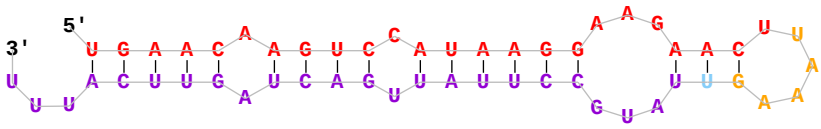
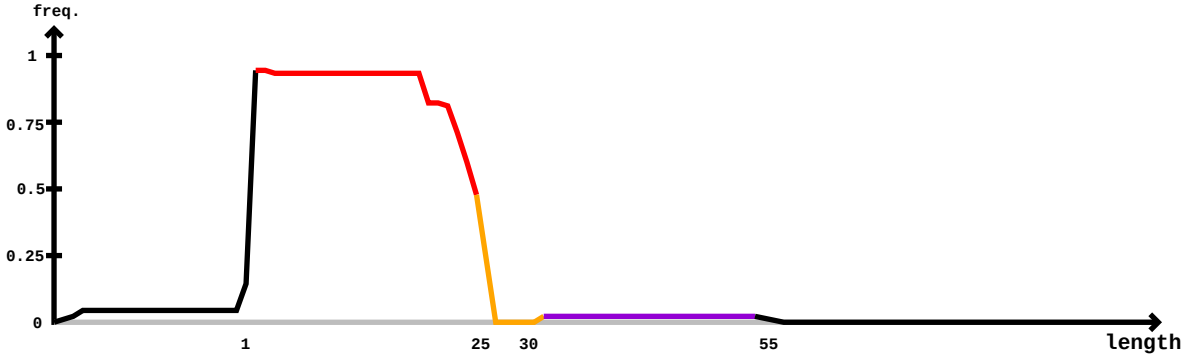
| | | |
|---|---|---|
| Provisional ID | : | NC_058067.1_Acropora_millepora_isolate_JS-1_chromosome_2_Amil_v2.1_whole_genome_shotgun_sequence_115437 |
| Score total | : | 46.6 |
| Score for star read(s) | : | 3.9 |
| Score for read counts | : | 37.9 |
| Score for mfe | : | 1.9 |
| Score for randfold | : | 0 |
| Score for cons. seed | : | 3 |
| Total read count | : | 86 |
| Mature read count | : | 84 |
| Loop read count | : | 0 |
| Star read count | : | 2 |




**Mature** **Star**

```
5'- gcgugguauaucugaaaaaugaacaaguccauaaggaagaacuuaaaguuaugccuuauugacuaguucauuuuucagauauauauacauauauuuugguguacucaaauuau  -3'   obs
    gcgugguauaucugaaaaaugaacaaguccauaaggaagaacuuaaaguuaugccuuauugacuaguucauuuucagauauauauacauauauuuugguguacucaaauuau        exp
    ..(((((((((((((((((((((.((((.((((((...(((.....)))...)))))).))))).))))))))))))))))))).))))..(((.(((((.....)))))).))))  reads   mm   sample
    gcgugguauaucugaaaa.......................................................................................  2       0    seq
    .cgugguauaucugaaaaa......................................................................................  1       0    seq
    .cgugguauaucugaaaaaug....................................................................................  1       0    seq
    ..................augaacaaguccauaaggUag..................................................................  1       1    seq
    ..................augaacaaguccauaaggUaga.................................................................  4       1    seq
    ..................augaacaaguccauaaggaagaacu..............................................................  1       0    seq
    ..................augaacaaguccauaaggUagaacu..............................................................  4       1    seq
    ..................augaacaaguccauaaggaagaacG..............................................................  1       1    seq
    ...................ugaacaaguccauaaggU.....................................................................  8       1    seq
    ...................ugaacaaguccauaaggC.....................................................................  2       1    seq
    ...................ugaacaaguccauaaggUaga..................................................................  5       1    seq
    ...................ugaacaaguccauaaggUagaa.................................................................  10      1    seq
    ...................ugaacaaguccauaaggUagaac................................................................  10      1    seq
    ...................ugaacaaguccauaaggaagaac................................................................  1       0    seq
    ...................ugaacaaguccauaaggUagaacu...............................................................  35      1    seq
    ...................ugaacaaguccauaaggaagaacC...............................................................  2       1    seq
    ...............................................uaugUcuuauugacuaguucauuu..................................  2       1    seq
```