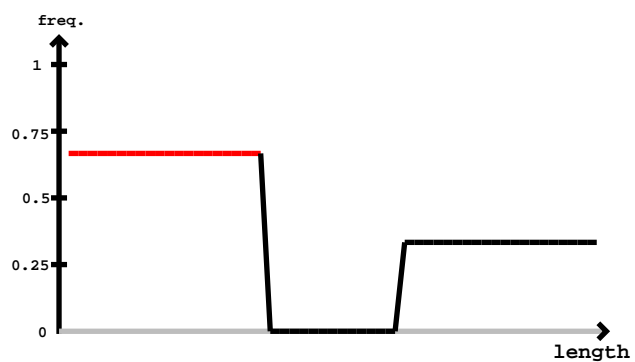
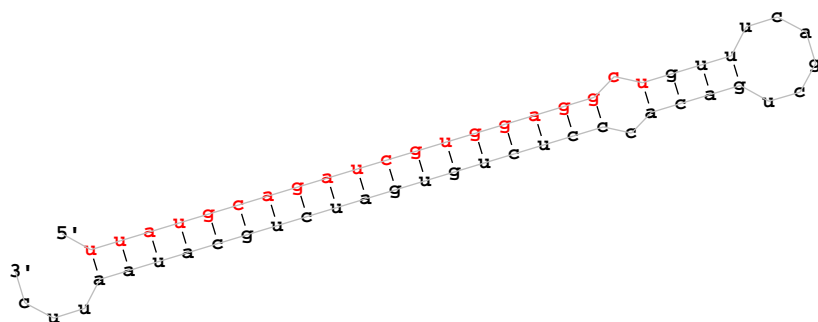


```

miRBase precursor      : chromosome_12_62633
Total read count       : 3
chromosome_12_62633 read count
remaining reads        : 1

```



chromosome_12_62633			
5'-		-3'	exp
uuauugcagauugcuggaggcguuuucagcugacacccucugugaucugcauaauuc	reads	mm	sample
(((((((((((((((((((((((.....)))))))))))))))))).....	2	0	seq
uuauugcagauugcuggaggcu.....	1	0	seq
.....ccucugugaucugcauaauuc			