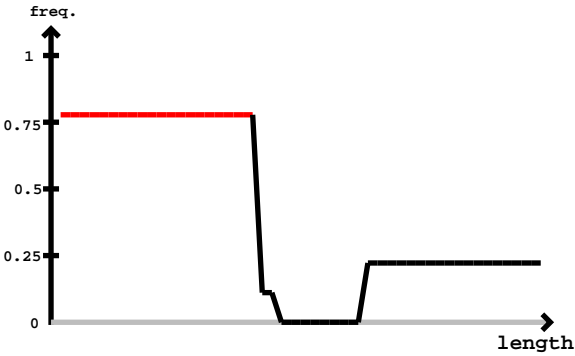
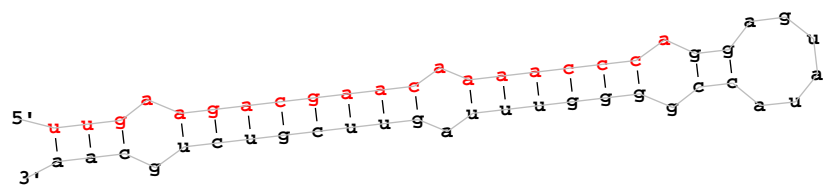


miRBase precursor : chromosome_12_59002
Total read count : 9
chromosome_12_59002 read count : 9
remaining reads : 2



chromosome_12_59002				
5'-	uugaagacgaacaaaaccca	ggaguauaccgggguuuaguucgucugcaa	-3'	exp
	(((.(((((((.(((((((.....)))))))).)))))).)).	reads	mm	sample
	uugCagacgaacaaaaccca.....	6	1	seq
	uugaagacgaacaaaacccagg.....	1	0	seq
ggguuuaguucgucugcaa	2	0	seq