$Provisional\ ID \\ : NW_025322663.1_A cropora_millepora_isolate_JS-1_unplaced_genomic_scaffold_A mil_v2.1_Sc0000161_whole_genome_shotgun_sequenter. \\ [-2.5]$

Score total : 0.6

Score for star read(s) : -1.3
Score for read counts : 0
Score for mfe : 2.5

5'CCGCGCGCAUUUCAAACUUCAACCGAAUUUUCAACCGAACUUCAACG

sample

 Score for mfe
 : 2.5

 Score for randfold
 : 0

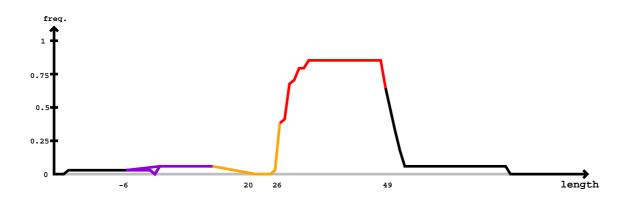
 Score for cons. seed
 : -0.6

 Total read count
 : 31

 Mature read count
 : 29

 Loop read count
 : 0

 Star read count
 : 2



		Star Mature	Star
ob	-3'	ccgcguguaagcguaaaaguugagcg <mark>agguucaacuuuuacguuuacgcgcggcu</mark> uuucuuacaucaccucuauuuuaauuuacgcacguaaagcc	5'- gauguaugaaaagccgcguguaagcguaa
ex		ccgcguguaagcguaaaaguugagcgagguuc <mark>aacuuuuacguuuacgcgcggcuuuucuuacaucaccucuauuuuaauuuacgcacguaaagcc</mark>	gauguaugaaaagccgcguguaagcguaa
ds	re	(((((((((((((((((((((((((((((((((((((((

uacaucaccucuauuuuaaC....

.uaagcguaaaaguugagcg	2	0	seq
.caacuuuuacgcucggggU	1	1	seq
aacuuuuacguuuacgcgcgggU	2	1	seq
aacuuuuacguuuacgcgcggc	1	0	seq
aacuuuuacguuuacgcgcggcu	5	0	seq
aacuuuuacguuuacgcgcgggcuu	4	0	seq
acuuuuacguuuacgcgcggcu	1	0	seq
.cuuuacguuuacgcgcgggU	2	1	seq
.cuuuacguuuacgcgcggc	1	0	seq
.cuuuacguuuacgcgcggcu	1	0	seq
.cuuuacguuuacgcgcAgcuuu	2	1	seq
.cuuuacguuuacgcgcgggcuuu	2	0	seq
.cuuuacguuuacgcgcggcuuuA	1	1	seq
uuuuacguuuacgcgcgggcuuC	1	1	seq
uuuacguuuacgcgcggcuuuu	3	0	seq
uacguuuacgcgcggcuuuuc	1	0	seq
uacguuuacgcgcggcuuuucu	1	0	seq
uuacaucGccucuauuuuaau	1	1	sea