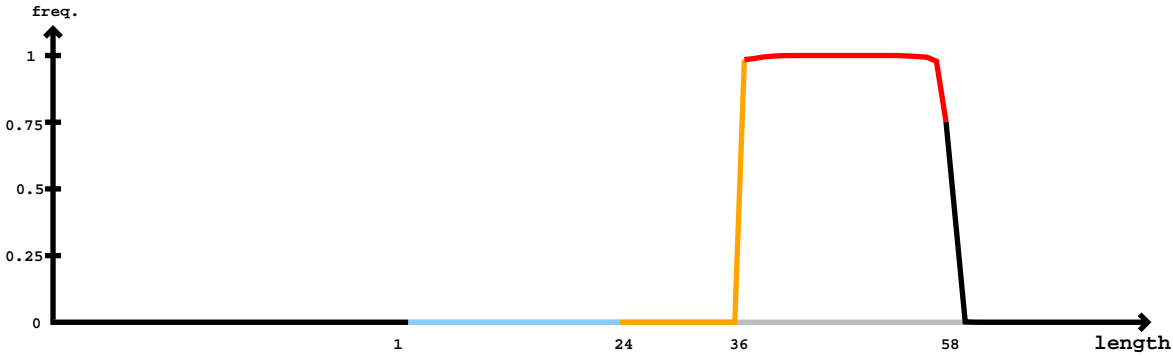
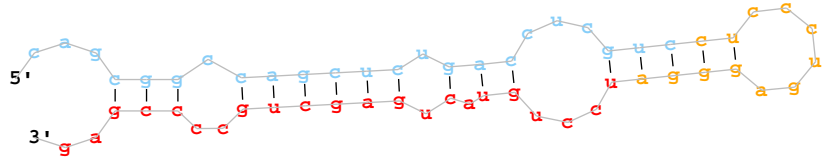


Provisional ID : DRO_27998_41892
 Score total : 2.5
 Score for star read(s) : -1.3
 Score for read counts : 0
 Score for mfe : 2.2
 Score for randfold : 1.6
 Score for cons. seed :
 Total read count : 402723
 Mature read count : 402723
 Loop read count : 0
 Star read count : 0



Star **Mature**

5' - aggcucuccagugggccaagcagugcugggggcuucagcggccagcucugaccucgucucccugagggauccuguacugagcugccccaggcucucacuggcgccgcu -3' exp

reads	mm	sample
1	1	seq
1	1	seq
2	1	seq
1	1	seq
7	1	seq
2	1	seq
51	1	seq
101	1	seq
1	1	seq
766	0	seq
1	1	seq
1	1	seq
1	1	seq
3	1	seq
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711	0	seq
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1	1	seq
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1	1	seq
1	1	seq
4	1	seq
1	1	seq
1	1	seq
4	1	seq

Star

Mature

Sequence	Count	Count	Label
aggcucuccagugggccaagcagugcugggggcuuagcggccagcucugaccucgucuccccagagggauccguacuagagcugccccagggcucucucacugcggcgcu	1006	0	seq
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.....uccguacuagagcugccccG.....	1	1	seq
.....uGcuguacuagagcugccccG.....	1	1	seq
.....uccguacuagagcucCccccG.....	1	1	seq
.....uccguacuagagcugGccccG.....	2	1	seq
.....uccgAacugagcugccccG.....	1	1	seq
.....uccguacuagaUcugccccG.....	1	1	seq
.....uccguacuagagcugccAcG.....	2	1	seq
.....uccgCacugagcugccccG.....	3	1	seq
.....uccguacuagagAugccccG.....	1	1	seq
.....ucUugacuagagcugccccG.....	1	1	seq
.....uccguacuagagcugccccA.....	8	1	seq
.....uccGguacuagagcugccccG.....	2	1	seq
.....uccgGacugagcugccccG.....	1	1	seq
.....ucGuguacuagagcugccccG.....	1	1	seq
.....uccguacuagagcugccccC.....	1	1	seq
.....uccguacuagagcugccccU.....	9	1	seq
.....uccguacuAagcugccccG.....	1	1	seq
.....uccguacuGugagcugccccG.....	1	1	seq
.....uccguacuagagcugccccG.....	5814	0	seq
.....uccguacuagagcugAcccG.....	6	1	seq
.....uccguacuagUgcugccccG.....	1	1	seq
.....GccuguacuagagcugccccG.....	3	1	seq
.....uccguacuagagcucUccccG.....	3	1	seq
.....AccguacuagagcugccccG.....	6	1	seq
.....uccguacuagagcugccccGU.....	292	1	seq
.....uccguacuagagcugccGcG.....	54	1	seq
.....uccguacuagagcugcAcccG.....	12	1	seq
.....uccguacuagagcucCccccG.....	6	1	seq
.....uccguacuagagcugccccCa.....	1	1	seq
.....uccguacuagagcugccccGG.....	777	1	seq
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.....uccguacuagagcugccccAga.....	2	1	seq
.....uccguacuUagcugccccG.....	2	1	seq
.....uccguacuagagcugccccUga.....	5	1	seq
.....uccguacuagagcugAcccG.....	51	1	seq
.....uccguacuUgagcugccccG.....	4	1	seq
.....uccguacuagagcCgccccG.....	1	1	seq
.....uccguacuagagcugGccccG.....	34	1	seq
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.....uccguacuagagcNgccccG.....	1	1	seq
.....ucUugacuagagcugccccG.....	3	1	seq
.....uccguacuagagUgccccG.....	5	1	seq
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.....uccguacuagagcugccAcG.....	24	1	seq
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Star

Mature

aggcucuccagugggccaacgagucgucgggggcuuacagggccagcucugaccucgucuccccgaggggauccuguaacugagcugccccgaggcucucacugcggcgcu

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.....uccuguGcugagcugccccga.....	10	1	seq
.....uAcuguaacugagcugccccga.....	4	1	seq
.....uccuguaacugCgugccccga.....	1	1	seq
.....uccuguaacugCgagcugccccga.....	3	1	seq
.....uccuguaacugagcugccUcga.....	6	1	seq
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.....uUcuguaacugagcugccccgag.....	10	1	seq
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Star

Mature

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.....uccguacuagagcNgccccgag.....	1	1	seq
.....uccguacuagagcugccccgagg.....	488	0	seq
.....uccguacuagagcugccccgagA.....	38396	1	seq
.....uccguacuagagcugccccgagAg.....	346	1	seq
.....uccguacuagagcugccccgagCg.....	28	1	seq
.....uccguacuagagcugcccGcgagg.....	2	1	seq
.....uccguacuagagcugcccAcgagg.....	1	1	seq
.....uccguacuagagcugccccgagU.....	79247	1	seq
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.....cuguGcugagcugccc.....	1	1	seq
.....cuguacuagagcugAcccg.....	1	1	seq
.....cuguacuagagcugccccg.....	53	0	seq
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.....cuUuacuagagcugccccga.....	1	1	seq
.....cuAuacuagagcugccccga.....	1	1	seq
.....cuguacuagagcugUcccga.....	1	1	seq
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.....cuguacuagagcugAcccga.....	1	1	seq
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Star

Mature

aggcucuccagugggccaacgacagucgugggggcuuacagggccagcucugaccucgucuccccgagggauccguacuagagcugccccgagggcucucacugcggcgcu

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.....cuUuacuagagcugccccgag.....	3	1	seq
.....cuguacuCagcugccccgag.....	2	1	seq
.....cuguacuagagcugcccGcgag.....	1	1	seq
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.....cuguacuagagcugccccgaA.....	22	1	seq
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.....cuguacuagagcugccccgagA.....	234	1	seq
.....cuguacuagagcugccccgaggU.....	120	1	seq
.....cuguacuagagcugccccgaggA.....	3	1	seq
.....cuguacuagagcugccccgagAc.....	2	1	seq
.....cuguacuagagcugccccgaggc.....	1	0	seq
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.....cuguacuagagcugccccgaggAu.....	1	1	seq
.....uguacuagagcugccccg.....	42	0	seq
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.....uacuagagcugccccgag.....	58	0	seq
.....uacuagagcugccccgagA.....	20	1	seq
.....uacuagagcugccccgagU.....	49	1	seq
.....acuagagcugccccgagg.....	1	0	seq
.....acuagagcugccccgagU.....	38	1	seq
.....acuagagcugccccgagA.....	15	1	seq
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