

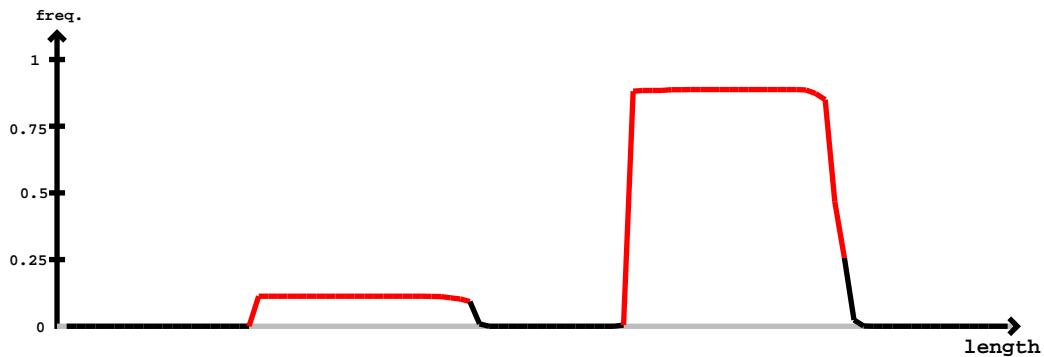
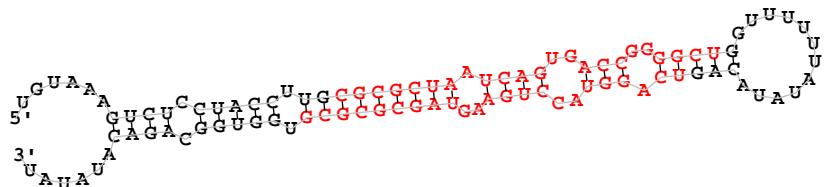
miRBase precursor : dme-mir-275_MI0000356

Total read count : 3118

dme-miR-275-3p MIMAT0000633 read count

dme-miR-275-5p MIMAT00239801 read count

remaining reads : 0



dme-miR-275-3p_MIMAT0000333

dme-miR-275-5p_MIMAT0020801			
	-3'	exp	
	reads	mm	sample
5'- uguuaagucuccuaccuugcgcgcuauacagugacccggggcugguuuuuuaauacag	ucaggauaccugaaguagcgcgcg	ugguggcagacauauau	
.....(((((((((.((((..(((.((.....))))))).))).))).))).)).).	
.....Acgcgcuaaucagugacccggggcu	1	1
.....cgcgcuauacagugacccg	1	0
.....cgcgcuauacagugacccg	1	1
.....cgcgcuauacagugacccg	2	0
.....cgcgcuauacagugacccg	1	1
.....cgcgcuauacagugacccg	1	1
.....cgcgcuauacagugacccg	13	0
.....cgcgcuauacagCgaccggg	1	1
.....cgcgcuauacagGgaccggg	1	1
.....cgcgcuauacagugacccggg	12	0
.....cgcgcuauacagugacccgggU	3	1
.....cgcgcuauacagugacccgggc	24	0
.....cgcgcuauacagugacccgggA	1	1
.....cgcgcuauacagugacccgggG	1	1
.....AgcgcuauacagugacccgggC	1	1
.....cgcgcuauacagugCccggggcu	1	1
.....cgcgcuauacaUugaccggggcu	1	1
.....cgcgcuauacagGgaccggggcu	2	1
.....cgcgcuauacagugacccggggA	1	1
.....cgcgcuauacagugacccggcu	1	1
.....Ugcgcuaaucagugacccggggcu	1	1
.....cgcgcuauacagugGccggggcu	1	1
.....cgcaAcuaaucagugacccggggcu	1	1
.....cgcgcuauacagugUccggggcu	1	1
.....cgcgcuauaUagugacccggggcu	1	1
.....cgcgcuauacagugacccggggcu	236	0
.....cgcgcuauacagugacccggggGu	2	1
.....cgcgcuauaAagugacccggggcu	1	1
.....cgcgcuauacagugacUggggcu	1	1
.....cgcgCaaucagugacccggggcu	1	1
.....cgcgcuauacagugacccggAgcu	1	1
.....cgcgcuauacagUUaccggggcu	1	1
.....cUgcgcuaaucagugacccggggcu	1	1
.....cgcgcuauacagugacccggggcu	2	1
.....cgcgcuauacagugacccggggcu	1	1

uguaaaagucuccuaccuuugcgcgcuauaucagugaccggggcugguuuuuuauauacagucagguaaccugaaguagcgcgcgugguggcagacauauau

.....cgcgcuauaucagCgaccggggcU.....	2	1	seq
.....cgcgcuauaucagugaccggggcU.....	11	1	seq
.....cgcgcuauaucagugaccggggcU.....	2	1	seq
.....cgcgcuauaucagugaccggggcU.....	15	1	seq
.....Aucagguaaccugaaguagcgcg.....	12	1	seq
.....Aucagguaaccugaaguagcgcgcg.....	1	1	seq
.....Cucagguaaccugaaguagcgcgcg.....	1	1	seq
.....ucagUuaccugaaguagc.....	1	1	seq
.....ucagguaaccugaaguagc.....	4	0	seq
.....Acagguaaccugaaguagcgcg.....	1	1	seq
.....ucagCuaccugaaguagcgcg.....	1	1	seq
.....ucagguaaccugaaguagcgcg.....	39	0	seq
.....ucagAuaccugaaguagcgc.....	2	1	seq
.....Acagguaaccugaaguagcgc.....	1	1	seq
.....ucCgguaaccugaaguagcgc.....	1	1	seq
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.....Ccagguaaccugaaguagcgc.....	1	1	seq
.....ucagguaaccugaauagcgc.....	1	1	seq
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.....ucagguaaccugaaguagAGcg.....	2	1	seq
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.....ucaUguaccugaaguagcgcg.....	7	1	seq
.....ucagguaaccugaaguagcgcU.....	1	1	seq
.....ucagguaaccCgaaguagcgcg.....	1	1	seq
.....uGagguaaccugaaguagcgcg.....	2	1	seq
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.....ucagguaaccugaaguaUcgcg.....	2	1	seq
.....ucagguaaccugaaguaAcgcg.....	1	1	seq
.....ucagCuaccugaaguagcgcg.....	2	1	seq
.....ucagguaaccuAaaguagcgcg.....	3	1	seq
.....ucagguaaccugaaguagUgcg.....	2	1	seq
.....ucaAguaaccugaaguagcgcg.....	7	1	seq
.....ucagguaaccugaagGaggcgcg.....	1	1	seq
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.....ucagguaaccugaagAagcgcg.....	3	1	seq
.....ucaggGaccugaaguagcgcg.....	6	1	seq
.....ucagguaaccugaCguagcgcg.....	1	1	seq
.....ucagguaaccugaaguagGGcg.....	2	1	seq
.....ucagguaUcugaaguagcgcg.....	3	1	seq
.....ucagUuaccugaaguagcgcg.....	2	1	seq
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.....ucaggAaccugaaguagcgcg.....	1	1	seq
.....ucagguaaccugaaguGgcgcg.....	3	1	seq
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.....uUagguaaccugaaguagcgcg.....	1	1	seq
.....ucagguaaccuUugaaguagcgcg.....	1	1	seq
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.....ucagguaaccugCaguagcgcgc.....	1	1	seq
.....Ccagguaaccugaaguagcgcgc.....	3	1	seq
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.....ucagguaaccugaauUaggcgcgc.....	1	1	seq
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dme-miR-275-5p_MIMAT0020801

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.....	ucagguaaccugaaguagcgcgc.....	2	1	seq
.....	ucagguaaccugaaguagcgcgc.....	4	1	seq
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.....	ucagguaaccugaaguagcgcgc.....	1	1	seq
.....	ucagguaaccCgaaguagcgcgc.....	1	1	seq
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.....	ucagguaaccugaaguagcgcgUg.....	1	1	seq
.....	ucagguaaccugaaguagcgcUcg.....	1	1	seq
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.....	ucagguaaccugaadGuagcgcgcg.....	2	1	seq
.....	ucaAguaccugaaguagcgcgcg.....	2	1	seq
.....	ucagguaaccugaaguagcgcgcgA.....	3	1	seq
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.....	ucagguaaccCgaaguagcgcgcg.....	3	1	seq
.....	ucGgguaaccugaaguagcgcgcg.....	2	1	seq
.....	ucagguaaccuCaaguagcgcgcg.....	1	1	seq
.....	ucagguaUcugaaguagcgcgcg.....	1	1	seq
.....	uAagguaaccugaaguagcgcgcg.....	1	1	seq
.....	ucagguaGccugaaguagcgcgcg.....	1	1	seq
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.....	ucagguaaccuAaaguagcgcgcg.....	2	1	seq
.....	Ccagguaaccugaaguagcgcgcg.....	1	1	seq
.....	Acagguaaccugaaguagcgcgcg.....	5	1	seq
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.....	ucagguaaccugaCguagcgcgcg.....	1	1	seq
.....	ucagguaaccugaaguagcgcgcU.....	2	1	seq
.....	ucagguaaccugaaguagcgcgcC.....	6	1	seq
.....	ucaAguaccugaaguagcgcgcu.....	1	1	seq
.....	ucaggGaccugaaguagcgcgcu.....	2	1	seq
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.....	ucagguaaccugaaguagcgcgcu.....	37	0	seq
.....	ucGgguaaccugaaguagcgcgcu.....	1	1	seq
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.....	Acagguaaccugaaguagcgcgcu.....	3	1	seq
.....	ucaCguaccugaaguagcgcgcu.....	1	1	seq
.....	ucagguaaccugaaguagcgcgcUu.....	1	1	seq

dme-miR-275-3p_MIMAT0000333

dme-miR-275-5p_MIMAT0020801

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.....cagguaaccugaaguagcgcg.....	3	0	seq
.....cagguaaccugaaguagcgcg.....	1	0	seq
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.....cagguaaccugaaguagcgcgug.....	1	1	seq
.....cagguaaccugaGguagcgcg.....	1	1	seq
.....cagguaaccugaaguagcgcg.....	1	0	seq
.....gguaccugaaguagcgcg.....	1	0	seq
.....guaccugaaguagcgcg.....	8	0	seq
.....accugaaguagcgcggu.....	2	0	seq