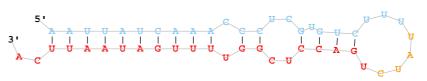
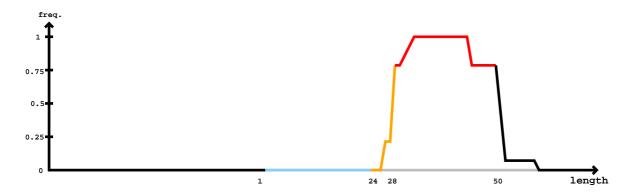
$Provisional\ ID \\ : NW_025322618.1_A cropora_millepora_isolate_JS-1_unplaced_genomic_scaffold_A mil_v2.1_Sc0000038_whole_genome_shotgun_sequenters. \\ [20pt] Provisional\ ID \\ : NW_025322618.1_A cropora_millepora_isolate_JS-1_unplaced_genomic_scaffold_A mil_v2.1_Sc0000038_whole_genome_shotgun_sequenters. \\ [20pt] Provisional\ ID \\ : NW_025322618.1_A cropora_millepora_isolate_JS-1_unplaced_genomic_scaffold_A mil_v2.1_Sc0000038_whole_genome_shotgun_sequenters. \\ [20pt] Provisional\ ID \\ : NW_025322618.1_A cropora_millepora_isolate_JS-1_unplaced_genomic_scaffold_A mil_v2.1_Sc00000038_whole_genome_shotgun_sequenters. \\ [20pt] Provisional\ ID \\ : NW_025322618.1_A cropora_millepora_isolate_JS-1_unplaced_genomic_scaffold_A mil_v2.1_Sc00000038_whole_genome_shotgun_sequenters. \\ [20pt] Provisional\ ID \\ : NW_025322618.1_A cropora_millepora_isolate_JS-1_unplaced_genomic_scaffold_A mil_v2.1_Sc00000038_whole_genome_shotgun_sequenters. \\ [20pt] Provisional\ ID \\ : NW_025322618.1_A cropora_millepora_isolate_JS-1_unplaced_genomic_scaffold_A cropora_isolate_JS-1_unplaced_genomic_scaffold_A cropora_isolate_JS-1_unplaced_genomic_scaffold_A cropora_isolate_JS-1_unplaced_genomic_A cropora_isolate_JS-1_unplaced_genomic_A cropora_isolate_JS-1_unplaced_genomic_A cropora_isolate_JS-1_unplaced_genomic_A cropora_isolate_A cropora_isolate_A c$

3.2 Score total Score for star read(s) -1.3 Score for read counts 0 1.5 Score for mfe Score for randfold 0 Score for cons. seed 3 Total read count 13 13 Mature read count Loop read count 0 0 Star read count





Star	Mature
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5'-	auaauaaacaauuauugaauucgguuuuuguaugauaucaugaauuaucaaacccucgugucuuu <mark>uaucugaccucgguuuugauaauucau</mark> gauaucacuuacucaaccu	-3' exp		
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	ucugaccucgguuuugau	3	0	seq
	ugaccucgguuuugauaauuca	7	0	seq
	.ugaccucgguuuugauaaCucau	1	1	seq
	accucgguuuugauaauucaugauauca	1	0	seq
	ccucgguuuugauaauuca	1	0	seq
	CucqquiiniqanaanicaA	1	1	sea