

Figure S2

ama-MIR1

GC- | CAA
5' CUUCUAUAGUUUAGGUAACUU A
3' GAAGAUAUCAAUCCAUGAA C
AUA^ ACA

ama-MIR2

miR2.5 UGGUCUUCAACGAGGAAUCCU
miR2.4 UUGGUCUUCAACGAGGAAUUC
miR2.3 GUUGGUCUUCAACGAGGAAUU
-- miR2.1 - | miR2.2 A AA G
5' GUGAUGGGGAUAGAUAUUGC AUUGUUGGUCUUCAACGAGGAU UCCU GU GC C
3' CCACUACCCCUAUCUAGUAACG UACAACCAGAAGUUGCUCUUA AGGA UA UG G
AG U^ miR2.2* C C- A
miR2.3* CAACCAGAAGUUGCUCUUA
miR2.4* AACCAGAAGUUGCUCUUAAG

ama-MIR3

miR3.4* GCCUGCCCAUGGAUUCAGCAGC
miR3.3* UCUUGCCUGCCCAUGGAUUCAG
-- miR3.1* UAGU-- C-- | GAAA C G-- AAAA A
5' GUUGUCUCUUGCCUGCCAU GGAUUCAGCAGC AAG GAC UG UUCGGGGA UUG UA G
3' CAACAGAGAACGGACGGGUA CCUAAGUCGUCGU UUC CUG AU AAGCCCUU GGC AU C
UC miR3.1 CAAACU CAA^ G--- - AGA CUAG G
miR3.2 AGAGAACGGACGGGUACCU
miR3.3 AGAACGGACGGGUACCUAAGUC
miR3.4 CGGACGGGUACCUAAGUCGUCG
miR3.5 GACGGGUACCUAAGUCGUCGU

ama-MIR4

UUA UC CAAUCACCAACCC AA | CCAAAAGUUU
5' AGUUUGUGCGUGAAUCUAAC AGAUU AGAUUC UCA \
3' UCAAUACGCACUUAAGAUUG UCUAG UCUGAG AGU U
UGC U- ACCC----- AC^ UUCUGUACC

ama-MIR5

A C UUU---- C U----- CAUAA----- AUAU- C---- C- - | U
5' GGUG CUUUUUUUUUUUUUUUUU UAU UGC UGAAA AGGAAAA UCCAUAU UG ACU GGU A
3' UUAC GAAAAGAAAGAGAAAAGA AUA AUG ACUUU UCCUUUU AGGUAA AC UGA CCA U
U A CACCACU C UUUAGU UUUCCA AUUCCUG GUGCU UUUUU UU U^ U

ama-MIR6

UC | U - U
5' GGGGUAUUA GA CUUUUUAUUC U
3' CUCCGUAGU CU GAAGGAUAAG U
UU^ - A U

ama-MIR7

AAG - | C GA- UA AUG
5' UGUUGAAUC AUGG AUGCCG GGC CU \
3' GCAACUAG UGCC UACGGC UCG GA C
CGA C^ - AAC UA GGA

G- | U GUUUAACAGC
 5' CGGGAAGAGUUAUC UUUCU \
 3' GCCCUUCUCAAUAG AAAGG C
 UA^ U UCCCACCCGC

UUU - GU A AUAAGUGGGA-- | G
 5' GUGAAAUGAC UUGAGAG GU GG GCC A
 3' CAUUUUUAUUG AAUUUUC CA CC CGG A
 AUU C AU - AUAAGUGAAAAG^ A

C--	C	CUCUC----	CU	CCGCCGUUCU	G-	-	A
5' UCUUCUCUCUCUCUCU	UCUCUCUCU	UCUCUCU	CUCUCUCUCU	AGUC	GC	GAGAGG	A
3' AGAGAGAGAGAGAGA	AGAGAGAGA	AGAGAGG	GAGAGAGAGA	UCAG	CG	CUCUUC	A
GAA	U	AAAAAUUAU	UU	C-----	ACUG	A^	G

GUU| U GU
 5' GCAUUAUUAUUUAUUUAUUG U U
 3' CGUAGUAGUAGUAGUAGU G G
 ACU^ U UU

GGA | A U
 5' UCGCGGCG CGUGGGCGG U
 3' AGCGCUGC GCGCCGUC C
 AAG ^ A G

miR13.2 CUCUCUCUCUCUCUCUCUCA
miR13.1 A -AUUCCGUAAA| C UUUUAUUUUAUCUCU
5' AG CA UCUCUCUCUCUCUCUCUCUCUC AGAGC CA U
3' UUGU AGAGGGGGAGAGAGAGAGAG UCUCG GU C
C A \ ----- ^ U UUAACUUAACUUU
(104 nt side loop)

CUU GAG ----- | A U
5' UUGAUUA UAGGGGUCGCGA GAGA GCAG GCG \
3' AACUGAU GUUCUCAGCGCU CUCU CGUC CGC A
AAC ACU UAUUUU GAGCACUGAUAC^ C C

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--                               CCG--- AA          .-G|          UCCGG      U  GUG--      C
5' GAGGUUGAAUAUUAAGUAUGAAUCAUAGUUCGA          CG  AUUUUAUUU      UUUCUGUAUU      AAUA GA          UGUGA U
3' CUCCAAACUUAUAAUUCAUACUUAAGUAUCAAGCU          GC  UAGGUAAA      AAAGACAUGA      UUAU CU          ACAUU U
CG                               UAUGAA  AC          \  -^          UAG--      C  AGUUA      G

                               (72 nt side loop)

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