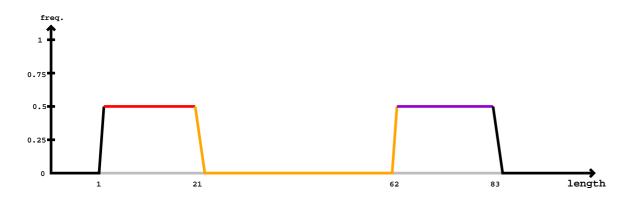
$Provisional\ ID \\ : NW_025322644.1_A cropora_millepora_isolate_JS-1_unplaced_genomic_scaffold_A mil_v2.1_Sc0000127_whole_genome_shotgun_sequenters. \\ [20pt] Provisional\ ID \\ : NW_025322644.1_A cropora_millepora_isolate_JS-1_unplaced_genomic_scaffold_A mil_v2.1_Sc0000127_whole_genome_shotgun_sequenters. \\ [20pt] Provisional\ ID \\ : NW_025322644.1_A cropora_millepora_isolate_JS-1_unplaced_genomic_scaffold_A mil_v2.1_Sc0000127_whole_genome_shotgun_sequenters. \\ [20pt] Provisional\ ID \\ : NW_025322644.1_A cropora_millepora_isolate_JS-1_unplaced_genomic_scaffold_A mil_v2.1_Sc0000127_whole_genome_shotgun_sequenters. \\ [20pt] Provisional\ ID \\ : NW_025322644.1_A cropora_millepora_isolate_JS-1_unplaced_genomic_scaffold_A mil_v2.1_Sc0000127_whole_genome_shotgun_sequenters. \\ [20pt] Provisional\ ID \\ : NW_025322644.1_A cropora_millepora_isolate_JS-1_unplaced_genomic_scaffold_A mil_v2.1_Sc0000127_whole_genome_shotgun_sequenters. \\ [20pt] Provisional\ ID \\ : NW_025322644.1_A cropora_millepora_isolate_JS-1_unplaced_genomic_scaffold_A mil_v2.1_Sc0000127_whole_genome_shotgun_sequenters. \\ [20pt] Provisional\ ID \\ : NW_025322644.1_A cropora_millepora_isolate_JS-1_unplaced_genomic_scaffold_A mil_v2.1_Sc0000127_whole_genome_shotgun_sequenters. \\ [20pt] Provisional\ ID \\ : NW_025322644.1_A cropora_millepora_isolate_JS-1_unplaced_genomic_scaffold_A mill_v2.1_Sc0000127_whole_genome_shotgun_sequenters. \\ [20pt] Provisional\ ID \\ : NW_025322644.1_A cropora_millepora_isolate_JS-1_unplaced_genome_shotgun_sequenters. \\ [20pt] Provisional\ ID \\ : NW_025322644.1_A cropora_millepora_isolate_JS-1_unplaced_genome_shotgun_sequenters. \\ [20pt] Provisional\ ID \\ : NW_025322644.1_A cropora_millepora_isolate_JS-1_unplaced_genome_shotgun_sequenters. \\ [20pt] Provisional\ ID \\ : NW_025322644.1_A cropora_isolate_JS-1_unplaced_genome_shotgun_sequenters. \\ [20pt] Provisional\ ID \\ : NW_025322644.1_A cropora_isolate_JS-1_unplaced_genome_shotgun_sequenters. \\ [20pt] Provisional\ ID \\ : NW_025322644.1_A cropora_isolate_JS-1_unplaced_genome_shotgun_seq$

0.2 Score total Score for star read(s) 3.9 Score for read counts -4.9 Score for mfe -1.6 Score for randfold 0 Score for cons. seed 3 Total read count 2 1 Mature read count Loop read count 0 Star read count 1





Mature Star

3 -	ageauuguaaaugugugugugugaagaeggguuauaagggaaaaeaggueaeuueegeuugeegueegeegueueuaaauegegegugeuuaageueeeuaauagaueagag	-3. ODS	,	
	agcauuguaaaugugauggugugaagacggguuauaagggaaaacaggucacuuccgcuugccgucgucgucuuaaaucgcgcgugcuuaagcucccuaauagaucagag	exp	,	
	(((.((((((.(((((((((((((((((((((((((reads	mm	sample
	aaugugauggugugaagacg	1	0	seq
	ucucuaaaucqcqcquqcuua	1	0	seq