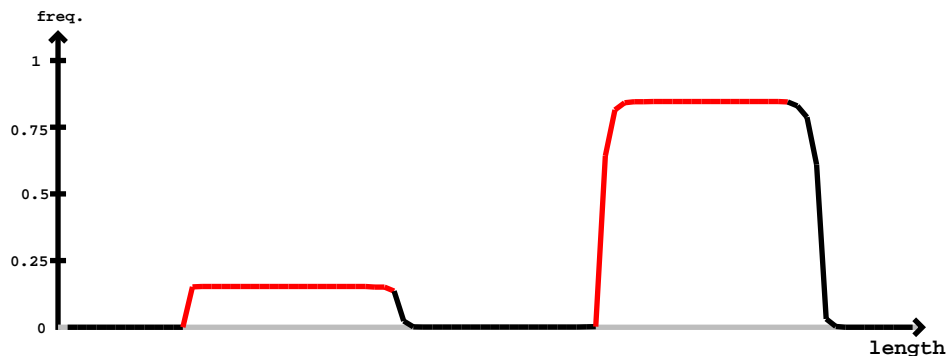
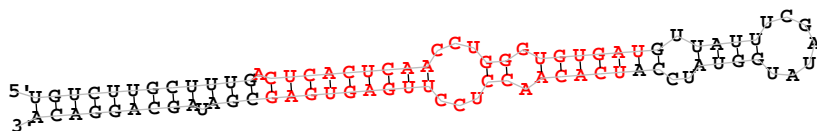


miRBase precursor : dme-mir-307a_MI0000418
 Total read count : 1275
 dme-miR-307a-3p_MIMAT0000398 read count
 dme-miR-307a-5p_MIMAT00020832 read count
 remaining reads : 1



dme-miR-307a-5p_MIMAT0020832

dme-miR-307a-3p_MIMAT0000398

5'				-3'	exp		
				reads	mm	sample	
5'-	ugucuuugcuuug <u>acucacucaaccuggggugugau</u> guuuuuucgauuagguaucca <u>uacaacaaccuccuugagugagc</u> gauagcaggaca						
.(((((((((((((((((((((((((((((.....)))))))).)))).)))).)))).)))).)))).)))).)))).)))).)))).)))).)))).				3	0	seq	
.....acucacucaaccuggggugu.....				1	1	seq	
.....acucacucaaccuCggguguga.....				18	0	seq	
.....acucacucaaccugggguguga.....				1	1	seq	
.....acucacucaaccuggggugGga.....				2	1	seq	
.....acucacucUaccuggggugugau.....				1	1	seq	
.....acucacucaaccuggggugugGu.....				1	1	seq	
.....acucacucaaccuggggugugCu.....				1	1	seq	
.....acucacucaaccAggggugugau.....				1	1	seq	
.....acucacucaaaAaccugggugugau.....				2	1	seq	
.....acucacucaaccuggggugugaA.....				1	1	seq	
.....acucacucaaccugggguguAAu.....				1	1	seq	
.....acucacucaaccugAgugugau.....				1	1	seq	
.....acucacucaaccugggugugUu.....				2	1	seq	
.....acucacucaaccuggggugugau.....				128	0	seq	
.....acucacGcaaccugggugugau.....				1	1	seq	
.....acucacucaaccuggggugugauA.....				1	1	seq	
.....acucacucaaccugggAugugaug.....				1	1	seq	
.....acucacucaaccugggugugauU.....				2	1	seq	
.....acucacucaaccuggggugugaug.....				23	0	seq	
.....acucacucaaccugAgugugaug.....				1	1	seq	
.....acucacucaaccugggugugaugu.....				1	0	seq	
.....cucacucaaccuggggugugau.....				2	0	seq	
.....uguuuuucgauuagguauccauc.....				1	0	seq	
.....Uaucacaaccuccuugagugag.....				1	1	seq	
.....aucacaaccuccuugagugagc.....				1	0	seq	
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.....ucacaaccuccuugagCgagc.....				1	1	seq	
.....ucacaaccuccuugagugagG.....				1	1	seq	
.....ucaGaacccuccuugagugagcg.....				1	1	seq	
.....ucacaaccuccuugagugUgcg.....				1	1	seq	

ugucuugcuuugacucaacucaaccugggugugauguuuuuucgauaugguauccaucacaaccuccuugagugagcgauagcaggaca

.....ucacaGccuccuugagugagcg.....	1	1	seq
.....uUacaaccuccuugagugagcg.....	1	1	seq
.....ucacaaccuccuugaAugagcg.....	2	1	seq
.....ucacaaccuccCugagugagcg.....	1	1	seq
.....ucacaaccuccuugagugagcg.....	194	0	seq
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.....cacaaccuccuugagugagcg.....	19	0	seq
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