$Provisional\ ID \\ : NW\_025322633.1\_A cropora\_millepora\_isolate\_JS-1\_unplaced\_genomic\_scaffold\_A mil\_v2.1\_Sc0000098\_whole\_genome\_shotgun\_sequenters. \\ [20pt] Provisional\ ID \\ : NW\_025322633.1\_A cropora\_millepora\_isolate\_JS-1\_unplaced\_genomic\_scaffold\_A mil\_v2.1\_Sc0000098\_whole\_genome\_shotgun\_sequenters. \\ [20pt] Provisional\ ID \\ : NW\_025322633.1\_A cropora\_millepora\_isolate\_JS-1\_unplaced\_genomic\_scaffold\_A mil\_v2.1\_Sc0000098\_whole\_genome\_shotgun\_sequenters. \\ [20pt] Provisional\ ID \\ : NW\_025322633.1\_A cropora\_millepora\_isolate\_JS-1\_unplaced\_genomic\_scaffold\_A mil\_v2.1\_Sc0000098\_whole\_genome\_shotgun\_sequenters. \\ [20pt] Provisional\ ID \\ : NW\_025322633.1\_A cropora\_millepora\_isolate\_JS-1\_unplaced\_genomic\_scaffold\_A mil\_v2.1\_Sc00000098\_whole\_genome\_shotgun\_sequenters. \\ [20pt] Provisional\ ID \\ : NW\_025322633.1\_A cropora\_millepora\_isolate\_JS-1\_unplaced\_genomic\_scaffold\_A mil\_v2.1\_Sc00000098\_whole\_genome\_shotgun\_sequenters. \\ [20pt] Provisional\ ID \\ : NW\_025322633.1\_A cropora\_millepora\_isolate\_JS-1\_unplaced\_genomic\_scaffold\_A mil\_v2.1\_Sc00000098\_whole\_genome\_shotgun\_sequenters. \\ [20pt] Provisional\ ID \\ : NW\_025322633.1\_A cropora\_millepora\_isolate\_JS-1\_unplaced\_genomic\_scaffold\_A mill\_v2.1\_Sc00000098\_whole\_genome\_shotgun\_sequenters. \\ [20pt] Provisional\ ID \\ : NW\_025322633.1\_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$ 

Score total : 0.5

Score for read counts : -1.3
Score for read counts : 0
Score for mfe : 2.4

Star

 Score for mfe
 : 2.4

 Score for randfold
 : 0

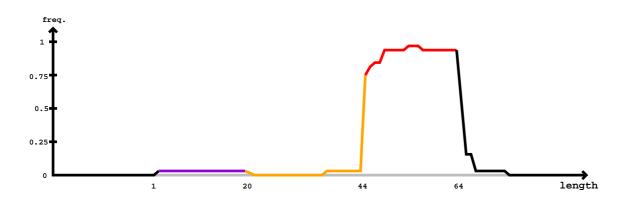
 Score for cons. seed
 : -0.6

 Total read count
 : 30

 Mature read count
 : 29

 Loop read count
 : 0

 Star read count
 : 1



.....aguucgacguuugaaccggGu.....

5'- ugucgaucccaguugauucasacgucgagcuuugcaugugccgaaccuaauuauuagguucgacacauguuaaaguucgacguuugaaccggaucuuaauuauuag	-3'	obs	
ugucgaucccaguugauucaaacgucgagcuuugcaugugccgaaccuaauuauuagguucgacacauguuaaaguucgacguuugaaccggaucuuaauuauuag		exp	
(((((.((((((((((((((((((((((((	reads	mm	sample
.aacgucgagcuuugcaugu	1	0	seq
agguucgGcacauguaaagu	1	1	seq
cacauguaaaguucgacguC	1	1	seq
cacauguaaaguucgacgCu	1	1	seq
	21	0	seq
acauguaaaguucgacguu	1	0	seq
acauguaaaguucgacguAu	1	1	seq
cauguaaaguucgacguuuga	1	0	seq
uguaaaguucgacguuugU	1	1	seq
uguaaaguucgacguuuga	2	0	seq

Mature