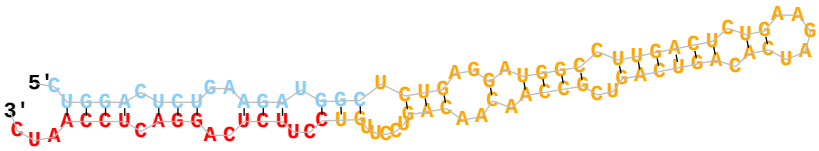
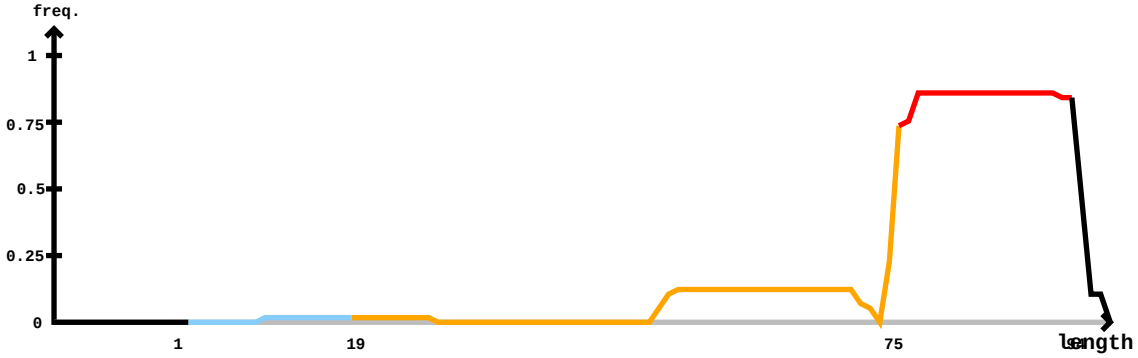
Provisional ID	:	NC_058067.1_Acropora_millepora_isolate_JS-1_chromosome_2_Amil_v2.1_whole_genome_shotgun_sequence_121417
Score total	:	2.9
Score for star read(s)	:	-1.3
Score for read counts	:	0
Score for mfe	:	1.2
Score for randfold	:	0
Score for cons. seed	:	3
Total read count	:	56
Mature read count	:	49
Loop read count	:	7
Star read count	:	0





Star Mature

```
5'- agucaagugagucuggacucugaagauggcucugaggauggccuugacucugaagaucacagucagucgccaacaacaguccuuguccuucucaggacuccaaucaca  -3'    exp
    ......((((...((((.(((..(((.(((.(((..(.((((.((((((.((.....)).)))))).))))).)..)))).....)))..)))..)))).)))).))))).   reads      mm         sample
    ....................ugaagauggcucugagga.......................................................................   1          0          seq
    ..........................................................gucagucgccaacaacaguccuu........................   3          0          seq
    ..........................................................ucagucgccaacaacagucc...........................   2          0          seq
    ..........................................................ucagucgccaacaacaguccA..........................   1          1          seq
    ...........................................................cagucgccaacaacagucc...........................   1          0          seq
    ...............................................................................uccuucucaggacuccaa......   1          0          seq
    ...............................................................................uUcuucucaggacuccaauc....   1          1          seq
    ...............................................................................uccuucucaggacuccaauc....   9          0          seq
    ...............................................................................uccuucucagCacuccaauc....   1          1          seq
    ...............................................................................uccuucucaggacuccaauca...   1          0          seq
    ...............................................................................ccuucuUaggacuccaauc....   1          1          seq
    ...............................................................................cAuucucaggacuccaauc....   2          1          seq
    ...............................................................................ccuucucaggacuccaauA....   3          1          seq
    ...............................................................................ccuucucaggacuccaauc....   23         0          seq
    ................................................................................cuucucaggacuccaauc....   1          0          seq
    ................................................................................uucucaggacuccUaucaca   2          1          seq
    ................................................................................uucucGggacuccaaucaca   1          1          seq
    ................................................................................uucucaggacuccaaucaca   3          0          seq
```