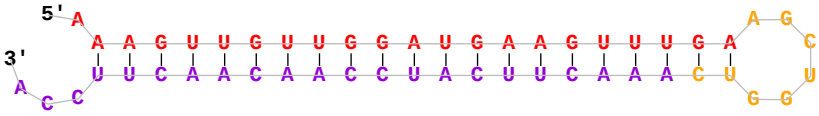
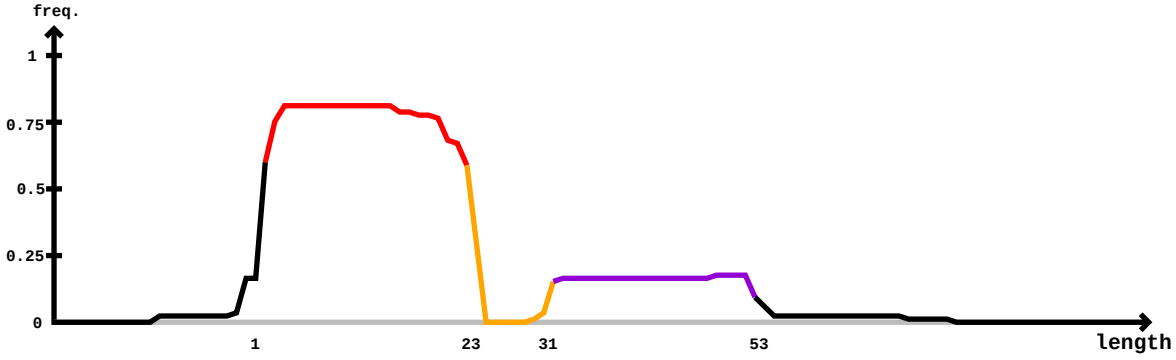
Provisional ID	:	NC_058067.1_Acropora_millepora_isolate_JS-1_chromosome_2_Amil_v2.1_whole_genome_shotgun_sequence_103691
Score total	:	44.3
Score for star read(s)	:	3.9
Score for read counts	:	34.8
Score for mfe	:	2.6
Score for randfold	:	0
Score for cons. seed	:	3
Total read count	:	80
Mature read count	:	66
Loop read count	:	0
Star read count	:	14





Mature Star

```
5'- cgcguuguugcgacuuguugaaaguuguuggaugaaguuugaagcugguccaaacuucauccaacaacuuccaacaagucgcaacaacacgcaacaucacacaacaugguuug -3'    obs
    cgcguuguugcgacuuguugaaaguuguuggaugaaguuugaagcugguccaaacuucauccaacaacuuccaacaagucgcaacaacacgcaacaucacacaacaugguuug        exp
    ((.(((((((((((((((.((((((((((((((((((.....)))))))))))))))))))).))))))))))))))))..))..................... reads     mm      sample
    .........gcgacuuguugGaaguuguuggaug.......................................................................... 2         1       seq
    ................uugGaaguuguuggaugaa.......................................................................... 1         1       seq
    .................ugGaaguuguuggaugaagu......................................................................... 1         1       seq
    .................ugGaaguuguuggaugaaguu........................................................................ 5         1       seq
    .................ugGaaguuguuggaugaaguuu....................................................................... 1         1       seq
    .................ugGaaguuguuggaugaaguuug...................................................................... 1         1       seq
    .................ugGaaguuguuggaugaaguuuga..................................................................... 3         1       seq
    ...................Gaaguuguuggaugaaguu........................................................................ 2         1       seq
    ...................Gaaguuguuggaugaaguuug...................................................................... 6         1       seq
    ...................Gaaguuguuggaugaaguuuga..................................................................... 28        1       seq
    ...................aaaguuguuggaugaaguuuga..................................................................... 1         0       seq
    ....................aGguuguuggaugaaguuuga..................................................................... 4         1       seq
    ....................aaguuguuggaugaaguuuga..................................................................... 9         0       seq
    .....................aguuguuggaugaaguuuga..................................................................... 5         0       seq
    ...............................................ucaaacuucauccaacaacuucU....................................... 1         1       seq
    ..............................................caaacuucauccaacaacuucU...................................... 2         1       seq
    ..............................................aaacuucauccaacaacuucc....................................... 3         0       seq
    ..............................................aaacuucauccaacaacuucA....................................... 2         1       seq
    ..............................................aaacuucauccaacaacuucca...................................... 4         0       seq
    ..............................................aaacuucauccaacaacuuccC...................................... 1         1       seq
    ...............................................aacuucauccaacaacuucca....................................... 1         0       seq
    ....................................................................uuccaacaagucgcaacaac....................... 1         0       seq
    ......................................................................aacaagucgcaacaacacgca.................... 1         0       seq
```