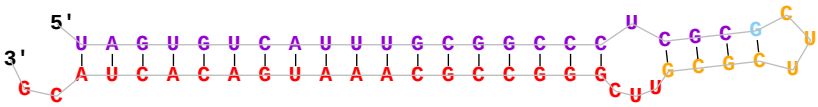
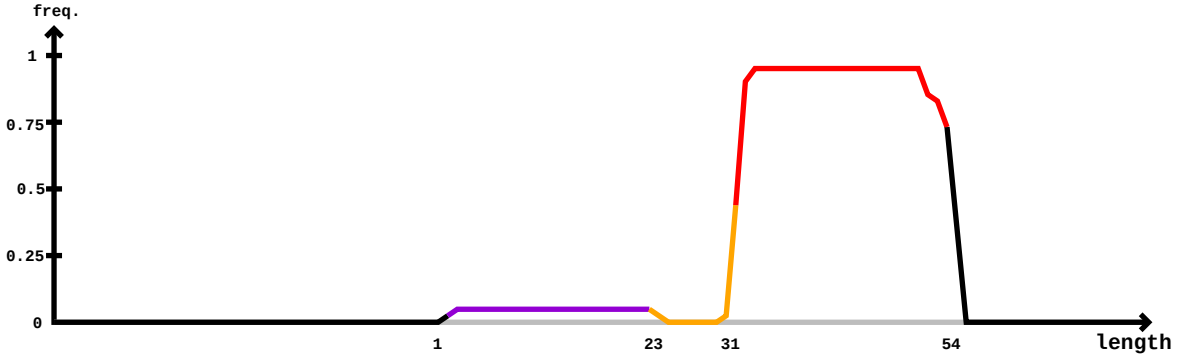
| | | |
|---|---|---|
| Provisional ID | : | NC_058068.1_Acropora_millepora_isolate_JS-1_chromosome_3_Amil_v2.1_whole_genome_shotgun_sequence_141124 |
| Score total | : | 24.5 |
| Score for star read(s) | : | 3.9 |
| Score for read counts | : | 14.9 |
| Score for mfe | : | 2.7 |
| Score for randfold | : | 0 |
| Score for cons. seed | : | 3 |
| Total read count | : | 41 |
| Mature read count | : | 39 |
| Loop read count | : | 0 |
| Star read count | : | 2 |




**Star** **Mature**

```
5'- uuguucgugacuccuccauugacaauauuugcgcucgcguagugucauuugcggcccucgcgcuucgcguucgggccgcaaaugacacuacgcgggcgcccaagcuauuaaa  -3'   obs
    uuguucgugacuccuccauugacaauauuugcgcucgcguagugucauuugcggcccucgcgcuucgcguucgggccgcaaaugacacuacgcgggcgcccaagcuauuaaa        exp
    (((((.(((.......))).)))))....(((((((((((((((((((((((((.((((...))))...)))))))))))))))))))))))))).............
    ......................................uagugucauuugcggcccucgU.............................................  reads   mm   sample
    .......................................agugucauuugcggcccucgc..............................................  1       1    seq
    ......................................................................guucgggccgcaaaugacacuacU...................  1       0    seq
    .......................................................................uucgggccgcaaaugacacC......................  1       1    seq
    .......................................................................uucgggccgcaaaugacacuac....................  1       1    seq
    .......................................................................uucgggccgcaaaugacacuac....................  1       0    seq
    .......................................................................uucgggcGgcaaaugacacuac....................  2       1    seq
    .......................................................................uucgggccgcaaaugacacuacA...................  2       1    seq
    .......................................................................uucgggccgcaaaugacacuacC...................  1       1    seq
    .......................................................................uucgggccgcaaaugacacuacU...................  10      1    seq
    .......................................................................ucgggccgcaaaugacacA......................  2       1    seq
    .......................................................................ucgggccgcaaaugacacu......................  1       0    seq
    .......................................................................ucgggccgcaaaugacacuU.....................  1       1    seq
    .......................................................................ucgggccgcaaaugacacuaU....................  1       1    seq
    .......................................................................ucgggccgcaaaugacacuacg...................  4       0    seq
    .......................................................................ucgggccgcaaaugacacuacU...................  10      1    seq
    ......................................................................cgggccgcaaaugacacuacg....................  1       0    seq
    ......................................................................cgggccgcaaaugacacuacC....................  1       1    seq
```