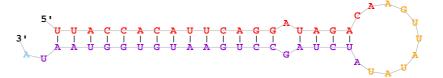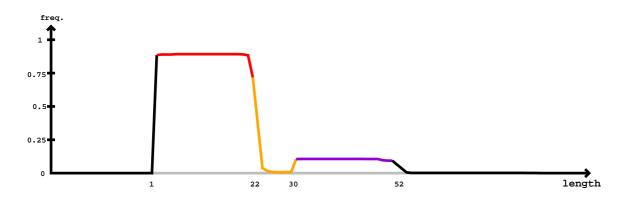| Provisional ID	: NC_058069.1_Acropora_millepora_isolate_JS-1_chromosome_4_Amil_v2.1_whole_genome_shotgun_sequence_158038
Score total	: 918.9
Score for star read(s)	: 3.9
Score for read counts	: 914.2
Score for mfe	: 1.3
Score for randfold	: 0
Score for cons. seed	: -0.6
Total read count	: 1805
Mature read count	: 1626
Loop read count	: 0
Star read count	: 179

Mature   **Star**

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

gaucuuucagcaguuuuaauuuaccacauucaggauagacaaguuauauaucuagccugaaugugguaauauuuaagcguuccaugauuuucaauguuauauuucauuu

Sequence	count	mm	
....................uuaccacGuucaggauagaca..	2	1	seq
....................uuaccacauucaggGuagaca..	5	1	seq
....................uuaccacauucaggauagacGa...	5	1	seq
....................uuaccacauucaggauagacaG...	10	1	seq
....................uuaccacauucaggaCagacaa...	1	1	seq
....................uuaccacauucaggauagacCa...	1	1	seq
....................uuaccacGuucaggauagacaa...	1	1	seq
....................uuaccacauucaAgauagacaa...	1	1	seq
....................uCaccacauucaggauagacaa...	1	1	seq
....................uuaccacauucaggauagacaC...	100	1	seq
....................uuacUacauucaggauagacaa...	1	1	seq
....................uuaccacauucaggauagacaa...	315	0	seq
....................Auaccacauucaggauagacaa...	1	1	seq
....................uuaccacauucaggauagacaaA..	3	1	seq
....................uuaccacauucaggauagacaaU..	30	1	seq
....................uuaccacauucaggauagacaaC..	4	1	seq
....................uuaccacauucaggauagacaag..	2	0	seq
....................uuaccacauucaggauagacaagC...	4	1	seq
....................uuaccacauucaggauagacaagu...	5	0	seq
....................uuaccacauucaggauagacaaUu...	4	1	seq
....................uuaccacauucaggauagacaaguC..	2	1	seq
....................uuaccacauucaggauagacaaguu..	2	0	seq
....................uuaccacauucaggauagacaaguuC...	2	1	seq
....................uuaccacauucaggauagacaaguuCu..	2	1	seq
.....................uaccacauucaggauagac...	1	0	seq
.....................uaccacauucaggauagaca..	4	0	seq
.....................uaccacauucaggauagacaC...	2	1	seq
.....................uaccacauucaggauagacaaU..	1	1	seq
........................cacauucaggauagacaC...	3	1	seq
........................cacauucaggauagacaa...	2	0	seq
...................................aaguuCuauaucuagccugaauguggu......................................	2	1	seq
....................................aguuCuauaucuagccugaaugu...	1	1	seq
.....................................guuCuauaucuagccugaauguggu......................................	2	1	seq
.....................................guuCuauaucuagccugaaugugguaauau.................................	4	1	seq
....uauaucuagccugaauguggu.....................................	1	0	seq
....auaucuagccugaaugugguaau.................................	2	0	seq
....auaucuagccugaaugCgguaau.................................	3	1	seq
....aucuagccugaaugugUu...................................	1	1	seq
....aucuagccugaauguggA...................................	1	1	seq
....aucuagccugaauguggC...................................	1	1	seq
....aucuagccugaauguggu...................................	8	0	seq
....aucuagccugaauguggua..................................	5	0	seq
....aucuagccugaauguggGa..................................	1	1	seq
....aucuagccugaaugugguaaC................................	6	1	seq
....aucCagccugaaugugguaau................................	2	1	seq
....aCcuagccugaaugugguaau................................	1	1	seq
....aucuagccugaaugugguaaG................................	1	1	seq
....aucuagccugaaugugguaau................................	74	0	seq
....aucuagccugaaugugguaaCa...............................	1	1	seq
....aCcuagccugaaugugguaaua...............................	1	1	seq
....aucuagccugaaugugguaaua...............................	17	0	seq
....aucuagccugaaugAgguaaua...............................	1	1	seq
....aucuagccugaaugugguaauU...............................	1	1	seq
....aucuagccugaaugugguaauaC..............................	7	1	seq
....aucuagccugaaugugguaauaG..............................	1	1	seq
....aucuagccugaaugugguaauau..............................	40	0	seq
....aucuagccugaaugugguaauauu.............................	5	0	seq
....aucuagUcugaaugugguaauauu.............................	1	1	seq
....aucuagccugaaugugguaauauC.............................	1	1	seq
....ucuagccugaaugugguaau.................................	1	0	seq
....ucuagccugaaugugguaauau...............................	1	0	seq
....uuaagcguuccaugauuuucGaug.....................	1	1	seq
....uuaagcguuccaugauuuucGauguua..................	2	1	seq