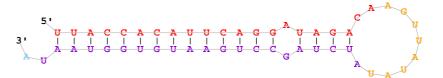
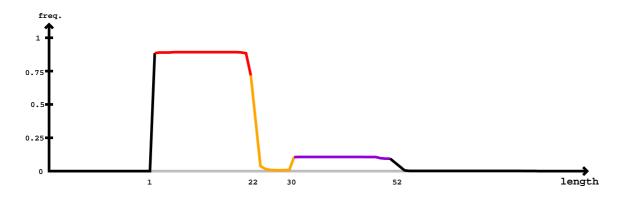
Provisional ID: NC_058069.1_Acropora_millepora_isolate_JS-1_chromosome_4_Amil_v2.1_whole_genome_shotgun_sequence_158038

918.9 Score total Score for star read(s) 3.9 914.2 Score for read counts 1.3 Score for mfe Score for randfold 0 Score for cons. seed -0.6 Total read count 1805 1626 Mature read count Loop read count 0 179 Star read count





Mature	Star
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gaucuuucagcaguuuuaau <mark>uuaccacauucaggauagaca</mark> guuauauaucuagccugaaugugguaauauuuaagcuguuccaugauuuucaauguuauauuucauuu	exp		
(((((((((((((((((((((((((((,)))).))))))))	reads	mm	sample
uuaccacauucaggauag	4	0	seq
uuaccacauucaggauaga	10	0	seq
Auaccacauucaggauaga	1	1	seq
Cuaccacauucaggauagac	2	1	seq
uuaccacauucaggauagaA	3	1	seq
uuaccacauucaggaCagac	1	1	seq
uuaccacauucaggauagac	254	0	seq
uuaccacauucGggauagac	2	1	seq
uuaccacaCucaggauagac	1	1	seq
uuaccacauucaggauagaU	37	1	seq
uuaccacauucaggauagaG	1	1	seq
uuaccacauucaAgauagac	1	1	seq
uuaccacGuucaggauagac	1	1	seq
uuaccGcauucaggauagac	2	1	seq
uuaccacauucUggauagaca	2	1	seq
uuaccacauucaggaCagaca	1	1	seq
uCaccacauucaggauagaca	5	1	seq
uuaccacaCucaggauagaca	1	1	seq
uuacVacauucaggauagaca	3	1	seq
uuaccacauucaggauagaUa	2	1	seq
uuaAcacauucaggauagaca	2	1	seq
uuaccGcauucaggauagaca	4	1	seq
uuaccacauucaggauagCca	5	1	seq
uuaccacauucaggauagacU	35	1	seq
uuaccacauucaggGuagaca	5	1	seq
uuaccacauucaggauagacC	1	1	seq
uuaccacauucaggauagaca	700	0	seq
uuaceUcauucaggauagaca	1	1	seq
uuaccacauCcaggauagaca	1	1	seq
uuaccacauucaggauCgaca	1	1	seq
uuaccacauucaggauagGca	5	1	seq
uuaccacauucaggauagaGa	1	1	seq
uuaccacauucaggauagacG	18	1	seq

Mature	Star
Matiira	ST27

Mature Star			
gaucuuucagcaguuuuaauuuaccacauucaggauagacaaguuauauaucuagccugaaugugguaauauuuaagcuguuccaugauuuucaauguuauauuucauuu			
.uuaccacGuucaggauagaca	2	1	seq
Cuaccacauucaggauagaca	3	1	seq
.uuaccacauucaggauagacaC	100	1	seq
.uuaccacauucaggauagacaG	10	1	seq
.Auaccacauucaggauagacaa.	1	1	seq
uuacUacauucaggauagacaa	1	1	seq
	315	0	seq
uuaccacauucaAgauagacaa	1	1	seq
uuaccacGuucaggauagacaa.	1	1	seq
uCaccacauucaggauagacaa.	1	1	seq
uuaccacauucaggauagacCa	1	1	seq
uuaccacauucaggaCagacaa	1	1	seq
uuaccacauucaggauagacGa.	5	1	seq
	2	0	seq
	3	1	seq
uuaccacauucaggauagacaaU	30	1	seq
uuaccacauucaggauagacaaC	4	1	seq
uuaccacauucaggauagacaaUu.	4	1	seq
uuaccacauucaggauagacaagC	4	1	seq
uuaccacauucaggauagacaagu	5	0	seq
uuaccacauucaggauagacaaguC	2	1	seq
uuaccacauucaggauagacaaguu	2	0	seq
	2	1	seq
uuaccacauucaggauagacaaguuCu	2	1	seq
uaccacauucaggauagac	1	0	seq
uaccacauucaggauagaca	4	0	seq
uaccacauucaggauagacaC	2	1	seq
uaccacauucaggauagacaaU	1	1	seq
cacauucaggauagacaC	3	1	seq
cacauucaggauagacaa	2	0	seq
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aguuCuauaucuagccugaaugu	1	1	seq
guuCuauaucuagccugaaugu	2	1	seq
guuCuauaucuagccugaaugugguaauau	4	1	seq
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auaucuagccugaaugCgguaauauaucuagccugaaugCgguaau	3	1	seq
auaucuagccugaaugugguaau	2	0	seq
aucuagccugaauguggu	8	0	seq
aucuagccugaaugugUu	1	1	seq
aucuagccugaauguggC	1	1	seq
	1	1	
aucuagccugaauguggAaucuagccugaauguggGa	1	1	seq seq
aucuagccugaauguggdaaucuagccugaauguggda	5	0	seq
	2	1	_
aucCagccugaaugugguaauaucCagccugaaugugguaaC	6	1	seq seq
aCcuagccugaaugugguaau	1	1	seq
aucuagccugaaugugguaau	74	0	seq
aucuagccugaaugugguaaG	1	1	seq
aucuagccugaaugAgguaaua	1	1	seq
aucuagccugaaugugguaauaaucuagccugaaugugguaaua	17	0	seq
accuagccugaaugugguaaua	1	1	seq
accuagccugaaugugguaauu	1	1	_
aucuagecugaaugugguaauo	1	1	seq
* * * * *	40	0	seq
aucuagccugaaugugguaauau	7	1	seq
	1	1	seq
aucuagccugaaugugguaauaG	5	0	seq
aucuagccugaaugugguaauauu	1	1	seq
aucuagUcugaaugugguaauauu			seq
aucuagccugaaugugguaauauC	1	1	seq
ucuagccugaaugugguaau	1	0	seq
ucuagccugaaugugguaauau			seq
uuaagcuguuccaugauuuucGaug	1	1	seq
uuaagcuguuccaugauuuucGauguua	2	1	seq