

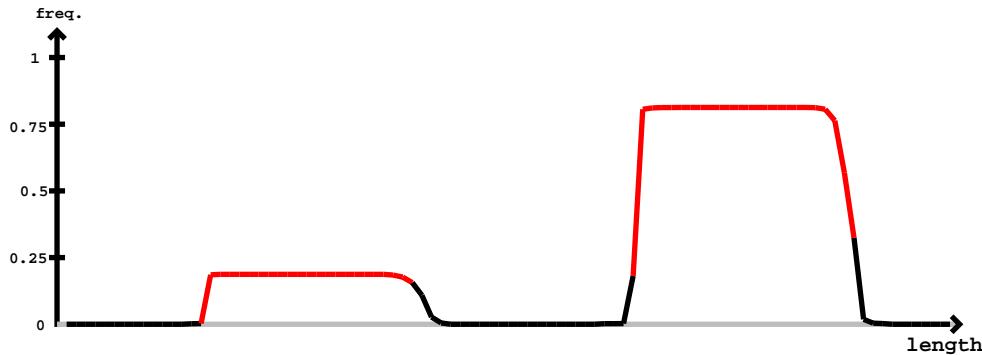
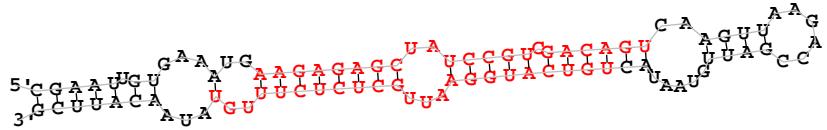
miRBase precursor : dme-mir-281-2_MI0000370
Total read count : 2217

Total read count : 2217

dme-miR-281-3p_MIMAT0000945 read count

dme-miR-281-2-5p_MIMAT0050349 read count

remaining reads : 5



dme-miR-281-2-5p_MIMAT0000349

dme-miR-281-3p_MIMAT0000345

5'- cggaaauugaaaaugaaagagacuauccgucgcacagucaaguuuaagaccgauuguaauac	-3'	exp	
	reads	mm	sample
.....(.....(.....(((((.....((((.....((.....))))....))))....))))....))))))	1	0	seq
.....ugaagagagacuauccgucgcac.....	3	0	seq
.....ugaagagagacuauccgucgcaca.....	1	0	seq
.....gaagagagacuauccgucgcacagu.....	1	0	seq
.....Aaagagagacuauccgucgcacagu.....	1	1	seq
.....aagagagacuauccgucgcac.....	4	0	seq
.....aagaAagacuauccgucgcac.....	1	1	seq
.....aagagagacuauccgucCaca.....	1	1	seq
.....aagagagacuauccgucgcacC.....	1	1	seq
.....aagagagacuauccgucgcaca.....	12	0	seq
.....aagaCagcuauccgucgcacag.....	1	1	seq
.....aGgagagacuauccgucgcacag.....	1	1	seq
.....aagagagacuauccgucgcacac.....	1	1	seq
.....aagagagacuauccgucgcacag.....	42	0	seq
.....aagagagacuauccgucgcacGg.....	1	1	seq
.....aagagagacuaucGgucgcacagu.....	1	1	seq
.....aagagagacuauccgucgcacagu.....	85	0	seq
.....aagagagacuauccgucgcacagA.....	3	1	seq
.....aagagGgcuauccgucgcacagu.....	1	1	seq
.....aagagagacuaucAgugcacagu.....	1	1	seq
.....aagagagacuauccgucgcacagC.....	3	1	seq
.....aagGgagacuauccgucgcacagu.....	1	1	seq
.....aGgagagacuauccgucgcacagu.....	1	1	seq
.....aagagagacuauccgugacagu.....	2	1	seq
.....aagagagacuauccCcgacagu.....	1	1	seq
.....aaAagagacuauccgucgcacagu.....	1	1	seq
.....aCgagagacuauccgucgcacagu.....	1	1	seq
.....aagagagacuauccUucgcacagu.....	1	1	seq
.....aagagagacuauccgCcgacagu.....	1	1	seq
.....aaAagagacuauccgucgcacagu.....	1	1	seq
.....aagagagacuauccgucgcacagu.....	2	1	seq
.....aagagagacuauccgucgcacAgc.....	4	1	seq
.....aagagagacuauccgucCacaguc.....	2	1	seq
.....aagagagAauccgucgcacagu.....	1	1	seq
.....aagagagacuauccgGgcacaguc.....	1	1	seq
.....aaqaaqacquaGccgucgcacagu.....	1	1	seq

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.....aagagagcuaCccgucgacaguc.....	1	1	seq
.....aagagagcuauccgucgacaguA.....	5	1	seq
.....aagagagcuauccgucgUcaguc.....	1	1	seq
.....aagagagcuauccgAcgacaguc.....	3	1	seq
.....aagagagcuauccgCcgcacaguc.....	6	1	seq
.....aagagagcuaucUgucgacaguc.....	1	1	seq
.....aagagagcuauccgucgaAaguc.....	1	1	seq
.....aagagagcuauccgucAacaguc.....	2	1	seq
.....aagagagcuauccgucgacagC.....	4	1	seq
.....aaAaagagcuauccgucgacaguc.....	1	1	seq
.....aagagagcuauccgucgacaguc.....	140	0	seq
.....aagagagcuauccgucgacagGc.....	2	1	seq
.....aagagagcuauccgucgacagucU.....	1	1	seq
.....Gagagagcuauccgucgacaguca.....	1	1	seq
.....aagagagcuauccgucgacagAca.....	1	1	seq
.....aagagagcuaUcguccgacaguca.....	1	1	seq
.....aagagagcuauccgucgacaguca.....	18	0	seq
.....aagagagcuauccgucgacagucC.....	27	1	seq
.....aagagagcuauccgucgacagucCa.....	5	1	seq
.....aagagagcuauccgucgacagucaa.....	4	0	seq
.....aagagagcuauccgucgacaguAaa.....	1	1	seq
.....agagagcuauccgucgacagu.....	1	0	seq
.....agagagcuauccgucgacaguc.....	2	0	seq
.....agagagcuauccgucgacagucaa.....	1	0	seq
.....auacugucauggaaauugcucuc.....	2	0	seq
.....auacugucauggaaauugcucucu.....	1	0	seq
.....auacugucauggaaauuAcucucu.....	1	1	seq
.....auacugucauggaaauugUucucu.....	1	1	seq
.....acugucauggaaauugcucucuuug.....	1	0	seq
.....cugucauggaaauugcucucu.....	3	0	seq
.....cugucauggaaauugcucuA.....	1	1	seq
.....cugucauggaaUuugcucucu.....	1	1	seq
.....cugucauggaaauugcucucuU.....	2	1	seq
.....cugucauggaaauugcucucu.....	55	0	seq
.....Augucauggaaauugcucucu.....	1	1	seq
.....cugucauggaaauugcAcucuu.....	2	1	seq
.....cugucauggaaauugcucuA.....	1	1	seq
.....cugucauggaaauugcUucuuu.....	3	1	seq
.....cugucaugUaaauugcucucuu.....	1	1	seq
.....cugucauggaaauugUucucuu.....	1	1	seq
.....cugCcuggaaauugcucucuu.....	2	1	seq
.....cugucauggaaauugcAcucuuu.....	2	1	seq
.....cugucGuggaaauugcucucuuu.....	1	1	seq
.....cugAcauggaaauugcucucuuu.....	1	1	seq
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.....cuguUauggaaauugcucucuuug.....	1	1	seq
.....cugucauggaaauugcucuAcuuiug.....	3	1	seq
.....cugucauggaaauugcAcuuiug.....	1	1	seq
.....Augucauggaaauugcucucuuug.....	2	1	seq
.....cugucauggaaauugcucucuCug.....	1	1	seq
.....cugucauggaaauugCcucucuuug.....	2	1	seq
.....cugucauggaaauugcAcucuuuug.....	2	1	seq
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.....cugucauggaaCugcucucuuug.....	1	1	seq
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.....Augucauggaaauugcucucuuuug.....	1	1	seq
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dme-miR-281-2-5p MIMAT0000349

dme-miR-281-3p MIMAT0000345

cggaaauugugaaaugagagagcuauccgucgcacagucaaguuuaagaccgauuguaauacugucaugggaauiugcucucuuugguaaacauucg

.....ugucauggaGuugcucucuu..... 2 1
.....ugucauggaaauugcucCuuu..... 2 1
.....ugucauggaaauugcucucuu..... 27 0
.....ugucauggaaauugcucucuA..... 1 1
.....ugucauggaaAugcucucuu..... 1 1
.....ugucaGggaaauugcucucuuu..... 2 1
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.....ugucauggaaCugcucucuuu..... 2 1
.....ugGcauggaaauugcucucuuu..... 1 1
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.....ugucGuggaaauugcucucuuu..... 2 1
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.....ugucauggaaAuGcucucuuuug..... 1 1
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.....ugucauggaaauuugcucucuuuug..... 1 1
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.....ugucauggaaauugcucucuCuug..... 2 1
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dme-miR-281-2-5p_MIMAT0000349

dme-miR-281-3p_MIMAT0000345

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.....	ugucauggaaauugcucucuuAu.....	1	1	seq
.....	ugucauggaaauugcucucuuUu.....	3	1	seq
.....	ugucaAggaauugcucucuuug.....	2	1	seq
.....	ugucauggaaauugcuaUcuuuug.....	1	1	seq
.....	Cgucauggaaauugcucucuuug.....	1	1	seq
.....	ugucauggaaauugcucuuugA.....	1	1	seq
.....	ugucauggaaAugcucucuuuug.....	7	1	seq
.....	ugucauggaaauugcucuuuug.....	564	0	seq
.....	ugucauggaaauugcucuuCgu.....	3	1	seq
.....	ugucau <u>Cgaaauugcucucuuuug</u>	1	1	seq
.....	ugucauggaaauugcucuuuugC.....	44	1	seq
.....	uAucauggaaauugcucucuuuug.....	1	1	seq
.....	ugucauggaaauugcucuuuugCa.....	3	1	seq
.....	ugucauggaaauugcucuuuugAa.....	2	1	seq
.....	ugucauggaaauugcucuuuugua.....	9	0	seq
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.....	ugucauggaaauugcucuuuuguaA.....	6	1	seq
.....	Uucauggaaauugcucucuuu.....	1	1	seq
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.....	gucauggaaauugcucucuuuug.....	4	0	seq
.....	ucauggaaauugcucucuuuug.....	1	0	seq
.....	ucauggaaauugcucucuuuugua.....	2	0	seq
.....	auggaaauugcucucuuuAg.....	1	1	seq