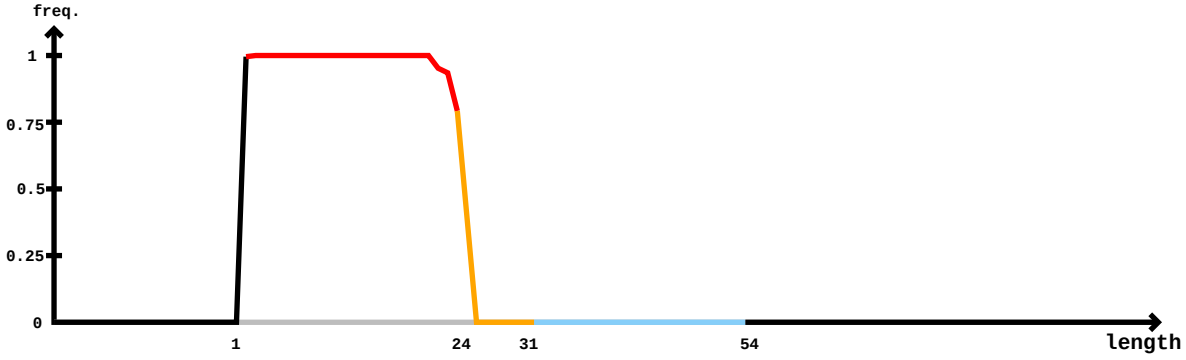
Provisional ID	:	NC_058067.1_Acropora_millepora_isolate_JS-1_chromosome_2_Amil_v2.1_whole_genome_shotgun_sequence_102453
Score total	:	0.1
Score for star read(s)	:	-1.3
Score for read counts	:	0
Score for mfe	:	2
Score for randfold	:	0
Score for cons. seed	:	-0.6
Total read count	:	231
Mature read count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



Mature Star

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