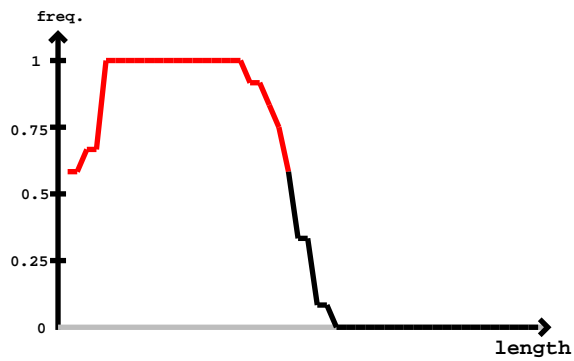
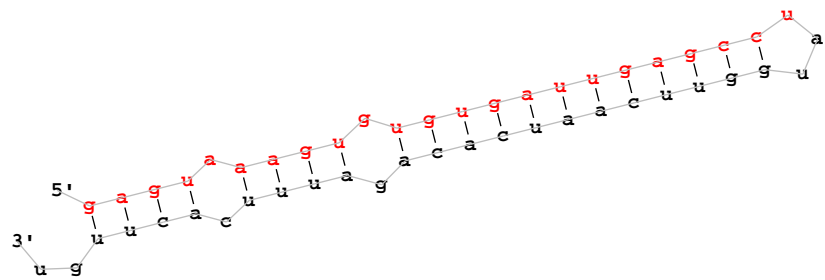


miRBase precursor : chromosome_1_1442
 Total read count : 12
 chromosome_1_1442 read count : 12
 remaining reads : 0



chromosome_1_1442

5'-	-3'	exp		
gaguaaaagugugauugagccuaugguucaaucagauuucacuugu			reads	mm
(((.((((.((((.((((.(...)))))))))).))))..				sample
gaguaaaagugugauug.....	1	0		seq
gaguaaaagugugauugagc.....	1	0		seq
gaguaaaagugugauugagcca.....	2	1		seq
gCguaaaagugugauugagccuau.....	1	1		seq
gaguaaaagugugauugagccuaC.....	1	1		seq
gaguaaaagugugauugagccuaugC.....	1	1		seq
.aguaaaagugugauugag.....	1	0		seq
...uaaaagugugauugagcA.....	2	1		seq
...uaaaagugugauugagccu.....	1	0		seq
...uaaaagugugauugagccuau.....	1	0		seq