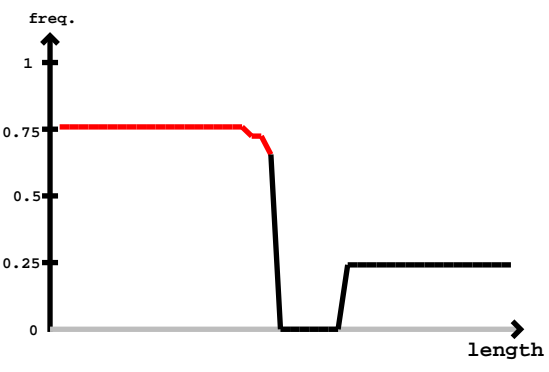
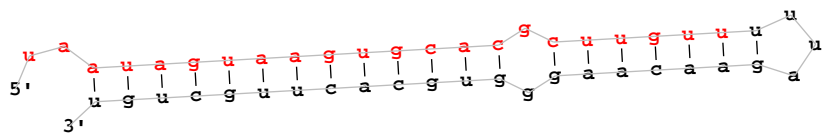


miRBase precursor : chromosome\_1\_216  
Total read count : 29  
chromosome\_1\_216 read count : 29  
remaining reads : 7



chromosome_1_216				
5'-	uaauaguaagucacgcuguuuuagaacaaggugcacuugcugu	-3'	exp	
	..((((((((((((((((((((((.....))))))))))))))))))	reads	mm	sample
	uaauaguaagucacgcuu.....	1	0	seq
	uaauaguaagucacgcugu.....	2	0	seq
	uaauaguaagucacgcuguuu.....	16	0	seq
	uaauaguaagucacgcuguuA.....	2	1	seq
	uaauaguaagucacgcuguuC.....	1	1	seq
	.....caaggugcacuugcugA	4	1	seq
	.....caaggugcacuugcugu	3	0	seq