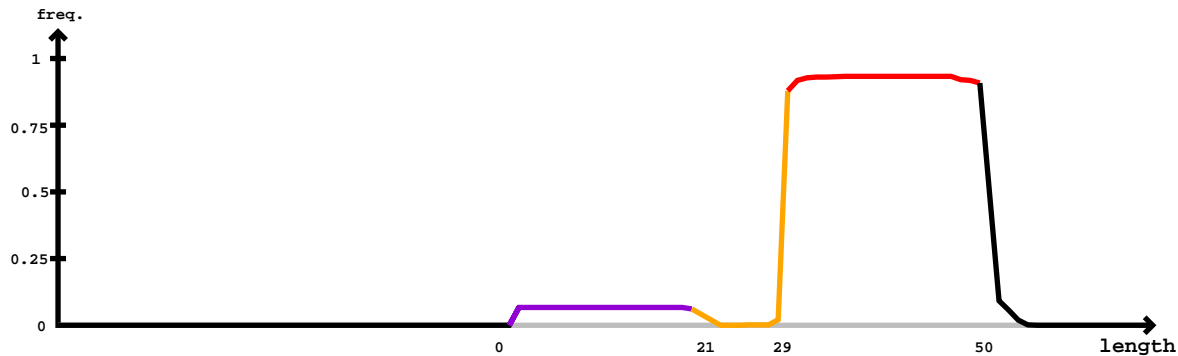
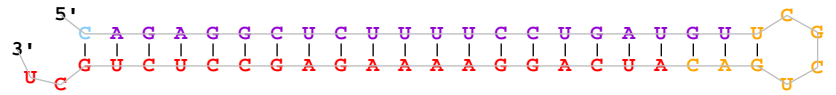


Provisional ID : NC_058076.1_Acropora_millepora_isolate_JS-1_chromosome_11_Amil_v2.1_whole_genome_shotgun_sequence_423809
Score total : 366.2
Score for star read(s) : 3.9
Score for read counts : 360.1
Score for mfe : 2.8
Score for randfold : 0
Score for cons. seed : -0.6
Total read count : 718
Mature read count : 670
Loop read count : 0
Star read count : 48



		Star	Mature			
5'	gucaaaagaugccuuggugacgaucgauugaccauucccugcuagcagaggcucuuuuccugauguucgcugacaucaggaaaagagccucugcuagcagggaagauugacc	-3'	obs			
	gucaaaagaugccuuggugacgaucgauugaccauucccugcuagcagaggcucuuuuccugauguucgcugacaucaggaaaagagccucugcuagcagggaagauugacc		exp			
	(((((.....(((.....))).....((.....))))))))))))))))))))))))))))))))))))).		reads	mm	sample	
agaggcucuuuuccugaC.....		3	1	seq	
agaggcucuuuuccugau.....		1	0	seq	
agaggcucuuuuccugaug.....		2	0	seq	
agaggcucuuuuccugaugG.....		2	1	seq	
Ggaggcucuuuuccugaugu.....		1	1	seq	
agaggcucuuuuccCgaugu.....		1	1	seq	
agaggcucuuuucccugaugu.....		23	0	seq	
agaggcucuuuuccugaugA.....		2	1	seq	
agaggcucuuuuccugaugC.....		6	1	seq	
agaggcucuuuuccugauguA.....		1	1	seq	
agaggcucuuuuccugauguu.....		3	0	seq	
agaggcucuuCuccugauguu.....		1	1	seq	
agGggcucuuuucccugauguu.....		2	1	seq	
ugacaucaggaaaagagccucu.....		1	0	seq	
caucaggaaaagagccucugcA.....		2	1	seq	
caucaggaaaagagccucugcu.....		9	0	seq	
caucaggaaaagagccucugcC.....		3	1	seq	
aucaggaaaagagccucu.....		7	0	seq	
aucaggaaaagagccucG.....		1	1	seq	
aucaggaaaagagccucuU.....		2	1	seq	
aucaggaaaagagccucugc.....		5	0	seq	
aucaggaaaagagccucugA.....		2	1	seq	
aucaggaaaagagccCcugcu.....		2	1	seq	
Gucaggaaaagagccucugcu.....		4	1	seq	
aucaggaaaUagccucugcu.....		1	1	seq	
Uucaggaaaagagccucugcu.....		1	1	seq	
aucaggaaaagagccucugcu.....		284	0	seq	
aucaggaaaagagccucugcA.....		57	1	seq	
aucaggaaaagagccucuAcu.....		1	1	seq	
aucaggaaaagagccucugcG.....		13	1	seq	
aucaggaaGagagccucugcu.....		3	1	seq	
aucGgggaaaagagccucugcu.....		1	1	seq	
aucaggaaaagagccucugcC.....		62	1	seq	

Star

Mature

gucaaaagaugccuuggugacgaucgauugaccauucccugcuagcagaggcucuuuccugauguucgcugacc	aucaggaaaagagccucugcuagcaggggaagauugacc	1	1	seq
.....	4	1	seq
.....	4	1	seq
.....	95	1	seq
.....	14	1	seq
.....	4	1	seq
.....	7	0	seq
.....	1	1	seq
.....	5	0	seq
.....	1	1	seq
.....	1	1	seq
.....	3	1	seq
.....	12	1	seq
.....	1	1	seq
.....	6	1	seq
.....	8	0	seq
.....	4	1	seq
.....	5	0	seq
.....	2	1	seq
.....	1	1	seq
.....	1	0	seq
.....	2	1	seq
.....	9	0	seq
.....	1	0	seq
.....	4	1	seq
.....	3	1	seq
.....	2	1	seq
.....	5	0	seq
.....	1	0	seq
.....	1	0	seq
.....	1	0	seq
.....	1	1	seq