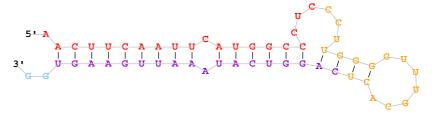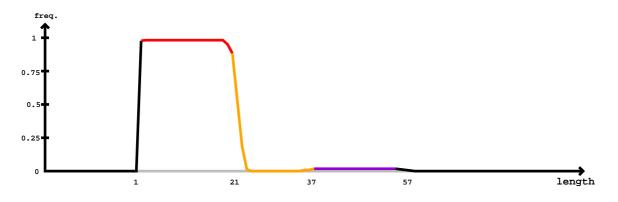| Provisional ID	: NC_058067.1_Acropora_millepora_isolate_JS-1_chromosome_2_Amil_v2.1_whole_genome_shotgun_sequence_99041
Score total	: 110.3
Score for star read(s)	: 3.9
Score for read counts	: 106.2
Score for mfe	: 0.8
Score for randfold	: 0
Score for cons. seed	: -0.6
Total read count	: 220
Mature read count	: 218
Loop read count	: 0
Star read count	: 2

Mature   **Star**

```
5'- acagcaccuuguuuucucaacuucaauucauggcccucccuuggggguuugcacucaggucauaaauugaagugggaaaagccucaucggugacugauaguacaacccauga  -3'   obs
    acagcaccuuguuuucucaacuucaauucauggcccucccuuggggguuugcacucaggucauaaauugaaguggaaaagccucaucggugacugauaguacaacccauga          exp
    .(((((((..(((((..((.((((((((.(((((......(((.......)))))))))).)))))))))..)))))......)))).)))                      reads   mm    sample
    .................aacuucaauucauggccc.......................................................................      1       0     seq
    .................aacuucaauucauggccU.......................................................................      2       1     seq
    .................aacuucaauucauggccA.......................................................................      4       1     seq
    .................aacuucaauucauggccAu......................................................................      1       1     seq
    .................aacuucaauucauggcccC......................................................................      1       1     seq
    .................aacuucaauucauggccUu......................................................................      1       1     seq
    .................aacuucaauucauggcccu......................................................................      12      0     seq
    .................aGcuucaauucauggcccuc.....................................................................      1       1     seq
    .................aacuucaauucauggcccuc.....................................................................      80      0     seq
    .................aacuucaauucauggcccuA.....................................................................      12      1     seq
    .................aacuucaauucauggcccGc.....................................................................      1       1     seq
    .................aacuucaauucauggccAuc.....................................................................      1       1     seq
    .................aacuucaauucauggUccuc.....................................................................      1       1     seq
    .................aacuucaauucauggcccCc.....................................................................      5       1     seq
    .................aacuucaauucauggcccuU.....................................................................      20      1     seq
    .................Gacuucaauucauggcccuc.....................................................................      2       1     seq
    .................aacuucaauAcauggcccuc.....................................................................      1       1     seq
    .................aacuucaauucauggcccucG....................................................................      2       1     seq
    .................aacuucaauucauggcccucc....................................................................      6       0     seq
    .................aacuucaauCcauggcccucc....................................................................      1       1     seq
    .................aacuucaauucauggcccuGc....................................................................      1       1     seq
    .................aacuucaauucauggcccucU....................................................................      20      1     seq
    .................aacuucaauucauggcccucAc...................................................................      38      1     seq
    .................aacuucaauucauggcccucAcu..................................................................      3       1     seq
    .................acuucaauucauggcccucGc....................................................................      1       1     seq
    .................................................cucaggCcauaaauugaagu.....................................      1       1     seq
    ..................................................cCcaggucauaaauugaaguggga.................................      1       1     seq
    .................................................caggucauaaauugaagu.......................................      2       0     seq
```