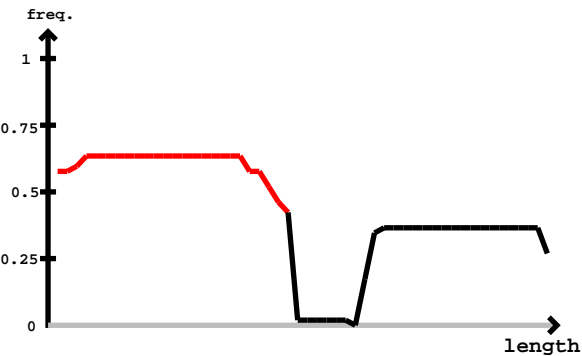
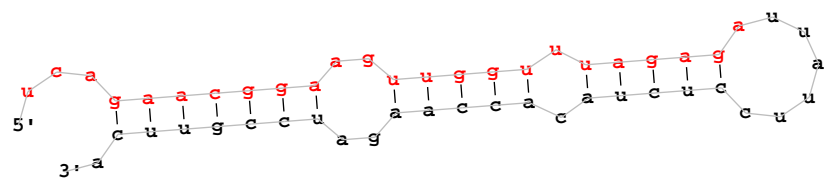


miRBase precursor : chromosome\_4\_12786  
Total read count : 52  
chromosome\_4\_12786 read count : 30  
remaining reads : 20



chromosome\_4\_12786

5'-	ucagaacggaaguugguuuagagauuauuccucacaccaagauccguuca	-3'	exp		
...	((((((((.....)))))).))))).))))).	reads	mm	sample	
ucagaacggaaguugguuu.....		3	0	seq	
ucagaacggaaguugguuuag.....		3	0	seq	
ucagaacggaaguugguuuagag.....		3	0	seq	
ucagaacggaaguugguuuagag.....		2	0	seq	
ucagaacggaaguugguuuagagC.....		2	1	seq	
ucagaacggaaguugguuuagagU.....		10	1	seq	
ucagaacggaaguugguuuagaga.....		4	0	seq	
ucagaacggaaguugguuuagagUa.....		2	1	seq	
ucagaacggaaguugguuuagagauuauuc.....		1	0	seq	
.cagaacggaaguugguuuagagU.....		1	1	seq	
.agaacggaaguugguuuagaga.....		1	0	seq	
.agaacggaaguugguuuagagU.....		1	1	seq	
.....ucacaccaagauccguuU.....		2	1	seq	
.....ucacaccaagauccguuU.....		1	1	seq	
.....ucacaccaagauccguuca.....		1	1	seq	
.....ucacaccaagauccguuca.....		5	0	seq	
.....cuacaccaagauccguuU.....		1	1	seq	
.....cuacaccaagauccguuc.....		2	0	seq	
.....cuacaccaagauccguuca.....		5	0	seq	
.....cuacaccaagauccguucG.....		1	1	seq	
.....uacaccaagauccguuca.....		1	0	seq	