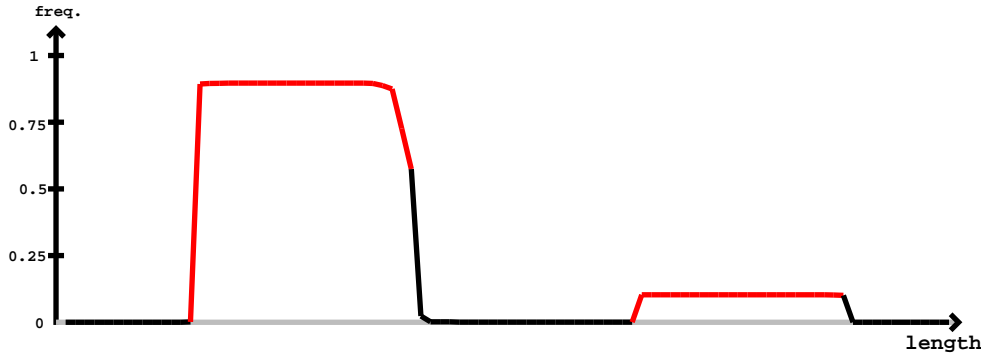
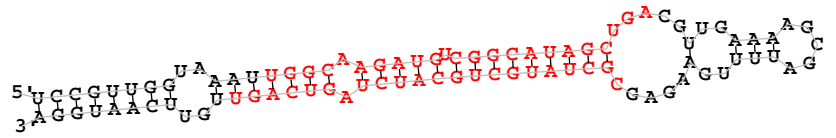


miRBase precursor : dme-mir-31a_MI0000420
 Total read count : 2398
 dme-miR-31a-5p_MIMAT000400 read count
 dme-miR-31a-3p_MIMAT0020834 read count
 remaining reads : 1



dme-miR-31a-5p_MIMAT000400

dme-miR-31a-3p_MIMAT0020834

5' -	reads	mm	exp	sample
uccguugguaaaauuggcaagaugucggcauagcugacguugaaaagcgauuuugaagagcgcuauugcugcaucauagcaguuguucaaugga	1	1		seq
.....Auggcaagaugucggcauagcu.....	1	1		seq
.....Auggcaagaugucggcauagcug.....	1	1		seq
.....Auggcaagaugucggcauagcuga.....	1	1		seq
.....uggcaagaugAcggcaua.....	1	1		seq
.....uggcaagauAucggcaua.....	1	1		seq
.....uggcaagaugucggcaua.....	1	0		seq
.....uggcaagaugucggcauag.....	19	0		seq
.....uggUaagaugucggcauagc.....	1	1		seq
.....uggcaagaugucggcauagU.....	1	1		seq
.....uggcaagaugucggcauagc.....	28	0		seq
.....uggcaagaugucggcauagA.....	1	1		seq
.....uggcaagaugucggcaAagcu.....	4	1		seq
.....uggcaagaugucggcauaAcu.....	2	1		seq
.....uggcaagaugucggcauagcu.....	304	0		seq
.....uggcaagaugucggcauagcG.....	1	1		seq
.....Cggcaagaugucggcauagcu.....	1	1		seq
.....uggcaagaugucCgcauagcu.....	1	1		seq
.....uggcaagauAucggcauagcu.....	1	1		seq
.....Aggcaagaugucggcauagcu.....	2	1		seq
.....uggcaaUaugucggcauagcu.....	1	1		seq
.....uggcaagaugucggUauagcu.....	3	1		seq
.....uggcaaAaugucggcauagcu.....	1	1		seq
.....uggcaagaugucggcauagcA.....	1	1		seq
.....uggcaagaugucggcaGagcu.....	1	1		seq
.....uggcaCgaugucggcauagcu.....	1	1		seq
.....uggcaagaugucggcauagGu.....	2	1		seq
.....uggcaagaugAcggcauagcu.....	2	1		seq
.....uggcaagaugcGAcuagcu.....	1	1		seq
.....uggcaagaugucggcauUgcu.....	4	1		seq
.....uggcaagaugucggcaCagcu.....	4	1		seq
.....uggcaagGugucggcauagcu.....	1	1		seq
.....uggcaagauguAaggcauagcu.....	1	1		seq
.....uggcaagaugucggcauaCcu.....	2	1		seq
.....uggcaagaugucGUcauagcu.....	2	1		seq

uccguugguaaaauuggcaagaugucggcauagcugacguugaaaagcgauuuugaagagcgcuauugcugcaucuaugucaguuguucaaugga

.....uggcaagaugucggcauagcugaG.....	3	1	seq
.....uggcaagaugucggcauagcugac.....	1	0	seq
.....uggcaagaugucggcauagcugacguu.....	3	0	seq
.....Ugcaagaugucggcauagcuga.....	3	1	seq
.....ggcaagaugucggcauagcuga.....	2	0	seq
.....gcaagaugucggcauagcu.....	1	0	seq
.....caagaugucggcauagcuga.....	2	0	seq
.....cguugaaaagcgauuuugaagag.....	1	0	seq
.....cgcuauugcugcaucuaugAca.....	1	1	seq
.....cgcuauugcugcaucuauguca.....	1	0	seq
.....cgcuauugcugcaucuaugucag.....	3	0	seq
.....cgcuAGcugcaucuaugucagu.....	2	1	seq
.....cgcuauugcugcaucuaugucagC.....	1	1	seq
.....Agcuauugcugcaucuaugucagu.....	38	1	seq
.....cgcuauugcugcaucuaugucagu.....	190	0	seq
.....cgcuauugcugcaucuaugucGgu.....	1	1	seq
.....cgcuauugcuUcaucuaugucagu.....	1	1	seq
.....cgcuauugcugcaucuCgucagu.....	1	1	seq
.....cgcuACgucgcaucuaugucagu.....	2	1	seq
.....cgcuauugcugcaucuaLucagu.....	1	1	seq
.....cgcuauugcugcaCcuagucagu.....	1	1	seq
.....cgcuauugcugcaucuaugucCgu.....	1	1	seq
.....cgcuAGucgcaucuaugucagu.....	1	1	seq
.....cgcuauugcugcaucuaUucagu.....	2	1	seq