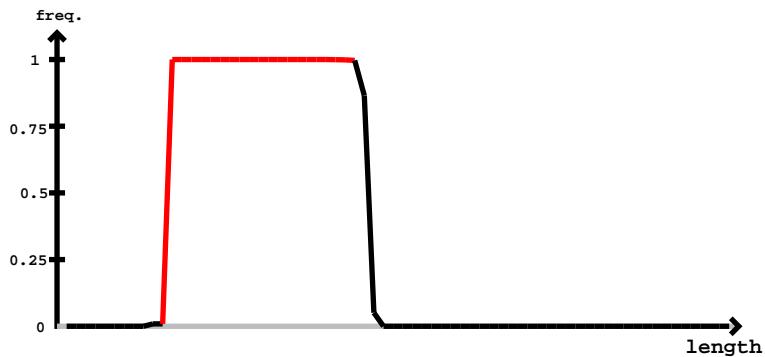
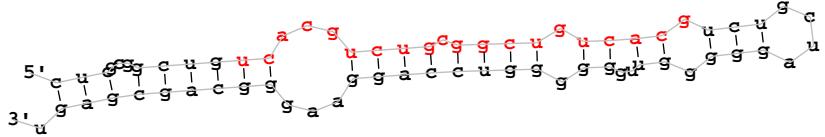


miRBase precursor : hsa-mir-11401
 Total read count : 1558
 hsa-miR-11401 read count : 1558
 remaining reads : 0



hsa-miR-11401

5' -		- 3'	exp	reads	mm	sample
(((.....(((((.....((((((.....))))....))))....))))....)).					
.....ucacgucugcggcuAucacg.....				1	1	Z3_-
.....ucacgucugcggcugucacA.....				2	1	Z3_-
.....ucacgucugcggcugucacg.....				17	0	Z3_-
.....ucacgucugcggcugucGgu.....				2	1	Z3_-
.....ucacgucAcgccugucacgu.....				1	1	Z3_-
.....ucacgucugcggcugucacgA.....				1	1	Z3_-
.....ucacgucugcggcugucacgC.....				8	1	Z3_-
.....uUacgucugcggcugucacgu.....				1	1	Z3_-
.....Ccacgucugcggcugucacgu.....				1	1	Z3_-
.....ucacgucugcggCgcucacgu.....				1	1	Z3_-
.....ucacgucugcggcugucacgG.....				1	1	Z3_-
.....ucacgucugcggcugucacgu.....				92	0	Z3_-
.....ucacgucugcggcugucacgA.....				4	1	Z3_-
.....ucacgucugcggcugucacgU.....				2	1	Z3_-
.....ucacgucugcggcugucacC.....				1	1	Z2_-
.....ucacgucugcggcuAucacg.....				1	1	Z2_-
.....ucacgucugcggUugucacg.....				2	1	Z2_-
.....ucacgucugcggcugucacA.....				2	1	Z2_-
.....ucacgucugcggcugucacU.....				3	1	Z2_-
.....ucacgucugcggcugucacg.....				20	0	Z2_-
.....ucacgucugcggcugacgG.....				1	1	Z2_-
.....ucGcgucugcggcugucacgu.....				1	1	Z2_-
.....uUacgucugcggcugucacgu.....				1	1	Z2_-
.....ucacgucugcggcugucacAu.....				1	1	Z2_-
.....ucacgucCcgccugucacgu.....				5	1	Z2_-
.....ucacgucugcggcugucacU.....				3	1	Z2_-
.....ucacgucugcUgcugucacgu.....				1	1	Z2_-
.....ucacgucugcggcugacgu.....				1	1	Z2_-
.....ucCcgucugcggcugucacgu.....				1	1	Z2_-
.....ucacgucugcggcugAcacgu.....				1	1	Z2_-
.....ucacgucugcggcugucacgC.....				17	1	Z2_-
.....ucacgucugcggcugucacgu.....				166	0	Z2_-
.....ucacgucugcggcugucacgU.....				5	1	Z2_-
.....ucacgucugcggcugucacgA.....				9	1	Z2_-

cucgcggcugucacgucugccggcugucacgucugcugaggguugggggguccaggaaaggcgagcggagu

.....ugucacgucugAggcuguc.....	1	1	GDM
.....ugucacgucugccggcuguc.....	1	0	GDM
.....ugucacgucugccggcugucacg.....	2	0	GDM
.....ucacgucugccggcugucacg.....	100	0	GDM
.....ucacgucugccggcugucacA.....	10	1	GDM
.....ucacgucugccggcugucacg.....	100	0	GDM
.....ucacgucugccggcugucacA.....	10	1	GDM
.....ucacgucugccggcugucacA.....	10	1	GDM
.....ucacgucugccggcugucacA.....	10	1	GDM
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.....ucacgucugccggcugucacg.....	1	1	GDM
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.....ucacgucCcgccugucacg.....	1	1	GDM
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.....ucacgucugccggAgucaacgu.....	1	1	GDM
.....Ccacgucugccggcugucacgu.....	4	1	GDM
.....ucacgucugccggcuguAacgu.....	1	1	GDM
.....ucacgucugccggCgucacgu.....	5	1	GDM
.....ucacgucugccggcugCcacgu.....	4	1	GDM
.....ucacgucugccggcugucaUgu.....	3	1	GDM
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.....ucaUgucugccggcugucacgu.....	2	1	GDM
.....ucacgucugccggcugucGogu.....	5	1	GDM
.....ucacgucugccggcugucacgA.....	19	1	GDM
.....ucacgucugccggcugucacgu.....	394	0	GDM
.....ucacgucugccggcugAcacgu.....	2	1	GDM
.....ucacgucugccggcugacgA.....	19	1	GDM
.....Acacgucugccggcugucacgu.....	1	1	GDM
.....ucGgucugccggcugucacgu.....	2	1	GDM
.....ucacgucugccggcugucacgC.....	43	1	GDM
.....ucacgucuAcgccugucacgu.....	2	1	GDM
.....ucacgucugccggcugucacgC.....	43	1	GDM
.....ucacgucugccggcugucaUgu.....	3	1	GDM
.....ucacgucugccggcugucacUu.....	2	1	GDM
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.....ucacgucCcgccugucacgu.....	3	1	GDM
.....ucacgucugccggcugacgC.....	43	1	GDM
.....ucacgucugccggcugucacgA.....	23	1	GDM
.....ucacgucugccggcugucacguU.....	7	1	GDM
.....ucacgucugccggcugucacguU.....	7	1	GDM
.....ucacgucugccggcugucacguA.....	23	1	GDM
.....ucacgucugccggcugucacguU.....	7	1	GDM
.....ucacgucugccggcugucacguA.....	23	1	GDM
.....ucacgucugccggcugucacguU.....	7	1	GDM
.....ucacgucugccggcugucacguA.....	23	1	GDM
.....ugucacgucugccggcugucacA.....	1	1	Z1_-
.....ugucacgucugccggcugucacg.....	4	0	Z1_-
.....ugucacgucugccggcugucacgu.....	3	0	Z1_-
.....ugAacacgucugccggcugucacgu.....	2	1	Z1_-
.....ucacgucugccggcugucaU.....	3	1	Z1_-
.....ucacgucugccggcugucacg.....	29	0	Z1_-
.....ucacgucugccggcugCcacg.....	1	1	Z1_-

Sequence	Count	Count	Count
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.....ucacgucugcgccugucacA.....	1	1	Z1_-
.....ucacgucugcgccugucGgu.....	2	1	Z1_-
.....ucacgucugcgccugucAgu.....	1	1	Z1_-
.....ucacgCcugcgccugucacgu.....	3	1	Z1_-
.....ucacgucugcgccugucacUu.....	1	1	Z1_-
.....ucacgucugcgccugCcacgu.....	1	1	Z1_-
.....ucacgucugUggcugucacgu.....	1	1	Z1_-
.....ucaUgucugcgccugucacgu.....	4	1	Z1_-
.....ucacgucugcgccCguacgu.....	2	1	Z1_-
.....ucGcgucugcgccugucacgu.....	1	1	Z1_-
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.....ucacgucugcgUcugucacgu.....	2	1	Z1_-
.....Ccacgucugcgccugucacgu.....	6	1	Z1_-
.....ucacgucuAcgccugucacgu.....	3	1	Z1_-
.....ucacgucugcgccugGcacgu.....	1	1	Z1_-
.....ucacgucugcgccugucacgA.....	13	1	Z1_-
.....ucacgucugcgccugucacgC.....	43	1	Z1_-
.....ucacgucugcgccugucacgAc.....	5	1	Z1_-
.....ucacgucugcgccugucacguA.....	24	1	Z1_-