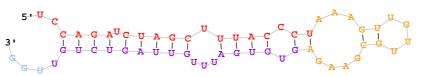
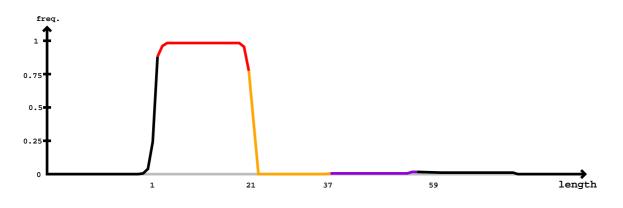
$Provisional\ ID \\ : NW\_025322624.1\_A cropora\_millepora\_isolate\_JS-1\_unplaced\_genomic\_scaffold\_A mil\_v2.1\_Sc0000070\_whole\_genome\_shotgun\_sequenters. \\ [20pt] Provisional\ ID \\ : NW\_025322624.1\_A cropora\_millepora\_isolate\_JS-1\_unplaced\_genomic\_scaffold\_A mil\_v2.1\_Sc0000070\_whole\_genome\_shotgun\_sequenters. \\ [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$ 

82.3 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 -0.6 Total read count 174 173 Mature read count Loop read count 0 Star read count 1





Mature	Star

ugauaucuugccucucugacc <mark>uccagaucuagcuuuacccu</mark> aaaguuguugcgaagagugugauuuguuagucuguuggaguauuuuguuaaaaauugauggugauuuau	exp		
((((((((((((((((((()))))))))	reads	mm	sample
.Cecuccagaucuagcuuuacc	1	1	seq
ccuccagaucuagcuuuaeU	2	1	seq
ccuccagaucuagcuuuacUcu	3	1	seq
Gcuccagaucuagcuuuacccu	1	1	seq
cuccagaucuagcuuuacccA	2	1	seq
Guccagaucuagcuuuacccu	4	1	seq
cuccagaucuagcuuuacccu	4	0	seq
cuccagaucuagcuuuacUcu	26	1	seq
.uccagaucuagcuuuacc	2	0	seq
.uccagaucuagcuuuaccc	6	0	seq
.uccagaucuagcuuuaccU	23	1	seq
uccagaucuagcuuwacccu	68	0	seq
.uAcagaucuagcuuwacccu	1	1	seq
.uccagaucuagcuuuaccUu	1	1	seq
.uccUgaucuagcuuuacccu	1	1	seq
.uccagaucuagcuuuacccA	10	1	seq
.uccagaucuagcuuuacccuU	1	1	seq
ccagaucuagcuuuaccU	3	1	seq
ccagaucuagcuuuacccu	4	0	seq
ccagGucuagcuuuacccu	1	1	seq
ccagaucuagcuuuacccA	6	1	seq
cagaucuagcuuuacccA	1	1	seq
cagaucuagcuuuacccu	3	0	seq
.guguUAuuuguuagucugu	1	1	seq
guuggaguauuuguuaaaaaC	1	1	seq