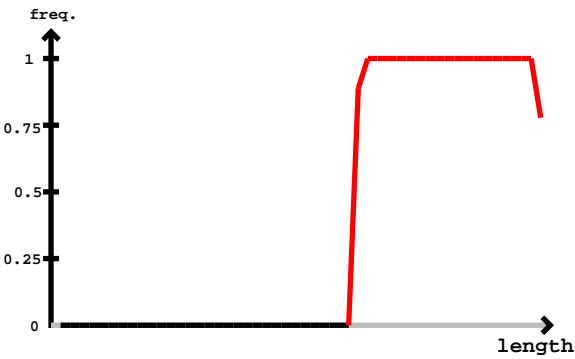
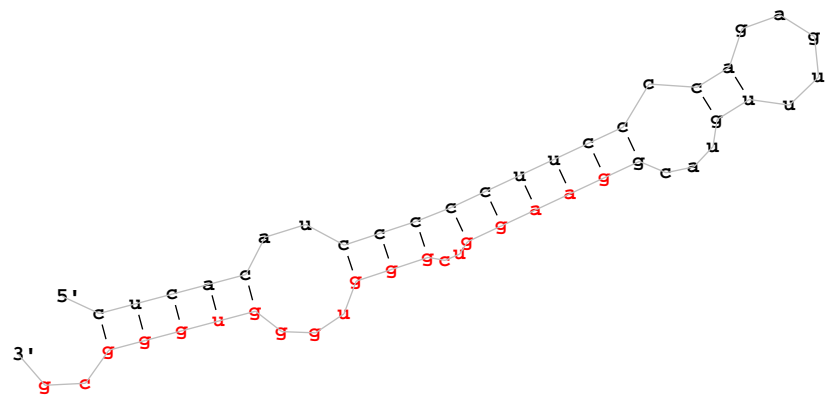


miRBase precursor : chromosome\_2\_4981  
Total read count : 9  
chromosome\_2\_4981 read count : 9  
remaining reads : 0



chromosome_2_4981			
5'- cucacaucccccuccccagaguuguacggaaggucgggugggugggcg -3'	exp		
((((((.....)))))).....	reads	mm	sample
.....Aaaggucgggugggugggc.	2	1	seq
.....Aaaggucgggugggugggcg	6	1	seq
.....aaggucgggugggugggcg	1	0	seq