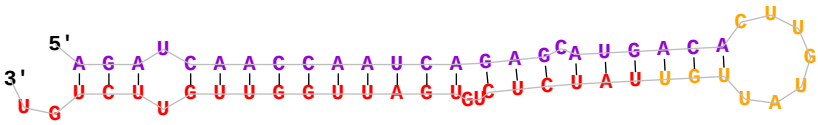
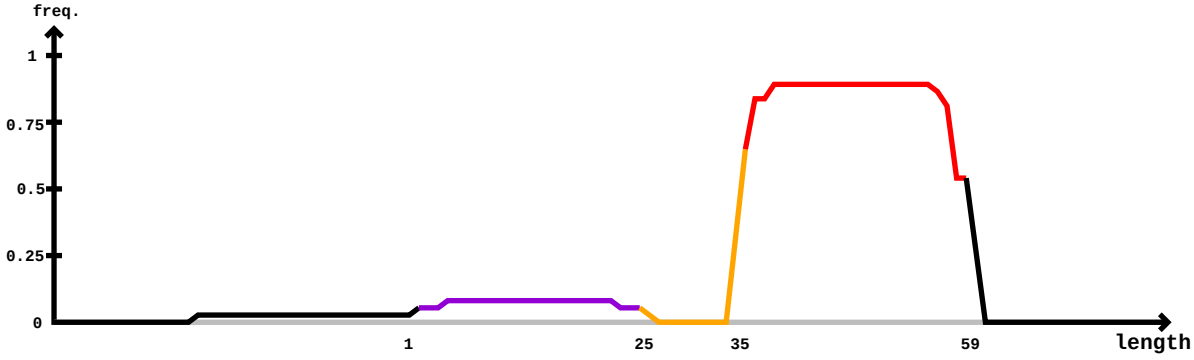
Provisional ID	:	NW_025322823.1_Acropora_millepora_isolate_JS-1_unplaced_genomic_scaffold_Amil_v2.1_xfSc0000167_whole_genome_shotgun_sequ
Score total	:	17.9
Score for star read(s)	:	3.9
Score for read counts	:	12.4
Score for mfe	:	2.3
Score for randfold	:	0
Score for cons. seed	:	-0.6
Total read count	:	36
Mature read count	:	33
Loop read count	:	0
Star read count	:	3





Star Mature

```
5'- cuuaugccaauuauuagguaauuugggcuaacacacagaucaaccaaucagagcaugacacuuguauuguuaucucugugauuggguuguucugugggguuggcccacuguaccuc  -3'   obs
    cuuaugccaauuauuagguaauuugggcuaacacacagaucaaccaaucagagcaugacacuuguauuguuaucucugugauuggguuguucugugggguuggcccacuguaccuc        exp
    ...............(((((...(((((((((.((((((.(((((((((((((.((((((.......)))))))))..)))))))))).)))))).)))))))))...)))))).   reads   mm   sample
    .............uuagguaauuugggcuaacacac.......................................................................................  1       0    seq
    ....................................agaucaaccaaucagagcauC.....................................................................  1       1    seq
    ....................................agaucaaccaaucagagcaugaca..................................................................  1       0    seq
    .....................................ucaaccaaucagagcaugacU....................................................................  1       1    seq
    ..........................................................uuaucucugugauuggguuguC.......................................  1       1    seq
    ..........................................................uuaucucugugauuggguuguucA.....................................  1       1    seq
    ..........................................................uuaucucugugauuggguuguucu.....................................  2       0    seq
    ..........................................................uuaucucugugauuggguuAuucugu...................................  2       1    seq
    ..........................................................uuaucucugugauuggguuguucugu...................................  5       0    seq
    ..........................................................uuaucucugugauuggguuguucuguU..................................  1       1    seq
    ..........................................................uaucucugugauuggguuguucu......................................  2       0    seq
    ..........................................................uaucucugugauuggguuguucugu....................................  5       0    seq
    ..........................................................uaucucugugauuggguuguucuguU...................................  2       1    seq
    ..........................................................uaucucugugauuggguuguucuguC...................................  2       1    seq
    ..........................................................uaucucugugauuggguuguucuguA...................................  1       1    seq
    ..........................................................aucucugugauuggguuguuc........................................  2       0    seq
    ..........................................................aucucugugauuggguuguucA......................................  1       1    seq
    ..........................................................aucucugugauuggguuguucu......................................  3       0    seq
    ..........................................................aucucugugauuggguuguucugC....................................  1       1    seq
    ........................................................cucugugauuggguuguucA.........................................  1       1    seq
    ........................................................cucugugauuggguuguucuguC.....................................  1       1    seq
```