Provisional ID : NC_058066.1_Acropora_millepora_isolate_JS-1_chromosome_1_Amil_v2.1_whole_genome_shotgun_sequence_39817

Score total : 0.6

Score for star read(s) : -1.3
Score for read counts : 0
Score for mfe : 2.5

 Score for mfe
 : 2.5

 Score for randfold
 : 0

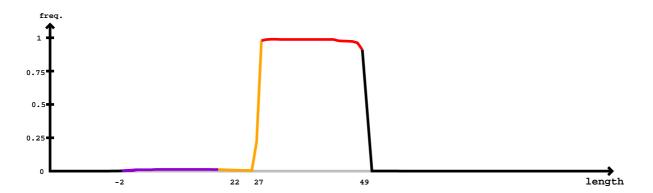
 Score for cons. seed
 : -0.6

 Total read count
 : 1245

 Mature read count
 : 1235

 Loop read count
 : 0

 Star read count
 : 10



5'- aucucguacccaagucucacucugucacuggaaaugucacauuuccagugacagagugagaucuggguacaagauuagaugccagcguaucuuucuaaauauaccucuugucucg

Star Mature

aucucguacccaagucucacucugucacuggaaaugucacauuuccagugacagagugagaucuggguacaagauuagaugccagcguaucuuucuaaauauaccucuugucucg	exp		
((((((((((((((((((((((())))))))	reads	mm	sample
.aGgucucacucugucacuggaaa	1	1	seq
gucucacucugucacuggaaU.	1	1	seq
gucucacucugucacuggaaUu.	1	1	seq
ucucacucugucacugga.	1	0	seq
ucacucugucacuggaaau.	1	0	seq
ucacucugucacuggaaaugu	3	0	seq
ucacucugucacuggaaaugucacauu	1	0	seq
ucacucugucacuggaaaugucacauC	1	1	seq
ucacucugucacuggaaaugucacauuucG	1	1	seq
ucacucugucacuggaaaugucacauuucc	1	0	seq
ucacucugucacuggaaaugucacauuucca.	1	0	seq
ucugucacuggaaauguA	1	1	seq
ucugucacuggaaaugucaU	1	1	seq
ucugucacuggaaaugucacauuuc	1	0	seq
.Cuuccagugacagagug	1	1	seq
.uuuccagugacagagug	11	0	seq
.uuuccagugacagagugU	2	1	seq
.uuuccagugacagagugaA	1	1	seq
.uuuccagugacagagugagU	1	1	seq
.uuuccagugacagagugaga	1	0	seq
.uuuccagugacagagugagaG	2	1	seq
.uuuccagugacagagugagUu	2	1	seq
.uuuccagugacagagugagaA	1	1	seq
.uuuccagugacagagugagau	3	0	seq
.uuuccagugacagagugagaAc	1	1	seq
.uuuccagugacagagugagauG	1	1	seq
.uuuccagugacagagugagauA	4	1	seq
.uuuccagugacagagugaAauc	2	1	seq
.uuuccagugacagagugagauU	44	1	seq
uuuccagugacagagugagaCcu	2	1	seq
.uuuccagugacagagugagGucu	2	1	seq
.uuuccagugacagagugagaucA	19	1	seq
.uuuccagugacagagugagaucu	151	0	seq

Star Mature

140410			
aucucguacccaagucucacucugucacuggaaaugucacau <mark>uuccagugacagagugagaucu</mark> ggguacaagauuagaugccagoguaucuuucuaaauauaccucuugucucg			
.uuuccagugacagagugGgaucu	2	1	seq
.uuuccagugacagagugagauUu	2	1	seq
.uuuccagugacagagugagaucC	14	1	seq
.uuuccagugacagagugUgaucu	1	1	seq
.uuuccagugacagagugagaucuU	1	1	seq
uuccagugacagagugG	1	1	seq
uuccagugacagagugaga	1	0	seq
uuccagugacagagugagau	4	0	seq
	1	1	seq
uuccagugacagagugagauh	2	1	seq
uuccagugacagagugGgauc	2	1	seq
uuccagugacagagugagauU	9	1	seq
uuccagugacagagugagauUu	6	1	seq
uuccagugacagagugagaCcu	4	1	seq
uuccagugacagagugagaucC	144	1	seq
uuccagugacagagugagGucu	2	1	seq
uuccagugacagagugagaucu	643	0	seq
uuccagugacagagugGgaucu	5	1	seq
uuccagugacagagCgagaucu	2	1	seq
uuccagugacagagugagaGcu	1	1	seq
uuccagugacUgagugagaucu	1	1	seq
uuccUgugacagagugagaucu	1	1	seq
uuccagugacagagugagaucA	109	1	seq
uuccagugacagagugagaucuA	4	1	seq
uuccagugacagagugagaucuC	1	1	seq
uuccagugacagagugagaucuU	4	1	seq
uuccagugacagagugagaucAg	1	1	seq
.uccagugacagagugagaucu	11	0	seq
.uccagugacagagugagaucC	1	1	seq
.ccagugacagagugagaucu	3	0	seq
	1	1	seq

seq