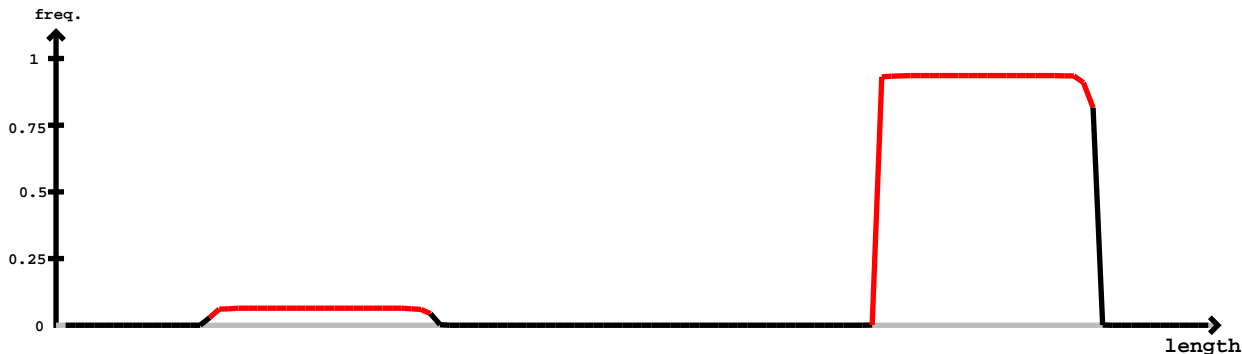
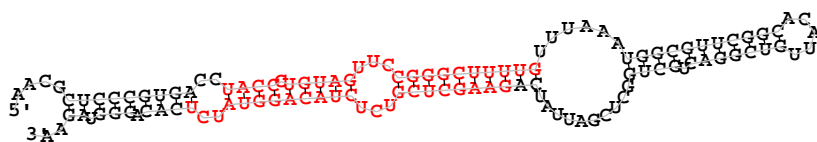


miRBase precursor : dme-mir-993\_MI0005854  
 Total read count : 875  
 dme-miR-993-5p\_MIMAT0020893 read count : 19  
 dme-miR-993-3p\_MIMAT0005512 read count : 685  
 remaining reads : 0



dme-miR-993-5p\_MIMAT0020893

dme-miR-993-3p\_MIMAT0005512

5'	reads	mm	exp	3'
aacgcuccgugaccuaccucguaguuccgggcuuuuguuuaaaggcgucgcacauugucggacucgucgucgauuaucagaagcucgucucacagguauucacagguagaa	2	0		-3'
.....cuaccucguaguuccgggcuuu.....	1	1		
.....cuaccucguaguuccgggcuuu.....	1	0		
.....cuaccucguaguuccgggcuuu.....	1	1		
.....cuaccucguaguuccgggcuuu.....	19	0		
.....uaccucguaguuccgggcuuu.....	2	0		
.....uaccucguaguuccgggcuuu.....	10	0		
.....uaccucguaguuccgggcuuu.....	1	1		
.....uaccucguaguuccgggcuuu.....	14	0		
.....uaccucguaguuccgggcuuu.....	1	1		
.....accucguaguuccgggcuuu.....	1	0		
.....cccucguaguuccgggcuuu.....	1	0		
.....cccucguaguuccgggcuuu.....	1	0		
.....agaagcucgucucacagguau.....	1	0		
.....gaagcucgucucacaggu.....	1	0		
.....gaagcucgucucacagguau.....	2	1		
.....gaagcucgucucacagguau.....	19	0		
.....gaagcucgucucacagguau.....	1	1		
.....gaagcucgucucacagguau.....	1	1		
.....gaagcucgucucacagguau.....	1	1		
.....gaagcucgucucacagguau.....	1	1		
.....gaagcucgucucacagguau.....	3	1		
.....gaagcucgucucacagguau.....	2	1		
.....gaagcucgucucacagguau.....	1	1		
.....aaagcucgucucacagguau.....	1	1		
.....gaagcucgucucacagguau.....	1	1		
.....gaagcucgucucacagguau.....	68	0		
.....gGagcucgucucacagguau.....	1	1		
.....gaagcucgucucacagguau.....	3	1		
.....gaagcucgucucacagguau.....	1	1		
.....gaagcucgucucacagguau.....	13	1		
.....gaagcucgucucacagguau.....	2	1		
.....gaagcucgucucacagguau.....	1	1		

aaagcucccgugaccuacccugaguuccgggcuuuuguuuaaaggcgguccgacacauugucggacucgucggcucgauuauca

.....gaagcucgucucCacagguaucu.....	5	1	seq
.....gaagcucgucucacaggGaucu.....	1	1	seq
.....gaagcucgucucacagAuauc.....	1	1	seq
.....Aaagcucgucucacagguaucu.....	1	1	seq
.....gaagcucgucucacaUguauc.....	1	1	seq
.....gaagcucgucucCcuacagguaucu.....	2	1	seq
.....gaagcucUucucacagguaucu.....	1	1	seq
.....gaagcAcgucucacagguaucu.....	1	1	seq
.....gaCgucgucucacagguaucu.....	1	1	seq
.....gaagcucgGcucacagguaucu.....	1	1	seq
.....gaagcucgucucacagguGcu.....	2	1	seq
.....gaagcucgucucacaggCauc.....	2	1	seq
.....gaagcucgucucacaggAauc.....	2	1	seq
.....gaagcucgucucacaCguauc.....	4	1	seq
.....gaagcGcgucucacagguaucu.....	3	1	seq
.....gaagcucgucucacagguCcu.....	2	1	seq
.....gaagcucgucucAacagguaucu.....	2	1	seq
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.....gaagcucgucucCacagguaucu.....	1	1	seq
.....gaagcucgucucacagguauC.....	1	1	seq
.....gaagcucgucucacGguauc.....	1	1	seq
.....gUagcucgucucacagguaucu.....	1	1	seq
.....gaagcucgucucacagguauU.....	4	1	seq
.....gaagcucgucucacGguauc.....	3	1	seq
.....gaagcucgucucacagguCuc.....	1	1	seq
.....gaagcucgucucacagguauU.....	1	1	seq
.....aagcucgucucacagguaucu.....	2	0	seq
.....agcucgucucacagguaucu.....	1	0	seq
.....gcucgucucacagguauc.....	1	0	seq