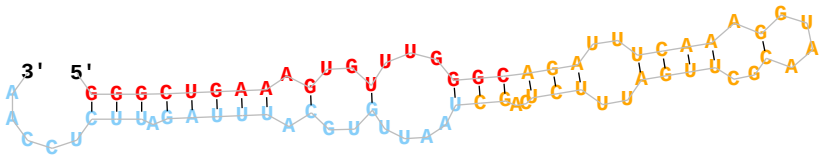
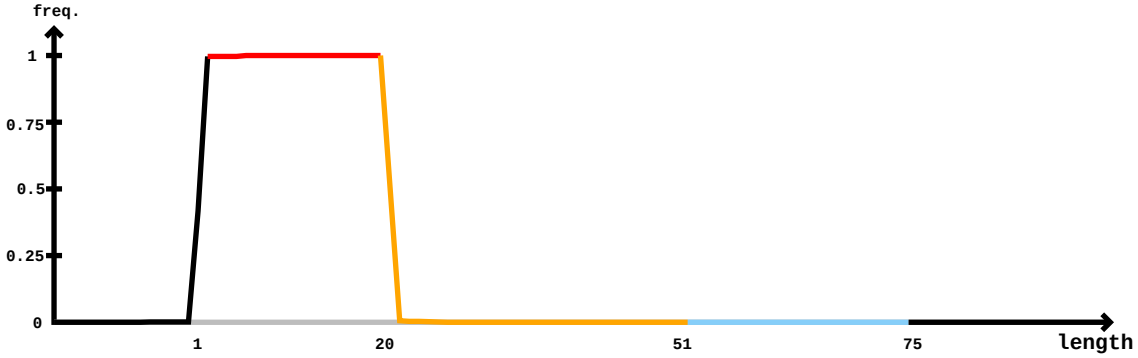
Provisional ID	:	NW_025323114.1_Acropora_millepora_isolate_JS-1_unplaced_genomic_scaffold_Amil_v2.1_xpSc0000551_whole_genome_shotgun_sequ
Score total	:	0.6
Score for star read(s)	:	-1.3
Score for read counts	:	0
Score for mfe	:	-1
Score for randfold	:	0
Score for cons. seed	:	3
Total read count	:	880
Mature read count	:	880
Loop read count	:	0
Star read count	:	0





Mature **Star**

```
5'- caauggcugugacugggcugaaaguguuugggcagauuucaaagguaacgcuugauuucucagcuaauugugcauuuagauucuccaaaaucgccugccaguuaucag  -3'    exp
   ...((((.((((.(((((((((.(..(...((((((..((((.(....)..))))..)))..)))....)..).)))))....))))...))))..)))).......   reads     mm        sample
   ........gugacuUggcugaaaguguuugggc.......................................................................   1         1         seq
   ............uAggcugaaaguguuugggc.......................................................................   1         1         seq
   ............uUggcugaaaguguuugggc.......................................................................   343       1         seq
   ............uCggcugaaaguguuugggc.......................................................................   4         1         seq
   ............uUggcugaaaguguuugggca......................................................................   17        1         seq
   ............uUggcugaaaguguuugggcag.....................................................................   2         1         seq
   ............gggcugaaaguguuugggc.........................................................................   1         0         seq
   ............Uggcugaaaguguuugggc.........................................................................   498       1         seq
   ............Cggcugaaaguguuugggc.........................................................................   3         1         seq
   ............Aggcugaaaguguuugggc.........................................................................   1         1         seq
   ............Uggcugaaaguguuugggca........................................................................   8         1         seq
   ................ugaaaguguuugggAagau.....................................................................   1         1         seq
   ................ugaaaguguuugggcagaCu....................................................................   1         1         seq
   ................ugaaaguguuugggcagaAuu...................................................................   1         1         seq
```