

NPM1mut-associated, downregulated genes and core miRNAs predicted to target the respective genes.
Gene symbols are sorted alphabetically.

Target gene symbol	Target mRNA, accession number	miRNA predicted to target gene	description
ABCB10	NM_012089	hsa-miR-369-3p	ABCB10 ATP-binding cassette, sub-family B (MDR/TAP), member 10 Hs.17614 R83875 23456 107773
ABCC5	NM_005688	hsa-let-7a	ABCC5 ATP-binding cassette, sub-family C (CFTR/MRP), member 5 Hs.368563 AA481269 Unknown UG Hs.178381 ESTs sc_id2627 10057 185889
ABCC5	NM_005688	hsa-let-7b	ABCC5 ATP-binding cassette, sub-family C (CFTR/MRP), member 5 Hs.368563 AA481269 Unknown UG Hs.178381 ESTs sc_id2627 10057 185889
ABCC5	NM_005688	hsa-let-7c	ABCC5 ATP-binding cassette, sub-family C (CFTR/MRP), member 5 Hs.368563 AA481269 Unknown UG Hs.178381 ESTs sc_id2627 10057 185889
ABCC5	NM_005688	hsa-let-7d	ABCC5 ATP-binding cassette, sub-family C (CFTR/MRP), member 5 Hs.368563 AA481269 Unknown UG Hs.178381 ESTs sc_id2627 10057 185889
ABCC5	NM_005688	hsa-let-7e	ABCC5 ATP-binding cassette, sub-family C (CFTR/MRP), member 5 Hs.368563 AA481269 Unknown UG Hs.178381 ESTs sc_id2627 10057 185889
ABCC5	NM_005688	hsa-let-7f	ABCC5 ATP-binding cassette, sub-family C (CFTR/MRP), member 5 Hs.368563 AA481269 Unknown UG Hs.178381 ESTs sc_id2627 10057 185889
ABCC5	NM_005688	hsa-miR-106a	ABCC5 ATP-binding cassette, sub-family C (CFTR/MRP), member 5 Hs.368563 AA481269 Unknown UG Hs.178381 ESTs sc_id2627 10057 185889
ABCC5	NM_005688	hsa-miR-142-5p	ABCC5 ATP-binding cassette, sub-family C (CFTR/MRP), member 5 Hs.368563 AA481269 Unknown UG Hs.178381 ESTs sc_id2627 10057 185889
ABCC5	NM_005688	hsa-miR-15a	ABCC5 ATP-binding cassette, sub-family C (CFTR/MRP), member 5 Hs.368563 AA481269 Unknown UG Hs.178381 ESTs sc_id2627 10057 185889
ABCC5	NM_005688	hsa-miR-15b	ABCC5 ATP-binding cassette, sub-family C (CFTR/MRP), member 5 Hs.368563 AA481269 Unknown UG Hs.178381 ESTs sc_id2627 10057 185889
ABCC5	NM_005688	hsa-miR-16	ABCC5 ATP-binding cassette, sub-family C (CFTR/MRP), member 5 Hs.368563 AA481269 Unknown UG Hs.178381 ESTs sc_id2627 10057 185889
ABCC5	NM_005688	hsa-miR-195	ABCC5 ATP-binding cassette, sub-family C (CFTR/MRP), member 5 Hs.368563 AA481269 Unknown UG Hs.178381 ESTs sc_id2627 10057 185889
ABCC5	NM_005688	hsa-miR-20a	ABCC5 ATP-binding cassette, sub-family C (CFTR/MRP), member 5 Hs.368563 AA481269 Unknown UG Hs.178381 ESTs sc_id2627 10057 185889
ABCC5	NM_005688	hsa-miR-23a	ABCC5 ATP-binding cassette, sub-family C (CFTR/MRP), member 5 Hs.368563 AA481269 Unknown UG Hs.178381 ESTs sc_id2627 10057 185889
ABCC5	NM_005688	hsa-miR-30c	ABCC5 ATP-binding cassette, sub-family C (CFTR/MRP), member 5 Hs.368563 AA481269 Unknown UG Hs.178381 ESTs sc_id2627 10057 185889
ABCC5	NM_005688	hsa-miR-369-3p	ABCC5 ATP-binding cassette, sub-family C (CFTR/MRP), member 5 Hs.368563 AA481269 Unknown UG Hs.178381 ESTs sc_id2627 10057 185889
ABCC5	NM_005688	hsa-miR-98	ABCC5 ATP-binding cassette, sub-family C (CFTR/MRP), member 5 Hs.368563 AA481269 Unknown UG Hs.178381 ESTs sc_id2627 10057 185889
ABCG2	NM_004827	hsa-miR-142-5p	ABCG2 ATP-binding cassette, sub-family G (WHITE), member 2 Hs.480218 W84773 9429 220202
ABO	NM_020469	hsa-miR-369-3p	AA974124 119062
ADCY3	NM_004036	hsa-miR-124	ADCY3 Adenylate cyclase 3 Hs.467898 AA400456 109 106954
ADCY3	NM_004036	hsa-miR-27a	ADCY3 Adenylate cyclase 3 Hs.467898 AA400456 109 106954
ADCY3	NM_004036	hsa-miR-92a	ADCY3 Adenylate cyclase 3 Hs.467898 AA400456 109 106954
ADIPOR1	NM_015999	hsa-miR-15b	ADIPOR1 Adiponectin receptor 1 Hs.5298 H48233 51094 108833
AGTPBP1	NM_015239	hsa-miR-18a	AGTPBP1 ATP/GTP binding protein 1 Hs.494321 N46882 23287 224959
AGTPBP1	NM_015239	hsa-miR-20a	AGTPBP1 ATP/GTP binding protein 1 Hs.494321 N46882 23287 224959
AGTPBP1	NM_015239	hsa-miR-369-3p	AGTPBP1 ATP/GTP binding protein 1 Hs.494321 N46882 23287 224959
AKAP13	NM_144767	hsa-let-7d	AKAP13 A kinase (PRKA) anchor protein 13 Hs.459211 AA156936 11214 104161
AKAP13	NM_144767	hsa-miR-106a	AKAP13 A kinase (PRKA) anchor protein 13 Hs.459211 AA156936 11214 104161
AKAP13	NM_144767	hsa-miR-15a	AKAP13 A kinase (PRKA) anchor protein 13 Hs.459211 AA156936 11214 104161
AKAP13	NM_144767	hsa-miR-17	AKAP13 A kinase (PRKA) anchor protein 13 Hs.459211 AA156936 11214 104161
AKAP13	NM_144767	hsa-miR-20a	AKAP13 A kinase (PRKA) anchor protein 13 Hs.459211 AA156936 11214 104161
AKAP13	NM_144767	hsa-miR-29a	AKAP13 A kinase (PRKA) anchor protein 13 Hs.459211 AA156936 11214 104161
AKAP13	NM_144767	hsa-miR-29b	AKAP13 A kinase (PRKA) anchor protein 13 Hs.459211 AA156936 11214 104161
AKAP13	NM_144767	hsa-miR-29c	AKAP13 A kinase (PRKA) anchor protein 13 Hs.459211 AA156936 11214 104161
AKAP13	NM_144767	hsa-miR-30c	AKAP13 A kinase (PRKA) anchor protein 13 Hs.459211 AA156936 11214 104161
ALAS2	NM_000032	hsa-miR-142-5p	ALAS2 Aminolevulinatase, delta-, synthase 2 (sideroblastic/hypochromic anemia) Hs.522666 AA406485 212 103595
ALS2CR2	NM_018571	hsa-miR-15a	ALS2CR2 **Amyotrophic lateral sclerosis 2 (juvenile) chromosome region, candidate 2 Hs.127519 H98822 55437 119965
ALS2CR2	NM_018571	hsa-miR-15b	ALS2CR2 **Amyotrophic lateral sclerosis 2 (juvenile) chromosome region, candidate 2 Hs.127519 H98822 55437 119965
ALS2CR2	NM_018571	hsa-miR-16	ALS2CR2 **Amyotrophic lateral sclerosis 2 (juvenile) chromosome region, candidate 2 Hs.127519 H98822 55437 119965
ALS2CR2	NM_018571	hsa-miR-195	ALS2CR2 **Amyotrophic lateral sclerosis 2 (juvenile) chromosome region, candidate 2 Hs.127519 H98822 55437 119965
ALS2CR3	NM_015049	hsa-miR-19a	ALS2CR3 **Amyotrophic lateral sclerosis 2 (juvenile) chromosome region, candidate 3 Hs.152774 AA487425 66008 101974
ALS2CR3	NM_015049	hsa-miR-19b	ALS2CR3 **Amyotrophic lateral sclerosis 2 (juvenile) chromosome region, candidate 3 Hs.152774 AA487425 66008 101974
ALS2CR3	NM_015049	hsa-miR-92a	ALS2CR3 **Amyotrophic lateral sclerosis 2 (juvenile) chromosome region, candidate 3 Hs.152774 AA487425 66008 101974
ANK1	NM_020477	hsa-miR-27a	ANK1 Ankyrin 1, erythrocytic Hs.491558 AA464049 286 112116
ANKRD9	NM_152326	hsa-miR-106a	ANKRD9 Ankyrin repeat domain 9 Hs.432945 AA402043 122416 220603
ANKRD9	NM_152326	hsa-miR-17	ANKRD9 Ankyrin repeat domain 9 Hs.432945 AA402043 122416 220603

ANKRD9	NM 152326	hsa-miR-20a	ANKRD9 Ankyrin repeat domain 9 Hs.432945 AA402043 122416 220603
APP	NM 000484	hsa-let-7a	APP Amyloid beta (A4) precursor protein (protease nexin-II, Alzheimer disease) Hs.434980 W42848 351 101388
APP	NM 000484	hsa-let-7d	APP Amyloid beta (A4) precursor protein (protease nexin-II, Alzheimer disease) Hs.434980 W42848 351 101388
APP	NM 000484	hsa-let-7f	APP Amyloid beta (A4) precursor protein (protease nexin-II, Alzheimer disease) Hs.434980 W42848 351 101388
APP	NM 000484	hsa-miR-106a	APP Amyloid beta (A4) precursor protein (protease nexin-II, Alzheimer disease) Hs.434980 W42848 351 101388
APP	NM 201414	hsa-miR-15a	APP Amyloid beta (A4) precursor protein (protease nexin-II, Alzheimer disease) Hs.434980 W42848 351 101388
APP	NM 201414	hsa-miR-15b	APP Amyloid beta (A4) precursor protein (protease nexin-II, Alzheimer disease) Hs.434980 W42848 351 101388
APP	NM 201414	hsa-miR-16	APP Amyloid beta (A4) precursor protein (protease nexin-II, Alzheimer disease) Hs.434980 W42848 351 101388
APP	NM 000484	hsa-miR-17	APP Amyloid beta (A4) precursor protein (protease nexin-II, Alzheimer disease) Hs.434980 W42848 351 101388
APP	NM 201414	hsa-miR-195	APP Amyloid beta (A4) precursor protein (protease nexin-II, Alzheimer disease) Hs.434980 W42848 351 101388
APP	NM 000484	hsa-miR-20a	APP Amyloid beta (A4) precursor protein (protease nexin-II, Alzheimer disease) Hs.434980 W42848 351 101388
APP	NM 000484	hsa-miR-23a	APP Amyloid beta (A4) precursor protein (protease nexin-II, Alzheimer disease) Hs.434980 W42848 351 101388
APP	NM 000484	hsa-miR-29a	APP Amyloid beta (A4) precursor protein (protease nexin-II, Alzheimer disease) Hs.434980 W42848 351 101388
APP	NM 000484	hsa-miR-29b	APP Amyloid beta (A4) precursor protein (protease nexin-II, Alzheimer disease) Hs.434980 W42848 351 101388
APP	NM 000484	hsa-miR-29c	APP Amyloid beta (A4) precursor protein (protease nexin-II, Alzheimer disease) Hs.434980 W42848 351 101388
APP	NM 201414	hsa-miR-369-3p	APP Amyloid beta (A4) precursor protein (protease nexin-II, Alzheimer disease) Hs.434980 W42848 351 101388
AR	NM 001011645	hsa-miR-124	AR Androgen receptor (dihydrotestosterone receptor; testicular feminization; spinal and bulbar muscular atrophy; Kennedy disease) Hs.496240 AI659563 367
ARF6	NM 001663	hsa-miR-18a	ARF6 ADP-ribosylation factor 6 Hs.525330 AA012866 382 108333
ARF6	NM 001663	hsa-miR-23a	ARF6 ADP-ribosylation factor 6 Hs.525330 AA012866 382 108333
ARG1	NM 000045	hsa-miR-23a	Arginase liver 245589
ARHGEF10	NM 014629	hsa-miR-106a	N53385 110081
ARHGEF10	NM 014629	hsa-miR-17	N53385 110081
ARHGEF10	NM 014629	hsa-miR-20a	N53385 110081
ARHGEF10	NM 014629	hsa-miR-369-3p	N53385 110081
ARHGEF10	NM 014629	hsa-miR-92a	N53385 110081
ARHGEF12	NM 015313	hsa-miR-19a	ARHGEF12 Rho guanine nucleotide exchange factor (GEF) 12 Hs.24598 AA479287 23365 103550
ARHGEF12	NM 015313	hsa-miR-19b	ARHGEF12 Rho guanine nucleotide exchange factor (GEF) 12 Hs.24598 AA479287 23365 103550
ARL2	NM 001667	hsa-let-7d	ARL2 ADP-ribosylation factor-like 2 Hs.502836 AW075431 402 331371
ARL2	NM 001667	hsa-miR-15a	ARL2 ADP-ribosylation factor-like 2 Hs.502836 AW075431 402 331371
ARL2	NM 001667	hsa-miR-15b	ARL2 ADP-ribosylation factor-like 2 Hs.502836 AW075431 402 331371
ARL2	NM 001667	hsa-miR-16	ARL2 ADP-ribosylation factor-like 2 Hs.502836 AW075431 402 331371
ARL2	NM 001667	hsa-miR-195	ARL2 ADP-ribosylation factor-like 2 Hs.502836 AW075431 402 331371
ASF1A	NM 014034	hsa-miR-106a	ASF1A ASF1 anti-silencing function 1 homolog A (S. cerevisiae) Hs.292316 T54544 25842 101047
ASF1A	NM 014034	hsa-miR-124	ASF1A ASF1 anti-silencing function 1 homolog A (S. cerevisiae) Hs.292316 T54544 25842 101047
ASF1A	NM 014034	hsa-miR-17	ASF1A ASF1 anti-silencing function 1 homolog A (S. cerevisiae) Hs.292316 T54544 25842 101047
ASF1A	NM 014034	hsa-miR-20a	ASF1A ASF1 anti-silencing function 1 homolog A (S. cerevisiae) Hs.292316 T54544 25842 101047
ASF1A	NM 014034	hsa-miR-21	ASF1A ASF1 anti-silencing function 1 homolog A (S. cerevisiae) Hs.292316 T54544 25842 101047
ASF1A	NM 014034	hsa-miR-23a	ASF1A ASF1 anti-silencing function 1 homolog A (S. cerevisiae) Hs.292316 T54544 25842 101047
ASF1A	NM 014034	hsa-miR-369-3p	ASF1A ASF1 anti-silencing function 1 homolog A (S. cerevisiae) Hs.292316 T54544 25842 101047
ATXN10	NM 013236	hsa-miR-21	ATXN10 Ataxin 10 Hs.475125 AI003963 25814 308243
ATXN10	NM 013236	hsa-miR-27a	ATXN10 Ataxin 10 Hs.475125 AI003963 25814 308243
BCL2L1	NM 138578	hsa-let-7a	BCL2L1 BCL2-like 1 Hs.516966 AA931820 598 310821
BCL2L1	NM 138578	hsa-let-7b	BCL2L1 BCL2-like 1 Hs.516966 AA931820 598 310821
BCL2L1	NM 138578	hsa-let-7c	BCL2L1 BCL2-like 1 Hs.516966 AA931820 598 310821
BCL2L1	NM 138578	hsa-let-7d	BCL2L1 BCL2-like 1 Hs.516966 AA931820 598 310821
BCL2L1	NM 138578	hsa-let-7e	BCL2L1 BCL2-like 1 Hs.516966 AA931820 598 310821
BCL2L1	NM 138578	hsa-let-7f	BCL2L1 BCL2-like 1 Hs.516966 AA931820 598 310821
BCL2L1	NM 138578	hsa-miR-30c	BCL2L1 BCL2-like 1 Hs.516966 AA931820 598 310821
BCL2L1	NM 138578	hsa-miR-98	BCL2L1 BCL2-like 1 Hs.516966 AA931820 598 310821
BMP7	NM 001719	hsa-let-7b	BMP7 Bone morphogenetic protein 7 (osteogenic protein 1) Hs.473163 AA029596 655 111581
BMP7	NM 001719	hsa-let-7c	BMP7 Bone morphogenetic protein 7 (osteogenic protein 1) Hs.473163 AA029596 655 111581
BMP7	NM 001719	hsa-let-7d	BMP7 Bone morphogenetic protein 7 (osteogenic protein 1) Hs.473163 AA029596 655 111581

BNIP3L	NM 004331	hsa-miR-23a	BNIP3L BCL2/adenovirus E1B 19kDa interacting protein 3-like Hs.131226 AA465697 665 307692
BNIP3L	NM 004331	hsa-miR-27a	BNIP3L BCL2/adenovirus E1B 19kDa interacting protein 3-like Hs.131226 AA465697 665 307692
BNIP3L	NM 004331	hsa-miR-30c	BNIP3L BCL2/adenovirus E1B 19kDa interacting protein 3-like Hs.131226 AA465697 665 307692
BNIP3L	NM 004331	hsa-miR-30e	BNIP3L BCL2/adenovirus E1B 19kDa interacting protein 3-like Hs.131226 AA465697 665 307692
BTBD7	NM 001002860	hsa-miR-106a	BTBD7 BTB (POZ) domain containing 7 Hs.525549 AI000138 55727 102852
BTBD7	NM 018167	hsa-miR-124	BTBD7 BTB (POZ) domain containing 7 Hs.525549 AI000138 55727 102852
BTBD7	NM 001002860	hsa-miR-142-5p	BTBD7 BTB (POZ) domain containing 7 Hs.525549 AI000138 55727 102852
BTBD7	NM 001002860	hsa-miR-17	BTBD7 BTB (POZ) domain containing 7 Hs.525549 AI000138 55727 102852
BTBD7	NM 001002860	hsa-miR-19a	BTBD7 BTB (POZ) domain containing 7 Hs.525549 AI000138 55727 102852
BTBD7	NM 001002860	hsa-miR-19b	BTBD7 BTB (POZ) domain containing 7 Hs.525549 AI000138 55727 102852
BTBD7	NM 001002860	hsa-miR-20a	BTBD7 BTB (POZ) domain containing 7 Hs.525549 AI000138 55727 102852
BTBD7	NM 018167	hsa-miR-23a	BTBD7 BTB (POZ) domain containing 7 Hs.525549 AI000138 55727 102852
BTBD7	NM 018167	hsa-miR-29a	BTBD7 BTB (POZ) domain containing 7 Hs.525549 AI000138 55727 102852
BTBD7	NM 018167	hsa-miR-29b	BTBD7 BTB (POZ) domain containing 7 Hs.525549 AI000138 55727 102852
BTBD7	NM 018167	hsa-miR-29c	BTBD7 BTB (POZ) domain containing 7 Hs.525549 AI000138 55727 102852
BTBD7	NM 018167	hsa-miR-30c	BTBD7 BTB (POZ) domain containing 7 Hs.525549 AI000138 55727 102852
BTBD7	NM 018167	hsa-miR-30e	BTBD7 BTB (POZ) domain containing 7 Hs.525549 AI000138 55727 102852
BTRC	NM 033637	hsa-miR-10a	BTRC Beta-transducin repeat containing Hs.500812 AA398420 8945 109640
BTRC	NM 033637	hsa-miR-10b	BTRC Beta-transducin repeat containing Hs.500812 AA398420 8945 109640
BTRC	NM 033637	hsa-miR-15a	BTRC Beta-transducin repeat containing Hs.500812 AA398420 8945 109640
BTRC	NM 033637	hsa-miR-15b	BTRC Beta-transducin repeat containing Hs.500812 AA398420 8945 109640
BTRC	NM 033637	hsa-miR-16	BTRC Beta-transducin repeat containing Hs.500812 AA398420 8945 109640
BTRC	NM 033637	hsa-miR-195	BTRC Beta-transducin repeat containing Hs.500812 AA398420 8945 109640
C10orf58	NM 032333	hsa-miR-106a	C10orf58 Chromosome 10 open reading frame 58 Hs.500333 AI339215 84293 311092
C10orf58	NM 032333	hsa-miR-16	C10orf58 Chromosome 10 open reading frame 58 Hs.500333 AI339215 84293 311092
C10orf58	NM 032333	hsa-miR-17	C10orf58 Chromosome 10 open reading frame 58 Hs.500333 AI339215 84293 311092
C10orf58	NM 032333	hsa-miR-195	C10orf58 Chromosome 10 open reading frame 58 Hs.500333 AI339215 84293 311092
C10orf58	NM 032333	hsa-miR-20a	C10orf58 Chromosome 10 open reading frame 58 Hs.500333 AI339215 84293 311092
C10orf58	NM 032333	hsa-miR-92a	C10orf58 Chromosome 10 open reading frame 58 Hs.500333 AI339215 84293 311092
C12orf22	NM 030809	hsa-miR-155	C12orf22 Chromosome 12 open reading frame 22 Hs.524425 AA018066 81566 102865
C12orf22	NM 030809	hsa-miR-29a	C12orf22 Chromosome 12 open reading frame 22 Hs.524425 AA018066 81566 102865
C12orf22	NM 030809	hsa-miR-29b	C12orf22 Chromosome 12 open reading frame 22 Hs.524425 AA018066 81566 102865
C12orf22	NM 030809	hsa-miR-29c	C12orf22 Chromosome 12 open reading frame 22 Hs.524425 AA018066 81566 102865
C12orf51	NM 173813	hsa-miR-30c	FLJ34154 Hypothetical protein FLJ34154 Hs.379848 H66282 283450 101747
C12orf51	NM 173813	hsa-miR-30e	FLJ34154 Hypothetical protein FLJ34154 Hs.379848 H66282 283450 101747
C13orf22	NM 005800	hsa-miR-23a	C13orf22 Chromosome 13 open reading frame 22 Hs.533831 H95641 10208 107727
C13orf22	NM 005800	hsa-miR-29b	C13orf22 Chromosome 13 open reading frame 22 Hs.533831 H95641 10208 107727
C13orf22	NM 005800	hsa-miR-30c	C13orf22 Chromosome 13 open reading frame 22 Hs.533831 H95641 10208 107727
C13orf22	NM 005800	hsa-miR-30e	C13orf22 Chromosome 13 open reading frame 22 Hs.533831 H95641 10208 107727
C13orf22	NM 005800	hsa-miR-92a	C13orf22 Chromosome 13 open reading frame 22 Hs.533831 H95641 10208 107727
C14orf31	NM 152330	hsa-miR-106a	C14orf31 Chromosome 14 open reading frame 31 Hs.434914 AA461078 122786 101690
C14orf31	NM 152330	hsa-miR-17	C14orf31 Chromosome 14 open reading frame 31 Hs.434914 AA461078 122786 101690
C14orf31	NM 152330	hsa-miR-20a	C14orf31 Chromosome 14 open reading frame 31 Hs.434914 AA461078 122786 101690
C14orf31	NM 152330	hsa-miR-30c	C14orf31 Chromosome 14 open reading frame 31 Hs.434914 AA461078 122786 101690
C14orf31	NM 152330	hsa-miR-30e	C14orf31 Chromosome 14 open reading frame 31 Hs.434914 AA461078 122786 101690
C14orf31	NM 152330	hsa-miR-369-3p	C14orf31 Chromosome 14 open reading frame 31 Hs.434914 AA461078 122786 101690
C14orf87	NM 016417	hsa-miR-27a	C14orf87 Chromosome 14 open reading frame 87 Hs.532683 AA677563 51218 225214
C14orf87	NM 016417	hsa-miR-369-3p	C14orf87 Chromosome 14 open reading frame 87 Hs.532683 AA677563 51218 225214
C18orf1	NM 181483	hsa-miR-92a	C18orf1 Chromosome 18 open reading frame 1 Hs.464697 AA489633 753 113788
C18orf10	NM 015476	hsa-let-7a	C18orf10 Chromosome 18 open reading frame 10 Hs.22981 AA465218 25941 221057
C18orf10	NM 015476	hsa-let-7d	C18orf10 Chromosome 18 open reading frame 10 Hs.22981 AA465218 25941 221057
C18orf10	NM 015476	hsa-let-7e	C18orf10 Chromosome 18 open reading frame 10 Hs.22981 AA465218 25941 221057
C18orf10	NM 015476	hsa-let-7f	C18orf10 Chromosome 18 open reading frame 10 Hs.22981 AA465218 25941 221057

C18orf10	NM 015476	hsa-miR-155	C18orf10 Chromosome 18 open reading frame 10 Hs.22981 AA465218 25941 221057
C1orf21	NM 030806	hsa-miR-124	C1orf21 Chromosome 1 open reading frame 21 Hs.497159 N26008 81563 224141
C1orf21	NM 030806	hsa-miR-15a	C1orf21 Chromosome 1 open reading frame 21 Hs.497159 N26008 81563 224141
C1orf21	NM 030806	hsa-miR-15b	C1orf21 Chromosome 1 open reading frame 21 Hs.497159 N26008 81563 224141
C1orf21	NM 030806	hsa-miR-16	C1orf21 Chromosome 1 open reading frame 21 Hs.497159 N26008 81563 224141
C1orf21	NM 030806	hsa-miR-195	C1orf21 Chromosome 1 open reading frame 21 Hs.497159 N26008 81563 224141
C1orf21	NM 030806	hsa-miR-27a	C1orf21 Chromosome 1 open reading frame 21 Hs.497159 N26008 81563 224141
C1orf21	NM 030806	hsa-miR-29a	C1orf21 Chromosome 1 open reading frame 21 Hs.497159 N26008 81563 224141
C1orf21	NM 030806	hsa-miR-29b	C1orf21 Chromosome 1 open reading frame 21 Hs.497159 N26008 81563 224141
C1orf21	NM 030806	hsa-miR-29c	C1orf21 Chromosome 1 open reading frame 21 Hs.497159 N26008 81563 224141
C20orf77	NM 021215	hsa-miR-15a	C20orf77 Chromosome 20 open reading frame 77 Hs.278839 N76944 58490 115132
C20orf77	NM 021215	hsa-miR-15b	C20orf77 Chromosome 20 open reading frame 77 Hs.278839 N76944 58490 115132
C22orf13	NM 031444	hsa-miR-124	C22orf13 Chromosome 22 open reading frame 13 Hs.9850 AA190998 83606 105333
C22orf13	NM 031444	hsa-miR-19a	C22orf13 Chromosome 22 open reading frame 13 Hs.9850 AA190998 83606 105333
C22orf13	NM 031444	hsa-miR-19b	C22orf13 Chromosome 22 open reading frame 13 Hs.9850 AA190998 83606 105333
C3orf6	NM 178335	hsa-miR-142-5p	C3orf6 Chromosome 3 open reading frame 6 Hs.478682 AA701978 152137 223512
C3orf6	NM 178335	hsa-miR-19a	C3orf6 Chromosome 3 open reading frame 6 Hs.478682 AA701978 152137 223512
C3orf6	NM 178335	hsa-miR-19b	C3orf6 Chromosome 3 open reading frame 6 Hs.478682 AA701978 152137 223512
C6orf204	NM 206921	hsa-miR-10a	AA621378 220606
C6orf204	NM 206921	hsa-miR-10b	AA621378 220606
C9orf78	NM 016520	hsa-miR-106a	C9orf78 Chromosome 9 open reading frame 78 Hs.278429 W78986 51759 116051
C9orf78	NM 016520	hsa-miR-155	C9orf78 Chromosome 9 open reading frame 78 Hs.278429 W78986 51759 116051
C9orf78	NM 016520	hsa-miR-17	C9orf78 Chromosome 9 open reading frame 78 Hs.278429 W78986 51759 116051
C9orf78	NM 016520	hsa-miR-20a	C9orf78 Chromosome 9 open reading frame 78 Hs.278429 W78986 51759 116051
CA1	NM 001738	hsa-miR-92a	CA1 Carbonic anhydrase I Hs.23118 AA911903 759 109290
CALM3	NM 005184	hsa-let-7a	CALM3 Calmodulin 3 (phosphorylase kinase, delta) Hs.515487 AA663828 808 119251
CALM3	NM 005184	hsa-let-7b	CALM3 Calmodulin 3 (phosphorylase kinase, delta) Hs.515487 AA663828 808 119251
CALM3	NM 005184	hsa-let-7c	CALM3 Calmodulin 3 (phosphorylase kinase, delta) Hs.515487 AA663828 808 119251
CALM3	NM 005184	hsa-let-7d	CALM3 Calmodulin 3 (phosphorylase kinase, delta) Hs.515487 AA663828 808 119251
CALM3	NM 005184	hsa-let-7e	CALM3 Calmodulin 3 (phosphorylase kinase, delta) Hs.515487 AA663828 808 119251
CALM3	NM 005184	hsa-let-7f	CALM3 Calmodulin 3 (phosphorylase kinase, delta) Hs.515487 AA663828 808 119251
CALM3	NM 005184	hsa-miR-15a	CALM3 Calmodulin 3 (phosphorylase kinase, delta) Hs.515487 AA663828 808 119251
CALM3	NM 005184	hsa-miR-15b	CALM3 Calmodulin 3 (phosphorylase kinase, delta) Hs.515487 AA663828 808 119251
CALM3	NM 005184	hsa-miR-16	CALM3 Calmodulin 3 (phosphorylase kinase, delta) Hs.515487 AA663828 808 119251
CALM3	NM 005184	hsa-miR-195	CALM3 Calmodulin 3 (phosphorylase kinase, delta) Hs.515487 AA663828 808 119251
CALM3	NM 005184	hsa-miR-19a	CALM3 Calmodulin 3 (phosphorylase kinase, delta) Hs.515487 AA663828 808 119251
CALM3	NM 005184	hsa-miR-19b	CALM3 Calmodulin 3 (phosphorylase kinase, delta) Hs.515487 AA663828 808 119251
CALM3	NM 005184	hsa-miR-27a	CALM3 Calmodulin 3 (phosphorylase kinase, delta) Hs.515487 AA663828 808 119251
CALM3	NM 005184	hsa-miR-29a	CALM3 Calmodulin 3 (phosphorylase kinase, delta) Hs.515487 AA663828 808 119251
CALM3	NM 005184	hsa-miR-29b	CALM3 Calmodulin 3 (phosphorylase kinase, delta) Hs.515487 AA663828 808 119251
CALM3	NM 005184	hsa-miR-29c	CALM3 Calmodulin 3 (phosphorylase kinase, delta) Hs.515487 AA663828 808 119251
CAND1	NM 018448	hsa-miR-30c	Transcribed locus Hs.172608 AI086239 317695
CAND1	NM 018448	hsa-miR-30e	Transcribed locus Hs.172608 AI086239 317695
CBR3	NM 001236	hsa-miR-29a	CBR3 Carbonyl reductase 3 Hs.154510 AI352345 874 309954
CBR3	NM 001236	hsa-miR-29b	CBR3 Carbonyl reductase 3 Hs.154510 AI352345 874 309954
CBR3	NM 001236	hsa-miR-29c	CBR3 Carbonyl reductase 3 Hs.154510 AI352345 874 309954
CBR3	NM 001236	hsa-miR-30e	CBR3 Carbonyl reductase 3 Hs.154510 AI352345 874 309954
CCL28	NM 019846	hsa-miR-155	CCL28 Chemokine (C-C motif) ligand 28 Hs.334633 R36539 56477 120584
CCM2	NM 031443	hsa-miR-18a	CCM2 Cerebral cavernous malformation 2 Hs.148272 AA890153 83605 224118
CCM2	NM 031443	hsa-miR-19a	CCM2 Cerebral cavernous malformation 2 Hs.148272 AA890153 83605 224118
CCM2	NM 031443	hsa-miR-19b	CCM2 Cerebral cavernous malformation 2 Hs.148272 AA890153 83605 224118
CCM2	NM 031443	hsa-miR-23a	CCM2 Cerebral cavernous malformation 2 Hs.148272 AA890153 83605 224118
CCM2	NM 031443	hsa-miR-27a	CCM2 Cerebral cavernous malformation 2 Hs.148272 AA890153 83605 224118

CCNB2	NM 004701	hsa-let-7e	CCNB2 Cyclin B2 Hs.194698 AA774665 9133 223334
CCNB2	NM 004701	hsa-miR-142-5p	CCNB2 Cyclin B2 Hs.194698 AA774665 9133 223334
CCNB2	NM 004701	hsa-miR-19a	CCNB2 Cyclin B2 Hs.194698 AA774665 9133 223334
CCNB2	NM 004701	hsa-miR-19b	CCNB2 Cyclin B2 Hs.194698 AA774665 9133 223334
CCNB2	NM 004701	hsa-miR-23a	CCNB2 Cyclin B2 Hs.194698 AA774665 9133 223334
CCND1	NM 053056	hsa-let-7a	CCND1 Cyclin D1 (PRAD1: parathyroid adenomatosis 1) Hs.523852 T77237 595 228984
CCND1	NM 053056	hsa-let-7b	CCND1 Cyclin D1 (PRAD1: parathyroid adenomatosis 1) Hs.523852 T77237 595 228984
CCND1	NM 053056	hsa-let-7c	CCND1 Cyclin D1 (PRAD1: parathyroid adenomatosis 1) Hs.523852 T77237 595 228984
CCND1	NM 053056	hsa-let-7d	CCND1 Cyclin D1 (PRAD1: parathyroid adenomatosis 1) Hs.523852 T77237 595 228984
CCND1	NM 053056	hsa-let-7e	CCND1 Cyclin D1 (PRAD1: parathyroid adenomatosis 1) Hs.523852 T77237 595 228984
CCND1	NM 053056	hsa-let-7f	CCND1 Cyclin D1 (PRAD1: parathyroid adenomatosis 1) Hs.523852 T77237 595 228984
CCND1	NM 053056	hsa-miR-106a	CCND1 Cyclin D1 (PRAD1: parathyroid adenomatosis 1) Hs.523852 T77237 595 228984
CCND1	NM 053056	hsa-miR-142-5p	CCND1 Cyclin D1 (PRAD1: parathyroid adenomatosis 1) Hs.523852 T77237 595 228984
CCND1	NM 053056	hsa-miR-155	CCND1 Cyclin D1 (PRAD1: parathyroid adenomatosis 1) Hs.523852 T77237 595 228984
CCND1	NM 053056	hsa-miR-15a	CCND1 Cyclin D1 (PRAD1: parathyroid adenomatosis 1) Hs.523852 T77237 595 228984
CCND1	NM 053056	hsa-miR-15b	CCND1 Cyclin D1 (PRAD1: parathyroid adenomatosis 1) Hs.523852 T77237 595 228984
CCND1	NM 053056	hsa-miR-16	CCND1 Cyclin D1 (PRAD1: parathyroid adenomatosis 1) Hs.523852 T77237 595 228984
CCND1	NM 053056	hsa-miR-17	CCND1 Cyclin D1 (PRAD1: parathyroid adenomatosis 1) Hs.523852 T77237 595 228984
CCND1	NM 053056	hsa-miR-19a	CCND1 Cyclin D1 (PRAD1: parathyroid adenomatosis 1) Hs.523852 T77237 595 228984
CCND1	NM 053056	hsa-miR-195	CCND1 Cyclin D1 (PRAD1: parathyroid adenomatosis 1) Hs.523852 T77237 595 228984
CCND1	NM 053056	hsa-miR-20a	CCND1 Cyclin D1 (PRAD1: parathyroid adenomatosis 1) Hs.523852 T77237 595 228984
CCND1	NM 053056	hsa-miR-23a	CCND1 Cyclin D1 (PRAD1: parathyroid adenomatosis 1) Hs.523852 T77237 595 228984
CCND1	NM 053056	hsa-miR-369-3p	CCND1 Cyclin D1 (PRAD1: parathyroid adenomatosis 1) Hs.523852 T77237 595 228984
CCND1	NM 053056	hsa-miR-98	CCND1 Cyclin D1 (PRAD1: parathyroid adenomatosis 1) Hs.523852 T77237 595 228984
CD34	NM 001773	hsa-miR-15a	CD34 CD34 antigen Hs.374990 H72113 CD34 947 105957
CD38	NM 001775	hsa-let-7a	CD38 CD38 antigen (p45) Hs.479214 R00276 CD38 952 106449
CD38	NM 001775	hsa-let-7c	CD38 CD38 antigen (p45) Hs.479214 R00276 CD38 952 106449
CD38	NM 001775	hsa-let-7f	CD38 CD38 antigen (p45) Hs.479214 R00276 CD38 952 106449
CDC27	NM 001256	hsa-miR-15a	CDC27 Cell division cycle 27 Hs.463295 T58431 996 312314
CDC27	NM 001256	hsa-miR-15b	CDC27 Cell division cycle 27 Hs.463295 T58431 996 312314
CDC27	NM 001256	hsa-miR-16	CDC27 Cell division cycle 27 Hs.463295 T58431 996 312314
CDC27	NM 001256	hsa-miR-195	CDC27 Cell division cycle 27 Hs.463295 T58431 996 312314
CDCA7	NM 031942	hsa-miR-106a	CDCA7 Cell division cycle associated 7 Hs.470654 AA406348 83879 111783
CDCA7	NM 031942	hsa-miR-20a	CDCA7 Cell division cycle associated 7 Hs.470654 AA406348 83879 111783
CENTD1	NM 139182	hsa-miR-10a	CENTD1 Centaurin, delta 1 Hs.479451 AA193671 116984 113987
CENTD1	NM 139182	hsa-miR-10b	CENTD1 Centaurin, delta 1 Hs.479451 AA193671 116984 113987
CENTD1	NM 139182	hsa-miR-17	CENTD1 Centaurin, delta 1 Hs.479451 AA193671 116984 113987
CENTD1	NM 015230	hsa-miR-17	CENTD1 Centaurin, delta 1 Hs.479451 AA193671 116984 113987
CEP350	NM 014810	hsa-miR-124	CAP350 Centrosome-associated protein 350 Hs.413045 AI001846 9857 310489
CEP350	NM 014810	hsa-miR-15a	CAP350 Centrosome-associated protein 350 Hs.413045 AI001846 9857 310489
CEP350	NM 014810	hsa-miR-15b	CAP350 Centrosome-associated protein 350 Hs.413045 AI001846 9857 310489
CEP350	NM 014810	hsa-miR-16	CAP350 Centrosome-associated protein 350 Hs.413045 AI001846 9857 310489
CEP350	NM 014810	hsa-miR-195	CAP350 Centrosome-associated protein 350 Hs.413045 AI001846 9857 310489
CEP350	NM 014810	hsa-miR-19a	CAP350 Centrosome-associated protein 350 Hs.413045 AI001846 9857 310489
CEP350	NM 014810	hsa-miR-19b	CAP350 Centrosome-associated protein 350 Hs.413045 AI001846 9857 310489
CHI3L2	NM 004000	hsa-miR-16	CHI3L2 Chitinase 3-like 2 Hs.514840 AA481250 chitinase precursor 1117 161172
CHI3L2	NM 004000	hsa-miR-98	CHI3L2 Chitinase 3-like 2 Hs.514840 AA481250 chitinase precursor 1117 161172
CHPT1	NM 020244	hsa-let-7a	SYCP3 **Synaptonemal complex protein 3 Hs.506504 AA009840 50511 119083
CHPT1	NM 020244	hsa-let-7b	SYCP3 **Synaptonemal complex protein 3 Hs.506504 AA009840 50511 119083
CHPT1	NM 020244	hsa-let-7c	SYCP3 **Synaptonemal complex protein 3 Hs.506504 AA009840 50511 119083
CHPT1	NM 020244	hsa-let-7d	SYCP3 **Synaptonemal complex protein 3 Hs.506504 AA009840 50511 119083
CHPT1	NM 020244	hsa-let-7e	SYCP3 **Synaptonemal complex protein 3 Hs.506504 AA009840 50511 119083
CHPT1	NM 020244	hsa-let-7f	SYCP3 **Synaptonemal complex protein 3 Hs.506504 AA009840 50511 119083

CHPT1	NM_020244	hsa-miR-15a	SYCP3	**Synaptonemal complex protein 3	Hs.506504 AA009840	50511	119083
CHPT1	NM_020244	hsa-miR-15b	SYCP3	**Synaptonemal complex protein 3	Hs.506504 AA009840	50511	119083
CHPT1	NM_020244	hsa-miR-16	SYCP3	**Synaptonemal complex protein 3	Hs.506504 AA009840	50511	119083
CHPT1	NM_020244	hsa-miR-195	SYCP3	**Synaptonemal complex protein 3	Hs.506504 AA009840	50511	119083
CHPT1	NM_020244	hsa-miR-369-3p	SYCP3	**Synaptonemal complex protein 3	Hs.506504 AA009840	50511	119083
CHPT1	NM_020244	hsa-miR-98	SYCP3	**Synaptonemal complex protein 3	Hs.506504 AA009840	50511	119083
CLCN3	NM_001829	hsa-miR-142-5p	CLCN3	Chloride channel 3	Hs.481186 AA459750	1182	115386
CLCN3	NM_173872	hsa-miR-155	CLCN3	Chloride channel 3	Hs.481186 AA459750	1182	115386
CLCN3	NM_173872	hsa-miR-15a	CLCN3	Chloride channel 3	Hs.481186 AA459750	1182	115386
CLCN3	NM_173872	hsa-miR-15b	CLCN3	Chloride channel 3	Hs.481186 AA459750	1182	115386
CLCN3	NM_173872	hsa-miR-23a	CLCN3	Chloride channel 3	Hs.481186 AA459750	1182	115386
CLCN3	NM_001829	hsa-miR-27a	CLCN3	Chloride channel 3	Hs.481186 AA459750	1182	115386
CLEC2B	NM_005127	hsa-miR-23a	CLECSF2	C-type lectin domain family 2, member B	Hs.85201 H11732 AICL=activation-induced C-type lectin	9976	104646
CLIC4	NM_013943	hsa-miR-369-3p	CLIC4	Chloride intracellular channel 4	Hs.440544 AA917861	25932	222968
CMKOR1	NM_020311	hsa-miR-10a	CMKOR1	Chemokine orphan receptor 1	Hs.471751 N58609 RDC1=orphan G protein-coupled receptor	57007	116170
CMKOR1	NM_020311	hsa-miR-10b	CMKOR1	Chemokine orphan receptor 1	Hs.471751 N58609 RDC1=orphan G protein-coupled receptor	57007	116170
CMKOR1	NM_020311	hsa-miR-142-5p	CMKOR1	Chemokine orphan receptor 1	Hs.471751 N58609 RDC1=orphan G protein-coupled receptor	57007	116170
CMKOR1	NM_020311	hsa-miR-18a	CMKOR1	Chemokine orphan receptor 1	Hs.471751 N58609 RDC1=orphan G protein-coupled receptor	57007	116170
CNOT6L	NM_144571	hsa-miR-18a	CNOT6L	CCR4-NOT transcription complex, subunit 6-like	Hs.183085 AI339743	246175	316467
COBLL1	NM_014900	hsa-let-7f	COBLL1	COBL-like 1	Hs.470457 AA099593	22837	226659
COBLL1	NM_014900	hsa-miR-15a	COBLL1	COBL-like 1	Hs.470457 AA099593	22837	226659
COBLL1	NM_014900	hsa-miR-15b	COBLL1	COBL-like 1	Hs.470457 AA099593	22837	226659
COBLL1	NM_014900	hsa-miR-16	COBLL1	COBL-like 1	Hs.470457 AA099593	22837	226659
COBLL1	NM_014900	hsa-miR-195	COBLL1	COBL-like 1	Hs.470457 AA099593	22837	226659
COBLL1	NM_014900	hsa-miR-98	COBLL1	COBL-like 1	Hs.470457 AA099593	22837	226659
COL8A1	NM_001850	hsa-miR-27a	COL8A1	Collagen, type VIII, alpha 1	Hs.134830 N51859	1295	222543
CPEB4	NM_030627	hsa-let-7a	CPEB4	Cytoplasmic polyadenylation element binding protein 4	Hs.127126 N47681	80315	223349
CPEB4	NM_030627	hsa-let-7b	CPEB4	Cytoplasmic polyadenylation element binding protein 4	Hs.127126 N47681	80315	223349
CPEB4	NM_030627	hsa-let-7c	CPEB4	Cytoplasmic polyadenylation element binding protein 4	Hs.127126 N47681	80315	223349
CPEB4	NM_030627	hsa-let-7d	CPEB4	Cytoplasmic polyadenylation element binding protein 4	Hs.127126 N47681	80315	223349
CPEB4	NM_030627	hsa-let-7e	CPEB4	Cytoplasmic polyadenylation element binding protein 4	Hs.127126 N47681	80315	223349
CPEB4	NM_030627	hsa-let-7f	CPEB4	Cytoplasmic polyadenylation element binding protein 4	Hs.127126 N47681	80315	223349
CPEB4	NM_030627	hsa-miR-142-5p	CPEB4	Cytoplasmic polyadenylation element binding protein 4	Hs.127126 N47681	80315	223349
CPEB4	NM_030627	hsa-miR-19a	CPEB4	Cytoplasmic polyadenylation element binding protein 4	Hs.127126 N47681	80315	223349
CPEB4	NM_030627	hsa-miR-19b	CPEB4	Cytoplasmic polyadenylation element binding protein 4	Hs.127126 N47681	80315	223349
CPEB4	NM_030627	hsa-miR-23a	CPEB4	Cytoplasmic polyadenylation element binding protein 4	Hs.127126 N47681	80315	223349
CPEB4	NM_030627	hsa-miR-27a	CPEB4	Cytoplasmic polyadenylation element binding protein 4	Hs.127126 N47681	80315	223349
CPEB4	NM_030627	hsa-miR-30c	CPEB4	Cytoplasmic polyadenylation element binding protein 4	Hs.127126 N47681	80315	223349
CPEB4	NM_030627	hsa-miR-30e	CPEB4	Cytoplasmic polyadenylation element binding protein 4	Hs.127126 N47681	80315	223349
CPEB4	NM_030627	hsa-miR-369-3p	CPEB4	Cytoplasmic polyadenylation element binding protein 4	Hs.127126 N47681	80315	223349
CPEB4	NM_030627	hsa-miR-92a	CPEB4	Cytoplasmic polyadenylation element binding protein 4	Hs.127126 N47681	80315	223349
CPEB4	NM_030627	hsa-miR-98	CPEB4	Cytoplasmic polyadenylation element binding protein 4	Hs.127126 N47681	80315	223349
CPOX	NM_000097	hsa-miR-124	CPOX	Coproporphyrinogen oxidase	Hs.476982 AA700808	1371	105144
CPOX	NM_000097	hsa-miR-27a	CPOX	Coproporphyrinogen oxidase	Hs.476982 AA700808	1371	105144
CPOX	NM_000097	hsa-miR-369-3p	CPOX	Coproporphyrinogen oxidase	Hs.476982 AA700808	1371	105144
CRIM1	NM_016441	hsa-miR-106a	CRIM1	Cysteine-rich motor neuron 1	Hs.332847 R70508	51232	109316
CRIM1	NM_016441	hsa-miR-15a	CRIM1	Cysteine-rich motor neuron 1	Hs.332847 R70508	51232	109316
CRIM1	NM_016441	hsa-miR-15b	CRIM1	Cysteine-rich motor neuron 1	Hs.332847 R70508	51232	109316
CRIM1	NM_016441	hsa-miR-16	CRIM1	Cysteine-rich motor neuron 1	Hs.332847 R70508	51232	109316
CRIM1	NM_016441	hsa-miR-17	CRIM1	Cysteine-rich motor neuron 1	Hs.332847 R70508	51232	109316
CRIM1	NM_016441	hsa-miR-18a	CRIM1	Cysteine-rich motor neuron 1	Hs.332847 R70508	51232	109316
CRIM1	NM_016441	hsa-miR-195	CRIM1	Cysteine-rich motor neuron 1	Hs.332847 R70508	51232	109316
CRIM1	NM_016441	hsa-miR-20a	CRIM1	Cysteine-rich motor neuron 1	Hs.332847 R70508	51232	109316

CRIM1	NM_016441	hsa-miR-21	CRIM1 Cysteine-rich motor neuron 1 Hs.332847 R70508 51232 109316
CRIM1	NM_016441	hsa-miR-369-3p	CRIM1 Cysteine-rich motor neuron 1 Hs.332847 R70508 51232 109316
CRKL	NM_005207	hsa-miR-15a	CRKL V-crk sarcoma virus CT10 oncogene homolog (avian)-like Hs.5613 AI017013 1399 224179
CRKL	NM_005207	hsa-miR-15b	CRKL V-crk sarcoma virus CT10 oncogene homolog (avian)-like Hs.5613 AI017013 1399 224179
CRKL	NM_005207	hsa-miR-16	CRKL V-crk sarcoma virus CT10 oncogene homolog (avian)-like Hs.5613 AI017013 1399 224179
CRKL	NM_005207	hsa-miR-195	CRKL V-crk sarcoma virus CT10 oncogene homolog (avian)-like Hs.5613 AI017013 1399 224179
CRKL	NM_005207	hsa-miR-29a	CRKL V-crk sarcoma virus CT10 oncogene homolog (avian)-like Hs.5613 AI017013 1399 224179
CRKL	NM_005207	hsa-miR-30c	CRKL V-crk sarcoma virus CT10 oncogene homolog (avian)-like Hs.5613 AI017013 1399 224179
CRKL	NM_005207	hsa-miR-30e	CRKL V-crk sarcoma virus CT10 oncogene homolog (avian)-like Hs.5613 AI017013 1399 224179
CRLF3	NM_015986	hsa-miR-10a	CRLF3 Cytokine receptor-like factor 3 Hs.370168 R92347 51379 99711
CRLF3	NM_015986	hsa-miR-10b	CRLF3 Cytokine receptor-like factor 3 Hs.370168 R92347 51379 99711
CRLF3	NM_015986	hsa-miR-23a	CRLF3 Cytokine receptor-like factor 3 Hs.370168 R92347 51379 99711
CTSE	NM_001910	hsa-miR-29a	CTSE Cathepsin E Hs.1355 H94487 1510 119852
CTSE	NM_001910	hsa-miR-29b	CTSE Cathepsin E Hs.1355 H94487 1510 119852
CTSE	NM_001910	hsa-miR-29c	CTSE Cathepsin E Hs.1355 H94487 1510 119852
CUL1	NM_003592	hsa-miR-106a	CUL1 Cullin 1 Hs.146806 AA682674 8454 313426
CUL1	NM_003592	hsa-miR-17	CUL1 Cullin 1 Hs.146806 AA682674 8454 313426
CUL1	NM_003592	hsa-miR-18a	CUL1 Cullin 1 Hs.146806 AA682674 8454 313426
CUL1	NM_003592	hsa-miR-20a	CUL1 Cullin 1 Hs.146806 AA682674 8454 313426
CYP19A1	NM_031226	hsa-let-7a	CYP19A1 Cytochrome P450, family 19, subfamily A, polypeptide 1 Hs.511367 R63948 1588 98921
CYP19A1	NM_031226	hsa-let-7b	CYP19A1 Cytochrome P450, family 19, subfamily A, polypeptide 1 Hs.511367 R63948 1588 98921
CYP19A1	NM_031226	hsa-let-7c	CYP19A1 Cytochrome P450, family 19, subfamily A, polypeptide 1 Hs.511367 R63948 1588 98921
CYP19A1	NM_031226	hsa-let-7d	CYP19A1 Cytochrome P450, family 19, subfamily A, polypeptide 1 Hs.511367 R63948 1588 98921
CYP19A1	NM_031226	hsa-let-7e	CYP19A1 Cytochrome P450, family 19, subfamily A, polypeptide 1 Hs.511367 R63948 1588 98921
CYP19A1	NM_031226	hsa-let-7f	CYP19A1 Cytochrome P450, family 19, subfamily A, polypeptide 1 Hs.511367 R63948 1588 98921
CYP19A1	NM_031226	hsa-miR-23a	CYP19A1 Cytochrome P450, family 19, subfamily A, polypeptide 1 Hs.511367 R63948 1588 98921
CYP19A1	NM_031226	hsa-miR-98	CYP19A1 Cytochrome P450, family 19, subfamily A, polypeptide 1 Hs.511367 R63948 1588 98921
CYP1B1	NM_000104	hsa-miR-27a	CYP1B1 Cytochrome P450, family 1, subfamily B, polypeptide 1 Hs.154654 N21019 1545 307443
CYP4F2	NM_001082	hsa-let-7a	AI364951 309221
CYP4F2	NM_001082	hsa-let-7b	AI364951 309221
CYP4F2	NM_001082	hsa-let-7c	AI364951 309221
CYP4F2	NM_001082	hsa-let-7d	AI364951 309221
CYP4F2	NM_001082	hsa-let-7e	AI364951 309221
CYP4F2	NM_001082	hsa-let-7f	AI364951 309221
CYP4F2	NM_001082	hsa-miR-98	AI364951 309221
DCBLD2	NM_080927	hsa-miR-106a	DCBLD2 Discoidin, CUB and LCCL domain containing 2 Hs.203691 AA431438 131566 115972
DCBLD2	NM_080927	hsa-miR-15a	DCBLD2 Discoidin, CUB and LCCL domain containing 2 Hs.203691 AA431438 131566 115972
DCBLD2	NM_080927	hsa-miR-15b	DCBLD2 Discoidin, CUB and LCCL domain containing 2 Hs.203691 AA431438 131566 115972
DCBLD2	NM_080927	hsa-miR-16	DCBLD2 Discoidin, CUB and LCCL domain containing 2 Hs.203691 AA431438 131566 115972
DCBLD2	NM_080927	hsa-miR-17	DCBLD2 Discoidin, CUB and LCCL domain containing 2 Hs.203691 AA431438 131566 115972
DCBLD2	NM_080927	hsa-miR-195	DCBLD2 Discoidin, CUB and LCCL domain containing 2 Hs.203691 AA431438 131566 115972
DCBLD2	NM_080927	hsa-miR-19a	DCBLD2 Discoidin, CUB and LCCL domain containing 2 Hs.203691 AA431438 131566 115972
DCBLD2	NM_080927	hsa-miR-19b	DCBLD2 Discoidin, CUB and LCCL domain containing 2 Hs.203691 AA431438 131566 115972
DCBLD2	NM_080927	hsa-miR-20a	DCBLD2 Discoidin, CUB and LCCL domain containing 2 Hs.203691 AA431438 131566 115972
DCBLD2	NM_080927	hsa-miR-23a	DCBLD2 Discoidin, CUB and LCCL domain containing 2 Hs.203691 AA431438 131566 115972
DCP2	NM_152624	hsa-miR-27a	DCP2 DCP2 decapping enzyme homolog (S. cerevisiae) Hs.443875 AI338325 167227 315232
DCP2	NM_152624	hsa-miR-29a	DCP2 DCP2 decapping enzyme homolog (S. cerevisiae) Hs.443875 AI338325 167227 315232
DCP2	NM_152624	hsa-miR-29b	DCP2 DCP2 decapping enzyme homolog (S. cerevisiae) Hs.443875 AI338325 167227 315232
DCP2	NM_152624	hsa-miR-29c	DCP2 DCP2 decapping enzyme homolog (S. cerevisiae) Hs.443875 AI338325 167227 315232
DCP2	NM_152624	hsa-miR-30c	DCP2 DCP2 decapping enzyme homolog (S. cerevisiae) Hs.443875 AI338325 167227 315232
DCP2	NM_152624	hsa-miR-30e	DCP2 DCP2 decapping enzyme homolog (S. cerevisiae) Hs.443875 AI338325 167227 315232
DDEF1	NM_018482	hsa-let-7a	DDEF1 Development and differentiation enhancing factor 1 Hs.106015 N72878 50807 107710
DDEF1	NM_018482	hsa-let-7b	DDEF1 Development and differentiation enhancing factor 1 Hs.106015 N72878 50807 107710

DDEF1	NM_018482	hsa-let-7c	DDEF1 Development and differentiation enhancing factor 1 Hs.106015 N72878 50807 107710
DDEF1	NM_018482	hsa-let-7d	DDEF1 Development and differentiation enhancing factor 1 Hs.106015 N72878 50807 107710
DDEF1	NM_018482	hsa-let-7e	DDEF1 Development and differentiation enhancing factor 1 Hs.106015 N72878 50807 107710
DDEF1	NM_018482	hsa-let-7f	DDEF1 Development and differentiation enhancing factor 1 Hs.106015 N72878 50807 107710
DDEF1	NM_018482	hsa-miR-30c	DDEF1 Development and differentiation enhancing factor 1 Hs.106015 N72878 50807 107710
DDEF1	NM_018482	hsa-miR-30e	DDEF1 Development and differentiation enhancing factor 1 Hs.106015 N72878 50807 107710
DDEF1	NM_018482	hsa-miR-98	DDEF1 Development and differentiation enhancing factor 1 Hs.106015 N72878 50807 107710
DHX36	NM_020865	hsa-miR-30c	DHX36 DEAH (Asp-Glu-Ala-His) box polypeptide 36 Hs.446270 AA169411 170506 116551
DHX36	NM_020865	hsa-miR-30e	DHX36 DEAH (Asp-Glu-Ala-His) box polypeptide 36 Hs.446270 AA169411 170506 116551
DNM2	NM_004945	hsa-miR-106a	DNM2 Dynamin 2 Hs.211463 AA780897 1785 108339
DNM2	NM_004945	hsa-miR-124	DNM2 Dynamin 2 Hs.211463 AA780897 1785 108339
DNM2	NM_004945	hsa-miR-17	DNM2 Dynamin 2 Hs.211463 AA780897 1785 108339
DNM2	NM_004945	hsa-miR-20a	DNM2 Dynamin 2 Hs.211463 AA780897 1785 108339
DSP	NM_004415	hsa-miR-142-5p	DSP Desmoplakin Hs.519873 H90899 1832 114333
DUSP16	NM_030640	hsa-let-7a	DUSP16 Dual specificity phosphatase 16 Hs.536535 N40952 80824 115398
DUSP16	NM_030640	hsa-let-7b	DUSP16 Dual specificity phosphatase 16 Hs.536535 N40952 80824 115398
DUSP16	NM_030640	hsa-let-7c	DUSP16 Dual specificity phosphatase 16 Hs.536535 N40952 80824 115398
DUSP16	NM_030640	hsa-let-7d	DUSP16 Dual specificity phosphatase 16 Hs.536535 N40952 80824 115398
DUSP16	NM_030640	hsa-let-7e	DUSP16 Dual specificity phosphatase 16 Hs.536535 N40952 80824 115398
DUSP16	NM_030640	hsa-let-7f	DUSP16 Dual specificity phosphatase 16 Hs.536535 N40952 80824 115398
DUSP16	NM_030640	hsa-miR-18a	DUSP16 Dual specificity phosphatase 16 Hs.536535 N40952 80824 115398
DUSP16	NM_030640	hsa-miR-29c	DUSP16 Dual specificity phosphatase 16 Hs.536535 N40952 80824 115398
DUSP16	NM_030640	hsa-miR-98	DUSP16 Dual specificity phosphatase 16 Hs.536535 N40952 80824 115398
EDG1	NM_001400	hsa-miR-106a	EDG1 Endothelial differentiation, sphingolipid G-protein-coupled receptor, 1 Hs.154210 N93476 EDG-1=endothelial differentiation protein=putative G-protein 1901 161545
EDG1	NM_001400	hsa-miR-155	EDG1 Endothelial differentiation, sphingolipid G-protein-coupled receptor, 1 Hs.154210 N93476 EDG-1=endothelial differentiation protein=putative G-protein 1901 161545
EDG1	NM_001400	hsa-miR-17	EDG1 Endothelial differentiation, sphingolipid G-protein-coupled receptor, 1 Hs.154210 N93476 EDG-1=endothelial differentiation protein=putative G-protein 1901 161545
EDG1	NM_001400	hsa-miR-19a	EDG1 Endothelial differentiation, sphingolipid G-protein-coupled receptor, 1 Hs.154210 N93476 EDG-1=endothelial differentiation protein=putative G-protein 1901 161545
EDG1	NM_001400	hsa-miR-19b	EDG1 Endothelial differentiation, sphingolipid G-protein-coupled receptor, 1 Hs.154210 N93476 EDG-1=endothelial differentiation protein=putative G-protein 1901 161545
EDG1	NM_001400	hsa-miR-20a	EDG1 Endothelial differentiation, sphingolipid G-protein-coupled receptor, 1 Hs.154210 N93476 EDG-1=endothelial differentiation protein=putative G-protein 1901 161545
EDG1	NM_001400	hsa-miR-92a	EDG1 Endothelial differentiation, sphingolipid G-protein-coupled receptor, 1 Hs.154210 N93476 EDG-1=endothelial differentiation protein=putative G-protein 1901 161545
EDIL3	NM_005711	hsa-miR-21	EDIL3 EGF-like repeats and discoidin I-like domains 3 Hs.482730 T57803 10085 113235
EGFR	NM_005228	hsa-miR-27a	EGFR Epidermal growth factor receptor (erythroblastic leukemia viral (v-erb-b) oncogene homolog, avian) Hs.488293 T58652 1956 108730
ELOVL2	NM_017770	hsa-miR-124	ELOVL2 Elongation of very long chain fatty acids (FEN1/Elo2, SUR4/Elo3, yeast)-like 2 Hs.408557 H06525 54898 98862
ELOVL5	NM_021814	hsa-miR-124	ELOVL5 ELOVL family member 5, elongation of long chain fatty acids (FEN1/Elo2, SUR4/Elo3-like, yeast) Hs.520189 AA992103 60481 310009
ELOVL5	NM_021814	hsa-miR-19a	ELOVL5 ELOVL family member 5, elongation of long chain fatty acids (FEN1/Elo2, SUR4/Elo3-like, yeast) Hs.520189 AA992103 60481 310009
ELOVL5	NM_021814	hsa-miR-19b	ELOVL5 ELOVL family member 5, elongation of long chain fatty acids (FEN1/Elo2, SUR4/Elo3-like, yeast) Hs.520189 AA992103 60481 310009
ELOVL5	NM_021814	hsa-miR-30c	ELOVL5 ELOVL family member 5, elongation of long chain fatty acids (FEN1/Elo2, SUR4/Elo3-like, yeast) Hs.520189 AA992103 60481 310009
ELOVL5	NM_021814	hsa-miR-30e	ELOVL5 ELOVL family member 5, elongation of long chain fatty acids (FEN1/Elo2, SUR4/Elo3-like, yeast) Hs.520189 AA992103 60481 310009
ELOVL6	NM_024090	hsa-miR-10a	ELOVL6 ELOVL family member 6, elongation of long chain fatty acids (FEN1/Elo2, SUR4/Elo3-like, yeast) Hs.412939 T98354 79071 118689
ELOVL6	NM_024090	hsa-miR-10b	ELOVL6 ELOVL family member 6, elongation of long chain fatty acids (FEN1/Elo2, SUR4/Elo3-like, yeast) Hs.412939 T98354 79071 118689
ELOVL6	NM_024090	hsa-miR-23a	ELOVL6 ELOVL family member 6, elongation of long chain fatty acids (FEN1/Elo2, SUR4/Elo3-like, yeast) Hs.412939 T98354 79071 118689
ELOVL6	NM_024090	hsa-miR-369-3p	ELOVL6 ELOVL family member 6, elongation of long chain fatty acids (FEN1/Elo2, SUR4/Elo3-like, yeast) Hs.412939 T98354 79071 118689
ENTPD5	NM_001249	hsa-miR-23a	ENTPD5 Ectonucleoside triphosphate diphosphohydrolase 5 Hs.131555 A1017442 957 310456
EPB41	NM_203343	hsa-miR-30c	EPB41 Erythrocyte membrane protein band 4.1 (elliptocytosis 1, RH-linked) Hs.175437 N93865 2035 115910
EPB41	NM_203343	hsa-miR-30e	EPB41 Erythrocyte membrane protein band 4.1 (elliptocytosis 1, RH-linked) Hs.175437 N93865 2035 115910
EPB49	NM_001978	hsa-miR-15a	EPB49 Erythrocyte membrane protein band 4.9 (dematin) Hs.274122 N76836 2039 103631

EPB49	NM 001978	hsa-miR-15b	EPB49 Erythrocyte membrane protein band 4.9 (dematin) Hs.274122 N76836 2039 103631
EPB49	NM 001978	hsa-miR-16	EPB49 Erythrocyte membrane protein band 4.9 (dematin) Hs.274122 N76836 2039 103631
EPB49	NM 001978	hsa-miR-195	EPB49 Erythrocyte membrane protein band 4.9 (dematin) Hs.274122 N76836 2039 103631
EPC2	NM 015630	hsa-miR-19a	EPC2 Enhancer of polycomb homolog 2 (Drosophila) Hs.23270 H05925 26122 119650
EPC2	NM 015630	hsa-miR-19b	EPC2 Enhancer of polycomb homolog 2 (Drosophila) Hs.23270 H05925 26122 119650
EPC2	NM 015630	hsa-miR-30c	EPC2 Enhancer of polycomb homolog 2 (Drosophila) Hs.23270 H05925 26122 119650
EPC2	NM 015630	hsa-miR-30e	EPC2 Enhancer of polycomb homolog 2 (Drosophila) Hs.23270 H05925 26122 119650
EPC2	NM 015630	hsa-miR-369-3p	EPC2 Enhancer of polycomb homolog 2 (Drosophila) Hs.23270 H05925 26122 119650
EPC2	NM 015630	hsa-miR-92a	EPC2 Enhancer of polycomb homolog 2 (Drosophila) Hs.23270 H05925 26122 119650
EPHA2	NM 004431	hsa-miR-124	EPHA2 EPH receptor A2 Hs.171596 H84480 1969 118742
EPS8	NM 004447	hsa-miR-124	EPS8 Epidermal growth factor receptor pathway substrate 8 Hs.26139 H13622 epidermal growth factor receptor kinase substrate (Eps8) 2059 99229
EPS8	NM 004447	hsa-miR-92a	EPS8 Epidermal growth factor receptor pathway substrate 8 Hs.26139 H13622 epidermal growth factor receptor kinase substrate (Eps8) 2059 99229
EPST11	NM 001002264	hsa-miR-10a	EPST11 Epithelial stromal interaction 1 (breast) Hs.546467 A1286247 94240 316724
EPST11	NM 001002264	hsa-miR-10b	EPST11 Epithelial stromal interaction 1 (breast) Hs.546467 A1286247 94240 316724
FAM104A	NM 032837	hsa-let-7a	FLJ14775 Hypothetical protein FLJ14775 Hs.103555 T58729 84923 117123
FAM104A	NM 032837	hsa-let-7b	FLJ14775 Hypothetical protein FLJ14775 Hs.103555 T58729 84923 117123
FAM104A	NM 032837	hsa-let-7c	FLJ14775 Hypothetical protein FLJ14775 Hs.103555 T58729 84923 117123
FAM104A	NM 032837	hsa-let-7d	FLJ14775 Hypothetical protein FLJ14775 Hs.103555 T58729 84923 117123
FAM104A	NM 032837	hsa-let-7e	FLJ14775 Hypothetical protein FLJ14775 Hs.103555 T58729 84923 117123
FAM104A	NM 032837	hsa-let-7f	FLJ14775 Hypothetical protein FLJ14775 Hs.103555 T58729 84923 117123
FAM104A	NM 032837	hsa-miR-124	FLJ14775 Hypothetical protein FLJ14775 Hs.103555 T58729 84923 117123
FAM104A	NM 032837	hsa-miR-19a	FLJ14775 Hypothetical protein FLJ14775 Hs.103555 T58729 84923 117123
FAM104A	NM 032837	hsa-miR-19b	FLJ14775 Hypothetical protein FLJ14775 Hs.103555 T58729 84923 117123
FAM104A	NM 032837	hsa-miR-27a	FLJ14775 Hypothetical protein FLJ14775 Hs.103555 T58729 84923 117123
FAM104A	NM 032837	hsa-miR-98	FLJ14775 Hypothetical protein FLJ14775 Hs.103555 T58729 84923 117123
FAM46C	NM 017709	hsa-miR-106a	FAM46C Family with sequence similarity 46, member C Hs.356216 AA056118 54855 107808
FAM46C	NM 017709	hsa-miR-17	FAM46C Family with sequence similarity 46, member C Hs.356216 AA056118 54855 107808
FAM46C	NM 017709	hsa-miR-19a	FAM46C Family with sequence similarity 46, member C Hs.356216 AA056118 54855 107808
FAM46C	NM 017709	hsa-miR-19b	FAM46C Family with sequence similarity 46, member C Hs.356216 AA056118 54855 107808
FAM46C	NM 017709	hsa-miR-20a	FAM46C Family with sequence similarity 46, member C Hs.356216 AA056118 54855 107808
FAM46C	NM 017709	hsa-miR-21	FAM46C Family with sequence similarity 46, member C Hs.356216 AA056118 54855 107808
FAM46C	NM 017709	hsa-miR-30c	FAM46C Family with sequence similarity 46, member C Hs.356216 AA056118 54855 107808
FAM46C	NM 017709	hsa-miR-30e	FAM46C Family with sequence similarity 46, member C Hs.356216 AA056118 54855 107808
FBXO30	NM 032145	hsa-let-7a	AA719040 223350
FBXO30	NM 032145	hsa-let-7b	AA719040 223350
FBXO30	NM 032145	hsa-let-7c	AA719040 223350
FBXO30	NM 032145	hsa-let-7d	AA719040 223350
FBXO30	NM 032145	hsa-let-7e	AA719040 223350
FBXO30	NM 032145	hsa-let-7f	AA719040 223350
FBXO30	NM 032145	hsa-miR-124	AA719040 223350
FBXO30	NM 032145	hsa-miR-142-5p	AA719040 223350
FBXO30	NM 032145	hsa-miR-98	AA719040 223350
FEM1B	NM 015322	hsa-miR-29a	FEM1B Fem-1 homolog b (C. elegans) Hs.362733 H82273 10116 114606
FEM1B	NM 015322	hsa-miR-29b	FEM1B Fem-1 homolog b (C. elegans) Hs.362733 H82273 10116 114606
FEM1B	NM 015322	hsa-miR-29c	FEM1B Fem-1 homolog b (C. elegans) Hs.362733 H82273 10116 114606
FGA	NM 000508	hsa-miR-18a	FGA Fibrinogen, A alpha polypeptide Hs.351593 AA011414 2243 116021
FGA	NM 000508	hsa-miR-29a	FGA Fibrinogen, A alpha polypeptide Hs.351593 AA011414 2243 116021
FGA	NM 000508	hsa-miR-29b	FGA Fibrinogen, A alpha polypeptide Hs.351593 AA011414 2243 116021
FGA	NM 000508	hsa-miR-29c	FGA Fibrinogen, A alpha polypeptide Hs.351593 AA011414 2243 116021
FGFR1	NM 000604	hsa-miR-16	FGFR1 Fibroblast growth factor receptor 1 (fms-related tyrosine kinase 2, Pfeiffer syndrome) Hs.264887 R54610 FGFR1=Fibroblast growth factor receptor 1 2260 101364
FGFR1	NM 023109	hsa-miR-27a	FGFR1 Fibroblast growth factor receptor 1 (fms-related tyrosine kinase 2, Pfeiffer syndrome) Hs.264887 R54610 FGFR1=Fibroblast growth factor receptor 1 2260 101364

FOXA1	NM 004496	hsa-miR-106a	FOXA1 Forkhead box A1 Hs.163484 T74639 3169 99032
FOXA1	NM 004496	hsa-miR-142-5p	FOXA1 Forkhead box A1 Hs.163484 T74639 3169 99032
FOXA1	NM 004496	hsa-miR-17	FOXA1 Forkhead box A1 Hs.163484 T74639 3169 99032
FOXA1	NM 004496	hsa-miR-20a	FOXA1 Forkhead box A1 Hs.163484 T74639 3169 99032
FOXA1	NM 004496	hsa-miR-23a	FOXA1 Forkhead box A1 Hs.163484 T74639 3169 99032
FOXA1	NM 004496	hsa-miR-30c	FOXA1 Forkhead box A1 Hs.163484 T74639 3169 99032
FOXA1	NM 004496	hsa-miR-30e	FOXA1 Forkhead box A1 Hs.163484 T74639 3169 99032
FOXO1A	NM 002015	hsa-miR-15a	FOXO1A Forkhead box O1A (rhabdomyosarcoma) Hs.370666 AA448277 2308 117043
FOXO1A	NM 002015	hsa-miR-17	FOXO1A Forkhead box O1A (rhabdomyosarcoma) Hs.370666 AA448277 2308 117043
FOXO1A	NM 002015	hsa-miR-27a	FOXO1A Forkhead box O1A (rhabdomyosarcoma) Hs.370666 AA448277 2308 117043
FOXO1A	NM 002015	hsa-miR-369-3p	FOXO1A Forkhead box O1A (rhabdomyosarcoma) Hs.370666 AA448277 2308 117043
FOXO3A	NM 201559	hsa-miR-10a	FOXO3A Forkhead box O3A Hs.220950 AA465236 2309 117277
FOXO3A	NM 201559	hsa-miR-10b	FOXO3A Forkhead box O3A Hs.220950 AA465236 2309 117277
FOXO3A	NM 201559	hsa-miR-142-5p	FOXO3A Forkhead box O3A Hs.220950 AA465236 2309 117277
FOXO3A	NM 201559	hsa-miR-155	FOXO3A Forkhead box O3A Hs.220950 AA465236 2309 117277
FOXO3A	NM 201559	hsa-miR-23a	FOXO3A Forkhead box O3A Hs.220950 AA465236 2309 117277
FOXO3A	NM 201559	hsa-miR-29a	FOXO3A Forkhead box O3A Hs.220950 AA465236 2309 117277
FOXO3A	NM 201559	hsa-miR-30c	FOXO3A Forkhead box O3A Hs.220950 AA465236 2309 117277
FOXO3A	NM 201559	hsa-miR-30e	FOXO3A Forkhead box O3A Hs.220950 AA465236 2309 117277
FZD6	NM 003506	hsa-miR-369-3p	FZD6 Frizzled homolog 6 (Drosophila) Hs.292464 T68333 8323 104207
FZD6	NM 003506	hsa-miR-92a	FZD6 Frizzled homolog 6 (Drosophila) Hs.292464 T68333 8323 104207
GAB1	NM 207123	hsa-miR-106a	GAB1 GRB2-associated binding protein 1 Hs.80720 H47315 2549 119933
GAB1	NM 207123	hsa-miR-142-5p	GAB1 GRB2-associated binding protein 1 Hs.80720 H47315 2549 119933
GAB1	NM 207123	hsa-miR-17	GAB1 GRB2-associated binding protein 1 Hs.80720 H47315 2549 119933
GAB1	NM 207123	hsa-miR-18a	GAB1 GRB2-associated binding protein 1 Hs.80720 H47315 2549 119933
GAB1	NM 207123	hsa-miR-20a	GAB1 GRB2-associated binding protein 1 Hs.80720 H47315 2549 119933
GAB1	NM 207123	hsa-miR-29a	GAB1 GRB2-associated binding protein 1 Hs.80720 H47315 2549 119933
GAB1	NM 207123	hsa-miR-29b	GAB1 GRB2-associated binding protein 1 Hs.80720 H47315 2549 119933
GAB1	NM 207123	hsa-miR-29c	GAB1 GRB2-associated binding protein 1 Hs.80720 H47315 2549 119933
GAB1	NM 207123	hsa-miR-30e	GAB1 GRB2-associated binding protein 1 Hs.80720 H47315 2549 119933
GAB1	NM 207123	hsa-miR-369-3p	GAB1 GRB2-associated binding protein 1 Hs.80720 H47315 2549 119933
GAB1	NM 207123	hsa-miR-98	GAB1 GRB2-associated binding protein 1 Hs.80720 H47315 2549 119933
GABARAPL2	NM 007285	hsa-miR-18a	GABARAPL2 GABA(A) receptor-associated protein-like 2 Hs.461379 AA455108 11345 98390
GALNT3	NM 004482	hsa-miR-27a	GALNT3 UDP-N-acetyl-alpha-D-galactosamine:polypeptide N-acetylgalactosaminyltransferase 3 (GalNAc-T3) Hs.170986 H13688 2591 107949
GALNT3	NM 004482	hsa-miR-30c	GALNT3 UDP-N-acetyl-alpha-D-galactosamine:polypeptide N-acetylgalactosaminyltransferase 3 (GalNAc-T3) Hs.170986 H13688 2591 107949
GALNT3	NM 004482	hsa-miR-30e	GALNT3 UDP-N-acetyl-alpha-D-galactosamine:polypeptide N-acetylgalactosaminyltransferase 3 (GalNAc-T3) Hs.170986 H13688 2591 107949
GBP1	NM 002053	hsa-miR-21	GBP1 Guanylate binding protein 1, interferon-inducible, 67kDa Hs.62661 AA280278 Guanylate binding protein 1, interferon-inducible, 67kD 2633 162324
GCA	NM 012198	hsa-miR-27a	GCA Grancalcin, EF-hand calcium binding protein Hs.377894 R19937 grancalcin=EF-hand calcium-binding protein abundant in neutr 25801 119966
GCLC	NM 001498	hsa-miR-18a	GCLC Glutamate-cysteine ligase, catalytic subunit Hs.271264 N45129 2729 220986
GCLC	NM 001498	hsa-miR-195	GCLC Glutamate-cysteine ligase, catalytic subunit Hs.271264 N45129 2729 220986
GCLC	NM 001498	hsa-miR-30c	GCLC Glutamate-cysteine ligase, catalytic subunit Hs.271264 N45129 2729 220986
GLMN	NM 007070	hsa-let-7a	GLMN Glomulin, FKBP associated protein Hs.49105 N95144 11146 107353
GLMN	NM 007070	hsa-let-7b	GLMN Glomulin, FKBP associated protein Hs.49105 N95144 11146 107353
GLMN	NM 007070	hsa-let-7c	GLMN Glomulin, FKBP associated protein Hs.49105 N95144 11146 107353
GLMN	NM 007070	hsa-let-7d	GLMN Glomulin, FKBP associated protein Hs.49105 N95144 11146 107353
GLMN	NM 007070	hsa-let-7e	GLMN Glomulin, FKBP associated protein Hs.49105 N95144 11146 107353
GLMN	NM 007070	hsa-let-7f	GLMN Glomulin, FKBP associated protein Hs.49105 N95144 11146 107353
GLMN	NM 053274	hsa-miR-124	GLMN Glomulin, FKBP associated protein Hs.49105 N95144 11146 107353
GLMN	NM 007070	hsa-miR-98	GLMN Glomulin, FKBP associated protein Hs.49105 N95144 11146 107353
GLS	NM 014905	hsa-miR-23a	GLS Glutaminase Hs.116448 R89349 2744 225594
GLT25D2	NM 015101	hsa-miR-27a	GLT25D2 Glycosyltransferase 25 domain containing 2 Hs.387995 H08862 23127 120717
GNAI1	NM 002069	hsa-miR-124	GNAI1 Guanine nucleotide binding protein (G protein), alpha inhibiting activity polypeptide 1 Hs.134587 W46769 2770 221953
GNAI1	NM 002069	hsa-miR-19a	GNAI1 Guanine nucleotide binding protein (G protein), alpha inhibiting activity polypeptide 1 Hs.134587 W46769 2770 221953

GNAI1	NM_002069	hsa-miR-19b	GNAI1 Guanine nucleotide binding protein (G protein), alpha inhibiting activity polypeptide 1 Hs.134587 W46769 2770 221953
GNG11	NM_004126	hsa-miR-15a	GNG11 Guanine nucleotide binding protein (G protein), gamma 11 Hs.83381 AA999901 2791 220964
GNG11	NM_004126	hsa-miR-15b	GNG11 Guanine nucleotide binding protein (G protein), gamma 11 Hs.83381 AA999901 2791 220964
GNG2	NM_053064	hsa-miR-18a	GNG2 Guanine nucleotide binding protein (G protein), gamma 2 Hs.187772 T80856 54331 112310
GNG2	NM_053064	hsa-miR-21	GNG2 Guanine nucleotide binding protein (G protein), gamma 2 Hs.187772 T80856 54331 112310
GNG2	NM_053064	hsa-miR-30e	GNG2 Guanine nucleotide binding protein (G protein), gamma 2 Hs.187772 T80856 54331 112310
GOLGA4	NM_002078	hsa-miR-16	AW058321 330343
GOLGA4	NM_002078	hsa-miR-19a	AW058321 330343
GOLGA4	NM_002078	hsa-miR-30c	AW058321 330343
GOLGA4	NM_002078	hsa-miR-30e	AW058321 330343
GOLGA4	NM_002078	hsa-miR-92a	AW058321 330343
GSPT1	NM_002094	hsa-let-7a	GSPT1 G1 to S phase transition 1 Hs.528780 AA962601 2935 308491
GSPT1	NM_002094	hsa-let-7b	GSPT1 G1 to S phase transition 1 Hs.528780 AA962601 2935 308491
GSPT1	NM_002094	hsa-let-7e	GSPT1 G1 to S phase transition 1 Hs.528780 AA962601 2935 308491
GSPT1	NM_002094	hsa-let-7f	GSPT1 G1 to S phase transition 1 Hs.528780 AA962601 2935 308491
GSPT1	NM_002094	hsa-miR-27a	GSPT1 G1 to S phase transition 1 Hs.528780 AA962601 2935 308491
H1FO	NM_005318	hsa-miR-19a	H1FO H1 histone family, member 0 Hs.226117 W69399 3005 119185
HAGH	NM_005326	hsa-miR-92a	HAGH Hydroxyacylglutathione hydrolase Hs.157394 AA931043 3029 309568
HECA	NM_016217	hsa-miR-106a	HECA Headcase homolog (Drosophila) Hs.197644 AA115382 51696 111756
HECA	NM_016217	hsa-miR-17	HECA Headcase homolog (Drosophila) Hs.197644 AA115382 51696 111756
HECA	NM_016217	hsa-miR-20a	HECA Headcase homolog (Drosophila) Hs.197644 AA115382 51696 111756
HMBS	NM_000190	hsa-miR-10a	HMBS Hydroxymethylbilane synthase Hs.82609 R06263 3145 119285
HMBS	NM_000190	hsa-miR-10b	HMBS Hydroxymethylbilane synthase Hs.82609 R06263 3145 119285
HMBS	NM_000190	hsa-miR-124	HMBS Hydroxymethylbilane synthase Hs.82609 R06263 3145 119285
HMBS	NM_000190	hsa-miR-15a	HMBS Hydroxymethylbilane synthase Hs.82609 R06263 3145 119285
HMBS	NM_000190	hsa-miR-15b	HMBS Hydroxymethylbilane synthase Hs.82609 R06263 3145 119285
HMBS	NM_000190	hsa-miR-16	HMBS Hydroxymethylbilane synthase Hs.82609 R06263 3145 119285
HMBS	NM_000190	hsa-miR-195	HMBS Hydroxymethylbilane synthase Hs.82609 R06263 3145 119285
HS3ST3B1	NM_006041	hsa-miR-124	HS3ST3B1 Heparan sulfate (glucosamine) 3-O-sulfotransferase 3B1 Hs.48384 W95040 9953 102818
HS3ST3B1	NM_006041	hsa-miR-29a	HS3ST3B1 Heparan sulfate (glucosamine) 3-O-sulfotransferase 3B1 Hs.48384 W95040 9953 102818
HS3ST3B1	NM_006041	hsa-miR-29b	HS3ST3B1 Heparan sulfate (glucosamine) 3-O-sulfotransferase 3B1 Hs.48384 W95040 9953 102818
HS3ST3B1	NM_006041	hsa-miR-29c	HS3ST3B1 Heparan sulfate (glucosamine) 3-O-sulfotransferase 3B1 Hs.48384 W95040 9953 102818
HSDL1	NM_031463	hsa-miR-27a	HSDL1 Hydroxysteroid dehydrogenase like 1 Hs.499483 H17130 83693 220509
ID2	NM_002166	hsa-miR-142-5p	ID2 Inhibitor of DNA binding 2, dominant negative helix-loop-helix protein Hs.180919 H89935 3398 318245
ID2	NM_002166	hsa-miR-19a	ID2 Inhibitor of DNA binding 2, dominant negative helix-loop-helix protein Hs.180919 H89935 3398 318245
ID2	NM_002166	hsa-miR-27a	ID2 Inhibitor of DNA binding 2, dominant negative helix-loop-helix protein Hs.180919 H89935 3398 318245
ID2	NM_002166	hsa-miR-30e	ID2 Inhibitor of DNA binding 2, dominant negative helix-loop-helix protein Hs.180919 H89935 3398 318245
IFIT5	NM_012420	hsa-miR-106a	IFIT5 Interferon-induced protein with tetratricopeptide repeats 5 Hs.252839 N98563 retinoic acid- and interferon-inducible 58K protein RI58 24138 109270
IFIT5	NM_012420	hsa-miR-17	IFIT5 Interferon-induced protein with tetratricopeptide repeats 5 Hs.252839 N98563 retinoic acid- and interferon-inducible 58K protein RI58 24138 109270
IFIT5	NM_012420	hsa-miR-20a	IFIT5 Interferon-induced protein with tetratricopeptide repeats 5 Hs.252839 N98563 retinoic acid- and interferon-inducible 58K protein RI58 24138 109270
IFITM1	NM_003641	hsa-miR-19a	IFITM1 Interferon induced transmembrane protein 1 (9-27) Hs.458414 AA058323 Interferon-inducible protein 9-27=interferon-induced 17kDa 8519 110694
IFITM1	NM_003641	hsa-miR-19b	IFITM1 Interferon induced transmembrane protein 1 (9-27) Hs.458414 AA058323 Interferon-inducible protein 9-27=interferon-induced 17kDa 8519 110694
IFITM1	NM_003641	hsa-miR-92a	IFITM1 Interferon induced transmembrane protein 1 (9-27) Hs.458414 AA058323 Interferon-inducible protein 9-27=interferon-induced 17kDa 8519 110694
IGFBP7	NM_001553	hsa-miR-21	POLR2B Polymerase (RNA) II (DNA directed) polypeptide B, 140kDa Hs.479814 AA704965 5431 222350
IL13RA2	NM_000640	hsa-miR-21	IL13RA2 Interleukin 13 receptor, alpha 2 Hs.336046 R52795 IL-13 receptor 3598 109672
IL18RAP	NM_003853	hsa-let-7a	IL18RAP Interleukin 18 receptor accessory protein Hs.158315 A1307682 8807 319041
IL18RAP	NM_003853	hsa-let-7b	IL18RAP Interleukin 18 receptor accessory protein Hs.158315 A1307682 8807 319041
IL18RAP	NM_003853	hsa-let-7c	IL18RAP Interleukin 18 receptor accessory protein Hs.158315 A1307682 8807 319041
IL18RAP	NM_003853	hsa-let-7e	IL18RAP Interleukin 18 receptor accessory protein Hs.158315 A1307682 8807 319041
IL18RAP	NM_003853	hsa-let-7f	IL18RAP Interleukin 18 receptor accessory protein Hs.158315 A1307682 8807 319041
IL18RAP	NM_003853	hsa-miR-124	IL18RAP Interleukin 18 receptor accessory protein Hs.158315 A1307682 8807 319041
IL18RAP	NM_003853	hsa-miR-142-5p	IL18RAP Interleukin 18 receptor accessory protein Hs.158315 A1307682 8807 319041
IL18RAP	NM_003853	hsa-miR-27a	IL18RAP Interleukin 18 receptor accessory protein Hs.158315 A1307682 8807 319041

IL18RAP	NM_003853	hsa-miR-369-3p	IL18RAP Interleukin 18 receptor accessory protein Hs.158315 AI307682 8807 319041
IL18RAP	NM_003853	hsa-miR-98	IL18RAP Interleukin 18 receptor accessory protein Hs.158315 AI307682 8807 319041
IL6ST	NM_175767	hsa-miR-142-5p	Transcribed locus, moderately similar to NP_002175.2 interleukin 6 signal transducer isoform 1 precursor; membrane glycoprotein gp130; oncostatin M receptor; CD130 antigen; interleukin receptor beta c Hs.318938 AA406546 116038
ILF3	NM_004516	hsa-miR-19a	CTL2 CTL2 protein Hs.515134 R39578 57153 100181
ILF3	NM_004516	hsa-miR-19b	CTL2 CTL2 protein Hs.515134 R39578 57153 100181
IREB2	NM_004136	hsa-miR-19a	IREB2 Iron-responsive element binding protein 2 Hs.436031 T49556 3658 120644
IREB2	NM_004136	hsa-miR-19b	IREB2 Iron-responsive element binding protein 2 Hs.436031 T49556 3658 120644
IREB2	NM_004136	hsa-miR-27a	IREB2 Iron-responsive element binding protein 2 Hs.436031 T49556 3658 120644
IREB2	NM_004136	hsa-miR-29a	IREB2 Iron-responsive element binding protein 2 Hs.436031 T49556 3658 120644
IREB2	NM_004136	hsa-miR-29b	IREB2 Iron-responsive element binding protein 2 Hs.436031 T49556 3658 120644
IREB2	NM_004136	hsa-miR-29c	IREB2 Iron-responsive element binding protein 2 Hs.436031 T49556 3658 120644
IREB2	NM_004136	hsa-miR-92a	IREB2 Iron-responsive element binding protein 2 Hs.436031 T49556 3658 120644
IRF2	NM_002199	hsa-miR-18a	IRF2 Interferon regulatory factor 2 Hs.374097 AA416883 IRF-2=interferon regulatory factor-2 3660 119240
IRF2	NM_002199	hsa-miR-20a	IRF2 Interferon regulatory factor 2 Hs.374097 AA416883 IRF-2=interferon regulatory factor-2 3660 119240
IRF2	NM_002199	hsa-miR-23a	IRF2 Interferon regulatory factor 2 Hs.374097 AA416883 IRF-2=interferon regulatory factor-2 3660 119240
IRF2	NM_002199	hsa-miR-27a	IRF2 Interferon regulatory factor 2 Hs.374097 AA416883 IRF-2=interferon regulatory factor-2 3660 119240
ITGA6	NM_000210	hsa-miR-142-5p	ITGA6 Integrin, alpha 6 Hs.133397 H06635 3655 101822
ITGA6	NM_000210	hsa-miR-19a	ITGA6 Integrin, alpha 6 Hs.133397 H06635 3655 101822
ITGA6	NM_000210	hsa-miR-19b	ITGA6 Integrin, alpha 6 Hs.133397 H06635 3655 101822
ITGA6	NM_000210	hsa-miR-23a	ITGA6 Integrin, alpha 6 Hs.133397 H06635 3655 101822
ITGA6	NM_000210	hsa-miR-29a	ITGA6 Integrin, alpha 6 Hs.133397 H06635 3655 101822
ITGA6	NM_000210	hsa-miR-29b	ITGA6 Integrin, alpha 6 Hs.133397 H06635 3655 101822
ITGA6	NM_000210	hsa-miR-29c	ITGA6 Integrin, alpha 6 Hs.133397 H06635 3655 101822
ITGA6	NM_000210	hsa-miR-30c	ITGA6 Integrin, alpha 6 Hs.133397 H06635 3655 101822
ITGA6	NM_000210	hsa-miR-30e	ITGA6 Integrin, alpha 6 Hs.133397 H06635 3655 101822
ITGB1BP2	NM_012278	hsa-miR-23a	ITGB1BP2 Integrin beta 1 binding protein (melusin) 2 Hs.109999 AA134808 26548 116944
ITM2A	NM_004867	hsa-miR-10a	ITM2A Integral membrane protein 2A Hs.17109 N72450 9452 118803
ITM2A	NM_004867	hsa-miR-10b	ITM2A Integral membrane protein 2A Hs.17109 N72450 9452 118803
ITM2A	NM_004867	hsa-miR-19a	ITM2A Integral membrane protein 2A Hs.17109 N72450 9452 118803
ITM2A	NM_004867	hsa-miR-19b	ITM2A Integral membrane protein 2A Hs.17109 N72450 9452 118803
ITPR3	NM_002224	hsa-miR-124	ITPR3 Inositol 1,4,5-triphosphate receptor, type 3 Hs.65758 AA701976 3710 105346
ITPR3	NM_002224	hsa-miR-142-5p	ITPR3 Inositol 1,4,5-triphosphate receptor, type 3 Hs.65758 AA701976 3710 105346
KATNA1	NM_007044	hsa-miR-124	PPIL4 Peptidylprolyl isomerase (cyclophilin)-like 4 Hs.551568 R25672 85313 222983
KATNA1	NM_007044	hsa-miR-15a	PPIL4 Peptidylprolyl isomerase (cyclophilin)-like 4 Hs.551568 R25672 85313 222983
KATNA1	NM_007044	hsa-miR-15b	PPIL4 Peptidylprolyl isomerase (cyclophilin)-like 4 Hs.551568 R25672 85313 222983
KATNA1	NM_007044	hsa-miR-18a	PPIL4 Peptidylprolyl isomerase (cyclophilin)-like 4 Hs.551568 R25672 85313 222983
KATNA1	NM_007044	hsa-miR-23a	PPIL4 Peptidylprolyl isomerase (cyclophilin)-like 4 Hs.551568 R25672 85313 222983
KATNA1	NM_007044	hsa-miR-27a	PPIL4 Peptidylprolyl isomerase (cyclophilin)-like 4 Hs.551568 R25672 85313 222983
KCNMB3	NM_171828	hsa-miR-21	KCNMB3 Potassium large conductance calcium-activated channel, subfamily M beta member 3 Hs.85701 AA993785 27094 310655
KCTD5	NM_018992	hsa-miR-29a	KCTD5 Potassium channel tetramerisation domain containing 5 Hs.61960 T99645 54442 99630
KCTD5	NM_018992	hsa-miR-29b	KCTD5 Potassium channel tetramerisation domain containing 5 Hs.61960 T99645 54442 99630
KCTD5	NM_018992	hsa-miR-29c	KCTD5 Potassium channel tetramerisation domain containing 5 Hs.61960 T99645 54442 99630
KCTD5	NM_018992	hsa-miR-30c	KCTD5 Potassium channel tetramerisation domain containing 5 Hs.61960 T99645 54442 99630
KCTD5	NM_018992	hsa-miR-30e	KCTD5 Potassium channel tetramerisation domain containing 5 Hs.61960 T99645 54442 99630
KIAA0329	NM_014844	hsa-let-7a	KIAA0329 KIAA0329 Hs.195667 H75628 9895 113791
KIAA0329	NM_014844	hsa-let-7b	KIAA0329 KIAA0329 Hs.195667 H75628 9895 113791
KIAA0329	NM_014844	hsa-let-7c	KIAA0329 KIAA0329 Hs.195667 H75628 9895 113791
KIAA0329	NM_014844	hsa-let-7d	KIAA0329 KIAA0329 Hs.195667 H75628 9895 113791
KIAA0329	NM_014844	hsa-let-7e	KIAA0329 KIAA0329 Hs.195667 H75628 9895 113791
KIAA0329	NM_014844	hsa-let-7f	KIAA0329 KIAA0329 Hs.195667 H75628 9895 113791
KIAA0329	NM_014844	hsa-miR-124	KIAA0329 KIAA0329 Hs.195667 H75628 9895 113791
KIAA0329	NM_014844	hsa-miR-92a	KIAA0329 KIAA0329 Hs.195667 H75628 9895 113791

KIAA0408	NM 014702	hsa-miR-30c	KIAA0408 KIAA0408 Hs.319247 AI674081 9729 316202
KIAA0408	NM 014702	hsa-miR-30e	KIAA0408 KIAA0408 Hs.319247 AI674081 9729 316202
KIT	NM 000222	hsa-miR-106a	KIT V-kit Hardy-Zuckerman 4 feline sarcoma viral oncogene homolog Hs.479754 N24824 3815 106224
KIT	NM 000222	hsa-miR-142-5p	KIT V-kit Hardy-Zuckerman 4 feline sarcoma viral oncogene homolog Hs.479754 N24824 3815 106224
KIT	NM 000222	hsa-miR-18a	KIT V-kit Hardy-Zuckerman 4 feline sarcoma viral oncogene homolog Hs.479754 N24824 3815 106224
KIT	NM 000222	hsa-miR-19a	KIT V-kit Hardy-Zuckerman 4 feline sarcoma viral oncogene homolog Hs.479754 N24824 3815 106224
KIT	NM 000222	hsa-miR-19b	KIT V-kit Hardy-Zuckerman 4 feline sarcoma viral oncogene homolog Hs.479754 N24824 3815 106224
KIT	NM 000222	hsa-miR-20a	KIT V-kit Hardy-Zuckerman 4 feline sarcoma viral oncogene homolog Hs.479754 N24824 3815 106224
KLF12	NM 007249	hsa-miR-106a	KLF12 Kruppel-like factor 12 Hs.373857 W84386 11278 115431
KLF12	NM 007249	hsa-miR-124	KLF12 Kruppel-like factor 12 Hs.373857 W84386 11278 115431
KLF12	NM 007249	hsa-miR-17	KLF12 Kruppel-like factor 12 Hs.373857 W84386 11278 115431
KLF12	NM 007249	hsa-miR-20a	KLF12 Kruppel-like factor 12 Hs.373857 W84386 11278 115431
KLF12	NM 007249	hsa-miR-23a	KLF12 Kruppel-like factor 12 Hs.373857 W84386 11278 115431
KLF12	NM 007249	hsa-miR-27a	KLF12 Kruppel-like factor 12 Hs.373857 W84386 11278 115431
KLF12	NM 007249	hsa-miR-29a	KLF12 Kruppel-like factor 12 Hs.373857 W84386 11278 115431
KLF12	NM 007249	hsa-miR-29b	KLF12 Kruppel-like factor 12 Hs.373857 W84386 11278 115431
KLF12	NM 007249	hsa-miR-29c	KLF12 Kruppel-like factor 12 Hs.373857 W84386 11278 115431
KLF12	NM 007249	hsa-miR-92a	KLF12 Kruppel-like factor 12 Hs.373857 W84386 11278 115431
KLF3	NM 016531	hsa-miR-23a	KLF3 Kruppel-like factor 3 (basic) Hs.298658 AA405550 51274 118316
LBH	NM 030915	hsa-let-7a	LBH Likely ortholog of mouse limb-bud and heart gene Hs.468044 AA630084 81606 225740
LBH	NM 030915	hsa-let-7b	LBH Likely ortholog of mouse limb-bud and heart gene Hs.468044 AA630084 81606 225740
LBH	NM 030915	hsa-let-7c	LBH Likely ortholog of mouse limb-bud and heart gene Hs.468044 AA630084 81606 225740
LBH	NM 030915	hsa-let-7e	LBH Likely ortholog of mouse limb-bud and heart gene Hs.468044 AA630084 81606 225740
LBH	NM 030915	hsa-let-7f	LBH Likely ortholog of mouse limb-bud and heart gene Hs.468044 AA630084 81606 225740
LBH	NM 030915	hsa-miR-19a	LBH Likely ortholog of mouse limb-bud and heart gene Hs.468044 AA630084 81606 225740
LBH	NM 030915	hsa-miR-19b	LBH Likely ortholog of mouse limb-bud and heart gene Hs.468044 AA630084 81606 225740
LBH	NM 030915	hsa-miR-27a	LBH Likely ortholog of mouse limb-bud and heart gene Hs.468044 AA630084 81606 225740
LBH	NM 030915	hsa-miR-98	LBH Likely ortholog of mouse limb-bud and heart gene Hs.468044 AA630084 81606 225740
LGMN	NM 001008530	hsa-miR-19b	LGMN Legumain Hs.18069 AA425938 5641 109698
LGMN	NM 001008530	hsa-miR-23a	LGMN Legumain Hs.18069 AA425938 5641 109698
LHFPL2	NM 005779	hsa-miR-23a	LHFPL2 Lipoma HMGIC fusion partner-like 2 Hs.79299 AA863469 10184 116059
LHFPL2	NM 005779	hsa-miR-92a	LHFPL2 Lipoma HMGIC fusion partner-like 2 Hs.79299 AA863469 10184 116059
LMO4	NM 006769	hsa-miR-10a	LMO4 LIM domain only 4 Hs.436792 H27986 8543 108294
LMO4	NM 006769	hsa-miR-10b	LMO4 LIM domain only 4 Hs.436792 H27986 8543 108294
LMO4	NM 006769	hsa-miR-124	LMO4 LIM domain only 4 Hs.436792 H27986 8543 108294
LPP	NM 005578	hsa-miR-124	LPP LIM domain containing preferred translocation partner in lipoma Hs.444362 AA047443 4026 114579
LPP	NM 005578	hsa-miR-142-5p	LPP LIM domain containing preferred translocation partner in lipoma Hs.444362 AA047443 4026 114579
LPP	NM 005578	hsa-miR-155	LPP LIM domain containing preferred translocation partner in lipoma Hs.444362 AA047443 4026 114579
LPP	NM 005578	hsa-miR-15a	LPP LIM domain containing preferred translocation partner in lipoma Hs.444362 AA047443 4026 114579
LPP	NM 005578	hsa-miR-15b	LPP LIM domain containing preferred translocation partner in lipoma Hs.444362 AA047443 4026 114579
LPP	NM 005578	hsa-miR-16	LPP LIM domain containing preferred translocation partner in lipoma Hs.444362 AA047443 4026 114579
LPP	NM 005578	hsa-miR-195	LPP LIM domain containing preferred translocation partner in lipoma Hs.444362 AA047443 4026 114579
LPP	NM 005578	hsa-miR-19a	LPP LIM domain containing preferred translocation partner in lipoma Hs.444362 AA047443 4026 114579
LPP	NM 005578	hsa-miR-19b	LPP LIM domain containing preferred translocation partner in lipoma Hs.444362 AA047443 4026 114579
LPP	NM 005578	hsa-miR-23a	LPP LIM domain containing preferred translocation partner in lipoma Hs.444362 AA047443 4026 114579
LPP	NM 005578	hsa-miR-30c	LPP LIM domain containing preferred translocation partner in lipoma Hs.444362 AA047443 4026 114579
LPP	NM 005578	hsa-miR-30e	LPP LIM domain containing preferred translocation partner in lipoma Hs.444362 AA047443 4026 114579
LRP10	NM 014045	hsa-miR-10b	Transcribed locus, weakly similar to XP_496299.1 PREDICTED: hypothetical protein LOC148206 [Homo sapiens] Hs.531057 AA775620 220506
LRP6	NM 002336	hsa-miR-106a	LRP6 Low density lipoprotein receptor-related protein 6 Hs.210343 AA126261 4040 223079
LRP6	NM 002336	hsa-miR-124	LRP6 Low density lipoprotein receptor-related protein 6 Hs.210343 AA126261 4040 223079
LRP6	NM 002336	hsa-miR-15a	LRP6 Low density lipoprotein receptor-related protein 6 Hs.210343 AA126261 4040 223079
LRP6	NM 002336	hsa-miR-15b	LRP6 Low density lipoprotein receptor-related protein 6 Hs.210343 AA126261 4040 223079
LRP6	NM 002336	hsa-miR-16	LRP6 Low density lipoprotein receptor-related protein 6 Hs.210343 AA126261 4040 223079

LRP6	NM_002336	hsa-miR-19a	LRP6	Low density lipoprotein receptor-related protein 6	Hs.210343	AA126261	4040	223079
LRP6	NM_002336	hsa-miR-195	LRP6	Low density lipoprotein receptor-related protein 6	Hs.210343	AA126261	4040	223079
LRP6	NM_002336	hsa-miR-20a	LRP6	Low density lipoprotein receptor-related protein 6	Hs.210343	AA126261	4040	223079
LRP6	NM_002336	hsa-miR-27a	LRP6	Low density lipoprotein receptor-related protein 6	Hs.210343	AA126261	4040	223079
LRP6	NM_002336	hsa-miR-29a	LRP6	Low density lipoprotein receptor-related protein 6	Hs.210343	AA126261	4040	223079
LRP6	NM_002336	hsa-miR-30c	LRP6	Low density lipoprotein receptor-related protein 6	Hs.210343	AA126261	4040	223079
LRRC28	NM_144598	hsa-miR-23a	LRRC28	Leucine rich repeat containing 28	Hs.459507	R95131	123355	100314
MALT1	NM_006785	hsa-miR-21		CDNA: FLJ22806 fis, clone KAI2845	Hs.33417	AA598828		99523
MAN1A2	NM_006699	hsa-miR-142-5p	MAN1A2	Mannosidase, alpha, class 1A, member 2	Hs.435938	AA455062	10905	119283
MAN1A2	NM_006699	hsa-miR-18a	MAN1A2	Mannosidase, alpha, class 1A, member 2	Hs.435938	AA455062	10905	119283
MAN1A2	NM_006699	hsa-miR-30c	MAN1A2	Mannosidase, alpha, class 1A, member 2	Hs.435938	AA455062	10905	119283
MAN1A2	NM_006699	hsa-miR-30e	MAN1A2	Mannosidase, alpha, class 1A, member 2	Hs.435938	AA455062	10905	119283
MAP3K2	NM_006609	hsa-miR-106a	MAP3K2	Mitogen-activated protein kinase kinase kinase 2	Hs.145605	H45390	10746	112380
MAP3K2	NM_006609	hsa-miR-17	MAP3K2	Mitogen-activated protein kinase kinase kinase 2	Hs.145605	H45390	10746	112380
MAP3K2	NM_006609	hsa-miR-20a	MAP3K2	Mitogen-activated protein kinase kinase kinase 2	Hs.145605	H45390	10746	112380
MAP3K7IP2	NM_015093	hsa-let-7a	MAP3K7IP2	Mitogen-activated protein kinase kinase kinase 7 interacting protein 2	Hs.269775	AA457253	23118	104333
MAP3K7IP2	NM_015093	hsa-let-7b	MAP3K7IP2	Mitogen-activated protein kinase kinase kinase 7 interacting protein 2	Hs.269775	AA457253	23118	104333
MAP3K7IP2	NM_015093	hsa-let-7c	MAP3K7IP2	Mitogen-activated protein kinase kinase kinase 7 interacting protein 2	Hs.269775	AA457253	23118	104333
MAP3K7IP2	NM_015093	hsa-let-7d	MAP3K7IP2	Mitogen-activated protein kinase kinase kinase 7 interacting protein 2	Hs.269775	AA457253	23118	104333
MAP3K7IP2	NM_015093	hsa-let-7e	MAP3K7IP2	Mitogen-activated protein kinase kinase kinase 7 interacting protein 2	Hs.269775	AA457253	23118	104333
MAP3K7IP2	NM_015093	hsa-let-7f	MAP3K7IP2	Mitogen-activated protein kinase kinase kinase 7 interacting protein 2	Hs.269775	AA457253	23118	104333
MAP3K7IP2	NM_015093	hsa-miR-142-5p	MAP3K7IP2	Mitogen-activated protein kinase kinase kinase 7 interacting protein 2	Hs.269775	AA457253	23118	104333
MAP3K7IP2	NM_015093	hsa-miR-155	MAP3K7IP2	Mitogen-activated protein kinase kinase kinase 7 interacting protein 2	Hs.269775	AA457253	23118	104333
MAP3K7IP2	NM_015093	hsa-miR-23a	MAP3K7IP2	Mitogen-activated protein kinase kinase kinase 7 interacting protein 2	Hs.269775	AA457253	23118	104333
MAP3K7IP2	NM_015093	hsa-miR-27a	MAP3K7IP2	Mitogen-activated protein kinase kinase kinase 7 interacting protein 2	Hs.269775	AA457253	23118	104333
MAP3K7IP2	NM_015093	hsa-miR-98	MAP3K7IP2	Mitogen-activated protein kinase kinase kinase 7 interacting protein 2	Hs.269775	AA457253	23118	104333
MAP4K4	NM_145687	hsa-let-7a	MAP4K4	Mitogen-activated protein kinase kinase kinase 4	Hs.431550	AA283001	9448	222151
MAP4K4	NM_145687	hsa-let-7b	MAP4K4	Mitogen-activated protein kinase kinase kinase 4	Hs.431550	AA283001	9448	222151
MAP4K4	NM_145687	hsa-let-7c	MAP4K4	Mitogen-activated protein kinase kinase kinase 4	Hs.431550	AA283001	9448	222151
MAP4K4	NM_145687	hsa-let-7d	MAP4K4	Mitogen-activated protein kinase kinase kinase 4	Hs.431550	AA283001	9448	222151
MAP4K4	NM_145687	hsa-let-7e	MAP4K4	Mitogen-activated protein kinase kinase kinase 4	Hs.431550	AA283001	9448	222151
MAP4K4	NM_145687	hsa-let-7f	MAP4K4	Mitogen-activated protein kinase kinase kinase 4	Hs.431550	AA283001	9448	222151
MAP4K4	NM_145687	hsa-miR-142-5p	MAP4K4	Mitogen-activated protein kinase kinase kinase 4	Hs.431550	AA283001	9448	222151
MAP4K4	NM_145687	hsa-miR-155	MAP4K4	Mitogen-activated protein kinase kinase kinase 4	Hs.431550	AA283001	9448	222151
MAP4K4	NM_145687	hsa-miR-23a	MAP4K4	Mitogen-activated protein kinase kinase kinase 4	Hs.431550	AA283001	9448	222151
MAP4K4	NM_145687	hsa-miR-29a	MAP4K4	Mitogen-activated protein kinase kinase kinase 4	Hs.431550	AA283001	9448	222151
MAP4K4	NM_145687	hsa-miR-29b	MAP4K4	Mitogen-activated protein kinase kinase kinase 4	Hs.431550	AA283001	9448	222151
MAP4K4	NM_145687	hsa-miR-29c	MAP4K4	Mitogen-activated protein kinase kinase kinase 4	Hs.431550	AA283001	9448	222151
MAP4K4	NM_145687	hsa-miR-30c	MAP4K4	Mitogen-activated protein kinase kinase kinase 4	Hs.431550	AA283001	9448	222151
MAP4K4	NM_145687	hsa-miR-30e	MAP4K4	Mitogen-activated protein kinase kinase kinase 4	Hs.431550	AA283001	9448	222151
MAP4K4	NM_145687	hsa-miR-369-3p	MAP4K4	Mitogen-activated protein kinase kinase kinase 4	Hs.431550	AA283001	9448	222151
MAP4K4	NM_145687	hsa-miR-98	MAP4K4	Mitogen-activated protein kinase kinase kinase 4	Hs.431550	AA283001	9448	222151
MAT2B	NM_182796	hsa-miR-106a	MAT2B	Methionine adenosyltransferase II, beta	Hs.54642	AI668657	27430	118774
MAT2B	NM_182796	hsa-miR-142-5p	MAT2B	Methionine adenosyltransferase II, beta	Hs.54642	AI668657	27430	118774
MAT2B	NM_182796	hsa-miR-17	MAT2B	Methionine adenosyltransferase II, beta	Hs.54642	AI668657	27430	118774
MAT2B	NM_182796	hsa-miR-20a	MAT2B	Methionine adenosyltransferase II, beta	Hs.54642	AI668657	27430	118774
MBNL2	NM_207304	hsa-miR-18a	MBNL2	Muscleblind-like 2 (Drosophila)	Hs.125715	W16832	10150	102644
MBNL2	NM_207304	hsa-miR-19a	MBNL2	Muscleblind-like 2 (Drosophila)	Hs.125715	W16832	10150	102644
MBNL2	NM_207304	hsa-miR-19b	MBNL2	Muscleblind-like 2 (Drosophila)	Hs.125715	W16832	10150	102644
MBNL2	NM_207304	hsa-miR-27a	MBNL2	Muscleblind-like 2 (Drosophila)	Hs.125715	W16832	10150	102644
MBNL2	NM_207304	hsa-miR-30c	MBNL2	Muscleblind-like 2 (Drosophila)	Hs.125715	W16832	10150	102644
MBNL2	NM_207304	hsa-miR-30e	MBNL2	Muscleblind-like 2 (Drosophila)	Hs.125715	W16832	10150	102644
MBNL2	NM_207304	hsa-miR-369-3p	MBNL2	Muscleblind-like 2 (Drosophila)	Hs.125715	W16832	10150	102644

MBNL3	NM 133486	hsa-miR-30c	MBNL3 Muscleblind-like 3 (Drosophila) Hs.105134 AA460149 55796 102468
MBNL3	NM 133486	hsa-miR-30e	MBNL3 Muscleblind-like 3 (Drosophila) Hs.105134 AA460149 55796 102468
MCAM	NM 006500	hsa-miR-106a	MCAM Melanoma cell adhesion molecule Hs.511397 AI888888 4162 331100
MDFIC	NM 199072	hsa-miR-19a	HIC MyoD family inhibitor domain containing Hs.546385 AI668639 29969 112705
MDFIC	NM 199072	hsa-miR-19b	HIC MyoD family inhibitor domain containing Hs.546385 AI668639 29969 112705
MDFIC	NM 199072	hsa-miR-23a	HIC MyoD family inhibitor domain containing Hs.546385 AI668639 29969 112705
MEF2A	NM 005587	hsa-miR-155	MEF2A MADS box transcription enhancer factor 2, polypeptide A (myocyte enhancer factor 2A) Hs.268675 AA480996 4205 309342
MEST	NM 177524	hsa-miR-155	MEST Mesoderm specific transcript homolog (mouse) Hs.270978 AA598610 4232 101074
MEST	NM 177524	hsa-miR-29a	MEST Mesoderm specific transcript homolog (mouse) Hs.270978 AA598610 4232 101074
MEST	NM 177524	hsa-miR-29b	MEST Mesoderm specific transcript homolog (mouse) Hs.270978 AA598610 4232 101074
MEST	NM 177524	hsa-miR-29c	MEST Mesoderm specific transcript homolog (mouse) Hs.270978 AA598610 4232 101074
MGAT4A	NM 012214	hsa-let-7a	MGC52110 Hypothetical protein MGC52110 Hs.177576 AA933892 493753 308139
MGAT4A	NM 012214	hsa-let-7b	MGC52110 Hypothetical protein MGC52110 Hs.177576 AA933892 493753 308139
MGAT4A	NM 012214	hsa-let-7c	MGC52110 Hypothetical protein MGC52110 Hs.177576 AA933892 493753 308139
MGAT4A	NM 012214	hsa-let-7d	MGC52110 Hypothetical protein MGC52110 Hs.177576 AA933892 493753 308139
MGAT4A	NM 012214	hsa-let-7e	MGC52110 Hypothetical protein MGC52110 Hs.177576 AA933892 493753 308139
MGAT4A	NM 012214	hsa-let-7f	MGC52110 Hypothetical protein MGC52110 Hs.177576 AA933892 493753 308139
MGAT4A	NM 012214	hsa-miR-142-5p	MGC52110 Hypothetical protein MGC52110 Hs.177576 AA933892 493753 308139
MGAT4A	NM 012214	hsa-miR-369-3p	MGC52110 Hypothetical protein MGC52110 Hs.177576 AA933892 493753 308139
MGAT4A	NM 012214	hsa-miR-98	MGC52110 Hypothetical protein MGC52110 Hs.177576 AA933892 493753 308139
MKLN1	NM 013255	hsa-miR-124	MKLN1 Muskelin 1, intracellular mediator containing kelch motifs Hs.44693 AI263113 4289 112075
MKLN1	NM 013255	hsa-miR-142-5p	MKLN1 Muskelin 1, intracellular mediator containing kelch motifs Hs.44693 AI263113 4289 112075
MLL	NM 005933	hsa-miR-124	MLL Myeloid/lymphoid or mixed-lineage leukemia (trithorax homolog, Drosophila) Hs.258855 AA701046 4297 221868
MLL	NM 005933	hsa-miR-142-5p	MLL Myeloid/lymphoid or mixed-lineage leukemia (trithorax homolog, Drosophila) Hs.258855 AA701046 4297 221868
MLL	NM 005933	hsa-miR-30c	MLL Myeloid/lymphoid or mixed-lineage leukemia (trithorax homolog, Drosophila) Hs.258855 AA701046 4297 221868
MLL	NM 005933	hsa-miR-30e	MLL Myeloid/lymphoid or mixed-lineage leukemia (trithorax homolog, Drosophila) Hs.258855 AA701046 4297 221868
MLL	NM 005933	hsa-miR-369-3p	MLL Myeloid/lymphoid or mixed-lineage leukemia (trithorax homolog, Drosophila) Hs.258855 AA701046 4297 221868
MLLT3	NM 004529	hsa-miR-124	MLLT3 Myeloid/lymphoid or mixed-lineage leukemia (trithorax homolog, Drosophila); translocated to, 3 Hs.493585 AA443284 4300 116155
MLLT7	NM 005938	hsa-miR-23a	MLLT7 Myeloid/lymphoid or mixed-lineage leukemia (trithorax homolog, Drosophila); translocated to, 7 Hs.239663 T54418 4303 108939
MLLT7	NM 005938	hsa-miR-29a	MLLT7 Myeloid/lymphoid or mixed-lineage leukemia (trithorax homolog, Drosophila); translocated to, 7 Hs.239663 T54418 4303 108939
MLLT7	NM 005938	hsa-miR-29b	MLLT7 Myeloid/lymphoid or mixed-lineage leukemia (trithorax homolog, Drosophila); translocated to, 7 Hs.239663 T54418 4303 108939
MLLT7	NM 005938	hsa-miR-29c	MLLT7 Myeloid/lymphoid or mixed-lineage leukemia (trithorax homolog, Drosophila); translocated to, 7 Hs.239663 T54418 4303 108939
MN1	NM 002430	hsa-miR-124	MN1 Meningioma (disrupted in balanced translocation) 1 Hs.268515 R59212 4330 101031
MN1	NM 002430	hsa-miR-15a	MN1 Meningioma (disrupted in balanced translocation) 1 Hs.268515 R59212 4330 101031
MN1	NM 002430	hsa-miR-19a	MN1 Meningioma (disrupted in balanced translocation) 1 Hs.268515 R59212 4330 101031
MN1	NM 002430	hsa-miR-19b	MN1 Meningioma (disrupted in balanced translocation) 1 Hs.268515 R59212 4330 101031
MX1	NM 002462	hsa-miR-98	MX1 Myxovirus (influenza virus) resistance 1, interferon-inducible protein p78 (mouse) Hs.517307 AA405416 4599 104876
MXI1	NM 130439	hsa-miR-106a	MXI1 MAX interactor 1 Hs.501023 N35022 Mad2=MXI-1=MAX-binding protein=antagonizer of myc transcript 4601 121528
MXI1	NM 130439	hsa-miR-142-5p	MXI1 MAX interactor 1 Hs.501023 N35022 Mad2=MXI-1=MAX-binding protein=antagonizer of myc transcript 4601 121528
MXI1	NM 130439	hsa-miR-20a	MXI1 MAX interactor 1 Hs.501023 N35022 Mad2=MXI-1=MAX-binding protein=antagonizer of myc transcript 4601 121528
MXI1	NM 130439	hsa-miR-29a	MXI1 MAX interactor 1 Hs.501023 N35022 Mad2=MXI-1=MAX-binding protein=antagonizer of myc transcript 4601 121528
MXI1	NM 130439	hsa-miR-29b	MXI1 MAX interactor 1 Hs.501023 N35022 Mad2=MXI-1=MAX-binding protein=antagonizer of myc transcript 4601 121528
MXI1	NM 130439	hsa-miR-29c	MXI1 MAX interactor 1 Hs.501023 N35022 Mad2=MXI-1=MAX-binding protein=antagonizer of myc transcript 4601 121528
NAALAD2	NM 005467	hsa-miR-15a	NAALAD2 N-acetylated alpha-linked acidic dipeptidase 2 Hs.503560 AI668656 10003 104195
NBL1	NM 182744	hsa-miR-106a	NBL1 Neuroblastoma, suppression of tumorigenicity 1 Hs.439671 AA598830 4681 103873
NBL1	NM 182744	hsa-miR-17	NBL1 Neuroblastoma, suppression of tumorigenicity 1 Hs.439671 AA598830 4681 103873
NBL1	NM 182744	hsa-miR-20a	NBL1 Neuroblastoma, suppression of tumorigenicity 1 Hs.439671 AA598830 4681 103873
NCOA4	NM 005437	hsa-miR-124	NCOA4 Nuclear receptor coactivator 4 Hs.522932 AA133212 8031 116750
NCOA4	NM 005437	hsa-miR-195	NCOA4 Nuclear receptor coactivator 4 Hs.522932 AA133212 8031 116750
NCOA4	NM 005437	hsa-miR-29a	NCOA4 Nuclear receptor coactivator 4 Hs.522932 AA133212 8031 116750
NCOA4	NM 005437	hsa-miR-29b	NCOA4 Nuclear receptor coactivator 4 Hs.522932 AA133212 8031 116750
NCOA4	NM 005437	hsa-miR-29c	NCOA4 Nuclear receptor coactivator 4 Hs.522932 AA133212 8031 116750
NCOA4	NM 005437	hsa-miR-369-3p	NCOA4 Nuclear receptor coactivator 4 Hs.522932 AA133212 8031 116750

NEDD4L	NM_015277	hsa-miR-106a	NEDD4L Neural precursor cell expressed, developmentally down-regulated 4-like Hs.185677 AA463445 23327 102546
NEDD4L	NM_015277	hsa-miR-17	NEDD4L Neural precursor cell expressed, developmentally down-regulated 4-like Hs.185677 AA463445 23327 102546
NEDD4L	NM_015277	hsa-miR-20a	NEDD4L Neural precursor cell expressed, developmentally down-regulated 4-like Hs.185677 AA463445 23327 102546
NEDD4L	NM_015277	hsa-miR-30c	NEDD4L Neural precursor cell expressed, developmentally down-regulated 4-like Hs.185677 AA463445 23327 102546
NEDD4L	NM_015277	hsa-miR-30e	NEDD4L Neural precursor cell expressed, developmentally down-regulated 4-like Hs.185677 AA463445 23327 102546
NEDD9	NM_006403	hsa-miR-18a	**MRNA full length insert cDNA clone EUROIIMAGE 966164 Hs.21177 R60924 100528
NEK2	NM_002497	hsa-miR-27a	NEK2 NIMA (never in mitosis gene a)-related kinase 2 Hs.153704 R94281 4751 115877
NID2	NM_007361	hsa-let-7a	NID2 Nidogen 2 (osteonidogen) Hs.369840 AA599763 22795 311780
NID2	NM_007361	hsa-let-7b	NID2 Nidogen 2 (osteonidogen) Hs.369840 AA599763 22795 311780
NID2	NM_007361	hsa-let-7c	NID2 Nidogen 2 (osteonidogen) Hs.369840 AA599763 22795 311780
NID2	NM_007361	hsa-let-7d	NID2 Nidogen 2 (osteonidogen) Hs.369840 AA599763 22795 311780
NID2	NM_007361	hsa-let-7e	NID2 Nidogen 2 (osteonidogen) Hs.369840 AA599763 22795 311780
NID2	NM_007361	hsa-let-7f	NID2 Nidogen 2 (osteonidogen) Hs.369840 AA599763 22795 311780
NID2	NM_007361	hsa-miR-29a	NID2 Nidogen 2 (osteonidogen) Hs.369840 AA599763 22795 311780
NID2	NM_007361	hsa-miR-29b	NID2 Nidogen 2 (osteonidogen) Hs.369840 AA599763 22795 311780
NID2	NM_007361	hsa-miR-29c	NID2 Nidogen 2 (osteonidogen) Hs.369840 AA599763 22795 311780
NID2	NM_007361	hsa-miR-30c	NID2 Nidogen 2 (osteonidogen) Hs.369840 AA599763 22795 311780
NID2	NM_007361	hsa-miR-30e	NID2 Nidogen 2 (osteonidogen) Hs.369840 AA599763 22795 311780
NID2	NM_007361	hsa-miR-98	NID2 Nidogen 2 (osteonidogen) Hs.369840 AA599763 22795 311780
NIPA1	NM_144599	hsa-let-7a	NIPA1 Non imprinted in Prader-Willi/Angelman syndrome 1 Hs.511797 R56814 123606 221666
NIPA1	NM_144599	hsa-let-7b	NIPA1 Non imprinted in Prader-Willi/Angelman syndrome 1 Hs.511797 R56814 123606 221666
NIPA1	NM_144599	hsa-let-7c	NIPA1 Non imprinted in Prader-Willi/Angelman syndrome 1 Hs.511797 R56814 123606 221666
NIPA1	NM_144599	hsa-let-7d	NIPA1 Non imprinted in Prader-Willi/Angelman syndrome 1 Hs.511797 R56814 123606 221666
NIPA1	NM_144599	hsa-let-7e	NIPA1 Non imprinted in Prader-Willi/Angelman syndrome 1 Hs.511797 R56814 123606 221666
NIPA1	NM_144599	hsa-let-7f	NIPA1 Non imprinted in Prader-Willi/Angelman syndrome 1 Hs.511797 R56814 123606 221666
NIPA1	NM_144599	hsa-miR-106a	NIPA1 Non imprinted in Prader-Willi/Angelman syndrome 1 Hs.511797 R56814 123606 221666
NIPA1	NM_144599	hsa-miR-124	NIPA1 Non imprinted in Prader-Willi/Angelman syndrome 1 Hs.511797 R56814 123606 221666
NIPA1	NM_144599	hsa-miR-17	NIPA1 Non imprinted in Prader-Willi/Angelman syndrome 1 Hs.511797 R56814 123606 221666
NIPA1	NM_144599	hsa-miR-19a	NIPA1 Non imprinted in Prader-Willi/Angelman syndrome 1 Hs.511797 R56814 123606 221666
NIPA1	NM_144599	hsa-miR-19b	NIPA1 Non imprinted in Prader-Willi/Angelman syndrome 1 Hs.511797 R56814 123606 221666
NIPA1	NM_144599	hsa-miR-20a	NIPA1 Non imprinted in Prader-Willi/Angelman syndrome 1 Hs.511797 R56814 123606 221666
NIPA1	NM_144599	hsa-miR-98	NIPA1 Non imprinted in Prader-Willi/Angelman syndrome 1 Hs.511797 R56814 123606 221666
NME5	NM_003551	hsa-miR-15a	NME5 Non-metastatic cells 5, protein expressed in (nucleoside-diphosphate kinase) Hs.519602 AA129736 8382 106388
NME5	NM_003551	hsa-miR-15b	NME5 Non-metastatic cells 5, protein expressed in (nucleoside-diphosphate kinase) Hs.519602 AA129736 8382 106388
NME5	NM_003551	hsa-miR-29a	NME5 Non-metastatic cells 5, protein expressed in (nucleoside-diphosphate kinase) Hs.519602 AA129736 8382 106388
NME5	NM_003551	hsa-miR-29b	NME5 Non-metastatic cells 5, protein expressed in (nucleoside-diphosphate kinase) Hs.519602 AA129736 8382 106388
NME5	NM_003551	hsa-miR-29c	NME5 Non-metastatic cells 5, protein expressed in (nucleoside-diphosphate kinase) Hs.519602 AA129736 8382 106388
NRBP	NM_013392	hsa-miR-15a	NRBP Nuclear receptor binding protein Hs.515876 R10018 29959 105435
NRBP	NM_013392	hsa-miR-15b	NRBP Nuclear receptor binding protein Hs.515876 R10018 29959 105435
NRBP	NM_013392	hsa-miR-16	NRBP Nuclear receptor binding protein Hs.515876 R10018 29959 105435
NRBP	NM_013392	hsa-miR-17	NRBP Nuclear receptor binding protein Hs.515876 R10018 29959 105435
NRBP	NM_013392	hsa-miR-195	NRBP Nuclear receptor binding protein Hs.515876 R10018 29959 105435
NRBP	NM_013392	hsa-miR-19a	NRBP Nuclear receptor binding protein Hs.515876 R10018 29959 105435
NRBP	NM_013392	hsa-miR-19b	NRBP Nuclear receptor binding protein Hs.515876 R10018 29959 105435
NRBP	NM_013392	hsa-miR-20a	NRBP Nuclear receptor binding protein Hs.515876 R10018 29959 105435
NRBP	NM_013392	hsa-miR-29a	NRBP Nuclear receptor binding protein Hs.515876 R10018 29959 105435
NRBP	NM_013392	hsa-miR-29b	NRBP Nuclear receptor binding protein Hs.515876 R10018 29959 105435
NRBP	NM_013392	hsa-miR-29c	NRBP Nuclear receptor binding protein Hs.515876 R10018 29959 105435
NRP1	NM_003873	hsa-miR-124	NRP1 Neuropilin 1 Hs.131704 AA098867 8829 103996
NRP1	NM_003873	hsa-miR-142-5p	NRP1 Neuropilin 1 Hs.131704 AA098867 8829 103996
NRP1	NM_003873	hsa-miR-29a	NRP1 Neuropilin 1 Hs.131704 AA098867 8829 103996
NRP1	NM_003873	hsa-miR-30c	NRP1 Neuropilin 1 Hs.131704 AA098867 8829 103996
NRP1	NM_003873	hsa-miR-30e	NRP1 Neuropilin 1 Hs.131704 AA098867 8829 103996

NSUN3	NM 022072	hsa-miR-17	NSUN3 NOL1/NOP2/Sun domain family, member 3 Hs.401062 AA682850 63899 222810
NSUN3	NM 022072	hsa-miR-29a	NSUN3 NOL1/NOP2/Sun domain family, member 3 Hs.401062 AA682850 63899 222810
NSUN3	NM 022072	hsa-miR-29b	NSUN3 NOL1/NOP2/Sun domain family, member 3 Hs.401062 AA682850 63899 222810
NSUN3	NM 022072	hsa-miR-29c	NSUN3 NOL1/NOP2/Sun domain family, member 3 Hs.401062 AA682850 63899 222810
NT5E	NM 002526	hsa-miR-30c	NT5E 5'-nucleotidase, ecto (CD73) Hs.153952 N35316 4907 310325
NT5E	NM 002526	hsa-miR-30e	NT5E 5'-nucleotidase, ecto (CD73) Hs.153952 N35316 4907 310325
ORC6L	NM 014321	hsa-miR-106a	ORC6L **Origin recognition complex, subunit 6 homolog-like (yeast) Hs.49760 N90667 23594 222560
ORC6L	NM 014321	hsa-miR-155	ORC6L **Origin recognition complex, subunit 6 homolog-like (yeast) Hs.49760 N90667 23594 222560
ORC6L	NM 014321	hsa-miR-17	ORC6L **Origin recognition complex, subunit 6 homolog-like (yeast) Hs.49760 N90667 23594 222560
ORC6L	NM 014321	hsa-miR-20a	ORC6L **Origin recognition complex, subunit 6 homolog-like (yeast) Hs.49760 N90667 23594 222560
OSBPL10	NM 017784	hsa-miR-124	OSBPL10 Oxysterol binding protein-like 10 Hs.150122 H03146 114884 106081
OTX2	NM 172337	hsa-let-7a	OTX2 Orthodenticle homolog 2 (Drosophila) Hs.288655 H17803 5015 120721
OTX2	NM 172337	hsa-let-7c	OTX2 Orthodenticle homolog 2 (Drosophila) Hs.288655 H17803 5015 120721
OTX2	NM 172337	hsa-let-7e	OTX2 Orthodenticle homolog 2 (Drosophila) Hs.288655 H17803 5015 120721
OTX2	NM 172337	hsa-miR-124	OTX2 Orthodenticle homolog 2 (Drosophila) Hs.288655 H17803 5015 120721
OTX2	NM 172337	hsa-miR-142-5p	OTX2 Orthodenticle homolog 2 (Drosophila) Hs.288655 H17803 5015 120721
OTX2	NM 172337	hsa-miR-27a	OTX2 Orthodenticle homolog 2 (Drosophila) Hs.288655 H17803 5015 120721
OTX2	NM 172337	hsa-miR-30c	OTX2 Orthodenticle homolog 2 (Drosophila) Hs.288655 H17803 5015 120721
P2RY5	NM 005767	hsa-miR-155	P2RY5 Purinergic receptor P2Y, G-protein coupled, 5 Hs.123464 R91539 purinergic receptor P2Y5=RB intorn-encoded putative G-protei 10161 112029
P2RY5	NM 005767	hsa-miR-23a	P2RY5 Purinergic receptor P2Y, G-protein coupled, 5 Hs.123464 R91539 purinergic receptor P2Y5=RB intorn-encoded putative G-protei 10161 112029
P2RY5	NM 005767	hsa-miR-27a	P2RY5 Purinergic receptor P2Y, G-protein coupled, 5 Hs.123464 R91539 purinergic receptor P2Y5=RB intorn-encoded putative G-protei 10161 112029
P2RY5	NM 005767	hsa-miR-369-3p	P2RY5 Purinergic receptor P2Y, G-protein coupled, 5 Hs.123464 R91539 purinergic receptor P2Y5=RB intorn-encoded putative G-protei 10161 112029
PAFAH1B1	NM 000430	hsa-miR-106a	PAFAH1B1 Platelet-activating factor acetylhydrolase, isoform lb, alpha subunit 45kDa Hs.77318 AA775445 5048 108551
PAFAH1B1	NM 000430	hsa-miR-10a	PAFAH1B1 Platelet-activating factor acetylhydrolase, isoform lb, alpha subunit 45kDa Hs.77318 AA775445 5048 108551
PAFAH1B1	NM 000430	hsa-miR-10b	PAFAH1B1 Platelet-activating factor acetylhydrolase, isoform lb, alpha subunit 45kDa Hs.77318 AA775445 5048 108551
PAFAH1B1	NM 000430	hsa-miR-124	PAFAH1B1 Platelet-activating factor acetylhydrolase, isoform lb, alpha subunit 45kDa Hs.77318 AA775445 5048 108551
PAFAH1B1	NM 000430	hsa-miR-15a	PAFAH1B1 Platelet-activating factor acetylhydrolase, isoform lb, alpha subunit 45kDa Hs.77318 AA775445 5048 108551
PAFAH1B1	NM 000430	hsa-miR-15b	PAFAH1B1 Platelet-activating factor acetylhydrolase, isoform lb, alpha subunit 45kDa Hs.77318 AA775445 5048 108551
PAFAH1B1	NM 000430	hsa-miR-16	PAFAH1B1 Platelet-activating factor acetylhydrolase, isoform lb, alpha subunit 45kDa Hs.77318 AA775445 5048 108551
PAFAH1B1	NM 000430	hsa-miR-17	PAFAH1B1 Platelet-activating factor acetylhydrolase, isoform lb, alpha subunit 45kDa Hs.77318 AA775445 5048 108551
PAFAH1B1	NM 000430	hsa-miR-195	PAFAH1B1 Platelet-activating factor acetylhydrolase, isoform lb, alpha subunit 45kDa Hs.77318 AA775445 5048 108551
PAFAH1B1	NM 000430	hsa-miR-20a	PAFAH1B1 Platelet-activating factor acetylhydrolase, isoform lb, alpha subunit 45kDa Hs.77318 AA775445 5048 108551
PAFAH1B1	NM 000430	hsa-miR-92a	PAFAH1B1 Platelet-activating factor acetylhydrolase, isoform lb, alpha subunit 45kDa Hs.77318 AA775445 5048 108551
PAG1	NM 018440	hsa-miR-124	PAG Phosphoprotein associated with glycosphingolipid microdomains 1 Hs.266175 H08999 55824 111523
PAG1	NM 018440	hsa-miR-21	PAG Phosphoprotein associated with glycosphingolipid microdomains 1 Hs.266175 H08999 55824 111523
PALM2-AKAP2	NM 007203	hsa-miR-15b	PALM2-AKAP2 PALM2-AKAP2 protein Hs.259461 H99415 445815 225677
PALM2-AKAP2	NM 147150	hsa-miR-15b	PALM2-AKAP2 PALM2-AKAP2 protein Hs.259461 H99415 445815 225677
PANX1	NM 015368	hsa-miR-20a	PANX1 Pannexin 1 Hs.503584 H41987 24145 110608
PAWR	NM 002583	hsa-miR-155	CDNA FLJ14942 fis, A-PLACE1011185 Hs.506182 R26405 307621
PAWR	NM 002583	hsa-miR-195	CDNA FLJ14942 fis, A-PLACE1011185 Hs.506182 R26405 307621
PAWR	NM 002583	hsa-miR-30c	CDNA FLJ14942 fis, A-PLACE1011185 Hs.506182 R26405 307621
PAWR	NM 002583	hsa-miR-30e	CDNA FLJ14942 fis, A-PLACE1011185 Hs.506182 R26405 307621
PCAF	NM 003884	hsa-miR-106a	PCAF P300/CBP-associated factor Hs.533055 N74637 P/CAF=p300/CBP-associated factor 8850 108991
PCAF	NM 003884	hsa-miR-142-5p	PCAF P300/CBP-associated factor Hs.533055 N74637 P/CAF=p300/CBP-associated factor 8850 108991
PCAF	NM 003884	hsa-miR-17	PCAF P300/CBP-associated factor Hs.533055 N74637 P/CAF=p300/CBP-associated factor 8850 108991
PCAF	NM 003884	hsa-miR-19a	PCAF P300/CBP-associated factor Hs.533055 N74637 P/CAF=p300/CBP-associated factor 8850 108991
PCAF	NM 003884	hsa-miR-19b	PCAF P300/CBP-associated factor Hs.533055 N74637 P/CAF=p300/CBP-associated factor 8850 108991
PCAF	NM 003884	hsa-miR-20a	PCAF P300/CBP-associated factor Hs.533055 N74637 P/CAF=p300/CBP-associated factor 8850 108991
PCAF	NM 003884	hsa-miR-23a	PCAF P300/CBP-associated factor Hs.533055 N74637 P/CAF=p300/CBP-associated factor 8850 108991
PCAF	NM 003884	hsa-miR-27a	PCAF P300/CBP-associated factor Hs.533055 N74637 P/CAF=p300/CBP-associated factor 8850 108991
PCAF	NM 003884	hsa-miR-92a	PCAF P300/CBP-associated factor Hs.533055 N74637 P/CAF=p300/CBP-associated factor 8850 108991
PCDH7	NM 032457	hsa-miR-142-5p	PCDH7 BH-protocadherin (brain-heart) Hs.479439 AA436456 5099 109841
PCGF5	NM 032373	hsa-miR-19a	PCGF5 Polycomb group ring finger 5 Hs.500512 AA136060 84333 104213

PCGF5	NM_032373	hsa-miR-19b	PCGF5 Polycomb group ring finger 5 Hs.500512 AA136060 84333 104213
PCGF5	NM_032373	hsa-miR-30c	PCGF5 Polycomb group ring finger 5 Hs.500512 AA136060 84333 104213
PCGF5	NM_032373	hsa-miR-30e	PCGF5 Polycomb group ring finger 5 Hs.500512 AA136060 84333 104213
PCYT2	NM_002861	hsa-miR-15b	PCYT2 **Phosphate cytidyltransferase 2, ethanolamine Hs.551512 AA477754 5833 316676
PDCL	NM_005388	hsa-miR-30c	PDCL Phosducin-like Hs.9302 AI149393 5082 315724
PDCL	NM_005388	hsa-miR-30e	PDCL Phosducin-like Hs.9302 AI149393 5082 315724
PDE3B	NM_000922	hsa-miR-17	PDE3B Phosphodiesterase 3B, cGMP-inhibited Hs.445711 W84658 5140 115428
PDE3B	NM_000922	hsa-miR-27a	PDE3B Phosphodiesterase 3B, cGMP-inhibited Hs.445711 W84658 5140 115428
PDE3B	NM_000922	hsa-miR-369-3p	PDE3B Phosphodiesterase 3B, cGMP-inhibited Hs.445711 W84658 5140 115428
PDE4DIP	NM_014644	hsa-miR-10a	NUDT4P1 Nudix (nucleoside diphosphate linked moiety X)-type motif 4 pseudogene 1 Hs.356699 AA883597 440672 224640
PDE4DIP	NM_014644	hsa-miR-10b	NUDT4P1 Nudix (nucleoside diphosphate linked moiety X)-type motif 4 pseudogene 1 Hs.356699 AA883597 440672 224640
PDE4DIP	NM_022359	hsa-miR-142-5p	NUDT4P1 Nudix (nucleoside diphosphate linked moiety X)-type motif 4 pseudogene 1 Hs.356699 AA883597 440672 224640
PDE4DIP	NM_014644	hsa-miR-15a	NUDT4P1 Nudix (nucleoside diphosphate linked moiety X)-type motif 4 pseudogene 1 Hs.356699 AA883597 440672 224640
PDE4DIP	NM_014644	hsa-miR-19a	NUDT4P1 Nudix (nucleoside diphosphate linked moiety X)-type motif 4 pseudogene 1 Hs.356699 AA883597 440672 224640
PDE4DIP	NM_014644	hsa-miR-19b	NUDT4P1 Nudix (nucleoside diphosphate linked moiety X)-type motif 4 pseudogene 1 Hs.356699 AA883597 440672 224640
PDE4DIP	NM_022359	hsa-miR-21	NUDT4P1 Nudix (nucleoside diphosphate linked moiety X)-type motif 4 pseudogene 1 Hs.356699 AA883597 440672 224640
PHF20L1	NM_198513	hsa-miR-10a	PHF20L1 PHD finger protein 20-like 1 Hs.304362 AI310971 51105 318933
PHF20L1	NM_198513	hsa-miR-10b	PHF20L1 PHD finger protein 20-like 1 Hs.304362 AI310971 51105 318933
PHF20L1	NM_198513	hsa-miR-124	PHF20L1 PHD finger protein 20-like 1 Hs.304362 AI310971 51105 318933
PHF20L1	NM_016018	hsa-miR-142-5p	PHF20L1 PHD finger protein 20-like 1 Hs.304362 AI310971 51105 318933
PHF20L1	NM_032205	hsa-miR-15a	PHF20L1 PHD finger protein 20-like 1 Hs.304362 AI310971 51105 318933
PHF20L1	NM_032205	hsa-miR-15b	PHF20L1 PHD finger protein 20-like 1 Hs.304362 AI310971 51105 318933
PHF20L1	NM_032205	hsa-miR-16	PHF20L1 PHD finger protein 20-like 1 Hs.304362 AI310971 51105 318933
PHF20L1	NM_198513	hsa-miR-18a	PHF20L1 PHD finger protein 20-like 1 Hs.304362 AI310971 51105 318933
PHF20L1	NM_032205	hsa-miR-195	PHF20L1 PHD finger protein 20-like 1 Hs.304362 AI310971 51105 318933
PHF20L1	NM_032205	hsa-miR-21	PHF20L1 PHD finger protein 20-like 1 Hs.304362 AI310971 51105 318933
PHF20L1	NM_198513	hsa-miR-23a	PHF20L1 PHD finger protein 20-like 1 Hs.304362 AI310971 51105 318933
PHF20L1	NM_016018	hsa-miR-369-3p	PHF20L1 PHD finger protein 20-like 1 Hs.304362 AI310971 51105 318933
PHLPPL	NM_015020	hsa-let-7a	KIAA0931 PH domain and leucine rich repeat protein phosphatase-like Hs.531564 AA417700 23035 307238
PHLPPL	NM_015020	hsa-let-7b	KIAA0931 PH domain and leucine rich repeat protein phosphatase-like Hs.531564 AA417700 23035 307238
PHLPPL	NM_015020	hsa-let-7c	KIAA0931 PH domain and leucine rich repeat protein phosphatase-like Hs.531564 AA417700 23035 307238
PHLPPL	NM_015020	hsa-let-7d	KIAA0931 PH domain and leucine rich repeat protein phosphatase-like Hs.531564 AA417700 23035 307238
PHLPPL	NM_015020	hsa-let-7f	KIAA0931 PH domain and leucine rich repeat protein phosphatase-like Hs.531564 AA417700 23035 307238
PHLPPL	NM_015020	hsa-miR-142-5p	KIAA0931 PH domain and leucine rich repeat protein phosphatase-like Hs.531564 AA417700 23035 307238
PIK3CA	NM_006218	hsa-miR-10a	PIK3CA Phosphoinositide-3-kinase, catalytic, alpha polypeptide Hs.553498 W72473 5290 113225
PIK3CA	NM_006218	hsa-miR-10b	PIK3CA Phosphoinositide-3-kinase, catalytic, alpha polypeptide Hs.553498 W72473 5290 113225
PIK3CA	NM_006218	hsa-miR-124	PIK3CA Phosphoinositide-3-kinase, catalytic, alpha polypeptide Hs.553498 W72473 5290 113225
PIK3CA	NM_006218	hsa-miR-155	PIK3CA Phosphoinositide-3-kinase, catalytic, alpha polypeptide Hs.553498 W72473 5290 113225
PIK3CA	NM_006218	hsa-miR-19a	PIK3CA Phosphoinositide-3-kinase, catalytic, alpha polypeptide Hs.553498 W72473 5290 113225
PIK3CA	NM_006218	hsa-miR-19b	PIK3CA Phosphoinositide-3-kinase, catalytic, alpha polypeptide Hs.553498 W72473 5290 113225
PIP5K2A	NM_005028	hsa-miR-30c	PIP5K2A Phosphatidylinositol-4-phosphate 5-kinase, type II, alpha Hs.499029 T39411 5305 118773
PIP5K2A	NM_005028	hsa-miR-30e	PIP5K2A Phosphatidylinositol-4-phosphate 5-kinase, type II, alpha Hs.499029 T39411 5305 118773
PLCL2	NM_015184	hsa-miR-27a	PLCL2 Phospholipase C-like 2 Hs.202010 AA677432 23228 222856
PLEKHA1	NM_021622	hsa-miR-15a	PLEKHA1 Pleckstrin homology domain containing, family A (phosphoinositide binding specific) member 1 Hs.287830 N95011 59338 119935
PLEKHA1	NM_021622	hsa-miR-15b	PLEKHA1 Pleckstrin homology domain containing, family A (phosphoinositide binding specific) member 1 Hs.287830 N95011 59338 119935
PLEKHA1	NM_021622	hsa-miR-16	PLEKHA1 Pleckstrin homology domain containing, family A (phosphoinositide binding specific) member 1 Hs.287830 N95011 59338 119935
PLEKHA1	NM_021622	hsa-miR-195	PLEKHA1 Pleckstrin homology domain containing, family A (phosphoinositide binding specific) member 1 Hs.287830 N95011 59338 119935
PLEKHA1	NM_021622	hsa-miR-21	PLEKHA1 Pleckstrin homology domain containing, family A (phosphoinositide binding specific) member 1 Hs.287830 N95011 59338 119935
PLEKHA1	NM_021622	hsa-miR-29a	PLEKHA1 Pleckstrin homology domain containing, family A (phosphoinositide binding specific) member 1 Hs.287830 N95011 59338 119935
PLEKHA1	NM_021622	hsa-miR-29b	PLEKHA1 Pleckstrin homology domain containing, family A (phosphoinositide binding specific) member 1 Hs.287830 N95011 59338 119935
PLEKHA1	NM_021622	hsa-miR-29c	PLEKHA1 Pleckstrin homology domain containing, family A (phosphoinositide binding specific) member 1 Hs.287830 N95011 59338 119935
PLEKHA1	NM_021622	hsa-miR-92a	PLEKHA1 Pleckstrin homology domain containing, family A (phosphoinositide binding specific) member 1 Hs.287830 N95011 59338 119935
PLEKHA8	NM_032639	hsa-miR-369-3p	PLEKHA8 **Pleckstrin homology domain containing, family A (phosphoinositide binding specific) member 8 Hs.233495 R25734 84725 104037

PLEKHC1	NM 006832	hsa-let-7e	PLEKHC1 Pleckstrin homology domain containing, family C (with FERM domain) member 1 Hs.509343 H29252 10979 114766
PLEKHC1	NM 006832	hsa-miR-124	PLEKHC1 Pleckstrin homology domain containing, family C (with FERM domain) member 1 Hs.509343 H29252 10979 114766
PLEKHC1	NM 006832	hsa-miR-15a	PLEKHC1 Pleckstrin homology domain containing, family C (with FERM domain) member 1 Hs.509343 H29252 10979 114766
PLEKHC1	NM 006832	hsa-miR-15b	PLEKHC1 Pleckstrin homology domain containing, family C (with FERM domain) member 1 Hs.509343 H29252 10979 114766
PLEKHC1	NM 006832	hsa-miR-16	PLEKHC1 Pleckstrin homology domain containing, family C (with FERM domain) member 1 Hs.509343 H29252 10979 114766
PLEKHC1	NM 006832	hsa-miR-195	PLEKHC1 Pleckstrin homology domain containing, family C (with FERM domain) member 1 Hs.509343 H29252 10979 114766
PLEKHC1	NM 006832	hsa-miR-29a	PLEKHC1 Pleckstrin homology domain containing, family C (with FERM domain) member 1 Hs.509343 H29252 10979 114766
PLEKHC1	NM 006832	hsa-miR-29b	PLEKHC1 Pleckstrin homology domain containing, family C (with FERM domain) member 1 Hs.509343 H29252 10979 114766
PLEKHC1	NM 006832	hsa-miR-29c	PLEKHC1 Pleckstrin homology domain containing, family C (with FERM domain) member 1 Hs.509343 H29252 10979 114766
PMP22	NM 000304	hsa-miR-29a	PMP22 Peripheral myelin protein 22 Hs.372031 H26086 5376 224197
PMP22	NM 000304	hsa-miR-29b	PMP22 Peripheral myelin protein 22 Hs.372031 H26086 5376 224197
PMP22	NM 000304	hsa-miR-29c	PMP22 Peripheral myelin protein 22 Hs.372031 H26086 5376 224197
POLS	NM 006999	hsa-miR-27a	POLS Polymerase (DNA directed) sigma Hs.481542 AA465382 11044 221354
POLS	NM 006999	hsa-miR-92a	POLS Polymerase (DNA directed) sigma Hs.481542 AA465382 11044 221354
POU4F1	NM 006237	hsa-miR-19a	POU4F1 POU domain, class 4, transcription factor 1 Hs.211588 AA428196 RDC-1=POU domain transcription factor 5457 107372
POU4F1	NM 006237	hsa-miR-19b	POU4F1 POU domain, class 4, transcription factor 1 Hs.211588 AA428196 RDC-1=POU domain transcription factor 5457 107372
PPM1A	NM 177954	hsa-miR-124	PPM1A Protein phosphatase 1A (formerly 2C), magnesium-dependent, alpha isoform Hs.130036 Al365364 5494 311430
PPM1A	NM 177952	hsa-miR-15a	PPM1A Protein phosphatase 1A (formerly 2C), magnesium-dependent, alpha isoform Hs.130036 Al365364 5494 311430
PPM1A	NM 177952	hsa-miR-15b	PPM1A Protein phosphatase 1A (formerly 2C), magnesium-dependent, alpha isoform Hs.130036 Al365364 5494 311430
PPM1A	NM 177952	hsa-miR-16	PPM1A Protein phosphatase 1A (formerly 2C), magnesium-dependent, alpha isoform Hs.130036 Al365364 5494 311430
PPM1A	NM 177952	hsa-miR-195	PPM1A Protein phosphatase 1A (formerly 2C), magnesium-dependent, alpha isoform Hs.130036 Al365364 5494 311430
PPM1A	NM 177954	hsa-miR-29a	PPM1A Protein phosphatase 1A (formerly 2C), magnesium-dependent, alpha isoform Hs.130036 Al365364 5494 311430
PPM1A	NM 177954	hsa-miR-29b	PPM1A Protein phosphatase 1A (formerly 2C), magnesium-dependent, alpha isoform Hs.130036 Al365364 5494 311430
PPM1A	NM 177954	hsa-miR-29c	PPM1A Protein phosphatase 1A (formerly 2C), magnesium-dependent, alpha isoform Hs.130036 Al365364 5494 311430
PRDX2	NM 005809	hsa-miR-369-3p	PRDX2 Peroxiredoxin 2 Hs.432121 H68845 7001 112207
PRKCE	NM 005400	hsa-let-7d	PRKCE Protein kinase C, epsilon Hs.97432 H23225 5581 120893
PRKCE	NM 005400	hsa-miR-19a	PRKCE Protein kinase C, epsilon Hs.97432 H23225 5581 120893
PRKCE	NM 005400	hsa-miR-23a	PRKCE Protein kinase C, epsilon Hs.97432 H23225 5581 120893
PRKCE	NM 005400	hsa-miR-29a	PRKCE Protein kinase C, epsilon Hs.97432 H23225 5581 120893
PRKCE	NM 005400	hsa-miR-369-3p	PRKCE Protein kinase C, epsilon Hs.97432 H23225 5581 120893
PRKCE	NM 005400	hsa-miR-92a	PRKCE Protein kinase C, epsilon Hs.97432 H23225 5581 120893
PROM1	NM 006017	hsa-miR-142-5p	PROM1 Prominin 1 Hs.479220 R40057 AC133 antigen=five-transmembrane hematopoietic stem cell ant 8842 118920
PROM1	NM 006017	hsa-miR-23a	PROM1 Prominin 1 Hs.479220 R40057 AC133 antigen=five-transmembrane hematopoietic stem cell ant 8842 118920
PROM1	NM 006017	hsa-miR-29a	PROM1 Prominin 1 Hs.479220 R40057 AC133 antigen=five-transmembrane hematopoietic stem cell ant 8842 118920
PROM1	NM 006017	hsa-miR-29b	PROM1 Prominin 1 Hs.479220 R40057 AC133 antigen=five-transmembrane hematopoietic stem cell ant 8842 118920
PROM1	NM 006017	hsa-miR-29c	PROM1 Prominin 1 Hs.479220 R40057 AC133 antigen=five-transmembrane hematopoietic stem cell ant 8842 118920
PROM1	NM 006017	hsa-miR-30c	PROM1 Prominin 1 Hs.479220 R40057 AC133 antigen=five-transmembrane hematopoietic stem cell ant 8842 118920
PROM1	NM 006017	hsa-miR-369-3p	PROM1 Prominin 1 Hs.479220 R40057 AC133 antigen=five-transmembrane hematopoietic stem cell ant 8842 118920
PTTG1	NM 004219	hsa-miR-19a	PTTG1 Pituitary tumor-transforming 1 Hs.350966 Al362866 9232 311002
PTTG1	NM 004219	hsa-miR-19b	PTTG1 Pituitary tumor-transforming 1 Hs.350966 Al362866 9232 311002
QKI	NM 206855	hsa-miR-106a	QKI **Quaking homolog, KH domain RNA binding (mouse) Hs.510324 AA452531 9444 226251
QKI	NM 206855	hsa-miR-124	QKI **Quaking homolog, KH domain RNA binding (mouse) Hs.510324 AA452531 9444 226251
QKI	NM 206855	hsa-miR-142-5p	QKI **Quaking homolog, KH domain RNA binding (mouse) Hs.510324 AA452531 9444 226251
QKI	NM 206855	hsa-miR-155	QKI **Quaking homolog, KH domain RNA binding (mouse) Hs.510324 AA452531 9444 226251
QKI	NM 206855	hsa-miR-15a	QKI **Quaking homolog, KH domain RNA binding (mouse) Hs.510324 AA452531 9444 226251
QKI	NM 206855	hsa-miR-15b	QKI **Quaking homolog, KH domain RNA binding (mouse) Hs.510324 AA452531 9444 226251
QKI	NM 206855	hsa-miR-16	QKI **Quaking homolog, KH domain RNA binding (mouse) Hs.510324 AA452531 9444 226251
QKI	NM 206855	hsa-miR-17	QKI **Quaking homolog, KH domain RNA binding (mouse) Hs.510324 AA452531 9444 226251
QKI	NM 206855	hsa-miR-18a	QKI **Quaking homolog, KH domain RNA binding (mouse) Hs.510324 AA452531 9444 226251
QKI	NM 206855	hsa-miR-195	QKI **Quaking homolog, KH domain RNA binding (mouse) Hs.510324 AA452531 9444 226251
QKI	NM 206855	hsa-miR-19a	QKI **Quaking homolog, KH domain RNA binding (mouse) Hs.510324 AA452531 9444 226251
QKI	NM 206855	hsa-miR-19b	QKI **Quaking homolog, KH domain RNA binding (mouse) Hs.510324 AA452531 9444 226251
QKI	NM 206855	hsa-miR-20a	QKI **Quaking homolog, KH domain RNA binding (mouse) Hs.510324 AA452531 9444 226251

QKI	NM 206855	hsa-miR-23a	QKI **Quaking homolog, KH domain RNA binding (mouse) Hs.510324 AA452531 9444 226251
QKI	NM 206855	hsa-miR-27a	QKI **Quaking homolog, KH domain RNA binding (mouse) Hs.510324 AA452531 9444 226251
QKI	NM 206855	hsa-miR-29a	QKI **Quaking homolog, KH domain RNA binding (mouse) Hs.510324 AA452531 9444 226251
QKI	NM 206855	hsa-miR-29b	QKI **Quaking homolog, KH domain RNA binding (mouse) Hs.510324 AA452531 9444 226251
QKI	NM 206855	hsa-miR-29c	QKI **Quaking homolog, KH domain RNA binding (mouse) Hs.510324 AA452531 9444 226251
QKI	NM 206855	hsa-miR-30c	QKI **Quaking homolog, KH domain RNA binding (mouse) Hs.510324 AA452531 9444 226251
QKI	NM 206855	hsa-miR-30e	QKI **Quaking homolog, KH domain RNA binding (mouse) Hs.510324 AA452531 9444 226251
QKI	NM 206855	hsa-miR-369-3p	QKI **Quaking homolog, KH domain RNA binding (mouse) Hs.510324 AA452531 9444 226251
QKI	NM 206855	hsa-miR-92a	QKI **Quaking homolog, KH domain RNA binding (mouse) Hs.510324 AA452531 9444 226251
RAB1A	NM 004161	hsa-miR-142-5p	RAB1A **RAB1A, member RAS oncogene family Hs.310645 AA463630 5861 106725
RAB1A	NM 004161	hsa-miR-155	RAB1A **RAB1A, member RAS oncogene family Hs.310645 AA463630 5861 106725
RAB1A	NM 004161	hsa-miR-15a	RAB1A **RAB1A, member RAS oncogene family Hs.310645 AA463630 5861 106725
RAB1A	NM 004161	hsa-miR-19a	RAB1A **RAB1A, member RAS oncogene family Hs.310645 AA463630 5861 106725
RAB1A	NM 004161	hsa-miR-19b	RAB1A **RAB1A, member RAS oncogene family Hs.310645 AA463630 5861 106725
RAB43	NM 198490	hsa-miR-124	RAB43 RAB43, member RAS oncogene family Hs.546542 AA155803 339122 224496
RAD50	NM 005732	hsa-miR-142-5p	Hypothetical gene supported by AL713721 Hs.483359 AA126482 441109 106162
RALA	NM 005402	hsa-miR-124	RALA V-ral simian leukemia viral oncogene homolog A (ras related) Hs.6906 H97948 5898 100960
RALA	NM 005402	hsa-miR-19a	RALA V-ral simian leukemia viral oncogene homolog A (ras related) Hs.6906 H97948 5898 100960
RALA	NM 005402	hsa-miR-27a	RALA V-ral simian leukemia viral oncogene homolog A (ras related) Hs.6906 H97948 5898 100960
RALA	NM 005402	hsa-miR-29a	RALA V-ral simian leukemia viral oncogene homolog A (ras related) Hs.6906 H97948 5898 100960
RALA	NM 005402	hsa-miR-29b	RALA V-ral simian leukemia viral oncogene homolog A (ras related) Hs.6906 H97948 5898 100960
RALA	NM 005402	hsa-miR-29c	RALA V-ral simian leukemia viral oncogene homolog A (ras related) Hs.6906 H97948 5898 100960
RALA	NM 005402	hsa-miR-30a	RALA V-ral simian leukemia viral oncogene homolog A (ras related) Hs.6906 H97948 5898 100960
RALA	NM 005402	hsa-miR-30e	RALA V-ral simian leukemia viral oncogene homolog A (ras related) Hs.6906 H97948 5898 100960
RALA	NM 005402	hsa-miR-369-3p	RALA V-ral simian leukemia viral oncogene homolog A (ras related) Hs.6906 H97948 5898 100960
RANBP10	NM 020850	hsa-miR-124	RANBP10 RAN binding protein 10 Hs.368569 H59628 57610 308519
RANBP10	NM 020850	hsa-miR-30c	RANBP10 RAN binding protein 10 Hs.368569 H59628 57610 308519
RANBP10	NM 020850	hsa-miR-30e	RANBP10 RAN binding protein 10 Hs.368569 H59628 57610 308519
RANBP10	NM 020850	hsa-miR-369-3p	RANBP10 RAN binding protein 10 Hs.368569 H59628 57610 308519
RAP2A	NM 021033	hsa-miR-10a	RAP2A RAP2A, member of RAS oncogene family Hs.508480 N70768 5911 115957
RAP2A	NM 021033	hsa-miR-10b	RAP2A RAP2A, member of RAS oncogene family Hs.508480 N70768 5911 115957
RAP2A	NM 021033	hsa-miR-124	RAP2A RAP2A, member of RAS oncogene family Hs.508480 N70768 5911 115957
RASD1	NM 016084	hsa-miR-106a	RASD1 RAS, dexamethasone-induced 1 Hs.25829 A1014398 51655 307161
RASD1	NM 016084	hsa-miR-17	RASD1 RAS, dexamethasone-induced 1 Hs.25829 A1014398 51655 307161
RASD1	NM 016084	hsa-miR-20a	RASD1 RAS, dexamethasone-induced 1 Hs.25829 A1014398 51655 307161
RASD1	NM 016084	hsa-miR-30c	RASD1 RAS, dexamethasone-induced 1 Hs.25829 A1014398 51655 307161
RASD1	NM 016084	hsa-miR-30e	RASD1 RAS, dexamethasone-induced 1 Hs.25829 A1014398 51655 307161
RGS16	NM 002928	hsa-let-7a	RGS16 Regulator of G-protein signalling 16 Hs.413297 AA453774 6004 115991
RGS16	NM 002928	hsa-let-7b	RGS16 Regulator of G-protein signalling 16 Hs.413297 AA453774 6004 115991
RGS16	NM 002928	hsa-let-7c	RGS16 Regulator of G-protein signalling 16 Hs.413297 AA453774 6004 115991
RGS16	NM 002928	hsa-let-7d	RGS16 Regulator of G-protein signalling 16 Hs.413297 AA453774 6004 115991
RGS16	NM 002928	hsa-let-7e	RGS16 Regulator of G-protein signalling 16 Hs.413297 AA453774 6004 115991
RGS16	NM 002928	hsa-let-7f	RGS16 Regulator of G-protein signalling 16 Hs.413297 AA453774 6004 115991
RGS16	NM 002928	hsa-miR-98	RGS16 Regulator of G-protein signalling 16 Hs.413297 AA453774 6004 115991
RIF1	NM 018151	hsa-miR-19a	RIF1 Receptor-interacting factor 1 Hs.25245 R37817 55791 105216
RIF1	NM 018151	hsa-miR-19b	RIF1 Receptor-interacting factor 1 Hs.25245 R37817 55791 105216
RIF1	NM 018151	hsa-miR-30c	RIF1 Receptor-interacting factor 1 Hs.25245 R37817 55791 105216
RIF1	NM 018151	hsa-miR-30e	RIF1 Receptor-interacting factor 1 Hs.25245 R37817 55791 105216
RIOK3	NM 145906	hsa-let-7a	RIOK3 RIO kinase 3 (yeast) Hs.445511 T54144 8780 111200
RIOK3	NM 145906	hsa-let-7b	RIOK3 RIO kinase 3 (yeast) Hs.445511 T54144 8780 111200
RIOK3	NM 145906	hsa-let-7c	RIOK3 RIO kinase 3 (yeast) Hs.445511 T54144 8780 111200
RIOK3	NM 145906	hsa-let-7d	RIOK3 RIO kinase 3 (yeast) Hs.445511 T54144 8780 111200
RIOK3	NM 145906	hsa-let-7e	RIOK3 RIO kinase 3 (yeast) Hs.445511 T54144 8780 111200

RIOK3	NM 145906	hsa-let-7f	RIOK3 RIO kinase 3 (yeast) Hs.445511 T54144 8780 111200
RIOK3	NM 003831	hsa-miR-98	RIOK3 RIO kinase 3 (yeast) Hs.445511 T54144 8780 111200
RKHD2	NM 016626	hsa-miR-15a	RKHD2 Ring finger and KH domain containing 2 Hs.465144 AI015297 51320 226384
RKHD2	NM 016626	hsa-miR-15b	RKHD2 Ring finger and KH domain containing 2 Hs.465144 AI015297 51320 226384
RKHD2	NM 016626	hsa-miR-16	RKHD2 Ring finger and KH domain containing 2 Hs.465144 AI015297 51320 226384
RKHD2	NM 016626	hsa-miR-195	RKHD2 Ring finger and KH domain containing 2 Hs.465144 AI015297 51320 226384
RKHD2	NM 016626	hsa-miR-19a	RKHD2 Ring finger and KH domain containing 2 Hs.465144 AI015297 51320 226384
RKHD2	NM 016626	hsa-miR-19b	RKHD2 Ring finger and KH domain containing 2 Hs.465144 AI015297 51320 226384
RKHD2	NM 016626	hsa-miR-23a	RKHD2 Ring finger and KH domain containing 2 Hs.465144 AI015297 51320 226384
RKHD2	NM 016626	hsa-miR-30c	RKHD2 Ring finger and KH domain containing 2 Hs.465144 AI015297 51320 226384
RKHD2	NM 016626	hsa-miR-30e	RKHD2 Ring finger and KH domain containing 2 Hs.465144 AI015297 51320 226384
RKHD2	NM 016626	hsa-miR-369-3p	RKHD2 Ring finger and KH domain containing 2 Hs.465144 AI015297 51320 226384
RNF24	NM 007219	hsa-miR-27a	RNF24 Ring finger protein 24 Hs.114180 AA995559 11237 309227
RNPC1	NM 183425	hsa-let-7a	RNPC1 RNA-binding region (RNP1, RRM) containing 1 Hs.236361 AW072104 55544 330915
RNPC1	NM 183425	hsa-let-7b	RNPC1 RNA-binding region (RNP1, RRM) containing 1 Hs.236361 AW072104 55544 330915
RNPC1	NM 183425	hsa-let-7c	RNPC1 RNA-binding region (RNP1, RRM) containing 1 Hs.236361 AW072104 55544 330915
RNPC1	NM 183425	hsa-let-7d	RNPC1 RNA-binding region (RNP1, RRM) containing 1 Hs.236361 AW072104 55544 330915
RNPC1	NM 183425	hsa-let-7e	RNPC1 RNA-binding region (RNP1, RRM) containing 1 Hs.236361 AW072104 55544 330915
RNPC1	NM 183425	hsa-let-7f	RNPC1 RNA-binding region (RNP1, RRM) containing 1 Hs.236361 AW072104 55544 330915
RNPC1	NM 183425	hsa-miR-98	RNPC1 RNA-binding region (RNP1, RRM) containing 1 Hs.236361 AW072104 55544 330915
RPIA	NM 144563	hsa-miR-124	RPIA Ribose 5-phosphate isomerase A (ribose 5-phosphate epimerase) Hs.469264 N20072 22934 111875
RSC1A1	NM 006511	hsa-miR-29b	RSC1A1 Regulatory solute carrier protein, family 1, member 1 Hs.239459 H02333 6248 109543
RUNX3	NM 004350	hsa-miR-19a	RUNX3 Runt-related transcription factor 3 Hs.170019 W94742 864 239268
RUNX3	NM 004350	hsa-miR-19b	RUNX3 Runt-related transcription factor 3 Hs.170019 W94742 864 239268
SACS	NM 014363	hsa-miR-106a	SACS Spastic ataxia of Charlevoix-Saguenay (sacsin) Hs.159492 AA481194 26278 308982
SACS	NM 014363	hsa-miR-142-5p	SACS Spastic ataxia of Charlevoix-Saguenay (sacsin) Hs.159492 AA481194 26278 308982
SACS	NM 014363	hsa-miR-17	SACS Spastic ataxia of Charlevoix-Saguenay (sacsin) Hs.159492 AA481194 26278 308982
SACS	NM 014363	hsa-miR-18a	SACS Spastic ataxia of Charlevoix-Saguenay (sacsin) Hs.159492 AA481194 26278 308982
SACS	NM 014363	hsa-miR-20a	SACS Spastic ataxia of Charlevoix-Saguenay (sacsin) Hs.159492 AA481194 26278 308982
SACS	NM 014363	hsa-miR-23a	SACS Spastic ataxia of Charlevoix-Saguenay (sacsin) Hs.159492 AA481194 26278 308982
SACS	NM 014363	hsa-miR-30c	SACS Spastic ataxia of Charlevoix-Saguenay (sacsin) Hs.159492 AA481194 26278 308982
SCAMP5	NM 138967	hsa-miR-29a	SCAMP5 Secretory carrier membrane protein 5 Hs.374180 AA663609 192683 223036
SCAMP5	NM 138967	hsa-miR-29b	SCAMP5 Secretory carrier membrane protein 5 Hs.374180 AA663609 192683 223036
SCAMP5	NM 138967	hsa-miR-29c	SCAMP5 Secretory carrier membrane protein 5 Hs.374180 AA663609 192683 223036
SELENBP1	NM 003944	hsa-miR-15a	SELENBP1 Selenium binding protein 1 Hs.334841 R90933 8991 221649
SELENBP1	NM 003944	hsa-miR-15b	SELENBP1 Selenium binding protein 1 Hs.334841 R90933 8991 221649
SELENBP1	NM 003944	hsa-miR-16	SELENBP1 Selenium binding protein 1 Hs.334841 R90933 8991 221649
SELENBP1	NM 003944	hsa-miR-195	SELENBP1 Selenium binding protein 1 Hs.334841 R90933 8991 221649
SELENBP1	NM 003944	hsa-miR-23a	SELENBP1 Selenium binding protein 1 Hs.334841 R90933 8991 221649
SELENBP1	NM 003944	hsa-miR-27a	SELENBP1 Selenium binding protein 1 Hs.334841 R90933 8991 221649
SELK	NM 021237	hsa-let-7d	SELK Selenoprotein K Hs.58471 AA064917 58515 118280
SELK	NM 021237	hsa-miR-155	SELK Selenoprotein K Hs.58471 AA064917 58515 118280
SELK	NM 021237	hsa-miR-21	SELK Selenoprotein K Hs.58471 AA064917 58515 118280
SERPINB9	NM 004155	hsa-let-7a	SERPINB9 Serine (or cysteine) proteinase inhibitor, clade B (ovalbumin), member 9 Hs.104879 AA147540 5272 103188
SERPINB9	NM 004155	hsa-miR-15a	SERPINB9 Serine (or cysteine) proteinase inhibitor, clade B (ovalbumin), member 9 Hs.104879 AA147540 5272 103188
SERPINB9	NM 004155	hsa-miR-29a	SERPINB9 Serine (or cysteine) proteinase inhibitor, clade B (ovalbumin), member 9 Hs.104879 AA147540 5272 103188
SERPINB9	NM 004155	hsa-miR-29b	SERPINB9 Serine (or cysteine) proteinase inhibitor, clade B (ovalbumin), member 9 Hs.104879 AA147540 5272 103188
SERPINB9	NM 004155	hsa-miR-29c	SERPINB9 Serine (or cysteine) proteinase inhibitor, clade B (ovalbumin), member 9 Hs.104879 AA147540 5272 103188
SERPINB9	NM 004155	hsa-miR-369-3p	SERPINB9 Serine (or cysteine) proteinase inhibitor, clade B (ovalbumin), member 9 Hs.104879 AA147540 5272 103188
SESN3	NM 144665	hsa-miR-155	SESN3 Sestrin 3 Hs.120633 W95428 143686 108178
SESN3	NM 144665	hsa-miR-92a	SESN3 Sestrin 3 Hs.120633 W95428 143686 108178
SGPP1	NM 030791	hsa-miR-124	SGPP1 Sphingosine-1-phosphate phosphatase 1 Hs.24678 H03177 81537 100315
SGPP1	NM 030791	hsa-miR-27a	SGPP1 Sphingosine-1-phosphate phosphatase 1 Hs.24678 H03177 81537 100315

SGPP1	NM_030791	hsa-miR-92a	SGPP1 Sphingosine-1-phosphate phosphatase 1 Hs.24678 H03177 81537 100315
SH3MD2	NM_020870	hsa-miR-18a	SH3MD2 SH3 multiple domains 2 Hs.301804 AA954262 57630 310191
SH3MD2	NM_020870	hsa-miR-23a	SH3MD2 SH3 multiple domains 2 Hs.301804 AA954262 57630 310191
SH3MD2	NM_020870	hsa-miR-27a	SH3MD2 SH3 multiple domains 2 Hs.301804 AA954262 57630 310191
SH3MD2	NM_020870	hsa-miR-30c	SH3MD2 SH3 multiple domains 2 Hs.301804 AA954262 57630 310191
SH3MD2	NM_020870	hsa-miR-30e	SH3MD2 SH3 multiple domains 2 Hs.301804 AA954262 57630 310191
SIPA1L1	NM_015556	hsa-miR-124	SIPA1L1 Signal-induced proliferation-associated 1 like 1 Hs.208846 R37938 26037 226364
SIPA1L1	NM_015556	hsa-miR-18a	SIPA1L1 Signal-induced proliferation-associated 1 like 1 Hs.208846 R37938 26037 226364
SIPA1L2	NM_020808	hsa-miR-15a	SIPA1L2 Signal-induced proliferation-associated 1 like 2 Hs.268774 AA464598 57568 101920
SIPA1L2	NM_020808	hsa-miR-15b	SIPA1L2 Signal-induced proliferation-associated 1 like 2 Hs.268774 AA464598 57568 101920
SIPA1L2	NM_020808	hsa-miR-16	SIPA1L2 Signal-induced proliferation-associated 1 like 2 Hs.268774 AA464598 57568 101920
SIPA1L2	NM_020808	hsa-miR-195	SIPA1L2 Signal-induced proliferation-associated 1 like 2 Hs.268774 AA464598 57568 101920
SIPA1L2	NM_020808	hsa-miR-19a	SIPA1L2 Signal-induced proliferation-associated 1 like 2 Hs.268774 AA464598 57568 101920
SIPA1L2	NM_020808	hsa-miR-19b	SIPA1L2 Signal-induced proliferation-associated 1 like 2 Hs.268774 AA464598 57568 101920
SIPA1L2	NM_020808	hsa-miR-23a	SIPA1L2 Signal-induced proliferation-associated 1 like 2 Hs.268774 AA464598 57568 101920
SIRT1	NM_012238	hsa-miR-124	SIRT1 Sirtuin (silent mating type information regulation 2 homolog) 1 (S. cerevisiae) Hs.369779 AA460952 23411 109813
SIRT1	NM_012238	hsa-miR-142-5p	SIRT1 Sirtuin (silent mating type information regulation 2 homolog) 1 (S. cerevisiae) Hs.369779 AA460952 23411 109813
SIRT1	NM_012238	hsa-miR-155	SIRT1 Sirtuin (silent mating type information regulation 2 homolog) 1 (S. cerevisiae) Hs.369779 AA460952 23411 109813
SIRT1	NM_012238	hsa-miR-19a	SIRT1 Sirtuin (silent mating type information regulation 2 homolog) 1 (S. cerevisiae) Hs.369779 AA460952 23411 109813
SIRT1	NM_012238	hsa-miR-23a	SIRT1 Sirtuin (silent mating type information regulation 2 homolog) 1 (S. cerevisiae) Hs.369779 AA460952 23411 109813
SIRT1	NM_012238	hsa-miR-29a	SIRT1 Sirtuin (silent mating type information regulation 2 homolog) 1 (S. cerevisiae) Hs.369779 AA460952 23411 109813
SIRT1	NM_012238	hsa-miR-30c	SIRT1 Sirtuin (silent mating type information regulation 2 homolog) 1 (S. cerevisiae) Hs.369779 AA460952 23411 109813
SIRT1	NM_012238	hsa-miR-30e	SIRT1 Sirtuin (silent mating type information regulation 2 homolog) 1 (S. cerevisiae) Hs.369779 AA460952 23411 109813
SIRT1	NM_012238	hsa-miR-369-3p	SIRT1 Sirtuin (silent mating type information regulation 2 homolog) 1 (S. cerevisiae) Hs.369779 AA460952 23411 109813
SIRT1	NM_012238	hsa-miR-92a	SIRT1 Sirtuin (silent mating type information regulation 2 homolog) 1 (S. cerevisiae) Hs.369779 AA460952 23411 109813
SLC11A2	NM_000617	hsa-let-7d	SLC11A2 Solute carrier family 11 (proton-coupled divalent metal ion transporters), member 2 Hs.505545 R09378 4891 106967
SLC11A2	NM_000617	hsa-miR-15a	SLC11A2 Solute carrier family 11 (proton-coupled divalent metal ion transporters), member 2 Hs.505545 R09378 4891 106967
SLC11A2	NM_000617	hsa-miR-15b	SLC11A2 Solute carrier family 11 (proton-coupled divalent metal ion transporters), member 2 Hs.505545 R09378 4891 106967
SLC11A2	NM_000617	hsa-miR-19a	SLC11A2 Solute carrier family 11 (proton-coupled divalent metal ion transporters), member 2 Hs.505545 R09378 4891 106967
SLC11A2	NM_000617	hsa-miR-19b	SLC11A2 Solute carrier family 11 (proton-coupled divalent metal ion transporters), member 2 Hs.505545 R09378 4891 106967
SLC38A2	NM_018976	hsa-miR-10a	SLC38A2 Solute carrier family 38, member 2 Hs.221847 AA081106 54407 315845
SLC38A2	NM_018976	hsa-miR-10b	SLC38A2 Solute carrier family 38, member 2 Hs.221847 AA081106 54407 315845
SLC38A2	NM_018976	hsa-miR-30c	SLC38A2 Solute carrier family 38, member 2 Hs.221847 AA081106 54407 315845
SLC38A2	NM_018976	hsa-miR-30e	SLC38A2 Solute carrier family 38, member 2 Hs.221847 AA081106 54407 315845
SLC38A2	NM_018976	hsa-miR-92a	SLC38A2 Solute carrier family 38, member 2 Hs.221847 AA081106 54407 315845
SORL1	NM_003105	hsa-miR-106a	SORL1 Sortilin-related receptor, L(DLR class) A repeats-containing Hs.368592 N50656 6653 111561
SORL1	NM_003105	hsa-miR-17	SORL1 Sortilin-related receptor, L(DLR class) A repeats-containing Hs.368592 N50656 6653 111561
SORL1	NM_003105	hsa-miR-20a	SORL1 Sortilin-related receptor, L(DLR class) A repeats-containing Hs.368592 N50656 6653 111561
SORL1	NM_003105	hsa-miR-92a	SORL1 Sortilin-related receptor, L(DLR class) A repeats-containing Hs.368592 N50656 6653 111561
SP100	NM_003113	hsa-miR-19a	SP140 SP140 nuclear body protein Hs.471576 H66483 LYSP100=SP140 11262 114358
SP100	NM_003113	hsa-miR-29a	SP140 SP140 nuclear body protein Hs.471576 H66483 LYSP100=SP140 11262 114358
SP100	NM_003113	hsa-miR-29b	SP140 SP140 nuclear body protein Hs.471576 H66483 LYSP100=SP140 11262 114358
SP100	NM_003113	hsa-miR-29c	SP140 SP140 nuclear body protein Hs.471576 H66483 LYSP100=SP140 11262 114358
SP100	NM_003113	hsa-miR-30c	SP140 SP140 nuclear body protein Hs.471576 H66483 LYSP100=SP140 11262 114358
SP140	NM_001005176	hsa-let-7a	SP140 SP140 nuclear body protein Hs.471576 H66483 LYSP100=SP140 11262 114358
SP140	NM_001005176	hsa-let-7b	SP140 SP140 nuclear body protein Hs.471576 H66483 LYSP100=SP140 11262 114358
SP140	NM_001005176	hsa-let-7c	SP140 SP140 nuclear body protein Hs.471576 H66483 LYSP100=SP140 11262 114358
SP140	NM_001005176	hsa-let-7d	SP140 SP140 nuclear body protein Hs.471576 H66483 LYSP100=SP140 11262 114358
SP140	NM_001005176	hsa-let-7e	SP140 SP140 nuclear body protein Hs.471576 H66483 LYSP100=SP140 11262 114358
SP140	NM_001005176	hsa-let-7f	SP140 SP140 nuclear body protein Hs.471576 H66483 LYSP100=SP140 11262 114358
SP140	NM_001005176	hsa-miR-98	SP140 SP140 nuclear body protein Hs.471576 H66483 LYSP100=SP140 11262 114358
SPARC	NM_003118	hsa-miR-106a	SPARC Secreted protein, acidic, cysteine-rich (osteonectin) Hs.111779 N95151 6678 112917
SPARC	NM_003118	hsa-miR-155	SPARC Secreted protein, acidic, cysteine-rich (osteonectin) Hs.111779 N95151 6678 112917

SPARC	NM 003118	hsa-miR-19a	SPARC Secreted protein, acidic, cysteine-rich (osteonectin) Hs.111779 N95151 6678 112917
SPARC	NM 003118	hsa-miR-20a	SPARC Secreted protein, acidic, cysteine-rich (osteonectin) Hs.111779 N95151 6678 112917
SPARC	NM 003118	hsa-miR-23a	SPARC Secreted protein, acidic, cysteine-rich (osteonectin) Hs.111779 N95151 6678 112917
SPARC	NM 003118	hsa-miR-29a	SPARC Secreted protein, acidic, cysteine-rich (osteonectin) Hs.111779 N95151 6678 112917
SPARC	NM 003118	hsa-miR-29b	SPARC Secreted protein, acidic, cysteine-rich (osteonectin) Hs.111779 N95151 6678 112917
SPARC	NM 003118	hsa-miR-29c	SPARC Secreted protein, acidic, cysteine-rich (osteonectin) Hs.111779 N95151 6678 112917
SPRY1	NM 199327	hsa-miR-124	SPRY1 Sprouty homolog 1, antagonist of FGF signaling (Drosophila) Hs.436944 AA055439 10252 119371
SPRY1	NM 199327	hsa-miR-142-5p	SPRY1 Sprouty homolog 1, antagonist of FGF signaling (Drosophila) Hs.436944 AA055439 10252 119371
SPRY1	NM 199327	hsa-miR-21	SPRY1 Sprouty homolog 1, antagonist of FGF signaling (Drosophila) Hs.436944 AA055439 10252 119371
SPRY1	NM 199327	hsa-miR-29a	SPRY1 Sprouty homolog 1, antagonist of FGF signaling (Drosophila) Hs.436944 AA055439 10252 119371
SPRY1	NM 199327	hsa-miR-29b	SPRY1 Sprouty homolog 1, antagonist of FGF signaling (Drosophila) Hs.436944 AA055439 10252 119371
SPRY1	NM 199327	hsa-miR-29c	SPRY1 Sprouty homolog 1, antagonist of FGF signaling (Drosophila) Hs.436944 AA055439 10252 119371
SPRY1	NM 199327	hsa-miR-369-3p	SPRY1 Sprouty homolog 1, antagonist of FGF signaling (Drosophila) Hs.436944 AA055439 10252 119371
SPTBN2	NM 006946	hsa-miR-124	SPTBN2 Spectrin, beta, non-erythrocytic 2 Hs.26915 H28127 6712 106530
SPTBN2	NM 006946	hsa-miR-15a	SPTBN2 Spectrin, beta, non-erythrocytic 2 Hs.26915 H28127 6712 106530
SPTBN2	NM 006946	hsa-miR-15b	SPTBN2 Spectrin, beta, non-erythrocytic 2 Hs.26915 H28127 6712 106530
SPTBN2	NM 006946	hsa-miR-16	SPTBN2 Spectrin, beta, non-erythrocytic 2 Hs.26915 H28127 6712 106530
SPTBN2	NM 006946	hsa-miR-195	SPTBN2 Spectrin, beta, non-erythrocytic 2 Hs.26915 H28127 6712 106530
ST18	NM 014682	hsa-miR-23a	ST18 Suppression of tumorigenicity 18 (breast carcinoma) (zinc finger protein) Hs.549092 AA082192 9705 103642
ST3GAL5	NM 003896	hsa-miR-19a	SIAT9 ST3 beta-galactoside alpha-2,3-sialyltransferase 5 Hs.415117 R88773 8869 308619
ST3GAL5	NM 003896	hsa-miR-19b	SIAT9 ST3 beta-galactoside alpha-2,3-sialyltransferase 5 Hs.415117 R88773 8869 308619
ST8SIA4	NM 005668	hsa-miR-106a	SIAT8D ST8 alpha-N-acetylneuraminide alpha-2,8-sialyltransferase 4 Hs.308628 AA994501 7903 307081
ST8SIA4	NM 175052	hsa-miR-124	SIAT8D ST8 alpha-N-acetylneuraminide alpha-2,8-sialyltransferase 4 Hs.308628 AA994501 7903 307081
ST8SIA4	NM 175052	hsa-miR-17	SIAT8D ST8 alpha-N-acetylneuraminide alpha-2,8-sialyltransferase 4 Hs.308628 AA994501 7903 307081
ST8SIA4	NM 005668	hsa-miR-19a	SIAT8D ST8 alpha-N-acetylneuraminide alpha-2,8-sialyltransferase 4 Hs.308628 AA994501 7903 307081
ST8SIA4	NM 005668	hsa-miR-19b	SIAT8D ST8 alpha-N-acetylneuraminide alpha-2,8-sialyltransferase 4 Hs.308628 AA994501 7903 307081
ST8SIA4	NM 005668	hsa-miR-20a	SIAT8D ST8 alpha-N-acetylneuraminide alpha-2,8-sialyltransferase 4 Hs.308628 AA994501 7903 307081
ST8SIA4	NM 005668	hsa-miR-30c	SIAT8D ST8 alpha-N-acetylneuraminide alpha-2,8-sialyltransferase 4 Hs.308628 AA994501 7903 307081
ST8SIA4	NM 005668	hsa-miR-30e	SIAT8D ST8 alpha-N-acetylneuraminide alpha-2,8-sialyltransferase 4 Hs.308628 AA994501 7903 307081
STAM	NM 003473	hsa-miR-142-5p	STAM Signal transducing adaptor molecule (SH3 domain and ITAM motif) 1 Hs.441498 AI359446 8027 308813
STX6	NM 005819	hsa-miR-106a	STX6 Syntaxin 6 Hs.518417 AI339089 10228 308307
STX6	NM 005819	hsa-miR-17	STX6 Syntaxin 6 Hs.518417 AI339089 10228 308307
STX6	NM 005819	hsa-miR-20a	STX6 Syntaxin 6 Hs.518417 AI339089 10228 308307
STX6	NM 005819	hsa-miR-27a	STX6 Syntaxin 6 Hs.518417 AI339089 10228 308307
SUZ12	NM 015355	hsa-miR-19a	SUZ12 Suppressor of zeste 12 homolog (Drosophila) Hs.462732 AA521347 23512 110576
SUZ12	NM 015355	hsa-miR-19b	SUZ12 Suppressor of zeste 12 homolog (Drosophila) Hs.462732 AA521347 23512 110576
TAL1	NM 003189	hsa-miR-106a	TAL1 T-cell acute lymphocytic leukemia 1 Hs.73828 T51236 scl=tal-1=T-cell acute lymphocytic leukemia 1 6886 113362
TAL1	NM 003189	hsa-miR-124	TAL1 T-cell acute lymphocytic leukemia 1 Hs.73828 T51236 scl=tal-1=T-cell acute lymphocytic leukemia 1 6886 113362
TAL1	NM 003189	hsa-miR-17	TAL1 T-cell acute lymphocytic leukemia 1 Hs.73828 T51236 scl=tal-1=T-cell acute lymphocytic leukemia 1 6886 113362
TAL1	NM 003189	hsa-miR-19a	TAL1 T-cell acute lymphocytic leukemia 1 Hs.73828 T51236 scl=tal-1=T-cell acute lymphocytic leukemia 1 6886 113362
TAL1	NM 003189	hsa-miR-20a	TAL1 T-cell acute lymphocytic leukemia 1 Hs.73828 T51236 scl=tal-1=T-cell acute lymphocytic leukemia 1 6886 113362
TAL1	NM 003189	hsa-miR-29a	TAL1 T-cell acute lymphocytic leukemia 1 Hs.73828 T51236 scl=tal-1=T-cell acute lymphocytic leukemia 1 6886 113362
TCF4	NM 003199	hsa-miR-155	TCF4 Transcription factor 4 Hs.200285 AA953301 6925 310651
TCTEL1	NM 006519	hsa-miR-106a	TCTEL1 T-complex-associated-testis-expressed 1-like 1 Hs.445999 N22889 6993 220883
TCTEL1	NM 006519	hsa-miR-124	TCTEL1 T-complex-associated-testis-expressed 1-like 1 Hs.445999 N22889 6993 220883
TCTEL1	NM 006519	hsa-miR-142-5p	TCTEL1 T-complex-associated-testis-expressed 1-like 1 Hs.445999 N22889 6993 220883
TCTEL1	NM 006519	hsa-miR-17	TCTEL1 T-complex-associated-testis-expressed 1-like 1 Hs.445999 N22889 6993 220883
TCTEL1	NM 006519	hsa-miR-18a	TCTEL1 T-complex-associated-testis-expressed 1-like 1 Hs.445999 N22889 6993 220883
TCTEL1	NM 006519	hsa-miR-20a	TCTEL1 T-complex-associated-testis-expressed 1-like 1 Hs.445999 N22889 6993 220883
TCTEL1	NM 006519	hsa-miR-23a	TCTEL1 T-complex-associated-testis-expressed 1-like 1 Hs.445999 N22889 6993 220883
TCTEL1	NM 006519	hsa-miR-27a	TCTEL1 T-complex-associated-testis-expressed 1-like 1 Hs.445999 N22889 6993 220883
TCTEL1	NM 006519	hsa-miR-29a	TCTEL1 T-complex-associated-testis-expressed 1-like 1 Hs.445999 N22889 6993 220883
TCTEL1	NM 006519	hsa-miR-29b	TCTEL1 T-complex-associated-testis-expressed 1-like 1 Hs.445999 N22889 6993 220883

TCTEL1	NM_006519	hsa-miR-29c	TCTEL1 T-complex-associated-testis-expressed 1-like 1 Hs.445999 N22889 6993 220883
TEAD1	NM_021961	hsa-miR-124	TEAD1 TEA domain family member 1 (SV40 transcriptional enhancer factor) Hs.546308 AA039369 7003 116352
TEAD1	NM_021961	hsa-miR-23a	TEAD1 TEA domain family member 1 (SV40 transcriptional enhancer factor) Hs.546308 AA039369 7003 116352
TEAD1	NM_021961	hsa-miR-27a	TEAD1 TEA domain family member 1 (SV40 transcriptional enhancer factor) Hs.546308 AA039369 7003 116352
TEAD1	NM_021961	hsa-miR-92a	TEAD1 TEA domain family member 1 (SV40 transcriptional enhancer factor) Hs.546308 AA039369 7003 116352
TGFB1	NM_000358	hsa-miR-21	TGFB1 Transforming growth factor, beta-induced, 68kDa Hs.369397 AA633901 7045 113722
THRAP1	NM_005121	hsa-miR-369-3p	THRAP1 Thyroid hormone receptor associated protein 1 Hs.282678 AA459383 9969 103199
TIMELESS	NM_003920	hsa-miR-155	TIMELESS Timeless homolog (Drosophila) Hs.118631 W87751 8914 310253
TIMELESS	NM_003920	hsa-miR-29a	TIMELESS Timeless homolog (Drosophila) Hs.118631 W87751 8914 310253
TIMELESS	NM_003920	hsa-miR-29b	TIMELESS Timeless homolog (Drosophila) Hs.118631 W87751 8914 310253
TIMELESS	NM_003920	hsa-miR-29c	TIMELESS Timeless homolog (Drosophila) Hs.118631 W87751 8914 310253
TM4SF13	NM_014399	hsa-miR-155	TM4SF13 Tetraspanin 13 Hs.364544 W86201 27075 226071
TM4SF13	NM_014399	hsa-miR-369-3p	TM4SF13 Tetraspanin 13 Hs.364544 W86201 27075 226071
TM4SF2	NM_004615	hsa-let-7d	TM4SF2 Tetraspanin 7 Hs.441664 N93505 TALLA=T-cell acute lymphoblastic leukemia associated antigen 7102 114192
TM4SF2	NM_004615	hsa-let-7f	TM4SF2 Tetraspanin 7 Hs.441664 N93505 TALLA=T-cell acute lymphoblastic leukemia associated antigen 7102 114192
TM4SF9	NM_005723	hsa-miR-15a	TM4SF9 Tetraspanin 5 Hs.118118 H23278 10098 117428
TM4SF9	NM_005723	hsa-miR-15b	TM4SF9 Tetraspanin 5 Hs.118118 H23278 10098 117428
TM4SF9	NM_005723	hsa-miR-16	TM4SF9 Tetraspanin 5 Hs.118118 H23278 10098 117428
TM4SF9	NM_005723	hsa-miR-195	TM4SF9 Tetraspanin 5 Hs.118118 H23278 10098 117428
TMEFF1	NM_003692	hsa-miR-29a	TMEFF1 Transmembrane protein with EGF-like and two follistatin-like domains 1 Hs.336224 AA431475 8577 116729
TMEFF1	NM_003692	hsa-miR-29b	TMEFF1 Transmembrane protein with EGF-like and two follistatin-like domains 1 Hs.336224 AA431475 8577 116729
TMEFF1	NM_003692	hsa-miR-29c	TMEFF1 Transmembrane protein with EGF-like and two follistatin-like domains 1 Hs.336224 AA431475 8577 116729
TMEFF1	NM_003692	hsa-miR-30c	TMEFF1 Transmembrane protein with EGF-like and two follistatin-like domains 1 Hs.336224 AA431475 8577 116729
TMEFF1	NM_003692	hsa-miR-30e	TMEFF1 Transmembrane protein with EGF-like and two follistatin-like domains 1 Hs.336224 AA431475 8577 116729
TMEPAI	NM_199171	hsa-miR-124	TMEPAI Transmembrane, prostate androgen induced RNA Hs.517155 AA486591 56937 118377
TMEPAI	NM_199171	hsa-miR-19a	TMEPAI Transmembrane, prostate androgen induced RNA Hs.517155 AA486591 56937 118377
TMEPAI	NM_199171	hsa-miR-19b	TMEPAI Transmembrane, prostate androgen induced RNA Hs.517155 AA486591 56937 118377
TMEPAI	NM_199171	hsa-miR-92a	TMEPAI Transmembrane, prostate androgen induced RNA Hs.517155 AA486591 56937 118377
TMOD1	NM_003275	hsa-miR-10a	TMOD1 Tropomodulin 1 Hs.404289 AA410680 7111 112356
TMOD1	NM_003275	hsa-miR-10b	TMOD1 Tropomodulin 1 Hs.404289 AA410680 7111 112356
TMOD1	NM_003275	hsa-miR-124	TMOD1 Tropomodulin 1 Hs.404289 AA410680 7111 112356
TMOD1	NM_003275	hsa-miR-23a	TMOD1 Tropomodulin 1 Hs.404289 AA410680 7111 112356
TNPO1	NM_153188	hsa-miR-23a	TNPO1 Transportin 1 Hs.482497 AA287917 3842 221461
TP53INP1	NM_033285	hsa-let-7a	TP53INP1 Tumor protein p53 inducible nuclear protein 1 Hs.492261 AA459364 94241 223961
TP53INP1	NM_033285	hsa-miR-106a	TP53INP1 Tumor protein p53 inducible nuclear protein 1 Hs.492261 AA459364 94241 223961
TP53INP1	NM_033285	hsa-miR-124	TP53INP1 Tumor protein p53 inducible nuclear protein 1 Hs.492261 AA459364 94241 223961
TP53INP1	NM_033285	hsa-miR-142-5p	TP53INP1 Tumor protein p53 inducible nuclear protein 1 Hs.492261 AA459364 94241 223961
TP53INP1	NM_033285	hsa-miR-155	TP53INP1 Tumor protein p53 inducible nuclear protein 1 Hs.492261 AA459364 94241 223961
TP53INP1	NM_033285	hsa-miR-17	TP53INP1 Tumor protein p53 inducible nuclear protein 1 Hs.492261 AA459364 94241 223961
TP53INP1	NM_033285	hsa-miR-19a	TP53INP1 Tumor protein p53 inducible nuclear protein 1 Hs.492261 AA459364 94241 223961
TP53INP1	NM_033285	hsa-miR-19b	TP53INP1 Tumor protein p53 inducible nuclear protein 1 Hs.492261 AA459364 94241 223961
TP53INP1	NM_033285	hsa-miR-20a	TP53INP1 Tumor protein p53 inducible nuclear protein 1 Hs.492261 AA459364 94241 223961
TP53INP1	NM_033285	hsa-miR-23a	TP53INP1 Tumor protein p53 inducible nuclear protein 1 Hs.492261 AA459364 94241 223961
TP53INP1	NM_033285	hsa-miR-29a	TP53INP1 Tumor protein p53 inducible nuclear protein 1 Hs.492261 AA459364 94241 223961
TP53INP1	NM_033285	hsa-miR-29b	TP53INP1 Tumor protein p53 inducible nuclear protein 1 Hs.492261 AA459364 94241 223961
TP53INP1	NM_033285	hsa-miR-29c	TP53INP1 Tumor protein p53 inducible nuclear protein 1 Hs.492261 AA459364 94241 223961
TP53INP1	NM_033285	hsa-miR-30c	TP53INP1 Tumor protein p53 inducible nuclear protein 1 Hs.492261 AA459364 94241 223961
TP53INP1	NM_033285	hsa-miR-30e	TP53INP1 Tumor protein p53 inducible nuclear protein 1 Hs.492261 AA459364 94241 223961
TP53INP1	NM_033285	hsa-miR-369-3p	TP53INP1 Tumor protein p53 inducible nuclear protein 1 Hs.492261 AA459364 94241 223961
TRIB2	NM_021643	hsa-let-7a	TRIB2 Tribbles homolog 2 (Drosophila) Hs.467751 AA053865 28951 236762
TRIB2	NM_021643	hsa-let-7b	TRIB2 Tribbles homolog 2 (Drosophila) Hs.467751 AA053865 28951 236762
TRIB2	NM_021643	hsa-let-7c	TRIB2 Tribbles homolog 2 (Drosophila) Hs.467751 AA053865 28951 236762
TRIB2	NM_021643	hsa-let-7d	TRIB2 Tribbles homolog 2 (Drosophila) Hs.467751 AA053865 28951 236762

TRIB2	NM_021643	hsa-let-7e	TRIB2 Tribbles homolog 2 (Drosophila) Hs.467751 AA053865 28951 236762
TRIB2	NM_021643	hsa-let-7f	TRIB2 Tribbles homolog 2 (Drosophila) Hs.467751 AA053865 28951 236762
TRIB2	NM_021643	hsa-miR-18a	TRIB2 Tribbles homolog 2 (Drosophila) Hs.467751 AA053865 28951 236762
TRIB2	NM_021643	hsa-miR-29a	TRIB2 Tribbles homolog 2 (Drosophila) Hs.467751 AA053865 28951 236762
TRIB2	NM_021643	hsa-miR-29b	TRIB2 Tribbles homolog 2 (Drosophila) Hs.467751 AA053865 28951 236762
TRIB2	NM_021643	hsa-miR-29c	TRIB2 Tribbles homolog 2 (Drosophila) Hs.467751 AA053865 28951 236762
TRIB2	NM_021643	hsa-miR-98	TRIB2 Tribbles homolog 2 (Drosophila) Hs.467751 AA053865 28951 236762
TRPS1	NM_014112	hsa-miR-106a	TRPS1 Trichorhinophalangeal syndrome I Hs.253594 R54192 7227 120838
TRPS1	NM_014112	hsa-miR-124	TRPS1 Trichorhinophalangeal syndrome I Hs.253594 R54192 7227 120838
TRPS1	NM_014112	hsa-miR-155	TRPS1 Trichorhinophalangeal syndrome I Hs.253594 R54192 7227 120838
TRPS1	NM_014112	hsa-miR-17	TRPS1 Trichorhinophalangeal syndrome I Hs.253594 R54192 7227 120838
TRPS1	NM_014112	hsa-miR-19a	TRPS1 Trichorhinophalangeal syndrome I Hs.253594 R54192 7227 120838
TRPS1	NM_014112	hsa-miR-19b	TRPS1 Trichorhinophalangeal syndrome I Hs.253594 R54192 7227 120838
TRPS1	NM_014112	hsa-miR-20a	TRPS1 Trichorhinophalangeal syndrome I Hs.253594 R54192 7227 120838
TRPS1	NM_014112	hsa-miR-23a	TRPS1 Trichorhinophalangeal syndrome I Hs.253594 R54192 7227 120838
TRPS1	NM_014112	hsa-miR-30c	TRPS1 Trichorhinophalangeal syndrome I Hs.253594 R54192 7227 120838
TRPS1	NM_014112	hsa-miR-30e	TRPS1 Trichorhinophalangeal syndrome I Hs.253594 R54192 7227 120838
TRPS1	NM_014112	hsa-miR-369-3p	TRPS1 Trichorhinophalangeal syndrome I Hs.253594 R54192 7227 120838
TTN	NM_133379	hsa-let-7b	TTN Titin Hs.134602 W96443 titin 7273 178074
TTN	NM_133379	hsa-let-7c	TTN Titin Hs.134602 W96443 titin 7273 178074
TTN	NM_133379	hsa-let-7f	TTN Titin Hs.134602 W96443 titin 7273 178074
TTN	NM_133437	hsa-miR-106a	TTN Titin Hs.134602 W96443 titin 7273 178074
TTN	NM_133379	hsa-miR-124	TTN Titin Hs.134602 W96443 titin 7273 178074
TTN	NM_133379	hsa-miR-155	TTN Titin Hs.134602 W96443 titin 7273 178074
TTN	NM_133437	hsa-miR-17	TTN Titin Hs.134602 W96443 titin 7273 178074
TTN	NM_133437	hsa-miR-20a	TTN Titin Hs.134602 W96443 titin 7273 178074
TTN	NM_133437	hsa-miR-98	TTN Titin Hs.134602 W96443 titin 7273 178074
UBB	NM_018955	hsa-miR-155	UBB Ubiquitin B Hs.356190 AW078798 7314 330827
UBE2D3	NM_181893	hsa-miR-155	UBE2D3 **Ubiquitin-conjugating enzyme E2D 3 (UBC4/5 homolog, yeast) Hs.518773 N95752 7323 115593
UBE2D3	NM_181893	hsa-miR-21	UBE2D3 **Ubiquitin-conjugating enzyme E2D 3 (UBC4/5 homolog, yeast) Hs.518773 N95752 7323 115593
UBE2D3	NM_181893	hsa-miR-23a	UBE2D3 **Ubiquitin-conjugating enzyme E2D 3 (UBC4/5 homolog, yeast) Hs.518773 N95752 7323 115593
UBE2D3	NM_181893	hsa-miR-30c	UBE2D3 **Ubiquitin-conjugating enzyme E2D 3 (UBC4/5 homolog, yeast) Hs.518773 N95752 7323 115593
UBE2D3	NM_181893	hsa-miR-369-3p	UBE2D3 **Ubiquitin-conjugating enzyme E2D 3 (UBC4/5 homolog, yeast) Hs.518773 N95752 7323 115593
UBE2H	NM_182697	hsa-miR-369-3p	UBE2H Ubiquitin-conjugating enzyme E2H (UBC8 homolog, yeast) Hs.344165 AA520978 7328 99360
UBE2O	NM_022066	hsa-miR-23a	E2-230K Likely ortholog of mouse ubiquitin-conjugating enzyme E2-230K Hs.16130 AA677190 63893 221357
UBE2O	NM_022066	hsa-miR-30c	E2-230K Likely ortholog of mouse ubiquitin-conjugating enzyme E2-230K Hs.16130 AA677190 63893 221357
UBE2O	NM_022066	hsa-miR-30e	E2-230K Likely ortholog of mouse ubiquitin-conjugating enzyme E2-230K Hs.16130 AA677190 63893 221357
UGCG	NM_003358	hsa-miR-30c	UGCG UDP-glucose ceramide glucosyltransferase Hs.304249 W38595 Human mRNA for ceramide glucosyltransferase complete cds 7357 247621
UGCG	NM_003358	hsa-miR-30e	UGCG UDP-glucose ceramide glucosyltransferase Hs.304249 W38595 Human mRNA for ceramide glucosyltransferase complete cds 7357 247621
UGCG	NM_003358	hsa-miR-369-3p	UGCG UDP-glucose ceramide glucosyltransferase Hs.304249 W38595 Human mRNA for ceramide glucosyltransferase complete cds 7357 247621
UGT2B7	NM_001074	hsa-miR-19a	UGT2B7 UDP glycosyltransferase 2 family, polypeptide B7 Hs.10319 AA746229 7364 105739
UGT2B7	NM_001074	hsa-miR-19b	UGT2B7 UDP glycosyltransferase 2 family, polypeptide B7 Hs.10319 AA746229 7364 105739
VCAM1	NM_001078	hsa-miR-142-5p	VCAM1 Vascular cell adhesion molecule 1 Hs.109225 H07071 7412 108899
VCAM1	NM_001078	hsa-miR-98	VCAM1 Vascular cell adhesion molecule 1 Hs.109225 H07071 7412 108899
VCPIP1	NM_025054	hsa-let-7a	VCIP135 Valosin containing protein (p97)/p47 complex interacting protein 1 Hs.205550 AA405241 80124 310092
VCPIP1	NM_025054	hsa-let-7b	VCIP135 Valosin containing protein (p97)/p47 complex interacting protein 1 Hs.205550 AA405241 80124 310092
VCPIP1	NM_025054	hsa-let-7c	VCIP135 Valosin containing protein (p97)/p47 complex interacting protein 1 Hs.205550 AA405241 80124 310092
VCPIP1	NM_025054	hsa-let-7d	VCIP135 Valosin containing protein (p97)/p47 complex interacting protein 1 Hs.205550 AA405241 80124 310092
VCPIP1	NM_025054	hsa-let-7e	VCIP135 Valosin containing protein (p97)/p47 complex interacting protein 1 Hs.205550 AA405241 80124 310092
VCPIP1	NM_025054	hsa-let-7f	VCIP135 Valosin containing protein (p97)/p47 complex interacting protein 1 Hs.205550 AA405241 80124 310092
VCPIP1	NM_025054	hsa-miR-18a	VCIP135 Valosin containing protein (p97)/p47 complex interacting protein 1 Hs.205550 AA405241 80124 310092
VCPIP1	NM_025054	hsa-miR-19a	VCIP135 Valosin containing protein (p97)/p47 complex interacting protein 1 Hs.205550 AA405241 80124 310092
VCPIP1	NM_025054	hsa-miR-19b	VCIP135 Valosin containing protein (p97)/p47 complex interacting protein 1 Hs.205550 AA405241 80124 310092

VCPIP1	NM_025054	hsa-miR-98	VCIP135 Valosin containing protein (p97)/p47 complex interacting protein 1 Hs.205550 AA405241 80124 310092
WAC	NM_100486	hsa-miR-106a	WAC WW domain containing adaptor with coiled-coil Hs.435610 AA633545 51322 221971
WAC	NM_100486	hsa-miR-17	WAC WW domain containing adaptor with coiled-coil Hs.435610 AA633545 51322 221971
WAC	NM_100486	hsa-miR-20a	WAC WW domain containing adaptor with coiled-coil Hs.435610 AA633545 51322 221971
WAC	NM_100486	hsa-miR-23a	WAC WW domain containing adaptor with coiled-coil Hs.435610 AA633545 51322 221971
WDR26	NM_025160	hsa-miR-10a	WDR26 WD repeat domain 26 Hs.497873 AA776903 80232 220221
WDR26	NM_025160	hsa-miR-10b	WDR26 WD repeat domain 26 Hs.497873 AA776903 80232 220221
WDR26	NM_025160	hsa-miR-142-5p	WDR26 WD repeat domain 26 Hs.497873 AA776903 80232 220221
WDR26	NM_025160	hsa-miR-19a	WDR26 WD repeat domain 26 Hs.497873 AA776903 80232 220221
WDR26	NM_025160	hsa-miR-19b	WDR26 WD repeat domain 26 Hs.497873 AA776903 80232 220221
WDR26	NM_025160	hsa-miR-29a	WDR26 WD repeat domain 26 Hs.497873 AA776903 80232 220221
WDR26	NM_025160	hsa-miR-29b	WDR26 WD repeat domain 26 Hs.497873 AA776903 80232 220221
WDR26	NM_025160	hsa-miR-29c	WDR26 WD repeat domain 26 Hs.497873 AA776903 80232 220221
WDR26	NM_025160	hsa-miR-92a	WDR26 WD repeat domain 26 Hs.497873 AA776903 80232 220221
WDR33	NM_018383	hsa-miR-23a	WDR33 WD repeat domain 33 Hs.516376 N47269 55339 110657
WDR40A	NM_015400	hsa-miR-15a	WDR40A WD repeat domain 40A Hs.493750 R98204 25853 308780
WDR40A	NM_015400	hsa-miR-15b	WDR40A WD repeat domain 40A Hs.493750 R98204 25853 308780
WDR40A	NM_015400	hsa-miR-16	WDR40A WD repeat domain 40A Hs.493750 R98204 25853 308780
WDR40A	NM_015400	hsa-miR-195	WDR40A WD repeat domain 40A Hs.493750 R98204 25853 308780
WDR40A	NM_015397	hsa-miR-29a	WDR40A WD repeat domain 40A Hs.493750 R98204 25853 308780
WDR40A	NM_015397	hsa-miR-29b	WDR40A WD repeat domain 40A Hs.493750 R98204 25853 308780
WDR40A	NM_015397	hsa-miR-29c	WDR40A WD repeat domain 40A Hs.493750 R98204 25853 308780
XK	NM_021083	hsa-let-7a	XK Kell blood group precursor (McLeod phenotype) Hs.78919 H18932 7504 120746
XK	NM_021083	hsa-let-7b	XK Kell blood group precursor (McLeod phenotype) Hs.78919 H18932 7504 120746
XK	NM_021083	hsa-let-7c	XK Kell blood group precursor (McLeod phenotype) Hs.78919 H18932 7504 120746
XK	NM_021083	hsa-let-7e	XK Kell blood group precursor (McLeod phenotype) Hs.78919 H18932 7504 120746
XK	NM_021083	hsa-let-7f	XK Kell blood group precursor (McLeod phenotype) Hs.78919 H18932 7504 120746
XK	NM_021083	hsa-miR-98	XK Kell blood group precursor (McLeod phenotype) Hs.78919 H18932 7504 120746
ZBTB26	NM_020924	hsa-miR-23a	ZNF482 Zinc finger protein 482 Hs.3053 AA292994 10773 100678
ZC3HDC1	NM_022750	hsa-miR-16	ZC3HDC1 Zinc finger CCCH type domain containing 1 Hs.12646 H57857 64761 101005
ZC3HDC1	NM_022750	hsa-miR-195	ZC3HDC1 Zinc finger CCCH type domain containing 1 Hs.12646 H57857 64761 101005
ZDHHC2	NM_016353	hsa-miR-124	ZDHHC2 Zinc finger, DHHC domain containing 2 Hs.443852 R09490 51201 111967
ZDHHC2	NM_016353	hsa-miR-142-5p	ZDHHC2 Zinc finger, DHHC domain containing 2 Hs.443852 R09490 51201 111967
ZHX2	NM_014943	hsa-miR-106a	ZHX2 Zinc fingers and homeoboxes 2 Hs.377090 AA478596 22882 110436
ZHX2	NM_014943	hsa-miR-17	ZHX2 Zinc fingers and homeoboxes 2 Hs.377090 AA478596 22882 110436
ZHX2	NM_014943	hsa-miR-20a	ZHX2 Zinc fingers and homeoboxes 2 Hs.377090 AA478596 22882 110436
ZNF184	NM_007149	hsa-miR-19a	ZNF184 Zinc finger protein 184 (Kruppel-like) Hs.158174 AA455657 7738 108017
ZNF184	NM_007149	hsa-miR-30c	ZNF184 Zinc finger protein 184 (Kruppel-like) Hs.158174 AA455657 7738 108017
ZNF184	NM_007149	hsa-miR-30e	ZNF184 Zinc finger protein 184 (Kruppel-like) Hs.158174 AA455657 7738 108017
ZNF482	NM_006626	hsa-miR-10a	ZNF482 Zinc finger protein 482 Hs.3053 AA292994 10773 100678
ZNF482	NM_006626	hsa-miR-10b	ZNF482 Zinc finger protein 482 Hs.3053 AA292994 10773 100678
ZNF626	NM_145297	hsa-miR-142-5p	Transcribed locus, weakly similar to XP_209041.2 PREDICTED: similar to KIAA1503 protein [Homo sapiens] Hs.128692 AA972792 308907