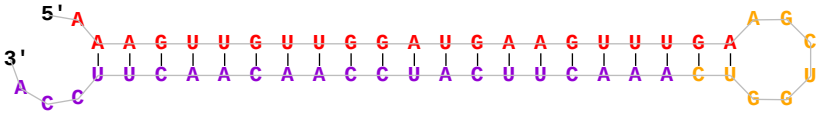
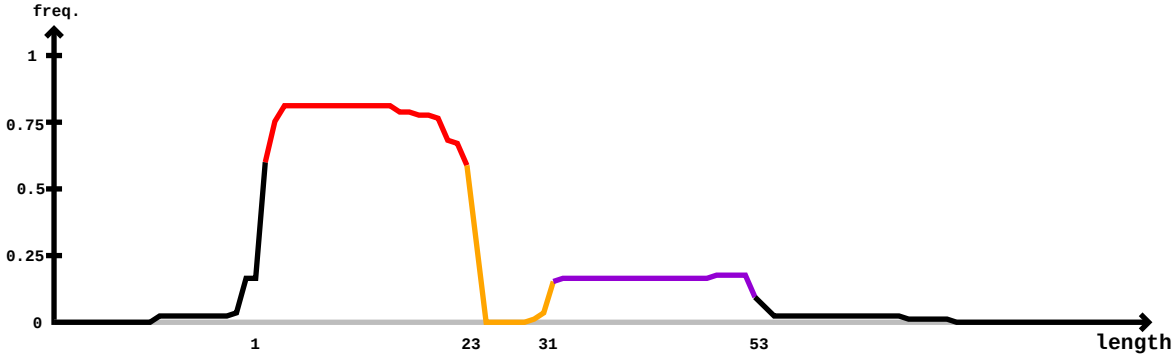
Provisional ID	:	NC_058067.1_Acropora_millepora_isolate_JS-1_chromosome_2_Amil_v2.1_whole_genome_shotgun_sequence_103691
Score total	:	40.7
Score for star read(s)	:	3.9
Score for read counts	:	34.8
Score for mfe	:	2.6
Score for randfold	:	0
Score for cons. seed	:	-0.6
Total read count	:	80
Mature read 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
    ...................aaaguuguuggaugaaguuuga................................................................  1         0         seq
    ...................Gaaguuguuggaugaaguuuga................................................................  28        1         seq
    ....................aaguuguuggaugaaguuuga................................................................  9         0         seq
    ....................aGguuguuggaugaaguuuga................................................................  4         1         seq
    .....................aguuguuggaugaaguuuga................................................................  5         0         seq
    ...............................................ucaaacuucauccaacaacuucU.....................................  1         1         seq
    ..............................................caaacuucauccaacaacuucU......................................  2         1         seq
    .............................................aaacuucauccaacaacuucA........................................  2         1         seq
    ............................................aaacuucauccaacaacuucc.........................................  3         0         seq
    ............................................aaacuucauccaacaacuuccC........................................  1         1         seq
    ............................................aaacuucauccaacaacuucca........................................  4         0         seq
    ...........................................aacuucauccaacaacuucca.........................................  1         0         seq
    ..................................................uuccaacaagucgcaacaac.....................................  1         0         seq
    ...............................................................aacaagucgcaacaacacgca......................  1         0         seq
```