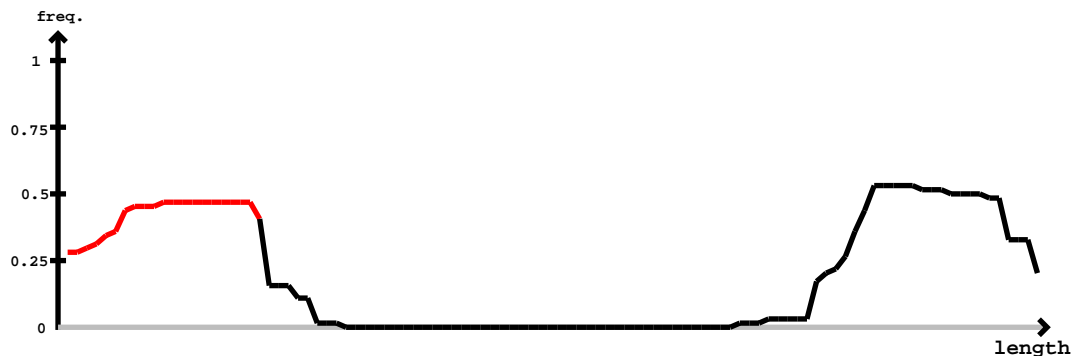
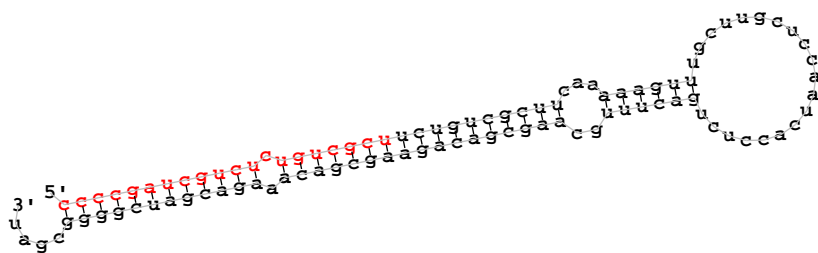


miRBase precursor : chromosome_7_31536
Total read count : 64
chromosome_7_31536 read count : 20
remaining reads : 35



chromosome_7_31536

5'-	reads	mm	sample
ccccgaucgucucugucguuucugucgcuucaaaaaguugcuugcuccaaucaccucugacuuugcaagcgacagaagcgacaaaagacgaucggggcgau	2	1	seq
Ucccgauucgucucugucgc.....	2	0	seq
ccccgaucgucucugucgc.....	1	1	seq
Ucccgauucgucucugucgc.....	2	1	seq
ccccgaucgucucugucgcA.....	10	0	seq
ccccgaucgucucugucgc.....	1	1	seq
.ccccgaucgucucugucgcA.....	1	1	seq
.ccgaucgucucugucgc.....	1	0	seq
..Ugaucgucucugucgcuuuc.....	2	1	seq
...Aaucgucucugucgcuuuc.....	1	1	seq
....aucgucucugucgcCucugu.....	5	1	seq
.....ucgucucugucgcuuucugu.....	1	0	seq
.....ucuUugucgcuuucugucgc.....	1	1	seq
.....gcgacagaagcgacaaaaga.....	1	0	seq
.....acagaagcgacagagacga.....	1	1	seq
.....agcgacagagacgaucgg.....	1	1	seq
.....agcgacagagacgaucgggg.....	8	1	seq
.....gcgacagagacgaucgggg.....	2	1	seq
.....cgacagagacgaucggggcgga.....	1	1	seq
.....gacagagacgaucggggcgga.....	3	1	seq
.....acaGagacgaucggggcgga.....	2	1	seq
.....acaGagacgaucggggcgau.....	4	1	seq
.....caGagacgaucggggcgga.....	2	1	seq
.....caGagacgaucggggcgau.....	3	1	seq
.....aaagacgaucggggcgau.....	2	0	seq
.....aGagacgaucggggcgau.....	4	1	seq