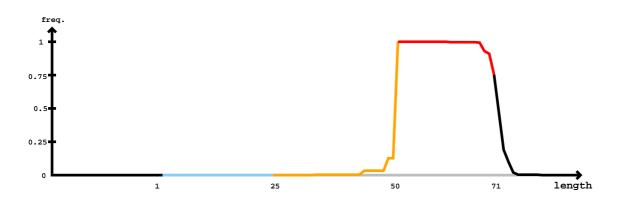|		
Provisional ID	:	NW_025322636.1_Acropora_millepora_isolate_JS-1_unplaced_genomic_scaffold_Amil_v2.1_Sc0000107_whole_genome_shotgun_sequen
Score total	:	2.4
Score for star read(s)	:	-1.3
Score for read counts	:	0
Score for mfe	:	0.7
Score for randfold	:	0
Score for cons. seed	:	3
Total read count	:	289
Mature read count	:	289
Loop read count	:	0
Star read count	:	0

Star   **Mature**

```
5'- ucgaagcacuaaaaaauccuauuuucaagcucucaguacuguccaaacacaguuuguaaagucuauguugguaguucagagccugagaauggaagcaagugugcaaua  -3'    exp
     .....((((.......(((.((((((((.(((((....(((((.((((.((.((.((....)).)).)))))))))))..))))).))))))))).......))))...    reads    mm         sample
     ....................................................uuguaaagucuauguugguaguucagag.................................  1       0          seq
     ...........................................................Aauguugguaguucagagccugagaau.................         2       1          seq
     ...........................................................uauguugguaguucagagccugagaaug................         2       0          seq
     ...........................................................uauguugguaguucGgagccugagaaug................         1       1          seq
     ...........................................................Aauguugguaguucagagccugagaaug................         1       1          seq
     ...........................................................uauguugguaguucagaAccugagaaug................         1       1          seq
     ...........................................................Aauguugguaguucagagccugagaaugg...............         1       1          seq
     ...........................................................uauguugguaguucagagccugagaaugA...............         1       1          seq
     ...............................................................ugguaguucagagccugagaau..................         2       0          seq
     ...............................................................ugguaguucagagccugagaauC.................         1       1          seq
     ...............................................................ugguaguucagagccugagaaCg.................         1       1          seq
     ...............................................................ugguaguucagagccugagaGug.................         1       1          seq
     ...............................................................ugguaguucagagccugagaaug.................         12      0          seq
     ...............................................................ugguaguucagagccugagaaugU................         1       1          seq
     ...............................................................ugguaguucagagccugagaaugA................         2       1          seq
     ...............................................................ugguaguucagagccugagaaugg................         1       0          seq
     ...............................................................ugguaguucagagccugagaauggG...............         1       1          seq
     ...............................................................ugguaguucagagccugagaaugga...............         4       0          seq
     ...............................................................ugguaguucagagccugagaauggaa..............         2       0          seq
     ...................................................................guaguucagagccugagag..................         1       0          seq
     ...................................................................guaguucagagccugagU...................         3       1          seq
     ...................................................................guaguucagagccugaga...................         16      0          seq
     ...................................................................guaguucagagccugagaa..................         5       0          seq
     ...................................................................guGguucagagccugagaa..................         1       1          seq
     ...................................................................guaguucagagccugagaau.................         32      0          seq
     ...................................................................guaguucagagccugagaGu.................         1       1          seq
     ...................................................................guaguucagagccugagGau.................         1       1          seq
     ...................................................................guaguucagagccGgagaau.................         1       1          seq
     ...................................................................guaguucagagccugagaaC.................         8       1          seq
     ...................................................................guaguucagagccCgagaaug................         1       1          seq
     ...................................................................guaguucagagccugagaGg................         1       1          seq
     ...................................................................guaguucagagccugagaaug................         114     0          seq
     ...................................................................guaguucagagccugagaauC................         2       1          seq
     ...................................................................guaguucagGgccugagaaug................         1       1          seq
```

ucgaagcacuaaaaaauccuauuuucaagcucucaguacuguccaaacacaguuuguaaagucuauguugguaguucagagccugagaugggaagcaagugugcaaua

....guaguucagagccugagaauU................	1	1	seq
....guaguucagagccugagaauA................	16	1	seq
....guaguucagagccugagaaugg...............	5	0	seq
....guaguucagagccugagaaugU...............	1	1	seq
....guaguucagagccugagaaugA...............	1	1	seq
....guagCucagagccugagaaugga..............	1	1	seq
....guaguucagagccugagaauggU.............	1	1	seq
....guaguucagagccugagaauggG.............	1	1	seq
....guaguucagagccugagaaugga.............	19	0	seq
....guaguucagagccugagaGuggaa............	2	1	seq
....guaguucagagUcugagaauggaa............	1	1	seq
....guaguucagagccugagaauggaa............	19	0	seq
....guaguucagagccugagaauggaaU...........	5	1	seq
....guaguucagagccugagaauggaagcaagC.......	1	1	seq