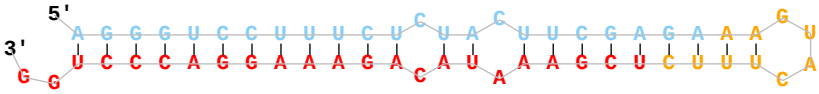
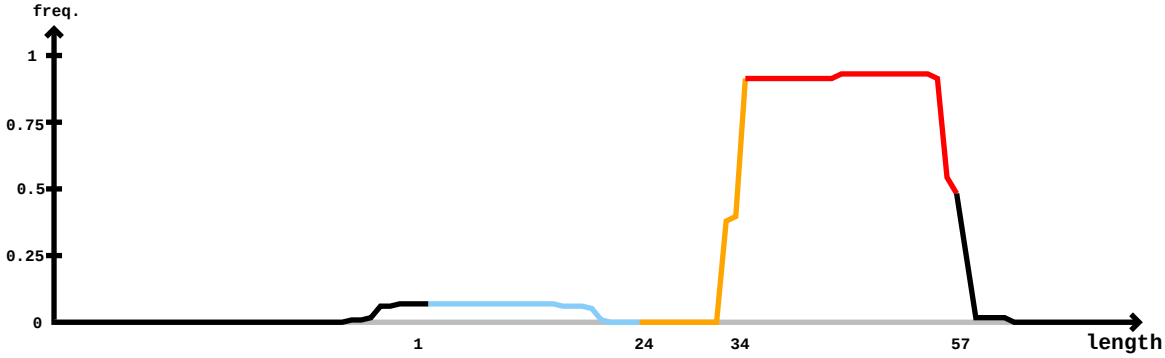
Provisional ID	:	NC_058076.1_Acropora_millepora_isolate_JS-1_chromosome_11_Amil_v2.1_whole_genome_shotgun_sequence_424051
Score total	:	4
Score for star read(s)	:	-1.3
Score for read counts	:	0
Score for mfe	:	2.3
Score for randfold	:	0
Score for cons. seed	:	3
Total read count	:	108
Mature read count	:	108
Loop read count	:	0
Star read count	:	0





Star Mature

```
5'- acucaaccgaagcacgcaaaaggacaaccucguucccaggguccuuucucuacuucgagaaaguacuuucucgaaauacagaaaggacccugggaacgaaguuagcaaaag  -3'    exp
    .......................(..(((.(((((((((((((((((((((.((.(((((((((....)))))))))).)).))))))))))))))))))).)))..).....   reads      mm         sample
    ............................ucguucccagggucccuuucucu.......................................................   1          0          seq
    .............................guucccagggucccuuucucuacC.......................................................   1          1          seq
    ............................uucccagggucccuuucucuacuu.......................................................   4          0          seq
    ............................uucccagggucccuuucucuacuuA......................................................   1          1          seq
    ...............................cccaggAuccuuucucuacuu.......................................................   1          1          seq
    ....................................................................ucucgaaauacagaaaggacAc...................   1          1          seq
    ....................................................................ucucgaaauacagaaaggacAcu..................   43         1          seq
    ...................................................................cucgaaauacagaaaggaccA....................   1          1          seq
    ...................................................................cucgaaauacagaaaggacccug..................   1          0          seq
    ..................................................................ucgaaauacagaaaggacccug...................   3          0          seq
    ..................................................................ucgaaauacagaaaggacccuU...................   2          1          seq
    ..................................................................ucgaaauacagaaaggacccuA...................   1          1          seq
    ..................................................................ucgaaauacagaaaggacccugC..................   8          1          seq
    ..................................................................ucgaaauacagaaaggacccugU..................   46         1          seq
    ...........................................................................gaaaggacccugUgaacg............   2          1          seq
```