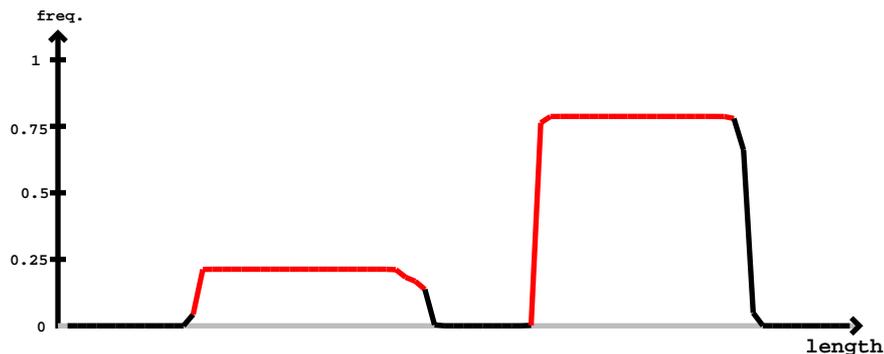
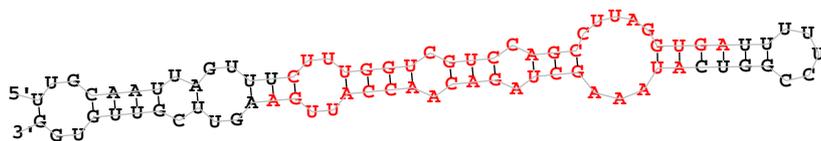


miRBase precursor : dme-mir-4_MI0000122
 Total read count : 18136
 dme-miR-4-5p_MIMAT0020785 read count : 3860
 dme-miR-4-3p_MIMAT0000109 read count : 1476
 remaining reads : 0



dme-miR-4-3p_MIMAT0000109

dme-miR-4-5p_MIMAT0020785

5'	3'	exp	reads	mm	sample
uugcaauuaguuuuuuuuggucuccagccuuaggugauuuuuuccgguc	auaaagcuagacaacc				
.....Aucuuuggucgucuccagccuuagg.....			1	1	seq
.....Aucuuuggucgucuccagccuuagguga.....			1	1	seq
.....ucuuuggucgucuccagccuuag.....			2	0	seq
.....ucuuuggucgucuccagccuuag.....			20	0	seq
.....Gcuuuggucgucuccagccuuag.....			8	1	seq
.....ucuuuggucgucuccagccuuagA.....			1	1	seq
.....ucuuuggucgucuccagccuuCgg.....			1	1	seq
.....ucuuuAagucgucuccagccuuagg.....			1	1	seq
.....ucuuuggucgucuccagccuuagg.....			1	1	seq
.....ucuuuggucgucuccagccuuAUG.....			2	1	seq
.....Acuuuggucgucuccagccuuagg.....			1	1	seq
.....ucuuuggCcgucuccagccuuagg.....			1	1	seq
.....ucuuugCugucuccagccuuagg.....			1	1	seq
.....ucuuuggucgucuccagcAuuagg.....			1	1	seq
.....ucuuugUcugucuccagccuuagg.....			2	1	seq
.....ucuCuggucgucuccagccuuagg.....			1	1	seq
.....Ccuuuggucgucuccagccuuagg.....			5	1	seq
.....ucuuuggucgCccagccuuagg.....			2	1	seq
.....ucUGuggucgucuccagccuuagg.....			1	1	seq
.....ucuuuggucgucuccagccuuagg.....			397	0	seq
.....ucuuuggGcgucuccagccuuagg.....			1	1	seq
.....ucuuuggucAuccagccuuagg.....			2	1	seq
.....Gcuuuggucgucuccagccuuagg.....			28	1	seq
.....ucuuuCGucgucuccagccuuagg.....			2	1	seq
.....ucuuuggucgucuccagccuuaggu.....			23	0	seq
.....ucuuuggucgCccagccuuaggu.....			1	1	seq
.....ucCuuggucgucuccagccuuaggu.....			1	1	seq
.....Gcuuuggucgucuccagccuuaggu.....			4	1	seq
.....ucuuuggucgucuccagccuuaggug.....			20	0	seq
.....ucuuuggucgucuccagccuuCggug.....			1	1	seq
.....ucuuuggucgucuccagccCuaggug.....			1	1	seq
.....Gcuuuggucgucuccagccuuaggug.....			4	1	seq
.....ucuuugAucgucuccagccuuaggug.....			1	1	seq
.....ucuuugguUgucuccagccuuagguga.....			1	1	seq

uugcaauuaguuuuuuugcuccagccuuaggugauuuuuucggucuaaagcuagacaaccauugaaguucguuggg

.....ucuuuggucguccagccuuaggugG.....	2	1	seq
.....ucuuuggucguccagccuuagguga.....	199	0	seq
.....ucuuCggucguccagccuuagguga.....	2	1	seq
.....Gcuuuggucguccagccuuagguga.....	18	1	seq
.....ucuuuggucguccagccuuaggugU.....	5	1	seq
.....ucuuuggucguccagccuGagguga.....	1	1	seq
.....ucuuuggucguccaAaccuuagguga.....	1	1	seq
.....ucuuuggucgucAagccuuagguga.....	2	1	seq
.....ucuuuggucguccagccCuagguga.....	1	1	seq
.....ucuuuggucguccagccuuaggAga.....	1	1	seq
.....ucuuuggucguccagccuuaggugCu.....	1	1	seq
.....ucuuuggucguccagccuuaggugaA.....	1	1	seq
.....ucuuuggucguccagccuuaggugau.....	2	0	seq
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.....cuuuggucguccCgccuuaggug.....	1	1	seq
.....cuuuggucguccagccCuaggug.....	1	1	seq
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.....cuuuggucguccagcAuuaggug.....	2	1	seq
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.....cuuuAgucguccagccuuaggug.....	1	1	seq
.....cuuuggucguccagccuAaggug.....	1	1	seq
.....cuuuggucguccagccuuCggug.....	3	1	seq
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dme-miR-4-3p_MIMAT000109

dme-miR-4-5p_MIMAT0020785

uugcaauuaguuuuuuuugugcuccagccuuaggugauuuuuuccggucuaaaagcuagacaaccauugaaguucguugugg

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