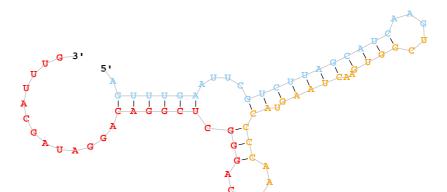
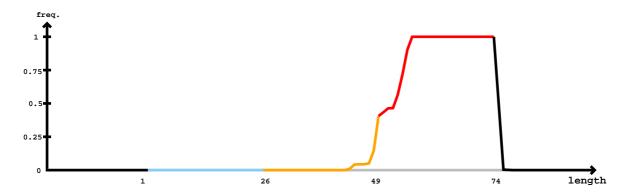
$Provisional\ ID \\ : NW_025322650.1_A cropora_millepora_isolate_JS-1_unplaced_genomic_scaffold_A mil_v2.1_Sc0000139_whole_genome_shotgun_sequential milliplaced_genomic_scaffold_A milliplaced_genomic_scaffold_genomic_scaffold_genomic_scaffold_genomic_scaffold_genomic_scaffold_genomic_scaffold_genomic_scaffold_genomic_scaffold_genomic_scaffold_genomic_scaffold_genomic_sc$

2 Score total Score for star read(s) -1.3 Score for read counts 0 Score for mfe 0.3 Score for randfold 0 Score for cons. seed 3 Total read count 547 547 Mature read count Loop read count 0 Star read count 0



ccagggcucggacaggauagcauuugaggugggaagagugcuccuc



Star Mature

, -	· uugeaaaueueueaeeeeaaguuugaauuegueuuageaueaagueggugaaeuaaguaeeeeaaueeagggeueggaaaggauuugaggugguggugeueeue	-3. exp		
	(((((((((((((((((((((((((((((((((((reads	mm	sample
	.ccaauccagggcucggacaggauugcauuuh	3	1	seq
	.ccaauccagggcucggacaggauagcauuug	3	0	seq
	caauccagUgcucggacaggauagcauuug	7	1	seq
		3	1	seq
	caauccaCggcucggacaggauagcauuug	1	1	seq
	caaCccagggcucggacaggauagcauuug	1	1	seq
	caauccagggcucggacaggauagcauuug	4	0	seq
		2	1	seq
	aauccagUgcucggacaggauagcauuug	1	1	seq
	uccagUgcucggacaggauagcauuug	4	1	seq
	ccagggcucggacaAgauagcauuug	1	1	seq
	caCggcucggacaggauagcauuug	5	1	seq
	ccagggcucggacaggauagcauuug	7	0	seq
	ccagggcucggacaggauagcauuuA	2	1	seq
	ccagUgcucggacaggauagcauuug	37	1	seq
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	.cagUgcucggacaggauagcauuug	105	1	seq
	cagCgcucggacaggauagcauuug	4	1	seq
	caCggcucggacaggauagcauuug	6	1	seq
	cagggcucggacaAgauagcauuug	8	1	seq
	cagggcucggacaggauagcauuug	24	0	seq
	cagggcucggacGggauagcauuug	1	1	seq
	cagUgcucggacaggauagcauuugag	1	1	seq
	agUgcucggacaggauagcauuug	12	1	seq
	agggcucggacaggauagcauuug	2	0	seq
	agggcucggacaggauagcauuuA	2	1	seq
	Cggcucggacaggauagcauuug	2	1	seq
	gUgcucggacaggauagcauuug	12	1	seq
	gggcucggacaggauagcauuuA	1	1	seq
	gggcucggacaggauagcauuug	1	0	seq
	gUgcucggacaggauagcauuugagg	1	1	seq
	ggcucggacaggauagcauuug	1	0	seq
	.gcucggacaggauagcCuuug	1	1	seq
	genegal Cagganageaning .	1	1	sea

Star Mature

uugcaaaucucucaccccaaguuugaauucgucuuagcaucaagucggugaacuaaguaccccaauccagggcucggacaggauagcauuugaggugggaagagugcuccuc			
	1	1	seq
gcucggacaggauagUauuug	2	1	seq
geueggacaggauaUcauuug	4	1	seq
gcuUggacaggauagcauuug	1	1	seq
geueggaeaggauageGuuug	1	1	seq
	3	1	seq
geueggacaggauagcauuug	43	0	seq
cucggacaUgauagcauuug	1	1	seq
cuUggacaggauagcauuug	2	1	seq
cucggacaggauaAcauuug	5	1	seq
cucggacaggauagcauuuC	1	1	seq
cucggacaggGuagcauuug	3	1	seq
cucggacaggauagcauuCg	1	1	seq
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cucggacaggauagUauuug	4	1	seq
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.uAggacaggauagcauuug	1	1	seq
ucggGcaggauagcauuug	3	1	seq
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.Ccggacaggauagcauuug	3	1	seq
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ucggacaggauagcauuug	84	0	seq
	1	1	seq
ucggacaggauagcauuuC	1	1	seq
ucggacaggauaUcauuug	1	1	seq
ucggacaggaAagcauuug	1	1	seq
.cggacaggauagcauuuC	2	1	seq
.cggacaggauagcauuug	48	0	seq
Aggacaggauagcauuug	1	1	seq
cggacaggauagUauuug	5	1	seq