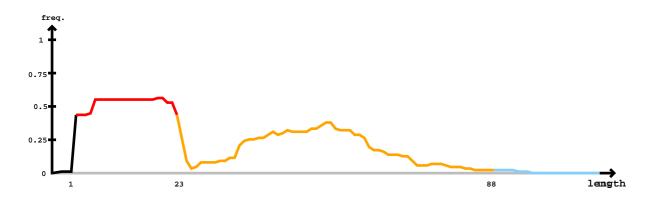
$Provisional\ ID \\ : NW_025322620.1_A cropora_millepora_isolate_JS-1_unplaced_genomic_scaffold_A mil_v2.1_Sc0000055_whole_genome_shotgun_sequenters and the provisional provi$

Score total : 3.3 Score for star read(s) : -1.3 Score for read counts : 0 Score for mfe : 1.6

5^{AAG}AU LUGAAAUGG^CGG^CGACUGGG<mark>GGACGCA</mark> LUGAAAUGG^CGGGACGGCUGGG_A

Score for mile : 1.6
Score for randfold : 0
Score for cons. seed : 3
Total read count : 83
Mature read count : 47
Loop read count : 36
Star read count : 0



Mature Star

	114 5 41 5			
5'	- aggaagauugaaauggcgcggacuggcuuggacgcaucuguguuuguu	-3'	exp	
	$((((\dots,(((((((((((((((((((((((((((((((($	reads	mm	sample
	aggaagauugaaauggcgGg	1	1	seq
	aagauugaaauggcgGgcg	3	1	seq
	aagauugaaauggcgGgcgac	8	1	seq
	aagauugaaauggegGgegacu	26	1	seq
	auugaaauggegGgegacu	1	1	seq
	uugaaauggegGgegacu	3	1	seq
	uugaaauggcgGgcgacugg.	3	1	seq
	uugaaauggegeggacugg	2	0	seq
	uugaaauggcgGgcgacuggc	1	1	seq
	egacuggcuuggacgcaucuguguu	2	0	seq
	uuggacgcaucuguguuu	1	0	seq
	.uuggaUgcaucuguguuuuguucugc	1	1	seq
	uggacgcaucuguguuuuguucugc	3	0	seq
	cgeGucuguguuuuuguucugegucucgu	1	1	seq
	.caucugugCuguuuuguucugcgucu	2	1	seq
	ucuguguuuuguucugcg	1	0	seq
	ucuguguuuuguucugcgucuc	1	0	seq
	ucuguguuguuuuguucugcgucucgC	1	1	seq
	ucuguguuuuguucugcgucucgu	2	0	seq
	ucuguguuuguuuuguucugcUucucgua	1	1	seq
	ucuguguuuuguucugcgucucgua	1	0	seq
	ucuguguuguuuuguucugcgucucgCaggu	1	1	seq
	.cugugCuguuuuguucugcgucucgu	2	1	seq
	.cugugCuguuuuguucugcgucucguaggu	1	1	seq
	uguguuguuuuguucugcgucucguagg	1	0	seq
	uguuguuuuguucugcgucucguagg	1	0	seq
	uguuuuguucugcgucu	1	0	seq
	uuguuuuguucugcUucucguagguguu	1	1	seq
	uguuuuguucugcgucucA	2	1	seq
	uuuuguucugcgucucguagguAuucgc	1	1	seq
	uuuguucugcgucuUguagguguucgc	2	1	seq
	ucugcgucucguagguguucg	2	0	seq
		1	1	seq
	ugcgucuUguagguguucgcggaagcgguc	1	1	seq

Mature Star