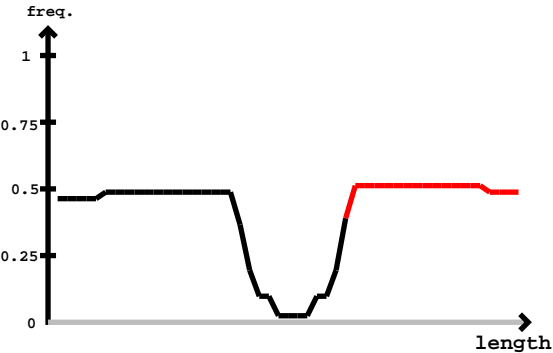
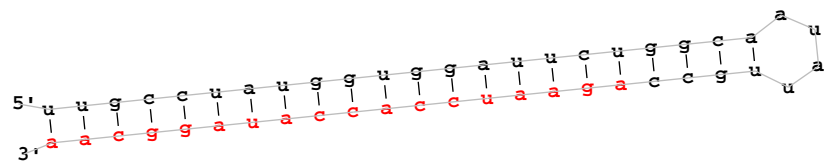


miRBase precursor : chromosome_1_1120
Total read count : 41
chromosome_1_1120 read count : 41
remaining reads : 24



chromosome_1_1120			
5'-	uugccuaugguggauucuggc	aaauugcc	agaauccaccuaggcaa
-3'	exp	reads	mm
	sample		
	seq	4	0
	seq	1	1
	seq	2	1
	seq	1	1
	seq	4	0
	seq	4	0
	seq	3	0
	seq	1	0
	seq	1	0
	seq	3	0
	seq	3	0
	seq	1	1
	seq	4	0
	seq	3	1
	seq	1	1
	seq	5	0