Provisional ID   : Pocillopora_meandrina_HIv1___Sc0000002_178665
Score total   :   290
Score for star read(s) :   3.9
Score for read counts :   279.5
Score for mfe   :   2
Score for randfold   :   1.6
Score for cons. seed :   3
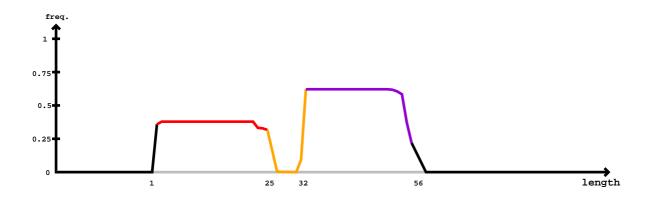Total read count   :   560
Mature read count   :   212
Loop read count   :   0
Star read count   :   348


```
5'- aacuuaaacucuggucauuuacgauguugucuugucugcaagcuugagguuugaagacgagacaacaguguaaaagaacaguguuuuucucuguuuggguggcgaaagcuucaa  -3'   obs
    aacuuaaacucuggucauuuacgauguugucuugucugcaagcuugagguuugaagacgagacaacaguguaaaagaacaguguuuuucucuguuuggguggcgaaagcuucaa        exp
    .....(((((.(((.((.(((((((.((((((((((((((.(((((.....)))))).))))))))))))))).))))))).)).))).)))).........(((((.(((....)))))))).     reads     mm     sample
    ................uacgauguugucuugucugca...................................................................................... 19      0     seq
    ................uacgauguugucuugucugcG...................................................................................... 5       1     seq
    ................uacgauguugucuugucugcaa..................................................................................... 2       0     seq
    ................uacgauguugucuugucugcaaA.................................................................................... 3       1     seq
    ................uacgauguugucuugucugcaag.................................................................................... 4       0     seq
    ................uacgauguugucuugucugcaGgc................................................................................... 1       1     seq
    ................uacgauguugucuugucugcaagG................................................................................... 1       1     seq
    ................Cacgauguugucuugucugcaagc................................................................................... 1       1     seq
    ................uacgauguugucuugucugcaagU................................................................................... 17      1     seq
    ................uacgauguugCcuugucugcaagc................................................................................... 1       1     seq
    ................uacgaugGugucuugucugcaagc................................................................................... 1       1     seq
    ................uacgGuguugucuugucugcaagc................................................................................... 1       1     seq
    ................uacgauguugucuugucCgcaagc................................................................................... 1       1     seq
    ................uacgauguugucuugucugcaaAc................................................................................... 3       1     seq
    ................uacgauguugucuugucugcaaCc................................................................................... 1       1     seq
    ................uGcgauguugucuugucugcaagc................................................................................... 1       1     seq
    ................Aacgauguugucuugucugcaagc................................................................................... 2       1     seq
    ................uacgauguugucuugucugcaagA................................................................................... 19      1     seq
    ................uacgauguugucuugCcugcaagc................................................................................... 1       1     seq
    ................uacgauguugucuugucugcaagc................................................................................... 108     0     seq
    ................uacgauguugucuugucugcaagcC.................................................................................. 1       1     seq
    ................uacgauguugucuugucugcaagcA.................................................................................. 2       1     seq
    ................uacgauguugucuugucugcaagcu.................................................................................. 3       0     seq
    ................uacgauguugucuugucugcaagcuu................................................................................. 2       0     seq
    ................uacgauguugucuugucugcaagcuuga.............................................................................. 1       0     seq
    .................acgauguugucCugucugca...................................................................................... 1       1     seq
    .................acgauguugucuugucugca...................................................................................... 1       0     seq
    .................Ccgauguugucuugucugcaagc.................................................................................. 1       1     seq
    .................acgauguugucuugucugcaagA.................................................................................. 2       1     seq
    .................acgauguugucuugucugcaagc.................................................................................. 6       0     seq
    .....................................uuugaagacgagacaacag.................................................................. 1       0     seq
    .....................................uuugaagacgagacaacagu................................................................. 1       0     seq
    .....................................uuugaagacgagacaacagug................................................................ 3       0     seq
```

aacuuaaacucuggucauuuacgauguugucuugucugcaagcuugagguuugaagacgagacaacaguguaaaagaacaguguuuuucucuguuugguggcgaaagcuucaa

Sequence	Count	Flag	
....uuugaagacgagacaacagugG...	4	1	seq
....uuugaagacgagacaacagugA...	3	1	seq
....uuugaagacgagacaacagugu...	28	0	seq
....uuugaagacgagacaacagugC...	3	1	seq
....uuugaagGcgagacaacagugu...	1	1	seq
....uuugaagacgagacaacaguguG...	1	1	seq
....uuugaagacgagacaacaguguC...	2	1	seq
....uuugaagacgagacaacagugua...	3	0	seq
....uuugaagacgagacaacaguguU...	2	1	seq
....uugaagacgagacaacag...	1	0	seq
....uugaagacgagacaacagu...	4	0	seq
....uugaagacgagacaacaAu...	1	1	seq
....uugaGgacgagacaacagu...	1	1	seq
....uugaagacAagacaacagug...	1	1	seq
....uugaagacgagacaacaguA...	2	1	seq
....uugaagacgagacaacagCg...	1	1	seq
....uugaagacgagacaacagug...	5	0	seq
....uugaagacgagacaacagCgu...	1	1	seq
....uugaagacgagacGacagugu...	1	1	seq
....Cugaagacgagacaacagugu...	1	1	seq
....uugaagacgagacaacagugu...	59	0	seq
....uugaagacgagacaacagugC...	13	1	seq
....uugaGgacgagacaacagugu...	1	1	seq
....uugaagacgagacaacagugA...	1	1	seq
....uugaagacgagacaacagugG...	2	1	seq
....uugaagacgagacGacagugua...	3	1	seq
....uugaagacgGacaacagugua...	2	1	seq
....Augaagacgagacaacagugua...	1	1	seq
....uugaGgacgagacaacagugua...	1	1	seq
....uugaagacgagacaacGgugua...	1	1	seq
....uugaagacgagacaacaguguG...	1	1	seq
....uugaagacgagacaacagAgua...	1	1	seq
....uugaagacgagacaacagugua...	68	0	seq
....uugaagaAgacaacagugua...	1	1	seq
....uugaagacgagacaacaguguaa...	81	0	seq
....uugaagacgagacaacaguguaU...	5	1	seq
....uugaagacgagacaacaguguGa...	3	1	seq
....uugaagacgagacaacagugAaa...	2	1	seq
....uugaagacgagacaacaguguaG...	1	1	seq
....uugaagacgagGcaacagugua...	1	1	seq
....uugaagacgagacaacaguguaC...	1	1	seq
....uugaagacgagacaacaguguaaa...	10	0	seq
....uugaagacgagacUacaguguaaa...	1	1	seq
....uugaagacgagacaacaguguaaC...	2	1	seq
....uugaagacgagacaacaguguaaU...	15	1	seq