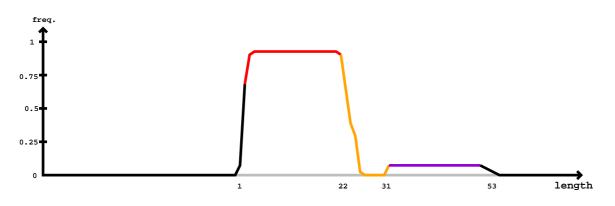
$Provisional\ ID \\ : NW_025322638.1_A cropora_millepora_isolate_JS-1_unplaced_genomic_scaffold_A mil_v2.1_Sc0000115_whole_genome_shotgun_sequenters. \\ [20pt] Provisional\ ID \\ : NW_025322638.1_A cropora_millepora_isolate_JS-1_unplaced_genomic_scaffold_A mil_v2.1_Sc0000115_whole_genome_shotgun_sequenters. \\ [20pt] Provisional\ ID \\ : NW_025322638.1_A cropora_millepora_isolate_JS-1_unplaced_genomic_scaffold_A mil_v2.1_Sc0000115_whole_genome_shotgun_sequenters. \\ [20pt] Provisional\ ID \\ : NW_025322638.1_A cropora_millepora_isolate_JS-1_unplaced_genomic_scaffold_A mil_v2.1_Sc0000115_whole_genome_shotgun_sequenters. \\ [20pt] Provisional\ ID \\ : NW_025322638.1_A cropora_millepora_isolate_JS-1_unplaced_genomic_scaffold_A mil_v2.1_Sc0000115_whole_genome_shotgun_sequenters. \\ [20pt] Provisional\ ID \\ : NW_025322638.1_A cropora_millepora_isolate_JS-1_unplaced_genomic_scaffold_A mil_v2.1_Sc0000115_whole_genome_shotgun_sequenters. \\ [20pt] Provisional\ ID \\ : NW_025322638.1_A cropora_millepora_isolate_JS-1_unplaced_genomic_scaffold_A mil_v2.1_Sc0000115_whole_genome_shotgun_sequenters. \\ [20pt] Provisional\ ID \\ : NW_025322638.1_A cropora_millepora_isolate_JS-1_unplaced_genomic_scaffold_A mil_v2.1_Sc0000115_whole_genome_shotgun_sequenters. \\ [20pt] Provisional\ ID \\ : NW_025322638.1_A cropora_millepora_isolate_JS-1_unplaced_genomic_scaffold_A mill_v2.1_Sc0000115_whole_genome_shotgun_sequenters. \\ [20pt] Provisional\ ID \\ : NW_025322638.1_A cropora_millepora_isolate_JS-1_unplaced_genome_shotgun_sequenters. \\ [20pt] Provisional\ ID \\ : NW_025322638.1_A cropora_millepora_isolate_JS-1_unplaced_genome_shotgun_sequenters. \\ [20pt] Provisional\ ID \\ : NW_025322638.1_A cropora_millepora_isolate_JS-1_unplaced_genome_shotgun_sequenters. \\ [20pt] Provisional\ ID \\ : NW_025322638.1_A cropora_isolate_JS-1_unplaced_genome_shotgun_sequenters. \\ [20pt] Provisional\ ID \\ : NW_025322638.1_A cropora_isolate_JS-1_unplaced_genome_shotgun_sequenters. \\ [20pt] Provisional\ ID \\ : NW_025322638.1_A cropora_isolate_JS-1_unplaced_genome_shotgun_seq$

Score total : 0.6
Score for star read(s) : -1.3
Score for read counts : 0
Score for mfe : -1
Score for randfold : 0

Score for randfold : 0
Score for cons. seed : 3
Total read count : 41
Mature read count : 38
Loop read count : 0
Star read count : 3



5'-	uaauaguguuaugauaauaaguacugagugaaguacaauu <mark>caaggaguaaucgggcgagugauuugaaau</mark> uacgagcaugcuuaccccugaauuaugcgacagcaagugu	-3' ob	s	
	uaauaguguuaugauaauaaguacugagugaaguacaauucaaggaguaaucgggcgagugauuugaaauuacgagcaugcuuaceccugaauuaugcgacagcaagugu	ex	p.	
	((((((((()))).)))).(((.(((((.(((.(((.((.	reads	mm	sample
	ucaaggaguaaucgggcgagugaC	1	1	seq
	ucaaggaguaaucgggcgagugau	1	0	seq
	ucaaggaguaaucgggcgagugauu	1	0	seq
	caaggaguaaucgggcgagu	1	0	seq
	caGggaguaaucgggcgagug	19	1	seq
		1	1	seq
		1	1	seq
		1	0	seq
	caGggaguaaucgggcgagugauu	1	1	seq
		1	0	seq
	aGggaguaaucgggcgagugau	1	1	seq
	aGggaguaaucgggcgagugauu	7	1	seq
	aGggaguaaucgggcgagugauuu	1	1	seq
	Gggaguaaucgggcgagugauu	1	1	seq
		_	_	