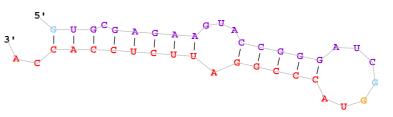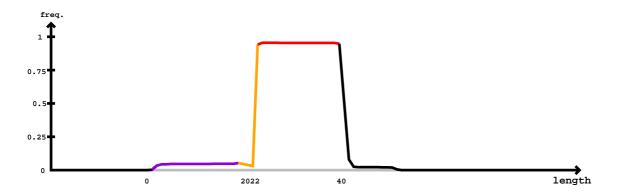| | | |
|---|---|---|
| Provisional ID | : | NW_025322653.1_Acropora_millepora_isolate_JS-1_unplaced_genomic_scaffold_Amil_v2.1_Sc0000146_whole_genome_shotgun_sequen |
| Score total | : | 349.7 |
| Score for star read(s) | : | 3.9 |
| Score for read counts | : | 344.8 |
| Score for mfe | : | 1.7 |
| Score for randfold | : | 0 |
| Score for cons. seed | : | -0.6 |
| Total read count | : | 688 |
| Mature read count | : | 656 |
| Loop read count | : | 0 |
| Star read count | : | 32 |





```
Star        Mature
5'- aauugcucuaacgguggguggugcgagaaguaccgggaucgguacccggauucuccaccaauuuuuguugagccuugcuugccuuagcgaaagguaccgagacucuaaa   -3'   obs
    aauugcucuaacgguggugugcgagaaguaccgggaucgguacccggauucuccaccaauuuuuguugagccuugcuugccuuagcgaaagguaccgagacucuaaa         exp
    ....(((((.(((((...(((((.(((((...(((((.....))))).)))))))))))....))))))))(((((..(((((((.....))))))).))))........   reads   mm       sample
.................gugcgagaaguaccgggauc.........................................................................    2       0        seq
................ugcgagaaguaccgggaC...........................................................................    2       1        seq
................ugcgagaaguaccgggau............................................................................    7       0        seq
................ugcgagaaguaccgggauU...........................................................................    1       1        seq
................ugcgagaaguaccCggauc...........................................................................    1       1        seq
................ugcgagaaguaccgggauA...........................................................................    2       1        seq
................ugcgagaaguaccgggauc...........................................................................    9       0        seq
.................gcgagGaguaccgggauc...........................................................................    1       1        seq
.................gcgagaaguaccgggauU...........................................................................    6       1        seq
..................gagaaguaccgggaucgg.........................................................................    1       0        seq
..................gagaaguaccgggaucgUuaccc....................................................................    1       1        seq
......................cgggaucgguacccggauucuccacca............................................................    1       0        seq
........................ucgguacccggauucuccacca..............................................................    4       0        seq
.......................cgguacccggauucuccacc.................................................................    1       0        seq
.......................cgguacccggauucuccacca................................................................    1       0        seq
.......................cgguacccggauucuccaccG................................................................    1       1        seq
........................gguacccggauucuccacca...............................................................    5       0        seq
........................gCuacccggauucuccacca...............................................................    2       1        seq
.........................guacccggauucuccacA...............................................................    2       1        seq
.........................guacccggauucuccacca..............................................................    2       0        seq
.........................Uuacccggauucuccaccaauuuuuguug....................................................    1       1        seq
..........................uacccggauucuccacc..............................................................    1       0        seq
..........................uacccggauucuccacAa.............................................................    2       1        seq
..........................Cacccggauucuccacca.............................................................    1       1        seq
..........................uacccAgauucuccacca.............................................................    1       1        seq
..........................uacccggauucCccacca.............................................................    2       1        seq
..........................uacccggauucuccaccU.............................................................    33      1        seq
..........................uacccggauucuccacca.............................................................    481     0        seq
..........................uacccggGuucuccacca.............................................................    1       1        seq
..........................uacccggauucuccaccC.............................................................    6       1        seq
..........................uacccggauCcuccacca.............................................................    2       1        seq
..........................uacUcggauucuccacca.............................................................    1       1        seq
..........................uacccggauucuccaccaU............................................................    64      1        seq
```

aauugcucuaacgguggugg**ugcgagaaguaccgggauc**gg**uacccggauucuccacca**auuuuuguugagccuugcuugccuuagcgaaagguaccgagacucuaaa

| | | | |
|---|---|---|---|
| ...........................................uacccggauucuccaccaa.................................................. | 2 | 0 | seq |
| ...........................................uacccggauucuccaccaUu................................................. | 35 | 1 | seq |
| ...........................................uacccggauucuccaccaaA................................................. | 1 | 1 | seq |
| ...........................................uacccggauucuccaccaau................................................. | 1 | 0 | seq |
| ...........................................uacccggauucuccaccaUuu................................................ | 1 | 1 | seq |
| ...........................................uacccggauucuccaUcaauuuuugu........................................... | 1 | 1 | seq |
| ...........................................uacccggauucuccaccaauuuuuguuga........................................ | 2 | 0 | seq |
| ...........................................uacccggauucuccaccaauuuuuguugG....................................... | 7 | 1 | seq |
| ...........................................uacccggauucuccaccGauuuuuguuga....................................... | 1 | 1 | seq |
| ...........................................uacccggauucuccaccaauuuuuguugGg..................................... | 4 | 1 | seq |
| ............................................acccggauucuccaccU.................................................... | 1 | 1 | seq |
| ............................................acccggauucuccacca................................................... | 6 | 0 | seq |
| ............................................acccggauucuccaccaUu................................................. | 2 | 1 | seq |
| ............................................acccggauucuccaccaUuu................................................ | 1 | 1 | seq |