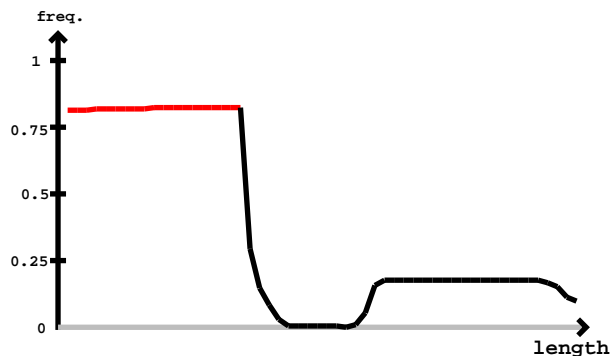
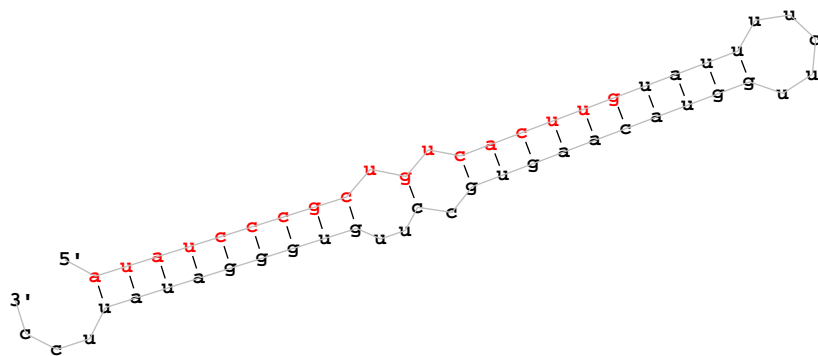


miRBase precursor : chromosome_12_56606
 Total read count : 204
 chromosome_12_56606 read length : 67nt
 remaining reads : 37



chromosome_12_56606			
5'-		-3'	exp
a-u-a-u-c-c-g-c-u-g-u-a-u-u-u-u-u-g-g-u-a-c-a-a-g-u-g-c-c-u-g-g-g-a-u-a-u-u-c	reads	mm	sample
U-u-a-u-c-c-g-c-u-g-u-a-u-g.....	1	1	seq
a-u-a-u-c-c-g-c-u-g-u-a-u-A.....	29	1	seq
a-u-a-u-c-c-U-g-c-u-g-u-a-u-g.....	1	1	seq
G-u-a-u-c-c-g-c-u-g-u-a-u-g.....	1	1	seq
a-u-a-u-c-c-g-c-u-g-u-a-u-g.....	61	0	seq
a-u-a-u-c-c-G-g-c-u-g-u-a-u-g.....	1	1	seq
a-u-a-u-c-c-g-c-u-g-u-a-u-U.....	14	1	seq
a-u-a-u-c-c-g-c-u-g-u-a-u-U-u.....	4	1	seq
a-u-a-u-c-c-g-c-u-g-u-a-u-g-A.....	5	1	seq
a-u-a-u-c-c-g-c-u-g-u-a-u-g-A-g-u.....	1	1	seq
a-u-a-u-c-c-g-c-u-g-u-a-u-g-u.....	12	0	seq
a-u-a-u-c-c-g-c-u-g-u-a-u-g-u-A.....	7	1	seq
a-u-a-u-c-c-g-c-u-g-C-c-a-u-g-u.....	1	1	seq
a-u-a-u-c-c-g-c-u-g-u-a-u-g-u-a.....	7	0	seq
G-u-a-u-c-c-g-c-u-g-u-a-u-g-u-a.....	1	1	seq
a-u-G-u-c-c-g-c-u-g-u-a-u-g-u-a.....	1	1	seq
a-u-a-u-c-c-g-c-u-g-u-a-u-U-u-a.....	1	1	seq
a-u-a-u-c-c-g-c-u-g-u-a-u-g-u-U.....	3	1	seq
C-u-a-u-c-c-g-c-u-g-u-a-u-g-u-a-u.....	1	1	seq
a-u-a-u-c-c-g-c-u-g-u-a-u-g-u-a-C.....	3	1	seq
a-u-a-u-c-c-g-c-u-g-u-a-u-g-u-a-u.....	7	0	seq
a-u-a-u-c-c-g-c-u-g-u-a-u-g-u-a-u-A.....	2	1	seq
a-u-a-u-c-c-g-c-u-g-u-a-u-g-u-a-u-A-u.....	1	1	seq
a-u-a-u-c-c-g-c-u-g-u-a-u-g-u-a-u-C.....	1	1	seq
..a-u-c-c-g-c-u-g-u-a-u-g-u-a-u-A-u.....	1	1	seq
.....U-g-u-a-c-u-g-u-a-u-u-u-u-c-u-g.....	1	1	seq
.....u-a-c-a-a-g-u-g-c-c-u-g-u-g-g-g-a-u-a-u-u-c-A.....	2	1	seq
.....a-c-a-a-g-u-g-c-c-u-g-u-g-g-g-a-u-a.....	2	0	seq
.....a-c-a-a-g-u-g-c-c-u-g-u-g-g-g-a-u-a-u.....	2	0	seq
.....a-c-a-a-g-u-g-c-c-u-g-u-g-g-g-a-u-a-u-u.....	2	0	seq
.....a-c-a-a-g-u-g-c-c-u-g-u-g-g-g-a-u-a-u-u-c.....	1	0	seq
.....a-c-a-a-g-u-g-c-c-u-g-u-g-g-g-a-u-a-u-u-c-A.....	2	1	seq
.....c-a-a-g-u-g-A-c-u-g-u-g-g-g-a-u-a-u.....	1	1	seq
.....c-a-a-g-u-g-c-c-u-g-u-g-g-g-a-u-a-u-u.....	5	0	seq
.....c-a-a-g-u-g-A-c-u-g-u-g-g-g-a-u-a-u-u.....	1	1	seq

auaucccgcuacacuuuguauuuucuugguacaagugccuugugggauauucc			
.....caagugccuugugggauauuc.	1	0	seq
.....caagugAcuugugggauauuc.	1	1	seq
.....caagugccuugugggauauucA	6	1	seq
.....caagugccuugugggauauucU	1	1	seq
.....caagugccuugugggauauucc	5	0	seq
.....aagugAcuugugggauauucc	1	1	seq
.....aagugccuugugggauauucA	3	1	seq