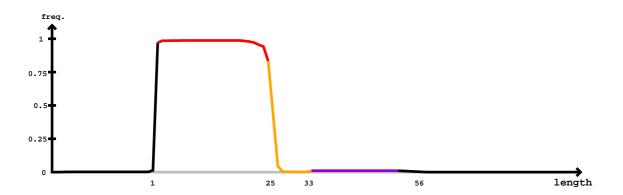| Provisional ID	: NC_058067.1_Acropora_millepora_isolate_JS-1_chromosome_2_Amil_v2.1_whole_genome_shotgun_sequence_84764
Score total	: 2769.9
Score for star read(s)	: 3.9
Score for read counts	: 2761.8
Score for mfe	: 1.1
Score for randfold	: 0
Score for cons. seed	: 3
Total read count	: 5429
Mature read count	: 5380
Loop read count	: 0
Star read count	: 49


```
                                                                                                                                          obs
5'- cagagauuuugcauugcauuuuucaaauuaggaagggagauguuuuacugaacaccgcucugcuaauuugaagagauacaguguaagaauuugcacuggaauaacagc  -3'
    cagagauuuugcauugcauuuuucaaauuaggaagggagauguuuuacugaacaccgcucugcuaauuugaagagauacaguguaagaauuugcacuggaauaacagc       exp
    (((((..(((((((((((..(((((((((((((..((((.(.(((((....))))).).))))·)))))))))))))...)))))))))).))))..(((......))).   reads    mm     sample
    cagagauuuugcauugcauu.................................................................................................   2       0      seq
    .agagauuuugcauugcaCu.................................................................................................   2       1      seq
    .agagauuuugcauugcauu.................................................................................................   6       0      seq
    ...................uuuucaaauuaggaagggagCu............................................................................   1       1      seq
    ...................uuuucaaauuaggaagggagGu............................................................................   3       1      seq
    ...................uuuucaaauuaggaagggagGug...........................................................................   13      1      seq
    ...................uuuucaaauuaggaagggagGugu..........................................................................   24      1      seq
    ...................uuuucaaauuaggaagggagGuguu.........................................................................   12      1      seq
    ...................uuuucaaauuaggaagggagGuguuu........................................................................   10      1      seq
    ...................uuuucaaauuaggaagggagGuguuuu.......................................................................   1       1      seq
    .......................uCucaaauuaggaaggga...........................................................................   1       1      seq
    .......................uuuucaaauuaggaaggga..........................................................................   13      0      seq
    .......................uuuucaaauuaggaagggG..........................................................................   3       1      seq
    .......................Cuucaaauuaggaaggga..........................................................................   1       1      seq
    .......................uuuucaaGuaggaaggga..........................................................................   1       1      seq
    .......................uuuucaaauuaggaagggag.........................................................................   21      0      seq
    .......................uuuucaaauuaggaagggaA.........................................................................   4       1      seq
    .......................uuuucaaauuaggaagggagG........................................................................   29      1      seq
    .......................uuuucaaauuaggaagggaga........................................................................   8       0      seq
    .......................uuuucaaauuaggaagggagU........................................................................   1       1      seq
    .......................uuuucaaauuaggaagggagGu.......................................................................   84      1      seq
    .......................uuuucaaauuaggaagggagCu.......................................................................   1       1      seq
    .......................uuuucaaauuaggaagggagUu.......................................................................   1       1      seq
    .......................uuuucaaauuaggaagggagGug......................................................................   59      1      seq
    .......................uuuucaaauuaggaagggagGugu.....................................................................   573     1      seq
    .......................uuuucaaauuaggaagggagaCgu.....................................................................   1       1      seq
    .......................uuuucaaauuaggaagggagauguC....................................................................   1       1      seq
    .......................uuuucaaauuaggaagggagauguu....................................................................   2       0      seq
    .......................uuuucaaauuaggaagggagGuguu....................................................................   3004    1      seq
    .......................uuuucaaauuaggaagggagauguuC...................................................................   1       1      seq
    .......................uuuucaaauuaggaagggagGuguuu...................................................................   1199    1      seq
    .......................uuuuCaaauuaggaagggagauguuu..................................................................   1       0      seq
    .......................uuuucaaauuaggaagggagauguuuA.................................................................   1       1      seq
```

cagagauuuugcauugcauuuuucaaauuaggaagggagauguuuuacugaacaccgcucugcuaauuugaagagauacaguguaagaauuugcacuggaauaacagc

Sequence	Count	Flag	
....................uuucaaauuaggaagggagGuguuuu..	215	1	seq
....................uuucaaauuaggaagggagGuguuuua...	8	1	seq
....................uuucaaauuaggaagggagGuguuuuacug...	2	1	seq
......................uucaaauuaggaagggagGu...	1	1	seq
......................uucaaauuaggaagggagGug..	3	1	seq
......................uucaaauuaggaagggagGugu...	3	1	seq
......................uucaaauuaggaagggagGuguu..	40	1	seq
......................uucaaauuaggaagggagGuguuu...	20	1	seq
......................uucaaauuaggaagggagGuguuuu..	5	1	seq
........................caaauuaggaagggagGuguuu...	3	1	seq
........................caaauuaggaagggagGuguuu...	2	1	seq
..........................aaauuaggaagggagGuguuu...	1	1	seq
...........................aauuaggaagggagGuguuu...	4	1	seq
............................auuaggaagggagGuguuu...	1	1	seq
....uuuacugaacaccgcucugcuaauuu..................................	1	0	seq
....uuuacugaacaccgcucugcuaauuugaag..............................	1	0	seq
....uuuacugaacaccgcucugcuaauuugaaga.............................	2	0	seq
....uuacugaacaccgcucugcuaauuuga................................	1	0	seq
....uuacugaacaccgcucugcuaauuugaag..............................	2	0	seq
....uuacugaacaccgcucugcuaauuugaagag............................	3	0	seq
....uuacugaacaccgcucugcuaauuugaagaA............................	1	1	seq
....caccgcucugcuaauuuga..........................	16	0	seq
....caccgcucugcuaauuugaa.........................	5	0	seq
....caccgcucugcuaauuugaaA........................	1	1	seq
....caccgcucugcuaauuugaaU........................	1	1	seq
....caccgcucugcuaauuugaag........................	8	0	seq
....caccgcucugcuaauuugaaga.......................	9	0	seq
....caccgcucugcuaauuugaagU.......................	1	1	seq
....caccgcucugUuaauuugaagag......................	1	1	seq
....caccgcucugcuaauuugaagaC......................	1	1	seq
....caccgcucugcuaauuugaagaga.....................	4	0	seq
....accgcucugcuaauuuga..........................	1	0	seq
....Ucgcucugcuaauuugaag......................	1	1	seq