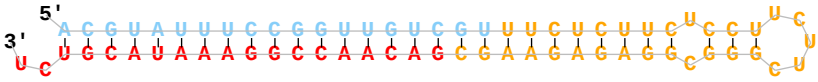
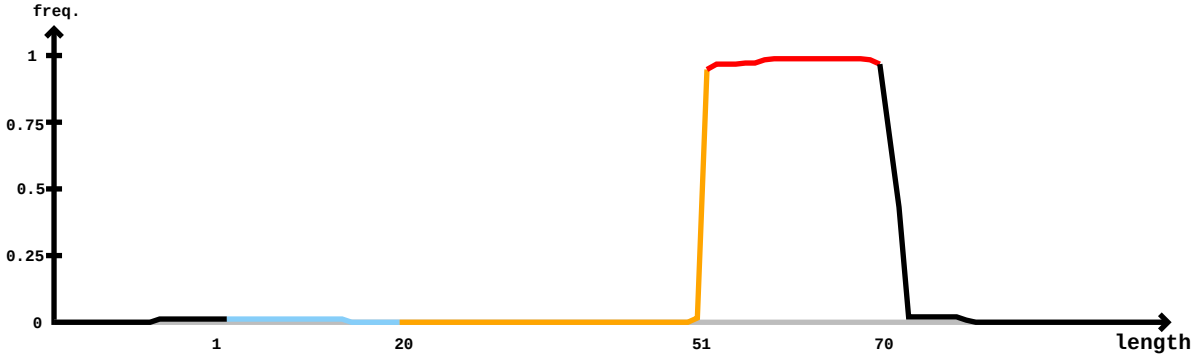
Provisional ID	:	NW_025322874.1_Acropora_millepora_isolate_JS-1_unplaced_genomic_scaffold_Amil_v2.1_xfSc0000223_whole_genome_shotgun_sequ
Score total	:	1.2
Score for star read(s)	:	-1.3
Score for read counts	:	0
Score for mfe	:	3.1
Score for randfold	:	0
Score for cons. seed	:	-0.6
Total read count	:	240
Mature read count	:	240
Loop read count	:	0
Star read count	:	0





Star Mature

```
5'- uaaccugcgaacgcagacguauuuccgguugucguuucucuucuccuucuucgggcggagagaagcgacaaccggaaauacgucugcguucgcaggcuagccaagcauacagca  -3'   exp
   ...(((((((((((((((((((((((((((((((((((((((.(((.....)))).))))))))))))))))))))))))))))))))))))))).........((.....)).   reads     mm        sample
   .........aacgcagacguauuuccggu.......................................................................................   3         0         seq
   ...........................................................cgacaaccggaaCuacgu..............................   1         1         seq
   ...........................................................cgacaaccggaaCuacgucugc..........................   3         1         seq
   ...........................................................gacaaccggaaUuacguc..............................   1         1         seq
   ...........................................................gacaaccggaaCuacguc..............................   3         1         seq
   ...........................................................gacaaccggaaCuacgucu.............................   123       1         seq
   ...........................................................gacaaccggaaCuacgucug............................   7         1         seq
   ...........................................................gacaaccggaaUuacgucug............................   2         1         seq
   ...........................................................gacaaccggaaCuacgucugc...........................   95        1         seq
   ............................................................acaaccggaaCuacgucu.............................   1         1         seq
   ............................................................acaaccggaaCuacgucugc...........................   4         1         seq
   .............................................................accggaaauacgucugcguucgca.......................   1         0         seq
   ..............................................................cggaaauacgucugcguucgU.........................   2         1         seq
   ..............................................................cggaaauacgucugcguucgca........................   1         0         seq
   .............................................................ggaaauacgucugcguucgc...........................   1         0         seq
```