Provisional ID : NC_058071.1_Acropora_millepora_isolate_JS-1_chromosome_6_Amil_v2.1_whole_genome_shotgun_sequence_255810

Score total : 2.6 Score for star read(s) : -1.3

Score for read counts : 0
Score for mfe : 0.9
Score for randfold : 0
Score for cons. seed : 3

gccgaauucaau

exp

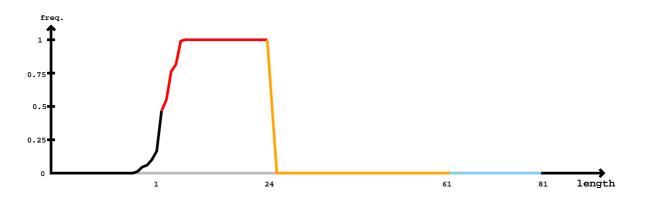
 Score for cons. seed
 :
 3

 Total read count
 :
 612

 Mature read count
 :
 612

 Loop read count
 :
 0

 Star read count
 :
 0



Mature Star

5'- uagcaauuauugaauaaggccgaguaggauaugaagaauucugcaaggcgacgaggggguguuauccacugaggucgaauu

((((((((((((((((((((((((((((((reads	mm	sample
aggcUgaguaggauaugaagaauucugc	8	1	seq
ggcUgaguaggauaugaagaauucugcggcUgaguaugagauucugc	22	1	seq
gcUgaguaggauaugaagaauucugc	9	1	seq
.cUgaguaggauaugaagaauucugc	27	1	seq
Agaguaggauaugaagaauucugc	1	1	seq
cgaguaggauaugaagaauucugU	3	1	seq
Ugaguaggauaugaagaauucugc	38	1	seq
gUguaggauaugaagaauucugc	2	1	seq
Caguaggauaugaagaauucugc	1	1	seq
gaguaggauaugaagaauucugc	167	0	seq
	8	1	seq
gaguaggauaugaagaaAucugc	1	1	seq
gCguaggauaugaagaauucugc	1	1	seq
gaUuaggauaugaagaauucugc	3	1	seq
gaguaggauaugaagaaCucugc	7	1	seq
gaguagUauaugaagaauucugc	1	1	seq
gaguaggauaugaagaGuucugc	1	1	seq
gaguaggauaugaagaauucugG	1	1	seq
gaguaggauaugaagaauCcugc	3	1	seq
aguaggauaugaagaauucCgc	2	1	seq
.aAuaggauaugaagaauucugc	2	1	seq
.Cguaggauaugaagaauucugc	1	1	seq
.aUuaggauaugaagaauucugc	1	1	seq
aguaggauaCgaagaauucugc	1	1	seq
aguaggauaugaagaauucugc	43	0	seq
aguaggauaugaagaaCucugc	3	1	seq
guaggauaugaagaauucugG	3	1	seq
guaggauGugaagaauucugcguaggauGugaagaauucugc	1	1	seq
guaggauaugaagaaAucugcguaggauaugaagaaAucugc	4	1	seq
guaggauaugaagaauucugcguaggauaugaagaauucugc	122	0	seq
guaggauaugaagaaCucugc	2	1	seq
guaggauaugaagGauucugc	2	1	seq
Uuaggauaugaagaauucugc	4	1	seq
guaggauauAaagaauucugca	1	1	seq

Mature Star

uagcaauuauugaauaaggccgaguaggauaugaagaauucugcaaggcgacgaggguguuauccacugaggucgaauucuacgaauucuauucuacgaaegccgaauucaau			
uaggauaugaagaauucugc	32	0	seq
uaggauaugaagaauucugG.	1	1	seq
uaggauaCgaagaauucugc	2	1	seq
aggauaugaagGauucugcaggauaugaagGauucugc	1	1	seq
aggauaugaagaauucugG	1	1	seq
aggauaugaagaauucGgcaggauaugaagaauucGgc	1	1	seq
aggauaugaagaauucugc	106	0	seq
aggauaugaagaGuucugcaggauaugaagaGuucugc	2	1	seq
aggauaugaagaahucugc	1	1	seq
Cggauaugaagaauucugc	1	1	seq
aggauaugaagaauCcugcaggauaugaagaauCcugc	1	1	seq
ggauaugaagaaCucugc	1	1	seq
ggauaugaagaauucugc	5	0	seq