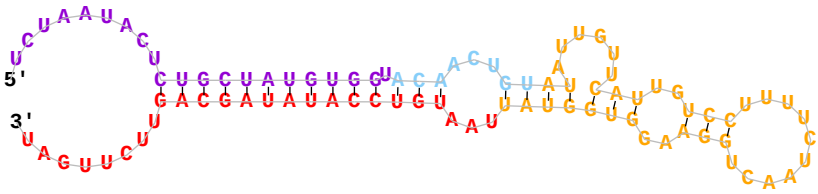
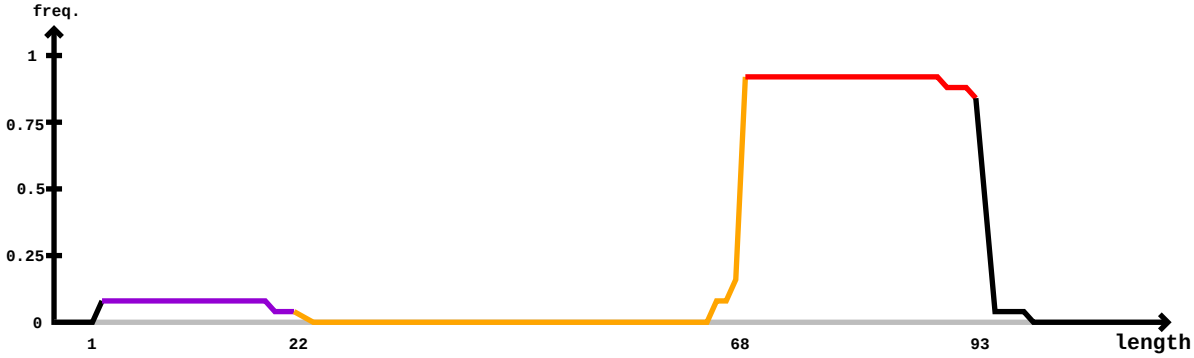
Provisional ID	:	NW_025322618.1_Acropora_millepora_isolate_JS-1_unplaced_genomic_scaffold_Amil_v2.1_Sc0000038_whole_genome_shotgun_sequen
Score total	:	12.4
Score for star read(s)	:	3.9
Score for read counts	:	5.7
Score for mfe	:	-0.1
Score for randfold	:	0
Score for cons. seed	:	3
Total read count	:	23
Mature read count	:	21
Loop read count	:	0
Star read count	:	2





Star **Mature**

```
5'- aagucuaauacucugcuaugugguacaacuguauauuguucauuguccuuuucuaacuggaagguggauuaauguccauauagcaguucuugauaagcauagaacgguugcag  -3'   obs
    aagucuaauacucugcuaugugguacaacuguauauuguucauuguccuuuucuaacuggaagguggauuaauguccauauagcaguucuugauaagcauagaacgguugcag        exp
    ..(((.......((((((((((.(((...((((......(((..(((..........)))..)))))))...))))))))))))))).....)))..(((.........)))..   reads   mm   sample
    ...ucuaauacucugcuaugu.......................................................................................   1       0    seq
    ...ucuaauacucugcuaugugCu....................................................................................   1       1    seq
    ..........................................................uauuaauguccauauagcaguucuugaC...................   1       1    seq
    ..........................................................uauuaauguccauauagcaguucuugaua..................   1       0    seq
    ...........................................................uuaauguccauauagcaguucuugau....................   1       0    seq
    ...........................................................uuaauguccauauagcaguucuugaua...................   1       0    seq
    ..........................................................uaauguUcauauagcaguucu.......................   1       1    seq
    ..........................................................uaauguccauauagcaguucuuga....................   1       0    seq
    ..........................................................uaauguccauauagcaguucuugaC...................   2       1    seq
    ..........................................................uaauguccauauagcaguucuugau...................   8       0    seq
    ..........................................................uaauguccauauagcaguucuugaua..................   6       0    seq
    ..........................................................uaauguccauauagGaguucuugauaagca..............   1       1    seq
```