

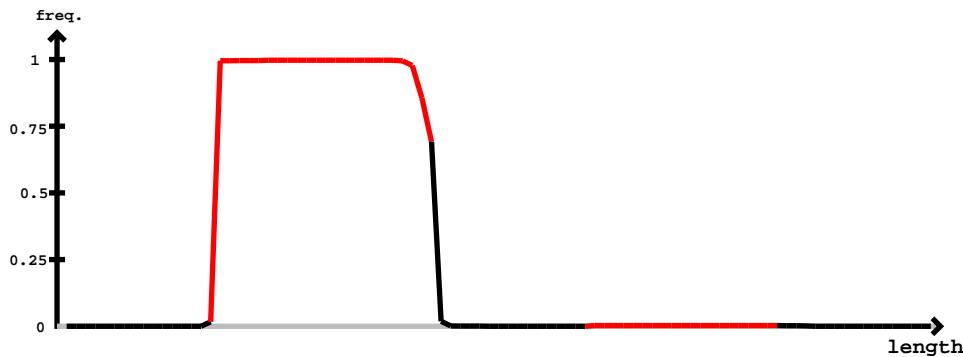
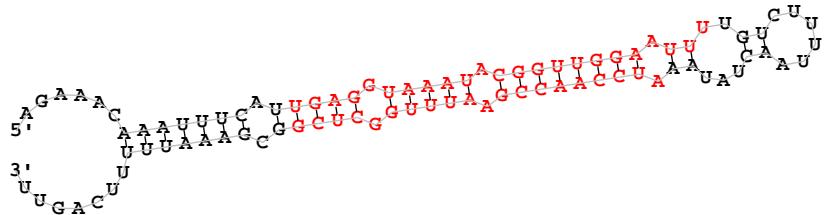
miRBase precursor : dme-mir-984_MI0005842

Total read count : 1507

dme-miR-984-3p_MIMAT0020881 read count

dme-miR-984-5p_MIMAT0005500 read count

remaining reads : 0



dme-miR-984-5p_MIMAT0005500

dme-miR-984-3p_MIMAT0020881

5'	-3'	exp	reads	mm	sample
.((((.....((((.(((.(((((.(((.((..((.....))....).))))))).)).))).))).))).....				
.....uugagguaaaauacgguuggaa.....			4	0	seq
.....uugagguaaaauacgguuggaa.....			7	0	seq
.....uugagguaaaauacgguuggGau.....			1	1	seq
.....uugagguaaaauacgguuggaa.....			3	0	seq
.....uugagguaaaauacgguuggaa.....			10	0	seq
.....uugagguaaaauacgguuggga.....			4	0	seq
.....ugagguaaaauacgguuggGa.....			1	1	seq
.....ugagguaaaauacgguuggaa.....			20	0	seq
.....ugagguaaaauacgguAggaa.....			1	1	seq
.....ugagguaaaauacggGuggaa.....			1	1	seq
.....ugagguaaaauacgguuggGa.....			2	1	seq
.....ugagguaaaauacgguuggaaC.....			1	1	seq
.....ugagguaaaauacgguuggGau.....			12	1	seq
.....ugagguaaaauacgguuggaa.....			1	1	seq
.....ugagguaaGuacgguuggaa.....			1	1	seq
.....ugagguaaaauacgguuggaa.....			150	0	seq
.....ugagguaaaauacgguuggaaA.....			1	1	seq
.....ugGgguaaaauacgguuggaa.....			1	1	seq
.....Cgagguaaaauacgguuggaa.....			1	1	seq
.....uaggguaaaauacgguugGaaau.....			2	1	seq
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.....ugagguaaaauacgCuuuggaa.....			1	1	seq
.....ugagguaaaauacgguuggGau.....			27	1	seq
.....ugagguaaaauacggGuggaa.....			2	1	seq
.....ugagguaaaGacgguuggaa.....			1	1	seq
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.....ugagguaaaauacAguuggaa.....			1	1	seq
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.....ugagguaaaauacggCuggaa.....			2	1	seq
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.....ugagguaaaauacgguuggCauu.....			1	1	seq

dme-miR-984-5p_MIMAT0005500

dme-miR-984-3p_MIMAT0020881

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.....Cgagguaauacgguuggaaau.....	1	1	seq
.....ugagguaauacgguugAaauu.....	3	1	seq
.....ugagguaauacAgugggaaau.....	1	1	seq
.....uUagguaauacgguuggaaau.....	2	1	seq
.....ugagguaauacgguuggaaau.....	882	0	seq
.....ugagguaauacgguuggaaau.....	1	1	seq
.....ugagguaauacgguuggaaau.....	5	1	seq
.....ugagguaauacgguuggaaauC.....	1	1	seq
.....ugagguaauacgguugAgaauu.....	2	1	seq
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.....ugagguaauacggUuggaaau.....	2	1	seq
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.....ugagguaauacgguuggaaCuu.....	1	1	seq
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.....auccaaccgaauuuuggcucggA.....	1	1	seq
.....auccaaccgaauuuuggcucggcA.....	2	1	seq