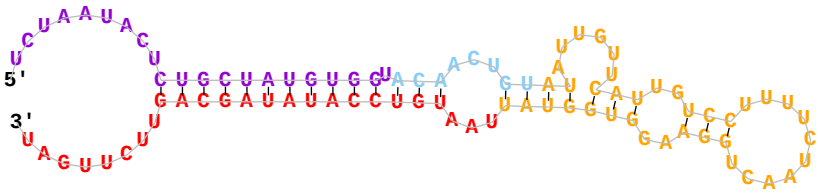
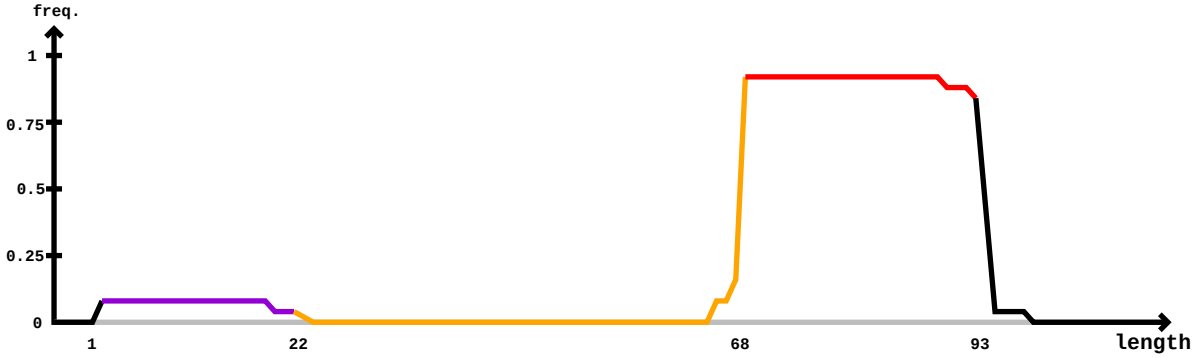
Provisional ID	:	NW_025322834.1_Acropora_millepora_isolate_JS-1_unplaced_genomic_scaffold_Amil_v2.1_xfSc0000181_whole_genome_shotgun_sequ
Score total	:	12.4
Score for star read(s)	:	3.9
Score for read counts	:	5.7
Score for mfe	:	-0.1
Score for randfold	:	0
Score for cons. seed	:	3
Total read count	:	23
Mature read count	:	21
Loop read count	:	0
Star read count	:	2





Star **Mature**

```
5'- aagucuaauacucugcuaugugguacaacuguauauuguucauuguccuuuucuaacuggaaggugguauuaauguccauauagcaguucuugauaagcauagaacgguugcag  -3'   obs
    aagucuaauacucugcuaugugguacaacuguauauuguucauuguccuuuucuaacuggaaggugguauuaauguccauauagcaguucuugauaagcauagaacgguugcag        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
```