$Provisional\ ID \\ : NW_025322877.1_A cropora_millepora_isolate_JS-1_unplaced_genomic_scaffold_A mil_v2.1_xfSc0000226_whole_genome_shotgun_sequences for the provisional ID \\ : NW_025322877.1_A cropora_millepora_isolate_JS-1_unplaced_genomic_scaffold_A mil_v2.1_xfSc0000226_whole_genome_shotgun_sequences for the provisional ID \\ : NW_025322877.1_A cropora_millepora_isolate_JS-1_unplaced_genomic_scaffold_A mil_v2.1_xfSc0000226_whole_genome_shotgun_sequences for the provisional ID \\ : NW_025322877.1_A cropora_millepora_isolate_JS-1_unplaced_genomic_scaffold_A mil_v2.1_xfSc00000226_whole_genome_shotgun_sequences for the provisional ID \\ : NW_025322877.1_A cropora_millepora_isolate_JS-1_unplaced_genomic_scaffold_A mil_v2.1_xfSc00000226_whole_genome_shotgun_sequences for the provisional ID \\ : NW_025322877.1_A cropora_millepora_isolate_JS-1_unplaced_genomic_scaffold_A mil_v2.1_xfSc00000226_whole_genome_shotgun_sequences for the provisional ID \\ : NW_025322877.1_A cropora_millepora_isolate_JS-1_unplaced_genomic_scaffold_A mil_v2.1_xfSc00000226_whole_genome_shotgun_sequences for the provisional ID \\ : NW_025322877.1_A cropora_millepora_isolate_JS-1_unplaced_genomic_scaffold_A mill_v2.1_xfSc00000226_whole_genome_shotgun_sequences for the provisional ID \\ : NW_025322877.1_A cropora_millepora_isolate_JS-1_unplaced_genomic_scaffold_A mill_v2.1_xfSc000000226_whole_genome_shotgun_sequences for the provisional ID \\ : NW_025322877.1_A cropora_millepora_isolate_JS-1_unplaced_genome_sequences for the provisional ID \\ : NW_025322877.1_A cropora_millepora_isolate_JS-1_unplaced_genome_sequences for the provisional ID \\ : NW_025322877.1_A cropora_millepora_isolate_JS-1_unplaced_genome_sequences for the provisional ID \\ : NW_025322877.1_A cropora_isolate_JS-1_unplaced_genome_sequences for the provisional ID \\ : NW_025322877.1_A cropora_isolate_JS-1_unplaced_genome_sequences for the provisional ID \\ : NW_025322877.1_A cropora_isolate_JS-1_unplaced_genome_sequences for the provisional ID \\ : NW_025322877.1_A cropora_isolate_JS-1_unplac$

Score total : 0

Total read count

Star read count

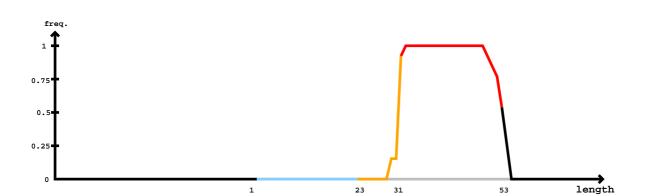
Mature read count Loop read count

Score for star read(s) : -1.3
Score for read counts : 0
Score for mfe : 1.8
Score for randfold : 0
Score for cons. seed : -0.6

13 13

 $0\\0$





Star		Ma	 ~~
o Lar		Ma	

		-3' exp	5'- uugugauaauacucuagaggauuuguccuuuguuugcucauuaaaagagcgcaugcaagcaugaauaugccugcaugca
sample	mm	reads	((((((((((((((((((((((
seq	0	1	ugccugcaugcacucuuuuuaa
seq	1	1	ugccugcaugcacucuuuuuaaC
seq	0	1	ccugcaugcacucuuuuu
seq	0	1	ccugcaugcacucuuuuua
seq	1	1	Ucugcaugcacucuuuuaau
seq	1	1	ccugcaugcacucuuuuulau
seq	1	1	ccugcaugcacucuuuuuaauU
seq	0	1	ccugcaugcacucuuuuuaaua
seq	1	4	ccugcaugcacucuuuuuaauC
sea	0	1	Cugcaugcacuciuiiiiiaaiia