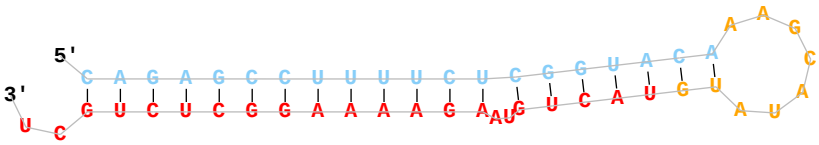
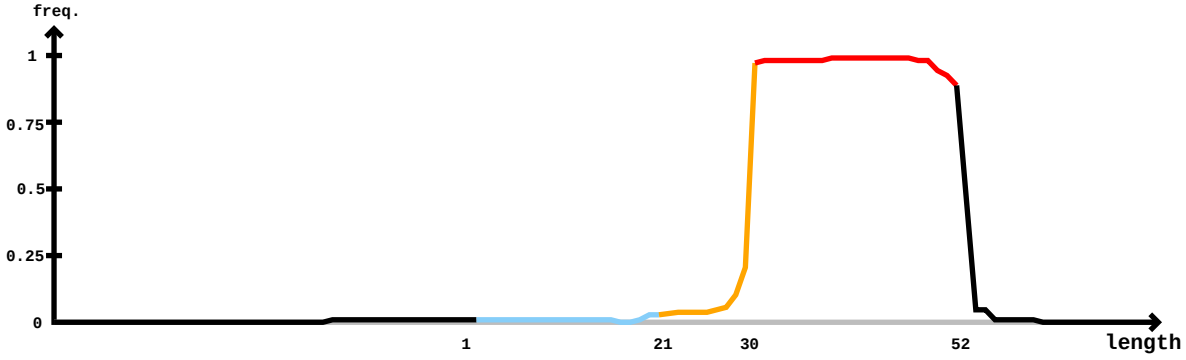
Provisional ID	:	NC_058074.1_Acropora_millepora_isolate_JS-1_chromosome_9_Amil_v2.1_whole_genome_shotgun_sequence_374842
Score total	:	0
Score for star read(s)	:	-1.3
Score for read counts	:	0
Score for mfe	:	1.9
Score for randfold	:	0
Score for cons. seed	:	-0.6
Total read count	:	99
Mature read count	:	99
Loop read count	:	0
Star read count	:	0





Star Mature

```
5'- aaugugcuaagcaugcucaacaagaacuuuacaggauucgugcagagccuuuucucgguacaaagcauauguacuguaagaaaaggcucugcuagcagggugacuggcuuguu  -3'    exp
    .....((((..(((.(((.....(((((.....)).)))..((((((((((((((((((((.......))))))).)))))))))))))..).)).)))..))))..... reads     mm        sample
    ........................uuCacaggauucgugcagagccuuuucucg.......................................................... 1         1         seq
    ..........................................................acaaagcauauguacuguaagaaaaggcC......................... 1         1         seq
    .........................................................caaagcauauguacCguaagaaaaggcucug...................... 2         1         seq
    ........................................................aagcauauguacuguaagaaaaggcucugcC.................... 1         1         seq
    ......................................................uauguacCguaagaaaaggcucu....................... 1         1         seq
    .....................................................auguacCguaagaaaaggcucu....................... 1         1         seq
    ....................................................uguacCguaagaaaaggcucugcu.................... 3         1         seq
    ....................................................uguacuguaagaaaaggcucugcu.................... 1         0         seq
    ....................................................uguGcuguaagaaaaggcucugcu.................... 1         1         seq
    ...................................................guacCguaagaaaaggcucugc..................... 3         1         seq
    ...................................................guacCguaagaaaaggcucugcu.................... 8         1         seq
    ..................................................uacCguaagaaaaggcucu....................... 1         1         seq
    ..................................................uacCguaagaaaaggcucugc..................... 1         1         seq
    ..................................................uacCguaagaaaaggcucugcu.................... 72        1         seq
    ..................................................uacCguaagaaaaggcucugcua................... 4         1         seq
    ..................................................uacCguaagaaaaggcucugcuagc................ 4         1         seq
    ...................................................acCguaagaaaaggcucu....................... 1         1         seq
    ...........................................................gaaaaggcuGugcuagcagggu........... 1         1         seq
```