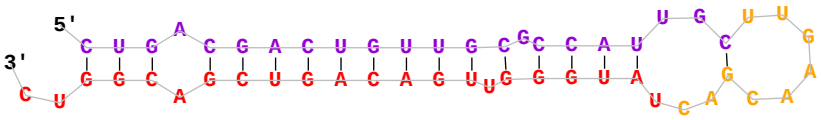
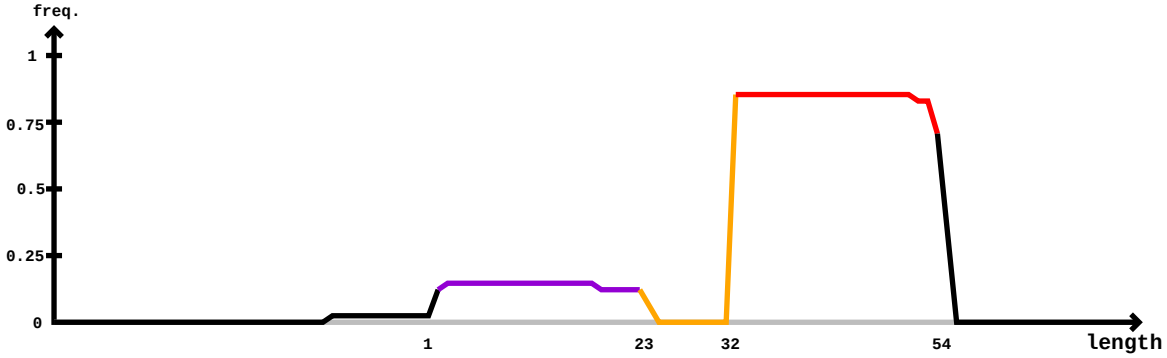
Provisional ID	:	NW_025322765.1_Acropora_millepora_isolate_JS-1_unplaced_genomic_scaffold_Amil_v2.1_xfSc0000041_whole_genome_shotgun_sequ
Score total	:	23.2
Score for star read(s)	:	3.9
Score for read counts	:	14.4
Score for mfe	:	1.9
Score for randfold	:	0
Score for cons. seed	:	3
Total read count	:	40
Mature read count	:	35
Loop read count	:	0
Star read count	:	5





Star Mature

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