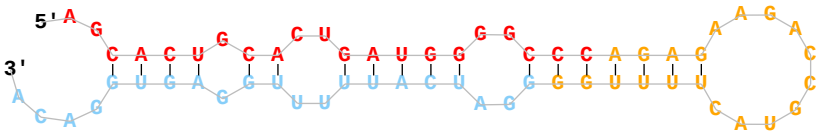
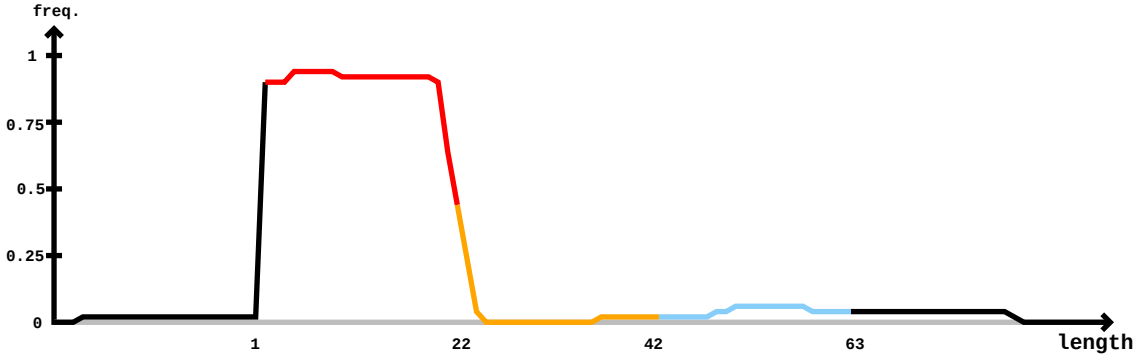
Provisional ID	:	NC_058066.1_Acropora_millepora_isolate_JS-1_chromosome_1_Amil_v2.1_whole_genome_shotgun_sequence_17842
Score total	:	0.7
Score for star read(s)	:	-1.3
Score for read counts	:	0
Score for mfe	:	-0.9
Score for randfold	:	0
Score for cons. seed	:	3
Total read count	:	46
Mature read count	:	46
Loop read count	:	0
Star read count	:	0





Mature Star

```
5'- uuaccuucauccuagcuaucagcacugcacugaugggcccagagaagaccguacuuuugggaucauuuuuggaguggacaaucuucuggaucauuuuuggagugga  -3'    exp
    .........(((..........((((.((..(((((..(((((((..........)))))))..)))))..)).)))).......(((..((.....))..))).)))   reads      mm         sample
    .uaccuucauccuagcuaucagcacugc.......................................................................................  1          0          seq
    ....................agcacugcacugaugggg.............................................................................  1          0          seq
    ....................agcacugcacugaugggc.............................................................................  6          0          seq
    ....................agcacugcacugauggguU............................................................................  5          1          seq
    ....................agcacugcacugUuggggc............................................................................  2          1          seq
    ....................agcacugcacugauggguU............................................................................  2          1          seq
    ....................agcacugcacugUuggggcc...........................................................................  2          1          seq
    ....................agcacugcacugauggguA............................................................................  1          1          seq
    ....................agcacugcacugauggggcc...........................................................................  3          0          seq
    ....................agcaAugcacugauggggcc...........................................................................  2          1          seq
    ....................agcacugcacugauggguU............................................................................  7          1          seq
    ....................agcacugcacugauggguA............................................................................  1          1          seq
    ....................agcaAugcacugauggggccc..........................................................................  1          1          seq
    ....................agcacugcacugUuggggccc..........................................................................  2          1          seq
    ....................agcacugcacugauggggcccU.........................................................................  5          1          seq
    ....................agcacugcacugauggggccca.........................................................................  1          0          seq
    ....................agcacugcacugauggggcccC.........................................................................  2          1          seq
    ....................agcaAugcacugauggggcccag........................................................................  1          1          seq
    .......................aAugcacugauggggccc...........................................................................  1          1          seq
    .......................acugcacuAauggggcccag........................................................................  1          1          seq
    ..........................................................uuuuggggaucauuuuuggagu...............................  1          0          seq
    ............................................................uuuuuggaguggacaaucuucuggaucauuu.......................  1          0          seq
    ..............................................................uuuggaguggacaaucuucuggaucauuuu......................  1          0          seq
```