$Provisional\ ID \\ \hspace*{0.5cm} : NC_058067.1_A cropora_millepora_isolate_JS-1_chromosome_2_Amil_v2.1_whole_genome_shotgun_sequence_99560$

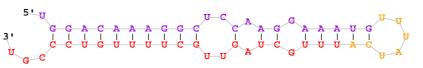
954.6 Score total Score for star read(s) 3.9 Score for read counts 945.9 1.8 Score for mfe Score for randfold 0 Score for cons. seed 3 Total read count 1867 1703 Mature read count

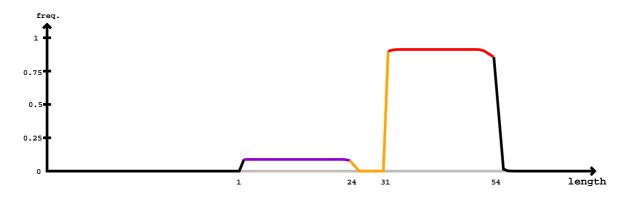
0

164

Loop read count

Star read count





Star	Mature
blai	Macure

5'- aaagguauugauaugccaguacuuuuucugguuaagaaauggacaaaggcuccaaggaaauguuuaucauuugcuaguugcuuuugucccguucuuaaacuagacauucaug

aaagguauugauaugccaguacuuuuucugguuaagaaauggacaaaggcuccaaggaaauguuuauca <mark>uuugcuaguugcuuuugucccgu</mark> ucuuaaacuagacauucaug	exp	exp	
(((((((())))))))))((((((((((((((((reads	mm	sample
	2	0	seq
	11	0	seq
uUgacaaaggcuccaaggaaau	1	1	seq
	1	1	seq
uggaUaaaggcuccaaggaaaug	1	1	seq
	2	1	seq
	1	1	seq
uggacaaaggcuccaaggaaauA	11	1	seq
uAgacaaaggcuccaaggaaaug	1	1	seq
uggacaaaggcuccaaggaaaCg	2	1	seq
	2	1	seq
	1	1	seq
	54	0	seq
	1	1	seq
uAgacaaaggcuccaaggaaaugu	1	1	seq
	1	1	seq
uggacaaaggcuccaaggGaaugu	1	1	seq
	4	1	seq
	54	0	seq
uggacaGaggcuccaaggaaaugu	1	1	seq
uggacGaaggcuccaaggaaaugu	1	1	seq
	10	1	seq
.uuugcuaguugcuuuuguc	1	0	seq
.uuugcuaguugcuuuuguU	1	1	seq
uuugcuaguugcuuuugucc	12	0	seq
.uuugcuaguugcuuuugucA	1	1	seq
uuugcuaguugcuuuugucU	3	1	seq
.uuugcuaguugcuuuuguccU	14	1	seq
.uuugcuaguugcuuuuguccA	11	1	seq
.uuugcuaguugcuuuuguccc	18	0	seq
uuugcuaguugcuuuugucccA	5	1	seq
uuugcuaguugcuuuugucccC	8	1	seq
uuugcuaguugcuuuugucccU	23	1	sea

Star Mature

Star Mature			
aaagguauugauaugccaguacuuuuucugguuaagaaauggacaaaggcuccaaggaaauguuuauca <mark>uuugcuaguugcuuuugucccguu</mark> cuuaaacuagacauucaug			
uuugcuaguugcuuuugucccg	6	0	seq
uuugcCaguuuguucccgu	2	1	seq
uuugcuaguugcuuuugucccUu	5	1	seq
	1	1	seq
	1	1	
uuugcuaguugcuuuAgucccgu			seq
uAugcuaguugcuuuuguccogu	1	1	seq
uuugcuagCugcuuuugucccgu.	7	1	seq
.uuugcuaguAgcuuuugucccgu	1	1	seq
.uuugcuaguugcuuuugucccCu	26	1	seq
uuugcuaguugcCuuuguccegu	4	1	seq
uuugcuaguuAcuuuugucccgu	1	1	seq
uuugcuaguugcuuuuguAccgu	1	1	seq
uuugcuaguugcuuuCgucccgu	8	1	seq
uuugUuaguugcuuuugucccgu	1	1	seq
uuugcuaguCgcuuuugucccgu	2	1	seq
uuugcuaguugcuuuugucccgu	912	0	seq
uuCgcuaguugcuuuugucccgu	5	1	seq
.uuugcuaguugcuuuugucccAu	5	1	seq
uuugcuaguugcuuuuguUccgu	3	1	seq
uuugcuaguugcuuuugucccgC	234	1	seq
uuugcuaguugcuuuugCcccgu	3	1	seq
uuugcuaAuugcuuuugucccgu	2	1	seq
	29	1	
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uuugcuaguugcuuuugucccgA	56	1	seq
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uuugcuaguugcuCuugucccgu	3	1	seq
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Cuugcuaguugcuuuugucccgu	3	1	seq
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	2	1	
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	62	1	seq
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uuugcuaguugcuuuugucceguu	102	0	seq
uCugcuaguugcuuuugucceguu.	1	1	seq
uugcuaguugcuuuugucccUuu	3	1	seq
uuugcAaguugcuuuugucccguu	1	1	seq
uAugcuaguugcuuuugucceguu	1	1	seq
uuugcuaguugcuuuuguceeguuG	2	1	seq
uuugcuaguuuguucccguuA.	4	1	seq
uuugcuaguuuguuccguuU	15	1	seq
uuugcuaquuqcuuuugucccquuc	1	0	seq
			_
uuugcuaguugcuuuugucccgCuc	2	1	seq
uuugcuaguugcuuuugucccguuUu.	2	1	seq
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.uugcuaguugcuuuugucceguA	2	1	seq
.uugcuaguuuguccoguu	4	0	seq
uugcuaguugcuuuugucccguC	1	1	seq
ugcuaguugcuuuugucc	1	0	seq
			_
ugcuaguugcuuuuguccU	1	1	seq
ugcuaguugcuuuugucccgu	4	0	seq