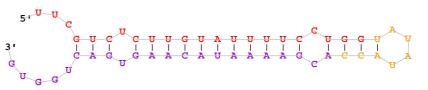
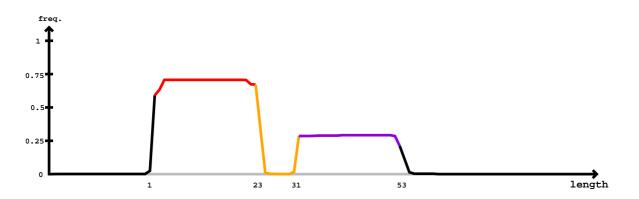
$Provisional\ ID \\ : NC_058078.1_A cropora_millepora_isolate_JS-1_chromosome_13_A mil_v2.1_whole_genome_shotgun_sequence_489676$

494.9 Score total Score for star read(s) 3.9 Score for read counts 486 Score for mfe 2 Score for randfold 0 Score for cons. seed 3 Total read count 965 685 Mature read count Loop read count 0 Star read count 280





Mature	Star
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auguaucguggaagauccca <u>uucgucucuuguauuuuccugguauauacc</u> acgaaaauacaagugacuggugggauauucuuugguuuaccacucaaaaguguggcauaacu	exp		
((((.((((((((((((((((((((((((((((((reads	mm	sample
auguaucguggaagauccc	1	0	seq
auuegucucuuguauuuueeCgg	1	1	seq
Guucgucucuuguauuuuccugg.	1	1	seq
auucgucucuuguauuuuccugA	1	1	seq
auCegucucuuguauuuuccugg	1	1	seq
auucgucucuuguauuuuccugg	13	0	seq
auuUgucucuuguauuuuccugg.	7	1	seq
uuegucucuuguauuuuce	1	0	seq
uuegucucuuguauuuuceA	5	1	seq
UUgucucuuguauuuuccu	2	1	seq
uuegucucuuguauuuuccu	15	0	seq
uuegucucuuguauuuuceG	3	1	seq
uucgucucuuguauuuuccC	4	1	seq
uuegucucuuguauuuuccuA	1	1	seq
uuegucucuuguauuuuccug	2	0	seq
uuegucucuuguauuuuccugC	5	1	seq
.uucgucucuuguGuuuuccugg.	2	1	seq
UuUgucucuuguauuuuccugg	43	1	seq
uuegucCcuuguauuuuccugg	2	1	seq
uucgucucCuguauuuuccugg	3	1	seq
uucgucucuAguauuuuccugg	1	1	seq
uuegucucuuguauuuuecCgg	2	1	seq
uuegucucuuguauuCuccugg	3	1	seq
uucgucucuuguauuuuccugg	256	0	seq
Auegucucuuguauuuuccugg	1	1	seq
uucgucucuCguauuuuccugg	1	1	seq
uuegucucuuguauuuuccugA	81	1	seq
uucgucucuuguauuuuccuAg	1	1	seq
uucgucucuuguauuuuccugU	14	1	seq
uucgucucuuguauuuuccuUg.	1	1	seq
uucgucucuuguaCuuuccugg.	1	1	seq
Cuegucucuuguauuuuccugg.	1	1	seq
	1	1	

Mature Star

Mature St	ar			
auguaucguggaagaucccauucgucuuguauuuuccugguauauaccacga	aaauacaagugacuggugggauauucuuugguuuaccacucaaaaguguggcauaacu			
uucgucucuuguauuuuccuggu		27	0	seq
uuUgucucuuguauuuuccuggu		56	1	seq
uucqucucuuquauuuuccuqqC		5	1	seq
uucgucucuuguauuuuccuggA		1	1	seq
	•••••	1	1	
uuUgucucuuguauuuuccuggua	••••••			seq
	•••••	4	1	seq
uucgucucuuguauuuuccuggua	•••••	2	0	seq
uucgucucuuguauuuuccugguaGa	•••••	1	1	seq
AgucucuuguauuuuccugA	•••••	3	1	seq
uUgucucuuguauuuuccugg	•••••	1	1	seq
ucgucucuuguauuuuccugg		10	0	seq
ucgucucuuguauuuuccuggC		4	1	seq
A		1	1	seq
ucgucucuuguauuuuccuggu		17	0	seq
ucgucucuuguauuuuccuggA		3	1	seq
uUgucucuuguauuuuccuggu		1	1	seq
ucgucucuuguauuuuccugguauU		1	1	seq
Ugucucuuguauuuuccu		2	1	seq
		1	1	seq
cgucucuuguauuuuccugC		1	1	seq
Ugucucuuguauuuuccugg		10	1	seq
		8	1	
	••••••			seq
gucucuuguauuuuccugg		10	0	seq
cgucucuuguauuuuccuCg	•••••	1	1	seq
Ugucucuuguauuuuccuggu	•••••	27	1	seq
gucucuuguauuuuccugCu	•••••	1	1	seq
cgucucuuguauuuuccuggu		8	0	seq
cgucucuuguauuuuccugguC		1	1	seq
gucucuuguauuuuccugguU		1	1	seq
caAga	aaauacaagugacugg	1	1	seq
cacga	aaauacaagugacuggu	7	0	seq
cacga	aaauacaagugacugCu	1	1	seq
caAga	aaauacaagugacuggu	4	1	seq
cacga	aaauacaagugacuggC	3	1	seq
cacga	aaauacaagugacuggug	1	0	seq
acga	aaauacaagugacugg	4	0	seq
	aaauacaagugacugA	1	1	seq
	aaauacaagugacuggu	39	0	seq
	aaauacGagugacuggu	1	1	seq
	aGauacaagugacuggu	1	1	seq
		1	1	
	aaauacaagugacuggu	8	1	seq
	aaauacaagugacuggC			seq
	aaauacaagugacuggu	5	1	seq
	aaauacaagugacuggA	2	1	seq
	aaauacCagugacuggug	2	1	seq
	aaauacaagugacuggug	25	1	seq
	aaauacaagugacuggug	3	1	seq
acga	aaauacaagugacugguU	6	1	seq
acga	aaauacaagugacuggCg	1	1	seq
acga	aaauacaagugacuggAg	1	1	seq
acga	aaauacaagugacugguA	18	1	seq
acga	aaauacaagugacugguC	4	1	seq
acga	aaauacaagugacuggug	56	0	seq
acga	aaauacaagugacuggugU	41	1	seq
aAga	aaauacaagugacuggugg	2	1	seq
acga	aaauacaagugacuggugA	9	1	seq
	aaauacaagugacAggugg	2	1	seq
	aaauacaagugacuggugC	6	1	seq
		1	1	_
	aaauacaagugacuggugg			seq
	aaauacaagugacuggugg	12	0	seq
	aaauacaagugacugguggA	1	1	seq
	aaauacaagugacugguggU	8	1	seq
	aaauacaagugacuCgu	1	1	seq
	aaauacaagugacuCguggg	2	1	seq
	caagugacuCgugggauauu	3	1	seq