

PicTar WEB INTERFACE

Choose Species:	vertebrate ▼
Choose Dataset:	target predictions for all human microRNAs based on conservation in mammals (human, chimp, mouse, rat, dog)
microRNA ID: Click above for all microRNAs linked to RFAM	hsa-miR-125a ▼
Gene ID: Click above for all RefSeq Id's linked to NCBI (Warning: may take ~20 secs)	<input type="text"/> vertebrates: use RefSeq identifiers, e.g. NM_003483 or Gene symbols (for example HK2).
Tissue:	▼

PicTar predictions

Rank Click here for detailed 3'utr alignments and location of predicted site	human Refseq Id	All miRNAs predicted to target the gene	PicTar score	microRNAs with Anchor sites	predicted sites for all microRNAs embedded in the UCSC genome browser	annotation
1	NM_006465	All miRNA predictions	15.15	hsa-miR-125a	Genome browser	Homo sapiens AT rich interactive domain 3B (BRIGHT- like) (ARID3B), mRNA.
2	NM_145185	All miRNA predictions	11.10	hsa-miR-125a	Genome browser	Homo sapiens mitogen-activated protein kinase kinase 7 (MAP2K7), mRNA.
3	NM_031232	All miRNA predictions	10.74	hsa-miR-125a	Genome browser	Homo sapiens amyloid beta (A4) precursor protein-binding, family A, member 2 binding protein (APBA2BP), transcript variant 2, mRNA.
4	NM_031231	All miRNA predictions	10.74	hsa-miR-125a	Genome browser	Homo sapiens amyloid beta (A4) precursor protein-binding, family A, member 2 binding protein (APBA2BP), transcript variant 1, mRNA.
5	NM_000449	All miRNA predictions	10.29	hsa-miR-125a	Genome browser	Homo sapiens regulatory factor X, 5 (influences HLA class II expression) (RFX5), mRNA.
6	NM_015094	All miRNA predictions	10.14	hsa-miR-125a	Genome browser	Homo sapiens hypermethylated in cancer 2 (HIC2), mRNA.
7	NM_152283	All miRNA predictions	10.11	hsa-miR-125a	Genome browser	Homo sapiens zinc finger protein 62 homolog (mouse) (ZFP62), mRNA.
8	NM_015898	All miRNA predictions	9.80	hsa-miR-125a	Genome browser	Homo sapiens zinc finger and BTB domain containing 7 (ZBTB7), mRNA.
9	NM_002537	All miRNA predictions	9.11	hsa-miR-125a	Genome browser	Homo sapiens ornithine decarboxylase antizyme 2 (OAZ2), mRNA.
10	NM_178007	All miRNA predictions	8.92	hsa-miR-125a	Genome browser	Homo sapiens START domain containing 13 (STARD13), transcript variant beta, mRNA.
11	NM_178006	All miRNA predictions	8.92	hsa-miR-125a	Genome browser	Homo sapiens START domain containing 13 (STARD13), transcript variant alpha, mRNA.
12	NM_052851	All miRNA predictions	8.92	hsa-miR-125a	Genome browser	Homo sapiens START domain containing 13 (STARD13), transcript variant gamma, mRNA.
13	NM_033503	All miRNA predictions	8.91	hsa-miR-125a	Genome browser	Homo sapiens Bcl2 modifying factor (BMF), transcript variant 2, mRNA.
14	NM_001003940	All miRNA predictions	8.91	hsa-miR-125a	Genome browser	Homo sapiens Bcl2 modifying factor (BMF), transcript variant 1, mRNA.
15	NM_001003942	All miRNA predictions	8.91	hsa-miR-125a	Genome browser	Homo sapiens Bcl2 modifying factor (BMF), transcript variant 3, mRNA.
16	NM_001003943	All miRNA predictions	8.90	hsa-miR-125a	Genome browser	Homo sapiens Bcl2 modifying factor (BMF), transcript variant 4, mRNA.
17	NM_004947	All miRNA predictions	8.67	hsa-miR-125a	Genome browser	Homo sapiens dedicator of cytokinesis 3 (DOCK3), mRNA.

18	NM_033387	All miRNA predictions	8.27	hsa-miR-125a	Genome browser	Homo sapiens chromosome 9 open reading frame 59 (C9orf59), mRNA.
19	NM_014674	All miRNA predictions	7.73	hsa-miR-125a	Genome browser	Homo sapiens ER degradation enhancer, mannosidase alpha-like 1 (EDEM1), mRNA.
20	NM_022484	All miRNA predictions	7.72	hsa-miR-125a	Genome browser	Homo sapiens hypothetical protein FLJ13576 (FLJ13576), mRNA.
21	NM_138769	All miRNA predictions	7.69	hsa-miR-125a	Genome browser	Homo sapiens ras homolog gene family, member T2 (RHOT2), mRNA.
22	NM_004283	All miRNA predictions	7.63	hsa-miR-125a	Genome browser	Homo sapiens RAB3D, member RAS oncogene family (RAB3D), mRNA.
23	NM_022465	All miRNA predictions	7.43	hsa-miR-125a	Genome browser	Homo sapiens zinc finger protein, subfamily 1A, 4 (Eos) (ZNFN1A4), mRNA.
24	NM_052830	All miRNA predictions	7.35	hsa-miR-125a	Genome browser	Homo sapiens gamma-glutamyltransferase-like 3 (GGTL3), transcript variant 1, mRNA.
25	NM_012429	All miRNA predictions	7.28	hsa-miR-125a	Genome browser	Homo sapiens SEC14-like 2 (S. cerevisiae) (SEC14L2), mRNA.
26	NM_032560	All miRNA predictions	7.23	hsa-miR-125a	Genome browser	Homo sapiens KIAA2010 (KIAA2010), transcript variant 1, mRNA.
27	NM_007345	All miRNA predictions	7.23	hsa-miR-125a	Genome browser	Homo sapiens zinc finger protein 236 (ZNF236), mRNA.
28	NM_004869	All miRNA predictions	7.17	hsa-miR-125a	Genome browser	Homo sapiens vacuolar protein sorting 4B (yeast) (VPS4B), mRNA.
29	NM_003077	All miRNA predictions	7.08	hsa-miR-125a	Genome browser	Homo sapiens SWI/SNF related, matrix associated, actin dependent regulator of chromatin, subfamily d, member 2 (SMARCD2), mRNA.
30	NM_002442	All miRNA predictions	7.08	hsa-miR-125a	Genome browser	Homo sapiens musashi homolog 1 (Drosophila) (MSI1), mRNA.
31	NM_014521	All miRNA predictions	7.00	hsa-miR-125a	Genome browser	Homo sapiens SH3-domain binding protein 4 (SH3BP4), mRNA.
32	NM_173217	All miRNA predictions	6.83	hsa-miR-125a	Genome browser	Homo sapiens sialyltransferase 1 (beta-galactoside alpha-2,6-sialyltransferase) (SIAT1), transcript variant 3, mRNA.
33	NM_173216	All miRNA predictions	6.83	hsa-miR-125a	Genome browser	Homo sapiens sialyltransferase 1 (beta-galactoside alpha-2,6-sialyltransferase) (SIAT1), transcript variant 1, mRNA.
34	NM_003032	All miRNA predictions	6.83	hsa-miR-125a	Genome browser	Homo sapiens sialyltransferase 1 (beta-galactoside alpha-2,6-sialyltransferase) (SIAT1), transcript variant 2, mRNA.
35	NM_005940	All miRNA predictions	6.78	hsa-miR-125a	Genome browser	Homo sapiens matrix metalloproteinase 11 (stromelysin 3) (MMP11), mRNA.
36	NM_020778	All miRNA predictions	6.66	hsa-miR-125a	Genome browser	Homo sapiens likely ortholog of mouse myocytic induction/differentiation originator (MIDORI), mRNA.
37	NM_005688	All miRNA predictions	6.63	hsa-miR-125a	Genome browser	Homo sapiens ATP-binding cassette, sub-family C (CFTR/MRP), member 5 (ABCC5), mRNA.
38	NM_005876	All miRNA predictions	6.49	hsa-miR-125a	Genome browser	Homo sapiens aortic preferentially expressed protein 1 (APEG1), mRNA.
39	NM_005397	All miRNA predictions	6.46	hsa-miR-125a	Genome browser	Homo sapiens podocalyxin-like (PODXL), mRNA.
40	NM_013368	All miRNA predictions	6.44	hsa-miR-125a	Genome browser	Homo sapiens SERTA domain containing 3 (SERTAD3), transcript variant 1, mRNA.
41	NM_203344	All miRNA predictions	6.44	hsa-miR-125a	Genome browser	Homo sapiens SERTA domain containing 3 (SERTAD3), transcript variant 2, mRNA.
42	NM_016097	All miRNA predictions	6.42	hsa-miR-125a	Genome browser	Homo sapiens immediate early response 3 interacting protein 1 (IER3IP1), mRNA.
43	NM_178026	All miRNA predictions	6.40	hsa-miR-125a	Genome browser	Homo sapiens gamma-glutamyltransferase-like 3 (GGTL3), transcript variant 3, mRNA.
44	NM_172217	All miRNA predictions	6.40	hsa-miR-125a	Genome browser	Homo sapiens interleukin 16 (lymphocyte chemoattractant factor) (IL16), transcript variant 2, mRNA.
45	NM_175709	All miRNA predictions	6.35	hsa-miR-125a	Genome browser	Homo sapiens chromobox homolog 7 (CBX7), mRNA.
46	NM_020770	All miRNA predictions	6.33	hsa-miR-125a	Genome browser	Homo sapiens cingulin (CGN), mRNA.
47	NM_000212	All miRNA predictions	6.25	hsa-miR-125a	Genome browser	Homo sapiens integrin, beta 3 (platelet glycoprotein IIIa, antigen CD61) (ITGB3), mRNA.
48	NM_016577	All miRNA	6.25	hsa-miR-	Genome	Homo sapiens RAB6B, member RAS oncogene family

		predictions		125a	browser	(RAB6B), mRNA.
49	NM_004656	All miRNA predictions	6.25	hsa-miR-125a	Genome browser	Homo sapiens BRCA1 associated protein-1 (ubiquitin carboxy-terminal hydrolase) (BAP1), mRNA.
50	NM_003326	All miRNA predictions	6.05	hsa-miR-125a	Genome browser	Homo sapiens tumor necrosis factor (ligand) superfamily, member 4 (tax-transcriptionally activated glycoprotein 1, 34kDa) (TNFSF4), mRNA.
51	NM_032242	All miRNA predictions	5.67	hsa-miR-125a	Genome browser	Homo sapiens plexin A1 (PLXNA1), mRNA.
52	NM_020777	All miRNA predictions	5.54	hsa-miR-125a	Genome browser	Homo sapiens sortilin-related VPS10 domain containing receptor 2 (SORCS2), mRNA.
53	NM_178126	All miRNA predictions	5.49	hsa-miR-125a	Genome browser	Homo sapiens hypothetical protein LOC162427 (LOC162427), mRNA.
54	NM_004423	All miRNA predictions	5.46	hsa-miR-125a	Genome browser	Homo sapiens dishevelled, dsh homolog 3 (Drosophila) (DVL3), mRNA.
55	NM_015995	All miRNA predictions	5.44	hsa-miR-125a	Genome browser	Homo sapiens Kruppel-like factor 13 (KLF13), mRNA.
56	NM_001150	All miRNA predictions	5.42	hsa-miR-125a	Genome browser	Homo sapiens alanyl (membrane) aminopeptidase (aminopeptidase N, aminopeptidase M, microsomal aminopeptidase, CD13, p150) (ANPEP), mRNA.
57	NM_012069	All miRNA predictions	5.42	hsa-miR-125a	Genome browser	Homo sapiens ATPase, (Na ⁺)/K ⁺ transporting, beta 4 polypeptide (ATP1B4), mRNA.
58	NM_014369	All miRNA predictions	5.39	hsa-miR-125a	Genome browser	Homo sapiens protein tyrosine phosphatase, non-receptor type 18 (brain-derived) (PTPN18), mRNA.
59	NM_007243	All miRNA predictions	5.35	hsa-miR-125a	Genome browser	Homo sapiens nurim (nuclear envelope membrane protein) (NRM), mRNA.
60	NM_006133	All miRNA predictions	5.34	hsa-miR-125a	Genome browser	Homo sapiens chromosome 11 open reading frame 11 (C11orf11), mRNA.
61	NM_004451	All miRNA predictions	5.18	hsa-miR-125a	Genome browser	Homo sapiens estrogen-related receptor alpha (ESRRA), mRNA.
62	NM_003901	All miRNA predictions	5.11	hsa-miR-125a	Genome browser	Homo sapiens sphingosine-1-phosphate lyase 1 (SGPL1), mRNA.
63	NM_005637	All miRNA predictions	5.10	hsa-miR-125a	Genome browser	Homo sapiens synovial sarcoma translocation, chromosome 18 (SS18), transcript variant 2, mRNA.
64	NM_022918	All miRNA predictions	5.08	hsa-miR-125a	Genome browser	Homo sapiens hypothetical protein FLJ22104 (FLJ22104), mRNA.
65	NM_080621	All miRNA predictions	5.08	hsa-miR-125a	Genome browser	Homo sapiens sterile alpha motif domain containing 10 (SAMD10), mRNA.
66	NM_004470	All miRNA predictions	5.00	hsa-miR-125a	Genome browser	Homo sapiens FK506 binding protein 2, 13kDa (FKBP2), transcript variant 1, mRNA.
67	NM_057092	All miRNA predictions	5.00	hsa-miR-125a	Genome browser	Homo sapiens FK506 binding protein 2, 13kDa (FKBP2), transcript variant 2, mRNA.
68	NM_014690	All miRNA predictions	4.99	hsa-miR-125a	Genome browser	Homo sapiens KIAA0773 gene product (KIAA0773), mRNA.
69	NM_003983	All miRNA predictions	4.97	hsa-miR-125a	Genome browser	Homo sapiens solute carrier family 7 (cationic amino acid transporter, y ⁺ system), member 6 (SLC7A6), mRNA.
70	NM_003893	All miRNA predictions	4.94	hsa-miR-125a	Genome browser	Homo sapiens LIM domain binding 1 (LDB1), mRNA.
71	NM_138340	All miRNA predictions	4.91	hsa-miR-125a	Genome browser	Homo sapiens abhydrolase domain containing 3 (ABHD3), mRNA.
72	NM_032857	All miRNA predictions	4.90	hsa-miR-125a	Genome browser	Homo sapiens lactamase, beta (LACTB), nuclear gene encoding mitochondrial protein, transcript variant 1, mRNA.
73	NM_002568	All miRNA predictions	4.90	hsa-miR-125a	Genome browser	Homo sapiens poly(A) binding protein, cytoplasmic 1 (PABPC1), mRNA.
74	NM_015899	All miRNA predictions	4.88	hsa-miR-125a	Genome browser	Homo sapiens pleckstrin homology domain containing, family A (phosphoinositide binding specific) member 9 (PLEKHA9), mRNA.
75	NM_004095	All miRNA predictions	4.87	hsa-miR-125a	Genome browser	Homo sapiens eukaryotic translation initiation factor 4E binding protein 1 (EIF4EBP1), mRNA.
76	NM_033272	All miRNA predictions	4.86	hsa-miR-125a	Genome browser	Homo sapiens potassium voltage-gated channel, subfamily H (eag-related), member 7 (KCNH7), transcript variant 1, mRNA.
77	NM_020309	All miRNA predictions	4.86	hsa-miR-125a	Genome browser	Homo sapiens solute carrier family 17 (sodium-dependent inorganic phosphate cotransporter),

						member 7 (SLC17A7), mRNA.
78	NM_199054	All miRNA predictions	4.84	hsa-miR-125a	Genome browser	Homo sapiens MAP kinase interacting serine/threonine kinase 2 (MKNK2), mRNA.
79	NM_004901	All miRNA predictions	4.84	hsa-miR-125a	Genome browser	Homo sapiens ectonucleoside triphosphate diphosphohydrolase 4 (ENTPD4), mRNA.
80	NM_014871	All miRNA predictions	4.84	hsa-miR-125a	Genome browser	Homo sapiens ubiquitin specific protease 52 (USP52), mRNA.
81	NM_004073	All miRNA predictions	4.82	hsa-miR-125a	Genome browser	Homo sapiens polo-like kinase 3 (Drosophila) (PLK3), mRNA.
82	NM_003427	All miRNA predictions	4.81	hsa-miR-125a	Genome browser	Homo sapiens zinc finger protein 76 (expressed in testis) (ZNF76), mRNA.
83	NM_030645	All miRNA predictions	4.81	hsa-miR-125a	Genome browser	Homo sapiens KIAA1720 protein (KIAA1720), mRNA.
84	NM_032325	All miRNA predictions	4.75	hsa-miR-125a	Genome browser	Homo sapiens hypothetical protein MGC11102 (MGC11102), mRNA.
85	NM_022778	All miRNA predictions	4.70	hsa-miR-125a	Genome browser	Homo sapiens hypothetical protein DKFZp434L0117 (DKFZP434L0117), mRNA.
86	NM_003211	All miRNA predictions	4.67	hsa-miR-125a	Genome browser	Homo sapiens thymine-DNA glycosylase (TDG), mRNA.
87	NM_005176	All miRNA predictions	4.63	hsa-miR-125a	Genome browser	Homo sapiens ATP synthase, H ⁺ transporting, mitochondrial F0 complex, subunit c (subunit 9), isoform 2 (ATP5G2), nuclear gene encoding mitochondrial protein, transcript variant 2, mRNA.
88	NM_001070	All miRNA predictions	4.63	hsa-miR-125a	Genome browser	Homo sapiens tubulin, gamma 1 (TUBG1), mRNA.
89	NM_001002031	All miRNA predictions	4.63	hsa-miR-125a	Genome browser	Homo sapiens ATP synthase, H ⁺ transporting, mitochondrial F0 complex, subunit c (subunit 9), isoform 2 (ATP5G2), nuclear gene encoding mitochondrial protein, transcript variant 1, mRNA.
90	NM_024789	All miRNA predictions	4.60	hsa-miR-125a	Genome browser	Homo sapiens chromosome 10 open reading frame 77 (C10orf77), mRNA.
91	NM_178441	All miRNA predictions	4.58	hsa-miR-125a	Genome browser	Homo sapiens zinc finger, FYVE domain containing 1 (ZFYVE1), transcript variant 2, mRNA.
92	NM_021260	All miRNA predictions	4.58	hsa-miR-125a	Genome browser	Homo sapiens zinc finger, FYVE domain containing 1 (ZFYVE1), transcript variant 1, mRNA.
93	NM_207335	All miRNA predictions	4.58	hsa-miR-125a	Genome browser	Homo sapiens FLJ46299 protein (FLJ46299), mRNA.
94	NM_024923	All miRNA predictions	4.57	hsa-miR-125a	Genome browser	Homo sapiens nucleoporin 210kDa (NUP210), mRNA.
95	NM_006481	All miRNA predictions	4.55	hsa-miR-125a	Genome browser	Homo sapiens transcription factor 2, hepatic; LF-B3; variant hepatic nuclear factor (TCF2), transcript variant b, mRNA.
96	NM_015288	All miRNA predictions	4.54	hsa-miR-125a	Genome browser	Homo sapiens PHD finger protein 15 (PHF15), mRNA.
97	NM_181313	All miRNA predictions	4.51	hsa-miR-125a	Genome browser	Homo sapiens tafazzin (cardiomyopathy, dilated 3A (X-linked); endocardial fibroelastosis 2; Barth syndrome) (TAZ), transcript variant 4, mRNA.
98	NM_000116	All miRNA predictions	4.51	hsa-miR-125a	Genome browser	Homo sapiens tafazzin (cardiomyopathy, dilated 3A (X-linked); endocardial fibroelastosis 2; Barth syndrome) (TAZ), transcript variant 1, mRNA.
99	NM_181312	All miRNA predictions	4.51	hsa-miR-125a	Genome browser	Homo sapiens tafazzin (cardiomyopathy, dilated 3A (X-linked); endocardial fibroelastosis 2; Barth syndrome) (TAZ), transcript variant 3, mRNA.
100	NM_181311	All miRNA predictions	4.51	hsa-miR-125a	Genome browser	Homo sapiens tafazzin (cardiomyopathy, dilated 3A (X-linked); endocardial fibroelastosis 2; Barth syndrome) (TAZ), transcript variant 2, mRNA.
101	NM_016538	All miRNA predictions	4.51	hsa-miR-125a	Genome browser	Homo sapiens sirtuin (silent mating type information regulation 2 homolog) 7 (<i>S. cerevisiae</i>) (SIRT7), mRNA.
102	NM_016147	All miRNA predictions	4.47	hsa-miR-125a	Genome browser	Homo sapiens protein phosphatase methylesterase-1 (PME-1), mRNA.
103	NM_198925	All miRNA predictions	4.46	hsa-miR-125a	Genome browser	Homo sapiens sema domain, immunoglobulin domain (Ig), transmembrane domain (TM) and short cytoplasmic domain, (semaphorin) 4B (SEMA4B), transcript variant 2, mRNA.
104	NM_020210	All miRNA	4.46	hsa-miR-	Genome	Homo sapiens sema domain, immunoglobulin

		predictions		125a	browser	
						domain (Ig), transmembrane domain (TM) and short cytoplasmic domain, (semaphorin) 4B (SEMA4B), transcript variant 1, mRNA.
105	NM_134440	All miRNA predictions	4.42	hsa-miR-125a	Genome browser	Homo sapiens regulatory factor X-associated ankyrin-containing protein (RFXANK), transcript variant 2, mRNA.
106	NM_003721	All miRNA predictions	4.42	hsa-miR-125a	Genome browser	Homo sapiens regulatory factor X-associated ankyrin-containing protein (RFXANK), transcript variant 1, mRNA.
107	NM_001655	All miRNA predictions	4.39	hsa-miR-125a	Genome browser	Homo sapiens archain 1 (ARCN1), mRNA.
108	NM_198279	All miRNA predictions	4.39	hsa-miR-125a	Genome browser	Homo sapiens chromosome X open reading frame 23 (CXorf23), mRNA.
109	NM_003924	All miRNA predictions	4.38	hsa-miR-125a	Genome browser	Homo sapiens paired-like homeobox 2b (PHOX2B), mRNA.
110	NM_016090	All miRNA predictions	4.37	hsa-miR-125a	Genome browser	Homo sapiens RNA binding motif protein 7 (RBM7), mRNA.
111	NM_033449	All miRNA predictions	4.36	hsa-miR-125a	Genome browser	Homo sapiens FCH and double SH3 domains 1 (FCHSD1), mRNA.
112	NM_006454	All miRNA predictions	4.36	hsa-miR-125a	Genome browser	Homo sapiens MAX dimerization protein 4 (MXD4), mRNA.
113	NM_016436	All miRNA predictions	4.35	hsa-miR-125a	Genome browser	Homo sapiens chromosome 20 open reading frame 104 (C20orf104), mRNA.
114	NM_002252	All miRNA predictions	4.31	hsa-miR-125a	Genome browser	Homo sapiens potassium voltage-gated channel, delayed-rectifier, subfamily S, member 3 (KCNS3), mRNA.
115	NM_016399	All miRNA predictions	4.29	hsa-miR-125a	Genome browser	Homo sapiens hypothetical protein HSPC132 (HSPC132), mRNA.
116	NM_014759	All miRNA predictions	4.26	hsa-miR-125a	Genome browser	Homo sapiens phytanoyl-CoA hydroxylase interacting protein (PHYHIP), mRNA.
117	NM_153267	All miRNA predictions	4.25	hsa-miR-125a	Genome browser	Homo sapiens MAM domain containing 2 (MAMDC2), mRNA.
118	NM_015292	All miRNA predictions	4.24	hsa-miR-125a	Genome browser	Homo sapiens likely ortholog of mouse membrane bound C2 domain containing protein (MBC2), mRNA.
119	NM_152236	All miRNA predictions	4.21	hsa-miR-125a	Genome browser	Homo sapiens growth arrest-specific 2 like 1 (GAS2L1), transcript variant 2, mRNA.
120	NM_006478	All miRNA predictions	4.21	hsa-miR-125a	Genome browser	Homo sapiens growth arrest-specific 2 like 1 (GAS2L1), transcript variant 1, mRNA.
121	NM_153354	All miRNA predictions	4.19	hsa-miR-125a	Genome browser	Homo sapiens hypothetical protein MGC33214 (MGC33214), mRNA.
122	NM_020453	All miRNA predictions	4.17	hsa-miR-125a	Genome browser	Homo sapiens ATPase, Class V, type 10D (ATP10D), mRNA.
123	NM_005238	All miRNA predictions	4.16	hsa-miR-125a	Genome browser	Homo sapiens v-ets erythroblastosis virus E26 oncogene homolog 1 (avian) (ETS1), mRNA.
124	NM_003369	All miRNA predictions	4.16	hsa-miR-125a	Genome browser	Homo sapiens UV radiation resistance associated gene (UVRAG), mRNA.
125	NM_003605	All miRNA predictions	4.15	hsa-miR-125a	Genome browser	Homo sapiens O-linked N-acetylglucosamine (GlcNAc) transferase (UDP-N-acetylglucosamine:polypeptide-N-acetylglucosaminyl transferase) (OGT), transcript variant 3, mRNA.
126	NM_001001933	All miRNA predictions	4.13	hsa-miR-125a	Genome browser	Homo sapiens LIM homeobox 8 (LHX8), mRNA.
127	NM_004716	All miRNA predictions	4.13	hsa-miR-125a	Genome browser	Homo sapiens proprotein convertase subtilisin/kexin type 7 (PCSK7), mRNA.
128	NM_005113	All miRNA predictions	4.13	hsa-miR-125a	Genome browser	Homo sapiens golgi autoantigen, golgin subfamily a, 5 (GOLGA5), mRNA.
129	NM_005740	All miRNA predictions	4.13	hsa-miR-125a	Genome browser	Homo sapiens dynein, axonemal, light polypeptide 4 (DNAL4), mRNA.
130	NM_015866	All miRNA predictions	4.13	hsa-miR-125a	Genome browser	Homo sapiens PR domain containing 2, with ZNF domain (PRDM2), transcript variant 2, mRNA.
131	NM_005845	All miRNA predictions	4.09	hsa-miR-125a	Genome browser	Homo sapiens ATP-binding cassette, sub-family C (CFTR/MRP), member 4 (ABCC4), mRNA.
132	NM_002033	All miRNA predictions	4.09	hsa-miR-125a	Genome browser	Homo sapiens fucosyltransferase 4 (alpha (1,3) fucosyltransferase, myeloid-specific) (FUT4), mRNA.
133	NM_004513	All miRNA predictions	4.07	hsa-miR-125a	Genome browser	Homo sapiens interleukin 16 (lymphocyte chemoattractant factor) (IL16), transcript variant 1,

						mRNA.
134	NM_207577	All miRNA predictions	4.06	hsa-miR-125a	Genome browser	Homo sapiens microtubule-associated protein 6 (MAP6), transcript variant 2, mRNA.
135	NM_005576	All miRNA predictions	4.03	hsa-miR-125a	Genome browser	Homo sapiens lysyl oxidase-like 1 (LOXL1), mRNA.
136	NM_024524	All miRNA predictions	4.02	hsa-miR-125a	Genome browser	Homo sapiens ATPase family homolog up-regulated in senescence cells (AFURS1), mRNA.
137	NM_005439	All miRNA predictions	4.02	hsa-miR-125a	Genome browser	Homo sapiens myeloid leukemia factor 2 (MLF2), mRNA.
138	NM_133455	All miRNA predictions	4.01	hsa-miR-125a	Genome browser	Homo sapiens EMI domain containing 1 (EMID1), mRNA.
139	NM_005575	All miRNA predictions	4.01	hsa-miR-125a	Genome browser	Homo sapiens leucyl/cystinyl aminopeptidase (LNPEP), mRNA.
140	NM_053029	All miRNA predictions	4.01	hsa-miR-125a	Genome browser	Homo sapiens myosin, light polypeptide kinase (MYLK), transcript variant 4, mRNA.
141	NM_002959	All miRNA predictions	3.99	hsa-miR-125a	Genome browser	Homo sapiens sortilin 1 (SORT1), mRNA.
142	NM_173514	All miRNA predictions	3.99	hsa-miR-125a	Genome browser	Homo sapiens hypothetical protein FLJ90709 (FLJ90709), mRNA.
143	NM_021213	All miRNA predictions	3.98	hsa-miR-125a	Genome browser	Homo sapiens phosphatidylcholine transfer protein (PCTP), mRNA.
144	NM_022140	All miRNA predictions	3.94	hsa-miR-125a	Genome browser	Homo sapiens erythrocyte membrane protein band 4.1 like 4A (EPB41L4A), mRNA.
145	NM_019005	All miRNA predictions	3.91	hsa-miR-125a	Genome browser	Homo sapiens hypothetical protein FLJ20323 (FLJ20323), mRNA.
146	NM_002460	All miRNA predictions	3.91	hsa-miR-125a	Genome browser	Homo sapiens interferon regulatory factor 4 (IRF4), mRNA.
147	NM_018243	All miRNA predictions	3.90	hsa-miR-125a	Genome browser	Homo sapiens septin 11 (SEPT11), mRNA.
148	NM_148907	All miRNA predictions	3.87	hsa-miR-125a	Genome browser	Homo sapiens oxysterol binding protein-like 9 (OSBPL9), transcript variant 4, mRNA.
149	NM_148905	All miRNA predictions	3.87	hsa-miR-125a	Genome browser	Homo sapiens oxysterol binding protein-like 9 (OSBPL9), transcript variant 2, mRNA.
150	NM_148904	All miRNA predictions	3.87	hsa-miR-125a	Genome browser	Homo sapiens oxysterol binding protein-like 9 (OSBPL9), transcript variant 1, mRNA.
151	NM_024586	All miRNA predictions	3.87	hsa-miR-125a	Genome browser	Homo sapiens oxysterol binding protein-like 9 (OSBPL9), transcript variant 6, mRNA.
152	NM_148909	All miRNA predictions	3.87	hsa-miR-125a	Genome browser	Homo sapiens oxysterol binding protein-like 9 (OSBPL9), transcript variant 7, mRNA.
153	NM_005382	All miRNA predictions	3.87	hsa-miR-125a	Genome browser	Homo sapiens neurofilament 3 (150kDa medium) (NEF3), mRNA.
154	NM_148906	All miRNA predictions	3.87	hsa-miR-125a	Genome browser	Homo sapiens oxysterol binding protein-like 9 (OSBPL9), transcript variant 3, mRNA.
155	NM_148908	All miRNA predictions	3.87	hsa-miR-125a	Genome browser	Homo sapiens oxysterol binding protein-like 9 (OSBPL9), transcript variant 5, mRNA.
156	NM_024701	All miRNA predictions	3.86	hsa-miR-125a	Genome browser	Homo sapiens ankyrin repeat and SOCS box-containing 13 (ASB13), mRNA.
157	NM_015955	All miRNA predictions	3.84	hsa-miR-125a	Genome browser	Homo sapiens chromosome 2 open reading frame 4 (C2orf4), mRNA.
158	NM_022058	All miRNA predictions	3.83	hsa-miR-125a	Genome browser	Homo sapiens solute carrier family 4, sodium bicarbonate transporter-like, member 10 (SLC4A10), mRNA.
159	NM_007260	All miRNA predictions	3.81	hsa-miR-125a	Genome browser	Homo sapiens lysophospholipase II (LYPLA2), mRNA.
160	NM_207446	All miRNA predictions	3.81	hsa-miR-125a	Genome browser	Homo sapiens hypothetical gene supported by AK075564; BC060873 (LOC400451), mRNA.
161	NM_000332	All miRNA predictions	3.79	hsa-miR-125a	Genome browser	Homo sapiens ataxin 1 (ATXN1), mRNA.
162	NM_015911	All miRNA predictions	3.79	hsa-miR-125a	Genome browser	Homo sapiens hypothetical protein LOC51058 (LOC51058), mRNA.
163	NM_000038	All miRNA predictions	3.77	hsa-miR-125a	Genome browser	Homo sapiens adenomatous polyposis coli (APC), mRNA.
164	NM_032154	All miRNA predictions	3.77	hsa-miR-125a	Genome browser	Homo sapiens ring finger protein 134 (RNF134), mRNA.
165	NM_023019	All miRNA	3.77	hsa-miR-	Genome	Homo sapiens dynactin 1 (p150, glued homolog,

		predictions		125a	browser	
						Drosophila (DCTN1), transcript variant 2, mRNA.
166	NM_152729	All miRNA predictions	3.77	hsa-miR-125a	Genome browser	Homo sapiens 5'-nucleotidase, cytosolic II-like 1 (NT5C2L1), mRNA.
167	NM_004082	All miRNA predictions	3.77	hsa-miR-125a	Genome browser	Homo sapiens dynactin 1 (p150, glued homolog, Drosophila) (DCTN1), transcript variant 1, mRNA.
168	NM_003458	All miRNA predictions	3.76	hsa-miR-125a	Genome browser	Homo sapiens bassoon (presynaptic cytomatrix protein) (BSN), mRNA.
169	NM_002112	All miRNA predictions	3.75	hsa-miR-125a	Genome browser	Homo sapiens histidine decarboxylase (HDC), mRNA.
170	NM_014771	All miRNA predictions	3.75	hsa-miR-125a	Genome browser	Homo sapiens ring finger protein 40 (RNF40), transcript variant 1, mRNA.
171	NM_001448	All miRNA predictions	3.74	hsa-miR-125a	Genome browser	Homo sapiens glypican 4 (GPC4), mRNA.
172	NM_024667	All miRNA predictions	3.74	hsa-miR-125a	Genome browser	Homo sapiens hypothetical protein FLJ12750 (FLJ12750), mRNA.
173	NM_032701	All miRNA predictions	3.71	hsa-miR-125a	Genome browser	Homo sapiens suppressor of variegation 4-20 homolog 2 (Drosophila) (SUV420H2), mRNA.
174	NM_152678	All miRNA predictions	3.71	hsa-miR-125a	Genome browser	Homo sapiens hypothetical protein FLJ34969 (FLJ34969), mRNA.
175	NM_033143	All miRNA predictions	3.71	hsa-miR-125a	Genome browser	Homo sapiens fibroblast growth factor 5 (FGF5), transcript variant 2, mRNA.
176	NM_130476	All miRNA predictions	3.70	hsa-miR-125a	Genome browser	Homo sapiens MAP-kinase activating death domain (MADD), transcript variant 8, mRNA.
177	NM_130470	All miRNA predictions	3.69	hsa-miR-125a	Genome browser	Homo sapiens MAP-kinase activating death domain (MADD), transcript variant 1, mRNA.
178	NM_130471	All miRNA predictions	3.69	hsa-miR-125a	Genome browser	Homo sapiens MAP-kinase activating death domain (MADD), transcript variant 2, mRNA.
179	NM_130472	All miRNA predictions	3.69	hsa-miR-125a	Genome browser	Homo sapiens MAP-kinase activating death domain (MADD), transcript variant 3, mRNA.
180	NM_130473	All miRNA predictions	3.69	hsa-miR-125a	Genome browser	Homo sapiens MAP-kinase activating death domain (MADD), transcript variant 5, mRNA.
181	NM_003682	All miRNA predictions	3.69	hsa-miR-125a	Genome browser	Homo sapiens MAP-kinase activating death domain (MADD), transcript variant 4, mRNA.
182	NM_014603	All miRNA predictions	3.68	hsa-miR-125a	Genome browser	Homo sapiens paraneoplastic antigen (HUMPPA), mRNA.
183	NM_002657	All miRNA predictions	3.67	hsa-miR-125a	Genome browser	Homo sapiens pleiomorphic adenoma gene-like 2 (PLAGL2), mRNA.
184	NM_012482	All miRNA predictions	3.67	hsa-miR-125a	Genome browser	Homo sapiens zinc finger protein 281 (ZNF281), mRNA.
185	NM_015076	All miRNA predictions	3.65	hsa-miR-125a	Genome browser	Homo sapiens cyclin-dependent kinase (CDC2-like) 11 (CDK11), mRNA.
186	NM_207346	All miRNA predictions	3.65	hsa-miR-125a	Genome browser	Homo sapiens likely homolog of yeast SEN54 (SEN54L), mRNA.
187	NM_014735	All miRNA predictions	3.65	hsa-miR-125a	Genome browser	Homo sapiens PHD finger protein 16 (PHF16), mRNA.
188	NM_181314	All miRNA predictions	3.64	hsa-miR-125a	Genome browser	Homo sapiens tafazzin (cardiomyopathy, dilated 3A (X-linked); endocardial fibroelastosis 2; Barth syndrome) (TAZ), transcript variant 5, mRNA.
189	NM_015074	All miRNA predictions	3.64	hsa-miR-125a	Genome browser	Homo sapiens kinesin family member 1B (KIF1B), transcript variant 1, mRNA.
190	NM_002446	All miRNA predictions	3.63	hsa-miR-125a	Genome browser	Homo sapiens mitogen-activated protein kinase kinase kinase 10 (MAP3K10), mRNA.
191	NM_001987	All miRNA predictions	3.63	hsa-miR-125a	Genome browser	Homo sapiens ets variant gene 6 (TEL oncogene) (ETV6), mRNA.
192	NM_015518	All miRNA predictions	3.61	hsa-miR-125a	Genome browser	Homo sapiens DKFZP434C131 protein (DKFZP434C131), mRNA.
193	NM_014552	All miRNA predictions	3.60	hsa-miR-125a	Genome browser	Homo sapiens transcription factor CP2-like 2 (TFCP2L2), transcript variant 1, mRNA.
194	NM_017510	All miRNA predictions	3.60	hsa-miR-125a	Genome browser	Homo sapiens gp25L2 protein (HSGP25L2G), mRNA.
195	NM_172027	All miRNA predictions	3.60	hsa-miR-125a	Genome browser	Homo sapiens ankyrin repeat and BTB (POZ) domain containing 1 (ABTB1), transcript variant 2, mRNA.
196	NM_032548	All miRNA predictions	3.60	hsa-miR-125a	Genome browser	Homo sapiens ankyrin repeat and BTB (POZ) domain containing 1 (ABTB1), transcript variant 1, mRNA.
197	NM_014901	All miRNA	3.60	hsa-miR-	Genome	Homo sapiens ring finger protein 44 (RNF44), mRNA.

		predictions		125a	browser	
198	NM_198182	All miRNA predictions	3.60	hsa-miR-125a	Genome browser	Homo sapiens transcription factor CP2-like 2 (TFCP2L2), transcript variant 2, mRNA.
199	NM_172028	All miRNA predictions	3.60	hsa-miR-125a	Genome browser	Homo sapiens ankyrin repeat and BTB (POZ) domain containing 1 (ABTB1), transcript variant 3, mRNA.
200	NM_023016	All miRNA predictions	3.59	hsa-miR-125a	Genome browser	Homo sapiens chromosome 2 open reading frame 26 (C2orf26), mRNA.
201	NM_020348	All miRNA predictions	3.59	hsa-miR-125a	Genome browser	Homo sapiens cyclin M1 (CNNM1), mRNA.
202	NM_152663	All miRNA predictions	3.56	hsa-miR-125a	Genome browser	Homo sapiens Ral GEF with PH domain and SH3 binding motif 2 (RALGPS2), mRNA.
203	NM_130475	All miRNA predictions	3.54	hsa-miR-125a	Genome browser	Homo sapiens MAP-kinase activating death domain (MADD), transcript variant 7, mRNA.
204	NM_004118	All miRNA predictions	3.54	hsa-miR-125a	Genome browser	Homo sapiens forkhead-like 18 (Drosophila) (FKHL18), mRNA.
205	NM_130474	All miRNA predictions	3.54	hsa-miR-125a	Genome browser	Homo sapiens MAP-kinase activating death domain (MADD), transcript variant 6, mRNA.
206	NM_052820	All miRNA predictions	3.53	hsa-miR-125a	Genome browser	Homo sapiens coronin, actin binding protein, 2A (CORO2A), transcript variant 2, mRNA.
207	NM_003389	All miRNA predictions	3.53	hsa-miR-125a	Genome browser	Homo sapiens coronin, actin binding protein, 2A (CORO2A), transcript variant 1, mRNA.
208	NM_183244	All miRNA predictions	3.53	hsa-miR-125a	Genome browser	Homo sapiens phosphatase and actin regulator 3 (PHACTR3), transcript variant 2, mRNA.
209	NM_183246	All miRNA predictions	3.53	hsa-miR-125a	Genome browser	Homo sapiens phosphatase and actin regulator 3 (PHACTR3), transcript variant 3, mRNA.
210	NM_021962	All miRNA predictions	3.52	hsa-miR-125a	Genome browser	Homo sapiens active BCR-related gene (ABR), transcript variant 1, mRNA.
211	NM_080672	All miRNA predictions	3.52	hsa-miR-125a	Genome browser	Homo sapiens phosphatase and actin regulator 3 (PHACTR3), transcript variant 1, mRNA.
212	NM_001092	All miRNA predictions	3.52	hsa-miR-125a	Genome browser	Homo sapiens active BCR-related gene (ABR), transcript variant 2, mRNA.
213	NM_014905	All miRNA predictions	3.50	hsa-miR-125a	Genome browser	Homo sapiens glutaminase (GLS), mRNA.
214	NM_017495	All miRNA predictions	3.50	hsa-miR-125a	Genome browser	Homo sapiens RNA-binding region (RNP1, RRM) containing 1 (RNPC1), transcript variant 1, mRNA.
215	NM_022833	All miRNA predictions	3.49	hsa-miR-125a	Genome browser	Homo sapiens chromosome 9 open reading frame 88 (C9orf88), mRNA.
216	NM_003690	All miRNA predictions	3.49	hsa-miR-125a	Genome browser	Homo sapiens protein kinase, interferon-inducible double stranded RNA dependent activator (PRKRA), mRNA.
217	NM_001204	All miRNA predictions	3.49	hsa-miR-125a	Genome browser	Homo sapiens bone morphogenetic protein receptor, type II (serine/threonine kinase) (BMP2), transcript variant 1, mRNA.
218	NM_017911	All miRNA predictions	3.48	hsa-miR-125a	Genome browser	Homo sapiens chromosome 22 open reading frame 8 (C22orf8), mRNA.
219	NM_003759	All miRNA predictions	3.48	hsa-miR-125a	Genome browser	Homo sapiens solute carrier family 4, sodium bicarbonate cotransporter, member 4 (SLC4A4), mRNA.
220	NM_033540	All miRNA predictions	3.47	hsa-miR-125a	Genome browser	Homo sapiens mitofusin 1 (MFN1), nuclear gene encoding mitochondrial protein, transcript variant 1, mRNA.
221	NM_013272	All miRNA predictions	3.46	hsa-miR-125a	Genome browser	Homo sapiens solute carrier organic anion transporter family, member 3A1 (SLCO3A1), mRNA.
222	NM_080491	All miRNA predictions	3.46	hsa-miR-125a	Genome browser	Homo sapiens GRB2-associated binding protein 2 (GAB2), transcript variant 1, mRNA.
223	NM_002207	All miRNA predictions	3.46	hsa-miR-125a	Genome browser	Homo sapiens integrin, alpha 9 (ITGA9), mRNA.
224	NM_003957	All miRNA predictions	3.46	hsa-miR-125a	Genome browser	Homo sapiens serine/threonine kinase 29 (STK29), mRNA.
225	NM_012296	All miRNA predictions	3.46	hsa-miR-125a	Genome browser	Homo sapiens GRB2-associated binding protein 2 (GAB2), transcript variant 2, mRNA.
226	NM_032557	All miRNA predictions	3.44	hsa-miR-125a	Genome browser	Homo sapiens ubiquitin specific protease 38 (USP38), mRNA.
227	NM_203403	All miRNA predictions	3.44	hsa-miR-125a	Genome browser	Homo sapiens chromosome 9 open reading frame 150 (C9orf150), mRNA.

228	NM_178454	All miRNA predictions	3.44	hsa-miR-125a	Genome browser	Homo sapiens hypothetical protein MGC54289 (MGC54289), mRNA.
229	NM_003330	All miRNA predictions	3.43	hsa-miR-125a	Genome browser	Homo sapiens thioredoxin reductase 1 (TXNRD1), transcript variant 1, mRNA.
230	NM_182743	All miRNA predictions	3.43	hsa-miR-125a	Genome browser	Homo sapiens thioredoxin reductase 1 (TXNRD1), transcript variant 4, mRNA.
231	NM_182729	All miRNA predictions	3.43	hsa-miR-125a	Genome browser	Homo sapiens thioredoxin reductase 1 (TXNRD1), transcript variant 5, mRNA.
232	NM_182742	All miRNA predictions	3.43	hsa-miR-125a	Genome browser	Homo sapiens thioredoxin reductase 1 (TXNRD1), transcript variant 2, mRNA.
233	NM_018323	All miRNA predictions	3.38	hsa-miR-125a	Genome browser	Homo sapiens phosphatidylinositol 4-kinase type-II beta (PI4K2B), mRNA.
234	NM_016339	All miRNA predictions	3.37	hsa-miR-125a	Genome browser	Homo sapiens Rap guanine nucleotide exchange factor (GEF)-like 1 (RAPGEFL1), mRNA.
235	NM_020935	All miRNA predictions	3.37	hsa-miR-125a	Genome browser	Homo sapiens ubiquitin specific protease 37 (USP37), mRNA.
236	NM_183425	All miRNA predictions	3.36	hsa-miR-125a	Genome browser	Homo sapiens RNA-binding region (RNP1, RRM) containing 1 (RNPC1), transcript variant 2, mRNA.
237	NM_033346	All miRNA predictions	3.34	hsa-miR-125a	Genome browser	Homo sapiens bone morphogenetic protein receptor, type II (serine/threonine kinase) (BMPR2), transcript variant 2, mRNA.
238	NM_182966	All miRNA predictions	3.32	hsa-miR-125a	Genome browser	Homo sapiens neural precursor cell expressed, developmentally down-regulated 9 (NEDD9), mRNA.
239	NM_004540	All miRNA predictions	3.32	hsa-miR-125a	Genome browser	Homo sapiens neural cell adhesion molecule 2 (NCAM2), mRNA.
240	NM_032046	All miRNA predictions	3.32	hsa-miR-125a	Genome browser	Homo sapiens mosaic serine protease (MSP), mRNA.
241	NM_002092	All miRNA predictions	3.32	hsa-miR-125a	Genome browser	Homo sapiens G-rich RNA sequence binding factor 1 (GRSF1), mRNA.
242	NM_015517	All miRNA predictions	3.31	hsa-miR-125a	Genome browser	Homo sapiens MBD2 (methyl-CpG-binding protein)-interacting zinc finger protein (MIZF), transcript variant 1, mRNA.
243	NM_002419	All miRNA predictions	3.31	hsa-miR-125a	Genome browser	Homo sapiens mitogen-activated protein kinase kinase kinase 11 (MAP3K11), mRNA.
244	NM_198971	All miRNA predictions	3.31	hsa-miR-125a	Genome browser	Homo sapiens MBD2 (methyl-CpG-binding protein)-interacting zinc finger protein (MIZF), transcript variant 2, mRNA.
245	NM_001955	All miRNA predictions	3.30	hsa-miR-125a	Genome browser	Homo sapiens endothelin 1 (EDN1), mRNA.
246	NM_003816	All miRNA predictions	3.30	hsa-miR-125a	Genome browser	Homo sapiens a disintegrin and metalloproteinase domain 9 (meltrin gamma) (ADAM9), transcript variant 1, mRNA.
247	NM_024718	All miRNA predictions	3.29	hsa-miR-125a	Genome browser	Homo sapiens chromosome 9 open reading frame 86 (C9orf86), mRNA.
248	NM_007372	All miRNA predictions	3.29	hsa-miR-125a	Genome browser	Homo sapiens DEAD (Asp-Glu-Ala-Asp) box polypeptide 42 (DDX42), transcript variant 1, mRNA.
249	NM_203499	All miRNA predictions	3.29	hsa-miR-125a	Genome browser	Homo sapiens DEAD (Asp-Glu-Ala-Asp) box polypeptide 42 (DDX42), transcript variant 2, mRNA.
250	NM_005342	All miRNA predictions	3.28	hsa-miR-125a	Genome browser	Homo sapiens high-mobility group box 3 (HMGB3), mRNA.
251	NM_182488	All miRNA predictions	3.28	hsa-miR-125a	Genome browser	Homo sapiens ubiquitin specific protease 12 (USP12), mRNA.
252	NM_000434	All miRNA predictions	3.27	hsa-miR-125a	Genome browser	Homo sapiens sialidase 1 (lysosomal sialidase) (NEU1), mRNA.
253	NM_024090	All miRNA predictions	3.27	hsa-miR-125a	Genome browser	Homo sapiens ELOVL family member 6, elongation of long chain fatty acids (FEN1/Elo2, SUR4/Elo3-like, yeast) (ELOVL6), mRNA.
254	NM_139276	All miRNA predictions	3.25	hsa-miR-125a	Genome browser	Homo sapiens signal transducer and activator of transcription 3 (acute-phase response factor) (STAT3), transcript variant 1, mRNA.
255	NM_017813	All miRNA predictions	3.25	hsa-miR-125a	Genome browser	Homo sapiens hypothetical protein FLJ20421 (FLJ20421), mRNA.
256	NM_018108	All miRNA predictions	3.25	hsa-miR-125a	Genome browser	Homo sapiens chromosome 14 open reading frame 130 (C14orf130), mRNA.
257	NM_003150	All miRNA predictions	3.25	hsa-miR-125a	Genome browser	Homo sapiens signal transducer and activator of transcription 3 (acute-phase response factor) (STAT3),

						transcript variant 2, mRNA.
258	NM_022039	All miRNA predictions	3.24	hsa-miR-125a	Genome browser	Homo sapiens split hand/foot malformation (ectrodactyly) type 3 (SHFM3), mRNA.
259	NM_016219	All miRNA predictions	3.22	hsa-miR-125a	Genome browser	Homo sapiens mannosidase, alpha, class 1B, member 1 (MAN1B1), mRNA.
260	NM_007230	All miRNA predictions	3.22	hsa-miR-125a	Genome browser	n/a
261	NM_012154	All miRNA predictions	3.21	hsa-miR-125a	Genome browser	Homo sapiens eukaryotic translation initiation factor 2C, 2 (EIF2C2), mRNA.
262	NM_020630	All miRNA predictions	3.21	hsa-miR-125a	Genome browser	Homo sapiens ret proto-oncogene (multiple endocrine neoplasia and medullary thyroid carcinoma 1, Hirschsprung disease) (RET), transcript variant 4, mRNA.
263	NM_030576	All miRNA predictions	3.20	hsa-miR-125a	Genome browser	Homo sapiens hypothetical protein MGC10986 (MGC10986), mRNA.
264	NM_005889	All miRNA predictions	3.20	hsa-miR-125a	Genome browser	Homo sapiens apolipoprotein B mRNA editing enzyme, catalytic polypeptide 1 (APOBEC1), transcript variant 2, mRNA.
265	NM_021991	All miRNA predictions	3.19	hsa-miR-125a	Genome browser	Homo sapiens junction plakoglobin (JUP), transcript variant 2, mRNA.
266	NM_017740	All miRNA predictions	3.19	hsa-miR-125a	Genome browser	Homo sapiens zinc finger, DHHC domain containing 7 (ZDHHC7), mRNA.
267	NM_002656	All miRNA predictions	3.17	hsa-miR-125a	Genome browser	Homo sapiens pleiomorphic adenoma gene-like 1 (PLAGL1), transcript variant 1, mRNA.
268	NM_006718	All miRNA predictions	3.17	hsa-miR-125a	Genome browser	Homo sapiens pleiomorphic adenoma gene-like 1 (PLAGL1), transcript variant 2, mRNA.
269	NM_020921	All miRNA predictions	3.16	hsa-miR-125a	Genome browser	Homo sapiens ninein (GSK3B interacting protein) (NIN), transcript variant 2, mRNA.
270	NM_183011	All miRNA predictions	3.16	hsa-miR-125a	Genome browser	Homo sapiens cAMP responsive element modulator (CREM), transcript variant 21, mRNA.
271	NM_145313	All miRNA predictions	3.16	hsa-miR-125a	Genome browser	Homo sapiens RasGEF domain family, member 1A (RASGEF1A), mRNA.
272	NM_182717	All miRNA predictions	3.16	hsa-miR-125a	Genome browser	Homo sapiens cAMP responsive element modulator (CREM), transcript variant 4, mRNA.
273	NM_032364	All miRNA predictions	3.16	hsa-miR-125a	Genome browser	Homo sapiens DnaJ (Hsp40) homolog, subfamily C, member 14 (DNAJC14), mRNA.
274	NM_182721	All miRNA predictions	3.16	hsa-miR-125a	Genome browser	Homo sapiens cAMP responsive element modulator (CREM), transcript variant 8, mRNA.
275	NM_183013	All miRNA predictions	3.16	hsa-miR-125a	Genome browser	Homo sapiens cAMP responsive element modulator (CREM), transcript variant 19, mRNA.
276	NM_182853	All miRNA predictions	3.16	hsa-miR-125a	Genome browser	Homo sapiens cAMP responsive element modulator (CREM), transcript variant 18, mRNA.
277	NM_017811	All miRNA predictions	3.16	hsa-miR-125a	Genome browser	Homo sapiens ubiquitin-conjugating enzyme E2R 2 (UBE2R2), mRNA.
278	NM_182720	All miRNA predictions	3.16	hsa-miR-125a	Genome browser	Homo sapiens cAMP responsive element modulator (CREM), transcript variant 7, mRNA.
279	NM_017423	All miRNA predictions	3.16	hsa-miR-125a	Genome browser	Homo sapiens UDP-N-acetyl-alpha-D-galactosamine:polypeptide N-acetylgalactosaminyltransferase 7 (GalNac-T7) (GALNT7), mRNA.
280	NM_182723	All miRNA predictions	3.16	hsa-miR-125a	Genome browser	Homo sapiens cAMP responsive element modulator (CREM), transcript variant 10, mRNA.
281	NM_182722	All miRNA predictions	3.16	hsa-miR-125a	Genome browser	Homo sapiens cAMP responsive element modulator (CREM), transcript variant 9, mRNA.
282	NM_183012	All miRNA predictions	3.16	hsa-miR-125a	Genome browser	Homo sapiens cAMP responsive element modulator (CREM), transcript variant 20, mRNA.
283	NM_213662	All miRNA predictions	3.15	hsa-miR-125a	Genome browser	Homo sapiens signal transducer and activator of transcription 3 (acute-phase response factor) (STAT3), transcript variant 3, mRNA.
284	NM_002715	All miRNA predictions	3.14	hsa-miR-125a	Genome browser	Homo sapiens protein phosphatase 2 (formerly 2A), catalytic subunit, alpha isoform (PPP2CA), mRNA.
285	NM_024572	All miRNA predictions	3.14	hsa-miR-125a	Genome browser	Homo sapiens UDP-N-acetyl-alpha-D-galactosamine:polypeptide N-acetylgalactosaminyltransferase 14 (GalNac-T14) (GALNT14), mRNA.
286	NM_139275	All miRNA	3.12	hsa-miR-	Genome	Homo sapiens A kinase (PRKA) anchor protein 1

		predictions		125a	browser	
						(AKAP1), nuclear gene encoding mitochondrial protein, transcript variant 2, mRNA.
287	NM_001315	All miRNA predictions	3.11	hsa-miR-125a	Genome browser	Homo sapiens mitogen-activated protein kinase 14 (MAPK14), transcript variant 1, mRNA.
288	NM_139012	All miRNA predictions	3.11	hsa-miR-125a	Genome browser	Homo sapiens mitogen-activated protein kinase 14 (MAPK14), transcript variant 2, mRNA.
289	NM_015481	All miRNA predictions	3.10	hsa-miR-125a	Genome browser	Homo sapiens zinc finger protein 385 (ZNF385), mRNA.
290	NM_004756	All miRNA predictions	3.10	hsa-miR-125a	Genome browser	Homo sapiens numb homolog (Drosophila)-like (NUMBL), mRNA.
291	NM_001001702	All miRNA predictions	3.09	hsa-miR-125a	Genome browser	Homo sapiens FLJ33360 protein (FLJ33360), mRNA.
292	NM_001003891	All miRNA predictions	3.09	hsa-miR-125a	Genome browser	Homo sapiens PC2 (positive cofactor 2, multiprotein complex) glutamine/Q-rich-associated protein (PCQAP), transcript variant 1, mRNA.
293	NM_003598	All miRNA predictions	3.09	hsa-miR-125a	Genome browser	Homo sapiens TEA domain family member 2 (TEAD2), mRNA.
294	NM_015889	All miRNA predictions	3.09	hsa-miR-125a	Genome browser	Homo sapiens PC2 (positive cofactor 2, multiprotein complex) glutamine/Q-rich-associated protein (PCQAP), transcript variant 2, mRNA.
295	NM_152860	All miRNA predictions	3.08	hsa-miR-125a	Genome browser	Homo sapiens Sp7 transcription factor (SP7), mRNA.
296	NM_002646	All miRNA predictions	3.08	hsa-miR-125a	Genome browser	Homo sapiens phosphoinositide-3-kinase, class 2, beta polypeptide (PIK3C2B), mRNA.
297	NM_032229	All miRNA predictions	3.07	hsa-miR-125a	Genome browser	Homo sapiens SLIT and NTRK-like family, member 6 (SLITRK6), mRNA.
298	NM_005433	All miRNA predictions	3.06	hsa-miR-125a	Genome browser	Homo sapiens v-yes-1 Yamaguchi sarcoma viral oncogene homolog 1 (YES1), mRNA.
299	NM_181701	All miRNA predictions	3.06	hsa-miR-125a	Genome browser	Homo sapiens quiescin Q6-like 1 (QSCN6L1), mRNA.
300	NM_024293	All miRNA predictions	3.06	hsa-miR-125a	Genome browser	Homo sapiens chromosome 2 open reading frame 17 (C2orf17), mRNA.
301	NM_033260	All miRNA predictions	3.06	hsa-miR-125a	Genome browser	Homo sapiens forkhead box Q1 (FOXQ1), mRNA.
302	NM_033315	All miRNA predictions	3.06	hsa-miR-125a	Genome browser	Homo sapiens RAS-like, family 10, member B (RASL10B), mRNA.
303	NM_006734	All miRNA predictions	3.05	hsa-miR-125a	Genome browser	Homo sapiens human immunodeficiency virus type I enhancer binding protein 2 (HIVEP2), mRNA.
304	NM_006426	All miRNA predictions	3.05	hsa-miR-125a	Genome browser	Homo sapiens dihydropyrimidinase-like 4 (DPYSL4), mRNA.
305	NM_139014	All miRNA predictions	3.05	hsa-miR-125a	Genome browser	Homo sapiens mitogen-activated protein kinase 14 (MAPK14), transcript variant 4, mRNA.
306	NM_003488	All miRNA predictions	3.05	hsa-miR-125a	Genome browser	Homo sapiens A kinase (PRKA) anchor protein 1 (AKAP1), nuclear gene encoding mitochondrial protein, transcript variant 1, mRNA.
307	NM_003728	All miRNA predictions	3.04	hsa-miR-125a	Genome browser	Homo sapiens unc-5 homolog C (C. elegans) (UNC5C), mRNA.
308	NM_012338	All miRNA predictions	3.04	hsa-miR-125a	Genome browser	Homo sapiens transmembrane 4 superfamily member 12 (TM4SF12), mRNA.
309	NM_133625	All miRNA predictions	3.03	hsa-miR-125a	Genome browser	Homo sapiens synapsin II (SYN2), transcript variant IIa, mRNA.
310	NM_080670	All miRNA predictions	3.02	hsa-miR-125a	Genome browser	Homo sapiens solute carrier family 35, member A4 (SLC35A4), mRNA.
311	NM_004475	All miRNA predictions	3.01	hsa-miR-125a	Genome browser	Homo sapiens flotillin 2 (FLOT2), mRNA.
312	NM_001002924	All miRNA predictions	3.01	hsa-miR-125a	Genome browser	Homo sapiens adaptor-related protein complex 3, sigma 1 subunit (AP3S1), transcript variant 2, mRNA.
313	NM_032632	All miRNA predictions	3.00	hsa-miR-125a	Genome browser	Homo sapiens poly(A) polymerase alpha (PAPOLA), mRNA.
314	NM_001005476	All miRNA predictions	3.00	hsa-miR-125a	Genome browser	Homo sapiens plakophilin 4 (PKP4), transcript variant 2, mRNA.
315	NM_003628	All miRNA predictions	3.00	hsa-miR-125a	Genome browser	Homo sapiens plakophilin 4 (PKP4), transcript variant 1, mRNA.
316	NM_177560	All miRNA predictions	2.99	hsa-miR-125a	Genome browser	Homo sapiens casein kinase 2, alpha 1 polypeptide (CSNK2A1), transcript variant 3, mRNA.

317	NM_001895	All miRNA predictions	2.99	hsa-miR-125a	Genome browser	Homo sapiens casein kinase 2, alpha 1 polypeptide (CSNK2A1), transcript variant 2, mRNA.
318	NM_177559	All miRNA predictions	2.99	hsa-miR-125a	Genome browser	Homo sapiens casein kinase 2, alpha 1 polypeptide (CSNK2A1), transcript variant 1, mRNA.
319	NM_182757	All miRNA predictions	2.98	hsa-miR-125a	Genome browser	Homo sapiens IBR domain containing 2 (IBRDC2), mRNA.
320	NM_004634	All miRNA predictions	2.96	hsa-miR-125a	Genome browser	Homo sapiens bromodomain and PHD finger containing, 1 (BRPF1), transcript variant 2, mRNA.
321	NM_175884	All miRNA predictions	2.96	hsa-miR-125a	Genome browser	Homo sapiens hypothetical protein FLJ36031 (FLJ36031), mRNA.
322	NM_001003694	All miRNA predictions	2.96	hsa-miR-125a	Genome browser	Homo sapiens bromodomain and PHD finger containing, 1 (BRPF1), transcript variant 1, mRNA.
323	NM_194352	All miRNA predictions	2.95	hsa-miR-125a	Genome browser	Homo sapiens ring finger protein 40 (RNF40), transcript variant 2, mRNA.
324	NM_015609	All miRNA predictions	2.95	hsa-miR-125a	Genome browser	Homo sapiens putative MAPK activating protein PM20,PM21 (DKFZP566C0424), mRNA.
325	NM_170674	All miRNA predictions	2.93	hsa-miR-125a	Genome browser	Homo sapiens Meis1, myeloid ecotropic viral integration site 1 homolog 2 (mouse) (MEIS2), transcript variant b, mRNA.
326	NM_172315	All miRNA predictions	2.93	hsa-miR-125a	Genome browser	Homo sapiens Meis1, myeloid ecotropic viral integration site 1 homolog 2 (mouse) (MEIS2), transcript variant g, mRNA.
327	NM_002399	All miRNA predictions	2.93	hsa-miR-125a	Genome browser	Homo sapiens Meis1, myeloid ecotropic viral integration site 1 homolog 2 (mouse) (MEIS2), transcript variant f, mRNA.
328	NM_006268	All miRNA predictions	2.93	hsa-miR-125a	Genome browser	Homo sapiens D4, zinc and double PHD fingers family 2 (DPF2), mRNA.
329	NM_170677	All miRNA predictions	2.93	hsa-miR-125a	Genome browser	Homo sapiens Meis1, myeloid ecotropic viral integration site 1 homolog 2 (mouse) (MEIS2), transcript variant a, mRNA.
330	NM_172316	All miRNA predictions	2.93	hsa-miR-125a	Genome browser	Homo sapiens Meis1, myeloid ecotropic viral integration site 1 homolog 2 (mouse) (MEIS2), transcript variant h, mRNA.
331	NM_130459	All miRNA predictions	2.92	hsa-miR-125a	Genome browser	Homo sapiens torsin family 2, member A (TOR2A), mRNA.
332	NM_018275	All miRNA predictions	2.92	hsa-miR-125a	Genome browser	Homo sapiens hypothetical protein FLJ10925 (FLJ10925), mRNA.
333	NM_018488	All miRNA predictions	2.92	hsa-miR-125a	Genome browser	Homo sapiens T-box 4 (TBX4), mRNA.
334	NM_001002847	All miRNA predictions	2.91	hsa-miR-125a	Genome browser	Homo sapiens decreased expression in renal and prostate (DERPC), transcript variant 2, mRNA.
335	NM_017804	All miRNA predictions	2.91	hsa-miR-125a	Genome browser	Homo sapiens decreased expression in renal and prostate (DERPC), transcript variant 1, mRNA.
336	NM_002230	All miRNA predictions	2.91	hsa-miR-125a	Genome browser	Homo sapiens junction plakoglobin (JUP), transcript variant 1, mRNA.
337	NM_015975	All miRNA predictions	2.90	hsa-miR-125a	Genome browser	Homo sapiens TAF9-like RNA polymerase II, TATA box binding protein (TBP)-associated factor, 31kDa (TAF9L), mRNA.
338	NM_018195	All miRNA predictions	2.89	hsa-miR-125a	Genome browser	Homo sapiens hypothetical protein FLJ10726 (FLJ10726), mRNA.
339	NM_019085	All miRNA predictions	2.89	hsa-miR-125a	Genome browser	Homo sapiens F-box and leucine-rich repeat protein 19 (FBXL19), mRNA.
340	NM_153051	All miRNA predictions	2.89	hsa-miR-125a	Genome browser	Homo sapiens myotubularin related protein 3 (MTMR3), transcript variant 2, mRNA.
341	NM_006852	All miRNA predictions	2.89	hsa-miR-125a	Genome browser	Homo sapiens tousled-like kinase 2 (TLK2), mRNA.
342	NM_021090	All miRNA predictions	2.89	hsa-miR-125a	Genome browser	Homo sapiens myotubularin related protein 3 (MTMR3), transcript variant 3, mRNA.
343	NM_153050	All miRNA predictions	2.89	hsa-miR-125a	Genome browser	Homo sapiens myotubularin related protein 3 (MTMR3), transcript variant 1, mRNA.
344	NM_152704	All miRNA predictions	2.87	hsa-miR-125a	Genome browser	Homo sapiens hypothetical protein FLJ25477 (FLJ25477), transcript variant 1, mRNA.
345	NM_014764	All miRNA predictions	2.87	hsa-miR-125a	Genome browser	Homo sapiens DAZ associated protein 2 (DAZAP2), mRNA.
346	NM_199138	All miRNA predictions	2.87	hsa-miR-125a	Genome browser	Homo sapiens hypothetical protein FLJ25477 (FLJ25477), transcript variant 2, mRNA.

347	NM_170692	All miRNA predictions	2.85	hsa-miR-125a	Genome browser	Homo sapiens RAS protein activator like 2 (RASAL2), transcript variant 2, mRNA.
348	NM_004841	All miRNA predictions	2.85	hsa-miR-125a	Genome browser	Homo sapiens RAS protein activator like 2 (RASAL2), transcript variant 1, mRNA.
349	NM_153693	All miRNA predictions	2.84	hsa-miR-125a	Genome browser	Homo sapiens homeo box C6 (HOXC6), transcript variant 2, mRNA.
350	NM_004503	All miRNA predictions	2.84	hsa-miR-125a	Genome browser	Homo sapiens homeo box C6 (HOXC6), transcript variant 1, mRNA.
351	NM_032558	All miRNA predictions	2.82	hsa-miR-125a	Genome browser	Homo sapiens hypothetical protein FLJ14753 (FLJ14753), mRNA.
352	NM_022845	All miRNA predictions	2.78	hsa-miR-125a	Genome browser	Homo sapiens core-binding factor, beta subunit (CBFB), transcript variant 1, mRNA.
353	NM_005112	All miRNA predictions	2.77	hsa-miR-125a	Genome browser	Homo sapiens WD repeat domain 1 (WDR1), transcript variant 2, mRNA.
354	NM_022652	All miRNA predictions	2.77	hsa-miR-125a	Genome browser	Homo sapiens dual specificity phosphatase 6 (DUSP6), transcript variant 2, mRNA.
355	NM_017491	All miRNA predictions	2.77	hsa-miR-125a	Genome browser	Homo sapiens WD repeat domain 1 (WDR1), transcript variant 1, mRNA.
356	NM_015173	All miRNA predictions	2.77	hsa-miR-125a	Genome browser	Homo sapiens TBC1 (tre-2/USP6, BUB2, cdc16) domain family, member 1 (TBC1D1), mRNA.
357	NM_001001481	All miRNA predictions	2.77	hsa-miR-125a	Genome browser	Homo sapiens hypothetical protein FLJ11011 (FLJ11011), transcript variant 1, mRNA.
358	NM_018299	All miRNA predictions	2.77	hsa-miR-125a	Genome browser	Homo sapiens hypothetical protein FLJ11011 (FLJ11011), transcript variant 2, mRNA.
359	NM_001946	All miRNA predictions	2.77	hsa-miR-125a	Genome browser	Homo sapiens dual specificity phosphatase 6 (DUSP6), transcript variant 1, mRNA.
360	NM_020676	All miRNA predictions	2.76	hsa-miR-125a	Genome browser	Homo sapiens abhydrolase domain containing 6 (ABHD6), mRNA.
361	NM_002719	All miRNA predictions	2.74	hsa-miR-125a	Genome browser	Homo sapiens protein phosphatase 2, regulatory subunit B (B56), gamma isoform (PPP2R5C), transcript variant 1, mRNA.
362	NM_005985	All miRNA predictions	2.74	hsa-miR-125a	Genome browser	Homo sapiens snail homolog 1 (Drosophila) (SNAI1), mRNA.
363	NM_178586	All miRNA predictions	2.74	hsa-miR-125a	Genome browser	Homo sapiens protein phosphatase 2, regulatory subunit B (B56), gamma isoform (PPP2R5C), transcript variant 2, mRNA.
364	NM_001755	All miRNA predictions	2.74	hsa-miR-125a	Genome browser	Homo sapiens core-binding factor, beta subunit (CBFB), transcript variant 2, mRNA.
365	NM_032431	All miRNA predictions	2.73	hsa-miR-125a	Genome browser	Homo sapiens synovial apoptosis inhibitor 1, synoviolin (SYVN1), transcript variant 1, mRNA.
366	NM_172230	All miRNA predictions	2.73	hsa-miR-125a	Genome browser	Homo sapiens synovial apoptosis inhibitor 1, synoviolin (SYVN1), transcript variant 2, mRNA.
367	NM_032181	All miRNA predictions	2.72	hsa-miR-125a	Genome browser	Homo sapiens hypothetical protein FLJ13391 (FLJ13391), mRNA.
368	NM_014112	All miRNA predictions	2.72	hsa-miR-125a	Genome browser	Homo sapiens trichorhinophalangeal syndrome I (TRPS1), mRNA.
369	NM_007007	All miRNA predictions	2.71	hsa-miR-125a	Genome browser	Homo sapiens cleavage and polyadenylation specific factor 6, 68kDa (CPSF6), mRNA.
370	NM_002355	All miRNA predictions	2.70	hsa-miR-125a	Genome browser	Homo sapiens mannose-6-phosphate receptor (cation dependent) (M6PR), mRNA.
371	NM_198513	All miRNA predictions	2.69	hsa-miR-125a	Genome browser	Homo sapiens CGI-72 protein (CGI-72), transcript variant 3, mRNA.
372	NM_024878	All miRNA predictions	2.69	hsa-miR-125a	Genome browser	Homo sapiens CGI-72 protein (CGI-72), transcript variant 4, mRNA.
373	NM_182907	All miRNA predictions	2.68	hsa-miR-125a	Genome browser	Homo sapiens PR domain containing 1, with ZNF domain (PRDM1), transcript variant 2, mRNA.
374	NM_001001482	All miRNA predictions	2.68	hsa-miR-125a	Genome browser	Homo sapiens hypothetical protein FLJ11011 (FLJ11011), transcript variant 3, mRNA.
375	NM_001198	All miRNA predictions	2.68	hsa-miR-125a	Genome browser	Homo sapiens PR domain containing 1, with ZNF domain (PRDM1), transcript variant 1, mRNA.
376	NM_006378	All miRNA predictions	2.67	hsa-miR-125a	Genome browser	Homo sapiens sema domain, immunoglobulin domain (Ig), transmembrane domain (TM) and short cytoplasmic domain, (semaphorin) 4D (SEMA4D), mRNA.
377	NM_004438	All miRNA	2.67	hsa-miR-	Genome	Homo sapiens EPH receptor A4 (EPHA4), mRNA.

		predictions		125a	browser	
378	NM_015265	All miRNA predictions	2.66	hsa-miR-125a	Genome browser	Homo sapiens SATB family member 2 (SATB2), mRNA.
379	NM_017927	All miRNA predictions	2.66	hsa-miR-125a	Genome browser	Homo sapiens mitofusin 1 (MFN1), nuclear gene encoding mitochondrial protein, transcript variant 2, mRNA.
380	NM_005477	All miRNA predictions	2.65	hsa-miR-125a	Genome browser	Homo sapiens hyperpolarization activated cyclic nucleotide-gated potassium channel 4 (HCN4), mRNA.
381	NM_001095	All miRNA predictions	2.65	hsa-miR-125a	Genome browser	Homo sapiens amiloride-sensitive cation channel 2, neuronal (ACCN2), transcript variant 2, mRNA.
382	NM_033083	All miRNA predictions	2.65	hsa-miR-125a	Genome browser	Homo sapiens ELL associated factor 1 (EAF1), mRNA.
383	NM_020834	All miRNA predictions	2.65	hsa-miR-125a	Genome browser	Homo sapiens KIAA1443 (KIAA1443), mRNA.
384	NM_020039	All miRNA predictions	2.65	hsa-miR-125a	Genome browser	Homo sapiens amiloride-sensitive cation channel 2, neuronal (ACCN2), transcript variant 1, mRNA.
385	NM_020149	All miRNA predictions	2.64	hsa-miR-125a	Genome browser	Homo sapiens Meis1, myeloid ecotropic viral integration site 1 homolog 2 (mouse) (MEIS2), transcript variant e, mRNA.
386	NM_002310	All miRNA predictions	2.64	hsa-miR-125a	Genome browser	Homo sapiens leukemia inhibitory factor receptor (LIFR), mRNA.
387	NM_014761	All miRNA predictions	2.64	hsa-miR-125a	Genome browser	Homo sapiens KIAA0174 gene product (KIAA0174), mRNA.
388	NM_005099	All miRNA predictions	2.61	hsa-miR-125a	Genome browser	Homo sapiens a disintegrin-like and metalloprotease (reprolysin type) with thrombospondin type 1 motif, 4 (ADAMTS4), mRNA.
389	NM_207313	All miRNA predictions	2.61	hsa-miR-125a	Genome browser	Homo sapiens hypothetical protein LOC124842 (LOC124842), mRNA.
390	NM_001702	All miRNA predictions	2.60	hsa-miR-125a	Genome browser	Homo sapiens brain-specific angiogenesis inhibitor 1 (BAI1), mRNA.
391	NM_016531	All miRNA predictions	2.59	hsa-miR-125a	Genome browser	Homo sapiens Kruppel-like factor 3 (basic) (KLF3), mRNA.
392	NM_006464	All miRNA predictions	2.59	hsa-miR-125a	Genome browser	Homo sapiens trans-golgi network protein 2 (TGOLN2), mRNA.
393	NM_005911	All miRNA predictions	2.59	hsa-miR-125a	Genome browser	Homo sapiens methionine adenosyltransferase II, alpha (MAT2A), mRNA.
394	NM_022822	All miRNA predictions	2.59	hsa-miR-125a	Genome browser	Homo sapiens likely ortholog of kinesin light chain 2 (KLC2), mRNA.
395	NM_002703	All miRNA predictions	2.58	hsa-miR-125a	Genome browser	Homo sapiens phosphoribosyl pyrophosphate amidotransferase (PPAT), mRNA.
396	NM_153020	All miRNA predictions	2.57	hsa-miR-125a	Genome browser	Homo sapiens RNA binding motif protein 24 (RBM24), mRNA.
397	NM_017789	All miRNA predictions	2.57	hsa-miR-125a	Genome browser	Homo sapiens sema domain, immunoglobulin domain (Ig), transmembrane domain (TM) and short cytoplasmic domain, (semaphorin) 4C (SEMA4C), mRNA.
398	NM_018650	All miRNA predictions	2.56	hsa-miR-125a	Genome browser	Homo sapiens MAP/microtubule affinity-regulating kinase 1 (MARK1), mRNA.
399	NM_002953	All miRNA predictions	2.56	hsa-miR-125a	Genome browser	Homo sapiens ribosomal protein S6 kinase, 90kDa, polypeptide 1 (RPS6KA1), transcript variant 1, mRNA.
400	NM_023029	All miRNA predictions	2.55	hsa-miR-125a	Genome browser	Homo sapiens fibroblast growth factor receptor 2 (bacteria-expressed kinase, keratinocyte growth factor receptor, craniofacial dysostosis 1, Crouzon syndrome, Pfeiffer syndrome, Jackson-Weiss syndrome) (FGFR2), transcript variant 11, mRNA.
401	NM_006091	All miRNA predictions	2.55	hsa-miR-125a	Genome browser	Homo sapiens coronin, actin binding protein, 2B (CORO2B), mRNA.
402	NM_023031	All miRNA predictions	2.55	hsa-miR-125a	Genome browser	Homo sapiens fibroblast growth factor receptor 2 (bacteria-expressed kinase, keratinocyte growth factor receptor, craniofacial dysostosis 1, Crouzon syndrome, Pfeiffer syndrome, Jackson-Weiss syndrome) (FGFR2), transcript variant 13, mRNA.
403	NM_000141	All miRNA predictions	2.55	hsa-miR-125a	Genome browser	Homo sapiens fibroblast growth factor receptor 2 (bacteria-expressed kinase, keratinocyte growth factor receptor, craniofacial dysostosis 1, Crouzon

						syndrome, Pfeiffer syndrome, Jackson-Weiss syndrome) (FGFR2), transcript variant 1, mRNA.
404	NM_023030	All miRNA predictions	2.55	hsa-miR-125a	Genome browser	Homo sapiens fibroblast growth factor receptor 2 (bacteria-expressed kinase, keratinocyte growth factor receptor, craniofacial dysostosis 1, Crouzon syndrome, Pfeiffer syndrome, Jackson-Weiss syndrome) (FGFR2), transcript variant 12, mRNA.
405	NM_022975	All miRNA predictions	2.55	hsa-miR-125a	Genome browser	Homo sapiens fibroblast growth factor receptor 2 (bacteria-expressed kinase, keratinocyte growth factor receptor, craniofacial dysostosis 1, Crouzon syndrome, Pfeiffer syndrome, Jackson-Weiss syndrome) (FGFR2), transcript variant 8, mRNA.
406	NM_022969	All miRNA predictions	2.55	hsa-miR-125a	Genome browser	Homo sapiens fibroblast growth factor receptor 2 (bacteria-expressed kinase, keratinocyte growth factor receptor, craniofacial dysostosis 1, Crouzon syndrome, Pfeiffer syndrome, Jackson-Weiss syndrome) (FGFR2), transcript variant 2, mRNA.
407	NM_022970	All miRNA predictions	2.55	hsa-miR-125a	Genome browser	Homo sapiens fibroblast growth factor receptor 2 (bacteria-expressed kinase, keratinocyte growth factor receptor, craniofacial dysostosis 1, Crouzon syndrome, Pfeiffer syndrome, Jackson-Weiss syndrome) (FGFR2), transcript variant 3, mRNA.
408	NM_022972	All miRNA predictions	2.55	hsa-miR-125a	Genome browser	Homo sapiens fibroblast growth factor receptor 2 (bacteria-expressed kinase, keratinocyte growth factor receptor, craniofacial dysostosis 1, Crouzon syndrome, Pfeiffer syndrome, Jackson-Weiss syndrome) (FGFR2), transcript variant 5, mRNA.
409	NM_000165	All miRNA predictions	2.53	hsa-miR-125a	Genome browser	Homo sapiens gap junction protein, alpha 1, 43kDa (connexin 43) (GJA1), mRNA.
410	NM_032982	All miRNA predictions	2.52	hsa-miR-125a	Genome browser	Homo sapiens caspase 2, apoptosis-related cysteine protease (neural precursor cell expressed, developmentally down-regulated 2) (CASP2), transcript variant 1, mRNA.
411	NM_058172	All miRNA predictions	2.52	hsa-miR-125a	Genome browser	Homo sapiens anthrax toxin receptor 2 (ANTXR2), mRNA.
412	NM_018963	All miRNA predictions	2.50	hsa-miR-125a	Genome browser	Homo sapiens WD repeat domain 9 (WDR9), transcript variant 1, mRNA.
413	NM_020390	All miRNA predictions	2.50	hsa-miR-125a	Genome browser	Homo sapiens eukaryotic translation initiation factor 5A2 (EIF5A2), mRNA.
414	NM_001949	All miRNA predictions	2.50	hsa-miR-125a	Genome browser	Homo sapiens E2F transcription factor 3 (E2F3), mRNA.
415	NM_005094	All miRNA predictions	2.50	hsa-miR-125a	Genome browser	Homo sapiens solute carrier family 27 (fatty acid transporter), member 4 (SLC27A4), mRNA.
416	NM_023028	All miRNA predictions	2.48	hsa-miR-125a	Genome browser	Homo sapiens fibroblast growth factor receptor 2 (bacteria-expressed kinase, keratinocyte growth factor receptor, craniofacial dysostosis 1, Crouzon syndrome, Pfeiffer syndrome, Jackson-Weiss syndrome) (FGFR2), transcript variant 10, mRNA.
417	NM_206855	All miRNA predictions	2.48	hsa-miR-125a	Genome browser	Homo sapiens quaking homolog, KH domain RNA binding (mouse) (QKI), transcript variant 4, mRNA.
418	NM_001773	All miRNA predictions	2.47	hsa-miR-125a	Genome browser	Homo sapiens CD34 antigen (CD34), mRNA.
419	NM_024954	All miRNA predictions	2.46	hsa-miR-125a	Genome browser	Homo sapiens ubiquitin domain containing 1 (UBTD1), mRNA.
420	NM_080862	All miRNA predictions	2.46	hsa-miR-125a	Genome browser	Homo sapiens SPRY domain-containing SOCS box protein SSB-4 (SSB4), mRNA.
421	NM_014006	All miRNA predictions	2.44	hsa-miR-125a	Genome browser	Homo sapiens PI-3-kinase-related kinase SMG-1 (SMG1), transcript variant 2, mRNA.
422	NM_014912	All miRNA predictions	2.44	hsa-miR-125a	Genome browser	Homo sapiens cytoplasmic polyadenylation element binding protein 3 (CPEB3), mRNA.
423	NM_015092	All miRNA predictions	2.44	hsa-miR-125a	Genome browser	Homo sapiens PI-3-kinase-related kinase SMG-1 (SMG1), transcript variant 1, mRNA.
424	NM_014682	All miRNA predictions	2.43	hsa-miR-125a	Genome browser	Homo sapiens suppression of tumorigenicity 18 (breast carcinoma) (zinc finger protein) (ST18), mRNA.
425	NM_153043	All miRNA predictions	2.43	hsa-miR-125a	Genome browser	Homo sapiens hypothetical protein FLJ37078 (FLJ37078), mRNA.
426	NM_022730	All miRNA predictions	2.43	hsa-miR-125a	Genome browser	Homo sapiens COP9 constitutive photomorphogenic homolog subunit 7B (Arabidopsis) (COPS7B), mRNA.

427	NM_025154	All miRNA predictions	2.42	hsa-miR-125a	Genome browser	Homo sapiens unc-84 homolog A (C. elegans) (UNC84A), mRNA.
428	NM_012294	All miRNA predictions	2.42	hsa-miR-125a	Genome browser	Homo sapiens Rap guanine nucleotide exchange factor (GEF) 5 (RAPGEF5), mRNA.
429	NM_003173	All miRNA predictions	2.42	hsa-miR-125a	Genome browser	Homo sapiens suppressor of variegation 3-9 homolog 1 (Drosophila) (SUV39H1), mRNA.
430	NM_014409	All miRNA predictions	2.42	hsa-miR-125a	Genome browser	Homo sapiens TAF5-like RNA polymerase II, p300/CBP-associated factor (PCAF)-associated factor, 65kDa (TAF5L), mRNA.
431	NM_173822	All miRNA predictions	2.42	hsa-miR-125a	Genome browser	Homo sapiens hypothetical protein MGC39518 (MGC39518), mRNA.
432	NM_006403	All miRNA predictions	2.40	hsa-miR-125a	Genome browser	Homo sapiens neural precursor cell expressed, developmentally down-regulated 9 (NEDD9), mRNA.
433	NM_001408	All miRNA predictions	2.40	hsa-miR-125a	Genome browser	Homo sapiens cadherin, EGF LAG seven-pass G-type receptor 2 (flamingo homolog, Drosophila) (CELSR2), mRNA.
434	NM_001001924	All miRNA predictions	2.38	hsa-miR-125a	Genome browser	Homo sapiens mitochondrial tumor suppressor 1 (MTUS1), nuclear gene encoding mitochondrial protein, transcript variant 1, mRNA.
435	NM_144599	All miRNA predictions	2.38	hsa-miR-125a	Genome browser	Homo sapiens non imprinted in Prader-Willi/Angelman syndrome 1 (NIPA1), mRNA.
436	NM_001001931	All miRNA predictions	2.38	hsa-miR-125a	Genome browser	Homo sapiens mitochondrial tumor suppressor 1 (MTUS1), nuclear gene encoding mitochondrial protein, transcript variant 4, mRNA.
437	NM_001001925	All miRNA predictions	2.38	hsa-miR-125a	Genome browser	Homo sapiens mitochondrial tumor suppressor 1 (MTUS1), nuclear gene encoding mitochondrial protein, transcript variant 2, mRNA.
438	NM_020749	All miRNA predictions	2.38	hsa-miR-125a	Genome browser	Homo sapiens mitochondrial tumor suppressor 1 (MTUS1), nuclear gene encoding mitochondrial protein, transcript variant 5, mRNA.
439	NM_001879	All miRNA predictions	2.37	hsa-miR-125a	Genome browser	Homo sapiens mannan-binding lectin serine protease 1 (C4/C2 activating component of Ra-reactive factor) (MASP1), transcript variant 1, mRNA.
440	NM_145818	All miRNA predictions	2.37	hsa-miR-125a	Genome browser	Homo sapiens component of oligomeric golgi complex 4 (COG4), transcript variant Cog4S, mRNA.
441	NM_012266	All miRNA predictions	2.37	hsa-miR-125a	Genome browser	Homo sapiens DnaJ (Hsp40) homolog, subfamily B, member 5 (DNAJB5), mRNA.
442	NM_017554	All miRNA predictions	2.36	hsa-miR-125a	Genome browser	Homo sapiens poly (ADP-ribose) polymerase family, member 14 (PARP14), mRNA.
443	NM_014729	All miRNA predictions	2.35	hsa-miR-125a	Genome browser	Homo sapiens thymus high mobility group box protein TOX (TOX), mRNA.
444	NM_018375	All miRNA predictions	2.35	hsa-miR-125a	Genome browser	Homo sapiens solute carrier family 39 (zinc transporter), member 9 (SLC39A9), mRNA.
445	NM_206835	All miRNA predictions	2.34	hsa-miR-125a	Genome browser	Homo sapiens TNF receptor-associated factor 7 (TRAF7), transcript variant 2, mRNA.
446	NM_007359	All miRNA predictions	2.33	hsa-miR-125a	Genome browser	Homo sapiens cancer susceptibility candidate 3 (CASC3), mRNA.
447	NM_000723	All miRNA predictions	2.33	hsa-miR-125a	Genome browser	Homo sapiens calcium channel, voltage-dependent, beta 1 subunit (CACNB1), transcript variant 1, mRNA.
448	NM_019074	All miRNA predictions	2.32	hsa-miR-125a	Genome browser	Homo sapiens delta-like 4 (Drosophila) (DLL4), mRNA.
449	NM_198549	All miRNA predictions	2.32	hsa-miR-125a	Genome browser	Homo sapiens FLJ35093 protein (FLJ35093), mRNA.
450	NM_194294	All miRNA predictions	2.31	hsa-miR-125a	Genome browser	Homo sapiens hypothetical protein LOC169355 (LOC169355), mRNA.
451	NM_017486	All miRNA predictions	2.31	hsa-miR-125a	Genome browser	Homo sapiens adducin 2 (beta) (ADD2), transcript variant beta-6a, mRNA.
452	NM_001617	All miRNA predictions	2.31	hsa-miR-125a	Genome browser	Homo sapiens adducin 2 (beta) (ADD2), transcript variant beta-1, mRNA.
453	NM_017483	All miRNA predictions	2.31	hsa-miR-125a	Genome browser	Homo sapiens adducin 2 (beta) (ADD2), transcript variant beta-3a, mRNA.
454	NM_017485	All miRNA predictions	2.31	hsa-miR-125a	Genome browser	Homo sapiens adducin 2 (beta) (ADD2), transcript variant beta-5a, mRNA.
455	NM_030927	All miRNA predictions	2.29	hsa-miR-125a	Genome browser	Homo sapiens transmembrane 4 superfamily member 14 (TM4SF14), mRNA.
456	NM_018330	All miRNA	2.27	hsa-miR-	Genome	Homo sapiens KIAA1598 (KIAA1598), mRNA.

		predictions		125a	browser	
457	NM_138494	All miRNA predictions	2.26	hsa-miR-125a	Genome browser	Homo sapiens vav-1 interacting Kruppel-like protein (VIK), transcript variant 1, mRNA.
458	NM_206902	All miRNA predictions	2.23	hsa-miR-125a	Genome browser	Homo sapiens reticulon 2 (RTN2), transcript variant 4, mRNA.
459	NM_003155	All miRNA predictions	2.22	hsa-miR-125a	Genome browser	Homo sapiens stanniocalcin 1 (STC1), mRNA.
460	NM_032827	All miRNA predictions	2.22	hsa-miR-125a	Genome browser	Homo sapiens atonal homolog 8 (Drosophila) (ATOH8), mRNA.
461	NM_018992	All miRNA predictions	2.21	hsa-miR-125a	Genome browser	Homo sapiens potassium channel tetramerisation domain containing 5 (KCTD5), mRNA.
462	NM_022763	All miRNA predictions	2.21	hsa-miR-125a	Genome browser	Homo sapiens factor for adipocyte differentiation 104 (FAD104), mRNA.
463	NM_004907	All miRNA predictions	2.20	hsa-miR-125a	Genome browser	Homo sapiens immediate early response 2 (IER2), mRNA.
464	NM_152383	All miRNA predictions	2.18	hsa-miR-125a	Genome browser	Homo sapiens hypothetical protein MGC42174 (MGC42174), mRNA.
465	NM_006275	All miRNA predictions	2.18	hsa-miR-125a	Genome browser	Homo sapiens splicing factor, arginine/serine-rich 6 (SFRS6), mRNA.
466	NM_018320	All miRNA predictions	2.16	hsa-miR-125a	Genome browser	Homo sapiens ring finger protein 121 (RNF121), transcript variant 1, mRNA.
467	NM_194453	All miRNA predictions	2.16	hsa-miR-125a	Genome browser	Homo sapiens ring finger protein 121 (RNF121), transcript variant 3, mRNA.
468	NM_194452	All miRNA predictions	2.16	hsa-miR-125a	Genome browser	Homo sapiens ring finger protein 121 (RNF121), transcript variant 2, mRNA.
469	NM_015192	All miRNA predictions	2.16	hsa-miR-125a	Genome browser	Homo sapiens phospholipase C, beta 1 (phosphoinositide-specific) (PLCB1), transcript variant 1, mRNA.
470	NM_182487	All miRNA predictions	2.15	hsa-miR-125a	Genome browser	Homo sapiens olfactomedin-like 2A (OLFML2A), mRNA.
471	NM_004535	All miRNA predictions	2.14	hsa-miR-125a	Genome browser	Homo sapiens myelin transcription factor 1 (MYT1), mRNA.
472	NM_001002909	All miRNA predictions	2.14	hsa-miR-125a	Genome browser	Homo sapiens KIAA0553 protein (KIAA0553), mRNA.
473	NM_033018	All miRNA predictions	2.13	hsa-miR-125a	Genome browser	Homo sapiens PCTAIRE protein kinase 1 (PCTK1), transcript variant 2, mRNA.
474	NM_021643	All miRNA predictions	2.13	hsa-miR-125a	Genome browser	Homo sapiens tribbles homolog 2 (Drosophila) (TRIB2), mRNA.
475	NM_006312	All miRNA predictions	2.13	hsa-miR-125a	Genome browser	Homo sapiens nuclear receptor co-repressor 2 (NCOR2), mRNA.
476	NM_033019	All miRNA predictions	2.13	hsa-miR-125a	Genome browser	n/a
477	NM_006201	All miRNA predictions	2.13	hsa-miR-125a	Genome browser	Homo sapiens PCTAIRE protein kinase 1 (PCTK1), transcript variant 1, mRNA.
478	NM_001224	All miRNA predictions	2.12	hsa-miR-125a	Genome browser	Homo sapiens caspase 2, apoptosis-related cysteine protease (neural precursor cell expressed, developmentally down-regulated 2) (CASP2), transcript variant 2, mRNA.
479	NM_003483	All miRNA predictions	2.09	hsa-miR-125a	Genome browser	Homo sapiens high mobility group AT-hook 2 (HMGA2), mRNA.
480	NM_005465	All miRNA predictions	2.09	hsa-miR-125a	Genome browser	Homo sapiens v-akt murine thymoma viral oncogene homolog 3 (protein kinase B, gamma) (AKT3), transcript variant 1, mRNA.
481	NM_016605	All miRNA predictions	2.08	hsa-miR-125a	Genome browser	Homo sapiens chromosome 5 open reading frame 6 (C5orf6), mRNA.
482	NM_004386	All miRNA predictions	2.07	hsa-miR-125a	Genome browser	Homo sapiens chondroitin sulfate proteoglycan 3 (neurocan) (CSPG3), mRNA.
483	NM_021116	All miRNA predictions	2.07	hsa-miR-125a	Genome browser	Homo sapiens adenylate cyclase 1 (brain) (ADCY1), mRNA.
484	NM_003347	All miRNA predictions	2.06	hsa-miR-125a	Genome browser	Homo sapiens ubiquitin-conjugating enzyme E2L 3 (UBE2L3), transcript variant 1, mRNA.
485	NM_003489	All miRNA predictions	2.05	hsa-miR-125a	Genome browser	Homo sapiens nuclear receptor interacting protein 1 (NRIP1), mRNA.
486	NM_004427	All miRNA predictions	2.03	hsa-miR-125a	Genome browser	Homo sapiens polyhomeotic-like 2 (Drosophila) (PHC2), transcript variant 2, mRNA.

487	NM_198040	All miRNA predictions	2.03	hsa-miR-125a	Genome browser	Homo sapiens polyhomeotic-like 2 (Drosophila) (PHC2), transcript variant 1, mRNA.
488	NM_033285	All miRNA predictions	2.01	hsa-miR-125a	Genome browser	Homo sapiens tumor protein p53 inducible nuclear protein 1 (TP53INP1), mRNA.
489	NM_025054	All miRNA predictions	2.00	hsa-miR-125a	Genome browser	Homo sapiens valosin-containing protein (p97)/p47 complex-interacting protein p135 (VCIP135), mRNA.
490	NM_003216	All miRNA predictions	2.00	hsa-miR-125a	Genome browser	Homo sapiens thyrotrophic embryonic factor (TEF), mRNA.
491	NM_017488	All miRNA predictions	1.99	hsa-miR-125a	Genome browser	Homo sapiens adducin 2 (beta) (ADD2), transcript variant beta-4, mRNA.
492	NM_177438	All miRNA predictions	1.99	hsa-miR-125a	Genome browser	Homo sapiens Dicer1, Dcr-1 homolog (Drosophila) (DICER1), transcript variant 1, mRNA.
493	NM_145166	All miRNA predictions	1.99	hsa-miR-125a	Genome browser	Homo sapiens hypothetical protein KIAA1190 (KIAA1190), mRNA.
494	NM_030621	All miRNA predictions	1.99	hsa-miR-125a	Genome browser	Homo sapiens Dicer1, Dcr-1 homolog (Drosophila) (DICER1), transcript variant 2, mRNA.
495	NM_182734	All miRNA predictions	1.97	hsa-miR-125a	Genome browser	Homo sapiens phospholipase C, beta 1 (phosphoinositide-specific) (PLCB1), transcript variant 2, mRNA.
496	NM_198157	All miRNA predictions	1.95	hsa-miR-125a	Genome browser	Homo sapiens ubiquitin-conjugating enzyme E2L 3 (UBE2L3), transcript variant 2, mRNA.
497	NM_183376	All miRNA predictions	1.94	hsa-miR-125a	Genome browser	Homo sapiens arrestin domain containing 4 (ARRDC4), mRNA.
498	NM_017873	All miRNA predictions	1.93	hsa-miR-125a	Genome browser	Homo sapiens ankyrin repeat and SOCS box-containing 6 (ASB6), transcript variant 1, mRNA.
499	NM_206853	All miRNA predictions	1.92	hsa-miR-125a	Genome browser	Homo sapiens quaking homolog, KH domain RNA binding (mouse) (QKI), transcript variant 2, mRNA.
500	NM_005920	All miRNA predictions	1.91	hsa-miR-125a	Genome browser	Homo sapiens MADS box transcription enhancer factor 2, polypeptide D (myocyte enhancer factor 2D) (MEF2D), mRNA.
501	NM_182763	All miRNA predictions	1.90	hsa-miR-125a	Genome browser	Homo sapiens myeloid cell leukemia sequence 1 (BCL2-related) (MCL1), transcript variant 2, mRNA.
502	NM_030627	All miRNA predictions	1.90	hsa-miR-125a	Genome browser	Homo sapiens cytoplasmic polyadenylation element binding protein 4 (CPEB4), mRNA.
503	NM_021960	All miRNA predictions	1.90	hsa-miR-125a	Genome browser	Homo sapiens myeloid cell leukemia sequence 1 (BCL2-related) (MCL1), transcript variant 1, mRNA.
504	NM_002840	All miRNA predictions	1.89	hsa-miR-125a	Genome browser	Homo sapiens protein tyrosine phosphatase, receptor type, F (PTPRF), transcript variant 1, mRNA.
505	NM_130440	All miRNA predictions	1.89	hsa-miR-125a	Genome browser	Homo sapiens protein tyrosine phosphatase, receptor type, F (PTPRF), transcript variant 2, mRNA.
506	NM_032984	All miRNA predictions	1.88	hsa-miR-125a	Genome browser	Homo sapiens caspase 2, apoptosis-related cysteine protease (neural precursor cell expressed, developmentally down-regulated 2) (CASP2), transcript variant 4, mRNA.
507	NM_032983	All miRNA predictions	1.88	hsa-miR-125a	Genome browser	Homo sapiens caspase 2, apoptosis-related cysteine protease (neural precursor cell expressed, developmentally down-regulated 2) (CASP2), transcript variant 3, mRNA.
508	NM_005197	All miRNA predictions	1.85	hsa-miR-125a	Genome browser	Homo sapiens checkpoint suppressor 1 (CHES1), mRNA.
509	NM_138473	All miRNA predictions	1.84	hsa-miR-125a	Genome browser	Homo sapiens Sp1 transcription factor (SP1), mRNA.
510	NM_019084	All miRNA predictions	1.83	hsa-miR-125a	Genome browser	Homo sapiens cyclin J (CCNJ), mRNA.
511	NM_013449	All miRNA predictions	1.83	hsa-miR-125a	Genome browser	Homo sapiens bromodomain adjacent to zinc finger domain, 2A (BAZ2A), mRNA.
512	NM_001438	All miRNA predictions	1.83	hsa-miR-125a	Genome browser	Homo sapiens estrogen-related receptor gamma (ESRRG), transcript variant 1, mRNA.
513	NM_206595	All miRNA predictions	1.83	hsa-miR-125a	Genome browser	Homo sapiens estrogen-related receptor gamma (ESRRG), transcript variant 3, mRNA.
514	NM_206594	All miRNA predictions	1.83	hsa-miR-125a	Genome browser	Homo sapiens estrogen-related receptor gamma (ESRRG), transcript variant 2, mRNA.
515	NM_003045	All miRNA predictions	1.82	hsa-miR-125a	Genome browser	Homo sapiens solute carrier family 7 (cationic amino acid transporter, y+ system), member 1 (SLC7A1), mRNA.
516	NM_015278	All miRNA	1.81	hsa-miR-	Genome	Homo sapiens SAM and SH3 domain containing 1

		predictions		125a	browser	(SASH1), mRNA.
517	NM_030915	All miRNA predictions	1.80	hsa-miR-125a	Genome browser	Homo sapiens likely ortholog of mouse limb-bud and heart gene (LBH), mRNA.
518	NM_005475	All miRNA predictions	1.76	hsa-miR-125a	Genome browser	Homo sapiens lymphocyte adaptor protein (LNK), mRNA.
519	NM_004393	All miRNA predictions	1.74	hsa-miR-125a	Genome browser	Homo sapiens dystroglycan 1 (dystrophin-associated glycoprotein 1) (DAG1), mRNA.
520	NM_005668	All miRNA predictions	1.73	hsa-miR-125a	Genome browser	Homo sapiens sialyltransferase 8D (alpha-2, 8-polysialyltransferase) (SIAT8D), transcript variant 1, mRNA.
521	NM_177999	All miRNA predictions	1.69	hsa-miR-125a	Genome browser	Homo sapiens ankyrin repeat and SOCS box-containing 6 (ASB6), transcript variant 2, mRNA.
522	NM_133333	All miRNA predictions	1.67	hsa-miR-125a	Genome browser	Homo sapiens Wolf-Hirschhorn syndrome candidate 1 (WHSC1), transcript variant 6, mRNA.
523	NM_000430	All miRNA predictions	1.67	hsa-miR-125a	Genome browser	Homo sapiens platelet-activating factor acetylhydrolase, isoform Ib, alpha subunit 45kDa (PAFAH1B1), mRNA.
524	NM_133332	All miRNA predictions	1.66	hsa-miR-125a	Genome browser	Homo sapiens Wolf-Hirschhorn syndrome candidate 1 (WHSC1), transcript variant 5, mRNA.
525	NM_206854	All miRNA predictions	1.64	hsa-miR-125a	Genome browser	Homo sapiens quaking homolog, KH domain RNA binding (mouse) (QKI), transcript variant 3, mRNA.
526	NM_021255	All miRNA predictions	1.64	hsa-miR-125a	Genome browser	Homo sapiens pellino homolog 2 (Drosophila) (PELI2), mRNA.
527	NM_014746	All miRNA predictions	1.40	hsa-miR-125a	Genome browser	Homo sapiens ring finger protein 144 (RNF144), mRNA.
528	NM_005065	All miRNA predictions	1.39	hsa-miR-125a	Genome browser	Homo sapiens sel-1 suppressor of lin-12-like (C. elegans) (SEL1L), mRNA.
529	NM_194278	All miRNA predictions	1.32	hsa-miR-125a	Genome browser	Homo sapiens chromosome 14 open reading frame 43 (C14orf43), mRNA.
530	NM_000793	All miRNA predictions	1.19	hsa-miR-125a	Genome browser	Homo sapiens deiodinase, iodothyronine, type II (DIO2), transcript variant 2, mRNA.
531	NM_013989	All miRNA predictions	1.19	hsa-miR-125a	Genome browser	Homo sapiens deiodinase, iodothyronine, type II (DIO2), transcript variant 1, mRNA.