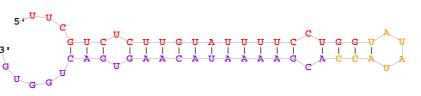
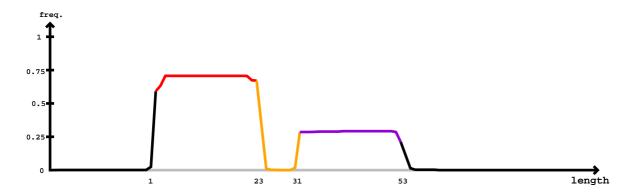
$Provisional\ ID \\ : NC_058078.1_A cropora_millepora_isolate_JS-1_chromosome_13_A mil_v2.1_whole_genome_shotgun_sequence_489676$

491.3 Score total Score for star read(s) 3.9 Score for read counts 486 2 Score for mfe Score for randfold 0 Score for cons. seed -0.6 Total read count 965 685 Mature read count Loop read count 0 Star read count 280





Mature St	tar
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auguaucguggaagauccca <mark>uucgucucuuguauuuccugguauauacc</mark> acgaaaauacaagugacuggugggauauucuuugguuuaccacucaaaaguguggcauaacu	exp		
((((.((((((((((((((((((((((((((((((reads	mm	sample
auguaucguggaagauccc	1	0	seq
auucgucucuuguauuuuccugA.	1	1	seq
auuUgucucuuguauuuuccugg.	7	1	seq
auucgucucuuguauuuuccCgg	1	1	seq
auCegucucuuguauuuuccugg.	1	1	seq
auucgucucuuguauuuuccugg.	13	0	seq
Guucgucucuuguauuuuccugg.	1	1	seq
uuegucucuuguauuuuce	1	0	seq
uucgucucuuguauuuuccG	3	1	seq
uucgucucuuguauuuuccC	4	1	seq
uuegucucuuguauuuuccu	15	0	seq
uuegucucuuguauuuuceA	5	1	seq
uuUgucucuuguauuuuccu	2	1	seq
uuegucucuuguauuuuccuA	1	1	seq
uucgucucuuguauuuuccug	2	0	seq
uucgucucuuguauuuuccuUg.	1	1	seq
.uucgucucuuguauuCuccugg.	3	1	seq
uucgCeucuuguauuuuceugg.	1	1	seq
uucgucucuuguauuuuccuAg.	1	1	seq
.uuegueCeuuguauuuuecugg	2	1	seq
uucgucucuuguauuuuccugA	81	1	seq
uucgucucuuguauuuuccugg	256	0	seq
uucgucucuuguGuuuuccugg.	2	1	seq
.uucgucucuuguauuuuccCgg	2	1	seq
Aucgucucuuguauuuuccugg.	1	1	seq
uucgucucuCguauuuuccugg.	1	1	seq
UuUgucucuuguauuuuccugg	43	1	seq
uucgucucuuguaCuuuccugg.	1	1	seq
uucgucucuAguauuuuccugg.	1	1	seq
uucgucucuuguauuuuccugC	5	1	seq
Cuegucucuuguauuuuccugg.	1	1	seq
uuegucucCuguauuuuccugg.	3	1	seq
uucgucucuuguauuuuccugU	14	1	seq

Mature Star

Mature Star				
auguaucguggaagaucccauucgucucuuguauuuuccugguauauaccacgaaaa	nacaagugacuggugggauauucuuugguuuaccacucaaaaguguggcauaacu			
uucgucucuuguauuuuccuggC		5	1	seq
gucucuuguauuuuccuggu		27	0	seq
uuUgucucuuguauuuuccuggu		56	1	seq
uucgucucuuguauuuuccuggA		1	1	seq
		4	1	
uucgucucuuguauuuuccugguU				seq
		1	1	seq
uucgucucuuguauuuuccuggua		2	0	seq
uucgucucuuguauuuuccugguaGa	••••••	1	1	seq
ucgucucuuguauuuuccugA		3	1	seq
uUgucucuuguauuuuccugg		1	1	seq
ucgucucuuguauuuuccugg		10	0	seq
uUgucucuuguauuuuccuggu		1	1	seq
ucgucucuuguauuuuccuggu		17	0	seq
ucgucucuuguauuuuccuggA		3	1	seq
ucgucucuuguauuuuccuggC		4	1	seq
AugucucuuguauuuuccugAu		1	1	seq
ucgucucuuguauuuuccugguauU		1	1	seq
Ugucucuuguauuuuccu		2	1	seq
gucucuuguauuuuccuCg		1	1	seq
cUucucuuquauuuuccuqq		1	1	
				seq
gucucuuguauuuuccugg	•••••	10	0	seq
gucucuuguauuuuccugA	•••••	8	1	seq
cgucucuuguauuuuccugC		1	1	seq
Ugucucuuguauuuuccugg		10	1	seq
cgucucuuguauuuuccuggu		8	0	seq
Ugucucuuguauuuuccuggu		27	1	seq
cgucucuuguauuuuccugCu		1	1	seq
cgucucuuguauuuuccugguC		1	1	seq
cgucucuuguauuuuccugguU		1	1	seq
caAgaaaa	uacaagugacugg	1	1	seq
	ıacaagugacuggu	4	1	seq
	aacaagugacuggC	3	1	seq
	acaagugacuggu	7	0	seq
	acaagugacugCu	1	1	seq
	acaagugacuggug	1	0	seq
		4	0	
	acaagugacugg			seq
	acaagugacugA	1	1	seq
	acaagugacuggC	8	1	seq
acgaaaat	acGagugacuggu	1	1	seq
acgaaaat	nacaagugacuggu	39	0	seq
acgaaaa	nacaagugacuggA	2	1	seq
aGgaaaat	nacaagugacuggu	1	1	seq
acgaaGat	nacaagugacuggu	1	1	seq
aAgaaaat	nacaagugacuggu	5	1	seq
acgaaaan	nacCagugacuggug	2	1	seq
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acgaaaa	ıacaagugacugguA	18	1	seq
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acgaaaa	ıacaagugacugguU	6	1	seq
	acaagugacuggug	56	0	seq
	acaagugacuggAg	1	1	seq
	acaagugacuggug	3	1	seq
	acaagugacuggug		1	_
-		25		seq
	acaagugacAggugg	2	1	seq
	acaagugacuggugg	1	1	seq
	acaagugacuggugg	12	0	seq
aAgaaaat	acaagugacuggugg	2	1	seq
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aaaaı	uacaagugacuCgu	1	1	seq
	nacaagugacuCguggg	2	1	seq
	caagugacuCgugggauauu	3	1	seq
				-