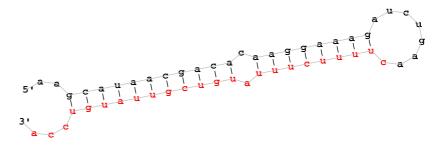
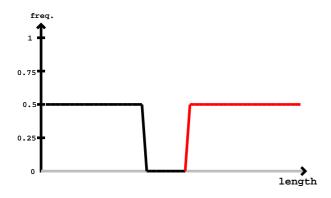
miRBase precursor : chromosome_1_784

Total read count : 2 chromosome_1_784 read count remaining reads : 1





chromosome_1_784

5'- aagcauaacgacacaaggaaagaucugaacuuuucuuuaugucguuaugucca	-3' exp		
(((((((((((((((((((((((((((((((((((((reads	mm	sample
aagcauaacgacacaaggaa	1	0	seq
cuuuucuuuaugucguuaugucca	1	0	seq