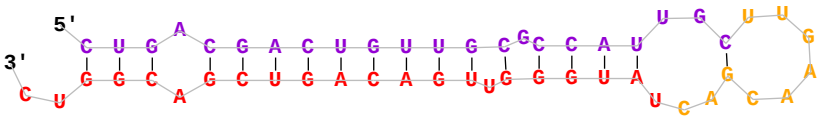
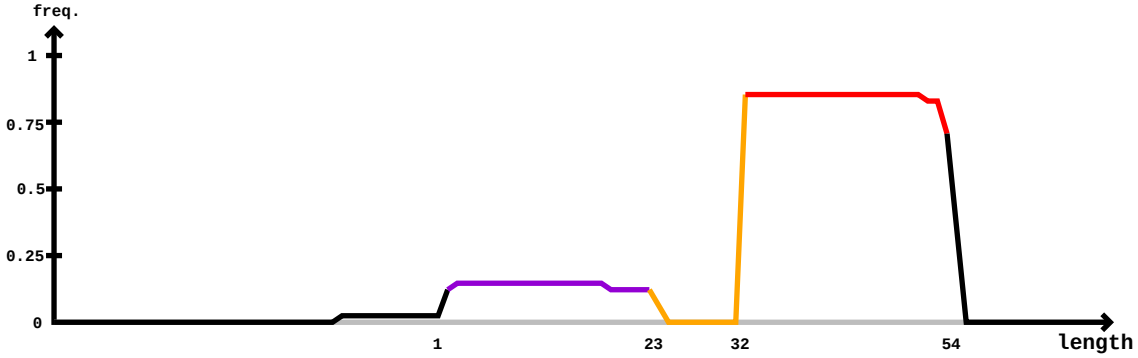
Provisional ID	:	NW_025322765.1_Acropora_millepora_isolate_JS-1_unplaced_genomic_scaffold_Amil_v2.1_xfSc0000041_whole_genome_shotgun_sequ
Score total	:	23.1
Score for star read(s)	:	3.9
Score for read counts	:	14.4
Score for mfe	:	1.8
Score for randfold	:	0
Score for cons. seed	:	3
Total read count	:	40
Mature read count	:	35
Loop read count	:	0
Star read count	:	5





Star Mature

```
5'- gaccuuguugcgccauuguuugaaugacugacugacugacugacgacguuugcgccauuugcuugaacgacuauggguugacagucgacggucagucggccgacaguug  -3'   obs
    gaccuuguugcgccauuguuugaaugacugacugacugacugacgacguuugcgccauuugcuugaacgacuauggguugacagucgacggucagucggccgacaguug        exp
    ....((((((.(((...(((.....)))).....(((((((((.((((((((.(((((((((.((((..(......)...)))))).)))))))))..))))))))))))))))...   reads   mm   sample
    ...........................ugacugacugacugacgacuguugcgcU.......................................................   1       1    seq
    ......................................cugacgacuguugcgccauugc.............................................   3       0    seq
    ......................................cugacgacuguugcgccauuguU............................................   1       1    seq
    .......................................ugacgacuguugcgccauuugcA...........................................   1       1    seq
    ....................................................................uauggguugacagucgacA..................   1       1    seq
    ....................................................................uauggguugacagucgacggu................   4       0    seq
    ....................................................................uauggguugacagucgacggG................   1       1    seq
    ....................................................................uauggguugacagucgacgguc...............   12      0    seq
    ....................................................................uauggguugacagucgacgguA...............   1       1    seq
    ....................................................................uauggguugacagucgacgguU...............   6       1    seq
    ....................................................................uauggguugacagucgacggucC..............   2       1    seq
    ....................................................................uauggguugacagucgacggucU..............   5       1    seq
    ....................................................................uauggguugacagucgacgguca..............   1       0    seq
    ....................................................................uauggguCgacagucgacgguca..............   1       1    seq
    ....................................................................uauggguugacagucgacggucG..............   1       1    seq
```