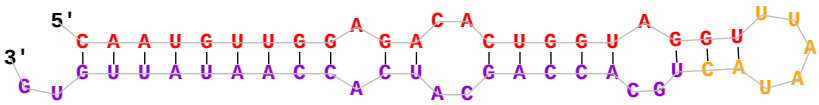
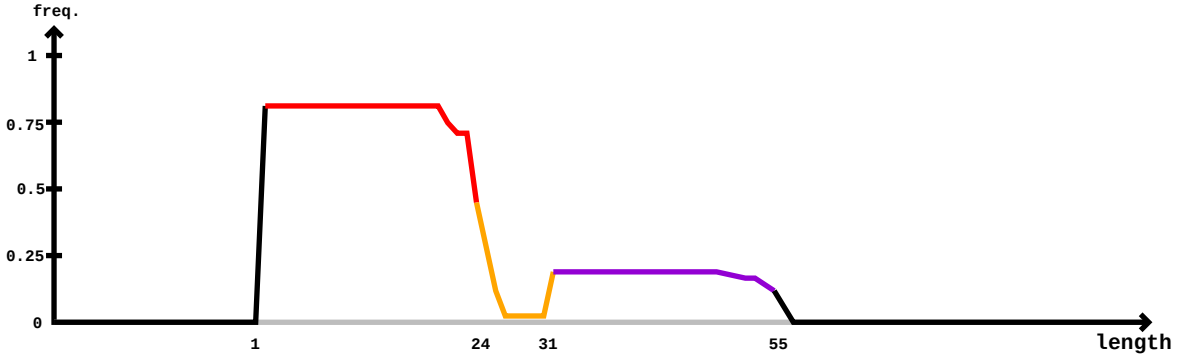
Provisional ID	:	NC_058067.1_Acropora_millepora_isolate_JS-1_chromosome_2_Amil_v2.1_whole_genome_shotgun_sequence_85610
Score total	:	65.4
Score for star read(s)	:	3.9
Score for read counts	:	57.2
Score for mfe	:	1.3
Score for randfold	:	0
Score for cons. seed	:	3
Total read count	:	124
Mature read count	:	103
Loop read count	:	0
Star read count	:	21






```
                    Mature                Star
5'- aguuucaguuuaugaugcugcaauguuggagacacugguaggguuuaauacugcaccagcaucaccaauauugugcugucauuaaacaugaaagaucuuuacugcugguuuug  -3'    obs
    aguuucaguuuaugaugcugcaauguuggagacacugguaggguuuaauacugcaccagcaucaccaauauugugcugucauuaaacaugaaagaucuuuacugcugguuuug         exp
    ..(((((((((((((((.((((((((((.((..(((((.(((.....)))..)))))..)).))))))))))))).))))).))))).))))).....................  reads     mm        sample
    ....................caauguuggagacacuggu.......................................................................  6         0         seq
    ....................caaCguuggagacacuggu.......................................................................  1         1         seq
    ....................caauguuggagacacuggC.......................................................................  1         1         seq
    ....................caauguuggagacacuggua......................................................................  4         0         seq
    ....................cGauguuggagacacuggua......................................................................  1         1         seq
    ....................caauguuggagacacugguagA....................................................................  4         1         seq
    ....................caaugCuggagacacugguagg....................................................................  1         1         seq
    ....................caauguuggagacacugguagg....................................................................  24        0         seq
    ....................caauguuggagacacugguagU....................................................................  3         1         seq
    ....................caauguuggagacacugguagC....................................................................  1         1         seq
    ....................caauguuggagacacugguaggA...................................................................  3         1         seq
    ....................caauguuggagacacugguaggC...................................................................  4         1         seq
    ....................caauguuggagacacugguaggu...................................................................  24        0         seq
    ....................caauguuggagacacugguagguA..................................................................  3         1         seq
    ....................caauguuggagacacugguagguu..................................................................  7         0         seq
    ....................caauguugAagacacugguagguu..................................................................  1         1         seq
    ....................caauguuggagacacugguagguG..................................................................  1         1         seq
    ....................caauguuggagacacugguagguC..................................................................  2         1         seq
    ....................caauguuggagacacugguaggguu.................................................................  11        0         seq
    ....................caauguuggagacacugguaggCuu.................................................................  1         1         seq
    ........................................uuaauacugcaccagcaucaccaauauugug.......................................  3         0         seq
    ...............................................ugcaccagcaucaccaau.............................................  1         0         seq
    ...............................................ugcaccagcaucaccaauU............................................  1         1         seq
    ...............................................ugcaccagcaucaccaauaC...........................................  1         1         seq
    ...............................................ugcaccagcaucaccaauauug.........................................  3         0         seq
    ...............................................ugcaccagcaucaccaauauugu........................................  3         0         seq
    ...............................................ugcaccagcaucaccaauauuguA.......................................  1         1         seq
    ...............................................ugcaccagcaucaccaauauugug.......................................  8         0         seq
    ...............................................ugcaccagcaucaccaauauugugU......................................  3         1         seq
```