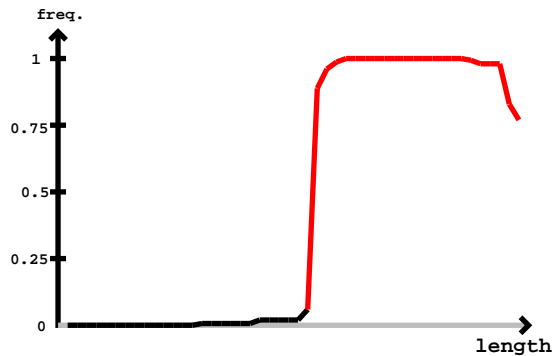
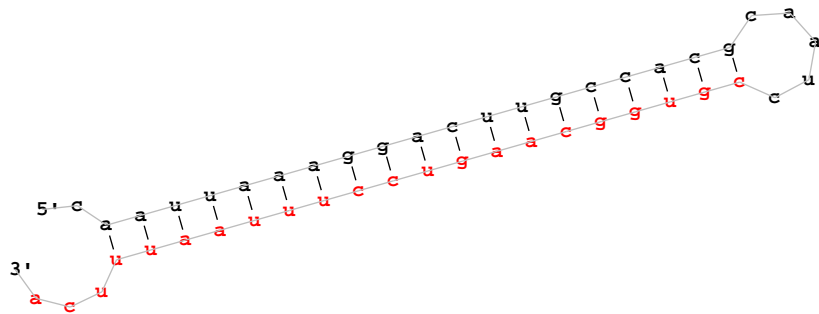


miRBase precursor : chromosome\_2\_5850  
 Total read count : 152  
 chromosome\_2\_5850 read count : 149  
 remaining reads : 3



chromosome_2_5850			
5'-		-3'	exp
caauuaaggacuugccacgcaaucguggcaaguccuuuaauuua	reads	mm	sample
.((((((((((((((((((((.....))))))))))))))....	1	0	seq
.....ugccacgcaauccguggcaaguccuuua.....	2	1	seq
.....Acaauccguggcaaguccuuua.....	2	1	seq
.....Ucuggcaaguccuuuaauuua	4	1	seq
.....Acguggcaaguccuuuaauuua	1	1	seq
.....cguggcaaguGcuuaauuu..	18	0	seq
.....cguggcaaguccuuuaauuu..	2	1	seq
.....Uguggcaaguccuuuaauuu..	1	1	seq
.....Aguggcaaguccuuuaauuu..	2	1	seq
.....Uguggcaaguccuuuaauuuc.	5	0	seq
.....cguggcaaguccuuuaauuuc.	1	1	seq
.....cguggcaaguccuuuaauGuca	1	1	seq
.....cguggUaaguccuuuaauuua	1	1	seq
.....cguggcGaguccuuuaauuua	1	1	seq
.....cguggcaaguUcuuaauuuua	1	1	seq
.....cgAggcaaguccuuuaauuua	2	1	seq
.....cguggcaaguccuuuaauuua	55	0	seq
.....cguggcaaguccuuuaauuuUa	1	1	seq
.....Uguggcaaguccuuuaauuua	27	1	seq
.....cguggcaaguccuuuaauuucU	1	1	seq
.....cguggcaaguccuuuUauuuua	1	1	seq
.....Aguggcaaguccuuuaauuua	2	1	seq
.....cguggcaaguccuuuCaauuua	1	1	seq
.....cguggcaaguccuuuaauuucG	1	1	seq
.....cguggcaaguccuCuauuuua	2	1	seq
.....guggcaaguccuuuaauuu..	1	0	seq
.....guggcaaguccuuuaauuuc.	2	0	seq
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.....guggcaaguccuuuaauuua	7	0	seq
.....uggcaaguccuuuaauuua	4	0	seq
.....ggcaaguccuuuaauuua	2	0	seq