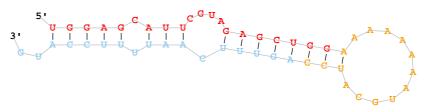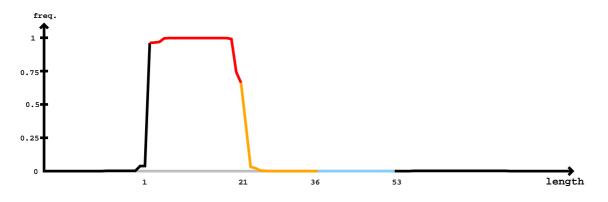| Provisional ID	: NC_058072.1_Acropora_millepora_isolate_JS-1_chromosome_7_Amil_v2.1_whole_genome_shotgun_sequence_300135
Score total	: 1.5
Score for star read(s)	: -1.3
Score for read counts	: 0
Score for mfe	: -0.1
Score for randfold	: 0
Score for cons. seed	: 3
Total read count	: 442
Mature read count	: 442
Loop read count	: 0
Star read count	: 0

Mature **Star**

```
5'- uagucgauuaucauacuauguggagcauucguagagcuggaaaaaaaaugcauccaguuucaauuuuccaugugaugugacacgcgacauugacaaggauuucaucag  -3'   exp
     ..(((((....(((((.(((((((((.(((....(((((((((...........))))))))).))).))))))))).))))))...))))....................   reads   mm      sample
     ..........cauacuauguggagcauucguagagcu...........................................................................   1       0       seq
     ...............ugugAagcauucguagagc.............................................................................   2       1       seq
     ...............uguggagcauucguGgagc.............................................................................   1       1       seq
     ...............uguggagcauucguagagcA............................................................................   1       1       seq
     ...............uguggagcauucguagagcu............................................................................   2       0       seq
     ...............uguggagcauucguagagcC............................................................................   2       1       seq
     ...............ugugAagcauucguagagcug...........................................................................   3       1       seq
     ...............ugugAagcauucguagagcugg..........................................................................   3       1       seq
     ...............uguggagcauucguagagcugg.........................................................................   2       0       seq
     .................uggagcauucguGgagcu...........................................................................   3       1       seq
     .................uggagcauucguagagcC...........................................................................   3       1       seq
     .................uggGgcauucguagagcu...........................................................................   2       1       seq
     .................uggagcauucguagagcA...........................................................................   3       1       seq
     .................ugCagcauucguagagcu...........................................................................   1       1       seq
     .................ugAagcauucguagagcu...........................................................................   85      1       seq
     .................uggagcauucguagagcu...........................................................................   7       0       seq
     .................uggagcauucguagagcuA..........................................................................   1       1       seq
     .................ugAagcauucguagagcug..........................................................................   30      1       seq
     .................uggGgcauucguagagcug..........................................................................   2       1       seq
     .................ugAagcauucguagagcugg.........................................................................   180     1       seq
     .................uggagcauucguagagcugg........................................................................   4       0       seq
     .................uggagcauucguagagcugA........................................................................   2       1       seq
     .................Cggagcauucguagagcugg........................................................................   1       1       seq
     .................uggagcGuucguagagcugg........................................................................   1       1       seq
     .................uggGgcauucguagagcugg........................................................................   3       1       seq
     .................uggagcauucguGgagcugg........................................................................   24      1       seq
     .................uggagcauucguagaAcugga.......................................................................   1       1       seq
     .................uggagcauucguagagcugga.......................................................................   2       0       seq
     .................uggagcauucguagagcuggU.......................................................................   1       1       seq
     .................ugAagcauucguagagcugga.......................................................................   42      1       seq
     .................uggGgcauucguagagcugga.......................................................................   4       1       seq
     .................uggagcauucguGgagcugga.......................................................................   5       1       seq
     .................uggagcauucguagagcuggaaU.....................................................................   1       1       seq
     .................ugAagcauucguagagcuggaaa.....................................................................   1       1       seq
```

uagucgauuaucauacuauguggagcauucguagagcuggaaaaaaaaugcauccaguuucaauuuuccaugugaugugacacgcgacauugacaaggauuucaucag

.....................uggagcauucguagagcuggaaaGa...	1	1	seq
....................gAagcauucguagagcugg...	1	1	seq
....................Aagcauucguagagcugga..	1	1	seq
....................gagcauucguGgagcuggaaa..	1	1	seq
....................agcauucguagagcugga...	2	0	seq
....................agcauucguagGgcuggaa..	2	1	seq
....................agcauucguGgagcuggaa..	2	1	seq
....................agcauucguGgagcuggaaa...	5	1	seq
....................agcauucguagagcuggaaaU..	1	1	seq
.....................gcauucguGgagcugga..	1	1	seq
....ugugacacgcgacauCgaca..............	1	1	seq