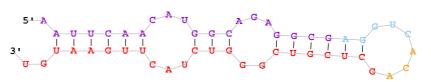
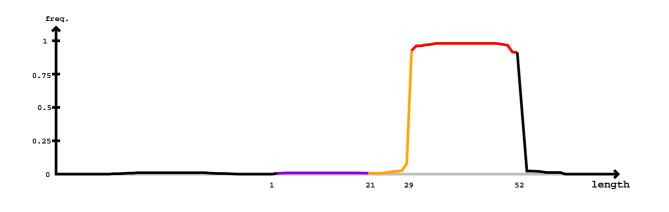
Provisional ID : NC_058078.1_Acropora_millepora_isolate_JS-1_chromosome_13_Amil_v2.1_whole_genome_shotgun_sequence_512504

313.4 Score total Score for star read(s) 3.9 Score for read counts 312.6 Score for mfe -2.4 Score for randfold 0 -0.6 Score for cons. seed Total read count 625 619 Mature read count Loop read count 0 Star read count 6



sample seq sea seq seq seq seq seq seq seq seq seq seq



Star

5'-	aauuuuaacaccguaucccacaaccgcgcgcgcgccuuauuuucaaauucaacauggcagaggcgaggucaca <mark>gcucgucgggucuacuugaaugu</mark> ucauucaguaacaggaaaug	-3'	obs	
	aauuuuaacaccguaucccacaaccgcgcgcgggccuuauuuucaaauucaacauggcagaggcgagguc <mark>acagcucgucgggucuacuugaaugu</mark> ucauucaguaacaggaaaug		exp	
	(((.((((()))((((((((((())))))	read	s n	nm
	cguaucccacaaccgcgcg	2	C)
	guaucccacaaccgcgcgcg	1	C)
	uaucccacaaccgcgcgA	1	1	L
	uaucccacaaccgcgcgcggc	1	C)
	ucccacaaccgcgcgcggcc	1	C)
	ucccacaaccgcgcgcggccu	1	C)
	aauucaacauggcagagg	1	C)
	aauucaacauggcagaggcg	2	()
	aauucaacauggcagaggcgU	1	1	L
		1	C)
	uucaacauggcagaggcgU	1	1	L
	aggucaGagcucgucgggucuacuugaaugu	2	1	L
	ggucaGagcucgucgggucuacuugaaugu	1	1	L
	.gucaGagcucgucgggucuacuugaaugu	2	1	L
	ucaGagcucgucgggucuacuugaaugu	1	1	L
	ucacagcucgucgggucuacuugaaugu	1	C)
	cacagcucgucgggucuacuugaaugu	2	C)
	caGagcucgucgggucuacuugaaugu	1	1	L
	caAagcucgucgggucuacuugaaugu	2	1	L
	acagcucgucgggucuacuugaaugu	2	C)
	cagcucgucgggucuacuugaaugu	3	C)
	agcucgucgggucuacuug	1	C)
	agcucgucgggucuacuuga	1	C)
	agcucgucgggucuacuugaa	3	()
	agcucgucgggucuacuugaU	1	1	L
	agcucgucgggucuacuugaau	2	C)
	agcucgucgggucuacuugaaugu	23	C)
	agcucgucggAucuacuugaaugu	2	1	L
	agcucgucgggucuacuugaaugG	1	1	L
	agcucgucgggucuacuugaaugA	1	1	L
	gcucgucgggucuacuugU	3	1	L
	geueguegggueuaeuugaau	24	C)
	.gcucgucgggucuacuugaUu	1	1	L

Mature

Star Mature

110.001			
aauuuuaacaccguaucccacaaccgcgcgcgcguuauuuucaaauuucagcayaggcgaggucaca <mark>gcucgucgggucuacuugaaugu</mark> ucauucaguaacaggaaaug			
geueguegggueuaeuugaaC	4	1	seq
gcucAucgggucuacuugaau	2	1	seq
.gcucgucgggucuacuugaauC	1	1	seq
gcucgucgggucuacuugGaugu	2	1	seq
geueguegggueuaeuugaaugu	430	0	seq
.gcucgucgggucCacuugaaugu	1	1	seq
.gcucgucgggucuacuugaaCgu	1	1	seq
.gcucgucgggucuacuugaUugu	6	1	seq
Acucgucgggucuacuugaaugu	9	1	seq
gcGcgucgggucuacuugaaugu	4	1	seq
.gcucgucgggucuacuCgaaugu	2	1	seq
.gcucgucggAucuacuugaaugu	21	1	seq
gcucgucgCgucuacuugaaugu	1	1	seq
.gcucgucgggucuacuugaaGgu	2	1	seq
gcucgucgggucuacAugaaugu	3	1	seq
gcucgucgggucuacuugaaugh	17	1	seq
gcucgucgggucuacuugaaugG	12	1	seq
.gcucguUgggucuacuugaaugu	1	1	seq
gcucgucgggucuacuGgaaugu	1	1	seq
.gcucgucgggucuacuugaauguu	1	0	seq
.geueguegggueuaeuugaauguuea	1	0	seq
.gcucgucgggucuacuugaauguucaC	1	1	seq
.gcucgucgggucuacuugaauguucau	1	0	seq
geueguegggueuaeuugaauguueauu	1	0	seq
geueguegggueuaeuugaauguueUuu	1	1	seq
.cucgucgggucuacuugaau	1	0	seq
.cucgucggAucuacuugaaugu	1	1	seq
.cucgucgggucuacuugaaugu	14	0	seq
cucgucgggucuacuugaauguucauucagu	7	0	seq
	5	1	seq
	1	1	seq
gucgggucuacuugaauguucauucagC	1	1	seq
ucgggucuacuugaaugu	2	0	seq
ucgggucuacuugaauguu	1	0	seq