$Provisional\ ID \\ : NW\_025322636.1\_A cropora\_millepora\_isolate\_JS-1\_unplaced\_genomic\_scaffold\_A mil\_v2.1\_Sc0000107\_whole\_genome\_shotgun\_sequenters and the provisional provi$ 

Score total : 2

Score for star read(s) : -1.3
Score for read counts : 0
Score for mfe : 0.3
Score for randfold : 0
Score for cons. seed : 3
Total read count : 344
Mature read count : 344

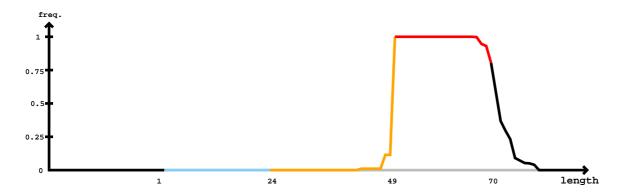
0

0

Loop read count

Star read count





Star	Mature

(((((((.((((((((((((((((((((	reads	mm	sample
Aauguugguaguucagagccugagaau	2	1	seq
Aauguugguaguucagagccugagaaug	1	1	seq
Aauguugguaguucagagccugagaaugg	1	1	seq
.ugguaguucagagccugagaau	2	0	seq
.ugguaguucagagccugagaaug	12	0	seq
.ugguaguucagagacugagaauC	1	1	seq
ugguaguucagagccugagaaCg	1	1	seq
ugguaguucagagacugagaGug	1	1	seq
ugguaguucagagacugagaaugU	1	1	seq
.ugguaguucagagccugagaaugg	1	0	seq
.ugguaguucagagccugagaaugA	2	1	seq
.ugguaguucagagccugagaaugga.	4	0	seq
.ugguaguucagagccugagaauggG.	1	1	seq
.ugguaguucagagaccugagaauggaa	2	0	seq
.ugguaguucagagccugagaauggaac	7	0	seq
.ugguaguucagagacugagaauggaacU	3	1	seq
.ugguaguucagagccugagaauggaacca	1	0	seq
guaguucagagccugag	1	0	seq
guaguucagagccugaga	16	0	seq
guaguucagagccugagU	3	1	seq
guGguucagagccugagaa	1	1	seq
guaguucagagccugagaa	5	0	seq
guaguucagagccGgagaau	1	1	seq
guaguucagagccugagGau	1	1	seq
guaguucagagccugagaaC	8	1	seq
guaguucagagacugagaGu	1	1	seq
guaguucagagccugagaau	32	0	seq
guaguucagagccugagaauU	1	1	seq
guaguucagGgccugagaaug	1	1	seq
guaguucagagccugagaauA	16	1	seq
guaguucagagccugagaauC	2	1	seq
guaguucagagccCgagaaug	1	1	seq
guaguucagagccugagaaug	114	0	seq
guaguucagagccugagaaGg	1	1	seq

Star Mature

uuccuagcacuaaaaaauccuauuuucaagcucuaguacuguccaaacacaguuuguaaaguucauguug <mark>guaguucagagccugagaaug</mark> gaaccaagugugcaaugua			
guaguucagagccugagaaugg	5	0	seq
guaguucagagccugagaaugA	1	1	seq
guaguucagagccugagaaugU	1	1	seq
guaguucagagccugagaauggG	1	1	seq
guaguucagagccugagaauggU	1	1	seq
guaguucagagccugagaaugga	19	0	seq
guagCucagagccugagaaugga	1	1	seq
guaguucagagccugagaauggaa.	19	0	seq
guaguucagagccugagaGuggaa	2	1	seq
guaguucagagUcugagaauggaa.	1	1	seq
guaguucagagccugagaauggaac	37	0	seq
guaguucagagccugagaauggaaU	5	1	seq
	2	1	seq
guGguucagagccugagaauggaac	2	1	seq
guaguucagagccugagaauggaacG.	2	1	seq
guaguucagagccugagaauggaacU	2	1	seq
guaguucagagccugagaauggaacUa	1	1	seq
guaguucagagccugagaauggaacGa	5	1	seq
guaguucagagccugagaauggaacGaa	1	1	seq
guaguucagagccugagaauggaacGaag	4	1	seq
quaquucaqaqccuqaqaauqqaacGaaqu	13	1	seq
guaguacagagccugagaauggaaccGaguguaguacagagccugagaauggaaccGagu	2	1	seq