Provisional ID	:	Pocillopora_meandrina_HIv1___Sc0000000_24086
Score total	:	4.8
Score for star read(s)	:	-1.3
Score for read counts	:	0
Score for mfe	:	1.5
Score for randfold	:	1.6
Score for cons. seed	:	3
Total read count	:	3297
Mature read count	:	3297
Loop read count	:	0
Star read count	:	0

Mature   **Star**

```
5'- ugcaguuguggguugugagaugcagcaaugucguauccuuuuaauuuauuuaaaggcauacgaaauugcuucacuuuacaacccucucuaaaaucucaauuuuguuac  -3'    exp
    ...((..(.((((((((((.((.((((((.(((((.(((((((.....)))))))).))))))..))))))).)).)))))))).).))................    reads    mm        sample
    ugcaguuguggguugugaga.............................................................................................    4        0          seq
    .gcaguugugggguugugaga............................................................................................    5        0          seq
    ..caguugugggguugugaga............................................................................................    2        0          seq
    ..................Aaugcagcaaugucguauccuuuu.......................................................................    1        1          seq
    ..................Cugcagcaaugucguauccuuuu.......................................................................    1        1          seq
    ...................ugcagcaaugucguaucc...........................................................................    1        0          seq
    ...................ugcagcaaugucguaucA...........................................................................    1        1          seq
    ...................ugcagcaaugucguaucAu..........................................................................    1        1          seq
    ...................ugcagcaaugucguauccG..........................................................................    3        1          seq
    ...................ugcagcGaugucguauccu..........................................................................    1        1          seq
    ...................ugcagcaaugucguauccu..........................................................................    8        0          seq
    ...................ugcagcaaugucguauccC..........................................................................    1        1          seq
    ...................ugcagcaaugucguauccA..........................................................................    5        1          seq
    ...................ugcagcaaugucguauccuA.........................................................................    6        1          seq
    ...................ugcagcaaugucguauccuC.........................................................................    2        1          seq
    ...................ugcagcaaugucguauccGu.........................................................................    1        1          seq
    ...................ugcagcaaugucguauccuu.........................................................................    12       0          seq
    ...................ugcagcaaugucguGuccuuu........................................................................    1        1          seq
    ...................ugcagcaaugucgCauccuuu........................................................................    1        1          seq
    ...................ugcagcaaugucguauccuAu........................................................................    3        1          seq
    ...................ugcagcaaugucguauccuuG........................................................................    1        1          seq
    ...................ugcagcaaugucguauccuuu........................................................................    167      0          seq
    ...................Cgcagcaaugucguauccuuu........................................................................    3        1          seq
    ...................ugcagcaaugucguauccuuA........................................................................    7        1          seq
    ...................ugcagcaaugucguauccCuu........................................................................    3        1          seq
    ...................Agcagcaaugucguauccuuu........................................................................    1        1          seq
    ...................ugcagcaaugucguauccuGu........................................................................    2        1          seq
    ...................Ggcagcaaugucguauccuuu........................................................................    1        1          seq
    ...................ugcagcaaugucguauccuuC........................................................................    25       1          seq
    ...................ugcagcaaugucguaGccuuu........................................................................    1        1          seq
    ...................ugcagcaaugucguauccuCu........................................................................    1        1          seq
    ...................ugcagcaaugucguauccuuCu.......................................................................    1        1          seq
    ...................ugUagcaaugucguauccuuuu.......................................................................    1        1          seq
    ...................ugcaUcaaugucguauccuuuu.......................................................................    1        1          seq
```

ugcaguugugggguugugaga ugcagcaaugucguauccuuua auuuauuua aggcauacgaaauugcuucacu uuacaacccucucuaaaaucucaauuuuguuac

Sequence	Count	Mismatch	
...................ugcGgcaaugucguauccuuuu...	4	1	seq
...................ugcagcaGgucguauccuuuu...	2	1	seq
...................ugcagcaaugucguauccuuuu...	664	0	seq
...................ugcagcaaugucAuauccuuuu...	14	1	seq
...................ugcagcaaugucguauccuuAu...	5	1	seq
...................ugcagcaauguUguauccuuuu...	2	1	seq
...................ugcagcaCgucguauccuuuu...	1	1	seq
...................Agcagcaaugucguauccuuuu...	2	1	seq
...................ugcagcaaugucguauAcuuuu...	1	1	seq
...................ugcaAcaaugucguauccuuuu...	1	1	seq
...................ugcagUaaugucguauccuuuu...	4	1	seq
...................ugcagcaaugucguauccuuuC...	66	1	seq
...................ugAagcaaugucguauccuuuu...	2	1	seq
...................ugcagcaaugucguauccuuuA...	24	1	seq
...................ugcagcaaugucguauccuuuG...	2	1	seq
...................ugcagcaaugucguauccCuuu...	4	1	seq
...................ugcagcaaugCcguauccuuuu...	2	1	seq
...................ugcagcaaugucguauccuAuu...	1	1	seq
...................ugcagcaaugucguauccuuGu...	9	1	seq
...................ugcagcaaugucguauccuGuu...	1	1	seq
...................ugcagcGaugucguauccuuuu...	3	1	seq
...................ugcagcaaugucguGuccuuuu...	2	1	seq
...................Cgcagcaaugucguauccuuuu...	17	1	seq
...................ugcagcaaugucgAauccuuuu...	1	1	seq
...................ugcagcaauCucguauccuuuu...	1	1	seq
...................ugcagcaaugucguaucAuuuu...	1	1	seq
...................ugcagcaaugucguaucUuuuu...	1	1	seq
...................ugcagcaaugucguGuccuuuua..	1	1	seq
...................ugcagcaaugucguaGccuuuua..	1	1	seq
...................ugcaUcaaugucguauccuuuuU...	1	1	seq
...................ugcagcaaugucUuauccuuuua..	2	1	seq
...................ugcagcaaugCcguauccuuuua..	2	1	seq
...................ugcagcaaugucguauccCuuua..	6	1	seq
...................ugcagcaaugucCuauccuuuua..	1	1	seq
...................Cgcagcaaugucguauccuuuua..	13	1	seq
...................ugcGgcaaugucguauccuuuua..	3	1	seq
...................ugcagUaaugucguauccuuuua..	1	1	seq
...................ugcagcaaugucguaCccuuuua..	2	1	seq
...................ugcagcaaugucAuauccuuuua..	4	1	seq
...................ugcagcaGgucguauccuuuua..	1	1	seq
...................ugAagcaaugucguauccuuuua..	1	1	seq
...................ugcagcaaugucguauUcuuuua..	5	1	seq
...................ugcagcaauguUguauccuuuua..	2	1	seq
...................Agcagcaaugucguauccuuuua..	7	1	seq
...................ugcagcaaugucguauccuuuAa..	1	1	seq
...................ugcagcaaugucgCauccuuuua..	1	1	seq
...................ugcagcaaugucguauccuuuuU...	22	1	seq
...................uAcagcaaugucguauccuuuua..	2	1	seq
...................ugcagcaaugucguauccuuuuG...	5	1	seq
...................ugcagcGaugucguauccuuuua..	7	1	seq
...................ugcagcaaugucguauccuCuua..	1	1	seq
...................ugcagcaauguAguauccuuuua..	1	1	seq
...................ugcagcaaugucguaucAuuuua..	1	1	seq
...................ugcagcaaugucguauccuuCua..	1	1	seq
...................ugcagcaaugucguauccuuuua..	924	0	seq
...................ugcagcaaugucgGauccuuuuaa...	3	1	seq
...................ugUagcaaugucguauccuuuuaa...	2	1	seq
...................ugcagcaaCgucguauccuuuuaa...	3	1	seq
...................ugcagcaaugucguaucUuuuuaa...	13	1	seq
...................ugcagUaaugucguauccuuuuaa...	1	1	seq
...................ugcagcaaugucguaGccuuuuaa...	2	1	seq
...................ugcagcaaugucguauccuuuuaa...	561	0	seq
...................ugcagcaaugucguauccuuuuaU..	13	1	seq
...................ugcagcaaugucguauccuuCuaa...	1	1	seq
...................ugcagcaaugucguauUcuuuuaa...	1	1	seq
...................ugcagcaaugucguauccCuuuaa...	4	1	seq
...................ugcagcaaugCcguauccuuuuaa...	3	1	seq
...................ugcAgcaaugucguauccuuuuUa..	2	1	seq
ugcaguugugggguugugaga Agcagcaaugucguauccuuuuaa auuuauuua aggcauacgaaauugcuucacu uuacaacccucucuaaaaucucaauuuuguuac	7	1	seq
...................ugcagcaaugucguauccuuAuaa...	1	1	seq

ugcaguuguggguugugagaugcagcaaugucguauccuuuuaauuuauuuaaaggcauacgaaauugcuucacuuuacaacccucucuaaaaucucaauuuuguuac

Sequence	Count	Flag	
...................ugAagcaaugucguauccuuuuaa..	1	1	seq
...................ugcagcaaugucguauccuuuuGa..	4	1	seq
...................ugcagcGaugucguauccuuuuaa..	3	1	seq
...................ugcagcaGugucguauccuuuuaa..	2	1	seq
...................ugcagcaaugucguauccuuuuaG..	2	1	seq
...................ugcagcaaugucgCauccuuuuaa..	1	1	seq
...................uAcagcaaugucguauccuuuuaa..	2	1	seq
...................ugcaAcaaugucguauccuuuuaa..	2	1	seq
...................ugcagcaaugucAuauccuuuuaa..	2	1	seq
...................ugcGgcaaugucguauccuuuuaa..	1	1	seq
...................ugcagcaaCgucguauccuuuuaau...	1	1	seq
...................ugcagcaaugucguaucUuuuuaau...	3	1	seq
...................ugcagcaaGgucguauccuuuuaau...	1	1	seq
...................ugcagcaaugCcguauccuuuuaau...	2	1	seq
...................ugcagcGaugucguauccuuuuaau...	1	1	seq
...................ugcagcaaugucguauccuuuuaGu...	3	1	seq
...................ugcagcaaugucgCauccuuuuaau...	1	1	seq
...................Cgcagcaaugucguauccuuuuaau...	4	1	seq
...................ugcagcaaugucguauccuGuuaau...	3	1	seq
...................ugcagcaGugucguauccuuuuaau...	1	1	seq
...................ugcagcaaugucguauccuuuuaaG...	8	1	seq
...................ugcagcaaugucguauccuuuuaUu...	7	1	seq
...................ugcagcaaugucguauccuuuuaau...	374	0	seq
...................ugcagcaaugucguGuccuuuuaau...	1	1	seq
...................ugcagcaaugucguauccuuuuaaC...	70	1	seq
...................ugcagcaaugucguauccCuuuaau...	2	1	seq
...................ugcagcaaugucguauccuuuuaaA...	15	1	seq
...................ugcagcaaugucguauccuuuuaaCu..	3	1	seq
...................ugcagcaaugucguauccuuuuaauu..	18	0	seq
...................ugcagcaaugucguauccuuuuaaAu..	1	1	seq
...................ugcagcaaugucguauccuuCuaauu..	1	1	seq
...................ugcagcaaugucguauccuuuuaauC...	3	1	seq
...................ugcagcaaugucguauccuuuuaauuA..	2	1	seq
...................ugcagcaaugucguauccuuuuaauuuauu...	2	0	seq
....................gcagcaaugucguauccuuuuaau..	1	0	seq
....................gcagcaaugucguauccuuuuaaC..	1	1	seq
......................cagcaaugucguauccuC..	1	1	seq
......................cagcaaugucguauccuuuuu...	3	0	seq
......................cagcaaugucguauccuuuuua..	7	0	seq
......................cagcaaugucguauccuuuuaa..	2	0	seq
....uuacaacccucucuaaaaucucaauuu......	2	0	seq