

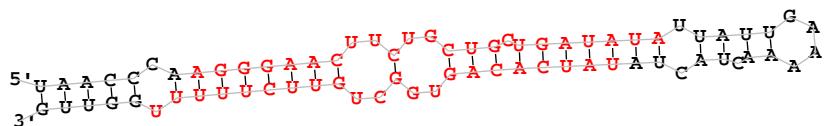
miRBase precursor : dme-mir-6-2\_MI0000125

Total read count : 2813

dme-miR-6-3p\_MIMAT0000190 read count

dme-miR-6-2-5p\_MIMAT0026288 read count

remaining reads : 0



dme-miR-6-2-5p\_MIMAT0020788

dme-miR-6-3p\_MIMAT0000111

5'	-3'	exp	sample
reads	mm		
(((((.....((((((.....((((((.....)).))).))).))).))).)).))	1	1	seq
....Aagggaacuuucugcugcugauua	9	0	seq
....aagggaacuuucugcugcugauua	24	0	seq
....aagggaacuuucugcugcugauua	2	1	seq
....aagggaacuuucugcugcugauua	4	0	seq
....aagggaacuuucugcugcugauua	1	1	seq
....aagggaacuuucugcugcugauua	9	0	seq
....aagggaacuuucugcugcugauu	1	1	seq
....aagggaacuuucugcugcugauu	1	1	seq
....aCggAACUUUCUGCUGCUGAUUA	1	1	seq
....agggaacCUCUGCUGCUGAUUA	1	1	seq
....aAgggAACUUUCUGCUGCUGAUUA	1	1	seq
....agggaacACUGCUGCUGAUUA	1	1	seq
....agUgaacuuucugcugcugauua	1	1	seq
....agggaacuuucugCgcugauua	1	1	seq
....agggaacuuucugcugcugauua	219	0	seq
....agggaacuuucugcugcugGuau	1	1	seq
....agggaacuuucugcugcugauua	1	1	seq
....agggaacuuucugcugcugauaG	3	1	seq
....agggaacuuucugcuAcugauua	1	1	seq
....aUgggaacuuucugcugcugauua	2	1	seq
....Ggggaacuuucugcugcugauua	2	1	seq
....agggaacuuucugcugcugauua	1	1	seq
....Ugggaacuuucugcugcugauua	1	1	seq
....agggaacuuucugcugcugauGu	2	1	seq
....agggaacuuucugcugcugCgauua	2	1	seq
....agggCACUUUCUGCUGCUGAUUA	3	1	seq
....agggaacuuucugGugcugauua	1	1	seq
....agAgAACUUUCUGCUGCUGAUUA	1	1	seq
....agggaacuuucugcugcugauua	2178	0	seq
....agggaacuuucugcugcugCuaua	7	1	seq
....agggGACUUUCUGCUGCUGAUUA	5	1	seq
....agggaacuuucugcugcugGuaua	4	1	seq

dme-miR-6-2-5p MIMAT0020788

dme-miR-6-3p MIMAT0000111

**uaacccaaggaaacuuucugcugcugauauuuauugaaaaacuacuaauacaguggcuguuucuuuuugguuu**

.....agggaacuuucugcugcugaCaua.....	4	1	seq
.....agggaauUucugcugcugauua.....	5	1	seq
.....aggUaacuuucugcugcugauua.....	3	1	seq
.....agggaacuuucugAugcugauua.....	3	1	seq
.....agggaacuuucugcugcugauua.....	3	1	seq
.....agggaacuuucugcugJugauua.....	3	1	seq
.....agggaacuuucugcugcugauaCa.....	2	1	seq
.....aCgaaacuuucugcugcugauua.....	1	1	seq
.....aAggaacuuucugcugcugauua.....	1	1	seq
.....agggaacuuucugcugAugauua.....	5	1	seq
.....agggaacuuucugcugcugauua.....	1	1	seq
.....agggaacuuCcgcugcugauua.....	4	1	seq
.....agggaacuuucugcugcugJuaua.....	6	1	seq
.....agggUacuuucugcugcugauua.....	1	1	seq
.....agggaacuuucugcGgcugauua.....	2	1	seq
.....agggaacCucugcugcugauua.....	3	1	seq
.....agggaCcucugcugcugauua.....	2	1	seq
.....agggaacuCcugcugcugauua.....	1	1	seq
.....agggaacuuCuUcugcugauua.....	1	1	seq
.....agCgaaacuuucugcugcugauua.....	1	1	seq
.....agggaGcuucugcugcugauua.....	2	1	seq
.....agggaacuuucugcugcugauua.....	1	1	seq
.....agggaacuuGcugcugcugauua.....	1	1	seq
.....agggaacuuucugcGcgugauua.....	3	1	seq
.....agggaacuuAugcugcugauua.....	1	1	seq
.....agggaacuuucugJugcugauua.....	1	1	seq
.....agggaacuuucugcugcugJuaau.....	1	1	seq
.....agggaacuuucugcugcugauua.....	34	0	seq
.....agggaacuuucugcugcugauuaA.....	6	1	seq
.....agggaacuuucugcugJugauua.....	1	1	seq
.....agggaacuuCcgcugcugauua.....	1	1	seq
.....agggaacuuucugcugcugauauu.....	1	0	seq
.....gggaacuuucugcugcugauau.....	2	0	seq
.....gggaacuuucugcugcugauua.....	20	0	seq
.....ggaacuuucugcugcugauua.....	1	0	seq
.....gaacuuucugcugcugauua.....	1	0	seq
.....uaucacaguUgcuguuuccuu.....	1	1	seq
.....uaucacagugCcuguuuccuu.....	1	1	seq
.....uaucacaguAgcuguuuccuu.....	4	1	seq
.....uaucacaguggcuUuuuccuu.....	1	1	seq
.....uaucacagagGgcuguuuccuu.....	1	1	seq
.....uaucacagugAcuguuuccuu.....	10	1	seq
.....uaucacagugUcguguuccuu.....	1	1	seq
.....uaucacagugguCuuuccuu.....	2	1	seq
.....uaucacaguUcguguuccuu.....	1	1	seq
.....uaucacaguggeCguuuccuu.....	1	1	seq
.....uaucacaguggcCguuuccuu.....	1	1	seq
.....uaucacaguggcguuuccuu.....	1	1	seq
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.....uaucacaguggcguuUuuuccuu.....	5	1	seq
.....uaucacaguggcguuAuuuccuu.....	3	1	seq
.....uaucacaguggeUuuuccuu.....	28	1	seq
.....uaucacaguggeCuuuccuu.....	3	1	seq
.....uaucGcaguggeCguuuccuu.....	1	1	seq
.....uaucacaguggeCguuuccuu.....	5	0	seq
.....uaucacagugggUguuuccuu.....	8	1	seq
.....uaucacaguAgcuguuuccuu.....	25	1	seq
.....uaucacaguUgcuguuuccuu.....	14	1	seq
.....uaucacaguggcguuuccuuCuu.....	1	1	seq
.....uaucacCguggcguuuccuu.....	1	1	seq
.....uaucacaguggeCguuuccuu.....	1	1	seq
.....uaucacaguggcguuGuuuuccuu.....	4	1	seq
.....uaucacagagggcguuuccuu.....	1	1	seq
.....uaucacaguggeCguuuccuu.....	6	1	seq
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.....uaucacaguCcuguuuccuu.....	2	1	seq
.....uaucacaguggcguuuccuu.....	1	1	seq
.....uaucacaguggGguuuccuu.....	1	1	seq
.....uaucacGguuaccuuuccuu.....	6	1	seq

dme-miR-6-2-5p\_MIMAT0020788

dme-miR-6-3p\_MIMAT0000111

uaacc  
.....  
.....  
.....  
.....  
.....

			seq	
.....	uaucacaguggcCguucuuuuu.....	9	1	seq
.....	.aucacagugUcuguucuuuu.....	1	1	seq
.....	.ucacGguggcuguuucuuuu.....	1	1	seq
.....	.ucacaguAgcuguuucuuuu.....	1	1	seq
.....	.ucacaguggUuguuucuuuu.....	1	1	seq