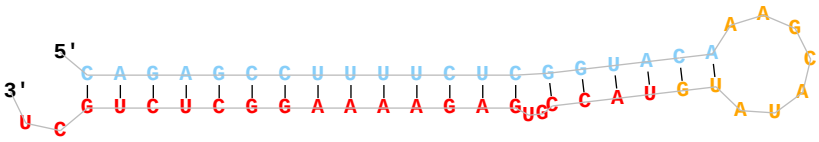
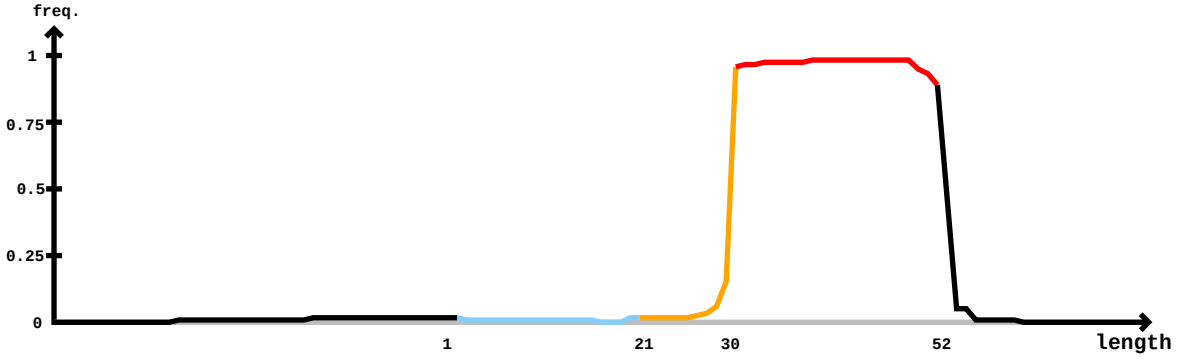
Provisional ID	:	NC_058071.1_Acropora_millepora_isolate_JS-1_chromosome_6_Amil_v2.1_whole_genome_shotgun_sequence_266806
Score total	:	0.2
Score for star read(s)	:	-1.3
Score for read counts	:	0
Score for mfe	:	2.1
Score for randfold	:	0
Score for cons. seed	:	-0.6
Total read count	:	111
Mature read count	:	111
Loop read count	:	0
Star read count	:	0





Star Mature

```
5'- cgcgcugaggaugcucaacaagaacuugacaggauucgugcagagccuuuucucgguacaaagcauauguaccgugagaaaaggcucugcuagcaggguggccagaacgaac  -3'    exp
    (((.(((.((((.((...(((....)))..)).)))).)).((((((((((((((((((((.......)))))))..)))))))))))))))...))).))).........  reads    mm    sample
    ..........ugcucaacaagaacuugacaggauucgugc.....................................................................  1        0     seq
    ......................uuCacaggauucgugcagagccuuuucucg.........................................................  1        1     seq
    ...........................................................caaagcauauguaccguAagaaaaggcucug.....................  2        1     seq
    ..............................................................uauguaccguAagaaaaggcucu.......................  1        1     seq
    .............................................................auguaccguAagaaaaggcucu........................  1        1     seq
    ............................................................uguaccguAagaaaaggcucugcu......................  3        1     seq
    ...........................................................guaccguAagaaaaggcucugc........................  3        1     seq
    ...........................................................guaccguAagaaaaggcucugcu.......................  8        1     seq
    ............................................................uaccguAagaaaaggcucu...........................  1        1     seq
    ............................................................uaccguAagaaaaggcucugc.........................  1        1     seq
    ............................................................uaccguAagaaaaggcucugcu........................  72       1     seq
    ............................................................uaccgugagaaaaggcucugcC........................  2        1     seq
    ............................................................uaccgugagaaaaggcucugcA........................  1        1     seq
    ............................................................uaccgugagaaaaggcucugcu........................  4        0     seq
    ............................................................uaccgugagaaaaggcucugcuC.......................  1        1     seq
    ............................................................uaccgugagaaaaggcucugcuU.......................  4        1     seq
    ............................................................uaccguAagaaaaggcucugcua.......................  4        1     seq
    ............................................................uaccguAagaaaaggcucugcuagc.....................  4        1     seq
    ............................................................uaccgugagaaaaggcucugcuagA.....................  1        1     seq
    ...........................................................accguAagaaaaggcucu............................  1        1     seq
    ..........................................................cgugagaaaaggcucugc.............................  1        0     seq
    ..........................................................gaaaaggcuGugcuagcagggu.........................  1        1     seq
```