Supplemental Table C. miRNAs Found in Exons of Non-Coding TUs

microRNA	miR Copy Number (identical)	miR Chr Location	Host Gene/TU	Host Gene/TU Details	microRNA Mapping Position Within Host Gene/TU	Host Gene/TU Expression	Expression Source/Ref
hsa-let-7b	1	22	OTTHUMG0000030111/ 22_44720862/speytey AceView OTTHUMG00000030111/	Vega mlncRNA (dJ695O20B.C22.9)	Within Exon 1	Well expressed	Aceview/Acembly
			22_44720862/speytey .				
hsa-let-7c-2	2	22	AceView	Vega mlncRNA (dJ695O20B.C22.9)		Well expressed	Aceview/Acembly
mmu-let-7b	1	15			undefined		
mmu-let-7c-2	2	15			undefined		
hsa-let-7i	1	12			undefined		
mmu-let-7i	1	10	D630033A02 RIKEN	FANTOM2 mlncRNA	Within Exon 1		
			ENSESTG00000028158/		Within Exon 2 (evidence for	Everageed at year, high	
hsa-miR-22	1	17	17_1826145/snoree AceView	Possible mlncRNA	alternative splicing)	Expressed at very high levels	AceView/Acembly
							, , , , , , , , , , , , , , , , , , , ,
					Within Exon 2 (evidence for	Highest expression seen	RIKEN Expression
mmu-miR-22	1	11	2210403K04 RIKEN	FANTOM2 mlncRNA	alternative splicing)	in neonate skin and lung	•
					, 37	5	,
hsa-miR-29b-			1 205058566/glawnabu				
2	2	1	AceView	Possible mlncRNA	Within Exon 1		
				MEMBRANE COFACTOR PROTEIN PRECURSOR (CD46 ANTIGEN) (TROPHOBLAST LEUCOCYTE	miRNA precursor maps close to this	Expressed in almost all cells except erythrocytes	
hsa-miR-29b-	2	1	ENSG00000117335	COMMON ANTIGEN) (TLX) (SWISS PROT - MCP)	gene on coding opposite strand	and some bone marrow cells	SWISSPROT
mmu-miR-				,	, p 10.01 00.01.0		
29b-2	2	1	C030002C11 RIKEN	Possible mlncRNA	Within Exon 1		
hsa-miR-34b	1	11	ENSESTG00000013662	Possible mlncRNA	Within Exon 2		
hsa-miR-34c	1	11	ENSESTG00000013662	Possible mlncRNA	Within Exon 2		

mmu-miR- 34b	1	9			undefined		
mmu-miR-							
34c	1	9			undefined		
hsa-miR-99b	2	19	19_56888521/blorbla AceView	Possible mlncRNA	Within Exon 1 on opposite strand Within Exon 1 on	Expressed at high levels	Aceview/Acembly
hsa-let-7e	1	19	19_56888521/blorbla AceView	Possible mlncRNA	opposite strand Within Exon 1 on	Expressed at high levels	Aceview/Acembly
hsa-miR-125a	1	19	19_56888521/blorbla AceView	Possible mlncRNA	opposite strand	Expressed at high levels	Aceview/Acembly
mmu-miR- 99b mmu-let-7e mmu-miR-	2 1	17 17			undefined undefined		
125a	1	17			undefined		
hsa-miR-101-	2	1			undefined		
mmu-miR-							
101	2	4	ENSMUSESTG00000022081	Possible mlncRNA	Within Exon 2		
hsa-miR-122a	1	18	18_54266208/slawso AceView	Possible mlncRNA	Within Exon 2	Well Expressed	AceView/Acembly
hsa-miR-122a mmu-miR- 122a	1	18	18_54266208/slawso AceView	Possible mlncRNA	Within Exon 2 undefined (note that mouse Build 32 is incorrectly assembled in the contig containing this miRNA)		AceView/Acembly
mmu-miR-			18_54266208/slawso AceView	Possible mlncRNA	undefined (note that mouse Build 32 is incorrectly assembled in the contig containing this		AceView/Acembly
mmu-miR-			18_54266208/slawso AceView	Possible mlncRNA Retrotransposon-like gene 1 (Rtl1)	undefined (note that mouse Build 32 is incorrectly assembled in the contig containing this miRNA) Within Exon 1 on opposite strand		AceView/Acembly
mmu-miR- 122a	1	18		Retrotransposon-like gene 1 (Rtl1) Retrotransposon-like gene 1 (Rtl1)	undefined (note that mouse Build 32 is incorrectly assembled in the contig containing this miRNA) Within Exon 1 on opposite strand Within Exon 1 on opposite strand		AceView/Acembly
mmu-miR- 122a hsa-miR-127 hsa-miR-136 mmu-miR- 127	1	18	Rtil NCBI	Retrotransposon-like gene 1 (Rtl1) Retrotransposon-like gene 1 (Rtl1) Retrotransposon-like gene 1 (Rtl1) (BK001261)	undefined (note that mouse Build 32 is incorrectly assembled in the contig containing this miRNA) Within Exon 1 on opposite strand		AceView/Acembly
mmu-miR- 122a hsa-miR-127 hsa-miR-136 mmu-miR-	1 1 1	18 14 14	Rtl1 NCBI Rtl1 NCBI	Retrotransposon-like gene 1 (Rtl1) Retrotransposon-like gene 1 (Rtl1) Retrotransposon-like gene 1 (Rtl1)	undefined (note that mouse Build 32 is incorrectly assembled in the contig containing this miRNA) Within Exon 1 on opposite strand		AceView/Acembly
mmu-miR- 122a hsa-miR-127 hsa-miR-136 mmu-miR- 127 mmu-miR-	1 1 1 1	18 14 14 12	Rtl1 NCBI Rtl1 NCBI Rtl1 NCBI	Retrotransposon-like gene 1 (Rtl1) Retrotransposon-like gene 1 (Rtl1) Retrotransposon-like gene 1 (Rtl1) (BK001261) Retrotransposon-like gene 1 (Rtl1)	undefined (note that mouse Build 32 is incorrectly assembled in the contig containing this miRNA) Within Exon 1 on opposite strand Within Exon 1 on		AceView/Acembly

					Within Intron 2 on		
hsa-miR-133b mmu-miR-	1	6	OTTHUMG00000014842	Vega mlncRNA	opposite strand		
133b	1	1			undefined		
hsa-miR-137	1	1	1_97978084/toyda Acembly	Possible mlncRNA	Within Exon 2	Moderately expressed	Aceview/Acembly
mmu-miR- 137	1	1	ENSMUSESTG00000012545	Possible mlncRNA	Within Exon 3		
hsa-miR-143	1	5	5_148836826/foygor Aceview	Possible mlncRNA	Within Exon 1	Moderately expressed	Aceview/Acembly
hsa-miR-145	1	5	5_148836826/foygor Aceview	Possible mlncRNA	Within Exon 1	Moderately expressed	Aceview/Acembly
mmu-miR- 143	1	18			undefined		
mmu-miR- 145	1	18			undefined		
hsa-miR-146	1	5	ENSESTG00000012459	Possible mlncRNA	Within Exon 2		
mmu-miR- 146	1	11	5830402E17 RIKEN	FANTOM2 mlncRNA	Within Exon 1		
					Within Exon 5 on		
hsa-miR-150				40S RIBOSOMAL PROTEIN S11	opposite strand (note miR maps within	e	
predicted mmu-miR-	1	19	19_54691494 AceView	(Swiss Prot: RPS11)	3'UTR)		
150	1	7			undefined		
							Tam W. Identification
							and characterization of human BIC, a gene
							on chromosome 21
			ENSESTG00000018742;				that encodes a noncoding RNA.
hsa-miR-155 predicted		21	21_25856092 Aceview; BIC NCBI	PIC mincDNA (AE402776)	Within Exon 3	Highest expression in	Gene. 2001 Aug
mmu-miR-	1	21	NCDI	BIC mlncRNA (AF402776)	WILTIIII EXOII 3	spleen, thymus	22;274(1-2):157-67
155	1	16	Bic NCBI	Bic mlncRNA (AY096003)	Within Exon 2		

			11_64434198/snymuby				
hsa-miR-192	1	11	AceView	Possible mlncRNA	Within Exon 1	Moderately expressed	Aceview/Acembly
mmu-miR- 192	1	19			undefined		
132	-	13			undenned		
hsa-miR-195		47	LOC284112/peeror NCBI				
predicted mmu-miR-	1	17	Aceview	Possible mlncRNA	Within Exon 1	Well expressed	Aceview/Acembly
195	1	11			undefined		
hsa-miR-196-							
4 predicted	4	7	ENSESTG00000015802	Possible mlncRNA	Within Exon 2		
mmu-miR- 196-4							
predicted	4	6			undefined		
hsa-miR-199a	2		1 160252004/bire: 4 View	Dansible valueDNA	Within From 4	Commercial at high lawyle	0) (; / 0
2 hsa-miR-	3	1	1_169353084/timi AceView	Possible mlncRNA	Within Exon 1	Expressed at high levels	
199a*-2	2	1	1_169353084/timi AceView	Possible mlncRNA	Within Exon 1	Expressed at high levels	AceView/Acembly
hsa-miR-214	1	1	1_169353084/timi AceView	Possible mlncRNA	Within Exon 2	Expressed at high levels	AceView/Acembly
							A novel dynamin III isoform is up-regulated in the central nervous system in hypothyroidism. dynamin III transcript is brain-specific and selectively regulated
hsa-miR-199a 2	3	1	ENSG00000053059	DYNAMIN 3 (EC 3.6.1.50) (DYNAMIN, TESTICULAR) (T- DYNAMIN). DYN3_HUMAN [Source:SWISSPROT;Acc:Q9UQ16]	Within Intron 14 on opposite strand	Highly expressed in testes. Also expressed in brain and lung.	by thyroid hormone status
hsa-miR- 199a*-2	2	1	ENSG00000053059	DYNAMIN 3 (EC 3.6.1.50) (DYNAMIN, TESTICULAR) (T- DYNAMIN). DYN3_HUMAN [Source:SWISSPROT;Acc:Q9UQ16]	Within Intron 14 on opposite strand	same	same

hsa-miR-214	1	1	ENSG00000053059	DYNAMIN 3 (EC 3.6.1.50) (DYNAMIN, TESTICULAR) (T- DYNAMIN). DYN3_HUMAN [Source:SWISSPROT;Acc:Q9UQ16]	Within Intron 14 opposite strand	same	same
mmu-miR- 199a-2	3	1	ENSMUSG00000040265	DYNAMIN 3 (EC 3.6.1.50) (DYNAMIN, TESTICULAR) (T- DYNAMIN)	Within Intron 3 on opposite strand	Highly expressed in testes. Also expressed in brain and lung.	SWISS PROT
mmu-miR- 199a*-2	2	1	ENSMUSG00000040265	DYNAMIN 3 (EC 3.6.1.50) (DYNAMIN, TESTICULAR) (T- DYNAMIN)	Within Intron 3 on opposite strand	Highly expressed in testes. Also expressed in brain and lung.	SWISS PROT
mmu-miR- 214	1	1	D230012004 RIKEN	Possible mlncRNA	Within Exon 1		
mmu-miR- 214	1	1	ENSMUSG00000040265	DYNAMIN 3 (EC 3.6.1.50) (DYNAMIN, TESTICULAR) (T- DYNAMIN)	Within Intron 3 on opposite strand	Highly expressed in testes. Also expressed in brain and lung.	SWISS PROT
hsa-miR-202 doe	es not app	ear conserve	ed				
mmu-miR- 202 mmu-miR-	1	7	G630041I22 RIKEN	FANTOM mlncRNA	Within Exon 2 Within Intron 1 on		
202	1	7	ENSMUSESTG00000019381	Possible mlncRNA	opposite strand		
hsa-miR-206	1	6			undefined		
mmu-miR- 206	1	1	4831414J02/D930049J19 RIKEN	FANTOM2 mlncRNA (Mouse homolog of Rat Synapse-Specific 7H4 RNA) (BB613730)	Within Exon 1	Rat Synapse-Specific 7H4 expression is restricted to skeletal muscle. In cultured cells, expressed in differentiated muscle cells and not expressed in Schwann cells, fibroblasts, undifferentiated muscle cells	Velleca MA, Wallace MC, Merlie JP. A novel synapse-
hsa-miR-215	1	1	ENSG00000067704	PROBABLE ISOLEUCYL TRNA SYNTHETASE (Swiss Prot: NM_018060)	Within Intron 12 on opposite strand		
mmu-miR- 215	1	1	2010103J01 RIKEN	FANTOM2 mlncRNA	Within Exon 2	Highest expression in neonate head	RIKEN Expression Array Database

mmu-miR- 215	1	1	ENSMUSG00000026618	SIMILAR TO MITOCHONDRIAL ISOLEUCINE TRNA SYNTHETASE. [Source: SPTREMBL (Q8BIP3)]	Within Intron 11 on opposite strand		
hsa-miR-221	1	X	X_44651888/rytor AceView	Possible mlncRNA	Within Exon 1 Within Exon 2 on	Expressed at high levels	AceView/Acembly
hsa-miR-221	1	X	ENSESTG00000020404	Possible mlncRNA	opposite strand		
mmu-miR- 221	1	Х			undefined		
hsa-miR-223	1	X	X_64102086/rertoy AceView	Possible mlncRNA	Within Exon 3	Expressed at high levels	AceView/Acembly
mmu-miR- 223	1	X	9830169G21 RIKEN	Possible mlncRNA	Within Exon 2		
hsa-miR-296	1	20	20_58078195/flartu AceView	Possible mlncRNA	Within Exon 1	Expressed at moderate levels	AceView/Acembly
mmu-miR- 296 mmu-miR-	1	2	D330038P10 RIKEN	FANTOM2 mlncRNA	Within Exon 1 Within Intron 1 on		
296	1	2	ENSMUSESTG00000021919	Possible mlncRNA	opposite strand		
hsa-miR-324- 5p hsa-miR-324-	1	17	17_7329130/tuwer AceView	Possible mlncRNA	Within Exon 3	Well Expressed	AceView/Acembly
3p	1	17	17_7329130/tuwer AceView	Possible mlncRNA	Within Exon 3	Well Expressed	AceView/Acembly
hsa-miR-324- 5p	1	17	ENSG00000072778	ACYL-COA DEHYDROGENASE, VERY LONG-CHAIN SPECIFIC, MITOCHONDRIAL PRECURSOR (EC 1.3.99) (VLCAD) (MVLCAD). [Source:SWISSPROT;Acc:P50544]	Within Intron 11 on opposite strand		
hsa-miR-324- 3p	1	17	ENSG00000072778	ACYL-COA DEHYDROGENASE, VERY LONG-CHAIN SPECIFIC, MITOCHONDRIAL PRECURSOR (EC 1.3.99) (VLCAD) (MVLCAD). [Source:SWISSPROT;Acc:P50544]	Within Intron 11 on opposite strand		

mmu-miR- 324-5p mmu-miR- 324-3p	1	11	ENSMUSG00000018574 ENSMUSG00000018574	ACYL-COA DEHYDROGENASE, VERY LONG-CHAIN SPECIFIC, MITOCHONDRIAL PRECURSOR (EC 1.3.99) (VLCAD) (MVLCAD). [Source:SWISSPROT;Acc:P50544] ACYL-COA DEHYDROGENASE, VERY LONG-CHAIN SPECIFIC, MITOCHONDRIAL PRECURSOR (EC 1.3.99) (VLCAD) (MVLCAD). [Source:SWISSPROT;Acc:P50544]	Within Intron 11 on opposite strand		
hsa-miR-331	1	12			undefined		
mmu-miR- 331	1	10	A630071D13 RIKEN	FANTOM2 mlncRNA	Within Exon 1	Expression seen in bone marrow, amygdala, eye	Affy U74A: 106501_at; 105366_at