

Table S2

miRNA	Sequence	4 species	rhesus	chimpanzee
hsa-miR-638	AGGGAUCGCGGGCGGUGGCGGCCU	NC	C	NC
hsa-miR-302b	UAAGUGCUUCCAUGUUUAGUAG	C	C	C
hsa-miR-302d	UAAGUGCUUCCAUGUUUGAGUGU	C	C	C
hsa-miR-302a	UAAGUGCUUCCAUGUUUUGGUGA	NC	C	C
hsa-miR-302a*	UAAACGUGGAUGUACUUGCUUU	C	C	C
hsa-miR-20a	UAAAGUGCUUAUAGUGCAGGUAG	C	NC	C
hsa-miR-17-5p	CAAAGUGCUUACAGUGCAGGUAGU	C	C	C
hsa-miR-302c	UAAGUGCUUCCAUGUUUCAGUGG	C	C	C
hsa-miR-663	AGGCGGGGCGCCGCGGGACCGC	NC	NC	C
hsa-miR-20b	CAAAGUGCUCAUAGUGCAGGUAG	C	C	C
hsa-miR-106a	AAAAGUGCUUACAGUGCAGGUAGC	C	C	C
hsa-miR-25	CAUUGCACUUGUCUCGGUCUGA	C	C	NC
hsa-miR-92	UAUUGCACUUGUCCCGGCCUG	C	C	C
hsa-miR-21	UAGCUUAUCAGACUGAUGUUGA	C	C	C
hsa-miR-92b	UAUUGCACUCGUCCCGGCCUC	C	C	NC
hsa-miR-26a	UUCAGUAAUCCAGGAUAGGC	C	C	C
hsa-miR-15b	UAGCAGCACAUCAUGGUUUACA	C	C	C
hsa-miR-513	UUCACAGGGAGGUGUCAUUUAU	NC	C	C
hsa-miR-19b	UGUGCAAAUCCAUGCAAAACUGA	C	C	C
hsa-miR-130a	CAGUGCAAUGUUAAAAGGGCAU	C	C	C
hsa-miR-200c	UAAUACUGCCGGGUAAUGAUGG	C	C	NC
hsa-miR-93	AAAGUGCUGUUCGUGCAGGUAG	C	C	C
hsa-miR-612	GCUGGGCAGGGCUUCUGAGCUCCUU	NC	NC	C
hsa-miR-16	UAGCAGCACGUAAAUUUGGCG	C	C	C
hsa-miR-103	AGCAGCAUUGUACAGGGCUAUGA	C	C	C
hsa-miR-107	AGCAGCAUUGUACAGGGCUAUC	C	C	C
hsa-miR-221	AGCUACAUGUCUGCUGGGUUUC	C	C	C
hsa-miR-363	AAUUGCACGGUAUCCAUCUGUA	C	C	NC
hsa-miR-575	GAGCCAGUUGGACAGGAGC	NC	NC	C
hsa-miR-373	GAAGUGCUUCGAUUUUGGGGUGU	NC	C	C
hsa-miR-342	UCUCACACAGAAUCGCACCCGUC	C	C	C
hsa-miR-191	CAACGGAAUCCCAAAGCAGCU	C	C	C
hsa-miR-302c*	UUUAACAUGGGGUACCUGCUG	C	C	C
hsa-miR-130b	CAGUGCAAUGAUGAAAGGGCAU	C	C	C
hsa-miR-494	UGAAACAUAACACGGGAAACCUCUU	C	C	C
hsa-miR-106b	UAAAGUGCUGACAGUGCAGAU	C	C	C
hsa-miR-320	AAAAGCUGGGUUGAGAGGGCGAA	C	C	C
hsa-miR-30c	UGUAAACAUCUACACUCUCAGC	C	C	C
hsa-miR-594	CCCAUCUGGGGUGGCCUGUGACUUU	NC	NC	C
hsa-miR-200b	UAAUACUGCCUGGUAAUGAUGAC	C	NC	NC
hsa-miR-26b	UUCAAGUAAUUCAGGAUAGGUU	C	C	C
hsa-miR-182	UUUGGCAAUGGUAGAACUCACA	C	C	C
hsa-miR-374	UUAUAAUACAACCUGAUAGUG	C	C	C
hsa-miR-30b	UGUAAACAUCUACACUCAGCU	C	C	C
hsa-miR-222	AGCUACAUCUGGCUACUGGGUCUC	C	C	NC

hsa-miR-148a	UCAGUGCACUACAGAACUUUGU	C	C	C
hsa-miR-361	UUAUCAGAAUCUCCAGGGGUAC	C	C	C
hsa-miR-367	AAUUGCACUUUAGCAAUGGUGA	C	C	C
hsa-miR-18a	UAAGGUGCAUCUAGUGCAGAU	C	C	C
hsa-miR-517a	AUCGUGCAUCCCUUAGAGUGUU	NC	C	C
hsa-miR-195	UAGCAGCACAGAAAUUUGGC	C	C	C
hsa-miR-637	ACUGGGGGCUUUCGGGCUCUGCGU	NC	NC	C
hsa-miR-371	GUGCCGCCAUCUUUUGAGUGU	NC	C	C
hsa-miR-183	UAUGGCACUGGUAGAAUUCACUG	C	C	C
hsa-miR-421	AUCAACAGACAUUAAUUGGGCGC	C	C	NC
hsa-miR-517b	UCGUGCAUCCCUUAGAGUGUU	NC	C	C
hsa-miR-23b	AUCACAUUGCCAGGGAUUACC	C	C	C
hsa-miR-498	UUUCAAGCCAGGGGCGUUUUUC	NC	C	C
hsa-miR-526a	CUCUAGAGGGAAGCACUUUCU	NC	C	C
hsa-miR-23a	AUCACAUUGCCAGGGAUUUC	C	C	C
hsa-miR-372	AAAGUGCUGCGACAUUUGAGCGU	NC	C	C
hsa-miR-602	GACACGGGCGACAGCUGCGGCC	NC	NC	NC
hsa-miR-18b	UAAGGUGCAUCUAGUGCAGUUA	C	C	C
hsa-miR-517*	CCUCUAGAUGGAAGCACUGUCU	NC	NC	NC
hsa-miR-302b*	ACUUUAACAUGGAAGUGCUUUCU	NC	C	C
hsa-miR-335	UCAAGAGCAAUAACGAAAAUGU	C	C	C
hsa-miR-129	CUUUUUGCGGUCUGGGCUUGC	C	C	C
hsa-miR-19a	UGUGCAAAUCUAUGCAAAACUGA	C	NC	C
hsa-miR-518c*	CAAAGCGCUUCUCUUUAGAGUG	NC	C	C
hsa-miR-524*	CUACAAAGGAAGCACUUUCUC	NC	C	C
hsa-miR-516-5p	CAUCUGGAGGUAAGAAGCACUUU	NC	NC	C
hsa-miR-185	UGGAGAGAAAGGCAGUUC	C	C	C
hsa-miR-658	GGCGGAGGGAAGUAGGUCGUGGU	NC	NC	C
hsa-miR-425-5p	AAUGACACGAUCACUCCGUUGA	C	C	C
hsa-miR-125a	UCCUGAGACCCUUUAACUGUG	C	C	C
hsa-miR-128b	UCACAGUGAACCGGUCUUUUC	C	C	C
hsa-miR-128a	UCACAGUGAACCGGUCUUUUU	C	C	C
hsa-miR-29a	UAGCACCAUCUGAAUUCGGUU	C	C	NC
hsa-miR-198	GGUCCAGAGGGGAGAUAGG	NC	C	C
hsa-miR-422b	CUGGACUUGGAGUCAGAAGGCC	NC	C	C
hsa-miR-151	ACUAGACUGAAGCUCCUUGAGG	C	C	C
hsa-miR-148b	UCAGUGCAUCACAGAACUUUGU	C	C	C
hsa-miR-28	AAGGAGCUCACAGUCUAUUGAG	C	C	C
hsa-miR-518f*	CUCUAGAGGGAAGCACUUUCUCU	NC	C	C
hsa-miR-30d	UGUAAACAUCCCCGACUGGAAG	C	C	C
hsa-miR-126	UCGUACCGUGAGUAAUAUGC	C	C	C
hsa-miR-27b	UUCACAGUGGCUAAGUUCUGC	C	C	C
hsa-miR-24	UGGCUAGUUCAGCAGGAACAG	C	C	C
hsa-miR-187	UCGUGUCUUGUGUUGCAGCCG	C	C	NC
hsa-miR-512-3p	AAGUGCUGUCAUAGCUGAGGUC	NC	NC	C
hsa-miR-492	AGGACCUGCGGGACAAGAUUCUU	NC	C	C
hsa-miR-526b	CUCUUGAGGGAAGCACUUUCUGUU	NC	C	C
hsa-miR-650	AGGAGGCAGCGCUCUCAGGAC	NC	C	C

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hsa-miR-101	UACAGUACUGUGAUAAACUGAAG	C	C	C
hsa-miR-564	AGGCACGGUGUCAGCAGGC	NC	NC	C
hsa-miR-574	CACGCUCAUGCACACACCCAC	C	NC	NC
hsa-miR-520h	ACAAAGUGCUUCCCUUAGAGU	NC	C	C
hsa-miR-520g	ACAAAGUGCUUCCCUUAGAGUGU	NC	C	C
hsa-miR-30a-5p	UGUAAACAUCUCGACUGGAAG	C	C	C
hsa-miR-193b	AACUGGCCCUCAAAGUCCCGCUUU	C	C	NC
hsa-miR-301	CAGUGCAAUAGUAUUGUCAAGC	C	C	C
hsa-miR-565	GGCUGGCUCGCGAUGUCUGUUU	NC	NC	NC
hsa-miR-7	UGGAAGACUAGUGAUUUUGUUG	C	C	C
hsa-miR-135a	UAUGGCUUUUAUUCUAUGUGA	C	C	C
hsa-miR-422a	CUGGACUUAGGGUCAGAAGGCC	NC	NC	C
hsa-miR-99b	CACCCGUAGAACCGACCUUGCG	C	C	C
hsa-miR-206	UGGAAUGUAAGGAAGUGUGUGG	C	C	C
hsa-miR-365	UAAUGCCCCUAAAAUCCUUAU	C	C	C
hsa-miR-572	GUCCGCUCGGCGGUGGCCCA	NC	C	C
hsa-miR-505	GUCAACACUUGCUGGUUCCUC	C	C	C
hsa-miR-146b	UGAGAACUGAAUCCAUAGGCU	C	NC	C
hsa-miR-517c	AUCGUGCAUCCUUUAGAGUGU	NC	C	C
hsa-miR-124a	UUAAGGCACGCGUGAAUGCCA	C	C	C
hsa-miR-27a	UUCACAGUGGCUAAGUUCGCG	C	C	C
hsa-miR-519d	CAAAGUGCCUCCCUUAGAGUGU	NC	C	C
hsa-miR-324-3p	CCACUGCCCCAGGUGCUGCUGG	C	C	C
hsa-miR-210	CUGUGCGUGUGACAGCGGUGA	C	C	C
hsa-miR-373*	ACUCAAAUGGGGGCGCUUCC	NC	NC	C
hsa-miR-497	CAGCAGCACACUGUGGUUGU	C	C	C
hsa-miR-636	UGUGCUUGCUCGUCCCGCCGCGAG	NC	C	C
hsa-miR-484	UCAGGCUCAGUCCCCUCCGAU	C	C	C
hsa-miR-205	UCCUUCAUUCCACCGGAGUCUG	C	C	C
hsa-miR-296	AGGGCCCCCCCUCAAUCCUGU	C	C	C
hsa-miR-186	CAAAGAAUUCUCCUUUUGGCUU	C	C	C
hsa-miR-519c	AAAGUGCAUCUUUUUAGAGGAU	NC	C	C
hsa-miR-30e-3p	CUUUCAGUCGGAUGUUUACAGC	C	C	C
hsa-miR-15a	UAGCAGCACAUAAUGGUUUGUG	C	C	C
hsa-miR-214	ACAGCAGGCACAGACAGGCAG	C	C	C
hsa-miR-192	CUGACCUAUGAAUUGACAGCC	C	C	C
hsa-miR-331	GCCCCUGGGCCUAUCCUAGAA	C	C	C
hsa-miR-223	UGUCAGUUUGUCAAUACCCC	C	C	C
hsa-miR-34a	UGGCAGUGUCUUAGCUGGUUGUU	C	C	C
hsa-miR-197	UUCACCACCUUCUCCACCCAGC	C	C	C
hsa-miR-520d	AAAGUGCUUCUUCUUUGGUGGUU	NC	C	C
hsa-miR-601	UGGUCUAGGAUUGUUGGAGGAG	NC	C	C
hsa-miR-598	UACGUCAUCGUUGUCAUCGUCA	NC	NC	C
hsa-miR-191*	GCUGCGCUUGGAUUUCGUCCCC	C	C	C
hsa-miR-515-5p	UUCUCCAAAAGAAAGCACUUUCUG	NC	NC	C
hsa-miR-557	GUUUGCACGGGUGGGCCUUGUCU	NC	NC	C
hsa-miR-520c	AAAGUGCUUCUUUUAGAGGUU	NC	C	C
hsa-miR-149	UCUGGCUCCGUGUCUUCACUCC	C	C	C

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hsa-miR-489	AGUGACAUCACAUUACGGCAGC	NC	C	NC
hsa-miR-31	GGCAAGAUGCUGGCAUAGCUG	C	C	C
hsa-miR-525	CUCCAGAGGGAUGCACUUUCU	NC	C	C
hsa-miR-18a*	ACUGCCCUAAGUGCUCUUUCU	C	NC	C
hsa-miR-519e*	AAAGUGCCUCCUUUAGAGUGU	NC	NC	C
hsa-miR-520f	AAGUGCUUCCUUUAGAGGGU	NC	NC	C
hsa-miR-593	AGGCACCAGCCAGGCAUUGCUCAGC	NC	NC	NC
hsa-miR-125b	UCCCUAGAGCCCUAACUUGUGA	C	C	C
hsa-miR-204	UCCCCUUUGUCAUCCUAUGCCU	C	C	C
hsa-miR-623	AUCCCUUGCAGGGGUGUUGGGU	NC	C	C
hsa-miR-519b	AAAGUGCAUCCUUUAGAGGUU	NC	C	C
hsa-miR-628	UCUAGUAAGAGUGGCAGUCG	C	C	C
hsa-miR-22	AAGCUGCCAGUUGAAGAACUGU	C	NC	C
hsa-miR-194	UGUAACAGCAACUCCAUGUGGA	C	C	C
hsa-miR-199a*	UACAGUAGUCUGCACAUUGGUU	C	C	C
hsa-miR-340	UCCGUCUCAGUUACUUUAUAGCC	C	C	C
hsa-miR-520a*	AAAGUGCUUCCCUUUGGACUGU	NC	NC	NC
hsa-miR-615	UCCGAGCCUGGGUCUCCUCU	C	C	C
hsa-miR-200a	UAACACUGUCUGGUAACGAUGU	C	C	C
hsa-miR-155	UUAUUGCUAAUCGUGAUAGGGG	C	C	C
hsa-miR-660	UACCAUUGCAUAUCGGAGUUG	C	C	C
hsa-miR-181b	AACAUCUUGCUGUCGGUGGG	C	C	C
hsa-miR-518b	CAAAGCGCUCCCCUUAGAGGU	NC	C	C
hsa-miR-532	CAUGCCUUGAGUGUAGGACCGU	C	C	C
hsa-miR-30e-5p	UGUAAACAUCUUGACUGGA	C	C	C
hsa-miR-560	GCGUGCGCCGGCCGCGCC	NC	NC	C
hsa-miR-362	AAUCCUUGGAACCUAGGUGUGAGU	C	C	C
hsa-miR-425-3p	AUCGGGAUUGUCGUGUCCGCC	C	C	C
hsa-miR-9*	UAAAGCUAGAUAAACCGAAAGU	C	C	C
hsa-miR-625	AGGGGGAAAGUUCUAUAGUCCU	NC	NC	C
hsa-miR-518f	AAAGCGCUUCUCUUUAGAGGA	NC	NC	C
hsa-miR-429	UAAUACUGUCUGGUAAGACCGU	NC	C	C
hsa-miR-141	UAACACUGUCUGGUAAGAUAGG	C	C	C
hsa-miR-345	UGCUGACUCCUAGUCCAGGGC	NC	C	C
hsa-miR-424	CAGCAGCAAUUAUGUUUUGAA	C	C	C
hsa-miR-150	UCUCCCAACCCUUGUACCAGUG	C	C	C
hsa-miR-522	AAAUGGUUCCCUUAGAGUGUU	NC	C	C
hsa-miR-224	CAAGUCACUAGUGGUUCCGUUUA	C	C	C
hsa-miR-520b	AAAGUGCUUCCUUUAGAGGG	NC	C	C
hsa-miR-9	UCUUUGGUUAUCUAGCUGUAUGA	C	C	C
hsa-miR-202	AGAGGUUAUAGGGCAUGGGAAAA	NC	NC	C
hsa-miR-188	CAUCCCUUGCAUGGUGGAGGGU	C	C	C
hsa-miR-200a*	CAUCUUAACGGACAGUCUGGA	C	C	C
hsa-miR-518e	AAAGCGCUUCCCUUCAGAGUGU	NC	NC	C
hsa-miR-142-5p	CAUAAAGUAGAAAGCACUAC	C	C	C
hsa-miR-423	AGCUCGGUCUGAGGCCCCUCAG	C	C	C
hsa-miR-181a	AACAUUAACGCUGUCGGUGAGU	C	C	C
hsa-miR-526c	CUCUAGAGGGAAGCGCUUUCUGUU	NC	C	C

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hsa-miR-17-3p	ACUGCAGUGAAGGCACUUGU	C	NC	C
hsa-miR-181d	AACAUUCAUUGUUGUCGGUGGGUU	C	C	C
hsa-miR-563	AGGUUGACAUACGUUUCCC	NC	NC	C
hsa-miR-629	GUUCUCCCAACGUAAGCCCAGC	NC	NC	NC
hsa-miR-203	GUGAAAUGUUUAGGACCACUAG	C	C	C
hsa-miR-589	UCAGAACAAAUGCCGGUUCCCAGA	NC	NC	C
hsa-miR-452	UGUUUGCAGAGGAAACUGAGAC	C	C	C
hsa-miR-654	UGGUGGGCCGCAGAACAUUGUC	NC	C	C
hsa-miR-145	GUCCAGUUUUCCCAGGAAUCCCUU	C	C	C
hsa-miR-526b*	AAAGUGCUUCCUUUUAGAGGC	NC	C	C
hsa-miR-324-5p	CGCAUCCCCUAGGGCAUUGGUGU	C	C	C
hsa-miR-146a	UGAGAACUGAAUCCAUGGGUU	C	C	C
hsa-miR-652	AAUGGCGCCACUAGGGUUGUGCA	C	C	C
hsa-miR-515-3p	GAGUGCCUUCUUUUGGAGCGU	NC	NC	C
hsa-miR-519a	AAAGUGCAUCCUUUUAGAGUGUAC	NC	NC	C
hsa-miR-455	UAUGUGCCUUUGGACUACAUCG	C	C	C
hsa-miR-105	UCAA AUGCUCAGACUCCUGU	C	C	C
hsa-miR-143	UGAGAUGAAGCACUGUAGCUC A	C	C	C
hsa-miR-486	UCCUGUACUGAGCUGCCCCGAG	C	C	C
hsa-miR-542-5p	UCGGGGAUCAUCAUGUCACGAG	C	C	NC
hsa-miR-363*	CGGGUGGAUCACGAUGCAAUUU	C	C	NC
hsa-miR-132	UACAGUCUACAGCCAUGGUCG	C	C	NC
hsa-miR-152	UCAGUGCAUGACAGAACUUGGG	C	C	C
hsa-miR-500	AUGCACCUGGGCAAGGAUUCUG	NC	C	C
hsa-miR-122a	UGGAGUGUGACAAUGGUGUUUGU	C	C	C
hsa-miR-659	CUUGGUUCAGGGAGGGUCCCCA	NC	NC	C
hsa-miR-451	AAACCGUUACCAUUACUGAGUUU	C	C	C
hsa-miR-30a-3p	CUUUCAGUCGGAUGUUUGCAGC	C	C	C
hsa-miR-590	GAGCUUAUUCAUAAAAGUGCAG	NC	C	C
hsa-miR-346	UGUCUGCCCGCAUGCCUGCCUCU	C	C	C
hsa-miR-512-5p	CACUCAGCCUUGAGGGCACUUUC	NC	NC	C
hsa-miR-641	AAAGACAUAGGAUAGAGUACCUC	NC	NC	C
hsa-miR-330	GCAAAGCACACGGCCUGCAGAGA	C	C	C
hsa-miR-502	AUCCUUGCUAUCUGGGUGCUA	C	C	C
hsa-miR-96	UUUGGCACUAGCACAUUUUUGC	C	C	C
hsa-miR-634	AACCAGCACCCCAACUUUGGAC	NC	NC	C
hsa-miR-215	AUGACCUAUGAAUUGACAGAC	C	C	C
hsa-miR-133b	UUGGUCCCCUUAACCAGCUA	C	C	C
hsa-miR-501	AAUCCUUUGUCCCUUGGGUGAGA	NC	C	C
hsa-miR-550	UGUCUUACUCCUCAGGCACAU	NC	C	C
hsa-miR-519e	AAAGUGCCUCCUUUUAGAGUGU	NC	NC	C
hsa-miR-218	UUGUGCUUGAUUAACCAUGU	C	C	C
hsa-miR-212	UACAGUCUCCAGUCACGGCC	C	C	NC
hsa-miR-133a	UUGGUCCCCUUAACCAGCUGU	C	C	C
hsa-miR-503	UAGCAGCGGGAACAGUUCUGCAG	C	C	C
hsa-miR-378	CUCCUGACUCCAGGUCCUGUGU	C	C	C
hsa-miR-382	GAAGUUGUUCGUGGUGGAUUCG	C	C	C
hsa-miR-520e	AAAGUGCUUCCUUUUUGAGGG	C	NC	NC

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hsa-miR-592	UUGUGUCAAU AUGCGAUGAUGU	C	C	C
hsa-miR-631	AGACCUGGCC CAGACCUCAGC	NC	C	C
hsa-miR-527	CUGCAAAGGGAAGCCCUUUCU	NC	NC	C
hsa-miR-450	UUUUUGCGAUGUGUCCUAUA	C	C	C
hsa-miR-127	UCGGAUCCGUCUGAGCUUGGCU	C	C	C
hsa-miR-495	AAACAAACAUGGUGCACUUCUU	C	C	C
hsa-miR-100	AACCCGUAGA UCCGAACUUGUG	C	C	C
hsa-miR-584	UUAUGGUUUGCCUGGGACUGAG	NC	C	C
hsa-miR-409-3p	CGAAUGUUGCUCGGUGAACCCCU	C	C	C
hsa-miR-504	AGACCCUGGUCUGCACUCUAU	C	C	C
hsa-miR-647	GUGGCUGCACUCACUUCUUC	NC	NC	NC
hsa-miR-328	CUGGCCCUCUCUGCCCUUCCGU	C	NC	C
hsa-miR-29c	UAGCACCAU UUGAAAUCGGU	C	C	C
hsa-miR-617	AGACU UCCCAUUGAAGGUGGC	NC	NC	C
hsa-miR-370	GCCUGCUGGGUGGAACCUGG	C	C	C
hsa-miR-624	UAGUACCAGUACCUUGUGUUA	NC	C	C
hsa-miR-524	GAAGGCGCUUCCCUUUGGAGU	NC	NC	C
hsa-miR-608	AGGGGUGUGUUGGGACAGCUCCGU	NC	NC	NC
hsa-miR-520d*	UCUACAAAGGGAAGCCCUUUCUG	NC	C	C
hsa-miR-453	GAGGUUGUCCGUGGUGAGUUCG	C	C	C
hsa-miR-510	UACUCAGGAGAGUGGCAAUCACA	NC	NC	NC
hsa-miR-551b	GCGACCCAUACUUGGUUUCAG	C	C	C
hsa-miR-199a	CCCAGUGUUCAGACUACCUGUUC	C	C	C
hsa-miR-622	ACAGUCUGCUGAGGUUGGAGC	NC	NC	C
hsa-miR-339	UCCUGUCCUCCAGGAGCUCA	C	C	C
hsa-miR-448	UUGCAUAUGUAGGAUGUCCCAU	C	C	C
hsa-miR-483	UCACUCCUCUCCUCCGUCUUCU	NC	NC	NC
hsa-miR-596	AAGCCUGCCCGGCUCUCCGGG	NC	NC	C
hsa-miR-431	UGUCUUGCAGGCCGUAUGCA	C	C	NC
hsa-miR-211	UUCCCUUUGUCAUCCUUCGCCU	C	C	C
hsa-miR-491	AGUGGGGAACCCUCCAUGAGGA	C	C	C
hsa-let-7a	UGAGGUAGUAGGUUGUAUAGUU	C	C	C
hsa-let-7f	UGAGGUAGUAGAUUGUAUAGUU	C	C	C
hsa-let-7e	UGAGGUAGGAGGUUGUAUAGU	C	C	C
hsa-let-7b	UGAGGUAGUAGGUUGUGUGGUU	C	C	C
hsa-let-7i	UGAGGUAGUAGUUUGUGCUGU	C	C	NC
hsa-let-7c	UGAGGUAGUAGGUUGUAUGGUU	C	C	C
hsa-let-7d	AGAGGUAGUAGGUUGCAUAGU	C	C	C
hsa-let-7g	UGAGGUAGUAGUUUGUACAGU	C	C	C