGGA
Plate 1	F15	hsa-miR-195	TAGCAGCACAGAAATATTGGC
Plate 1	F16	hsa-miR-196a	TAGGTAGTTTCATGTTGTTGGG
Plate 1	F17	hsa-miR-196b	TAGGTAGTTTCCTGTTGTTGGG
Plate 1	F18	hsa-miR-197	TTCACCACCTTCTCCACCCAGC
Plate 1	F19	hsa-miR-198	GGTCCAGAGGGGAGATAGGTTT
Plate 1	F20	hsa-miR-199a-3p	ACAGTAGTCTGCACATTGGTTA
Plate 1	F21	hsa-miR-199a-5p	CCCAGTGTTCCAGACTACCTGTTT
Plate 1	F22	hsa-miR-199b-3p	ACAGTAGTCTGCACATTGGTTA
Plate 1	F23	hsa-miR-199b-5p	CCCAGTGTTCAGACTATCTGTTT
Plate 1	F24	hsa-miR-200a	TAACACTGTCTGGTAACGATGT
Plate 1	G1	hsa-miR-200b	TAATACTGCCTGGTAATGATGA
Plate 1	G2	hsa-miR-200c	TAATACTGCCGGTAATGATGGA
Plate 1	G3	hsa-miR-202	AGAGGTATAGGGCATGGGAA
Plate 1	G4	hsa-miR-203	GTGAAATGTTTAGGACCACTAG
Plate 1	G5	hsa-miR-204	TTCCCTTTGTCATCCTATGCCT
Plate 1	G6	hsa-miR-205	TCCTTCATTCCACCGAGTCTG
Plate 1	G7	hsa-miR-206	TGGAATGTAAGGAAGTGTGTGG
Plate 1	G8	hsa-miR-208a	ATAAGACGAGCAAAAAGCTTGT
Plate 1	G9	hsa-miR-208b	ATAAGACGAACAAAAGTTTTGT
Plate 1	G10	hsa-miR-210	CTGTGCGTGTGACAGCGGCTGA
Plate 1	G11	hsa-miR-211	TTCCCTTTGTCATCCTTCGCCT
Plate 1	G12	hsa-miR-212	TAACAGTCTCCAGTCACGGCC
Plate 1	G13	hsa-miR-214	ACAGCAGGCACAGACAGGCAGT
Plate 1	G14	hsa-miR-215	ATGACCTATGAATTGACAGAC
Plate 1	G15	hsa-miR-216a	TAATCTCAGCTGGCAACTGTGA
Plate 1	G16	hsa-miR-216b	AAATCTCTGCAGGCAAATGTGA
Plate 1	G17	hsa-miR-217	TACTGCATCAGGAACTGATTGGA
Plate 1	G18	hsa-miR-218	TTGTGCTTGATCTAACCATGT
Plate 1	G19	hsa-miR-219-1-3p	AGAGTTGAGTCTGGACGTCCCG
Plate 1	G20	hsa-miR-219-2-3p	AGAATTGTGGCTGGACATCTGT
Plate 1	G21	hsa-miR-219-5p	TGATTGTCCAACGCAATTCT
Plate 1	G22	hsa-miR-220a	CCACACCGTATCTGACACTTT
Plate 1	G23	hsa-miR-220b	CCACCACCGTGTCTGACACTT
Plate 1	G24	hsa-miR-220c	ACACAGGGCTGTTGTGAAGACT
Plate 1	H1	hsa-miR-221	AGCTACATTGTCTGCTGGGTTTC
Plate 1	H2	hsa-miR-222	AGCTACATCTGGCTACTGGGT
Plate 1	H3	hsa-miR-223	TGTCAGTTTGTCAAATACCCCA
Plate 1	H4	hsa-miR-224	CAAGTCACTAGTGGTTCCGTT
Plate 1	H5	hsa-miR-296-3p	GAGGGTTGGGTGGAGGCTCTCC
Plate 1	H6	hsa-miR-296-5p	AGGGCCCCCCTCAATCCTGT
Plate 1	H7	hsa-miR-297	ATGTATGTGTGCATGTGCATG
Plate 1	H8	hsa-miR-298	AGCAGAAGCAGGGAGGTTCTCCCA
Plate 1	H9	hsa-miR-299-3p	TATGTGGGATGGTAAACCGCTT
Plate 1	H10	hsa-miR-299-5p	TGGTTTACCGTCCCACATACAT
Plate 1	H11	hsa-miR-300	TATACAAGGGCAGACTCTCTCT
Plate 1	H12	hsa-miR-301a	CAGTGCAATAGTATTGTCAAAGC
Plate 1	H13	hsa-miR-301b	CAGTGCAATGATATTGTCAAAGC
Plate 1	H14	hsa-miR-302a	TAAGTGCTTCCATGTTTTGGTGA
Plate 1	H15	hsa-miR-302b	TAAGTGCTTCCATGTTTTAGTAG
Plate 1	H16	hsa-miR-302c	TAAGTGCTTCCATGTTTCAGTGG
Plate 1	H17	hsa-miR-302d	TAAGTGCTTCCATGTTTGAGTGT
Plate 1	H18	hsa-miR-302e	TAAGTGCTTCCATGCCTT
Plate 1	H19	hsa-miR-302f	TAATTGCTTCCATGTTT
Plate 1	H20	hsa-miR-320a	AAAAGCTGGGTTGAGAGGGCGA
Plate 1	H21	hsa-miR-320b	AAAAGCTGGGTTGAGAGGGCAA
Plate 1	H22	hsa-miR-320c	AAAAGCTGGGTTGAGAGGGT
Plate 1	H23	hsa-miR-320d	AAAAGCTGGGTTGAGAGGA
Plate 1	H24	hsa-miR-323-3p	CACATTACACGGTCGACCTCT
Plate 1	I1	hsa-miR-323-5p	AGGTGGTCCGTGGCGGTTTCGC
Plate 1	I2	hsa-miR-324-3p	ACTGCCCCAGGTGCTGCTGG

Plate 1	I3	hsa-miR-324-5p	CGCATCCCCTAGGGCATTGGTGT
Plate 1	I4	hsa-miR-325	CCTAGTAGGTGTCCAGTAAGTGT
Plate 1	I5	hsa-miR-326	CCTCTGGGCCCTTCCTCCAG
Plate 1	I6	hsa-miR-328	CTGGCCCTCTCTGCCCTCCGT
Plate 1	I7	hsa-miR-329	AACACACCTGGTTAACCTCTTT
Plate 1	I8	hsa-miR-330-3p	GCAAAGCACACGGCCTGCAGAGA
Plate 1	I9	hsa-miR-330-5p	TCTCTGGGCCTGTGTCTTAGGC
Plate 1	I10	hsa-miR-331-3p	GCCCCTGGGCCTATCCTAGAA
Plate 1	I11	hsa-miR-331-5p	CTAGGTATGGTCCCAGGGATCC
Plate 1	I12	hsa-miR-335	TCAAGAGCAATAACGAAAAATGT
Plate 1	I13	hsa-miR-337-3p	CTCCTATATGATGCCTTTCTTC
Plate 1	I14	hsa-miR-337-5p	GAACGGCTTCATACAGGAGTT
Plate 1	I15	hsa-miR-338-3p	TCCAGCATCAGTGATTTTGTTG
Plate 1	I16	hsa-miR-338-5p	ACAATATCCTGGTGTGAGTG
Plate 1	I17	hsa-miR-339-3p	TGAGCGCTCGACGACAGAGCCG
Plate 1	I18	hsa-miR-339-5p	TCCCTGTCTCCAGGAGCTCACG
Plate 1	I19	hsa-miR-340	TTATAAGCAATGAGACTGATT
Plate 1	I20	hsa-miR-342-3p	TCTCACACAGAAATCGACCCGT
Plate 1	I21	hsa-miR-342-5p	AGGGGTGTATCTGTGATTGA
Plate 1	I22	hsa-miR-345	GCTGACTCCTAGTCCAGGGCTC
Plate 1	I23	hsa-miR-346	TGTCTGCCCGCATGCCTGCCTCT
Plate 1	I24	hsa-miR-361-3p	TCCCCAGGTGTGATTCTGATTT
Plate 1	J1	hsa-miR-361-5p	TTATCAGAACTCCAGGGGTAC
Plate 1	J2	hsa-miR-362-3p	AACACACCTATTCAAGGATTCA
Plate 1	J3	hsa-miR-362-5p	AATCCTTGGAACCTAGGTGTGAGT
Plate 1	J4	hsa-miR-363	AATTGCACGGTATCCATCTGTA
Plate 1	J5	hsa-miR-365	TAATGCCCTAAAAATCCTTAT
Plate 1	J6	hsa-miR-367	AATTGCACTTTAGCAATGGTGA
Plate 1	J7	hsa-miR-369-3p	ATAATACATGGTTGATCTTT
Plate 1	J8	hsa-miR-369-5p	AGATCGACCGTGTATATTCGC
Plate 1	J9	hsa-miR-370	GCCTGCTGGGGTGAACCTGGT
Plate 1	J10	hsa-miR-371-3p	AAGTGCCGCCATCTTTTGAGTGT
Plate 1	J11	hsa-miR-371-5p	ACTCAAAGTGGGGGCACT
Plate 1	J12	hsa-miR-372	AAAGTGCTGCGACATTTGAGCGT
Plate 1	J13	hsa-miR-373	GAAGTGCTTCGATTTTGGGGTGT
Plate 1	J14	hsa-miR-374a	TTATAATACAACCTGATAAGTG
Plate 1	J15	hsa-miR-374b	ATATAATACAACCTGCTAAGTG
Plate 1	J16	hsa-miR-375	TTTGTTTCGTTTCGGCTCGCGTGA
Plate 1	J17	hsa-miR-376a	ATCATAGAGGAAAAATCCACGT
Plate 1	J18	hsa-miR-376b	ATCATAGAGGAAAAATCCATGTT
Plate 1	J19	hsa-miR-376c	AACATAGAGGAAATCCACGT
Plate 1	J20	hsa-miR-377	ATCACACAAAGGCAACTTTTGT
Plate 1	J21	hsa-miR-378	ACTGGACTTGGAGTCAGAAGG
Plate 1	J22	hsa-miR-379	TGGTAGACTATGGAACGTAGG
Plate 1	J23	hsa-miR-380	TATGTAATATGGTCCACATCTT
Plate 1	J24	hsa-miR-381	TATACAAGGGCAAGCTCTCTGT
Plate 1	K1	hsa-miR-382	GAAGTTGTTTCGTGGTGGATTCCG
Plate 1	K2	hsa-miR-383	AGATCAGAAGGTGATTGTGGCT
Plate 1	K3	hsa-miR-384	ATTCCTAGAAATTGTTTCATA
Plate 1	K4	hsa-miR-409-3p	GAATGTTGCTCGGTGAACCCCT
Plate 1	K5	hsa-miR-409-5p	AGGTTACCCGAGCAACTTTGCAT
Plate 1	K6	hsa-miR-410	AATATAACACAGATGGCCTGT
Plate 1	K7	hsa-miR-411	TAGTAGACCGTATAGCGTACG
Plate 1	K8	hsa-miR-412	ACTTACCTGGTCCACTAGCCGT
Plate 1	K9	hsa-miR-421	ATCAACAGACATTAATTGGGCGC
Plate 1	K10	hsa-miR-422a	ACTGGACTTAGGGTCAGAAGGC
Plate 1	K11	hsa-miR-423-3p	AGCTCGGTCTGAGGCCCTCAGT
Plate 1	K12	hsa-miR-423-5p	TGAGGGGCAGAGAGCGAGACTTT
Plate 1	K13	hsa-miR-424	CAGCAGCAATTCATGTTTTGAA
Plate 1	K14	hsa-miR-425	AATGACACGATCACTCCGTTGA
Plate 1	K15	hsa-miR-429	TAATACTGTCTGGTAAAACCGT
Plate 1	K16	hsa-miR-431	TGTCTTGACAGGCCGTCATGCA
Plate 1	K17	hsa-miR-432	TCTTGGAGTAGGTTCATTGGGTGG
Plate 1	K18	hsa-miR-433	ATCATGATGGGCTCCTCGGTGT
Plate 1	K19	hsa-miR-448	TTGCATATGTAGGATGTCCCAT
Plate 1	K20	hsa-miR-449a	TGGCAGTGTATTGTTAGCTGGT

Plate 1	K21	hsa-miR-449b	AGGCAGTGATTGTTAGCTGGC
Plate 1	K22	hsa-miR-450a	TTTTGCGATGTGTTCTTAATAT
Plate 1	K23	hsa-miR-450b-3p	TTGGGATCATTTTGCATCCATA
Plate 1	K24	hsa-miR-450b-5p	TTTTGCAATATGTTCTGAATA
Plate 1	L1	hsa-miR-451	AAACCGTTACCATTACTGAGTT
Plate 1	L2	hsa-miR-452	AACTGTTTGCAGAGGAAACTGA
Plate 1	L3	hsa-miR-453	AGGTTGTCCGTGGTGAGTTCGCA
Plate 1	L4	hsa-miR-454	TAGTGCAATATTGCTTATAGGGT
Plate 1	L5	hsa-miR-455-3p	GCAGTCCATGGGCATATACAC
Plate 1	L6	hsa-miR-455-5p	TATGTGCCTTTGGACTACATCG
Plate 1	L7	hsa-miR-483-3p	TCACTCTCTCCTCCCGTCTT
Plate 1	L8	hsa-miR-483-5p	AAGACGGGAGGAAAGAAGGGAG
Plate 1	L9	hsa-miR-484	TCAGGCTCAGTCCCCTCCGAT
Plate 1	L10	hsa-miR-485-3p	GTCATACACGGCTCTCTCTCT
Plate 1	L11	hsa-miR-485-5p	AGAGGCTGGCCGTGATGAATTC
Plate 1	L12	hsa-miR-486-3p	CGGGGCAGCTCAGTACAGGAT
Plate 1	L13	hsa-miR-486-5p	TCCTGTACTGAGCTGCCCGAG
Plate 1	L14	hsa-miR-487a	AATCATACAGGGACATCCAGTT
Plate 1	L15	hsa-miR-487b	AATCGTACAGGGTCATCCACTT
Plate 1	L16	hsa-miR-488	TTGAAAGGCTATTTCTTGGTC
Plate 1	L17	hsa-miR-489	GTGACATCACATATACGGCAGC
Plate 1	L18	hsa-miR-490-3p	CAACCTGGAGGACTCCATGCTG
Plate 1	L19	hsa-miR-490-5p	CCATGGATCTCCAGGTGGGT
Plate 1	L20	hsa-miR-491-3p	CTTATGCAAGATTCCTTCTAC
Plate 1	L21	hsa-miR-491-5p	AGTGGGGAACCCCTCCATGAGG
Plate 1	L22	hsa-miR-492	AGGACCTGCGGGACAAGATTCTT
Plate 1	L23	hsa-miR-493	TGAAGGTCTACTGTGTCCAGG
Plate 1	L24	hsa-miR-494	TGAAACATACACGGGAAACCTC
Plate 1	M1	hsa-miR-495	AAACAACATGGTGCCTTCTT
Plate 1	M2	hsa-miR-496	TGAGTATTACATGGCCAATCTC
Plate 1	M3	hsa-miR-497	CAGCAGCACACTGTGGTTTGT
Plate 1	M4	hsa-miR-498	TTTCAAGCCAGGGGGCGTTTTTC
Plate 1	M5	hsa-miR-499-3p	AACATCACAGCAAGTCTGTGCT
Plate 1	M6	hsa-miR-499-5p	TTAAGACTTGCAAGTATGTTT
Plate 1	M7	hsa-miR-500	TAATCCTTGCTACCTGGGTGAGA
Plate 1	M8	hsa-miR-501-3p	AATGCACCCGGGCAAGGATTCT
Plate 1	M9	hsa-miR-501-5p	AATCCTTTGTCCCTGGGTGAGA
Plate 1	M10	hsa-miR-502-3p	AATGCACCTGGGCAAGGATTCA
Plate 1	M11	hsa-miR-502-5p	ATCCTTGCTATCTGGGTGCTA
Plate 1	M12	hsa-miR-503	TAGCAGCGGGAACAGTTCTGCAG
Plate 1	M13	hsa-miR-504	AGACCCTGGTCTGCACTCTATC
Plate 1	M14	hsa-miR-505	CGTCAACACTTGCTGGTTTCCT
Plate 1	M15	hsa-miR-506	TAAGGCACCCTTCTGAGTAGA
Plate 1	M16	hsa-miR-507	TTTTGCACCTTTTGGAGTGAA
Plate 1	M17	hsa-miR-508-3p	TGATTGTAGCCTTTTGGAGTAGA
Plate 1	M18	hsa-miR-508-5p	TACTCCAGAGGGCGTCACTCATG
Plate 1	M19	hsa-miR-509-3-5p	TACTGCAGACGTGGCAATCATG
Plate 1	M20	hsa-miR-509-3p	TGATTGGTACGTCTGTGGGTAG
Plate 1	M21	hsa-miR-509-5p	TACTGCAGACAGTGGCAATCA
Plate 1	M22	hsa-miR-510	TACTCAGGAGAGTGGCAATCAC
Plate 1	M23	hsa-miR-511	GTGTCTTTTGCTCTGCAGTCA
Plate 1	M24	hsa-miR-512-3p	AAGTGCTGTATAGCTGAGGTC
Plate 1	N1	hsa-miR-512-5p	CACTCAGCCTTGAGGGCACTTTC
Plate 1	N2	hsa-miR-513a-3p	TAAATTTACCTTTCTGAGAAGG
Plate 1	N3	hsa-miR-513a-5p	TTCAAGGGAGGTGTCAT
Plate 1	N4	hsa-miR-513b	TTCAAGGAGGTGTCATTTAT
Plate 1	N5	hsa-miR-513c	TTCTCAAGGAGGTGTCGTTTAT
Plate 1	N6	hsa-miR-514	ATTGACACTTCTGTGAGTAGA
Plate 1	N7	hsa-miR-515-3p	GAGTGCCTTCTTTTGGAGCGTT
Plate 1	N8	hsa-miR-515-5p	TTCTCCAAAAGAAAGCACTTCTG
Plate 1	N9	hsa-miR-516a-3p	TGCTTCCTTTCAGAGGGT
Plate 1	N10	hsa-miR-516a-5p	TTCTCGAGGAAAGAAGCACTTTC
Plate 1	N11	hsa-miR-516b	ATCTGGAGGTAAGAAGCACTTT
Plate 1	N12	hsa-miR-517a	ATCGTGCATCCCTTTAGAGTGT
Plate 1	N13	hsa-miR-517b	TCGTGCATCCCTTTAGAGTGT
Plate 1	N14	hsa-miR-517c	ATCGTGCATCCCTTTAGAGTGT

Plate 1	N15	hsa-miR-518a-3p	GAAAGCGCTTCCCTTTGCTGGA
Plate 1	N16	hsa-miR-518a-5p	CTGCAAAGGGAAGCCCTTTC
Plate 1	N17	hsa-miR-518b	CAAAGCGCTCCCTTTAGAGGT
Plate 1	N18	hsa-miR-518c	CAAAGCGCTTCTCTTTAGAGTGT
Plate 1	N19	hsa-miR-518d-3p	CAAAGCGCTTCCCTTTGGAGC
Plate 1	N20	hsa-miR-518d-5p	CTCTAGAGGGAAGCACTTTCTG
Plate 1	N21	hsa-miR-518e	AAAGCGCTTCCCTCAGAGTG
Plate 1	N22	hsa-miR-518f	GAAAGCGCTTCTCTTTAGAGG
Plate 1	N23	hsa-miR-519a	AAAGTGCATCCTTTTAGAGTGT
Plate 1	N24	hsa-miR-519b-3p	AAAGTGCATCCTTTTAGAGGTT
Plate 1	O1	hsa-miR-519b-5p	CTCTAGAGGGAAGCGCTTCTG
Plate 1	O2	hsa-miR-519c-3p	AAAGTGCATCTTTTAGAGGAT
Plate 1	O3	hsa-miR-519c-5p	CTCTAGAGGGAAGCGCTTCTG
Plate 1	O4	hsa-miR-519d	CAAAGTGCCTCCCTTTAGAGTG
Plate 1	O5	hsa-miR-519e	AAGTGCCTCCTTTTAGAGTGT
Plate 1	O6	hsa-miR-520a-3p	AAAGTGCCTCCCTTTGGACTGT
Plate 1	O7	hsa-miR-520a-5p	CTCCAGAGGGAAGTACTTTCT
Plate 1	O8	hsa-miR-520b	AAAGTGCCTTCTTTTAGAGGG
Plate 1	O9	hsa-miR-520c-3p	AAAGTGCCTTCTTTTAGAGGGT
Plate 1	O10	hsa-miR-520c-5p	CTCTAGAGGGAAGCACTTTCTG
Plate 1	O11	hsa-miR-520d-3p	AAAGTGCCTTCTCTTTGGTGGGT
Plate 1	O12	hsa-miR-520d-5p	CTACAAAGGGAAGCCCTTTC
Plate 1	O13	hsa-miR-520e	AAAGTGCCTTCTTTTAGAGGG
Plate 1	O14	hsa-miR-520f	AAGTGCCTTCTTTTAGAGGGT
Plate 1	O15	hsa-miR-520g	ACAAAGTGCCTCCCTTTAGAGTGT
Plate 1	O16	hsa-miR-520h	ACAAAGTGCCTCCCTTTAGAGT
Plate 1	O17	hsa-miR-521	AACGCCTTCCCTTTAGAGTGT
Plate 1	O18	hsa-miR-522	AAAATGTTCCCTTTAGAGTGT
Plate 1	O19	hsa-miR-523	GAACGCGCTTCCCTATAGAGGGT
Plate 1	O20	hsa-miR-524-3p	GAAGGCGCTTCCCTTTGGAGT
Plate 1	O21	hsa-miR-524-5p	CTACAAAGGGAAGCACTTCTC
Plate 1	O22	hsa-miR-525-3p	GAAGGCGCTTCCCTTTAGAGCG
Plate 1	O23	hsa-miR-525-5p	CTCCAGAGGGATGCACCTTCT
Plate 1	O24	hsa-miR-526a	CTCTAGAGGGAAGCACTTCTG
Plate 1	P1	hsa-miR-526b	CTCTTGAGGGAAGCACTTCTGT
Plate 1	P2	hsa-miR-527	CTGCAAAGGGAAGCCCTTTC
Plate 1	P3	hsa-miR-532-3p	CCTCCACACCCAAGGCTTGCA
Plate 1	P4	hsa-miR-532-5p	CATGCCTTGAGGTAGGACCGT
Plate 1	P5	hsa-miR-539	GGAGAAATTATCCTTGGTGTGT
Plate 1	P6	hsa-miR-541	TGGTGGGCACAGAATCTGGACT
Plate 1	P7	hsa-miR-542-3p	TGTGACAGATTGATAACTGAAA
Plate 1	P8	hsa-miR-542-5p	TCGGGGATCATCATGTCACGAGA
Plate 1	P9	hsa-miR-543	AAACATTCGCGGTGCACTTCTT
Plate 1	P10	hsa-miR-544	ATTCTGCATTTTATGCAAGTTC
Plate 1	P11	hsa-miR-545	TCAGCAAACATTTATTGTGTGC
Plate 1	P12	hsa-miR-548a-3p	CAAACTGGCAATTACTTTTGC
Plate 1	P13	hsa-miR-548a-5p	AAAAGTAATTGCGAGTTTACC
Plate 1	P14	hsa-miR-548b-3p	CAAGAACCTCAGTTGCTTTTGT
Plate 1	P15	hsa-miR-548b-5p	AAAAGTAATTGTGGTTTTGGCC
Plate 1	P16	hsa-miR-548c-3p	CAAAAATCTCAATTACTTTTGC
Plate 1	P17	hsa-miR-548c-5p	AAAAGTAATTGCGGTTTTTGC
Plate 1	P18	hsa-miR-548d-3p	CAAAAACCACAGTTTCTTTTGC
Plate 1	P19	hsa-miR-548d-5p	AAAAGTAATTGTGGTTTTTGC
Plate 1	P20	hsa-miR-548e	AAAAACTGAGACTACTTTTGC
Plate 1	P21	Human U6 snRNA	CGCAAGGATGACACGCAAATTC
Plate 1	P22	RNU43 snoRNA	CTTATTGACGGGCGGACAGAAAC
Plate 1	P23	Hm/Ms/Rt U1 snRNA	CGACTGCATAATTTGTGGTAGTGG
Plate 1	P24	empty	empty