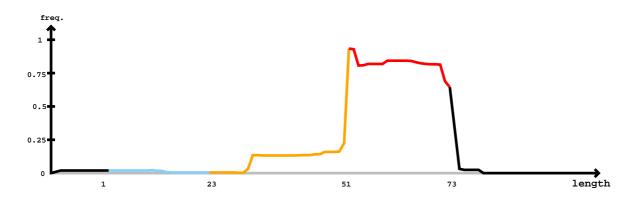
$Provisional\ ID \\ : NC_058078.1_A cropora_millepora_isolate_JS-1_chromosome_13_A mil_v2.1_whole_genome_shotgun_sequence_512651$

Score total : 0.7

Score for star read(s) : -1.3
Score for read counts : 0
Score for mfe : 2.6
Score for randfold : 0

5.A. UAUAAGUUGGGAUGGGA[©]UUAUAGGUUGG^GA AAUAUUCAGCGCUAGCGUU_GAAUAUUGAACG



Star Mature

5'- ugcgaucgcascuuauaaguugcgaucgcgacuuauagcuugcgaucgcaaguuauaacu <mark>ugcgaucgcgacuuauaacuug</mark> cgaucguaacuuauaacuuguaaccugcga	-3'	exp	
(((((((((((((((((((((((((((((((((((((((read	ls mm	sample
ugcgaucgcaacuuauaagu	3	0	seq
ugcgaucgcGacuuauaaguug	1	1	seq
ugcgaucgcaacuuauaaguuA	1	1	seq
ugcgaucgcaacuuauaagAug	1	1	seq
ugcgaucgcaacuuauaaguug	1	0	seq
ugegauegeaacuuauaagAuge	1	1	seq
uugcgaucgegacuuaua	2	0	seq
ugcgaucgcgacuuauagcuug	1	0	seq
uugcgaucgcaaguuauaacuu	2	0	seq
	2	1	seq
uCgcgaucgcaaguuauaacuug	2	1	seq
	1	1	seq
	1	1	seq
	3	0	seq
gggaucgcaaguuauaacuugc	1	0	seq
.ugcgaucgcaaguuauaacA	1	1	seq
.ugcgaucgcaaguuauaacuug	28	0	seq
.ugcgaucgcaaguuauGacuug	1	1	seq
.ugcgaucgcaaguuauaacuuU	3	1	seq
.uCcgaucgcaaguuauaacuug	1	1	seq
.ugcgaucgcaaguuGuaacuug	2	1	seq
.ugcgaucgcaaguuauaacuuA	6	1	seq
aguuauaacuugcgaucgcgacu	1	0	seq
uauaacuugcgaucgcgacuu	3	0	seq
	1	0	seq
uaacuugcgaucgcgacuua	1	0	seq
.uaacuugcgaucgcgacuuU	1	1	seq
uaGcuugcgaucgcgacuua	1	1	seq
uaacuugcgaucgcgacuuUu	1	1	seq
uaacuugcgaucgcgacuuau	1	0	seq
uaacuugcgaucAcgacuuauaacuugcg	1	1	seq
cuugcgaucgcgacuuauaaG	1	1	seq
uugcgaucgcgacuuaua	2	0	seq
uuqqaauqqqacuuauaacu	5	0	sea

Star Mature

Flactic			
ugcgaucgcascuuauaaguugcgaucgcgac <mark>uuauagcuugcgaucgcaaguuauaacuugcgaucgcgacuuauaacuug</mark> cgaucguaacuuauaacuuguaaccugcga			
uugcgaucgegacuuauaacuC	3	1	seq
uugcgaucgcgacuuauaacuG	1	1	seq
uugcgaucgcgacuuauaacCu	1	1	seq
uugcgaucgegacuuauaacuu	4	0	seq
uugcgaucgcgacuuauaacuA	1	1	seq
uugcgaucgcgacuuauaacuuU	1	1	seq
uugcgaucgcgacuuauaacuug	6	0	seq
.uugcgaucgcgacuuauaacuAg	1	1	seq
uugcgaucgcgacuuauaacuugc	2	0	seq
.ugcgaucgcgacuuauaacA	2	1	seq
.ugcgaucgcgacuuauaacC	5	1	seq
.ugcgaucgcgacuuauaacu	39	0	seq
.ugcgaucgcgacuuauaacuA	4	1	seq
.ugcgaucgcgacAuauaacuu	1	1	seq
.ugcgaucgcgacuuauaacuu	3	0	seq
.uUcgaucgcgacuuauaacuug.	1	1	seq
.ugcgaucgcgacuuauaacuug	192	0	seq
.ugcgaucgcgacuuauaacCug	1	1	seq
.ugcgaucgcgacuuauaacuuC	2	1	seq
.ugcgaucgcgacuuauGacuug.	2	1	seq
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.ugcgaucgcgacuuauaacuCg	1	1	seq
ugcgaucgcgacuuauaaUuug	1	1	seq
ugcgaucgcgacuuauaacAug	1	1	seq
ugcgaucgcgacuuauaacuuU	2	1	seq
.ugcgaucgcgacuuauaacuugA	1	1	seq
	14	1	seq
ugcgaucgcgacuuauaacuugc	12	0	seq
.ugcgGucgcgacuuauaacuugc	3	1	seq
.ugcgaucgcgacuuauaacuugcA	1	1	seq
gaucgcgacuuauaacuug	1	0	seq
gaucgcgacuuauaacuugc	1	0	seq
aucgcgacuuauaacuug	1	0	seq
Gucgcgacuuauaacuug	2	1	seq
aucgcgacuuauaacuugcA	1	1	seq
cgacuuauaacuugcgaucU	1	1	seq
cgacuuauaacuugcgaucA	9	1	seq