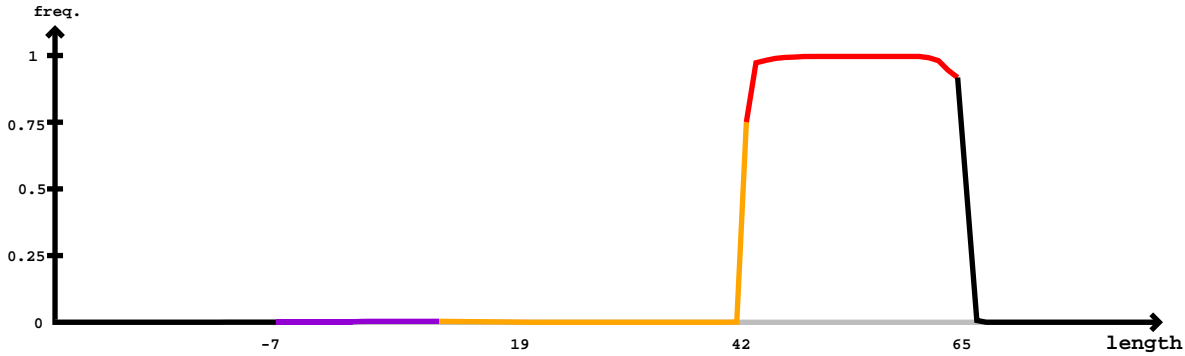
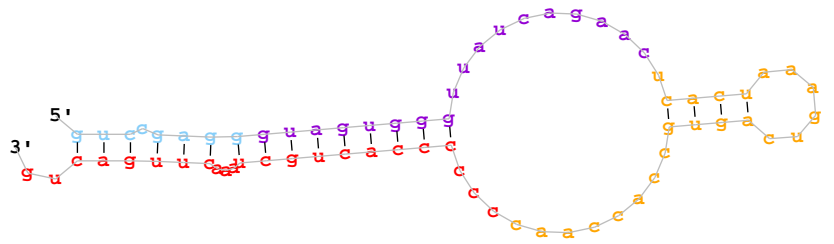


Provisional ID : MNA_308_5501
 Score total : 1.7
 Score for star read(s) : -1.3
 Score for read counts : 0
 Score for mfe : 1.4
 Score for randfold : 1.6
 Score for cons. seed :
 Total read count : 787270
 Mature read count : 785362
 Loop read count : 0
 Star read count : 1908



Star

Mature

5'		-3'	obs	mm	sample
ugaaaguggguuucucggcugcaggguagugguuauucagaacucacuaaagucagugccaccaacccccacugcuaaaacuugacuggccuuuuuaaaccacuggac					
ugaaaguggguuucucggcugcaggguagugguuauucagaacucacuaaagucagugccaccaacccccacugcuaaaacuugacuggccuuuuuaaaccacuggac					
.....((((.....((((((((((((((((((((.....)))))))))....)))))))))....)))))).....		reads			
.....ggcugguuccgaAgguag.....		6	1		seq
.....ggcugguuccgaggCag.....		3	1		seq
.....ggcugguuccgaCgguaugu.....		1	1		seq
.....ggcugguuccgaggCagu.....		2	1		seq
.....ggcugguuccgaAgguagug.....		1	1		seq
.....ggcugguuccgaggCagug.....		3	1		seq
.....ggcugguuccgaggCaguggguuuau.....		1	1		seq
.....ggcugguuccgaUgguaguggguuuauuc.....		1	1		seq
.....ggcugguuccgaggCaguggguuuauucaga.....		3	1		seq
.....ggcugguuccgaUgguaguggguuuauucaga.....		2	1		seq
.....ggcugguuccgaUgguaguggguuuauucagaa.....		10	1		seq
.....ggcugguuccgaggCaguggguuuauucagaa.....		2	1		seq
.....ggcugguuccgaUgguaguggguuuauucagaaac.....		142	1		seq
.....ggcugguuccgaggCaguggguuuauucagaaac.....		28	1		seq
.....ggcugguuccgaUgguaguggguuuauucagaaacu.....		46	1		seq
.....ggcugguuccgaggCaguggguuuauucagaaacu.....		3	1		seq
.....ggcugguuccgaUgguaguggguuuauucagaaac.....		4	1		seq
.....ggcugguuccgaggCaguggguuuauucagaaac.....		1	1		seq
.....ggcugguuccgaUgguaguggguuuauuc.....		1	1		seq
.....ggcugguuccgaggCaguggguuuauuca.....		1	1		seq
.....ggcugguuccgaggCaguggguuuauucagaa.....		1	1		seq
.....ggcugguuccgaggCaguggguuuauucagaaac.....		1	1		seq
.....ggcugguuccgaggCaguggguuuauucagaaacu.....		1	1		seq
.....ggcugguuccgaUgguaguggguuu.....		1	1		seq
.....ggcugguuccgaggCaguggguuuauucaga.....		1	1		seq
.....ggcugguuccgaUgguaguggguuuauucagaac.....		2	1		seq
.....ggcugguuccgaggCaguggguuuauucaga.....		7	1		seq
.....ggcugguuccgaUgguaguggguuuauucagaa.....		18	1		seq
.....ggcugguuccgaggCaguggguuuauucagaac.....		282	1		seq
.....ggcugguuccgaggCaguggguuuauucagaaacu.....		33	1		seq
.....ggcugguuccgaUgguaguggguuuauucagaa.....		32	1		seq
.....ggcugguuccgaggCaguggguuuauucagaac.....		1	1		seq
.....ggcugguuccgaggCaguggguuuauucagaac.....		859	1		seq
.....ggcugguuccgaggCaguggguuuauucagaaacu.....		109	1		seq

Star

Mature

ugaaaguggcuucucggcugguccgaggguaguggguuauucagaacucacuaaagucagugcccaacccccacugcuuaacuugacuggccuuuaaaaccacuggac			
.....uGuggguuauucagaac.....	1	1	seq
.....Caguggguuauucagaac.....	699	1	seq
.....uaguggguuaCcagaac.....	2	1	seq
.....Caguggguuauucagaacu.....	248	1	seq
.....aguggguuauucagaCcu.....	3	1	seq
.....aguggguuauucagaacA.....	34	1	seq
.....aguggguuauucagaacu.....	144	0	seq
.....aguggguuauucagaacG.....	1	1	seq
.....aguggguuauucCgaacu.....	8	1	seq
.....aguggguuauucagaacuU.....	7	1	seq
.....guggguuauucagaacuG.....	1	1	seq
.....guggguuauucagaacuA.....	1	1	seq
.....guggguuauucagaacuU.....	5	1	seq
.....guggguuauucagaacuUa.....	2	1	seq
.....accccccacugcuGaacuugacug.....	3	1	seq
.....acccccacugcuGaacuugacug.....	2	1	seq
.....acccccacugcuGaacuugacugg.....	1	1	seq
.....ccccccacugcuGaac.....	3	1	seq
.....ccccccacugcuGaacuu.....	2	1	seq
.....ccccccacugcuGaacuug.....	5	1	seq
.....ccccccacugcuGaacuuga.....	15	1	seq
.....ccccccacugcuGaacuugac.....	32	1	seq
.....ccccccacugcuGaacuugacu.....	18	1	seq
.....ccccccacugcuGaacuugacug.....	583	1	seq
.....ccccccacugcuGaacuugacugg.....	56	1	seq
.....ccccccacugcuGaacuugacuggc.....	6	1	seq
.....ccccccacugcuuaacu.....	1	0	seq
.....ccccccacugcuUaacuu.....	1	1	seq
.....ccccccacugcuUaacuug.....	1	1	seq
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.....ccccccacugcuuaacuug.....	1	0	seq
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.....ccccccacugcuUaacuugacug.....	16	1	seq
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.....ccccccacugcuuaacuugacugU.....	3	1	seq
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.....ccccccacugcuuaacuugacug.....	7	0	seq
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Star

Mature

Sequence	Count	Count	Label
ugaaaguggcuucucggcugguccgaggguaguggguuaucaagaacucacuaaagucagugcccaacccccacugcuuaacuugacuggc			
.....ccccacugcuUaacuugacugg.....	1	1	seq
.....ccccacugcuNaacuugacugg.....	1	1	seq
.....ccccacugcuCaacuugacugg.....	2	1	seq
.....ccccacugcuuaacuugacugg.....	1	0	seq
.....ccccacugcuGaacuugacugg.....	19023	1	seq
.....ccccacugcuGaacuugacuggc.....	1158	1	seq
.....ccccacugcuCaacuugacuggc.....	1	1	seq
.....ccccacugcuGaacuugacuggcc.....	8	1	seq
.....ccccacugcuGaacuug.....	100	1	seq
.....ccccacugcuGaacuuga.....	129	1	seq
.....ccccacugcuGaacuugac.....	215	1	seq
.....ccccacugcuGaacuugacu.....	214	1	seq
.....ccccacugcuUaacuugacu.....	1	1	seq
.....ccccacugcuUaacuugacug.....	1	1	seq
.....ccccacugcuGaacuugacug.....	5669	1	seq
.....ccccacugcuGaacuugacugg.....	545	1	seq
.....ccccacugcuuaacuugacugg.....	1	0	seq
.....ccccacugcuGaacuugacuggc.....	61	1	seq
.....cccacugcuGaacuuga.....	90	1	seq
.....cccacugcuGaacuugac.....	199	1	seq
.....cccacugcuGaacuugacu.....	172	1	seq
.....cccacugcuaaUcuugacug.....	1	1	seq
.....cccacugcuGaacuugacug.....	4856	1	seq
.....cccacugcuGaacuugacugg.....	463	1	seq
.....cccacugcuGaacuugacuggc.....	32	1	seq
.....cccacugcuaaaUuugacuggc.....	1	1	seq
.....ccacugcuGaacuugac.....	134	1	seq
.....ccacugcuGaacuugacu.....	108	1	seq
.....ccacugcuUaacuugacug.....	1	1	seq
.....ccacugcuGaacuugacug.....	2466	1	seq
.....ccacugcuGaacuugacugg.....	243	1	seq
.....ccacugcuGaacuugacuggc.....	30	1	seq
.....cacugcuGaacuugacu.....	42	1	seq
.....cacugcuGaacuugacug.....	850	1	seq
.....cacugcuCaacuugacug.....	1	1	seq
.....cacugcuGaacuugacugg.....	158	1	seq
.....cacugcuGaacuugacuggc.....	19	1	seq
.....acugcuGaacuugacug.....	1629	1	seq
.....acugcuGaacuugacugg.....	86	1	seq
.....acugcuCaacuugacugg.....	1	1	seq
.....acugcuGaacuugacuggc.....	11	1	seq
.....cugcuGaacuugacugg.....	130	1	seq
.....cugcuGaacuugacuggc.....	10	1	seq
.....cugcuGaacuugacuggcc.....	1	1	seq
.....ugcuGaacuugacuggc.....	4	1	seq