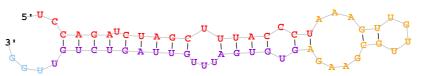
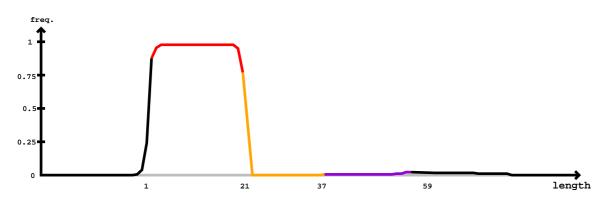
$Provisional\ ID \\ : NW_025322624.1_A cropora_millepora_isolate_JS-1_unplaced_genomic_scaffold_A mil_v2.1_Sc0000070_whole_genome_shotgun_sequenters. \\ [20pt] Provisional\ ID \\ : NW_025322624.1_A cropora_millepora_isolate_JS-1_unplaced_genomic_scaffold_A mil_v2.1_Sc0000070_whole_genome_shotgun_sequenters. \\ [20pt] Provisional\ ID \\ : NW_025322624.1_A cropora_millepora_isolate_JS-1_unplaced_genomic_scaffold_A mil_v2.1_Sc0000070_whole_genome_shotgun_sequenters. \\ [20pt] Provisional\ ID \\ : NW_025322624.1_A cropora_millepora_isolate_JS-1_unplaced_genomic_scaffold_A mil_v2.1_Sc0000070_whole_genome_shotgun_sequenters. \\ [20pt] Provisional\ ID \\ : NW_025322624.1_A cropora_millepora_isolate_JS-1_unplaced_genomic_scaffold_A mil_v2.1_Sc00000070_whole_genome_shotgun_sequenters. \\ [20pt] Provisional\ ID \\ : NW_025322624.1_A cropora_millepora_isolate_JS-1_unplaced_genomic_scaffold_A mil_v2.1_Sc00000070_whole_genome_shotgun_sequenters. \\ [20pt] Provisional\ ID \\ : NW_025322624.1_A cropora_millepora_isolate_JS-1_unplaced_genomic_scaffold_A mil_v2.1_Sc00000070_whole_genome_shotgun_sequenters. \\ [20pt] Provisional\ ID \\ : NW_025322624.1_A cropora_millepora_isolate_JS-1_unplaced_genomic_scaffold_A mill_v2.1_Sc00000070_whole_genome_shotgun_sequenters. \\ [20pt] Provisional\ ID \\ : NW_025322624.1_A cropora_millepora_isolate_JS-1_unplaced_genomic_scaffold_A cropora_millepora_isolate_JS-1_unplaced_genomic_scaffold_A cropora_millepora_isolate_JS-1_unplaced_genomic_scaffold_A cropora_millepora_isolate_JS-1_unplaced_genomic_scaffold_A cropora_millepora_isolate_JS-1_unplaced_genomic_scaffold_A cropora_millepora_isolate_JS-1_unplaced_genomic_scaffold_A cropora_millepora_isolate_JS-1_unplaced_genomic_scaffold_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_A cropora_isolate_A cropora_isolate_A cropora_isolate_A cropora_isolate_A cropora_isolate_A cropora_isolate_A cropora_isolate_A cropora_is$

85.9 Score total Score for star read(s) 3.9 Score for read counts 82.7 Score for mfe -3.6 Score for randfold 0 Score for cons. seed 3 Total read count 174 173 Mature read count Loop read count 0 Star read count 1





| ugauaucuugccucucugacc <mark>uccagaucuagcuuuacccu</mark> aaaguuguugcgaagagugugauuuguuagucuguugg aguauuuuguuaaaaauugauggugauuuau | exp | | |
|---|-------|----|--------|
| ((((((((((((.(((.((((())))).)) | reads | mm | sample |
| Cecuccagaucuagcuuuacc | 1 | 1 | seq |
| ccuccagaucuagcuuuacU | 2 | 1 | seq |
| ccuccagaucuagcuuuacUcu | 3 | 1 | seq |
| Geuecagaucuageuuuaceeu | 1 | 1 | seq |
| .cuccagaucuagcuuuacUcu | 26 | 1 | seq |
| | 4 | 1 | seq |
| cuccagaucuagcuuuacccu | 4 | 0 | seq |
| .cuccagaucuagcuuuacccA | 2 | 1 | seq |
| .uccagaucuagcuuuacc | 2 | 0 | seq |
| .uccagaucuagcuuuaccU | 23 | 1 | seq |
| .uccagaucuagcuuuaccc | 6 | 0 | seq |
| .uceUgaucuagcuuuacccu | 1 | 1 | seq |
| .uccagaucuagcuuuacccu | 68 | 0 | seq |
| .uAcagaucuagcuuuacccu | 1 | 1 | seq |
| .uccagaucuagcuuuacccA | 10 | 1 | seq |
| .uccagaucuagcuuuaccUu. | 1 | 1 | seq |
| .uccagaucuagcuuuacccuU | 1 | 1 | seq |
| ccagaucuagcuuuaccU | 3 | 1 | seq |
| ccagaucuagcuuuacccA | 6 | 1 | seq |
| ccagGucuagcuuuacccu | 1 | 1 | seq |
| ccagaucuagcuuuacccu | 4 | 0 | seq |
| cagaucuagcuuuacccu | 3 | 0 | seq |
| cagaucuagcuuuacceA | 1 | 1 | seq |
| guguUauuuguuagucugu | 1 | 1 | seq |
| cuVuuggaguauuuugu | 1 | 1 | seq |
| guuggaguauuuuguuaaaaaC | 1 | 1 | seq |
| guuggaguauuuCguuaaaaau | 1 | 1 | seq |
| | | | |