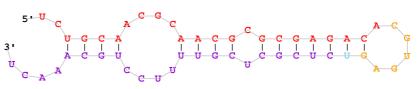
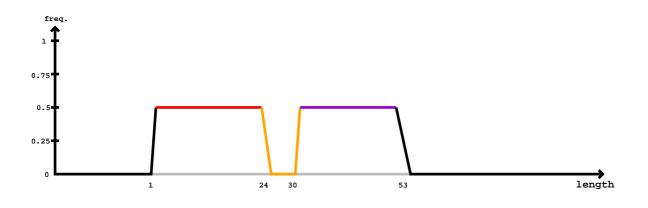
$Provisional\ ID \\ : NW_025322616.1_A cropora_millepora_isolate_JS-1_unplaced_genomic_scaffold_A mil_v2.1_Sc0000028_whole_genome_shotgun_sequenters. \\ [20pt] Provisional\ ID \\ : NW_025322616.1_A cropora_millepora_isolate_JS-1_unplaced_genomic_scaffold_A mil_v2.1_Sc0000028_whole_genome_shotgun_sequenters.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

0.3 Score total Score for star read(s) 3.9 Score for read counts -4.9 1.9 Score for mfe Score for randfold 0 Score for cons. seed -0.6 Total read count 2 1 Mature read count Loop read count 0 Star read count 1

Mature





Star

5'- caacaggcguuagugggagu <mark>cugcaacgcaacgcgagacacg</mark> ugagucucgcucguuuuccugcaaacucucaguaacaccuauugugcgugc	-3'	obs		
caacaggcguuagugggagu <mark>cugcaacgcaacgcgagacacg</mark> ugagucucgcucguuuuccugcaaacucucaguaacaccuauugugcgugc	exp			
(((-(((-((((-((((((((((((((((((((((((((read	s	mm	sample
ucugcaacgcgagacC	1		1	seq
anaganamunifangasasan	1		1	500