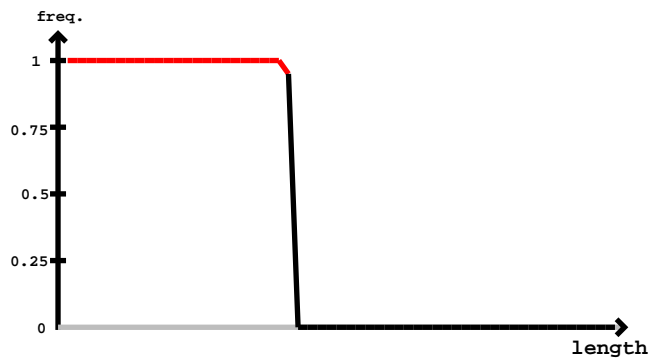
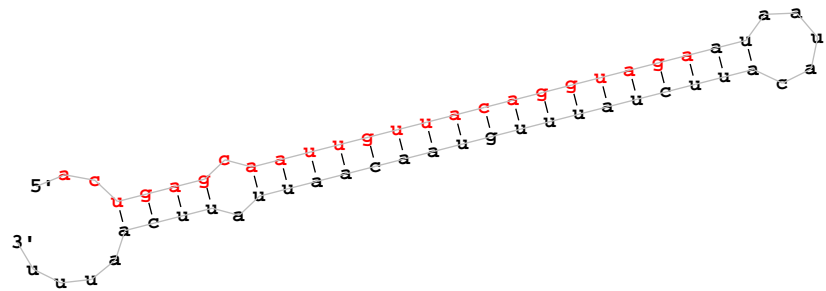


miRBase precursor : chromosome_12_62475
 Total read count : 20
 chromosome_12_62475 read count : 20
 remaining reads : 0



chromosome_12_62475			
5'-		-3'	exp
acugagcaauuguuacagguaga	..(((.((((((((((((((((((((.....))))))))))))))))))....	reads	mm
acugagcaauuguuacagguag	1	1
acugagcaauuguuacagguaga	11	1
acugagcaauuguuacagguaga	4	0
Gcugagcaauuguuacagguaga	3	1
acCgagcaauuguuacagguaga	1	1
			sample
			seq
			seq
			seq
			seq
			seq