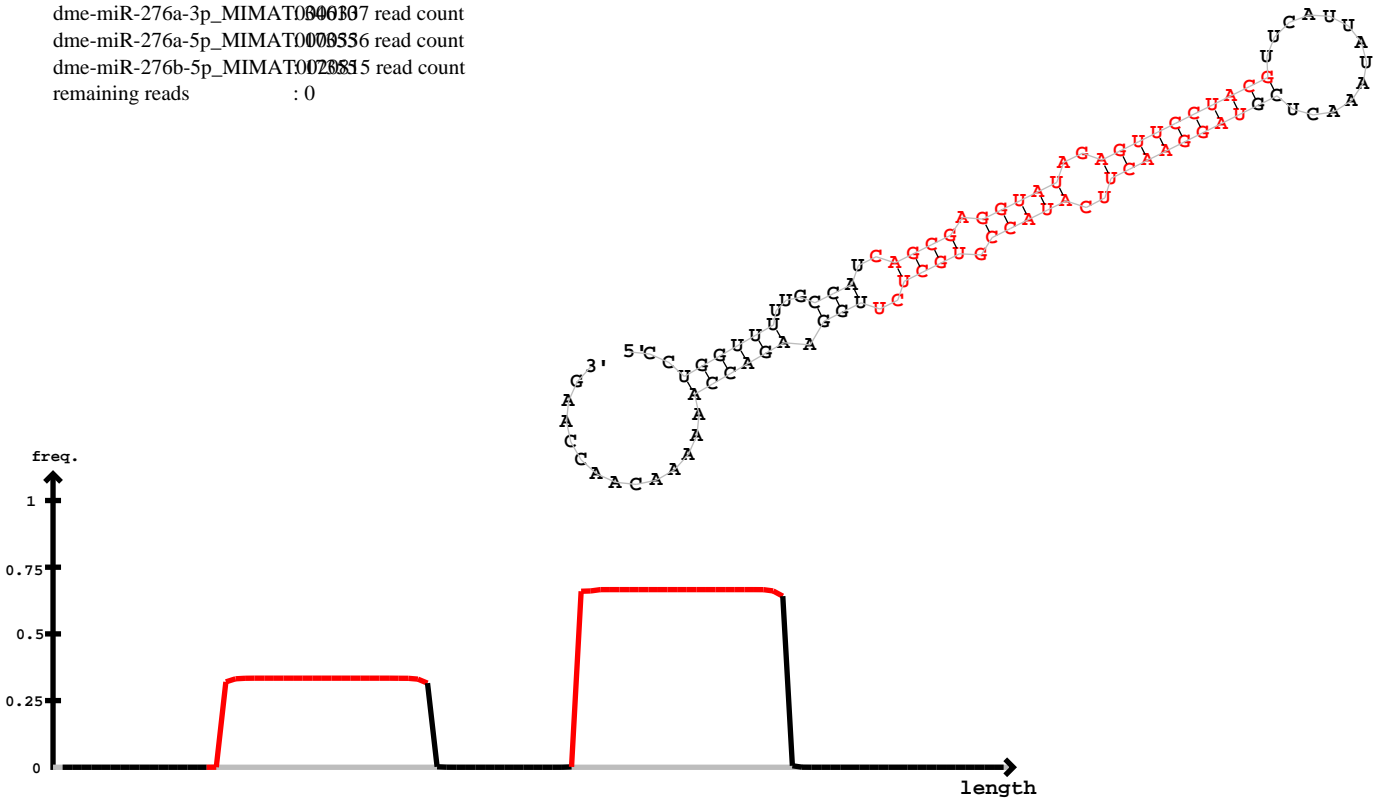


miRBase precursor : dme-mir-276a_MI0000359
 Total read count : 69320
 dme-miR-276a-3p_MIMAT0000337 read count
 dme-miR-276a-5p_MIMAT0000336 read count
 dme-miR-276b-5p_MIMAT0020815 read count
 remaining reads : 0



dme-miR-276b-5p_MIMAT0020815
 dme-miR-276a-5p_MIMAT0000336

dme-miR-276a-3p_MIMAT0000337

5'-	3'	exp	reads	mm	sample
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.....Aagcgaggguuagaguuccuacg.....			13	1	seq
.....cagcgaggguuagaguuccuacg.....			3	0	seq
.....agcgaggguuagaguucc.....			11	0	seq
.....agcgaggguuagaguuccA.....			1	1	seq
.....agcgaggguuagaguuccu.....			1	1	seq
.....agcgaggguuagaguuccu.....			12	0	seq
.....agcgaggguuagaguuccu.....			1	1	seq
.....agcgaggguuagaguuccu.....			2	1	seq
.....agcgaggguuagaguuccu.....			126	0	seq
.....agcgaggguuagaguuccu.....			1	1	seq
.....agcgaggguuagaguuccu.....			2	1	seq
.....Cgagggguuagaguuccu.....			8	1	seq
.....agcgaggguuagaguuccu.....			1	1	seq
.....agcgaggguuagaguuccu.....			1	1	seq
.....agcgaggguuagaguuccu.....			3	1	seq
.....agcgaggguuagaguuccu.....			1	1	seq
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.....agcgaggguuagaguuccu.....			4	1	seq
.....agcgaggguuagaguuccu.....			653	0	seq
.....agcgaggguuagaguuccu.....			1	1	seq
.....agcgaggguuagaguuccu.....			1	1	seq
.....agcgaggguuagaguuccu.....			4	1	seq
.....agcgaggguuagaguuccu.....			1	1	seq
.....agcgaggguuagaguuccu.....			2	1	seq
.....agcgaggguuagaguuccu.....			3	1	seq
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.....agcgaggguuagaguuccu.....			3	1	seq
.....agcgaggguuagaguuccu.....			2	1	seq
.....agcgaggguuagaguuccu.....			1	1	seq
.....agcgaggguuagaguuccu.....			7	1	seq

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..... agcAaggguuagagauccuac	4	1	seq
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..... agcgaggguuagagauccCac	3	1	seq
..... agcgaggguauaUaguuccuac	1	1	seq
..... agcgaggguuagagauGccuac	1	1	seq
..... agcgaggguuUcagauccuac	2	1	seq
..... agcgaggguuagagauccuGc	1	1	seq
..... agcgaCguuagagauccuac	1	1	seq
..... agcgaggguuAagagauccuac	2	1	seq
..... agcgGggguuagagauccuac	2	1	seq
..... agUgaggguuagagauccuac	5	1	seq
..... agcgaggguuagagauCccuac	2	1	seq
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..... agcgaggGauagagauccuac	2	1	seq
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..... agcgaggguuagauAuuuccuac	1	1	seq
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..... agcgaggguuagagauUuac	3	1	seq
..... Ccgaggguuagagauccuac	5	1	seq
..... Gcgaggguuagagauccuac	1	1	seq
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..... agcgaggguuagaguuAcuacg	12	1	seq
..... agcgaggguUuagagauccuacg	5	1	seq
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..... agcgaggguGuaagauccuacg	19	1	seq
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.....agcCagguauagaguuccuacg.....	7	1	seq
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.....gCgagguauagaguuccuUcg.....	1	1	seq
.....gCgagguauagaguucccCcuacg.....	3	1	seq
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.....gcgagguauagaguuccuacgU.....	3	0	seq
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dme-miR-276b-5p_MIMAT0020815

dme-miR-276a-5p_MIMAT0000336

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.....uaggaacucauaccguAcuc.....	6	1	seq
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.Caggaacucauaccgugcucu.....	79	1	seq
.uaggaacucauaccgugcuUu.....	28	1	seq
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.uaggaCcucauaccgugcucu.....	9	1	seq
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.uaggaacucauaccgugcucu.....	6	1	seq
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.uaggaacucauaccgAgcucu.....	5	1	seq
.uagUaacucauaccgugcucu.....	1	1	seq
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.aggaacucauaccgugcucu.....	51	0	seq
.aggaacucauaccgugcucu.....	1	1	seq
.aCgaacucauaccgugcucu.....	1	1	seq
.aggaacucauaccgugcucu.....	1	1	seq
.Uggaacucauaccgugcucu.....	2	1	seq
.aggaacuCcauaccgugcucu.....	1	1	seq
.aggaacucauaccgugcucu.....	1	0	seq
.ggaacucauaccgugcuc.....	1	0	seq
.ggaacucauaccgCgcuc.....	1	1	seq
.ggaacucauaccgGgcuc.....	1	1	seq
.ggaacucauaccgugcuc.....	4	0	seq
.Agaacucauaccgugcucu.....	1	1	seq
.ggaacucauaccgCgcucu.....	1	1	seq
.ggaacucauaccgugGucu.....	1	1	seq
.ggaacucauaccAugcucu.....	1	1	seq
.ggaacucauaccgugUucu.....	2	1	seq
.ggaacucauaccgugcCcu.....	1	1	seq
.ggaacucauaccgugcucu.....	1	1	seq
.Ugaacucauaccgugcucu.....	7	1	seq
.ggaacucauaccgugcucG.....	1	1	seq
.ggaacucauaccUgugcucu.....	1	1	seq
.ggaacuUauaccgugcucu.....	1	1	seq
.ggaacucauaccgulaucu.....	3	1	seq

dme-miR-276b-5p_MIMAT0020815

dme-miR-276a-5p_MIMAT000336

ccugguuuugccaucagcgagguauagaguuccuacguucauuuaaacucgdmggmRu026a0pgMgM006aa0accaaaaacaaccaag

.....ggGacuucauaccgugcucu.....	1	1	seq
.....ggaacuucauaccgugcucu.....	220	0	seq
.....ggaacuucauaccgugcucuC.....	1	1	seq
.....ggaacuucauaccgugcucu.....	1	0	seq
.....ggaacuucauaccgugcucuA.....	1	1	seq
.....gaacuucauaccgugcuc.....	1	0	seq
.....gaacuucauaccgugcucC.....	1	1	seq
.....gaacuucauaccgugcucu.....	4	0	seq
.....aacuucauaccgugcucu.....	2	0	seq