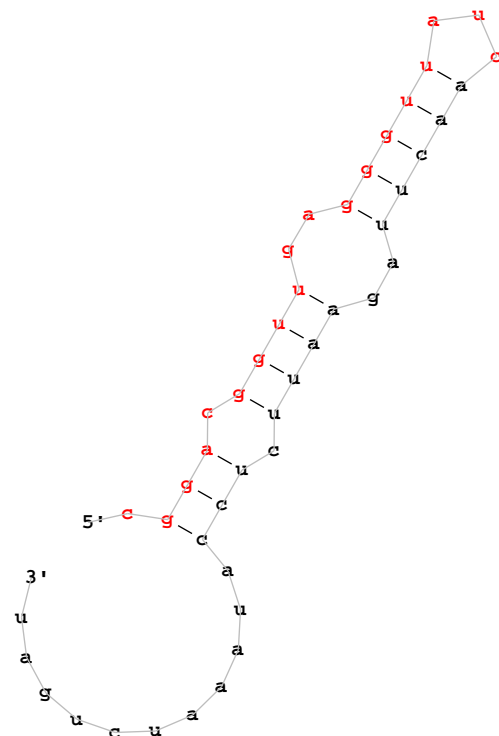
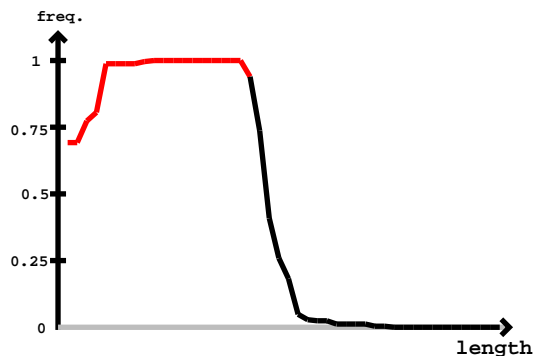


miRBase precursor : chromosome_14_73322
 Total read count : 247
 chromosome_14_73322 read count : 240
 remaining reads : 7



chromosome_14_73322			
5'-	-3'	exp	
cggacggguugagggguuaucaacuuagaauucuccauaaaucugau	reads	mm	sample
.(((.((((.((((...))))))..))))).)).....	12	0	seq
cggacggguugagggguuau.....	2	1	seq
cggacggguugagggguuac.....	1	1	seq
cggacggguugagggguuauA.....	1	1	seq
cggacggguugagggguuauC.....	1	1	seq
cggacggguugagggguuauGc.....	39	0	seq
cggacggguugagggguuauuc.....	8	1	seq
cggacggguugagggguuauU.....	1	1	seq
cggacggguugagggguuauG.....	1	1	seq
cggacggguugagggguuauucG.....	1	1	seq
cggacgAuugagggguuauca.....	1	1	seq
cggGcggguugagggguuauca.....	1	1	seq
cggacggguugagggguuauca.....	31	0	seq
cggacggguugagggguuGuca.....	1	1	seq
cggacggguugagggguuauuC.....	16	1	seq
cggacggguugagggguuauucU.....	9	1	seq
cggacggguugagggguuauCca.....	1	1	seq
cggacggguugagggguuauGca.....	1	1	seq
cggacggguugagggguuauCa.....	2	1	seq
Uggacggguugagggguuaucaa.....	1	1	seq
cggacggguugagggguuaucaa.....	1	0	seq
cggacggguugagggguuaucaac.....	1	0	seq
cggacggguugagggguuaucaaU.....	2	1	seq
Uggacggguugagggguuaucaacu.....	1	1	seq
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cggacggguugagggguuaucaaUu.....	30	1	seq
cggacggguugagggguuaucaacuC.....	3	1	seq
cggacggguugagggguuaucaauUuu.....	1	1	seq
cggacggguugagggguuaucaacuCa.....	1	1	seq
.Ugacggguugagggguuauca.....	5	1	seq
.ggacggguugagggguuauuC.....	4	1	seq
.ggacggguugagggguuauU.....	1	1	seq
.ggacggguugagggguuauca.....	4	0	seq
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.ggacggguugagggguuaucaa.....	1	0	seq
.ggacggguugagggguuaucaaUu.....	1	1	seq

cggaaggguaggguuaucaacuuagaauuccauaaaucugau

.ggacggguaggguuaucaacuC.....	1	1	seq
.ggacggguaggguuaucaaaUuuagaauuc.....	2	1	seq
.gacggguaggguuauucU.....	1	1	seq
.gacggguaggguuauca.....	3	0	seq
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.gacggguaggguuauuCa.....	1	1	seq
...acggguaggguuauucaa.....	20	0	seq
...acggguaggguuauuCa.....	4	1	seq
...Gcggguaggguuauucaa.....	1	1	seq
...acggguaggguuauucaa.....	1	1	seq
...Ucggguaggguuauucaa.....	3	1	seq
...acggguaggguuauucaaU.....	15	1	seq
...acggguaggguuauucaaUuuagaauucuc.....	1	1	seq
.....uaggguuauucaaUuuaga.....	2	1	seq
.....uaggguuauucaaUuuaga.....	1	1	seq