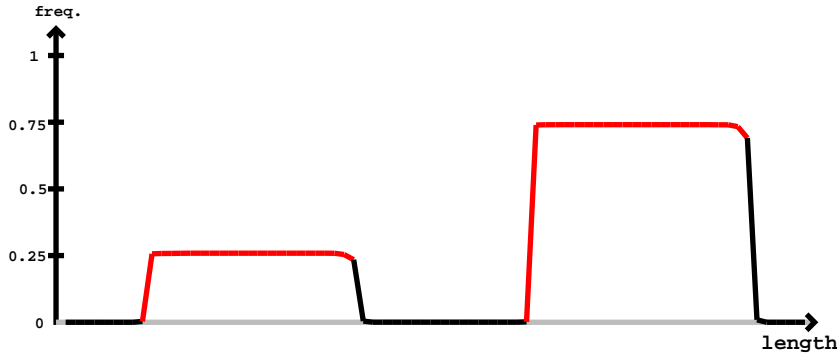
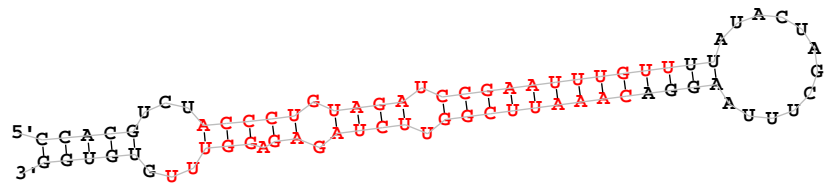


miRBase precursor : dme-mir-10_MI0000130
 Total read count : 5863
 dme-miR-10-3p_MIMAT0012531 read count
 dme-miR-10-5p_MIMAT0000115 read count
 remaining reads : 2



dme-miR-10-3p_MIMAT0012531

dme-miR-10-5p_MIMAT0000115

5'	3'	exp	reads	mm	sample
ccacgucua <u>acc</u> cug <u>uaga</u> accgaa <u>uuuguuuuu</u> uuacuagcuuaagga <u>caaa</u> u <u>ucgg</u> u <u>ucua</u> gag <u>aggguuu</u> gugugg					
.....uacc <u>c</u> gu <u>a</u> ga <u>acc</u> gaa <u>uuugu</u>			5	1	seq
.....uacc <u>c</u> gu <u>a</u> ga <u>acc</u> gaa <u>uuugu</u>			5	0	seq
.....Aacc <u>c</u> gu <u>a</u> ga <u>acc</u> gaa <u>uuuguu</u>			1	1	seq
.....uacc <u>c</u> gu <u>a</u> ga <u>acc</u> gaa <u>uuuguu</u>			2	0	seq
.....uacc <u>c</u> gu <u>a</u> ga <u>acc</u> gaa <u>uuugu</u> G.....			4	1	seq
.....uacc <u>c</u> gu <u>a</u> ga <u>acc</u> gaa <u>uuugu</u> C.....			1	1	seq
.....acc <u>c</u> gu <u>a</u> ga <u>acc</u> gaa <u>uuu</u>			2	0	seq
.....acc <u>c</u> gu <u>a</u> ga <u>acc</u> gaa <u>uuug</u>			27	0	seq
.....acc <u>c</u> gu <u>a</u> ga <u>acc</u> Aaa <u>uuugu</u>			1	1	seq
.....acc <u>c</u> gu <u>a</u> ga <u>acc</u> gCaa <u>uuugu</u>			1	1	seq
.....acc <u>c</u> gu <u>a</u> ga <u>acc</u> gaa <u>uu</u> Cgu.....			1	1	seq
.....acc <u>c</u> gu <u>a</u> ga <u>acc</u> gaa <u>uu</u> Cuugu.....			1	1	seq
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.....acc <u>c</u> gGag <u>a</u> ccgaa <u>uuuguu</u>			1	1	seq
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.....aAcc <u>c</u> gu <u>a</u> ga <u>acc</u> gaa <u>uuuguu</u>			10	1	seq
.....acc <u>c</u> gu <u>a</u> ga <u>acc</u> gaa <u>u</u> Guguu.....			3	1	seq
.....ac <u>A</u> c <u>u</u> gu <u>a</u> ga <u>acc</u> gaa <u>uuuguu</u>			7	1	seq
.....acc <u>c</u> gu <u>a</u> ga <u>acc</u> gaa <u>uuuguu</u>			2	1	seq
.....acc <u>c</u> gu <u>a</u> ga <u>acc</u> gaa <u>u</u> Cuguu.....			1	1	seq
.....acc <u>c</u> gu <u>a</u> ga <u>acc</u> gaa <u>uuuguu</u>			1	1	seq
.....acc <u>c</u> gu <u>a</u> ga <u>acc</u> gaa <u>uuuguu</u>			2	1	seq
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.....acc <u>c</u> gu <u>a</u> gC <u>uc</u> cgaa <u>uuuguu</u>			1	1	seq
.....acc <u>c</u> gu <u>a</u> ga <u>acc</u> gaa <u>uuugu</u> C.....			1	1	seq
.....acc <u>c</u> gu <u>a</u> ga <u>acc</u> gUaa <u>uuuguu</u>			3	1	seq
.....acc <u>c</u> gu <u>a</u> ga <u>acc</u> gaa <u>uu</u> Cguu.....			1	1	seq
.....acc <u>c</u> gu <u>a</u> Ca <u>uc</u> cgaa <u>uuuguu</u>			3	1	seq
.....acc <u>c</u> gu <u>a</u> ga <u>acc</u> gCaa <u>uuuguu</u>			16	1	seq

ccacgucuccccugagauccggaauuuguuuuuauacuagcuuaagga caaaucgguucuaagagagguuuugugugg

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..... acccugCagauccggaauuuguu.....	3	1	seq
..... acccCguagauccggaauuuguu.....	3	1	seq
..... acccuguCgauccggaauuuguu.....	1	1	seq
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..... acccuguagauccgCuuuuguu.....	5	1	seq
..... acccuguaAauccggaauuuguu.....	3	1	seq
..... acccuguagauccggaauuugCu.....	4	1	seq
..... acccCguagauccggaauuuguu.....	3	1	seq
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..... acccuguagGuccggaauuuguu.....	1	1	seq
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