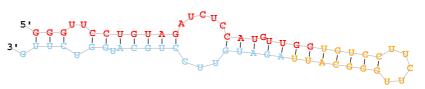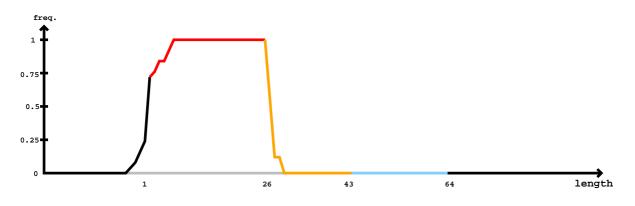| Provisional ID	: NW_025323209.1_Acropora_millepora_isolate_JS-1_unplaced_genomic_scaffold_Amil_v2.1_xpSc0000659_whole_genome_shotgun_sequ
Score total	: 2.7
Score for star read(s)	: -1.3
Score for read counts	: 0
Score for mfe	: 1
Score for randfold	: 0
Score for cons. seed	: 3
Total read count	: 23
Mature read count	: 23
Loop read count	: 0
Star read count	: 0

Mature   **Star**

```
5'- ucuucagcagcgaaaacacugggguuccuguagaucuccauguugguguccuucuugggcauuagauguuccugcaugugucuuguguuuaucugguaagacgcucuguaguuuca  -3'    exp
    .((.((((.((((.((((((((.(((..(((((((((.....(((.(((((((((((.....)))))))))))...)))))).))))).))))))).((((...))))))))))).))...   reads   mm    sample
    ...............cacuggguuccuguagaucuccauguugg.........................................................................   1       0     seq
    ................acuggguuccuguagaucuccauguugg........................................................................   1       0     seq
    .................cuggguuccuguagaucuccauguugA.......................................................................   1       1     seq
    .................cuggguuccuguagaucuccauguugg.......................................................................   1       0     seq
    .................ugggguuccuguagaucuccauguugg.......................................................................   2       0     seq
    ..................gggguuccuguagaucuccauguugg......................................................................   6       0     seq
    ..................gggguuccuguagaucuccauguugA......................................................................   3       1     seq
    ..................gggguuccuguagaucuccauguuggugu...................................................................   3       0     seq
    ...................ggguuccuguagaucuccauguugg.....................................................................   1       0     seq
    ....................gguuccuguagaucuccauguugg....................................................................   2       0     seq
    .....................guuccuguagaucuccauguugA...................................................................   2       1     seq
    ......................uccuguagaucuccauguugA..................................................................   1       1     seq
    .......................ccuguagaucuccauguugg.................................................................   1       0     seq
```