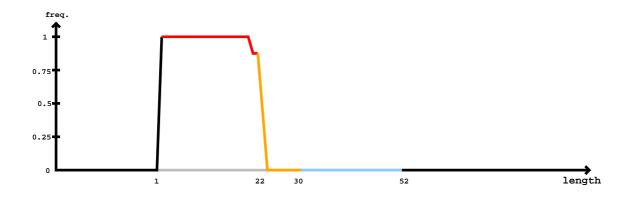
$Provisional\ ID \\ : NW_025322689.1_A cropora_millepora_isolate_JS-1_unplaced_genomic_scaffold_A mil_v2.1_Sc0000203_whole_genome_shotgun_sequenters. \\ [20pt] Provisional\ ID \\ : NW_025322689.1_A cropora_millepora_isolate_JS-1_unplaced_genomic_scaffold_A mil_v2.1_Sc0000203_whole_genome_shotgun_sequenters. \\ [20pt] Provisional\ ID \\ : NW_025322689.1_A cropora_millepora_isolate_JS-1_unplaced_genomic_scaffold_A mil_v2.1_Sc0000203_whole_genome_shotgun_sequenters. \\ [20pt] Provisional\ ID \\ : NW_025322689.1_A cropora_millepora_isolate_JS-1_unplaced_genomic_scaffold_A mil_v2.1_Sc0000203_whole_genome_shotgun_sequenters. \\ [20pt] Provisional\ ID \\ : NW_025322689.1_A cropora_millepora_isolate_JS-1_unplaced_genomic_scaffold_A mil_v2.1_Sc0000203_whole_genome_shotgun_sequenters. \\ [20pt] Provisional\ ID \\ : NW_025322689.1_A cropora_millepora_isolate_JS-1_unplaced_genomic_scaffold_A mil_v2.1_Sc0000203_whole_genome_shotgun_sequenters. \\ [20pt] Provisional\ ID \\ : NW_025322689.1_A cropora_millepora_isolate_JS-1_unplaced_genomic_scaffold_A mil_v2.1_Sc0000203_whole_genome_shotgun_sequenters. \\ [20pt] Provisional\ ID \\ : NW_025322689.1_A cropora_millepora_isolate_JS-1_unplaced_genomic_scaffold_A mil_v2.1_Sc0000203_whole_genome_shotgun_sequenters. \\ [20pt] Provisional\ ID \\ : NW_025322689.1_A cropora_millepora_isolate_JS-1_unplaced_genomic_scaffold_A mill_v2.1_Sc0000203_whole_genome_shotgun_sequenters. \\ [20pt] Provisional\ ID \\ : NW_025322689.1_A cropora_millepora_isolate_JS-1_unplaced_genome_shotgun_sequenters. \\ [20pt] Provisional\ ID \\ : NW_025322689.1_A cropora_millepora_isolate_JS-1_unplaced_genome_shotgun_sequenters. \\ [20pt] Provisional\ ID \\ : NW_025322689.1_A cropora_millepora_isolate_JS-1_unplaced_genome_shotgun_sequenters. \\ [20pt] Provisional\ ID \\ : NW_025322689.1_A cropora_isolate_JS-1_unplaced_genome_shotgun_sequenters. \\ [20pt] Provisional\ ID \\ : NW_025322689.1_A cropora_isolate_JS-1_unplaced_genome_shotgun_sequenters. \\ [20pt] Provisional\ ID \\ : NW_025322689.1_A cropora_isolate_JS-1_unplaced_genome_shotgun_seq$

1.4 Score total Score for star read(s) -1.3 Score for read counts -1.8 Score for mfe 1.6 Score for randfold 0 Score for cons. seed 3 Total read count 8 8 Mature read count Loop read count 0 Star read count 0

Mature





Star

.....cuuuugauuuauggcccacAug.

	exp	-3'	5'- uuaugggucuucauuuuucc cuuuugauuuauggcccacgugcuuugca cuuggcgcauaaaucaaaggguuaggguuaggguugguccguaagugucauuuuccacug
m sample	ls	read	.(((((.((((((((((((((((((((((,))))))).)))
) seq		1	.cuuugauuuauggcccac