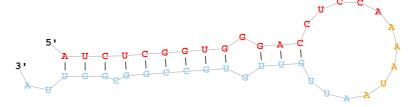
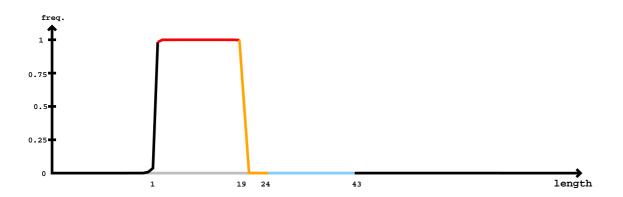
Provisional ID: NC_058074.1_Acropora_millepora_isolate_JS-1_chromosome_9_Amil_v2.1_whole_genome_shotgun_sequence_372527

Score total 1.1 Score for star read(s) -1.3 Score for read counts 0 Score for mfe -0.5 Score for randfold 0 Score for cons. seed 3 Total read count 3796 3796 Mature read count Loop read count 0 Star read count 0





Mature Star

5 '	- agccaggcaucgugguucugaucucggugggaccuccaaaauaauuguuguggcgggguuaucccaugugcaagcugacgaccgagaauacuugaucaucuaauuc	-3'	exp	
	((((((((((((((((((((((((((((((((((((read	s mm	sample
	guucAgaucucgguggaccucca	1	1	seq
	.cAgaucucgguggaccucca	1	1	seq
	Agaucucgguggaccucc	2	1	seq
	uCaucucgguggaccucc	1	1	seq
	uCaucucgguggaccucca	16	1	seq
		9	1	seq
	Uaucucgguggaccucc	1	1	seq
	Caucucggugggaccucc	2	1	seq
	Aaucucgguggaccucc	1	1	seq
	Caucucgguggaccucca	55	1	seq
	Aaucucgguggaccucca	50	1	seq
	Uaucucgguggaccucca	1	1	seq
	.aucucggCgggaccucc	1	1	seq
	.auAucggugggaccucc	1	1	seq
	.aucucgguggaccucc	2	0	seq
	.aCcucgguggaccucca.	3	1	seq
	Cucucgguggaccucca.	6	1	seq
	.aucucggugggGccucca.	10	1	seq
	.auAucggugggaccucca.	1	1	seq
	.aucucggAgggaccucca.	1	1	seq
	.aucAcgguggaccucca.	2	1	seq
	.aucucgguggaccuUca.	4	1	seq
	.aucucgguggaccuccU	107	1	seq
	.aucucgguggaccucca.	2038	0	seq
	.aucucgguggacUucca.	3	1	seq
	aucucgguggaccCcca.	21	1	seq
	.aucuUgguggaccucca.	1	1	seq
	.aucuAgguggaccucca.	2	1	seq
	.aucucgguggUaccucca.	1	1	seq
	.aucucggugAgaccucca.	3	1	seq
	aucucggugggaccucUa	11	1	seq
	.aucucgguggacAucca.	31	1	seq
	.aucucggugUgaccucca.	1	1	seq
	aucucaquaaficcucca	5	1	sea

Mature	d L
	Star

agccaggcaucgugguucug <mark>aucucggugggaccucca</mark> aaauaauuguugugccggcgguuaucccaugugcaagcugacgacccgagaauacuugaucaucuaauuc			
aucucggugggaccucAa	3	1	seq
aucucgguAggaccuccaaucucgguAggaccucca	1	1	seq
aucucgguggAaccucca	102	1	seq
aucucggugggaUcucca	8	1	seq
aucucgAugggaccucca	1	1	seq
aucucggugggaccuceG	84	1	seq
aucucgguggaccAcca	27	1	seq
aucucggugggaccGcca	18	1	seq
aucucggGgggaccucca	1	1	seq
aucucggCgggaccucca	1003	1	seq
aucucggugggaccuceC	32	1	seq
	9	1	seq
aucucgguUggaccucca	1	1	seq
aAcucggugggaccucca	2	1	seq
aucCcgguggaccucca	10	1	seq
Gucucgguggaccucca	34	1	seq
aucucgguggaccuccaa	1	0	seq
.ucCcgguggaccucca	1	1	seq
ucucggCgggaccucca	9	1	seq
.ucucgguggacUucca	1	1	seq
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.ucucggugggaccuceU	3	1	seq
.ucucggCgggaccuccaa	2	1	seq
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ucucggugggaccuceCa	1	1	seq
.ucucgguggaccuccaU	13	1	seq
.ucucgguggaccuccaVa.	7	1	seq
uGccaugugcaagcugacgcgagaaua	1	1	seq