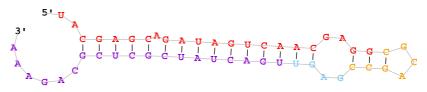
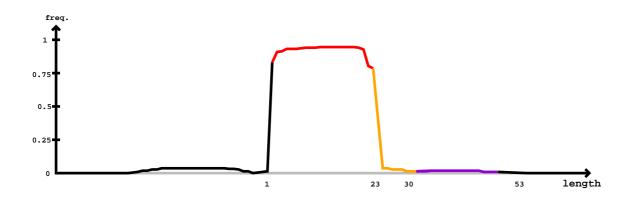
Provisional ID : NC_058066.1_Acropora_millepora_isolate_JS-1_chromosome_1_Amil_v2.1_whole_genome_shotgun_sequence_41109

0.3 Score total Score for star read(s) -1.3 Score for read counts 0 2.2 Score for mfe Score for randfold 0 Score for cons. seed -0.6 Total read count 205 203 Mature read count Loop read count 0 Star read count 2



sample



	Mature	Star		
5'- ugguaaauacuaauuaacaauuagacuacgagcccgacuu	uucuacgagcagauagucaacgaggcgcagccgag	uugacuaucgcucgcagaaaacgagggcgaguag	-3'	obs
ugguaaauacuaauuaacaauuagacuacgagcccgacuu	uucuacgagcagauagucaacgaggcgcagccgag	guugacuaucgcucgcagaaaacgagggcgaguag		exp
(((((())))).(((((((((((((.(((((((((((((((((((((((((((((((((())))))))))))))))))))))))))))))	read	s
1122022111202011202000001			1	

.....acgagcagauagucaacgagA.....

uaacaauuagacuacgagcccgU	1	1	seq
aacaauuagacuacgagcc	1	0	seq
acaauuagacuacgagcccU	1	1	seq
acaauuagacuacgagcccgU	1	1	seq
aauuagacuacgagcccga	1	0	seq
aauuagacuacgagcccgacC	1	1	seq
uuagacuacgagecegacu	2	0	seq
.uucuacgagcagauagucaacgaggcgcagc	1	0	seq
ucuacgagcagauagucaacgaggU	1	1	seq
cuacgagcagauagucaacgaCg	1	1	seq
.uacgagcagauagucaac	1	0	seq
uacgagcagauagucaacg	3	0	seq
.uacgagcagauagCcaacga	1	1	seq
.uacgagcagauagucaacga	22	0	seq
.uacgagcagauagucaacgG	1	1	seq
.uUcgagcagauagucaacgag	1	1	seq
.uacgagcagauagucaacgaC	1	1	seq
uacgagcagauagucaacgag	2	0	seq
uacgagcagauagucaacgagC	1	1	seq
.uacgagcagauagucaacgaCg	28	1	seq
uacgagcagauagucaacgGgg	1	1	seq
.uacgagcagauagucaacgagA	25	1	seq
uacgagcagauagCcaacgagg	1	1	seq
.uacgagUagauagucaacgagg	1	1	seq
uacgagcagauagucaacgagg	84	0	seq
uaUgagcagauagucaacgagg	1	1	seq
.uGcgagcagauagucaacgagg	2	1	seq
uacgagcagauagucaacgaggA	2	1	seq
uacgagcagauagucaacgaggcgcagc	1	0	seq
.acgagcagauagucaacga	3	0	seq
.acgagcagauagucaacgaUg	1	1	seq
acgagcagauagucaacgagg	4	0	seq

Mature	Star
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ugguaaauacuaauuaacaauuagacuacgagcccgacuuuucu <mark>acgagcagauagucaacgaggc</mark> gcagccgaguugacuaucgcucgcagaaaacgagggcgaguag			
.acgagcagauagucaacgagC	1	1	seq
acgagcagauagucaacgaCg	6	1	seq
acgagcagauagucaacgaggU	1	1	seq
	1	0	seq
gagcagauagucaacgagg	1	0	seq
gagcagauagucaacgagA	2	1	seq
gagcagauagucaacgaggcgc	1	0	seq
cagauagucaacgaggcgU	1	1	seq
agauagucaacgaggcgcagU	1	1	seq
uagucaacgaggcgcagccgagu	1	0	seq
.cgcagccgaguugacuaucgcC	2	1	seq
ugacuaucgcucgcagaaa	2	0	seq