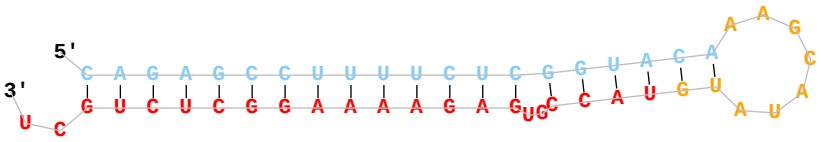
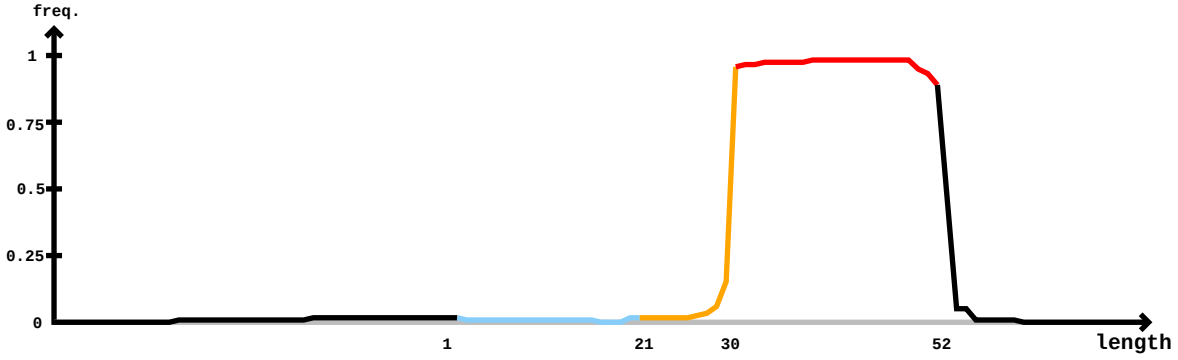
| | | |
|---|---|---|
| Provisional ID | : | NC_058077.1_Acropora_millepora_isolate_JS-1_chromosome_12_Amil_v2.1_whole_genome_shotgun_sequence_475648 |
| Score total | : | 0.2 |
| Score for star read(s) | : | -1.3 |
| Score for read counts | : | 0 |
| Score for mfe | : | 2.1 |
| Score for randfold | : | 0 |
| Score for cons. seed | : | -0.6 |
| Total read count | : | 111 |
| Mature read count | : | 111 |
| Loop read count | : | 0 |
| Star read count | : | 0 |




Star Mature

```
5'- cgcgcugaggaugcucaacaagaacuugacaggauucgugcagagccuuuucucgguacaaagcauaguuaccgugagaaaaggcucugcuagcaggguguuaauuacacug  -3'    exp
    ...(((..((((.((...(((....)))..)).)))).)..((((((((((((((((((((.......))))))..))))))))))))))).)))..(((((.....))))).   reads      mm         sample
    ...........ugcucaacaagaacuugacaggauucgugc.....................................................................   1          0          seq
    ........................uuCacaggauucgugcagagccuuuucucg.........................................................   1          1          seq
    ..............................................................caaagcauauguaccguAagaaaaggcucug.......................   2          1          seq
    ...............................................................uauguaccguAagaaaaggcucu.......................   1          1          seq
    ................................................................auguaccguAagaaaaggcucu.......................   1          1          seq
    ..............................................................uguaccguAagaaaaggcucugcu.....................   3          1          seq
    .............................................................guaccguAagaaaaggcucugc.......................   3          1          seq
    .............................................................guaccguAagaaaaggcucugcu......................   8          1          seq
    ..............................................................uaccguAagaaaaggcucu.......................   1          1          seq
    ..............................................................uaccguAagaaaaggcucugc.....................   1          1          seq
    ..............................................................uaccguAagaaaaggcucugcu....................   72         1          seq
    ..............................................................uaccgugagaaaaggcucugcA....................   1          1          seq
    ..............................................................uaccgugagaaaaggcucugcC....................   2          1          seq
    ..............................................................uaccgugagaaaaggcucugcu....................   4          0          seq
    ..............................................................uaccgugagaaaaggcucugcuC...................   1          1          seq
    ..............................................................uaccgugagaaaaggcucugcuU...................   4          1          seq
    ..............................................................uaccguAagaaaaggcucugcua....................   4          1          seq
    ..............................................................uaccgugagaaaaggcucugcuagA.................   1          1          seq
    ..............................................................uaccguAagaaaaggcucugcuagc.................   4          1          seq
    ...............................................................accguAagaaaaggcucu.......................   1          1          seq
    ................................................................cgugagaaaaggcucugc......................   1          0          seq
    ...........................................................................gaaaaggcuGugcuagcagggu............   1          1          seq
```