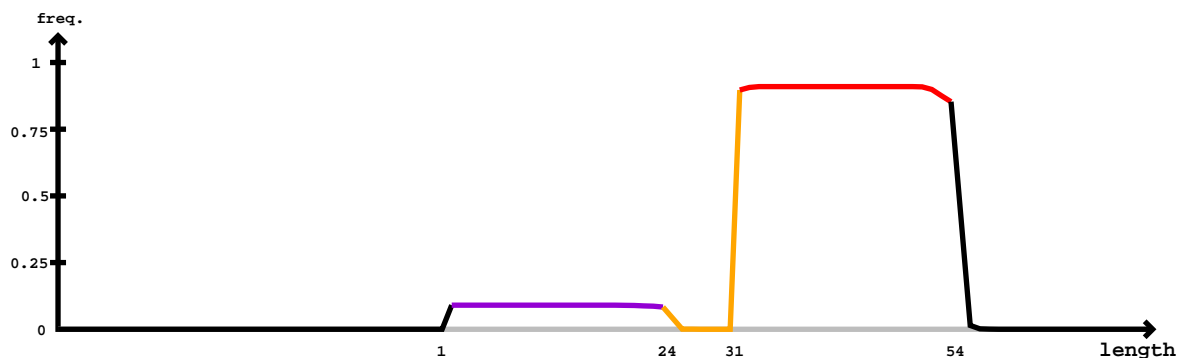
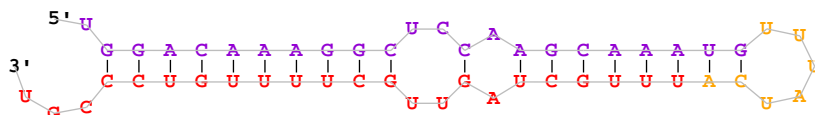


Provisional ID : NC_058067.1_Acropora_millepora_isolate_JS-1_chromosome_2_Amil_v2.1_whole_genome_shotgun_sequence_99544
 Score total : 957
 Score for star read(s) : 3.9
 Score for read counts : 948.4
 Score for mfe : 1.7
 Score for randfold : 0
 Score for cons. seed : 3
 Total read count : 1872
 Mature read count : 1703
 Loop read count : 0
 Star read count : 169



Star Mature

5' -	obs	reads	mm	sample
aaagauauugauauaccaggacuuuuucugguuaagaaauggacaaaggcuccaagcaaauguuuaucauuugcuaguugcuuuuugucccgucuuuaaacuagauauucaua	exp			
...(((((((.....(((((((.....)))))))))(((((((.....(((((((.....))))))))).....)))))).....)))))).....)))))).....				
.....uggacaaaggcuccaagc.....		1	0	seq
.....uggacaaaggcuccaagca.....		1	0	seq
.....uggacaaaggcuccaagcaa.....		3	0	seq
.....uggacaaaggcuccaagcaaa.....		2	0	seq
.....uggacaaaggcuccaagcaaaU.....		4	0	seq
.....uggacaaaggcCccaagcaaaU.....		2	1	seq
.....uggaUaaaggcuccaagcaaaug.....		1	1	seq
.....uggacaaaggcAccaagcaaaug.....		1	1	seq
.....uggacaaaggcuccaagcaaaUC.....		2	1	seq
.....Cggacaaaggcuccaagcaaaug.....		1	1	seq
.....uggacaaaggcuccaagcaaaUA.....		13	1	seq
.....uggacaaaggcCccaagcaaaug.....		1	1	seq
.....uggacaaaggcuccaagcaaaU.....		2	1	seq
.....uggacaaaggcuccGagcaaaug.....		1	1	seq
.....uggacaaaggUuccaagcaaaug.....		1	1	seq
.....uggacaaaggcuccaagcaaaug.....		126	0	seq
.....uggacaaaggcuccaagcaaaugu.....		5	0	seq
.....Aggacaaaggcuccaagcaaaugu.....		1	1	seq
.....uggacaaaggcuccaagcaaauguu.....		1	0	seq
.....uuugcuaguugcuuuuguc.....		1	0	seq
.....uuugcuaguugcuuuuguU.....		1	1	seq
.....uuugcuaguugcuuuugucU.....		3	1	seq
.....uuugcuaguugcuuuugucA.....		1	1	seq
.....uuugcuaguugcuuuugucc.....		12	0	seq
.....uuugcuaguugcuuuuguccU.....		14	1	seq
.....uuugcuaguugcuuuuguccA.....		11	1	seq
.....uuugcuaguugcuuuuguccc.....		18	0	seq
.....uuugcuaguugcuuuugucccC.....		8	1	seq
.....uuugcuaguugcuuuugucccU.....		23	1	seq
.....uuugcuaguugcuuuugucccA.....		5	1	seq
.....uuugcuaguugcuuuugucccg.....		6	0	seq
.....uuugcuaguugcCuugucccg.....		4	1	seq
.....uuugcuaguugcCuugucccg.....		3	1	seq

Mature

uuugcuaguugcuuuugucccgA	56	1	seq
uuugcuaguugcuuuuguccAgu	6	1	seq
uuugcAaguugcuuuugucccg	1	1	seq
uuugUuaguugcuuuugucccg	1	1	seq
uAugcuaguugcuuuugucccg	1	1	seq
uuugcuaguugcuuuuguccAcg	1	1	seq
Auugcuaguugcuuuugucccg	1	1	seq
uuugcuaguCgcuuuugucccg	2	1	seq
uuugcuagCugcuuuugucccg	7	1	seq
uuCgcuaguugcuuuugucccg	5	1	seq
uuugcuaguugcuuuAgucccg	1	1	seq
uuugcuaguugcuuuuguUccg	3	1	seq
uuugcuaguugcuuuCgucccg	8	1	seq
uuugcuaguugcuuuugucccU	5	1	seq
uuugcuaguugcuuuugucccgC	234	1	seq
uuugcuCguugcuuuugucccg	2	1	seq
uuugcuaguugcuuuugucccg	1	1	seq
uuugcuaguuuAcuuuugucccg	1	1	seq
uuugcuaguugcuuuuguccUgu	8	1	seq
uuugcuaguugcuuugucccg	2	1	seq
uuugcuaguugcuuuugucccCu	26	1	seq
uuugcuaguugcuuuugucccgG	29	1	seq
Cuugcuaguugcuuuugucccg	3	1	seq
uuugcuAAuugcuuuugucccg	2	1	seq
uuugcuaguugcuuuugucccAu	5	1	seq
uuugcuaguugcuuuugCcccg	3	1	seq
uuugcuaguugcuuuuguAccg	1	1	seq
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uuugcuagAugcuuuugucccg	1	1	seq
uuugcuaguugcuuuugucccg	912	0	seq
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uuugcuaguugcuuuuguccUguu	2	1	seq
uuugcuUguugcuuuugucccguu	2	1	seq
uuugcuaguugcuuuugucccUuu	3	1	seq
uuugcuaguugcuuuugucccgCu	62	1	seq
uuugcAAguugcuuuugucccguu	1	1	seq
uuugcuUguugcuuuugucccguu	1	1	seq
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uuugcuaguugcuuuugucccguu	102	0	seq
uuugcuaguugcuuuugucccguC	19	1	seq
uAugcuaguugcuuuugucccguu	1	1	seq
uCuugcuaguugcuuuugucccguu	1	1	seq
uuugcuaguugcuuuCgucccguu	1	1	seq
uuugcuaguugcuuuugucccgGu	3	1	seq
uuugcuaguugcuuuugucccgA	12	1	seq
uuugcuaguugcuuuugucccgCuc	2	1	seq
uuugcuaguugcuuuugucccguuc	1	0	seq
uuugcuaguugcuuuugucccguuA	4	1	seq
uuugcuaguugcuuuugucccguuG	2	1	seq
uuugcuaguugcuuuugucccguuU	15	1	seq
uuugcuaguugcuuuugucccguuUu	2	1	seq
uuugcuaguugcuuuugucccguuAuu	1	1	seq
uugcuaguugcuuuugucc	1	0	seq
uugcuaguugcuuuugucccg	10	0	seq
uugcuaguugcuuuugucccguu	4	0	seq
uugcuaguugcuuuugucccguC	1	1	seq
uugcuaguugcuuuugucccgA	2	1	seq
ugcuaguugcuuuugucc	1	0	seq
ugcuaguugcuuuuguccU	1	1	seq
ugcuaguugcuuuugucccg	4	0	seq