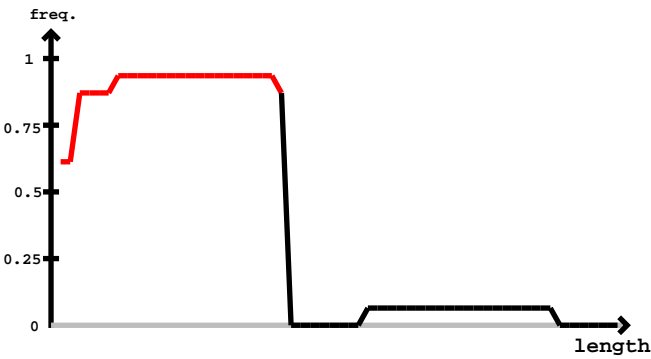
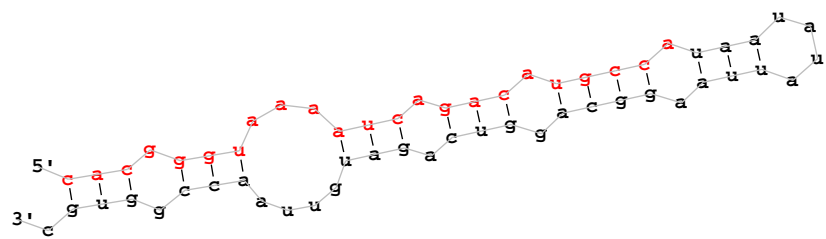


miRBase precursor : chromosome\_3\_7672  
Total read count : 31  
chromosome\_3\_7672 read count : 31  
remaining reads : 2



chromosome\_3\_7672

5'-	cacggguaaaaucagacaugcca	uaauauuuuaggcaggucagauguuaaccggugc	-3'	exp	
	(((((.....(((.....(((.....(((.....))))).))))).))))).))))).		reads	mm	sample
	Aacggguaaaaucagacaugcca.....		17	1	seq
	Gacggguaaaaucagacaugcca.....		2	1	seq
	.acggguaaaaucagacaugcc.....		1	0	seq
	.Gcggguaaaaucagacaugcc.....		1	1	seq
	.Gcggguaaaaucagacaugcca.....		1	1	seq
	.acggguaaaaucagacaugcca.....		5	0	seq
	....Aaaaaucagacaugcca.....		1	1	seq
	....guaaaaucagacaugcca.....		1	0	seq
	.....aaaggcaggucagauguua.....		1	1	seq
	.....uaaggcaggucagauguua.....		1	0	seq