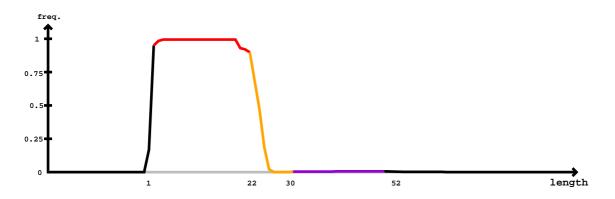
Provisional ID : NC_058067.1_Acropora_millepora_isolate_JS-1_chromosome_2_Amil_v2.1_whole_genome_shotgun_sequence_94158

255 Score total Score for star read(s) 3.9 249.4 Score for read counts 2.2 Score for mfe Score for randfold 0 Score for cons. seed -0.6 Total read count 501 499 Mature read count Loop read count 0

2

Star read count





uucucguaaaauuaaggauu <mark>auugucacguguguuuucaga</mark> aguugguucugaaaacacgcgugauauuaauccuuaauuuuacucggccccaugcaauuac	cuacag	exp		
$\dots (((((((((((((((((((((((((((((((((((($	• • • • • •	reads	mm	sample
.uauugucacguguguuuuc	• • • • • •	3	0	seq
.uauugucacguguguuuuca	• • • • • •	2	0	seq
.uauugucacguguguuuucaA	• • • • • •	1	1	seq
uauugucacguguuuucaga		36	0	seq
uauugucacguguuuCcaga	• • • • • •	1	1	seq
uauugucacguguuuucagC		2	1	seq
uauugucacguguuuucagU		1	1	seq
uauugucacguguuuucagaa	• • • • • •	8	0	seq
uauugucacguguuuucagaaU	• • • • • •	21	1	seq
uauugucacguguuuucagaaC		3	1	seq
uauugucacguguuuucagaaDu	• • • • • •	6	1	seq
auugucaUguguguuuc	• • • • • •	2	1	seq
auugucacguguuuuc		27	0	seq
	• • • • • •	3	0	seq
	• • • • • •	10	0	seq
auugucacgCguguuuucaga	• • • • • •	1	1	seq
	• • • • • •	111	0	seq
Guugucacguguuuucaga	• • • • • •	6	1	seq
auugucacgugCguuuucaga	• • • • • •	2	1	seq
	• • • • • •	2	1	seq
	• • • • • •	1	1	seq
	• • • • • •	2	1	seq
	• • • • • •	1	1	seq
auugucacgugugCuuucaga	• • • • •	2	1	seq
	• • • • • •	1	1	seq
auugucacguguuuucagaU	• • • • • •	1	1	seq
Guugucacguguuuucagaa	• • • • •	2	1	seq
	• • • • • •	26	0	seq
	• • • • • •	1	1	seq
auugucacgCguguuuucagaa	• • • • •	1	1	seq
auugucacguguguuuucagaaC	• • • • • •	10	1	seq
auugucacguguguuuucagaaU	• • • • • •	97	1	seq
auugucacguguguuuucagaaUu	• • • • •	73	1	seq

Mature	Star
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uucucguaaaauuaaggauu <mark>auugucacguguguuuucaga</mark> aguugguucugaaaacacgcgugauauuaauccuuaauuuuacucggccccaugcaauuaccuacag			
auugucacguguguuuucagaagu	1	0	seq
auugucacguguguuuucagaaUuu.	10	1	seq
uugucacguguguuuucaga	4	0	seq
uugucacguguguuuucagaa	2	0	seq
.uugucacguguguuuucagaaC	1	1	seq
uugucacguguguuuucagaaU	4	1	seq
uugucacguguguuuucagaaA	2	1	seq
.uugucacguguguuuucagaaUu	3	1	seq
uugucacguguguuuucagaaUuu.	1	1	seq
ugucacguguguuuucagaaU	4	1	seq
ugucacgCguguuuucagaagu	1	1	seq
.cugaaaacacgcgugauauu	2	0	seq
cgcgugauauuaauccuuaaAuu	1	1	seq