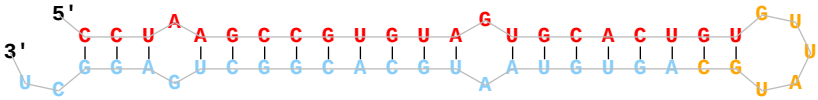
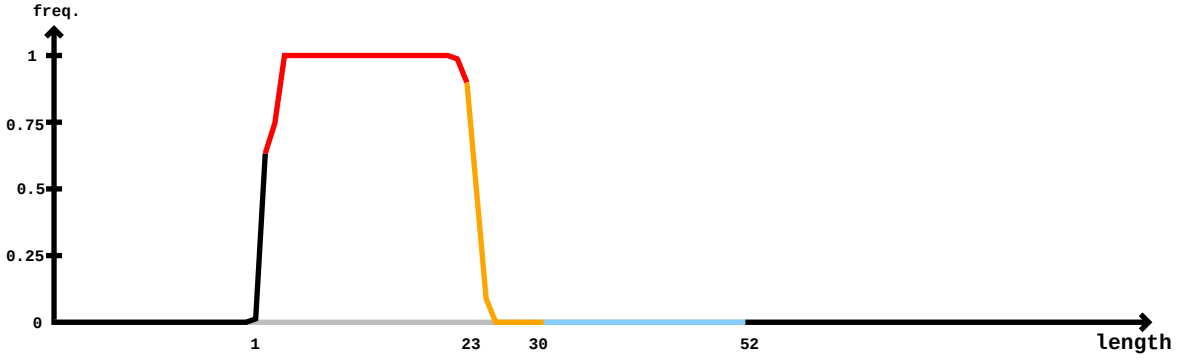
Provisional ID	:	NC_058077.1_Acropora_millepora_isolate_JS-1_chromosome_12_Amil_v2.1_whole_genome_shotgun_sequence_443230
Score total	:	0.2
Score for star read(s)	:	-1.3
Score for read counts	:	0
Score for mfe	:	2.1
Score for randfold	:	0
Score for cons. seed	:	-0.6
Total read count	:	79
Mature read count	:	79
Loop read count	:	0
Star read count	:	0





Mature Star

```
5'- aucauaaacuaaucaagaagccuaagccguguagugcacugugguuaugcaguguaaugcacggcugaggcuucuuuauuuguuaaauuauauugguaauuaaaaugguucuc  -3'    exp
    (((((.(((.(((.(((((((((.(((((((((.(((((((((.....)))))))).)))))))))).))))))))).))).))).((((((....)))))).....)))))....   reads      mm         sample
    ...................gccuaagccguguagugcacugu.....................................................................   1          0          seq
    ...................ccuaagccguguagugcacC.......................................................................   1          1          seq
    ...................ccuaagccguguagugcacuA......................................................................   2          1          seq
    ...................ccuaagccguguagugcacug......................................................................   5          0          seq
    ...................ccuaagccguguagugcacugC.....................................................................   2          1          seq
    ...................ccuaagccguguagugcacuAu.....................................................................   1          1          seq
    ...................ccuaagccguguagugcacugu.....................................................................   26         0          seq
    ...................ccuaagccguguagugcacugA.....................................................................   1          1          seq
    ...................ccuaagccguguagugcacuguC....................................................................   2          1          seq
    ...................ccuaagccguguagugcacuguU....................................................................   5          1          seq
    ...................ccuaagccguguagugcacuguA....................................................................   2          1          seq
    ...................ccuaagccguguagugcacuguUu...................................................................   2          1          seq
    ....................cuaagccguguagugcacugG.....................................................................   1          1          seq
    ....................cuaagccguguagugcacuguC....................................................................   1          1          seq
    ....................cuaagccguguagugcacuguU....................................................................   6          1          seq
    ....................cuaagccguguagugcacuguUu...................................................................   1          1          seq
    .....................uaagccguguagugcacugu.....................................................................   2          0          seq
    .....................uaagccguguagCgcacugu.....................................................................   1          1          seq
    .....................uaagccguguagugcacuguU....................................................................   6          1          seq
    .....................uaagccguguagugcacuguA....................................................................   1          1          seq
    .....................uaagccguguagugcacuguC....................................................................   6          1          seq
    .....................uaagccguguagugcacuguAu...................................................................   2          1          seq
    .....................uaagccguguagugcacuguUu...................................................................   2          1          seq
```