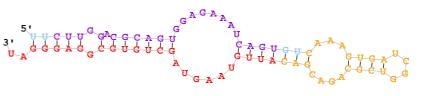
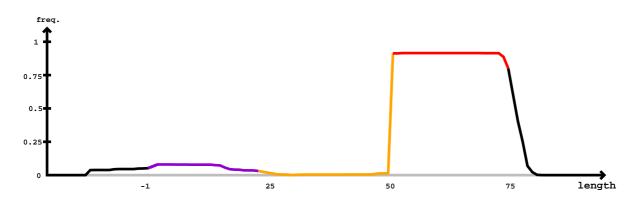
$Provisional\ ID \\ : NW_025322671.1_A cropora_millepora_isolate_JS-1_unplaced_genomic_scaffold_A mil_v2.1_Sc0000176_whole_genome_shotgun_sequenters and the provisional provi$

2.8 Score total Score for star read(s) -1.3 Score for read counts 0 Score for mfe 1.1 Score for randfold 0 Score for cons. seed 3 Total read count 1323 1268 Mature read count Loop read count 6 Star read count 49





Star Mature
5'- ucucgaauagcgugguauuuucuuggacgcaguggagaaaucagugucaaagugaucggucgcagacgacauuguaaguagcugugcggagggauggagguuccaugccaguua

ucucgaauagcgugguauu:ucuuggacgcaguggagaaaucagugucaaagugaucggucgcagacgacauuguaaguagcugugcggagggauggagguuccaugccaguua	exp		
((((((((((((((((((((((((((((((((((reads	mm	sample
uagcgugguauuuucuuC	2	1	seq
uagcgugguauuuucuuggUc	1	1	seq
uaUcgugguauuuucuuggacgcagu.	1	1	seq
uagcgugguauuuucuuggacgcagu	2	0	seq
uagcgugguauuuucuuggacgcagC	1	1	seq
uagcgugguauuuucuuggacgcagug.	2	0	seq
uagcgugguauuuucuuggacgcagugA	9	1	seq
uagcgugguauuuucuuggacgcagugg.	13	0	seq
uagcgugguauuuucuuggacgcagugga.	15	0	seq
uagcguhguauuuucuuggacgcagugga.	1	1	seq
uagcgugguauuuucuuggacgcaguggU	1	1	seq
uagcgugguauuuucuuggacgcaguggaU	1	1	seq
uagcgugguauuuucuuggacgcaguggaA	1	1	seq
uagcgugguauuuucuuggacgcaguggag	2	0	seq
uagcgugguauuuucuuggaUgcaguggag	1	1	seq
agcgugguauuuucuuggacgcagug	1	0	seq
gguauuuucuuggacgcagC	1	1	seq
gguauuCucuuggacgcaguggaga.	1	1	seq
gguauuuucuuggacgcaguggagaa	5	0	seq
gguauuuucuuggacgcaguggagaaau	1	0	seq
gguauuuucuuggacgcaguggagaa	2	0	seq
uuuucuuggacgcaguggagaaauc	1	0	seq
uuuucuuggacgcaguggagaaaucagu.	3	0	seq
uuuucuuggacgcaguggagaaaucaguA	1	1	seq
uuucuuggacgcaguggagaaaucaguU	1	1	seq
uucuuggacgcaguggagaaauc	2	0	seq
uucuuggacgcaguggagaaaucaUuguca	1	1	seq
ucuuggacgcaguggagaaauc	2	0	seq
ucuuggacgcaguggagaaaucag.	5	0	seq
ucuuggacgcaguggagaaaucagu.	5	0	seq
ucuuggacgcaguggagaaaucaguA	1	1	seq
ucuuggacgcaguggagaaaucagugu	1	0	seq
ucuuggacgcaguggagaaaucaguguc	1	0	seq

Star Mature

nature in the second se			
ucucgaauagcgugguauuuucuuggacgcaguggagaaaucagugucaaagugaucggucgcagacgac <mark>auuguaaguagcugugcggagggauggagguuccaugcca</mark> gua			
ucuuggacgcaguggagaaaucaguguca	1	0	seq
ucuuggacgcaguggagaaaucaUuguca	2	1	seq
ucuuggacgcaguggagaaaucaguguUa	1	1	seq
.cuuggacgcaguggagaaauc	1	0	seq
.cuuggacgcaguggagaaGucagu	1	1	seq
.cuuggacgcaguggagaaaucagA	1	1	seq
.cuuggacgcaguggagaaaucagC	3	1	seq
cuuggacgcaguggagaaaucagu	12	0	seq
cuuggacgcaguggagaaaucagug	1	0	seq
cuuggacgcaguggagaaaucaguguca	1	0	seq
gacgcaguggŪgaaaucagu	1	1	seq
agugaucggucgcagacgaU	1	1	seq
agugaucggucgcagacgac	3	0	seq
gugaucggucgcagacgac	1	0	seq
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.cgacauuguaagCagcugugcggaggg	2	1	seq
	1	1	seq
gacauuguaagCagcugugcggaggga	1	1	seq
gacauuguaagCagcugugcggagggau	4	1	seq
gacauuguaagCagcugugcggaagggaug	3	1	seq
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auuguaagCagcugugcggaggg	36	1	seq
auuguaagCagcugugcggaggga	121	1	seq
auuguaagCagcugugcggagggau	439	1	seq
auuguaaguagcugugcggagggaC	1	1	seq
auuguaagCagcugugcggagggaug	105	1	seq
auuguaagCagcugugcggaagggaugg	217	1	seq
auuguaagCagcugugcggaugga	255	1	seq
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uguaagCagcugugcggaauggaaggu	3	1	seq