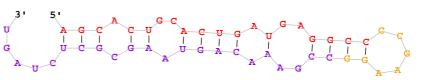
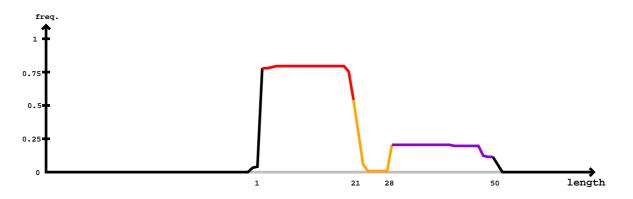
$Provisional\ ID \\ : NW_025322643.1_A cropora_millepora_isolate_JS-1_unplaced_genomic_scaffold_A mil_v2.1_Sc0000126_whole_genome_shotgun_sequenter. \\ [-2.5]$

62.2 Score total Score for star read(s) 3.9 Score for read counts 55.7 -0.3 Score for mfe Score for randfold 0 Score for cons. seed 3 Total read count 121 97 Mature read count Loop read count 0 Star read count 24





Mature	Star

-	aagcucgagguggcuuuuaaagcuucaccuucaucucagcaucagcacugcacugaugaggccccgaaggccgaaacaguaagcgcucuaguaaguucucguuuguuauuu	-3'	obs	
	aagcucgagguggcuuuuaaagcuucaccuucaucucagcauc <mark>agcacugcacu</mark>		exp	
	(((.((((((((())))))))))))((.((read	s mm	sample
	.ucagcacugcacugagg	2	0	seq
	.ucagcacugcacugaggc	1	0	seq
	.ucagcacugcacugagggcc	1	0	seq
	cagcacugcacugagggcc	1	0	seq
	agcacugcacugaugagg	2	0	seq
	agcacugcacugaugagC	1	1	seq
	agcacugcacugaugaggA	2	1	seq
	agcacugcacugaugaggc	23	0	seq
	agcacugcacugaugaggcc	25	0	seq
	agcacugcacugaugaggcG	4	1	seq
	agcacugcacugaugaggccc	8	0	seq
	agcacugcacugaugaggccG	1	1	seq
	agcacugcacugaugaggccA	18	1	seq
	agcacugGacugUugaggcccc	1	1	seq
	agcacugcacugaugaggccAc	3	1	seq
	agcacugcacugaugGggcccc	2	1	seq
	cacugcacugaugaggccA	1	1	seq
	acugcacugaugaggccc	1	0	seq
	ccgaaggccgaaacaguaCg	1	1	seq
	ccgaaacaguaagcgcucu	7	0	seq
	ccgaaacaguaCgcgcucu	1	1	seq
	ccgaaacaguaagcgcucG	1	1	seq
	ccgaaacaguaCgcgcucua	1	1	seq
	ccgaaacaguaagcgcucuagu	10	0	seq
	ccgaaacaguaagcgcucuagA	2	1	seq
	ccgaaacaguaagcgcucuagC	1	1	seq