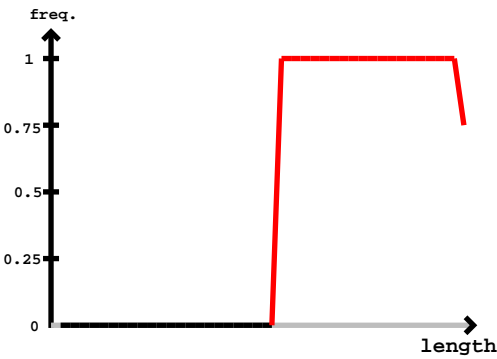
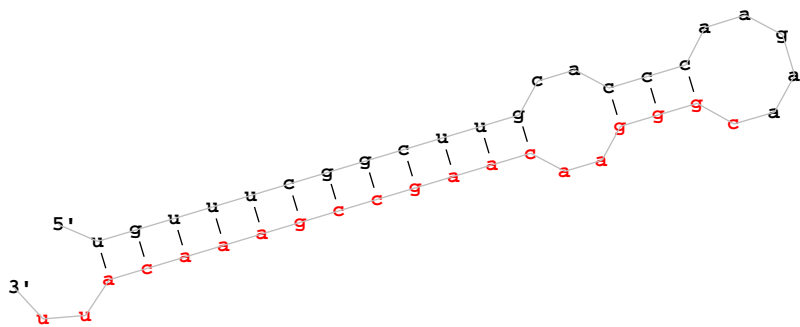


miRBase precursor : chromosome_12_61354
Total read count : 4
chromosome_12_61354 read count : 4
remaining reads : 0



chromosome_12_61354				
5'-	uguuucggcuugcacccaagaac	cg	ggaacaagccgaaacauu	-3'
(((((((((((.....))))))))))..		reads	mm	sample
.....cggaacaagccgaaacau.		1	0	seq
.....cggaacaagccgaaacauu		3	0	seq