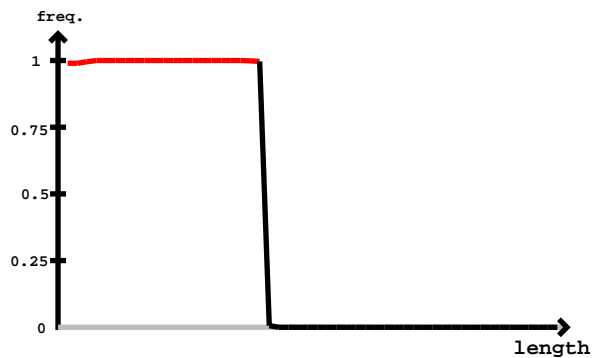
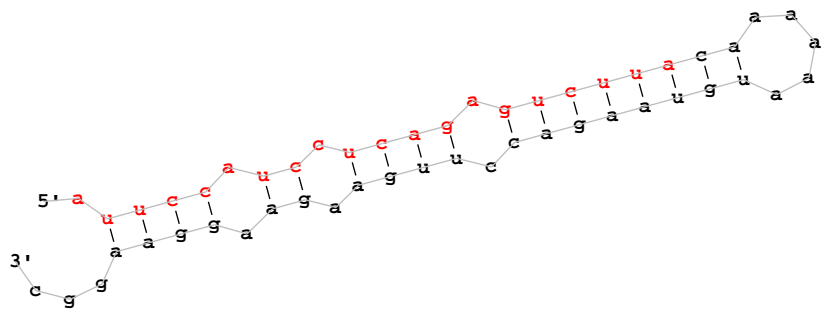


miRBase precursor : chromosome\_1\_1579  
 Total read count : 2529  
 chromosome\_1\_1579 read count : 2529  
 remaining reads : 0



chromosome_1_1579			
5'-	-3'	exp	
auuccauccucagagucuucaaaaaaugaaagccuugaagaaggaaggc			
.(((.(.(((.((((.(.....)))))))).)).)).))....	reads	mm	sample
Guuccauccucagagucu.....	3	1	seq
auuccauccucagagucuu.....	6	0	seq
auuccaucAucagagucuu.....	2	1	seq
auuccauccGcagagucuu.....	1	1	seq
auuccaAccucagagucuu.....	2	1	seq
auuAcauccucagagucuu.....	4	1	seq
auuccauccucagagucuuC.....	1	1	seq
auuccauccucagCgucuu.....	2	1	seq
auGccauccucagagucuu.....	2	1	seq
auuccauccucagaAucuu.....	1	1	seq
auuccauccucagagAucuu.....	4	1	seq
auuGcauccucagagucuu.....	1	1	seq
auuccauccucagagucuuU.....	9	1	seq
auuccauccuAagagucuu.....	6	1	seq
auuccaucGucagagucuu.....	4	1	seq
auucUauccucagagucuu.....	1	1	seq
auuccauccucagagucuu.....	2055	0	seq
auuccauccucagagucuC.....	13	1	seq
auucGauccucagagucuu.....	4	1	seq
auuccauccucagagucuuA.....	1	1	seq
auuUcauccucagagucuu.....	2	1	seq
auuccGuccucagagucuu.....	14	1	seq
auuccauGcucagagucuu.....	4	1	seq
auuccauccucagagucuuG.....	1	1	seq
auuccauccucUgagucuu.....	2	1	seq
Uuuccauccucagagucuu.....	14	1	seq
Cuuccauccucagagucuu.....	8	1	seq
auuccauccucagagucuuG.....	9	1	seq
Guuccauccucagagucuu.....	169	1	seq
auuccauccucagUgucuu.....	2	1	seq
auCccauccucagagucuu.....	12	1	seq
auuccauccucagaguAuu.....	3	1	seq
auuccauccucGgagucuu.....	13	1	seq
auuccauccucagagCuu.....	5	1	seq
auuccauccucagagucGua.....	3	1	seq

auuccauccucagagucuuacaaaaaaguaagaccuugaagaaggaaggc

auuccauccucagGgucuuu.....	9	1	seq
aCuccauccucagagucuuu.....	23	1	seq
auuccauccCcagagucuuu.....	11	1	seq
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auucAauccucagagucuuu.....	4	1	seq
auuccUuccucagagucuuu.....	2	1	seq
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auuccaCccucagagucuuu.....	5	1	seq
auuccauccuUagagucuuu.....	4	1	seq
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auuccaucUucagagucuuu.....	4	1	seq
aAuuccauccucagagucuuu.....	2	1	seq
auuccauccucagagucCuu.....	6	1	seq
auuccauccucagaUucuuu.....	2	1	seq
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auuccauAcucagagucuuu.....	2	1	seq
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