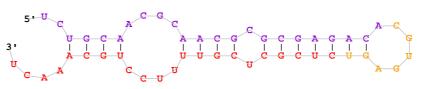
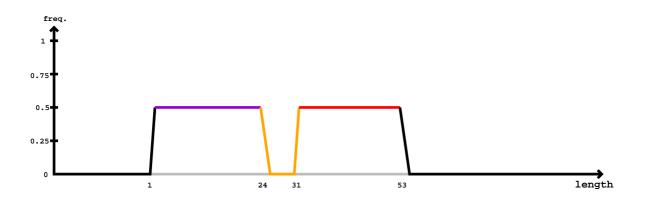
$Provisional\ ID \\ : NW_025322616.1_A cropora_millepora_isolate_JS-1_unplaced_genomic_scaffold_A mil_v2.1_Sc0000028_whole_genome_shotgun_sequenters. \\ [20pt] Provisional\ ID \\ : NW_025322616.1_A cropora_millepora_isolate_JS-1_unplaced_genomic_scaffold_A mil_v2.1_Sc0000028_whole_genome_shotgun_sequenters.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

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

Star





Mature

5'- caacaggcguuagugggagucugcaacgcaacgcgagacac <mark>gugagucucgcucguuuuccugcaaacu</mark> cucaguaacaccuauugugcgugcuaauacaauuuuucaac	-3'	obs		
caacaggcguuagugggagucugcaacgcaacgcgagac <mark>acgugagucucgcucguuuuccugcaaacu</mark> cucaguaacaccuauugugcgugcuaauacaauuuuucaac	exp			
(((.(((.((((.((((.(((((())))))	read	ls	mm	sample
ucugcaacgcaacgcggagacC	1		1	seq
avagavamus (avagava av	1		1	202