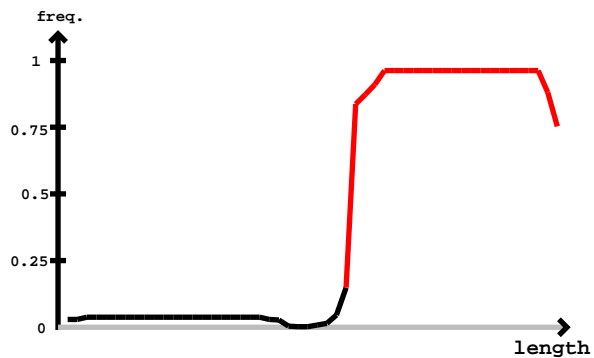
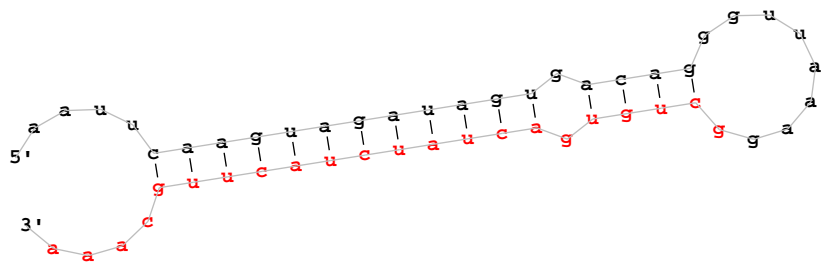


miRBase precursor : chromosome\_5\_16991  
 Total read count : 478  
 chromosome\_5\_16991 read count : 451  
 remaining reads : 25



chromosome_5_16991			
5'-	reads	mm	sample
aaaucaaguagauagugacaggguaaaagcgugugacuaucuaucugcaaa	1	1	seq
....((((((((((((((((.....))))))))))))))....	1	0	seq
aaaucaaaAuagauagugacag.....	5	1	seq
aaaucaaguagaCagugacagg.....	1	0	seq
Uauucaaguagauagugacagg.....	2	1	seq
Gauucaaguagauagugacagg.....	1	1	seq
aaaucaaaAuagauagugacagg.....	2	1	seq
aaaucaaguagaCagugacagg.....	1	1	seq
.auucaaguagauagugaca.....	3	0	seq
.Guucaaguagauagugaca.....	1	1	seq
.....Uguuaaaggcgugugacuaucuaucugca..	1	1	seq
.....aaaggcgugugacuaucuaucugca..	2	0	seq
.....aaaggcgugugacuaucuaucugcaaa	1	0	seq
.....aUggcgugugacuaucuaucugcaaa	3	1	seq
.....Uggcgugugacuaucuaucugca..	4	1	seq
.....aggcgugugacuaucuaucugcaa..	1	0	seq
.....aggcgugugacuaucuaucugcaaU	3	1	seq
.....aggcgugugacuaucuaucugcaaa	1	0	seq
.....Uggcgugugacuaucuaucugcaaa	5	1	seq
.....Gggcgugugacuaucuaucugcaaa	1	1	seq
.....ggcgugugacuaucuaucugca..	2	0	seq
.....Agcgugugacuaucuaucugca..	2	1	seq
.....ggcgugugacuaucuaucugcaa..	1	0	seq
.....Agcgugugacuaucuaucugcaa..	1	1	seq
.....Cgcgugugacuaucuaucugcaaa	1	1	seq
.....ggcgugugGcuaucaucugcaaa	2	1	seq
.....ggcgugugacuaucuaucugcaaU	18	1	seq
.....ggcgugugacuaucuaucugcaaa	15	0	seq
.....Agcgugugacuaucuaucugcaaa	7	1	seq
.....Acugugacuaucuaucugca..	1	1	seq
.....Ucugugacuaucuaucugca..	1	1	seq
.....gcuugugacuUucuaucugca..	1	1	seq
.....gcuugugacuaucuaucugca..	16	0	seq
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.....gAugugacuaucuaucugcaa..	1	1	seq

aaaucaaguagauagugacaggguaaaggcugugacuaucuaucugcaaa

.....gcugugacuaucCacuugcaa.	1	1	seq
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.....gcugugacuaucuaucugcaa.	22	0	seq
.....gcugugacuaAcuaucugcaa.	1	1	seq
.....gcugugacAaucuaucugcaaa	2	1	seq
.....gcugugacuaucuaucCugcaaa	1	1	seq
.....Ucugugacuaucuaucugcaaa	4	1	seq
.....gcugugacuaGcuacuugcaaa	1	1	seq
.....gUugugacuaucuaucugcaaa	1	1	seq
.....gcugugacuaCcuacuugcaaa	5	1	seq
.....gcugCgacuaucuaucugcaaa	2	1	seq
.....gcGgugacuaucuaucugcaaa	1	1	seq
.....gcugugacuaucuaucAugcaaa	1	1	seq
.....gcugugacuaucuaucugcaGa	1	1	seq
.....Acugugacuaucuaucugcaaa	46	1	seq
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.....ugugacuaucuaucugcaaU	1	1	seq
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.....gCgacuaucuaucugcaaa	1	1	seq
.....gugacuaAuacuugcaaa	1	1	seq
.....Augacuaucuaucugcaaa	3	1	seq
.....gugacuaucuaucugcaaU	4	1	seq
.....gugacuaucuaucugcaaa	7	0	seq