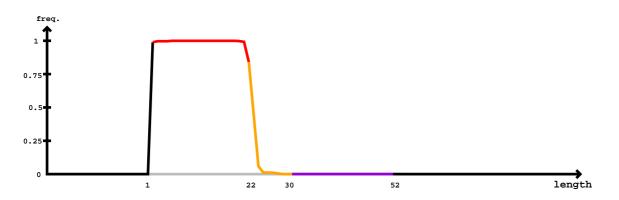| Provisional ID	: NC_058069.1_Acropora_millepora_isolate_JS-1_chromosome_4_Amil_v2.1_whole_genome_shotgun_sequence_158036
Score total	: 1020.8
Score for star read(s)	: 3.9
Score for read counts	: 1016.2
Score for mfe	: 1.3
Score for randfold	: 0
Score for cons. seed	: -0.6
Total read count	: 2005
Mature read count	: 2004
Loop read count	: 0
Star read count	: 1


```
                              Mature                    Star
5'- cugcuuucaauaguuuggauuuaccacauucaggauagacauguucuaucucuagccugaaugaggguaauauuuaagcuguuaaauugcauuuaaauugcauaaaauaug  -3'    obs
    cugcuuucaauaguuuggauuuaccacauucaggauagacauguucuaucucuagccugaaugaggguaauauuuaagcuguuaaauugcauuuaaauugcauaaaauaug           exp
    ........(((((((((((((((((.(((((((((.((((.(((...))).)))).))))))))).))))).)))))))))))....(((((....)))))........     reads    mm     sample
...................uuaccacauucaggauag.......................................................................             4       0      seq
...................uuaccacauucaggauaga......................................................................            10       0      seq
...................Auaccacauucaggauaga......................................................................             1       1      seq
...................uuaccacauucaAgauagac.....................................................................             1       1      seq
...................uuaccacauucGggauagac.....................................................................             2       1      seq
...................uuaccacGuucaggauagac.....................................................................             1       1      seq
...................uuaccacauucaggauagac.....................................................................           254       0      seq
...................uuaccacauucaggauagaG.....................................................................             1       1      seq
...................uuaccacauucaggauagaU.....................................................................            37       1      seq
...................uuaccacauucaggauagaA.....................................................................             3       1      seq
...................uuaccacaCucaggauagac.....................................................................             1       1      seq
...................uuaccacauucaggaCagac.....................................................................             1       1      seq
...................uuaccGcauucaggauagac.....................................................................             2       1      seq
...................Cuaccacauucaggauagac.....................................................................             2       1      seq
...................uuaccacauucaggauagGca....................................................................             5       1      seq
...................uuaccacauucUggauagaca....................................................................             2       1      seq
...................uuaccGcauucaggauagaca....................................................................             4       1      seq
...................uuaccacauucaggauagaGa....................................................................             1       1      seq
...................uuaAcacauucaggauagaca....................................................................             2       1      seq
...................Cuaccacauucaggauagaca....................................................................             3       1      seq
...................uCacUccacauucaggauagaca..................................................................             5       1      seq
...................uuacUacauucaggauagaca....................................................................             3       1      seq
...................uuaccUcauucaggauagaca....................................................................             1       1      seq
...................uuaccacauucaggauagacC....................................................................             1       1      seq
...................uuaccacauucaggauagacU....................................................................            35       1      seq
...................uuaccacauucaggauagaUa....................................................................             2       1      seq
...................uuaccacauucaggaCagaca....................................................................             1       1      seq
...................uuaccacauCcaggauagaca....................................................................             1       1      seq
...................uuaccacauucaggauagaca....................................................................           700       0      seq
...................uuaccacaCucaggauagaca....................................................................             1       1      seq
...................uuaccacauucaggauagacG....................................................................            18       1      seq
...................uuaccacauucaggauCgaca....................................................................             1       1      seq
...................uuaccacGuucaggauagaca....................................................................             2       1      seq
```

Mature   **Star**

cugcuuucaauaguuuggauuuaccacauucaggauagacauguucuaucucuagccugaaugagguaauauuuaagcguuaaauugcauuuaaaugcauaaaauaug

Sequence	Count	Flag	
....................uuaccacauucaggGuagaca...	5	1	seq
....................uuaccacauucaggauagCca...	5	1	seq
....................uuaccacauucaAgauagacau..	1	1	seq
....................uuaccacauucaggauagacaC..	100	1	seq
....................uuaccacauucGggauagacau..	3	1	seq
....................uuaccacauCcaggauagacau..	1	1	seq
....................uuaccacGuucaggauagacau..	1	1	seq
....................uuaccacauucaggauagGcau..	4	1	seq
....................uuaccacauucaggaCagacau..	1	1	seq
....................uuaccacauucaggauagacaG..	10	1	seq
....................uuGccacauucaggauagacau..	2	1	seq
....................Auaccacauucaggauagacau..	3	1	seq
....................uuacUacauucaggauagacau..	1	1	seq
....................uuaccacauucaggauagacau..	616	0	seq
....................uuCccacauucaggauagacau..	1	1	seq
....................uuaccacaCucaggauagacau..	1	1	seq
....................uCaccacauucaggauagacau..	3	1	seq
....................uuaccacauucaggauagacGu..	10	1	seq
....................uuaccacauucaggUuagacau..	1	1	seq
....................uuaccGcauucaggauagacau..	1	1	seq
....................Cuaccacauucaggauagacau..	3	1	seq
....................uuaUcacauucaggauagacau..	2	1	seq
....................uuaccacauucaggauagacauU...	47	1	seq
....................uuaccacauucaggauagacauA...	18	1	seq
....................uuaccacauucaggauagacauC...	6	1	seq
....................uuaccacauucaggauagacaug...	16	0	seq
....................uuaccUcauucaggauagacaug...	2	1	seq
....................uuaccacauucaggauagacauguu...	2	0	seq
....................uuaccacauucaggauagacauguuU..	1	1	seq
....................uuaccacauucaggauagacauguuc..	6	0	seq
....................uuaccacauucaggauagacaAguuc..	2	1	seq
....................uuaccacauucaggauagacaAguucu...	2	1	seq
....................uuaccacauucaggauagacauguucA...	6	1	seq
....................uuaccacauucaggauagacauguucG...	1	1	seq
....................uuaccacauucaggauagacauguucu...	2	0	seq
....................uuaccacauucaggauagacauguucC...	1	1	seq
.....................uaccacauucaggauagac...	1	0	seq
.....................uaccacauucaggauagaca..	4	0	seq
.....................uaccacauucaggauagacaC...	2	1	seq
.....................uaccacauucaggauagacau...	2	0	seq
.....................uaccacauucaggauagacauU..	4	1	seq
.....................uaccacauucaggauagacaug..	1	0	seq
.....................uaccacauucaggauagacauguu..	1	0	seq
.........................cacauucaggauagacaC..	3	1	seq
.........................cacauucaggauagacau..	2	0	seq
.........................cacauucaggauagacauA...	1	1	seq
....Aucuagccugaaugagguaaua......................................	1	1	seq