

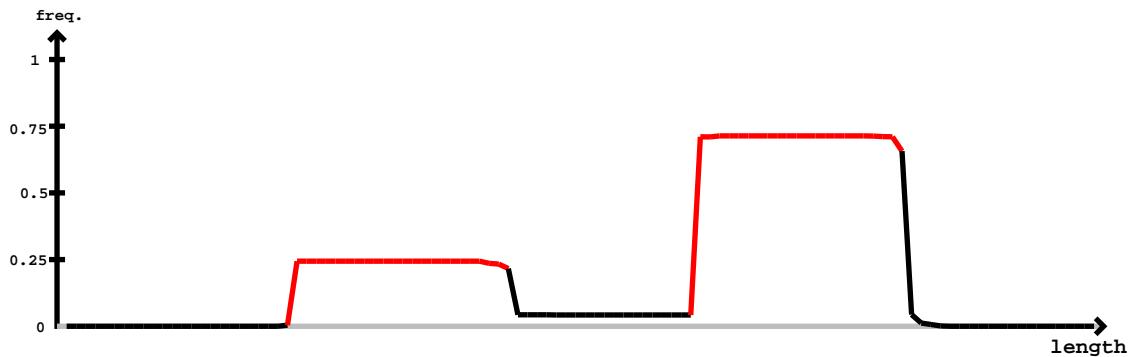
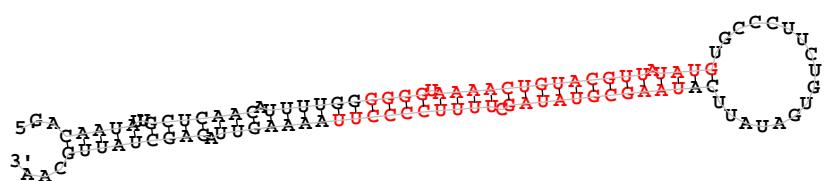
miRBase precursor : dme-mir-965_MI0005821

Total read count : 1488

dme-miR-965-3p MIMAT0006480 read count

dme-miR-965-5p MIMAT0036361 read count

remaining reads : 63



dme-miR-965-3p_MIMAT0005480

dme-miR-965-5p_MIMAT0020861

		-3'	exp	
		reads	mm	sample
.....	gacaauuuugcuacaacauuuugggguaaaaaacuguacguuaauugugccuuucugugauauucauaagcguaugcguaucuuuccccuu	1	0	seq
.....	ggggguaaaaacuguacguuuau.....	2	0	seq
.....	.Agggguaaaaacuguacguuuau.....	4	1	seq
.....	gggguaaaaacuguacguuu.....	12	0	seq
.....	gggguaaaaacuguacguuuau.....	4	0	seq
.....	gggguaaaaacuguacguuuA.....	1	1	seq
.....	gggguaaaaacuguacguuuau.....	22	0	seq
.....	gggguaaaaAuguacguuuau.....	2	1	seq
.....	gggguaaaaacuguacguuuau.....	291	0	seq
.....	gggguaaaaacuguacguuuauG.....	1	1	seq
.....	gggAuaaaaacuguacguuuau.....	4	1	seq
.....	gggguaaaaacuguacgAuau.....	1	1	seq
.....	gggguaUaacuguacguuuau.....	1	1	seq
.....	gggguGaaacuguacguuuau.....	1	1	seq
.....	gAgguaaaaacuguacguuuau.....	1	1	seq
.....	gggguaaaaacuguacgCuaua.....	2	1	seq
.....	gggguaaaaacugCacguuuau.....	3	1	seq
.....	gggguaaaaacuguacguuCua.....	2	1	seq
.....	gggguaaaaacuguCcguuuau.....	1	1	seq
.....	gggguaaaaacuguacguuuauCg.....	2	1	seq
.....	gggguaCaacuguacguuuau.....	1	1	seq
.....	ggggCaaacuguacguuuau.....	1	1	seq
.....	gggguaaaaacuguaAguuau.....	1	1	seq
.....	gggguaaaCcuguacguuuau.....	1	1	seq
.....	gggguaaaaacuguacguuuauGug.....	1	1	seq
.....	gggguaaaaacuguacguuuauuggc.....	1	0	seq
.....ugccuuucugugauauuc.....	9	0	seq
.....	ugccuuucugugauauca.....	45	0	seq
.....Ggccuuucugugauauca.....	1	1	seq
.....	ugGccuuucugugauauca.....	1	1	seq
.....	ugccuuucugugGuauucau.....	1	1	seq
.....	ugccuuucugugauaucaA.....	5	1	seq
.....	ugccuuucugugauaucaAa.....	1	1	seq
.....auaagcguaugcguaucuuuccccuu.....	9	0	seq
.....uaagcguaugcguaucuuucc.....	1	0	seq

dme-miR-965-5p_MIMAT0020861

gacaauauugcuacaacauuuugggguaaaaacuguacguauaugugcccuucugugauauucauaagcguauagcguuuucccuuuaaaaguuuagaggcuauugcaa

.....uaagcguauagcguuuuccc.....	3	0	seq
.....uaagcguauagcguuuuccc.....	1	0	seq
.....Gaagcguauagcguuuuccc.....	1	1	seq
.....Aaagcguauagcguuuuccc.....	3	1	seq
.....uaagcguauagcguuuucccU.....	3	1	seq
.....uaagcguauagcguuuuccc.....	70	0	seq
.....uCagcguauagcguuuuccc.....	2	1	seq
.....uaagcguauagcguuuucccui.....	1	1	seq
.....uaagcguauagcguuuucccui.....	3	1	seq
.....Aaagcguauagcguuuucccui.....	59	1	seq
.....uaagcguauagcguuuucccui.....	4	1	seq
.....uaagcguauagcguuuAccccui.....	1	1	seq
.....uaagcguauagcguuuAccccui.....	1	1	seq
.....uaagcguauagcguuuGccccui.....	1	1	seq
.....uaagcguauagcguuuAccccui.....	1	1	seq
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.....Caagcguauagcguuuucccui.....	4	1	seq
.....uaagcguauagcguuucccui.....	4	1	seq
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.....uaagcguauagcguuuucccui.....	1	1	seq
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.....uaagcguauagcguuuucccui.....	1	1	seq
.....uaagcguauGgcuuucccui.....	1	1	seq
.....Gaagcguauagcguuuucccui.....	1	1	seq
.....uaagcguauagcguuuucccui.....	1	1	seq
.....uaagcguauagcguuuucccuiA.....	9	1	seq
.....uaagcguauagcguuuucccuiA.....	1	1	seq
.....uaagcguauagcguuuucccuiC.....	2	1	seq
.....uaagcguauagcguuuucccuiA.....	15	0	seq
.....uaagcguauagcguuuucccuiA.....	1	1	seq
.....uaagcguauagcguuuucccuiU.....	28	1	seq
.....uaagcguauagcguuuucccuiCaa.....	1	1	seq
.....uaagcguauagcguuuucccuiCa.....	1	1	seq
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.....uaagcguauagcguuuucccuiUaa.....	5	1	seq
.....uaagcguauagcguuuucccuiaaa.....	3	0	seq
.....uaagcguauagcguuuucccuiUaaa.....	2	1	seq
.....aagcguauagcguuuucccuiGu.....	1	1	seq
.....aagcguauagcguuuucccui.....	4	0	seq
.....agcguauagcguuuucccui.....	3	0	seq
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