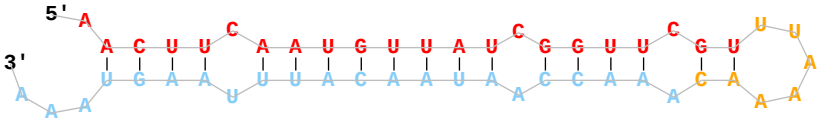
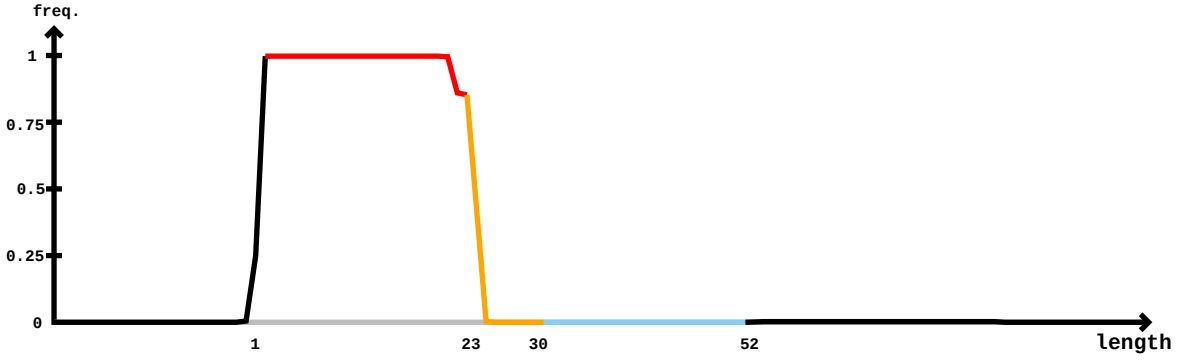
Provisional ID	:	Pocillopora_meandrina_HIv1___Sc0000010_468765
Score total	:	4.3
Score for star read(s)	:	-1.3
Score for read counts	:	0
Score for mfe	:	1
Score for randfold	:	1.6
Score for cons. seed	:	3
Total read count	:	413
Mature read count	:	413
Loop read count	:	0
Star read count	:	0





Mature Star

```
5'- guuuacaaguuuuguguuuuaacuucaauguuaucgguucguuuaaaacaaaccaauaacauuuaaguaaaaacacaaaacuuacugcauagaaaacaaacaaaccucggac  -3'    exp
    ((....(((((((((((((.((((.(((((((.((((.((.....)).)))).)))))))).)))).)))))))))))))...)).....                          reads      mm         sample
    ..................uuUacuucaauguuaucgguuc.....                                                                        2          1          seq
    ..................uUacuucaauguuaucgguuc.....                                                                        11         1          seq
    ..................uaacuucaauguuaucgguuc.....                                                                        1          0          seq
    ..................uaacuucaauguuaucgguucg.....                                                                       1          0          seq
    ..................uaacuucaauguuaucgguucgC.....                                                                      10         1          seq
    ..................uaacuucaauguuaucgguuUgu.....                                                                      16         1          seq
    ..................uUacuucaauguuaucgguucgu.....                                                                      8          1          seq
    ..................uaacuucaaugGuaucgguucgu.....                                                                      1          1          seq
    ..................uaacuucaauguuaucgguucAu.....                                                                      2          1          seq
    ..................uaacuucaauguuaucgguucgu.....                                                                      38         0          seq
    ..................uaacuucaauguCaucgguucgu.....                                                                      1          1          seq
    ..................uaacuucaauAuuaucgguucgu.....                                                                      1          1          seq
    ..................uaacuucaauguuaucgguucgA.....                                                                      2          1          seq
    ..................uaacuucaauguuaucgguucgG.....                                                                      3          1          seq
    ..................uaacuucaauguuaucgguucguA.....                                                                      2          1          seq
    ..................uUacuucaauguuaucgguucguu.....                                                                      2          1          seq
    ..................uaacuucaauguuaucgguucguu.....                                                                      1          0          seq
    ..................uaacuucaauguuaucgguucguGu.....                                                                     1          1          seq
    ...................Cacuucaauguuaucgguu.....                                                                          1          1          seq
    ...................aacuucaauguuaucgguU.....                                                                          4          1          seq
    ...................Uacuucaauguuaucgguuc.....                                                                         37         1          seq
    ...................Cacuucaauguuaucgguuc.....                                                                         1          1          seq
    ...................Uacuucaauguuaucgguucg.....                                                                        1          1          seq
    ...................aacuucaauguuaucgguuUg.....                                                                        1          1          seq
    ...................Uacuucaauguuaucgguucgu.....                                                                       10         1          seq
    ...................aacuucaauguuaucgguuUgu.....                                                                       235        1          seq
    ...................aacuucaauguuaucgguuUguu.....                                                                      18         1          seq
    ...................aacuucaauguuaucgguucgAu.....                                                                      1          1          seq
    ...................Uacuucaauguuaucgguucguu.....                                                                      1          1          seq
    ..............................................Ccacaaaacuuacugcauagaaaac.....                                       1          1          seq
```