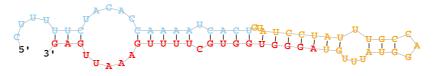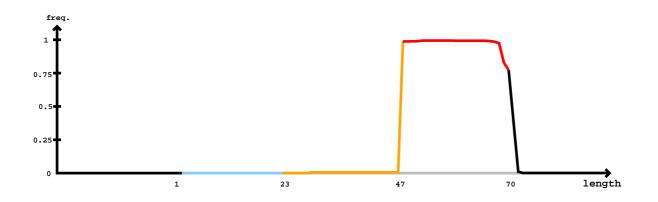| Provisional ID	: Pocillopora_meandrina_HIv1___Sc0000018_677916
Score total	: 2.8
Score for star read(s)	: -1.3
Score for read counts	: 0
Score for mfe	: -0.4
Score for randfold	: 1.6
Score for cons. seed	: 3
Total read count	: 958
Mature read count	: 953
Loop read count	: 5
Star read count	: 0

Star   **Mature**

```
5'- uggacugauaacacagacaaaauucuuuuucuacaccaaaaucacuguauccuauuugccagguauuuguaggguggugcuuuugaaauugaguuuauucugauaucaggguc  -3'   exp
    ....(((((((...((((..(((((((..(((((..(((((...((((.((((.(((.......)))))..)).)).)))))).....)))).))))))..))))...)))))))....   reads   mm    sample
    ...............................................cuauuugccagguauuuguC...........................................     2     1     seq
    ...............................................cuauuugccagguauuugu............................................     3     0     seq
    ...............................................cuauuugccagguauuuguaggguggugcu..................................     1     0     seq
    ....................................................agggguggugcuuuugaaa........................................     2     0     seq
    ....................................................aggguggugcuuuugaaau.......................................     5     0     seq
    ....................................................aggguggugcuuuugaaauu......................................     9     0     seq
    ....................................................agggAggugcuuuugaaauu......................................     1     1     seq
    ....................................................aggguggugcuuuugaaauug.....................................   113     0     seq
    ....................................................agggugguCcuuuugaaauug.....................................     1     1     seq
    ....................................................aggguggugcuuuugGaauug.....................................     1     1     seq
    ....................................................aggguggugcuuuugaaaCg......................................     4     1     seq
    ....................................................aggguggugcuuuugaaauuU.....................................     1     1     seq
    ....................................................aAgguggugcuuuugaaauug.....................................     2     1     seq
    ....................................................aggguggugcuuuugaaauuA.....................................    15     1     seq
    ....................................................aggguggugcuuuugaGauug.....................................     1     1     seq
    ....................................................aggguggugcuuuCgaaauug.....................................     1     1     seq
    ....................................................aggguggugcuuuugaaauGg.....................................     1     1     seq
    ....................................................agggCggugcuuuugaaauuga....................................     1     1     seq
    ....................................................aggguggugcuuuugaaauugG....................................     2     1     seq
    ....................................................aggguggugcuuuuCgaaauuga...................................     1     1     seq
    ....................................................Ggguggugcuuuugaaauuga....................................     3     1     seq
    ....................................................aggguggugcuuuugaaauugU....................................     3     1     seq
    ....................................................aggguggugcuuuugaaauugC....................................     2     1     seq
    ....................................................aggguggugcuuuugGaauuga....................................     1     1     seq
    ....................................................aggguggugcuuuugaaauuga....................................    42     0     seq
    ....................................................aggguggugcuuuugaaauCgag...................................     1     1     seq
    ....................................................aggguggugcuuuugaGauugag...................................     2     1     seq
    ....................................................aUgguggugcuuuugaaauugag...................................     1     1     seq
    ....................................................aggguggCgcuuuugaaauugag...................................     1     1     seq
    ....................................................aggguggugcuuuugaaGuugag...................................     2     1     seq
    ....................................................aggCuggugcuuuugaaauugag...................................     2     1     seq
    ....................................................aAgguggugcuuuugaaauugag...................................     4     1     seq
    ....................................................Cgguggugcuuuugaaauugag....................................     8     1     seq
    ....................................................Ugguggugcuuuugaaauugag....................................     2     1     seq
```

uggacugauaacacagacaaaauucuuuuucuacaccaaaaucacuguauccuauuugccagguauuuguaggguggugcuuuugaaauugaguuuauucugauaucaggguc

Sequence	Count	Flag	
....aggguggugcuuGugaaauugag....................	1	1	seq
....aggguggugcuuuugGaauugag....................	3	1	seq
....agUguggugcuuuugaaauugag....................	1	1	seq
....aggguggugcuuuugaaauugUg....................	1	1	seq
....aggguggugcuuuugaaCuugag....................	1	1	seq
....Gggguggugcuuuugaaauugag....................	2	1	seq
....aggguggugcuuuAgaaauugag....................	1	1	seq
....aggguggugcuuuugaaauuAag....................	1	1	seq
....aggguggugcuuuugaaauugaU....................	10	1	seq
....aggguggugcuuuugaaauugag....................	569	0	seq
....aggguggugcuuuugaaauugaC....................	10	1	seq
....aggguggugcuuuugaaauugGg....................	2	1	seq
....aggguggugcuuuugaaauugaA...................	58	1	seq
....aggguggugcuuuugaaaCugagu...................	1	1	seq
....aggguggugcuuuugaaauugagu..................	22	0	seq
....aggguggugcuuuugaaauugagC..................	3	1	seq
....aggguggugcuuuugaaauugaCu..................	1	1	seq
....aggguggugcuuuugaaauugaUu..................	1	1	seq
....aggguggugcuuuugaaauugagA..................	15	1	seq
....aggguggugcuuuugaaauugaUuu.................	1	1	seq
....aggguggugcuuuugaaauugaguu.................	3	0	seq
....aggguggugcuuuugaaauugagAu.................	2	1	seq
....aggguggugcuuuugaaauugaguC.................	1	1	seq
....aggguggugcuuuugaaauugaguA.................	2	1	seq
....Aguggugcuuuugaaauugag....................	1	1	seq
....uggugcuuuugaaauugaU....................	1	1	seq
....uggugcuuuugaaauugagA...................	4	1	seq
....uuuauucugauaucaggguc	1	0	seq