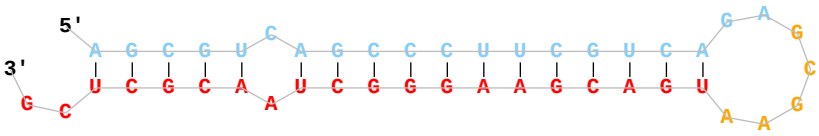
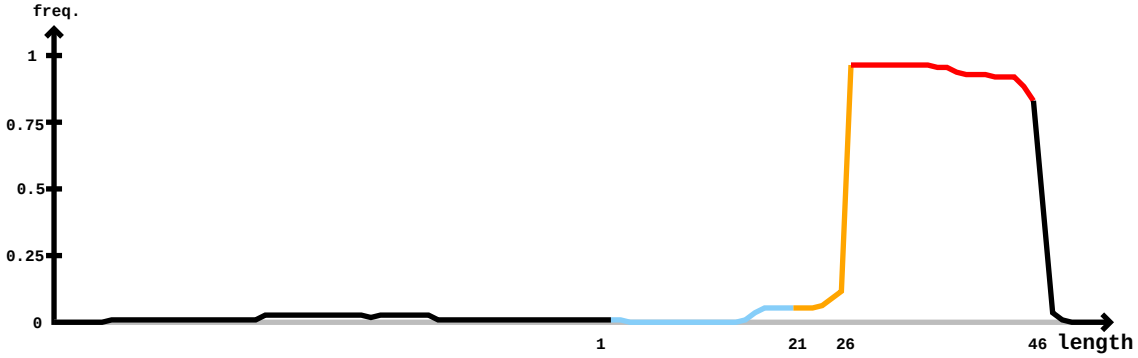
Provisional ID	:	NC_058067.1_Acropora_millepora_isolate_JS-1_chromosome_2_Amil_v2.1_whole_genome_shotgun_sequence_92488
Score total	:	0.2
Score for star read(s)	:	-1.3
Score for read counts	:	0
Score for mfe	:	2.1
Score for randfold	:	0
Score for cons. seed	:	-0.6
Total read count	:	101
Mature read count	:	101
Loop read count	:	0
Star read count	:	0





Star Mature

```
5'- acgaguugauaaagguggaauuaccaccgugaaagaacuggaaagcugacguuccgagcgucagcccuucgucagagcgaaugacgaagggcuaacgcucgaaauucg  -3'    exp
    .((((((......(((((.....)))))).........................(((((((.(((((((((((.......)))))))))))).)))))))..))))))   reads      mm         sample
    ....guugauaaagguAgaauuaccaccgug.......................................................................   1          1          seq
    ...................uuaccaccgugaaagaac...................................................................   2          0          seq
    ..............................aagaacuggaaagcugacguGccgag..............................................   1          1          seq
    ...................................................................gucagagcgaaGgacgaagggcuaacgcu.........   1          1          seq
    ..................................................................ucagagcgaaugacgaagA..................   1          1          seq
    ..................................................................ucagagcgaaugacgaagggc................   2          0          seq
    .................................................................cagagcAaaugacgaagggcu...............   1          1          seq
    .................................................................cagagcgaaGgacgaagggcuaac............   1          1          seq
    ...............................................................gaaugacgaagggcuaacgcucgU......   1          1          seq
    ...............................................................aaugacAaagggcuaacgcu.........   1          1          seq
    ...............................................................aaugacgaagggcuaacgcucg.......   1          0          seq
    ...............................................................aaGgacgaagggcuaacgcucgaaa....   1          1          seq
    ..............................................................aGgacgaagggcuaacgcucga......   1          1          seq
    ..............................................................aGgacgaagggcuaacgcucgaa.....   2          1          seq
    ..............................................................ugacgaagggcuaacgcu.........   1          0          seq
    ..............................................................ugacgaagggcuaacgcA.........   1          1          seq
    ..............................................................ugacgaagggcuaacgcuc........   5          0          seq
    ..............................................................ugacgaagggcuaacgcCc........   1          1          seq
    ..............................................................ugacgaagggcuGacgcucg.......   5          1          seq
    ..............................................................ugacgaGggcuaacgcucg.......   2          1          seq
    ..............................................................ugacgaagggcuaacgcCcg.......   1          1          seq
    ..............................................................ugacgaagggcuaacgcucg.......   68         0          seq
    ..............................................................ugacgaagggcuaacgcucgU......   8          1          seq
    ..............................................................ugacgaagggcuaacgcucga......   2          0          seq
    ..............................................................ugacgaagggcuaacgcucgaU.....   1          1          seq
```