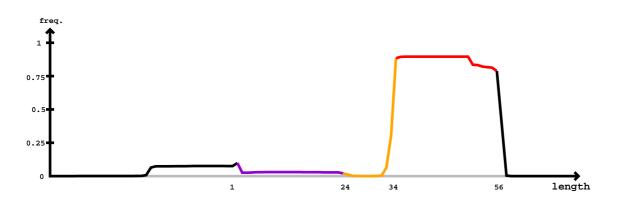
$: NC_058073.1_A cropora_millepora_isolate_JS-1_chromosome_8_A mil_v2.1_whole_genome_shotgun_sequence_328215$ Provisional ID

637.6 Score total

Score for star read(s) 3.9 629.8 Score for read counts 0.9 Score for mfe 0

Score for randfold Score for cons. seed 3 Total read count 1247 1209 Mature read count Loop read count 0 Star read count 38



Star Mature 5'- auuugaccugcuccucaaguuuugaaugaaauuguauuggcucuuuuaaguaguugaauaauuaaauuuacucacugcuuaaaagagucacuacagcuuugucuuacu

auuugaccugcuccucaaguuuugaaugaaauuguauuggcucuuuuaaguugaauaauuaaauuaacucacugcuuaaaagagucacuacagcuuugucuuacu	exp		
(((((())))))(((.((((((((((((reads	mm	sample
gaceugcuccuAaaguuuugaaugaaauugu	1	1	seq
aaguuuugaaugaaauuguaGu	1	1	seq
guuugaaugaaauuguaGu	8	1	seq
uuuugaaugaaauuguaGu	77	1	seq
uuugaaugaaauuguaGu	13	1	seq
.aaugaaauuguaGuggcucuuuuaaguaguu	1	1	seq
gaaauuguaGuggcucuuuuaagu	1	1	seq
.aaauuguaGuggcucuuuuaaguaguu	1	1	seq
uggcucuuuaaguaguugaaC	4	1	seq
uggcucuuuaaguaguugaau	4	0	seq
uggcucuuuaaguaguCgaau	1	1	seq
uggcucuuuaaguaguugaauU	3	1	seq
uggcucuuuaaguaguugaaua	9	0	seq
uggcucuuuaaguaguugaauC	1	1	seq
uggcucuuuaaguaUuugaaua	1	1	seq
uggcucuuuaaguaguugaauaa	4	0	seq
uggcucuuuaaguGguugaauaa	1	1	seq
uggcucuuuaaguaguugaauaaA	1	1	seq
uggcucuuuaaguaguugaauaauu.	2	0	seq
.ggcucuuuuaaguaguugaau	1	0	seq
cucuuuaaguaguugGauaa.	2	1	seq
.ucuuuaaguaguugGauaa	3	1	seq
uuuuaaguaguugGauaa	1	1	seq
uuuacucacuAcuuaaaagagu	2	1	seq
uuacucacuAcuuaaaaga	1	1	seq
uuacucacugcuuaaaaga	2	0	seq
.uagucaguhguuaaaaga	79	1	seq
.uacucacuAcuuaaaagagu	1	1	seq
acucacugcuuaaaagaA	1	1	seq
acucacugcuuaaaagGg	3	1	seq
acucacugcuuaaaagGgu	3	1	seq
acucacugcuuaaaagaAu	1	1	seq
acucacugcuuaaaagaguca	1	0	seq

Star Mature

Star Ma	ature			
auuugaccugcuccucaaguuuugaaugaaauuguauuggcucuuuuaaguaguugaauaauuaaauuuacuc	cacugcuuaaaagagucacuacagcuuugucuuacu			
acuc	cacugcuuaaaagagucaU	1	1	seq
acAc	cacugcuuaaaagagucac	2	1	seq
acuc	cacugcuuaaaagagucac	3	0	seq
acuc	cacugcuuaaaagagucaG	1	1	seq
acCc.	cacugcuuaaaagagucacu	1	1	seq
acuc	• • • •	1	1	seq
acuc		1	1	seq
	cacugcuuaaaagagucacu	1	1	
				seq
acuc		1	1	seq
acuc		1	1	seq
acuc	cacugcuuaaaagagucacG	10	1	seq
acuc	cacugcuuaaaagagucUcu	1	1	seq
acuc	cacugcuuaaaagaCucacu	2	1	seq
acuc	cacugcuuaaaGgagucacu	1	1	seq
acuc	cacugcuuaaaagagucacu	192	0	seq
Gcuc	cacugcuuaaaagagucacu	10	1	seq
acuc	cacugcuuaaaagagucacA	23	1	seq
acuc	cacugcuuaaaagagucacC	44	1	seq
acuc	cacugcuuaaaagagCcacu	1	1	seq
acuc	cacCgcuuaaaagagucacu	1	1	seq
acuc		3	1	seq
acuc		13	1	seq
	cacugcuuaaaagagucacuC	5	1	seq
	cacugcuuaaaagagucacua	1	0	seq
acuc		2	1	seq
cuc	cacugcuuaaaagagu	1	0	seq
cuc	cacugcuuaaaagGgu	7	1	seq
cuc	cacugcuuaaaagaguc	4	0	seq
cuc	cacugcuuaaaagaguA	1	1	seq
	cacugcuuaaaagaguU	1	1	seq
	cacugcuuaaaagaguca	1	0	seq
cuc	cacugcuuaaaagagucU	3	1	seq
cCc	cacugcuuaaaagagucac	1	1	seq
cuc	cacugcuuaaaagagucaU	3	1	seq
	cacugcuuaaaagagucac	1	1	seq
cuc	cacugcuuaaaagGgucac	1	1	seq
cuc	cacugcuuaaaagagucac	20	0	seq
cuc	cacugcuuaaaagagucaA	1	1	seq
cuc	cacugcuuaaaagagucacu	502	0	seq
	cacugcuuaaaagagucacG	13	1	seq
	cacugcuuaaaUgagucacu	2	1	seq
cuc		6	1	seq
cuc	• • • •	7	1	seq
	caUugcuuaaaagagucacu	1	1	seq
		2	1	seq
cuc	• • •	1	1	seq
cuc		2	1	seq
cuc	cacugcuuaaaagagAcacu	1	1	seq
cuc	cacugcuuaaaagaUucacu	1	1	seq
cuc	cacugcuuaaaagagucGcu	1	1	seq
cuc	cacugcuuaaaagagucacC	107	1	seq
	cGcugcuuaaaagagucacu	3	1	seq
cuc	cacugcuuaGaagagucacu	1	1	seq
cuc	cacugcuuGaaagagucacu	1	1	seq
cuc	cacugcuuaaaagagucacA	58	1	seq
	cacugcuuaaaagagucacu	3	1	seq
cuc	cacugcuuaaaagaCucacu	1	1	seq
cuc		6	1	seq
cuc		3	1	seq
cuc		1	1	seq
cuc		10	1	seq
cuc		4	1	seq
		1	1	
		2	1	seq
uc				seq
uc		1	1	seq
uc		10	0	seq
		1	1	seq
	cGcugcuuaaaagagucacu	1	1	seq