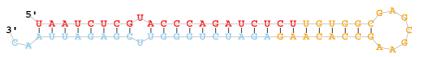
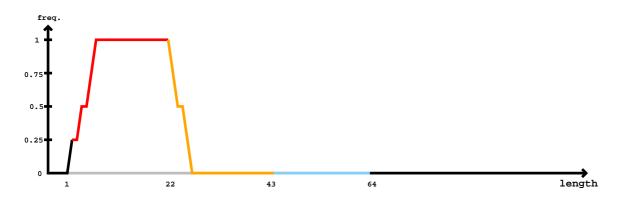
$Provisional\ ID \\ : NW_025322616.1_A cropora_millepora_isolate_JS-1_unplaced_genomic_scaffold_A mil_v2.1_Sc0000028_whole_genome_shotgun_sequenters. \\ [20pt] Provisional\ ID \\ : NW_025322616.1_A cropora_millepora_isolate_JS-1_unplaced_genomic_scaffold_A mil_v2.1_Sc0000028_whole_genome_shotgun_sequenters. \\ [20pt] Provisional\ ID \\ : NW_025322616.1_A cropora_millepora_isolate_JS-1_unplaced_genomic_scaffold_A mil_v2.1_Sc0000028_whole_genome_shotgun_sequenters. \\ [20pt] Provisional\ ID \\ : NW_025322616.1_A cropora_millepora_isolate_JS-1_unplaced_genomic_scaffold_A mil_v2.1_Sc00000028_whole_genome_shotgun_sequenters. \\ [20pt] Provisional\ ID \\ : NW_025322616.1_A cropora_millepora_isolate_JS-1_unplaced_genomic_scaffold_A mil_v2.1_Sc00000028_whole_genome_shotgun_sequenters. \\ [20pt] Provisional\ ID \\ : NW_025322616.1_A cropora_millepora_isolate_JS-1_unplaced_genomic_scaffold_A mil_v2.1_Sc00000028_whole_genome_shotgun_sequenters. \\ [20pt] Provisional\ ID \\ : NW_025322616.1_A cropora_millepora_isolate_JS-1_unplaced_genomic_scaffold_A cropora_isolate_JS-1_unplaced_genomic_scaffold_A cropora_isolate_JS-1_unplaced_genomic_scaffold_A cropora_isolate_JS-1_unplaced_genomic_Scaffold_A cropora_isolate_JS-1_unplaced_genomic_A cropora_isolate_JS-1_unplaced_genomic_A cropora_isolate_JS-1_unplaced_genomic_A cropora_isolate_JS-1_unplace$

0.1 Score total Score for star read(s) -1.3 Score for read counts -3.9 2.3 Score for mfe Score for randfold 0 Score for cons. seed 3 Total read count 4 Mature read count Loop read count 0

Star read count





Mature Star

0

5'- uccuaaucucguacccagaucucuuguggcgagcgaagccacaagagaucuggguucgagauuaacccugacccauuagcaauucgcuacuccucuggugcugugcuugc	-3' exp		
((((((((((((((((((((((((((((((((((((reads	mm	sample
uaaucucguacccagaucucu	1	0	seq
aucucguacccagaucucuA	1	1	seq
cucguacccagaucucuugu	1	0	seq
name and a condition of the condition of	1	1	202