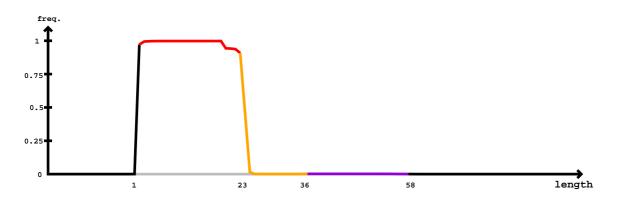|		
Provisional ID	:	NC_058067.1_Acropora_millepora_isolate_JS-1_chromosome_2_Amil_v2.1_whole_genome_shotgun_sequence_107317
Score total	:	4346.5
Score for star read(s)	:	3.9
Score for read counts	:	4337.2
Score for mfe	:	2.4
Score for randfold	:	0
Score for cons. seed	:	3
Total read count	:	8519
Mature read count	:	8506
Loop read count	:	0
Star read count	:	13

```
         5'    A A A A A U G U C G   G U   U G C U U   A   A G C U      C U   U A
                                                                                 A
         3' A  U U U U U G C A G C   A C   G A A   C U C   G A      A U A U   A U
            G                        A C                  A     A U
```

freq.

```
1
0.75
0.5
0.25
0
      1        23   36        58        length
```

Mature   **Star**

```
5'- guuucuguccguggcucaaaaaugucgguugcuuaagcucucuaauaguaagcucaagcacacgacguuuuugagccacgggcggcaaaagcaagugagcugcauuuuu -3'   obs
    guuucuguccguggcucaaaaaugucgguugcuuaagcucucuaauaguaagcucaagcacacgacguuuuugagccacgggcggcaaaagcaagugagcugcauuuuu       exp
    .(((((((((((((((((((((((((..(((((.((((..(.....)..)))).)))))..))))))))))))))))))))))).))).(((.(....)))).......   reads   mm    sample
    guuucuguccguggcuca.................................................................................................   1       0     seq
    ..............cUaaaaugucgguugcuuaagcu.............................................................................   2       1     seq
    ..............aaaaaugucgguugcuuG..................................................................................   1       1     seq
    ..............Uaaaaugucgguugcuua..................................................................................   464     1     seq
    ..............Caaaaugucgguugcuua..................................................................................   1       1     seq
    ..............Uaaaaugucgguugcuuaa.................................................................................   12      1     seq
    ..............Uaaaaugucgguugcuuaag................................................................................   35      1     seq
    ..............Gaaaaugucgguugcuuaagc...............................................................................   2       1     seq
    ..............Uaaaaugucgguugcuuaagc...............................................................................   243     1     seq
    ..............Caaaaugucgguugcuuaagc...............................................................................   2       1     seq
    ..............aaaaaugucgguugcuuaagcA..............................................................................   3       1     seq
    ..............aaaaaugucgguugcuuaagcu..............................................................................   7       0     seq
    ..............Uaaaaugucgguugcuuaagcu..............................................................................   7452    1     seq
    ..............Gaaaaugucgguugcuuaagcu..............................................................................   2       1     seq
    ..............Caaaaugucgguugcuuaagcu..............................................................................   22      1     seq
    ..............Uaaaaugucgguugcuuaagcuc.............................................................................   15      1     seq
    ..............aaaaaugucgguugcuuaagcCc.............................................................................   1       1     seq
    ..............Uaaaaugucgguugcuuaagcucu............................................................................   17      1     seq
    ..............Uaaaaugucgguugcuuaagcucucu..........................................................................   2       1     seq
    ..............Uaaaaugucgguugcuuaagcucucuaau.......................................................................   1       1     seq
    ..............Uaaaaugucgguugcuuaagcucucuaaua......................................................................   2       1     seq
    ...............aaaaugucgguugcuuaa.................................................................................   1       0     seq
    ...............aaaaugucgguugcuuGagc...............................................................................   3       1     seq
    ...............aaaaugucgguugcuuaagc...............................................................................   4       0     seq
    ...............aaaaugucgguugcuuGagcu..............................................................................   15      1     seq
    ...............aaaaugucgguugcuuaagcu..............................................................................   36      0     seq
    ...............Uaaaugucgguugcuuaagcu..............................................................................   3       1     seq
    ...............aaaaugucgguugcuuaagcA..............................................................................   8       1     seq
    ...............aaaaugucgguugcuuaagcC..............................................................................   11      1     seq
    ...............aaaaugucgguugcuuaagcuc.............................................................................   4       0     seq
    ...............aaaaugucgguugcuuaagcuU.............................................................................   4       1     seq
    ...............aaaaugucgguugcuuGagcuc.............................................................................   7       1     seq
    ...............aaaaugucgguugcuuaagcCcu............................................................................   2       1     seq
```

guuucuguccguggcucaaaaaugucgguugcuuaagcucucuaauaguaagcucaagcacacgacguuuugagccacgggcggcaaaagcaagugagcugcauuuuu

Sequence	Count	Mismatch	Type
................aaaaugucgguugcuuaagcuAu..	2	1	seq
................aaaaugucgguugcuuaagcucu..	34	0	seq
................aaaaugucgguugcuuaagcucG..	2	1	seq
................aaaaugucgguugcuuaagcucC..	15	1	seq
................aaaaugucgguugcuuaagcuUu..	1	1	seq
................aaaaugucgguugcuuGagcucu..	25	1	seq
................aaaaugucgguugcuuaagcucA..	13	1	seq
................aaaaugucgguugcuuaagcuAuc...	1	1	seq
................aaaaugucgguugcuuaagcucuA...	1	1	seq
................aaaaugucgguugcuuaagcucuU...	1	1	seq
................aaaugucgguugcuuaagcu...	4	0	seq
................aaaugucgguugcuuaagcC...	4	1	seq
................aaaugucgguugcuuaagcA...	1	1	seq
................aaaugucgguugUuuaagcu..	1	1	seq
................Uaaugucgguugcuuaagcu..	2	1	seq
................aaaugucgguugcuuaagcucu..	3	0	seq
................aaaugucgguugcuuaagcucA..	1	1	seq
................aaaugucgguugcuuaagcucuU...	1	1	seq
................aaugucgguugcuuaagc..	1	0	seq
................aaugucgguugcuuaagcC...	2	1	seq
................aaugucgguugcuuaagcu...	1	0	seq
................aaugucgguugcuuaagcA...	1	1	seq
................aaugucgguugcuuGagcu...	3	1	seq
................aaugucgguugcuuaagcucu...	2	0	seq
.................augucgguugcuuaagcucu...	1	0	seq
.................augucgguugcuuaagcucC...	1	1	seq
..................ugucgguugcuuaagcucucu..	1	0	seq
.................................cuaauaguaagcucaagcacacgacgG..	1	1	seq
.................................cuaauaguaagcucaagcacacgacguuuu.......................................	1	0	seq
...................................aauaguaagcucaagcaca...	1	0	seq
......................................uaagcucaagcacacgacguuC..	1	1	seq
......................................uaagcucaagcacacgacguuu..	1	0	seq
....gcucaagcacacgacguuuC...	1	1	seq
....gcucaagcacacgacguuu..	2	0	seq
....gcucaagcacacgacguuuG...	1	1	seq
....gcucaagcacacgacguuuGg..	2	1	seq
....cucaagcacacgacguuu..	1	0	seq
....cucaagcacacgacguuuGga.....................................	6	1	seq