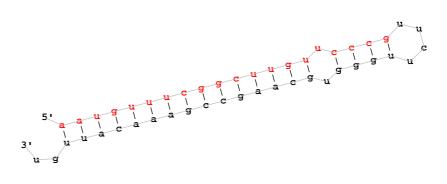
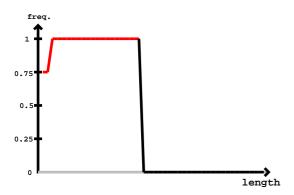
miRBase precursor : chromosome_12_57430

Total read count : 4 chromosome_12_57430 read &ount remaining reads : 0





chromosome_12_57430

5'- aauguuucggcuuguucccguucuugggugcaagccgaaacauugu	-3' exp		
(((((((((((((((((((((((((((((((((((((((reads	mm	sample
aauguuucggcuuguucccg	3	0	seq
.auguuucggcuuguucccg	1	0	seq