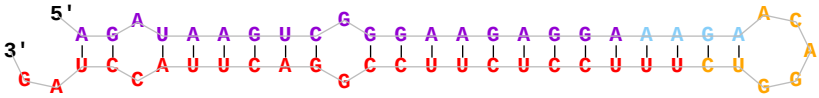
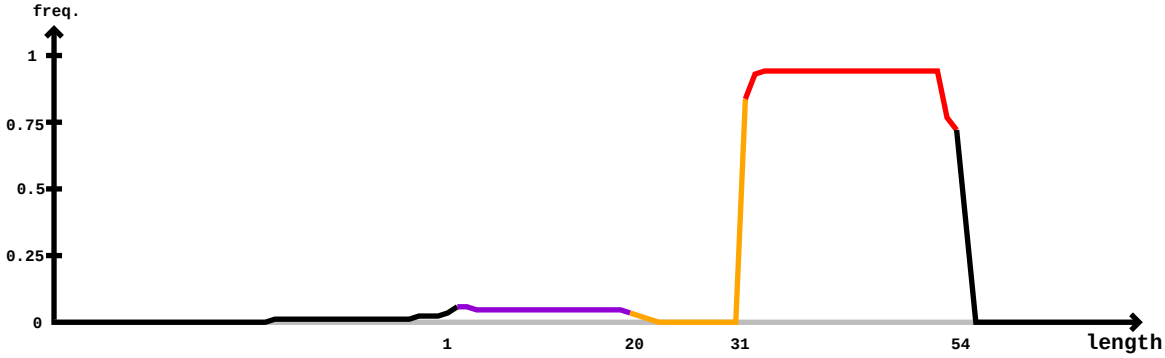
Provisional ID	:	NC_058072.1_Acropora_millepora_isolate_JS-1_chromosome_7_Amil_v2.1_whole_genome_shotgun_sequence_299173
Score total	:	46.5
Score for star read(s)	:	3.9
Score for read counts	:	36.8
Score for mfe	:	2.7
Score for randfold	:	0
Score for cons. seed	:	3
Total read count	:	84
Mature read count	:	81
Loop read count	:	0
Star read count	:	3





Star **Mature**

```
5'- uacaaccacuucucuaugagcagucucuuucgaucuuccuagauaagucgggaagaggaaagaacaggucuuuccucuuccggacuuaccuaggaagauugaaggagacug  -3'   obs
    uacaaccacuucucuaugagcagucucuuucgaucuuccuagauaagucgggaagaggaaagaacaggucuuuccucuuccggacuuaccuaggaagauugaaggagacug        exp
    ....................((((((((((((((((((((((.(((((((.(((((((((((((.....)))))))))))))).)))))))..)))))))))))))))))))))))  reads   mm   sample
    ....................agucucuuucgaucuuccuag.......................................................................  1       0    seq
    ....................................uccuagauaagucgggaagaggaaa...................................................  1       0    seq
    ....................................uagauaagucgggaagagA.........................................................  1       1    seq
    ....................................agauaagucgggaagagga.........................................................  2       0    seq
    ...................................................................uuuccucuuccggacuuaccC...................  1       1    seq
    ...................................................................uuuccucuuccggacuuaccG...................  3       1    seq
    ...................................................................uuuccucuuccggacuuaccu...................  11      0    seq
    ...................................................................uuuccucuuccggacuuaccuU..................  1       1    seq
    ...................................................................uuuccucuuccggacuuaccua..................  3       0    seq
    ...................................................................uuuccucuuccggacuuaccuaU.................  8       1    seq
    ...................................................................uuuccucuuccggacuuaccuag.................  17      0    seq
    ...................................................................uuuccucuuccggacuuaccuaA.................  5       1    seq
    ...................................................................uuuccucuuccggacuuaccuaC.................  9       1    seq
    ...................................................................uuuccucCuccggacuuaccuag.................  1       1    seq
    ...................................................................uuuccucuuccggacuuaccuagA................  1       1    seq
    ...................................................................uuuccucuuccggacuuaccuagU................  9       1    seq
    ...................................................................uuuccucuuccggacuuaccuagC................  2       1    seq
    ...................................................................uuuccucuuccggacuuaccuagg................  1       0    seq
    ..................................................................uuccCcuuccggacuuaccuag...................  1       1    seq
    ..................................................................uuccucuuccggacuuaccuagU..................  1       1    seq
    ..................................................................uuccucuuccggacuuaccuagg..................  5       0    seq
    ..................................................................uuccucuuccggacuuaccuagC..................  1       1    seq
    ..................................................................uccucuuccggacuuaccuagg...................  1       0    seq
```