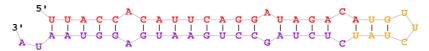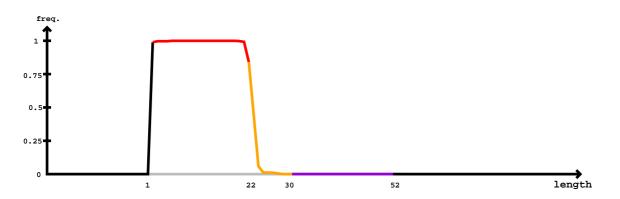| Provisional ID	: NC_058069.1_Acropora_millepora_isolate_JS-1_chromosome_4_Amil_v2.1_whole_genome_shotgun_sequence_158036
Score total	: 1020.8
Score for star read(s)	: 3.9
Score for read counts	: 1016.2
Score for mfe	: 1.3
Score for randfold	: 0
Score for cons. seed	: -0.6
Total read count	: 2005
Mature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

Mature   **Star**

```
5'- cugcuuucaauaguuuggauuuaccacauucaggauagacauguucuaucucuagccugaaugagguaauauuuaagcuguuaaauugcauuuaaaugcauaaaauaug  -3'   obs
    cugcuuucaauaguuuggauuuaccacauucaggauagacauguucuaucucuagccugaaugagguaauauuuaagcuguuaaauugcauuuaaaugcauaaaauaug        exp
    ........(((((((((((((((((((.(((((((((.((((.(((...))).)))).))))))))).))))).)))))))))))))....(((((....)))))........   reads   mm   sample
```

sequence	reads	mm	sample
....................uuaccacauucaggauag..	4	0	seq
...................Auaccacauucaggauaga...	1	1	seq
...................uuaccacauucaggauaga...	10	0	seq
...................uuaccacauucaAgauagac..	1	1	seq
...................uuaccacauucaggauagaA...	3	1	seq
...................uuaccacauucaggauagaU...	37	1	seq
...................uuaccacGuucaggauagac..	1	1	seq
...................uuaccacauucGggauagac..	2	1	seq
...................uuaccGcauucaggauagac..	2	1	seq
...................uuaccacauucaggaCagac..	1	1	seq
...................uuaccacaCucaggauagac..	1	1	seq
...................Cuaccacauucaggauagac..	2	1	seq
...................uuaccacauucaggauagac..	254	0	seq
...................uuaccacauucaggauagaG...	1	1	seq
...................uuaccacauCcaggauagaca..	1	1	seq
...................uuaccacauucaggauagacU..	35	1	seq
...................uuaccacauucaggGuagaca..	5	1	seq
...................uuaccacauucaggauagCca..	5	1	seq
...................uuaccacauucaggauagacC..	1	1	seq
...................uuaccacauucaggaCagaca..	1	1	seq
...................uCaccacauucaggauagaca..	5	1	seq
...................uuaccUcauucaggauagaca..	1	1	seq
...................Cuaccacauucaggauagaca..	3	1	seq
...................uuaccacauucaggauagaUa..	2	1	seq
...................uuaccacaucUggauagaca..	2	1	seq
...................uuaccacGuucaggauagaca..	2	1	seq
...................uuaccacauucaggauagaca..	700	0	seq
...................uuaccacaCucaggauagaca..	1	1	seq
...................uuaAcacauucaggauagaca..	2	1	seq
...................uuaccGcauucaggauagaca..	4	1	seq
...................uuacUacauucaggauagaca..	3	1	seq
...................uuaccacauucaggauagGca..	5	1	seq
...................uuaccacauucaggauagaGa..	1	1	seq

```
cugcuuucaauaguuuggauuuaccacauucaggauagacauguucuaucucuagccugaaugagguaauauuuaagcguuaaauugcauuuaaaugcauaaaauaug
```

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