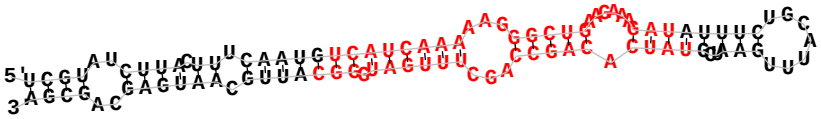
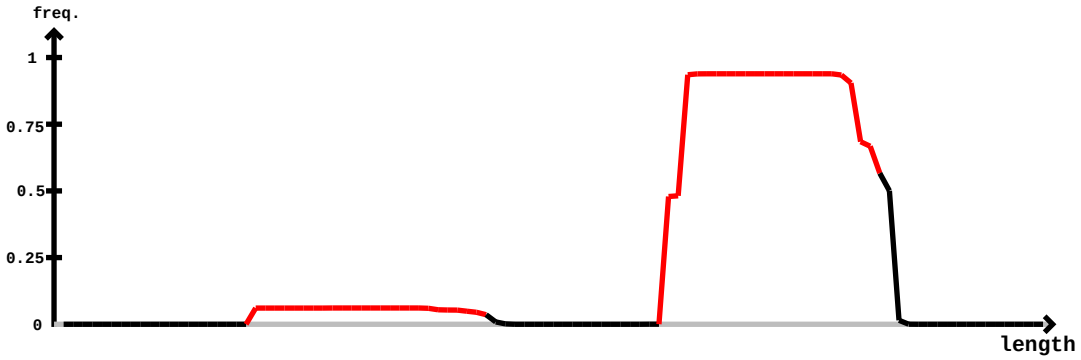
miRBase precursor : dme-mir-2c_MI0000431
Total read count : 3295
dme-miR-2c-3p_MIMAT0000411 : 3094 read count
dme-miR-2c-5p_MIMAT0020845 : 2084 read count
remaining reads : 0





dme-miR-2c-5p_MIMAT0020845

dme-miR-2c-3p_MIMAT0000411

5'- ucguaucuuacuuucaaugucaucaaaaagggcugaagaaagauauuucugcauuugaaucguaucacagccagcuuugaugggcauugcaaugagcagcga -3' exp

	reads	mm	sample
((((..((((.((.(((((((((((((...(((((.(((((....))))).....(((.....)))))))...)))))).))))))).)))))).))))			
.....ucaucaaaaagggcugaa.....	4	0	seq
.....ucaucaaaaagggcugaag.....	12	0	seq
.....Acaucaaaaagggcugaag.....	5	1	seq
.....ucaucaaaaGgggcugaag.....	1	1	seq
.....ucaucaaaaagggcugaaga.....	3	0	seq
.....ucaucaaaaagggcugaagaa.....	1	0	seq
.....Acaucaaaaagggcugaagaaa.....	1	1	seq
.....ucaucaaaaagggcugaagaaa.....	11	0	seq
.....ucaucaaaaagggcugaagCaa.....	1	1	seq
.....ucaucaaaaagggcuAaagaaa.....	1	1	seq
.....ucaucaaaaagggcugaagaaag.....	8	0	seq
.....ucaucaaaaagAgcugaagaaag.....	2	1	seq
.....Acaucaaaaagggcugaagaaag.....	2	1	seq
.....ucaucaaaaagggcGgaagaaag.....	1	1	seq
.....ucaucaaaaagggcugaagaaaga.....	22	0	seq
.....Acaucaaaaagggcugaagaaaga.....	8	1	seq
.....Acaucaaaaagggcugaagaaagau.....	25	1	seq
.....ucaucaaaaagggcugaagaaagau.....	58	0	seq
.....ucGucaaaaagggcugaagaaagau.....	2	1	seq
.....ucaucaaaaagggcugaagaaaUau.....	1	1	seq
.....ucaucaaaaagggcugaagaaagaA.....	2	1	seq
.....ucaucaaaaaAggcugaagaaagau.....	1	1	seq
.....ucaucaaaaagggcugaagaaagCua.....	1	1	seq
.....ucaucaaaaagggcugaagCaagaua.....	1	1	seq
.....Acaucaaaaagggcugaagaaagaua.....	4	1	seq
.....ucaucaaaaagggcugaagaaGgaua.....	1	1	seq
.....ucaucaaaaagggcugaagaaagaua.....	16	0	seq
.....ucaucaaaaagggcugaagaaagauau.....	2	0	seq
.....ucaucaaaaagggcugaagaaagauaA.....	3	1	seq
.....aagggcugaagaaagaua.....	1	0	seq
.....Aguaucacagccagcuuugaug.....	1	1	seq
.....Aaucacagccagcuuuga.....	1	1	seq
.....uaucacagccagcuuuga.....	13	0	seq
.....uaucaUagccagcuuuga.....	1	1	seq

dme-miR-2c-5p_MIMAT0020845

dme-miR-2c-3p_MIMAT0000411

ucguaucuuacuuucaaugucaucaaaaagggcugaagaaagauauuucugcauuugaaucguaucacagccagcuuugaugggcauugcaaugagcagcga

uaucacagccagcuuuAau	1	1	seq
uaucacagccagcuuugaA	1	1	seq
uaCcacagccagcuuugau	1	1	seq
uaucacagccagcCuugau	1	1	seq
uaucacagccagcuuugau	86	0	seq
uaucacagccagcuuCgau	2	1	seq
uaucaUagccagcuuugau	1	1	seq
Caucacagccagcuuugau	1	1	seq
uaucacagccagcuuugCu	1	1	seq
uaucacagccagcuuuCau	1	1	seq
uaucacagccagcuuugUu	1	1	seq
uaAcacagccagcuuugau	1	1	seq
uaucacagccagcuAugaug	1	1	seq
uaucacagccagcuuAgaug	5	1	seq
uaucacagccagcCuugaug	4	1	seq
uaucaGagccagcuuugaug	1	1	seq
uaucacagccagcuuugGug	2	1	seq
uaucacagccagcuuuUaug	2	1	seq
uaucacagccagcuuCgaug	3	1	seq
uaucacagccagcuuugaug	613	0	seq
uaucacagccagcuCugaug	1	1	seq
uaucacagccagcuuugaCg	1	1	seq
uaucaUagccagcuuugaug	1	1	seq
uaCcacagccagcuuugaug	2	1	seq
uaucCcagccagcuuugaug	1	1	seq
Aaucacagccagcuuugaug	4	1	seq
uaucacagccagcuuugaAg	2	1	seq
uaucacagccagcuuuCaug	2	1	seq
uaucacagccagcuuugauA	14	1	seq
uaucacagccagcAuugaug	1	1	seq
uauUacagccagcuuugaug	1	1	seq
uaucacagccagcuuugUug	1	1	seq
uaucacagccaAcuuugaug	5	1	seq
uaucacagccagcuuuAaug	4	1	seq
uaucacaAccagcuuugaug	1	1	seq
uaucacagccagcuuugCug	3	1	seq
uaucacagccagcuuugaugg	2	0	seq
uaucacagccagcuuugaugU	45	1	seq
uaucacagccagcuuugaGgg	3	1	seq
uaucacagccagcuuGgaugg	1	1	seq
uaucacagccagcuuugaugC	2	1	seq
uaucacagccagcuuugaugggg	11	0	seq
uaucacagccagcuuugaugUg	255	1	seq
Aaucacagccagcuuugaugggg	1	1	seq
uaucacagccagcuuugaugCg	13	1	seq
uaucacagAcagcuuugaugggg	1	1	seq
uaucacagccagcuuugaGggg	38	1	seq
uaucacagccagcuuugaugUgc	170	1	seq
uaucacagccagcGuugaugggc	1	1	seq
uaucacagccagcuuugaugCgc	2	1	seq
uaucacagccagcuuugaugggc	7	0	seq
uaucacagccagcuuugaGgggc	19	1	seq
uaucacagccagcuuugaugAgca	177	1	seq
uaucacagccagcuuugaugggcU	35	1	seq
uaucacagccagcuuugaugggca	2	0	seq
uaucacagccagcuuugaugUgca	1	1	seq
uaucacagccagcuuugaugAgcau	2	1	seq
uaucacagccagcuuugaugggcUu	1	1	seq
aucacagccagcuuugau	1	0	seq
aucacagccagcuuugaug	1	0	seq
aucacGgccagcuuugaug	1	1	seq
aucacagccagcuuugaugUgc	2	1	seq
aucacagccagcuuugaugggU	1	1	seq
aucacagccagcuuugaugggcU	1	1	seq
aucacagccagcuuugaugggca	2	0	seq
ucacagccagcuuugGug	1	1	seq
ucacagccaAcuuugaug	1	1	seq
ucacagccagcuuugUug	1	1	seq
ucacagccagcuuugaug	42	0	seq

dme-miR-2c-5p_MIMAT0020845

dme-miR-2c-3p_MIMAT0000411

ucguaucuuacuuucaaugucaucaaaaagggcugaagaaagauauuucugcauuugaaucguaucacagccagcuuugaugggcauugcaaugagcagcga

.....ucacagccagcCuugaug.....	1	1	seq
.....ucacCgccagcuuugaug.....	1	1	seq
.....ucacagccagcuuugaugg.....	3	0	seq
.....ucacagccagcuuugaugUg.....	6	1	seq
.....ucacagccagcuuugauggg.....	7	0	seq
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.....Acacagccagcuuugaugggca.....	51	1	seq
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.....Gcacagccagcuuugaugggca.....	3	1	seq
.....ucacagccagcuuCgaugggca.....	5	1	seq
.....ucacagccagcuuugaugggAa.....	1	1	seq
.....ucacagccagcuuugaugggcC.....	5	1	seq
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.....ucacagccagcuuugaugggca.....	1193	0	seq
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.....ucacGgccagcuuugaugggca.....	2	1	seq
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.....ucacagccagcuGugaugggca.....	3	1	seq
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.....ucacagccagcuuugauCggca.....	1	1	seq
.....Ccacagccagcuuugaugggca.....	5	1	seq
.....ucacagccagcuuugaugCgca.....	2	1	seq
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.....ucacagccagcuuugaugUgcau.....	1	1	seq
.....ucacagccagcuuugaugAgcau.....	1	1	seq
.....Acacagccagcuuugaugggcau.....	1	1	seq
.....ucacagccagcuuugaugggcUu.....	1	1	seq
.....ucacagccagcuuugaugggcaA.....	11	1	seq
.....ucacagccagcuuugaugggcaC.....	7	1	seq
.....ucacagccagcuuugaugggcau.....	19	0	seq
.....Ccacagccagcuuugaugggcau.....	1	1	seq
.....ucacagccagcuuugaugggcauu.....	1	0	seq
.....cacagccagcuuugaugggca.....	7	0	seq
.....Aacagccagcuuugaugggca.....	2	1	seq
.....cacagccagcuuugaugggcaA.....	2	1	seq
.....acagccagcuuugaugUgc.....	1	1	seq