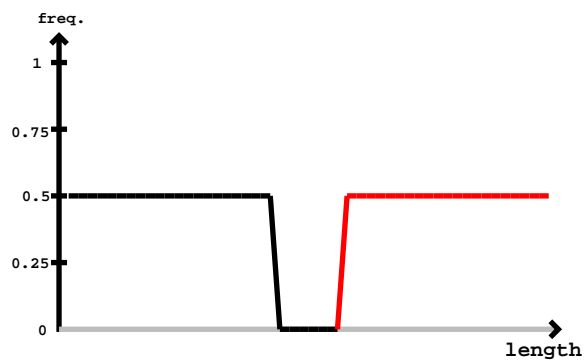
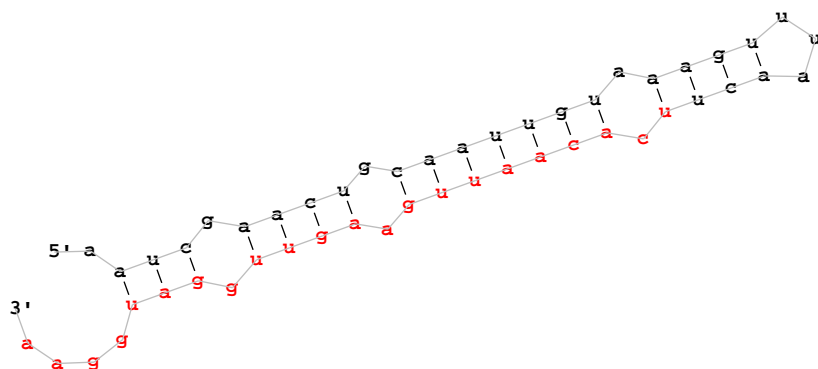


```

miRBase precursor      : chromosome_3_8020
Total read count       : 2
chromosome_3_8020 read count
remaining reads        : 1

```



chromosome_3_8020					
5'-	aau	cgaacugcaaauguuaaguuuaacuucaca	aaugaaguuggauggaa	-3'	exp
	.(((.((((.((((.((((.....)))))))))..))))).))....	reads	mm	sample	
	aaucgaacugcaaauguuaag.....	1	0	seq	
ucacaaCugaaguuggauggaa	1	1	seq	