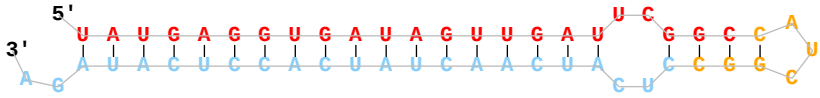
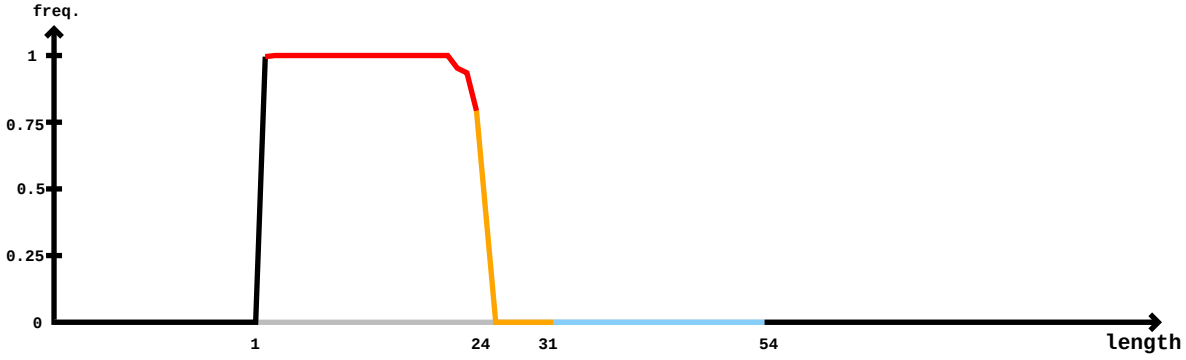
Provisional ID	:	NC_058067.1_Acropora_millepora_isolate_JS-1_chromosome_2_Amil_v2.1_whole_genome_shotgun_sequence_73358
Score total	:	0.2
Score for star read(s)	:	-1.3
Score for read counts	:	0
Score for mfe	:	2.1
Score for randfold	:	0
Score for cons. seed	:	-0.6
Total read count	:	231
Mature read count	:	231
Loop read count	:	0
Star read count	:	0





Mature Star

```
5'- ggguaguauggaggaaagucuaugaggugauaguugauucggccaucggccucaucaacuaucaccucauagaaaucucgaccucauaaucuaauuguuuaauagauauaucc  -3'    exp
    (((((.....((((....((((((((((((((((((((..((((...))))..))))))))))))))))))))).........))))...(((((.........)))))..)))))   reads     mm        sample
    ....................uaugaggugauaguugauuU.......................................................................   8         1         seq
    ....................uaugaggugauaguugauuc.......................................................................   3         0         seq
    ....................uaugaggugauaguugauucg......................................................................   2         0         seq
    ....................uaugaggugauaguugauucA......................................................................   1         1         seq
    ....................uaugaggugauaguugauuUg......................................................................   1         1         seq
    ....................uaugaggugauaguugauucgU.....................................................................   3         1         seq
    ....................uCugaggugauaguugauucgg.....................................................................   2         1         seq
    ....................uaugaggugauaguugauucgg.....................................................................   12        0         seq
    ....................uaugaggugauaguugauuUgg.....................................................................   13        1         seq
    ....................uaugaggugauaguugauucgA.....................................................................   2         1         seq
    ....................uaugaggugauaguugauucUg.....................................................................   1         1         seq
    ....................uaugaggugauaguugauucggU....................................................................   20        1         seq
    ....................uaugaggugauaguugauuUggc....................................................................   11        1         seq
    ....................uaugaggugauaguugauucgUc....................................................................   1         1         seq
    ....................uaugaggugauaguugauucggA....................................................................   8         1         seq
    ....................uaugaggugauaguugauucggc....................................................................   60        0         seq
    ....................uaugaggugauaguugGuucggc....................................................................   1         1         seq
    ....................uaugaggugauaguugauucgAc....................................................................   3         1         seq
    ....................uaugaggugauaguugauucggcG...................................................................   1         1         seq
    ....................uaugaggugauaguugauucggcA...................................................................   46        1         seq
    ....................uaugaggugauaguugauucggcU...................................................................   23        1         seq
    ....................uaugaggugauaguugauucggcc...................................................................   6         0         seq
    ....................uaugaggugauaguugauuUggcc...................................................................   2         1         seq
    ....................augaggugauaguugauucggU.....................................................................   1         1         seq
```