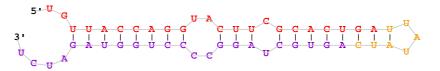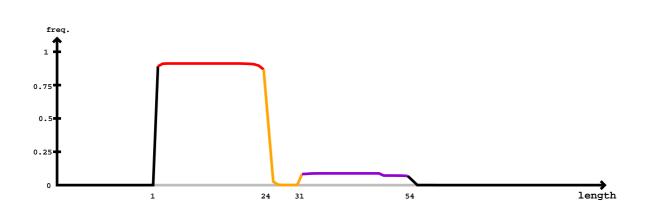| Provisional ID	: Pocillopora_meandrina_HIv1___Sc0000002_184721
Score total	: 1406.8
Score for star read(s)	: 3.9
Score for read counts	: 1400.1
Score for mfe	: 1.8
Score for randfold	: 1.6
Score for cons. seed	: -0.6
Total read count	: 2758
Mature read count	: 2515
Loop read count	: 0
Star read count	: 243

Mature　　**Star**

```
5'- uugaaacaagagggaagaguguuaccagguacuucgcacugauuauaucagugcuaggccccugguagaucucuugccagcuugcauugcuccaguauucauacuuuccuau  -3'   obs
    uugaaacaagagggaagaguguuaccagguacuucgcacugauuauaucagugcuaggccccugguagaucucuugccagcuugcauugcuccaguauucauacuuuccuau         exp
    .(((((((....((.((((((..((((((((((..(((.(((((((((..)))))))))..)))..)))))))))..)))))))).)))))).)).((.......))....))).)))))..........
```

	reads	mm	sample
...................Cguuaccagguacuucgcacuga.................................	1	1	seq
...................uguuaccagguacuucgA.....................................	2	1	seq
...................uguuaccagguacuucgc.....................................	1	0	seq
...................uguuaccagguacuucgca....................................	4	0	seq
...................uguuaccagguacuucgcaA...................................	2	1	seq
...................uguuaccagguacuucgcac...................................	4	0	seq
...................uguuaccagguacuucgcaG...................................	1	1	seq
...................uguuaccagguacuucgcGc...................................	1	1	seq
...................uguuaccagguacuucgcacC..................................	5	1	seq
...................uguuaccagguacuucgcacA..................................	5	1	seq
...................uguuaccagguacuucgcacG..................................	1	1	seq
...................uguuaccagguacuucgcacu..................................	14	0	seq
..................Cguuaccagguacuucgcacu...................................	1	1	seq
..................Cguuaccagguacuucgcacug..................................	1	1	seq
...................uguuaccagguacuucgcacAg.................................	1	1	seq
...................uguuaccagguacuucgcacug.................................	48	0	seq
...................uguuaccagguacuucgcacuA.................................	20	1	seq
...................uguuaccGgguacuucgcacug.................................	1	1	seq
...................uguuaccagguacuucgcacuga................................	1808	0	seq
..................Gguuaccagguacuucgcacuga.................................	1	1	seq
...................uguuaccagguacuucgcacCga................................	2	1	seq
...................uguuaccagAuacuucgcacuga................................	1	1	seq
...................uguuaccagguacuucgcaUuga................................	4	1	seq
...................uguuaccaggCacuucgcacuga................................	3	1	seq
...................uguuaccagguacuucgcUcuga................................	1	1	seq
...................uguuaccagguacuucAcacuga................................	3	1	seq
...................uguuaccagguacuuUgcacuga................................	1	1	seq
...................uguuaccagguacuucgcGcuga................................	2	1	seq
...................uguuGccagguacuucgcacuga................................	13	1	seq
...................uguuaccagguacuucgcacugU................................	97	1	seq
...................uguuaccagguacuCcgcacuga................................	5	1	seq
...................ugCuaccagguacuucgcacuga................................	6	1	seq
...................uguuaccGgguacuucgcacuga................................	12	1	seq

```
uugaaacaagagggaagaguguuaccagguacuucgcacugauuauaucagugcuaggccccugguagaucucuugccagcuugcauugcuccaguauucauacuuuccuau
```

Sequence	Count	Flag	
..................uguuaccagguacuucgAacuga..	1	1	seq
..................uguuaccagguacuucgcacAga..	1	1	seq
..................uguuaccagguacCucgcacuga..	3	1	seq
..................uguuaccagguacuucgcacugG..	51	1	seq
..................uguuaccagguacuucgcacuAa..	7	1	seq
..................uguuUccagguacuucgcacuga..	1	1	seq
..................Cguuaccagguacuucgcacuga..	28	1	seq
..................uguuacUagguacuucgcacuga..	2	1	seq
..................uguuaccagguacAucgcacuga..	1	1	seq
..................uguuacAagguacuucgcacuga..	1	1	seq
..................uCuuaccagguacuucgcacuga..	1	1	seq
..................uguuaccagguacuucgcacugC..	12	1	seq
..................ugAuaccagguacuucgcacuga..	1	1	seq
..................Aguuaccagguacuucgcacuga..	2	1	seq
..................uguuaccUgguacuucgcacuga..	1	1	seq
..................uguuaccagguacuucCcacuga..	1	1	seq
..................uguCaccagguacuucgcacuga..	1	1	seq
..................uguuaccagguacuucgUacugau..	1	1	seq
..................uguuaccagguacuCcgcacugau..	1	1	seq
..................Cguuaccagguacuucgcacugau..	1	1	seq
..................uguuaccagguacuucgcacugaC..	41	1	seq
..................uguuaccagguacuucgcacugUu..	2	1	seq
..................uguuaccagguacuucgcacugaA..	16	1	seq
..................uguuaccagguacuucgcacugaG..	2	1	seq
..................uguuaccagguacuucgcacugau..	131	0	seq
..................uguuaccagguacuucgcacugGu..	7	1	seq
..................ugCuaccagguacuucgcacugau..	1	1	seq
..................uguuaccagguacuucgcacugauA...	8	1	seq
..................uguuaccagguacuucgcacugauu...	41	0	seq
..................uguuaccagguacuucgcacugaAu...	1	1	seq
..................uguuaccagguacuucgcacugaCu...	2	1	seq
..................uguuaccagguacuucgcacugauC...	6	1	seq
..................Cguuaccagguacuucgcacugauu..	1	1	seq
..................uguuaccagguacuucgcacugaAua..	1	1	seq
..................uguuaccagguacuucgcacugauuU..	6	1	seq
..................uguuaccagguacuucgcacugauuGu..	1	1	seq
..................uguuaccagguacuucgcacugauuaua...	1	0	seq
..................uguuaccagguacuucgcacugauuauauc..	1	0	seq
...................guuaccagguacuucgcacu..	2	0	seq
...................Cuuaccagguacuucgcacu..	2	1	seq
...................guuaccagguacuucgcacug...	6	0	seq
...................guuaccagguacuucgcacuA...	1	1	seq
...................guuaccagguacuucgcacuga..	28	0	seq
...................Cuuaccagguacuucgcacuga..	3	1	seq
...................guuaccagguacuucgcacugU..	2	1	seq
...................Auuaccagguacuucgcacuga..	1	1	seq
...................guuaccagguacuucgcacugCu...	1	1	seq
...................Cuuaccagguacuucgcacugau..	1	1	seq
...................guuaccagguacuucgcacugau..	2	0	seq
...................guuaccagguacuucgcacugaC..	1	1	seq
...................guuaccagguacuucgcacugauu...	2	0	seq
...................guuaccagguacuucgcacugauA...	1	1	seq
....................uuaccagguacuucgcacug..	1	0	seq
....................uuaccagguacuucgcacuga...	2	0	seq
....................uuaccagguacuucgcacugauA...	1	1	seq
....cagugUuaggccccugguagaucu...............................	3	1	seq
....Ugugcuaggccccuggu.......................................	1	1	seq
....agugUuaggccccuggu.......................................	23	1	seq
....agugcuaggccccuggu.......................................	20	0	seq
....aAugcuaggccccuggu.......................................	1	1	seq
....agugUuaggccccugguaga...................................	1	1	seq
....agugUuaggccccugguagau..................................	1	1	seq
....agugcuaggccccugguagauc.................................	2	0	seq
....agugcuaggccccugguagauU.................................	1	1	seq
....agugcuaggccccugguGgauc.................................	2	1	seq
....agugcuaggccccugguagauA.................................	1	1	seq
....agugcuaggccccugguagaucu................................	51	0	seq
....agugGuaggccccugguagaucu................................	1	1	seq
uugaaacaagagggaagaguguuaccagguacuucgcacugauuauaucCgugcuaggccccugguagaucucuugccagcuugcauugcuccaguauucauacuuuccuau	2	1	seq
....agugUuaggccccugguagaucu................................	77	1	seq
```

uugaaacaagagggaagaguguuaccagguacuucgcacugauuauaucagugcuaggccccugguagaucucuugccagcuugcauugcuccaguauucauacuuuccuau

| | | | |
|---|---|---|---|
| .............................................................agugcuaggccccugguagaucC........................................... | 16 | 1 | seq |
| .............................................................agugcuaggccccugguagaucA........................................... | 20 | 1 | seq |
| .............................................................Ggugcuaggccccugguagaucu........................................... | 1 | 1 | seq |
| .............................................................agCgcuaggccccugguagaucu........................................... | 1 | 1 | seq |
| .............................................................agugcuaggccccugguagaucuA.......................................... | 3 | 1 | seq |
| .............................................................agugUuaggccccugguagaucuc.......................................... | 1 | 1 | seq |
| ..............................................................gugcuaggccccugguagaucu........................................... | 1 | 0 | seq |
| ..............................................................gugcuaggccccugguagaucC........................................... | 1 | 1 | seq |
| ..............................................................gugUuaggccccugguagaucu........................................... | 1 | 1 | seq |
| ..............................................................gugcuaggccccugguagaucucuugc..................................... | 1 | 0 | seq |
| ...............................................................ugcuaggccccugguagauA............................................ | 1 | 1 | seq |
| ...............................................................ugcuaggccccugguagaucA............................................ | 2 | 1 | seq |
| ...............................................................ugcuaggccccugguagaucC............................................ | 3 | 1 | seq |
| ...............................................................ugcuaggccccugguagaucu............................................ | 1 | 0 | seq |
| ...............................................................ugUuaggccccugguagaucu............................................ | 1 | 1 | seq |
| ................................................................gcuagCccccugguagaucu............................................ | 1 | 1 | seq |
| .................................................................cuaggccccugguagaucA............................................ | 1 | 1 | seq |