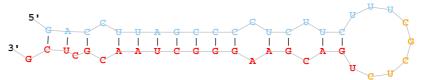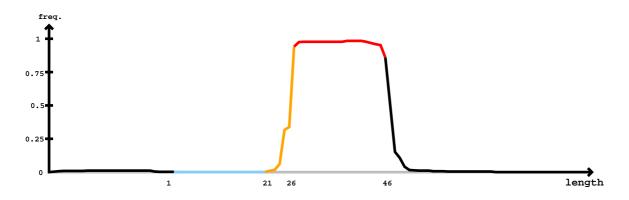Provisional ID   : Pocillopora_meandrina_HIv1___Sc0000022_747819
Score total   : 1.7
Score for star read(s)   : -1.3
Score for read counts   : 0
Score for mfe   : 2
Score for randfold   : 1.6
Score for cons. seed   : -0.6
Total read count   : 402
Mature read count   : 402
Loop read count   : 0
Star read count   : 0

Star   **Mature**

```
5'- aaggguuuuaaagcugacguuucggaccuuagccccucuucuuucgcucugacgaagggcuaacgcucgaaacuucagcuucgaaacucuuuuugguggccacuuuucguu  -3'   exp
    (((((((((..(((((((((.(((((((..(.(((((((((.((.((.........)).)).)))))))).).))))))).))))))))..)))))))))..................  reads    mm         sample
    aGgguuuuaaagcugacgu......................................................................................................   1      1          seq
    aaggguuuCaaagcugacgu.....................................................................................................   2      1          seq
    .agggguuuuaaagcugacgGu...................................................................................................   1      1          seq
    ......uuAaaagcugacguuucgg................................................................................................   1      1          seq
    ...............................ucgcucugacgaagggcuaacgcuc.................................................................   1      0          seq
    ...............................cgcucugacgaagggcuaacgcucg.................................................................   2      0          seq
    ...............................cgcucugacgaagggcuaacgcucgaa...............................................................   1      0          seq
    ...............................cgcucugacgaagggcuaacgcucgaaa..............................................................   1      0          seq
    ...............................cgcucugacgaaUggcuaacgcucgaaa..............................................................   1      1          seq
    ...............................gcucugacgaagggcuaacgcucU..................................................................   1      1          seq
    ...............................gcucugacgaagggcuaacgcucg..................................................................   1      0          seq
    ................................cucugacgaagggcuaac.......................................................................   3      0          seq
    ................................cucugacgaagggcuaacgc.....................................................................   1      0          seq
    ................................cucugacgaagggcuaacgcuc...................................................................   3      0          seq
    ................................cucugacgaagggcuaacgcucg..................................................................   3      0          seq
    ................................cucugacgaagggcuaacgcucga.................................................................   4      0          seq
    ................................cucugacgaagggcuaacgcucgaa................................................................   3      0          seq
    ................................cucugacgaagggcuaacgcucgaaac..............................................................   2      0          seq
    .................................ucugacgaagggcuaacg......................................................................   3      0          seq
    .................................ucugacgaagggcuaacU......................................................................   1      1          seq
    .................................ucugacgaagggcuaacgc.....................................................................   3      0          seq
    .................................ucugacgaagggcuaacgcu....................................................................   3      0          seq
    .................................ucugacgaagggcuaacgcAc...................................................................   1      1          seq
    .................................ucugacgaagggcuaacgcuc...................................................................   8      0          seq
    .................................ucugacgaagggcuaacgcucg..................................................................  81      0          seq
    .................................ucugacgaagggcuaacgcucga.................................................................  11      0          seq
    .................................ucugacgaagggcuaacgcucgaaac..............................................................   1      0          seq
    ..................................cugacgaagggcuaacgcuc...................................................................   3      0          seq
    ..................................cugacgaagggcuaacgcucg..................................................................   3      0          seq
    ..................................cugacgaagggcuaacgcucga.................................................................   1      0          seq
    ..................................cugacgaagggcuaacgcucgaaG...............................................................   1      1          seq
    ..................................cugacgaagggcuaacgcucgaaac..............................................................   1      0          seq
    ..................................cugacgaagggcuaGcgcucgaaacuu............................................................   1      1          seq
    ...................................ugacgaagggcuaacgcAc..................................................................   2      1          seq
```

aaggguuuuaaagcugacguuucggaccuuagcccucuucuuucgcucugacgaagggcuaacgcucgaaacuucagcuucgaaacucuuuuugguggccacuuuucguu

Sequence	Count	Mismatch	
....ugacgaagggcuaacgcuc...	21	0	seq
....ugacgaagggcuaacgcAcg...	1	1	seq
....ugacgaagggcuaacgcucg...	174	0	seq
....ugacgaUgggcuaacgcucg...	1	1	seq
....ugacgaagggcuaacgcucga..	21	0	seq
....ugacgaagggcuaacgcucgC..	2	1	seq
....ugacgaagggcuaacgcucgaa...	16	0	seq
....ugacgaagggcuaacgcucgaaa..	23	0	seq
....ugacgaagggcuaacgcucgaaG..	2	1	seq
....ugacgaagggcuaacgcucgaaac.......................................	2	0	seq
....gacgaagggcuaacgcuc...	2	0	seq
....gacgaagggcuaacgcucg..	6	0	seq
....gacgaagggcuaacgcucgaaa.......................................	1	0	seq
....gacgaagggcuaacgcucgaaac......................................	5	0	seq
....gacgaagggcuaacgcuUgaaacu.....................................	1	1	seq
....acgaagggcuaacgcucg...	1	0	seq
....uaacgcucgaaacuucaA................................	2	1	seq
....uaacgcucgaaacuucagcuu.............................	1	0	seq
....aacuucagcuucgaaacucuu..................	2	0	seq