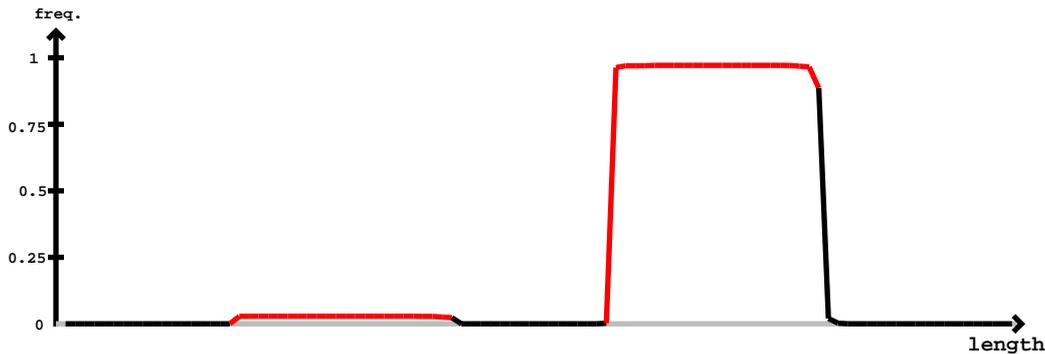
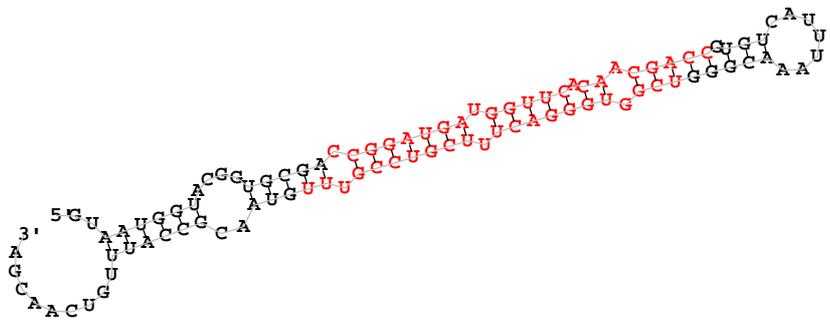


miRBase precursor : dme-mir-278_MI0000361
 Total read count : 2914
 dme-miR-278-5p_MIMAT0020805 read count : 2085
 dme-miR-278-3p_MIMAT0000339 read count : 829
 remaining reads : 0



dme-miR-278-5p_MIMAT0020805

dme-miR-278-3p_MIMAT0000339

5'	guaaugguacggugcgaccggaugaugguucacacaacgaccgugucauuuaaacgggucggugggacuuucgucgguuuuaacgcccauuugucaacga	-3'	exp	reads	mm	sample
.....	1	0	seq
.....	1	1	seq
.....	1	1	seq
.....	6	0	seq
.....	6	0	seq
.....	1	1	seq
.....	1	1	seq
.....	1	1	seq
.....	1	1	seq
.....	1	1	seq
.....	1	1	seq
.....	1	1	seq
.....	1	1	seq
.....	58	0	seq
.....	1	0	seq
.....	7	1	seq
.....	1	0	seq
.....	7	0	seq
.....	1	1	seq
.....	1	1	seq
.....	1	1	seq
.....	6	0	seq
.....	199	0	seq
.....	1	1	seq
.....	1	1	seq
.....	1	1	seq
.....	2	1	seq
.....	1	1	seq
.....	1	1	seq
.....	1	1	seq
.....	1	1	seq
.....	2	1	seq
.....	1	1	seq

guaaugguacggugcgacccggaugaugguucaacaacgaccgugucauuuaacgggucgguggggacuuucgucgguuuuaacgcacuuugucaacga

.....ucggugggacuuucCuccggu.....	1	1	seq
.....ucggugggacuuCcgucggu.....	1	1	seq
.....Ccguggggacuuucgucggu.....	1	1	seq
.....ucggugggacuuucgCccggu.....	4	1	seq
.....uAggggggacuuucgucggu.....	1	1	seq
.....ucggugggacuuucgucggu.....	3	1	seq
.....ucggCgggacuuucgucggu.....	1	1	seq
.....ucgguaAggacuuucgucggu.....	1	1	seq
.....ucggugggacuuucgucggu.....	2	1	seq
.....ucggugggacuuucgUcgu.....	2	1	seq
.....uUggugggacuuucgucggu.....	2	1	seq
.....ucggugggacuuucgucggu.....	3	1	seq
.....ucggAgggacuuucgucggu.....	1	1	seq
.....ucggugggacuuucAuucggu.....	3	1	seq
.....uUggugggacuuucgucggu.....	3	1	seq
.....ucggugggacuuucUucggu.....	2	1	seq
.....ucggugggacuuucGgucggu.....	1	1	seq
.....ucggugggacuuucCuucggu.....	11	1	seq
.....ucAgugggacuuucgucggu.....	1	1	seq
.....ucggugggacuuucgucCuuu.....	6	1	seq
.....ucggugggacuuucgucAuuu.....	1	1	seq
.....ucggugggacuuucgucggu.....	1	1	seq
.....ucggugUgacuuucgucggu.....	1	1	seq
.....ucggucGggacuuucgucggu.....	3	1	seq
.....ucggugggacuuucgUcgu.....	23	1	seq
.....ucggugggacuuucgUcgu.....	1	1	seq
.....ucggugggacuuucgucggu.....	3	1	seq
.....ucggugggacuuucgucggu.....	11	1	seq
.....ucggugggacuuucgucgGu.....	1	1	seq
.....ucggCgggacuuucgucggu.....	3	1	seq
.....ucggugggacuuucgucUggu.....	9	1	seq
.....ucggugggacuuucgCccggu.....	12	1	seq
.....ucggugggacuuucgUccggu.....	18	1	seq
.....ucgguaUggacuuucgucggu.....	1	1	seq
.....ucggugggacuuucgucggu.....	7	1	seq
.....Ccguggggacuuucgucggu.....	1	1	seq
.....ucggugggacuuucgucgGu.....	2	1	seq
.....ucggugggacuuucgucggu.....	3	1	seq
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.....ucggugggacuuucgucggu.....	2	1	seq
.....ucggugggacuuucgucggu.....	2	1	seq
.....ucggugggacuuucgucAguu.....	1	1	seq
.....ucggugggacuuucgucgGu.....	6	1	seq
.....ucggugggacuuucCuccggu.....	6	1	seq
.....ucggugggacuuucgucggu.....	7	1	seq
.....ucggugggacuuucgGccggu.....	5	1	seq
.....ucggugggacuuucgucggu.....	2	1	seq
.....ucgUugggacuuucgucggu.....	4	1	seq
.....ucggugggacuuucgucggu.....	4	1	seq
.....ucggugggacuuucgucgGu.....	1	1	seq
.....ucggugggacuuucgucggu.....	5	1	seq
.....ucggugggacuuucgucggu.....	5	1	seq
.....ucggugggacuuucgucggu.....	9	1	seq
.....ucUgggggacuuucgucggu.....	2	1	seq
.....ucgguaAggacuuucgucggu.....	6	1	seq
.....ucggugAgacuuucgucggu.....	3	1	seq
.....ucgCugggacuuucgucggu.....	2	1	seq
.....ucgAuugggacuuucgucggu.....	3	1	seq
.....ucggugggacuuucgucggu.....	2	1	seq
.....uGggugggacuuucgucggu.....	1	1	seq
.....Gcggugggacuuucgucggu.....	2	1	seq
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.....ucggugggacuuucgucggu.....	38	1	seq
.....ucggugggacuuucgucggu.....	1	0	seq
.....ucggugggacuuucgucggu.....	11	1	seq
.....ucggugggacuuucgucggu.....	1	1	seq

dme-miR-278-5p_MIMAT0020805

dme-miR-278-3p_MIMAT000339

guaaugguacggugcgaccggaugaugguucaacaagaccgugucauuuaaacgggucggugggacuuucgucceguuuuaacgccauuugucaacga

.....ucggugggacuuucgucceguuuUu.....	7	1	seq
.....cggugggacuuucgucceguu.....	2	0	seq
.....cggugggacuCucgucceguuu.....	1	1	seq
.....cggugggacuuucgucceguuu.....	14	0	seq
.....cggugggacuuucgCceguuu.....	2	1	seq
.....guggAacuuucgucceguuu.....	1	1	seq
.....ugggacuuucgucceguCu.....	1	1	seq
.....ugggacuuucgucceguuu.....	2	0	seq
.....ugggacuuucgucceguAu.....	1	1	seq