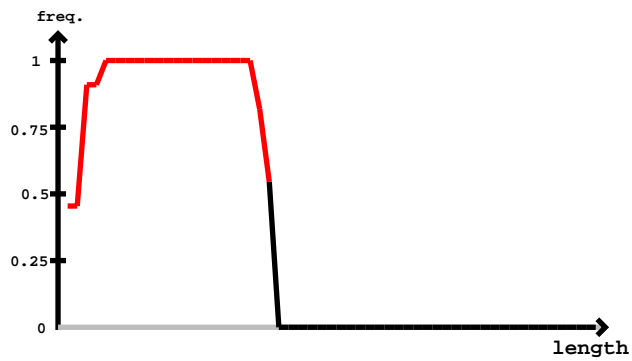
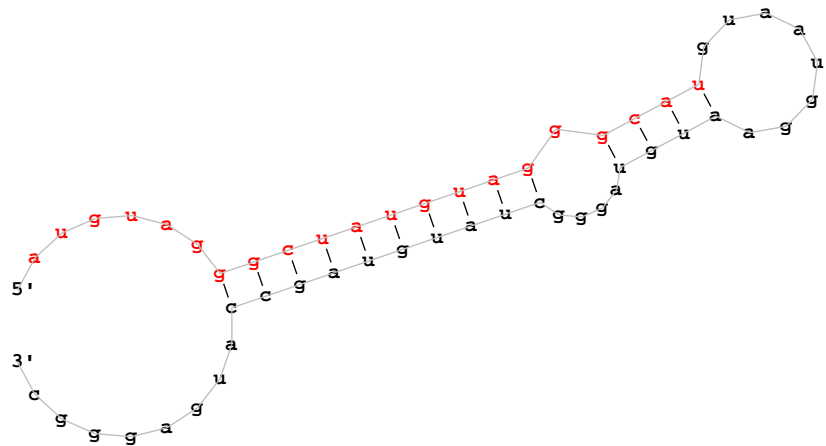


miRBase precursor : chromosome\_3\_8854  
 Total read count : 11  
 chromosome\_3\_8854 read count : 11  
 remaining reads : 0



chromosome_3_8854			
5'-	auguaggggcuau <u>guaggggca</u> guaauggaauguaggggcuau <u>guagcc</u> augagggc	-3'	exp
	.....(((((((((((.....)))))).)))))).....	reads	mm
	auguaggggcuau <u>guaggggca</u> .....	5	0
	. <u>guaggggcuau</u> aggggU.....	2	1
	. <u>guaggggcuau</u> agggAca.....	2	1
	. <u>guaggggcuau</u> aggggca.....	1	0
	... <u>uaggggcuau</u> aggggcau.....	1	0
		seq	seq