

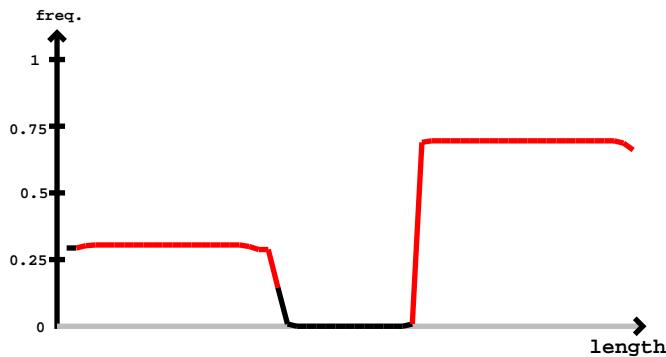
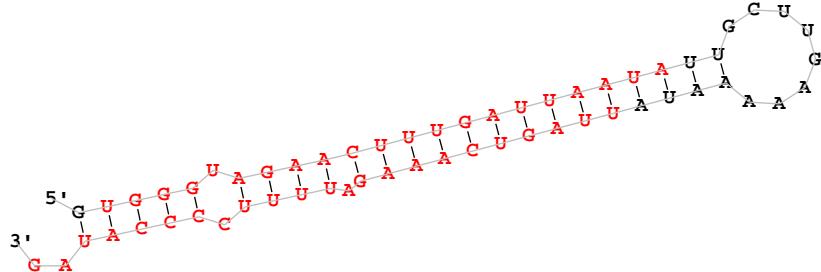
miRBase precursor : dme-mir-1012_MI0005873

Total read count : 351

dme-miR-1012-5p_MIMAT0020906 read count

dme-miR-1012-3p_MIMAT0005025 read count

remaining reads : 0



dme-miR-1012-3p_MIMAT0005025

5' -	-3'	exp	
reads	mm	sample	
(((((.....((((((.....))))))))....))).)).)).)).).	0	seq	
guggguagaacuuugauu.....	2	0	seq
guggguagaacuuugauu.....	4	0	seq
guggguagaacuuugauu.....	43	0	seq
guggguagaacuuugauu.....	1	1	seq
guggguagaacuuugauu.....	1	1	seq
guggguagaacuuugauu.....	1	1	seq
guggguagaacuuugauu.....	1	1	seq
guggguagaGcuuugauu.....	1	1	seq
guggguagaacuuugauu.....	45	0	seq
guggguuaacuuugauu.....	1	1	seq
guggguagaacuuugauu.....	1	1	seq
guggguagaacuuugauu.....	1	1	seq
gGggguagaacuuugauu.....	1	1	seq
guggguagaacuuugauu.....	1	1	seq
guggguagaacuuugauu.....	1	0	seq
.uggguagaacuuugauu.....	2	0	seq
.uggguagaacuuugauu.....	1	0	seq
..ggguagaacuuugauu.....	1	1	seq
.....auagucaaagauuuucccauag	3	0	seq
.....uuagucaaagauuuucccauag	3	0	seq
.....uuagucaaagauuuucccau.....	9	0	seq
.....uuagucaaagGuuuucccauag	1	1	seq
.....uuagucaaagauuuucccauA	2	1	seq
.....uuagucaaagauuuucccau.....	1	1	seq
.....uuagucaaagauuuucccauU	1	1	seq
.....uuagucaaagauuuucccauag	1	1	seq
.....uuagucaaagauuuucccauag	1	1	seq
.....uuagucaaagauuuucccauag	1	1	seq
.....uuagucaaagauuuucccauag	1	1	seq
.....uuagucaaagauuuucccauag	1	1	seq
.....Auagucaaagauuuucccauag	5	1	seq
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.....uuagucaaagauuuucccauag	1	1	seq
.....uuagucaaagauuuucccauag	1	1	seq
.....uuagucaaGgauuuucccauag	2	1	seq

dme-miR-1012-3p_MIMAT0005025

dme-miR-1012-5p_MIMAT0020906

guggguagaacuuugauuaauuugcuuugaaaaaauauuagucaaagauuuuccccauag

.....uuagucaaagauuuucccccaCag	1	1	seq
.....uuagucaaagauuuuccUcauag	1	1	seq
.....uuagucaaagauuuuccccGuag	1	1	seq
.....uagGcaaagauuuuccccauag	1	1	seq
.....uagucaaagauuuuccccauCg	1	1	seq