Provisional ID   : NC_058069.1_Acropora_millepora_isolate_JS-1_chromosome_4_Amil_v2.1_whole_genome_shotgun_sequence_171661
Score total   : 0.7
Score for star read(s)   : -1.3
Score for read counts   : 0
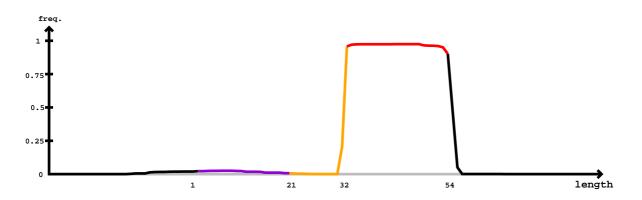Score for mfe   : 2.6
Score for randfold   : 0
Score for cons. seed   : -0.6
Total read count   : 1310
Mature read count   : 1302
Loop read count   : 0
Star read count   : 8

```
5' A G A U C U C A C U C U G U C A C U G G A A A U        C U
3' U C U A G A G U G A G A C A G U G A C C U U U A        C A
                                                    C   A
                                                      C
```



```
                                Star                    Mature

5'- acuugguccugaauccuaaucucguacccagaucucacucugucacuggaaaucucacauuuccagugacagagugagaucuguguacgagauuaccugaauccuaaccuuuu -3'   obs
    acuugguccugaauccuaaucucguacccagaucucacucugucacuggaaaucucacauuuccagugacagagugagaucuguguacgagauuaccugaauccuaaccuuuu       exp
    ....(((...((.((.((((((((((((.(((((((((((((((((((((((.....))))))))))))))))))))))))).))))))))...)).))...)))....   reads   mm   sample
..............cuaaucucguacccagaucucacucugu....................................................................   3    0   seq
..............uaaucucguacccagaucucacucugu.....................................................................   2    0   seq
..............uaaucucguacccagaucucacucugucacu.................................................................   1    0   seq
..............uaaucucguacccGgaucucacucugucacu.................................................................   1    1   seq
.................ucucguacccagaucuca............................................................................   2    0   seq
.................ucucguacccagaucucac...........................................................................   1    0   seq
.................ucucguacccagaucucacu..........................................................................   4    0   seq
.................ucucguacccagaucucacC..........................................................................   3    1   seq
.................ucucguacccagaucucacucugucacuA................................................................   1    1   seq
..................cucguacccagaucucacucugu.....................................................................   3    0   seq
...................ucguacccagaCcucacucug......................................................................   1    1   seq
....................guacccagaucucacucugCcacu..................................................................   2    1   seq
.....................uacccagaucucacucugucacu..................................................................   1    0   seq
.......................cccagaucucacucugucacugU................................................................   1    1   seq
........................agaucucacucugucacugg..................................................................   3    0   seq
.........................agGucucacucugucacuggaaa..............................................................   1    1   seq
...........................aucucacucugucacuggaaa..............................................................   1    0   seq
...........................aucucacucugucacuggaaau.............................................................   1    0   seq
.............................ucucacucugucacugga...............................................................   1    0   seq
..............................ucacucugucacuggaaau.............................................................   1    0   seq
.....................................................Cuuccagugacagagug........................................   1    1   seq
.....................................................uuuccagugacagagug........................................   11   0   seq
.....................................................uuuccagugacagagugU.......................................   2    1   seq
.....................................................uuuccagugacagagugaA......................................   1    1   seq
.....................................................uuuccagugacagagugaga.....................................   1    0   seq
.....................................................uuuccagugacagagugagU.....................................   1    1   seq
.....................................................uuuccagugacagagugagaA....................................   1    1   seq
.....................................................uuuccagugacagagugagUu....................................   2    1   seq
.....................................................uuuccagugacagagugagau....................................   3    0   seq
.....................................................uuuccagugacagagugagaG....................................   2    1   seq
.....................................................uuuccagugacagagugagauG...................................   1    1   seq
.....................................................uuuccagugacagagugagauA...................................   4    1   seq
.....................................................uuuccagugacagagugagauU...................................   44   1   seq
```

acuugguccugaauccuaaucucguaccc agaucucacucugucacugg aaaucucacau uuccagugacagagugagaucu guguacgagauuaccugaauccuaaccuuuu

Sequence	Count	Mismatch	
....uuccagugacagagugaAauc................................	2	1	seq
....uuccagugacagagugagaAc................................	1	1	seq
....uuccagugacagagugagaCcu...............................	2	1	seq
....uuccagugacagagugagaucC...............................	14	1	seq
....uuccagugacagagugagaucA...............................	19	1	seq
....uuccagugacagagugagGucu...............................	2	1	seq
....uuccagugacagagugUgaucu..............................	1	1	seq
....uuccagugacagagugagauUu...............................	2	1	seq
....uuccagugacagagugagaucu................................	151	0	seq
....uuccagugacagagugGgaucu...............................	2	1	seq
....uuccagugacagagugagaucuU..............................	1	1	seq
....uuccagugacagagugagaucuAu.............................	4	1	seq
....uuccagugacagagugG.....................................	1	1	seq
....uuccagugacagagugaga...................................	1	0	seq
....uuccagugacagagugagUu..................................	1	1	seq
....uuccagugacagagugagau..................................	4	0	seq
....uuccagugacagagugagauA.................................	2	1	seq
....uuccagugacagagugGgauc.................................	2	1	seq
....uuccagugacagagugagauU.................................	9	1	seq
....uuccagugacagagugagaCcu................................	4	1	seq
....uuccUgugacagagugagaucu................................	1	1	seq
....uuccagugacagagugGgaucu................................	5	1	seq
....uuccagugacagagugagaucA................................	109	1	seq
....uuccagugacagagugagGucu................................	2	1	seq
....uuccagugacagagCgagaucu................................	2	1	seq
....uuccagugacagagugagaucu................................	643	0	seq
....uuccagugacagagugagauUu...............................	6	1	seq
....uuccagugacagagugagaucC................................	144	1	seq
....uuccagugacagagugagaGcu................................	1	1	seq
....uuccagugacUgagugagaucu................................	1	1	seq
....uuccagugacagagugagaucuC...............................	1	1	seq
....uuccagugacagagugagaucuA...............................	4	1	seq
....uuccagugacagagugagaucAg..............................	1	1	seq
....uuccagugacagagugagaucuU..............................	4	1	seq
....uuccagugacagagugagaucuUu.............................	42	1	seq
....uuccagugacagagugagaucuAu.............................	1	1	seq
....uuccagugacagagugagaCcugu.............................	1	1	seq
....uuccagugacagagugagaucugu.............................	13	0	seq
....uccagugacagagugagaucu................................	11	0	seq
....uccagugacagagugagaucC...............................	1	1	seq
....uccagugacagagugagaucuUu.............................	6	1	seq
....ccagugacagagugagaucu...............................	3	0	seq
....cagugacagagugagaucC..............................	1	1	seq
....agagugagaucuguguacgagau...................	1	0	seq