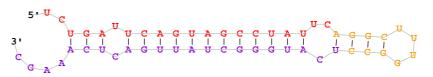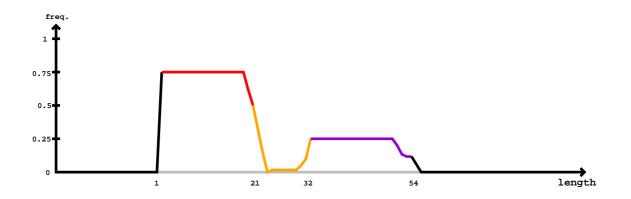| Provisional ID | : NC_058066.1_Acropora_millepora_isolate_JS-1_chromosome_1_Amil_v2.1_whole_genome_shotgun_sequence_5882 |
|---|---|
| Score total | : 3.4 |
| Score for star read(s) | : -1.3 |
| Score for read counts | : 0 |
| Score for mfe | : 1.7 |
| Score for randfold | : 0 |
| Score for cons. seed | : 3 |
| Total read count | : 54 |
| Mature read count | : 45 |
| Loop read count | : 0 |
| Star read count | : 9 |





```
                      Mature                    Star
5'- uaccucucacccucaugcccucugauucaguagccuauucaggcuuuggccucaugggcuauugacucaaagccauuuugggcucaaggaauaauuauugucaauua  -3'    obs
    uaccucucacccucaugcccucugauucaguagccuauucaggcuuuggccucaugggcuauugacucaaagccauuuugggcucaaggaauaauuauugucaauua          exp
    ..(((....(((..(((((.((.(((.((((((((((((..(((((....)))).)))))))))))).))).)).)))).))).))).................           reads    mm    sample
    ..................ucugauucaguagccuau............................................................................   6       0     seq
    ..................ucugauucaguagccuaC............................................................................   2       1     seq
    ..................ucugauucaguagccuauC...........................................................................   2       1     seq
    ..................ucugauucaguagccuauu...........................................................................   5       0     seq
    ..................ucugauucaguagccuauuc..........................................................................   17      0     seq
    ..................ucugauucaguagccuauucG.........................................................................   1       1     seq
    ..................ucugauucaguagccuauuca.........................................................................   3       0     seq
    ..................ucugauucaguagccuauucGg........................................................................   9       1     seq
    ...............................................cuuuggccucaugggcuauugacucaGagc....................................   1       1     seq
    ...............................................ccucaugggcuauugacucaa...........................................   2       0     seq
    ...............................................cucaugggcuauugacucaa............................................   2       0     seq
    ...............................................cucaugggcuauugacucaCa...........................................   1       1     seq
    .................................................ucaugggcuauugacuca............................................   1       0     seq
    .................................................ucaugCgcuauugacucaa...........................................   1       1     seq
    .................................................ucaugggcuauugacucGa...........................................   1       1     seq
    .................................................ucaugCgcuauugacucaaagc........................................   6       1     seq
```