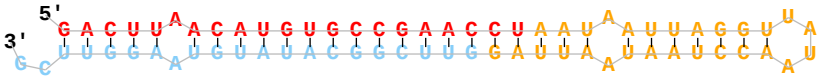
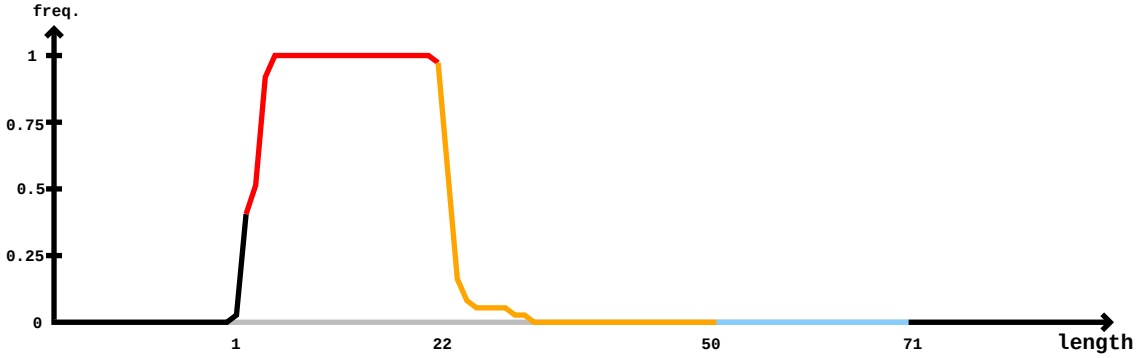
Provisional ID	:	NW_025323144.1_Acropora_millepora_isolate_JS-1_unplaced_genomic_scaffold_Amil_v2.1_xpSc0000587_whole_genome_shotgun_sequ
Score total	:	0.5
Score for star read(s)	:	-1.3
Score for read counts	:	0
Score for mfe	:	2.4
Score for randfold	:	0
Score for cons. seed	:	-0.6
Total read count	:	35
Mature read count	:	35
Loop read count	:	0
Star read count	:	0





Mature Star

```
5'- cugcggguucaaacguuggacuuaacaugugccgaaccuaauaauuagguuauaaccuaauaauuagguucggcauauguaagguucgacguuucaaucaauugggau  -3'    exp
    .....((((.(((((((((((((.((((((((((((((((((.(((((((....)))))))).))))))))))))))))))).)))))))))))))).)))).........   reads      mm         sample
    ................gAacuuaacaugugccgaaccu.....................................................................   1          1          seq
    ................Aacuuaacaugugccgaaccu......................................................................   10         1          seq
    ................Aacuuaacaugugccgaaccua.....................................................................   4          1          seq
    ..................Gcuuaacaugugccgaaccu.....................................................................   2          1          seq
    ..................acuuaacaugugccgaaccua....................................................................   1          0          seq
    ..................acuuaacaugugccgaaccuaUu..................................................................   1          1          seq
    ...................cuuaacaugugccgaacc......................................................................   1          0          seq
    ...................cuuaacaugugccgaaccG.....................................................................   1          1          seq
    ...................cuuaacaugugccgaaccu.....................................................................   4          0          seq
    ...................cuuaacaugugccgaaccua....................................................................   6          0          seq
    ...................cuuaacaugugccgaaccuaU...................................................................   3          1          seq
    ....................uuaacaugugccgaaccua....................................................................   1          0          seq
    ....................uuaacaugugccgaaccuaauaauu..............................................................   1          0          seq
    ....................uuaacaugugccgaaUcuaauaauuag............................................................   1          1          seq
```