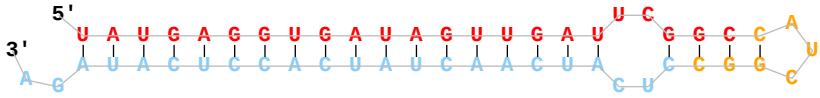
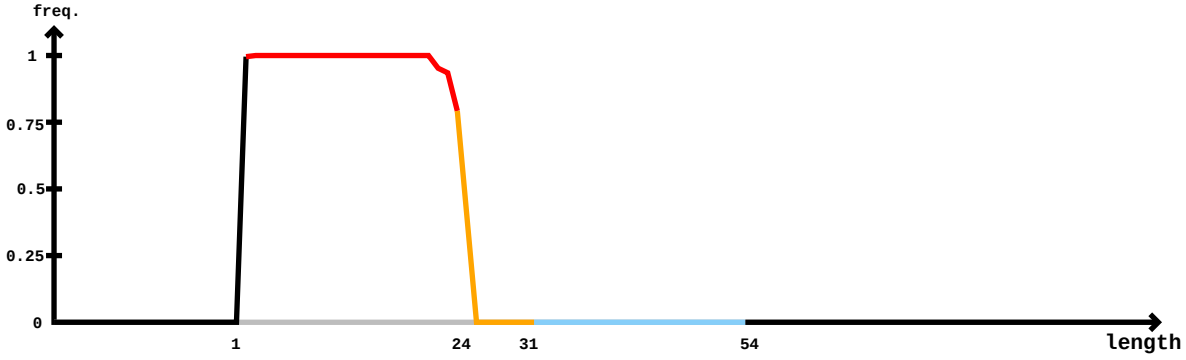
Provisional ID	:	NC_058067.1_Acropora_millepora_isolate_JS-1_chromosome_2_Amil_v2.1_whole_genome_shotgun_sequence_102453
Score total	:	0.1
Score for star read(s)	:	-1.3
Score for read counts	:	0
Score for mfe	:	2
Score for randfold	:	0
Score for cons. seed	:	-0.6
Total read count	:	231
Mature read count	:	231
Loop read count	:	0
Star read count	:	0





Mature Star

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