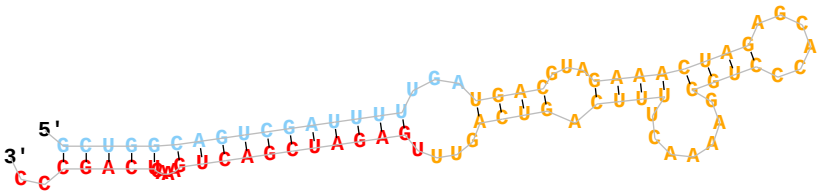
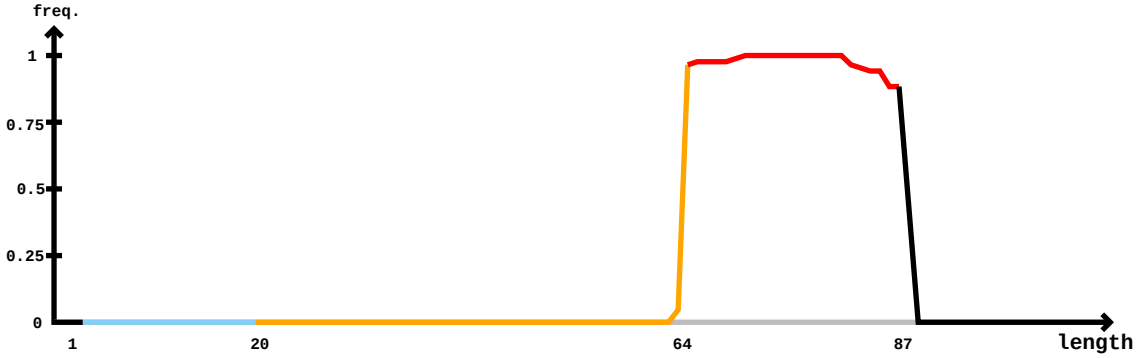
Provisional ID	:	NC_058077.1_Acropora_millepora_isolate_JS-1_chromosome_12_Amil_v2.1_whole_genome_shotgun_sequence_455813
Score total	:	2.3
Score for star read(s)	:	-1.3
Score for read counts	:	0
Score for mfe	:	0.6
Score for randfold	:	0
Score for cons. seed	:	3
Total read count	:	86
Mature read count	:	86
Loop read count	:	0
Star read count	:	0





Star Mature

```
5'- cgcuggcagucgauuuuugaugacguagaaacuagagcacccugggaaaacuuuucagucaguuugagaucgacugaaacucagcccagguccggacaaagucugua  -3'    exp
    .(((((((((((((((...((((...((((((((......))))).......)))).))))....)))))))))))....)))))........(((((...)))))..   reads     mm        sample
    ...........................................................uugagaucgacugaaacu.........................   2         0         seq
    ...........................................................uugagaucgacugaaacuc........................   1         0         seq
    ...........................................................uugagaucgacugaaacucagU.....................   1         1         seq
    ...........................................................ugagaucgacugaaacu..........................   1         0         seq
    ...........................................................ugagaucgacugaaacuca........................   1         0         seq
    ...........................................................ugagaucgacugaaacucagc......................   3         0         seq
    ...........................................................ugagaucgacugaaacucagU......................   1         1         seq
    ...........................................................ugagaucgacugaaacucagccA....................   20        1         seq
    ...........................................................ugagGucgacugaaacucagccc....................   1         1         seq
    ...........................................................ugagaucgacugaaacucagccU....................   22        1         seq
    ...........................................................ugagaucgacugaaacucagcUc....................   2         1         seq
    ...........................................................ugagaucgacugaaacucagccc....................   25        0         seq
    ...........................................................ugagaucgacugaaacucagccG....................   1         1         seq
    ..........................................................Agagaucgacugaaacucagccc....................   1         1         seq
    ...........................................................ugagaucgacugaaacucagcccc...................   1         0         seq
    ..........................................................gagaucgacugaaacucagccc.....................   1         0         seq
    .........................................................ucgacugaaacucagccc.........................   1         0         seq
    ........................................................cgacugaaacucagcccc.........................   1         0         seq
```