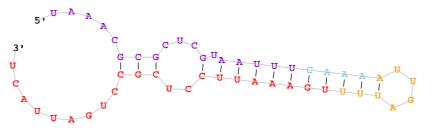
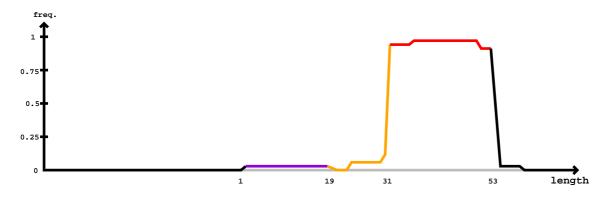
$Provisional\ ID \\ : NW_025322636.1_A cropora_millepora_isolate_JS-1_unplaced_genomic_scaffold_A mil_v2.1_Sc0000107_whole_genome_shotgun_sequential mil_v2.1_Sc0000107_whole_genome_shotgun_sequential mil_v2.1_Sc0000107_whole_genome_sequential mil_v2.1_Sc0000107_whole_genome_sequential mil_v2.1_Sc0000107_whole_genome_sequential mil_v2.1_Sc0000107_whole_genome_sequential mil_v2.1_Sc00000107_whole_genome_sequential mil_v2.1_Sc00000107_whole_g$

13.6 Score total Score for star read(s) 3.9 Score for read counts 9.8 0.5 Score for mfe Score for randfold 0 Score for cons. seed -0.6 Total read count 31 30 Mature read count Loop read count 0 Star read count 1





Star Mature

-	uuaauuaguaauuggaeuueaugueeuaeaauueagggaauaaaegegeueguaauuueaaaauugauuuugaaauueeuegeeugauuaeueeegaauuguaeueea	-3.	DDS	
	uuaauuaguaauuggacuucauguccuacaauucagggaauaaacgcgoucguaauuucaaa <mark>auugauuuugaaauuccucgccugauuacu</mark> ccccgaauuguacucca	exp		
	((()))).(((((.((.((.((.((.((.((reads	mm	sample
	uaaacgGgcucguaauuu	1	1	seq
	auugauuuugaaauuccucgccugauuacA	2	1	seq
	uugaaauuccucgccugauuacu	2	0	seq
	ugaaauuccucgccugauu	2	0	seq
	ugaaauuccucgccugauuacA	3	1	seq
	ugGaauuccucgccugauuacu	1	1	seq
	ugaaauuccucgccugauuacC	3	1	seq
	ugaaauuccucgccugauuacu	18	0	seq
	ugaaauuccucgccugauuacG	1	1	seq
	miganeaccidamia and cacada	1	1	500