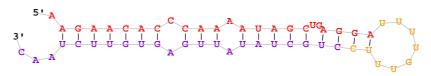
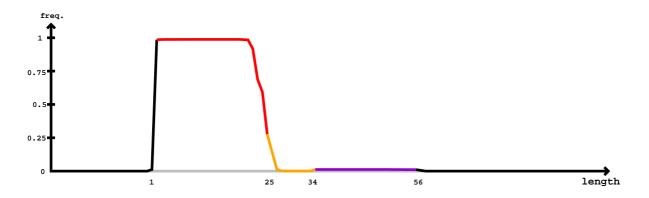
Provisional ID : NC_058070.1_Acropora_millepora_isolate_JS-1_chromosome_5_Amil_v2.1_whole_genome_shotgun_sequence_226769

2323 Score total Score for star read(s) 3.9 2314.2 Score for read counts 1.9 Score for mfe Score for randfold 0 Score for cons. seed 3 Total read count 4551 Mature read count 4497 Loop read count 0

54

Star read count





5'- uugcaaguucagcuuggaguaagaacacccaaaauagcugaggauuuuguuuccugcuaauugaguguucuaacuccugcgagagccuagugcaaggaauauugugccaggaa

Mature	Star

uugcaaguucagcuuggaguaagaacacccaaaauagcugaggauuuuguuuccugcuauauugaguguucuaacuccugcgagagccuagugcaaggaauauugugccaggaa	exp		
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uaagaacGcccaaaauagcug	1	1	seq
uaagaacacccaaaauagcuga	9	0	seq
uaagaacacccaaaauagcugag	4	0	seq
uaagaacacccaaaauagcugaU	1	1	seq
uaagaacacccaaaauagcugaA	2	1	seq
uaagaacacccaaaauagcugagA	4	1	seq
uaagaacacccaaaauagcugagU	2	1	seq
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Caagaacacccaaaauagcugagg	1	1	seq
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aagaacacccaaaauagc	3	0	seq
aagaacacccaaaauagU	3	1	seq
aagaacacccaaaauagcA	1	1	seq
aagaacacccaaaauagcC	1	1	seq
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aGgaacacccaaaauagcug	4	1	seq
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Gagaacacccaaaauagcug	9	1	seq
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aagaacacAcaaaauagcug	1	1	seq
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aagaacacccaaaauagcuUaagaacacccaaaauagcuU	14	1	seq

Mature Star

Mature	Star			
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	•••••	46	1	seq
		3 2	1	seq
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		4	1	seq
		1	1	seq
		3	1	seq
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aagGacacccaaaauagcuga		3	1	seq
aagaacacccaaaGuagcuga		1	1	seq
aagaacacccaaaauagcugG		17	1	seq
		1	1	seq
	•••••	16	1	seq
	•••••	1	1	seq
		1	1	seq
		4	1	seq
		8	1	seq
		1	1	seq
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		5 24	1	seq
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	•••••	1	1	seq
		2 1	1	seq
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		2	1	seq
		3	1	seq
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		9	1	seq
		2	1	seq
	•••••	1	1	seq
		1 276	1	seq
		3	1	seq
		4	1	seq
		4	1	seq
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aGgaacacccaaaauagcugagg		10	1	seq
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		977	0	seq
		14	1	seq
		2	1	seq
		1 2	1	seq
		1	1	seq
		10	1	seq
		3	1	seq
		2	1	seq
		4	1	seq

Mature Star

Mature Star			
uugcaaguucagcuuggaguaagaacacccaaaauagcugaggauuuuguuuccugcuauauugaguguucuaacuccugcgagagccuagugcaaggaauauugugccaggaa			
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aagUacacccaaaauagcugagg	1	1	seq
aagaacacccaaGauagcugagg	4	1	seq
	18 1	1 1	seq
aaUaacacccaaaauagcugagg	3	1	seq
aagaacacccaaaauagcugGggaagaacacccaaaauagcugAgg	1	1	seq
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gaacacccaaaauagcugag	1	0	seq
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Ma	ture	Star

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.uuuguuuccugcuauauugagugu	1	0	seq
cugcuauauugaguguuc	1	0	seq
cugcuauauugaguguucC	1	1	seq
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cugcuauaCugaguguucuaac	1	1	seq
cugcuauauugaguguucuaac	32	0	seq
cugcuauauugaguguucuaaU	2	1	seq
cugcuauauugaguguucuaacA	1	1	seq
cugcuauauugaguguucuaaUu	1	1	seq
cugcuauauugaguguucuaacu	12	0	seq
cugcuauauugaguguucuaacC	1	1	seq
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