$Provisional\ ID \\ \hspace*{0.5cm}: NC\_058067.1\_A cropora\_millepora\_isolate\_JS-1\_chromosome\_2\_Amil\_v2.1\_whole\_genome\_shotgun\_sequence\_93467.1\_A cropora\_isolate\_JS-1\_Chromosome\_3\_Amil\_v2.1\_A cropora\_millepora\_isolate\_JS-1\_Chromosome\_3\_Amil\_v2.1\_A cropora\_millepora\_isolate\_JS-1\_Chromosome\_3\_Amil\_v2.1\_A cropora\_millepora\_isolate\_JS-1\_Chromosome\_3\_Amil\_v3.1\_A cropora\_millepora\_isolate\_JS-1\_Chromosome\_3\_Amil\_v3.1\_A cropora\_millepora\_isolate\_JS-1\_Chromosome\_3\_Amil\_v3.1\_A cropora\_millepora\_isolate\_JS-1\_Chromosome\_3\_Amil\_v3.1\_A cropora\_millepora\_isolate\_3\_A cropora\_isolate\_3\_A cropora\_isolate\_3\_A$ 

Score total : 18.1 Score for star read(s) : 3.9 Score for read counts : 12.4 Score for mfe : 2.4 Score for randfold : 0 Score for cons. seed : -0.6

> 36 32

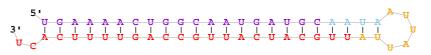
0

4

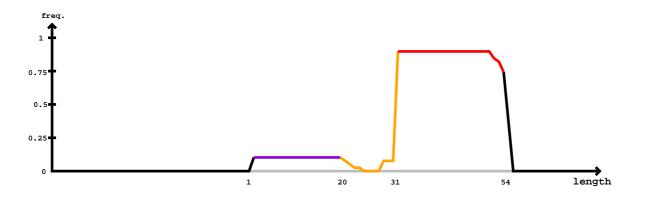
Total read count

Star read count

Mature read count Loop read count



0



Star

5°- ugutageaageaataacaataatugutacagugugacagugaaaacuggeaatgaugeaataatuatuatugeatcatugecagututteacugucacaceateuteaaaac	-3 ODS		
uguuagcaagcaaauaacaauaauuguuacagugugacagugaaaacuggcaaugaugcaaua <mark>auuauugcaucauugccaguuuucacu</mark> gucacaccaucuucaaaac	exp		
(((()))((((()))))((((((((((	reads	mm	sample
ugaaaacuggcaaugaugc	2	0	seq
	1	1	seq
ugaaaacuggcaaugaugcaaua	1	0	seq
.uuauugcaucGuugccaguuuuc	2	1	seq
	1	1	seq

Mature