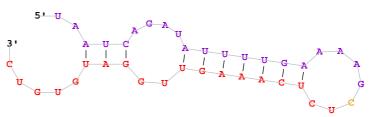
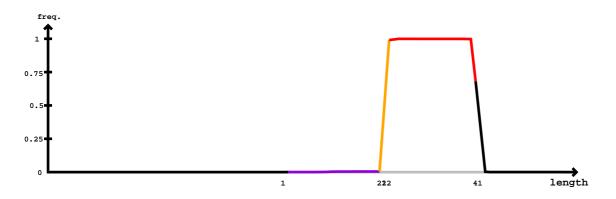
$Provisional\ ID \\ : NC\_058078.1\_A cropora\_millepora\_isolate\_JS-1\_chromosome\_13\_A mil\_v2.1\_whole\_genome\_shotgun\_sequence\_502316$ 

2268.9 Score total Score for star read(s) 3.9 Score for read counts 2269.4 Score for mfe -3.7 Score for randfold 0 Score for cons. seed -0.6 Total read count 4463 4462 Mature read count Loop read count 0 Star read count 1



obs



5'- uguuaguaaacuuuaguucaccgcaauuagagucauuauaagagaggauaaucagauauuuugaaaag<mark>cucucaaaguuggaugucc</mark>aaggucucagcucggcuau

## Star Mature

-55			
uguuaguaaacuuuaguucaccgcaauuagagucauuauaagagagga.uaauuagaaaagcucucaaaguuggauguguccaaggucucagcucggcuau	exp		
(((((((((((()))))))))((((((	reads	mm	sample
.uaGucagauauuuugaaaag	1	1	seq
	1	0	seq
auauuuugaaaagcucucaaaguuggaugu	3	0	seq
uauuuugaaaagcucucaaaguuggaugug	2	0	seq
auuuugaaaagcucucaaaguuggGugug	1	1	seq
auuuugaaaagcucucaaaguuggaugug	7	0	seq
auuuugaaaagcucucaaaguuggaugugG	1	1	seq
uuuugaaaagcucucaaaguuggaugugC	1	1	seq
uuuugaaaagcucucaaaguuggaugug	1	0	seq
uuugaaaagcucucaaaguuggaugugGc	1	1	seq
gaaaagcucucaaaguuggauguguŪ	2	1	seq
.cucucaaaguuggaugugu	1	1	seq
.cucucaaaguuggaugugucA	1	1	seq
ucucGaaguuggaugug	15	1	seq
ucucaaaguuggaugugG	32	1	seq
ucucaaaguuggauAugu	56	1	seq
ucucaaCguuggaugugu	1	1	seq
ucucaaaguuggaugGgu	2	1	seq
ucucaaaguuggaugAgu	1	1	seq
ucucaaaguuggaugugu	1200	0	seq
ucucaaaguuggaugugA	75	1	seq
ucucaaaguuggaugCgu	4	1	seq
ucucaaaguuggauguUu	1	1	seq
ucucaaaguCggaugugu	3	1	seq
ucucaaaguuggauguCu	4	1	seq
ucucaaaguuggauguku	4	1	seq
ucCcaaaguuggaugugu	5	1	seq
ucucaaaguuggGugugu	6	1	seq
ucucaaaguuggaCgugu	3	1	seq
ucCcaaaguuggauguc	3	1	seq
ucucGaaguuggauguguc	9	1	seq
ucucUaaguuggauguguc	1	1	seq
ucucaaaguuggauguA	1588	1	seq

## Star Mature

2001 1100010			
uguuaguaaacuuuaguucaccgcaauuagagucauuauaagagaggauaaucagauauuuugaaaag <mark>cucucaaaguuggaugugc</mark> caaggucucagcucggcuau			
ucucaaaguuggauAuguc	23	1	seq
.ucucaaaguuggauCuguc	1	1	seq
ucucaaaguuggaugAguc	1	1	seq
ucucaaaguuggaGguguc	1	1	seq
.ucucaaaguuggaCguguc	1	1	seq
ucucaaaguAggauguguc	1	1	seq
ucucaaaguuggaugugGc	1	1	seq
ucucaaaguuggauguCuc	9	1	seq
.ucucaaaCuuggauguguc	1	1	seq
ucGcaaaguuggauguguc	4	1	seq
ucucaaaguCggauguguc	7	1	seq
.ucucaaagAuggauguguc	1	1	seq
ucucaaaguuggauguAuc	1	1	seq
.ucucaUaguuggauguguc	1	1	seq
ucucaaaguuggauguguc	568	0	seq
.ucAcaaaguuggauguguc	2	1	seq
ucucaaaguuggaugugAc	2	1	seq
ucucaaaguuggaugCguc	4	1	seq
ucucaaaguuggGuguguc	4	1	seq
ucucaaaguuggauguguG	28	1	seq
ucucaaaguuggaugugucc	37	0	seq
ucucaaaguuggaugugucG	17	1	seq
ucucaaaguuggauguguAc	27	1	seq
ucucaaaguuggaugugucA	512	1	seq
ucucGaaguuggaugugucc	1	1	seq
ucCcaaaguuggaugugucc	1	1	seq
ucucaaaguuggauguguGc	1	1	seq
ucucaaaguuggaugugucU	141	1	seq
ucucaaaguuggaugugucAa	1	1	seq
ucucaaaguuggaugugucUa	7	1	seq
ucucaaaguuggauguguceU	1	1	seq
cucaaaguuggauguaA	5	1	seq
cucaaaguuggauguguc	8	0	seq
cucaaaguuggaugugucU	1	1	seq
cucaaaguuggGugugcc	1	1	seq
cucaaaguuggaugugucA	2	1	seq
cucaaaguuggauguwac	3	1	seq
ucaaaguuggaugugcc	3	0	seq
ucaaaguuggaugugucA	11	1	seq
ucaaaguuggaugugucU	5	1	seq
ucaaaguuggaugugucAa	1	1	seq