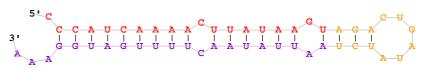
$Provisional\ ID \\ : NW_025322774.1_A cropora_millepora_isolate_JS-1_unplaced_genomic_scaffold_A mil_v2.1_xfSc0000084_whole_genome_shotgun_sequences for the provisional ID \\ : NW_025322774.1_A cropora_millepora_isolate_JS-1_unplaced_genomic_scaffold_A mil_v2.1_xfSc0000084_whole_genome_shotgun_sequences for the provisional ID \\ : NW_025322774.1_A cropora_millepora_isolate_JS-1_unplaced_genomic_scaffold_A mil_v2.1_xfSc0000084_whole_genome_shotgun_sequences for the provisional ID \\ : NW_025322774.1_A cropora_millepora_isolate_JS-1_unplaced_genomic_scaffold_A mil_v2.1_xfSc0000084_whole_genome_shotgun_sequences for the provisional ID \\ : NW_025322774.1_A cropora_millepora_isolate_JS-1_unplaced_genomic_scaffold_A mil_v2.1_xfSc0000084_whole_genome_shotgun_sequences for the provisional ID \\ : NW_025322774.1_A cropora_millepora_isolate_JS-1_unplaced_genomic_scaffold_A mil_v2.1_xfSc00000084_whole_genome_shotgun_sequences for the provisional ID \\ : NW_025322774.1_A cropora_millepora_isolate_JS-1_unplaced_genomic_scaffold_A mil_v2.1_xfSc00000084_whole_genome_shotgun_sequences for the provisional ID \\ : NW_025322774.1_A cropora_millepora_isolate_JS-1_unplaced_genomic_scaffold_A mil_v2.1_xfSc00000084_whole_genome_shotgun_sequences for the provisional ID \\ : NW_025322774.1_A cropora_millepora_isolate_JS-1_unplaced_genomic_scaffold_A millepora_isolate_JS-1_unplaced_genomic_scaffold_A millepora_isolate_JS-1_unplaced_genomic_scaffold_A millepora_isolate_JS-1_unplaced_genomic_scaffold_A millepora_isolate_JS-1_unplaced_genomic_scaffold_A millepora_isolate_JS-1_unplaced_genomic_scaffold_A millepora_isolate_JS-1_unplaced_genomic_scaffold_A millepora_isolate_JS-1_unplaced_genomic_scaffold_A millepora_isolate_JS-1_unplaced_genomic_scaffold_A millepora_isolate_A millepora_isolate$

2.8 Score total Score for star read(s) -1.3 Score for read counts 0 Score for mfe 1.1 Score for randfold 0 Score for cons. seed 3 Total read count 20 Mature read count 16 Loop read count 0

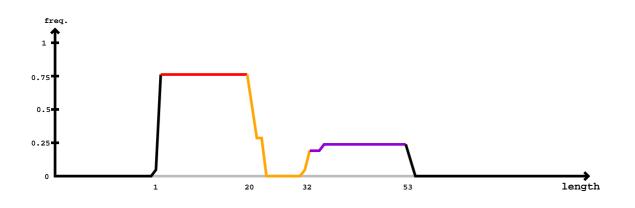
4

Mature

Star read count



seq



Star

.....uauaacuuuugauggaaU.......

5'-	auuguacucauucauuaguucccaucaaaacuuauaaguagacugauaucuaauuauaacuuuugauggaaacaaaugaaug	-3'	obs		
	auuguacucauucauuaguucccaucaaaacuuauaaguagacugauaucuaauuaaegaugaugaugaagaaacaaaugaacuuugaucuugauuuuaguaauacg	exp			
	((((((((((((((((((((((((((((((((((reads	s	mm	sample
	ucccaucaaaacuuauaaguaga	1		0	seq
	cccaucaaaGcuuauaagu	1		1	seq
		7		0	seq
	cccaucaaaacuuauaagC	2		1	seq
		5		0	seq