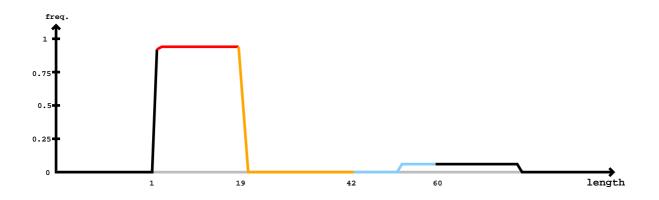
$Provisional\ ID \\ : NW_025322685.1_A cropora_millepora_isolate_JS-1_unplaced_genomic_scaffold_A mil_v2.1_Sc0000197_whole_genome_shotgun_sequenters. \\ [20pt] Provisional\ ID \\ : NW_025322685.1_A cropora_millepora_isolate_JS-1_unplaced_genomic_scaffold_A mil_v2.1_Sc0000197_whole_genome_shotgun_sequenters. \\ [20pt] Provisional\ ID \\ : NW_025322685.1_A cropora_millepora_isolate_JS-1_unplaced_genomic_scaffold_A mil_v2.1_Sc0000197_whole_genome_shotgun_sequenters. \\ [20pt] Provisional\ ID \\ : NW_025322685.1_A cropora_millepora_isolate_JS-1_unplaced_genomic_scaffold_A mil_v2.1_Sc0000197_whole_genome_shotgun_sequenters. \\ [20pt] Provisional\ ID \\ : NW_025322685.1_A cropora_millepora_isolate_JS-1_unplaced_genomic_scaffold_A mil_v2.1_Sc0000197_whole_genome_shotgun_sequenters. \\ [20pt] Provisional\ ID \\ : NW_025322685.1_A cropora_millepora_isolate_JS-1_unplaced_genomic_scaffold_A mil_v2.1_Sc0000197_whole_genome_shotgun_sequenters. \\ [20pt] Provisional\ ID \\ : NW_025322685.1_A cropora_millepora_isolate_JS-1_unplaced_genomic_scaffold_A mil_v2.1_Sc0000197_whole_genome_shotgun_sequenters. \\ [20pt] Provisional\ ID \\ : NW_025322685.1_A cropora_millepora_isolate_JS-1_unplaced_genomic_scaffold_A mil_v2.1_Sc0000197_whole_genome_shotgun_sequenters. \\ [20pt] Provisional\ ID \\ : NW_025322685.1_A cropora_millepora_isolate_JS-1_unplaced_genomic_scaffold_A mill_v2.1_Sc0000197_whole_genome_shotgun_sequenters. \\ [20pt] Provisional\ ID \\ : NW_025322685.1_A cropora_millepora_isolate_JS-1_unplaced_genome_shotgun_sequenters. \\ [20pt] Provisional\ ID \\ : NW_025322685.1_A cropora_millepora_isolate_JS-1_unplaced_genome_shotgun_sequenters. \\ [20pt] Provisional\ ID \\ : NW_025322685.1_A cropora_millepora_isolate_JS-1_unplaced_genome_shotgun_sequenters. \\ [20pt] Provisional\ ID \\ : NW_025322685.1_A cropora_isolate_JS-1_unplaced_genome_shotgun_sequenters. \\ [20pt] Provisional\ ID \\ : NW_025322685.1_A cropora_isolate_JS-1_unplaced_genome_shotgun_sequenters. \\ [20pt] Provisional\ ID \\ : NW_025322685.1_A cropora_isolate_JS-1_unplaced_genome_shotgun_seq$

0.2 Score total Score for star read(s) -1.3 Score for read counts 0 Score for mfe -1.4 Score for randfold 0 Score for cons. seed 3 Total read count 47 47 Mature read count Loop read count 0 Star read count 0

Mature





5'- aaugacacuagcaccaggg <mark>cuguggugauuacuuauu</mark> gguuuguuacauucacagauaacaggauaagcuucaauccaauaccuuccauggcuaggcaguagccagaacaca	-3' exp	•	
(((((((((((((((())))))	reads	mm	sample
cuguggugauGacuuauu	46	1	seq
uguggugauGacuuauu	1	1	seq
cuucaauccauuccauggcu	2	0	seq
	1	1	

Star