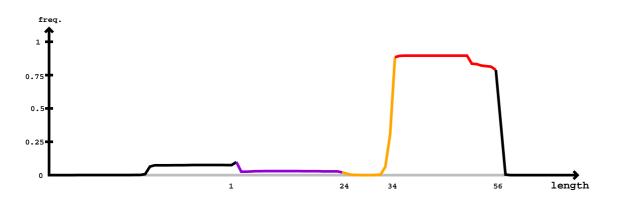
Provisional ID : NC_058073.1_Acropora_millepora_isolate_JS-1_chromosome_8_Amil_v2.1_whole_genome_shotgun_sequence_328215

Score total : 634

Score for star read(s): 3.9
Score for read counts: 629.8
Score for mfe: 0.9
Score for randfold: 0
Score for cons seed: -0.6

5' U G G C U C U U U A A G U A G U G A G U A A U C U G A G A A A A U U C G U C A C U C A U A A



5'- auuugaccugcuccucaaguuuugaaugaauuguauuggcucuuuuaaguaguugaauaauuaaauuuacucacugcuuaaaagagucacu

Star Mature

auuugaccugcuccucaaguuuugaaugaaauuguauuggcucuuuuaaguaguugaauaauuuaauuuacucacugcuuaaaagagucacuacagcuuugucuuacu	exp		
(((((())))))(.((((((((((((((((reads	mm	sample
gaccugcuccuAaaguuuugaaugaaauugu	1	1	seq
aaguuuugaaugaaauuguaGuaaguuuugaaugaauuguaGu	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
uggcucuuuuaaguaguCgaau	1	1	seq
uggcucuuuuaaguaguugaau	4	0	seq
uggcucuuuaaguaguugaaC	4	1	seq
uggcucuuuuaaguaguugaauC	1	1	seq
uggcucuuuuaaguaguugaaua	9	0	seq
uggcucuuuaaguaguugaauU	3	1	seq
uggcucuuuuaaguaUuugaaua	1	1	seq
.uggcucuuuaaguaguugaauaa	4	0	seq
uggcucuuuaaguGguugaauaa	1	1	seq
uggcucuuuuaaguaguugaauaaA	1	1	seq
	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
uacucacuAcuuaaaaga	79	1	seq
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acucacugcuuaaaagaAu	1	1	seq
acucacugcuuaaaagGgu	3	1	seq
acucacuqcuuaaaaqaquca.	1	0	seq

Star Mature

Star Ma	ature			
auuugaccugcuccucaaguuuugaaugaaauuguauuggcucuuuuaaguaguugaauaauuaaauuuacuca	acugcuuaaaagagucacuacagcuuugucuuacu			
acuca	acugcuuaaaagagucac	3	0	seq
acuca	acugcuuaaaagagucaG	1	1	seq
acuca	acugcuuaaaagagucaU	1	1	seq
acAca	acugcuuaaaagagucac	2	1	seq
	acugcuuaaaagagucacu	1	1	seq
acuca		10	1	seq
acuca		1	1	seq
		1	1	_
acuca				seq
acuca		1	1	seq
acuca		192	0	seq
acuca	acCgcuuaaaagagucacu	1	1	seq
acuca	acugcuuaaaagaUucacu	3	1	seq
acuca	acugcuuaaaagGgucacu	1	1	seq
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acuca	acugcuuaaaagagucacA	23	1	seq
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	acugcuuaaaagagucacu	10	1	seq
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	acugcuuaaaagagucacuU	13	1	seq
	acugcuuaaaagagucacuC	5	1	seq
acuca acuca		2	1	seq
		7	1	_
cuca				seq
cuca		1	0	seq
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	acugcuuaaaagaguU	1	1	seq
	acugcuuaaaagaguA	1	1	seq
cuca	acugcuuaaaagagucU	3	1	seq
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cuca	acugcuuaaaagagucaA	1	1	seq
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cCca	acugcuuaaaagagucac	1	1	seq
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cAca	acugcuuaaaagagucac	1	1	seq
cuca	acCgcuuaaaagagucacu	2	1	seq
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	acugcuuaaaagaUucacu	1	1	seq
cuca	acugcuuaaaagagucGcu	1	1	seq
cuca	acugcuuGaaagagucacu	1	1	seq
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cuca	acugcuuaaaagagucacu	502	0	seq
cuca		1	1	seq
cuca		6	1	seq
cuca		107	1	seq
cuca		6	1	_
		3	1	seq
		2	1	
				seq
cuca		58	1	seq
cucG	•	3	1	seq
cuca	• • • •	1	1	seq
cuca		2	1	seq
cuca		7	1	seq
cuca		13	1	seq
cuca	acugcuuaaaagaCucacu	1	1	seq
cuca	acugcuuaaaagagucacuC	3	1	seq
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cuca	acugcuuaaaagagucacuU	10	1	seq
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	acugcuuaaaagagucacG	1	1	seq
cg	Gcugcuuaaaagagucacu	1	1	seq