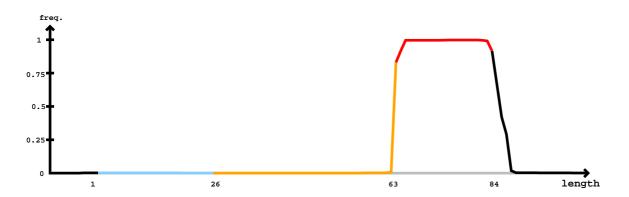| Provisional ID	: NC_058072.1_Acropora_millepora_isolate_JS-1_chromosome_7_Amil_v2.1_whole_genome_shotgun_sequence_278807
Score total	: 1.4
Score for star read(s)	: -1.3
Score for read counts	: 0
Score for mfe	: -0.2
Score for randfold	: 0
Score for cons. seed	: 3
Total read count	: 1000
Mature read count	: 1000
Loop read count	: 0
Star read count	: 0


```
                 Star                                        Mature
5'- uccacuaaaaggaucacuucugguuguucuuggcgcugaucgucugaguucgaauucaaaugaucagcauaaaacugccgauuguccaugaauauuuuuccaucucuagc  -3'   exp
    ..........((((....((.(((.(((.(((.(((((((((((.(((((....))))).)))))))))).))))))).)))))..)))).....................   reads   mm    sample
    ....uaaaGggaucacuucugguug..........................................................................................     2     1      seq
    ..............................................................ucagcauaaaacugccgauuguccaugaau..................     1     0      seq
    ...............................................................cagcauaaaacugccgauuguccaugaau..................     1     0      seq
    ..............................................................uaaaacugccgauuAuccaug.........................     1     1      seq
    ..............................................................uaaaacugccgauuCuccaug.........................     1     1      seq
    ..............................................................uaaaacugccgauuCuccauga........................     2     1      seq
    ..............................................................uaaaacugccgauuguccauga........................     1     0      seq
    ..............................................................uaaaacugccgauuguccaugaa.......................     1     0      seq
    ..............................................................uaaaacugccgauuUuccaugaaua.....................     1     1      seq
    ..............................................................aaaaacugccgauuguccU...........................     2     1      seq
    ..............................................................aaaacugccgauuCuccau..........................     5     1      seq
    ..............................................................aaaacugccgauuCuccaug.........................    55     1      seq
    ..............................................................aaaacugccgauuAuccaug.........................     5     1      seq
    ..............................................................aaaacugccgauuguccaug.........................     6     0      seq
    ..............................................................aaaacugccgauuguccaGga........................     1     1      seq
    ..............................................................aaaacugccgauuCuccauga........................   241     1      seq
    ..............................................................aaaacugccgauuguccauga........................    23     0      seq
    ..............................................................aaaacugccgauuguccGuga........................     1     1      seq
    ..............................................................aaaGcugccgauuguccauga........................     1     1      seq
    ..............................................................aaaacugccgauuAuccauga........................    19     1      seq
    ..............................................................aaaacugccgauuguccaugaa.......................     5     0      seq
    ..............................................................aaaacugccgauuAuccaugaa.......................     5     1      seq
    ..............................................................aaaacugccgauuCuccaugaa.......................   132     1      seq
    ..............................................................aaGacugccgauuguccaugaa.......................     1     1      seq
    ..............................................................aaaacugccgauuguccaugaau......................     3     0      seq
    ..............................................................aaaacugccgauuCuccaugaau......................    84     1      seq
    ..............................................................aaaacugccgauuguccaugaaC......................     1     1      seq
    ..............................................................aaaacugccgauuAuccaugaau......................    10     1      seq
    ..............................................................aaaacugccgauuguccaugaaA......................     1     1      seq
    ..............................................................aaaacugccgauuguccaugaUu......................     1     1      seq
    ..............................................................aaaacugccgauuUuccaugaaua.....................   217     1      seq
    ..............................................................aaaacugccgauuUuccaugaauau....................     6     1      seq
    ..............................................................aaaacugccgauuguccaugaauUu....................     1     1      seq
    ..............................................................aaaacugccgauuguccaugaauauuCuuc................     1     1      seq
```

uccacuaaaaggaucacuucugguuguucuuggcgcugaucgucugaguucgaauucaaaugaucagcauaaaacugccgauuguccaugaauauuuuuccaucucuagc

Sequence	Count	Flag	
....aaacugccgauuCuccaug...................	4	1	seq
....aaacugccgauuCuccauga..................	13	1	seq
....aaacugccgauuCuccaugaa.................	11	1	seq
....aaacugccgauuguccaugaau................	1	0	seq
....aaacugccgauuAuccaugaau................	1	1	seq
....aaacugccgauuCuccaugaau................	7	1	seq
....aaacugccgauuUuccaugaaua...............	45	1	seq
....aaacugccgauuUuccaugaauau..............	5	1	seq
....aacugccgauuCuccaug...................	3	1	seq
....aacugccgauuguccauA...................	1	1	seq
....aacugccgauuCuccauga..................	10	1	seq
....aacugccgauuAuccauga..................	2	1	seq
....aacugccgauuAuccaugaa.................	3	1	seq
....aacugccgauuCuccaugaa.................	22	1	seq
....aacugccgauuguccaugaa.................	4	0	seq
....Gacugccgauuguccaugaau................	1	1	seq
....aacugccgauuCuccaugaau................	17	1	seq
....aacugccgauuguccaugaau................	1	0	seq
....aacugccgauuguccaugaaC................	1	1	seq
....aacugccgauuUuccaugaaua...............	12	1	seq
....aacugccgauuUuccaugaauau..............	3	1	seq
....auuguccaugaauauuCuuccaucucuagc	1	1	seq
....uuguccaugaauauuCuuccaucucu...	1	1	seq