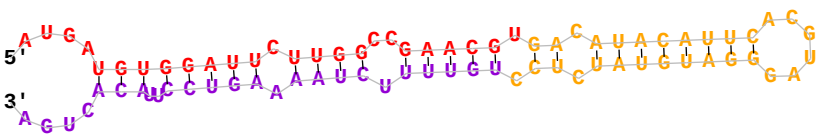
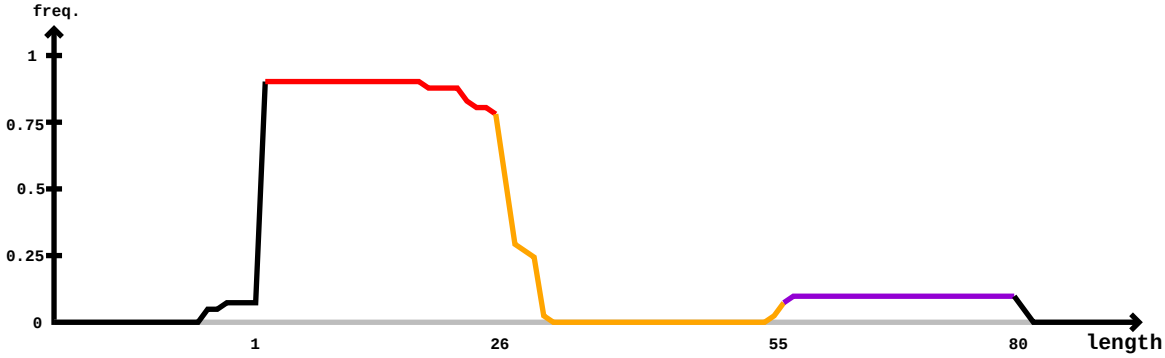
| | | |
|---|---|---|
| Provisional ID | : | NC_058079.1_Acropora_millepora_isolate_JS-1_chromosome_14_Amil_v2.1_whole_genome_shotgun_sequence_518700 |
| Score total | : | 16.8 |
| Score for star read(s) | : | 3.9 |
| Score for read counts | : | 13.4 |
| Score for mfe | : | 0.1 |
| Score for randfold | : | 0 |
| Score for cons. seed | : | -0.6 |
| Total read count | : | 38 |
| Mature read count | : | 34 |
| Loop read count | : | 0 |
| Star read count | : | 4 |




**Mature** **Star**

```
5'- uccgcgaaacaaguggacagaugauguggauucuuggccgaacgugacauacauucacguagggauguaucuccuguuuucuaaaaguccuuacacugaacaaguagccaa -3'   obs
    uccgcgaaacaaguggacagaugauguggauucuuggccgaacgugacauacauucacguagggauguaucuccuguuuucuaaaaguccuuacacugaacaaguagccaa        exp
    (((((.......)))))(((.((.((.(((((.((((..(((((.((.(((((((((.......)))))))).)).)))))).)))).)))))).)))))))............  reads   mm   sample
    ..............ggacagaugauguggauucuugg..............................................................................  1       0    seq
    ..............ggacagaugauguggauucuuggccgaa.........................................................................  1       0    seq
    ................acagaugauguggauucuuggccgaacgu.......................................................................  1       0    seq
    ....................augauguggauucuuggccga...........................................................................  2       0    seq
    ....................augauguggauucuuggccgaacA........................................................................  1       1    seq
    ....................augauguggauucuuggccgaacgA.......................................................................  1       1    seq
    ....................augauguggauucuuggccgaacCu.......................................................................  3       1    seq
    ....................augauguggauucuuggccgaacgC.......................................................................  2       1    seq
    ....................augauguggauucuuggccgaacgu.......................................................................  9       0    seq
    ....................augauguggauucuuggccgaacgug......................................................................  3       0    seq
    ....................augauguggauucuuggccgaacguA......................................................................  1       1    seq
    ....................augauguggauucuuggccgaacguga.....................................................................  1       0    seq
    ....................augauguggauucuuggccgaacgugac....................................................................  1       0    seq
    ....................augauguggauucuuggccgaacgugacG...................................................................  1       1    seq
    ....................augauguggauucuCggccgaacgugaca...................................................................  1       1    seq
    ....................augauguggauucuuggccgaacgugaca...................................................................  7       0    seq
    ....................augauguggauucuuggccgaacgugacau..................................................................  1       0    seq
    ...........................................................cuguuuucuaaaaguccuuacacugaa...........................  1       0    seq
    ..............................................................uguuuucuaaaaguccuuacacuga............................  2       0    seq
    ...........................................................guuuucuaaaaguccuuacacugaa...........................  1       0    seq
```