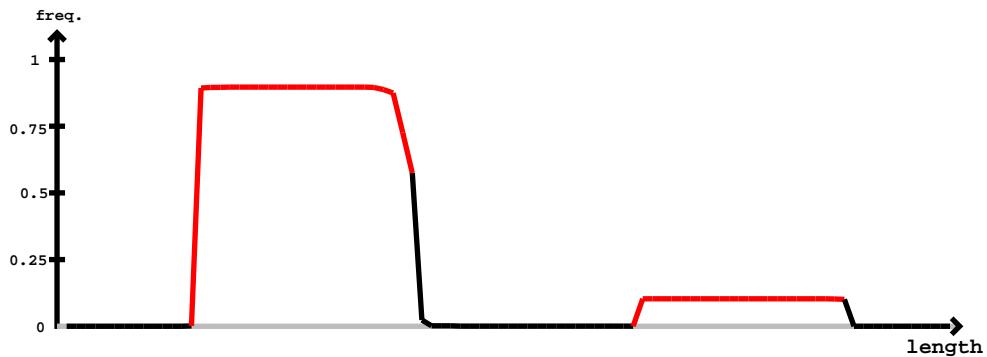


miRBase precursor : dme-mir-31a_MI0000420
Total read count : 2398
dme-miR-31a-5p_MIMAT0000500 read count
dme-miR-31a-3p_MIMAT0020834 read count
remaining reads : 1



dme-miR-31a-5p_MIMAT0000400

dme-miR-31a-3p_MIMAT0020834

uccguugguaauuggcaagaaugucggcauagcuga~~cguugaaaaagcgauuuuugaagag~~cgcuaugcugcaucuagucaguuguucaaugga

.....uggcaagaaugucggcUuagcu.....	1	1	seq
.....uggcaagaugucAgcuaugcu.....	5	1	seq
.....uggcaagaugucggcauCgcu.....	1	1	seq
.....uggcaagaaugCcgccauagcu.....	2	1	seq
.....uggcaagaaugucgcCcauagcu.....	1	1	seq
.....uggcaagaaugucggcauagAug.....	1	1	seq
.....uggcaagaugucgcUcauagcug.....	2	1	seq
.....ugUcaagaugucggcauagcug.....	1	1	seq
.....Cggcaagaugucggcauagcug.....	2	1	seq
.....uggcaagaugucggcauagcua.....	2	1	seq
.....Aggcaagaugucggcauagcug.....	7	1	seq
.....uggcaagaugucggcauagcug.....	321	0	seq
.....uggcaagaugucggcauUgcug.....	1	1	seq
.....uggcaagaugucggGauagcug.....	2	1	seq
.....uggcaagaugucgcAcauagcug.....	2	1	seq
.....uggcaagaugucgcAcauagcug.....	5	1	seq
.....uggcaaaaugucggcauagcug.....	1	1	seq
.....uggcaagaugucggcauagcCg.....	1	1	seq
.....uggcaagaugucggcaCagcug.....	1	1	seq
.....uggcaagaugucggcauagUug.....	1	1	seq
.....uggcaagaugucggcauagcuU.....	9	1	seq
.....uggcaagaugucggcaAagcug.....	3	1	seq
.....uggcaGgaugucggcauagcug.....	1	1	seq
.....uggcaagaugAcggcauagcug.....	1	1	seq
.....uggcaagaugucggcaCgcuga.....	5	1	seq
.....uggcaagaugucggcauUcuga.....	2	1	seq
.....uggcaagaugucggcauagcAga.....	4	1	seq
.....uggcaagaugucggcauagcuga.....	1125	0	seq
.....uggcaagGugucggcauagcuga.....	1	1	seq
.....uggcaagaugucggcauagcuA.....	5	1	seq
.....uggcaagaugucCcgaauagcuga.....	2	1	seq
.....uggcaagaugucggcauagcugU.....	21	1	seq
.....uggcaagaugCggcauagcuga.....	2	1	seq
.....uggcaagaugucggcaCagcuga.....	3	1	seq
.....uggcaagaAguucggcauagcuga.....	1	1	seq
.....uggcaagaugucgcAcauagcuga.....	1	1	seq
.....uggcaagaugucggcauagcUa.....	2	1	seq
.....Aggcaagaugucggcauagcuga.....	58	1	seq
.....uggcaagauUucggcauagcuga.....	6	1	seq
.....uggcaagaugucCcuaagcuga.....	1	1	seq
.....uggcaagaugucggcUuagcuga.....	2	1	seq
.....uggcGagaugucggcauagcuga.....	1	1	seq
.....uggcaagaCgucggcauagcuga.....	2	1	seq
.....uggcaagaugucggcauAcuga.....	2	1	seq
.....uggcaagaugucUgcuaagcuga.....	1	1	seq
.....uggcaagaugucggGauagcuga.....	1	1	seq
.....uCgcagaugucggcauagcuga.....	1	1	seq
.....uggcaagaugucggcauagAuga.....	1	1	seq
.....uggcaagaugucggCuaagcuga.....	5	1	seq
.....uggcaagaugAcggcauagcuga.....	3	1	seq
.....uggcaagaugucggcaAagcuga.....	1	1	seq
.....Cggcaagaugucggcauagcuga.....	8	1	seq
.....Gggcaagaugucggcauagcuga.....	1	1	seq
.....uUgcaagaugucggcauagcuga.....	2	1	seq
.....uggcaagaGgucggcauagcuga.....	1	1	seq
.....uggcaaaAaugucggcauagcuga.....	2	1	seq
.....uggcaagauAucggcauagcuga.....	1	1	seq
.....uggcaagaugucggcauagCg.....	3	1	seq
.....uggcaagaugucggcauagCga.....	2	1	seq
.....uggcaagaugucggcauagUuga.....	2	1	seq
.....uggcaagaugucgcAcauagcuga.....	8	1	seq
.....uggcaagaugucggcGuagcuga.....	3	1	seq
.....uggcaagaugucggcauagcugC.....	8	1	seq
.....ugUcaagaugucggcauagcuga.....	2	1	seq
.....uggcaagaugucggcaGgcuga.....	2	1	seq
.....uggcaagaugucggcauagcugG.....	7	1	seq
.....uggcaagauCucggcauagcuga.....	1	1	seq
.....uggUaagaugucggcauagcuga.....	4	1	seq
.....uggcaagaugucggcauagcugaA.....	26	1	seq
.....uggcaagaugucggcauagcugaU.....	22	1	seq

dme-miR-31a-5p_MIMAT0000400

dme-miR-31a-3p_MIMAT0020834

uccguugguaauuggcaagaugucggcauagcugacguugaaaagcgauuuuugaagcgcuaugcugcaucuaguguucaaugga

.....uggcaagaugucggcauagcugaG.....	3	1	seq
.....uggcaagaugucggcauagcugac.....	1	0	seq
.....uggcaagaugucggcauagcugacuu.....	3	0	seq
.....Ugcaagaugucggcauagcuga.....	3	1	seq
.....ggcaagaugucggcauagcuga.....	2	0	seq
.....gcaagaugucggcauagcu.....	1	0	seq
.....caagaugucggcauagcuga.....	2	0	seq
.....cguugaaaagcgauuuuugaag.....	1	0	seq
.....cgcuaugcug <u>caucuag</u> Aca.....	1	1	seq
.....cgcuaugcug <u>caucuagu</u> ca.....	1	0	seq
.....cgcuaugcug <u>caucuagu</u> ag.....	3	0	seq
.....cgcuauGgcug <u>caucuagu</u>	2	1	seq
.....cgcuaugcug <u>caucuagu</u> gc.....	1	1	seq
.....Agcuaugcug <u>caucuagu</u>	38	1	seq
.....cgcuaugcug <u>caucuagu</u> ag.....	190	0	seq
.....cgcuaugcug <u>caucuagu</u> Ggu.....	1	1	seq
.....cgcuaugcuUcaucuagu <u>ca</u>	1	1	seq
.....cgcuaugcug <u>caucuGu</u> ca.....	1	1	seq
.....cgcuauCgcug <u>caucuagu</u>	2	1	seq
.....cgcuaugcug <u>caucuaA</u> ucagu.....	1	1	seq
.....cgcuaugcug <u>caC</u> cuagu <u>ca</u>	1	1	seq
.....cgcuaugcug <u>caucuagu</u> Cgu.....	1	1	seq
.....cgcGaugcug <u>caucuagu</u>	1	1	seq
.....cgcuaugcug <u>caucuaU</u> ucagu.....	2	1	seq