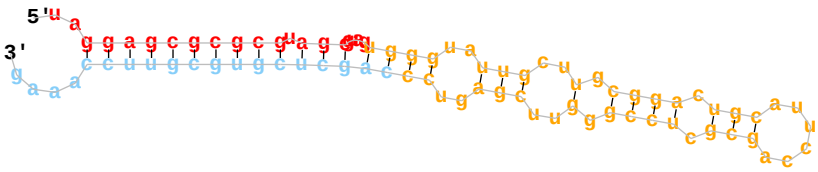
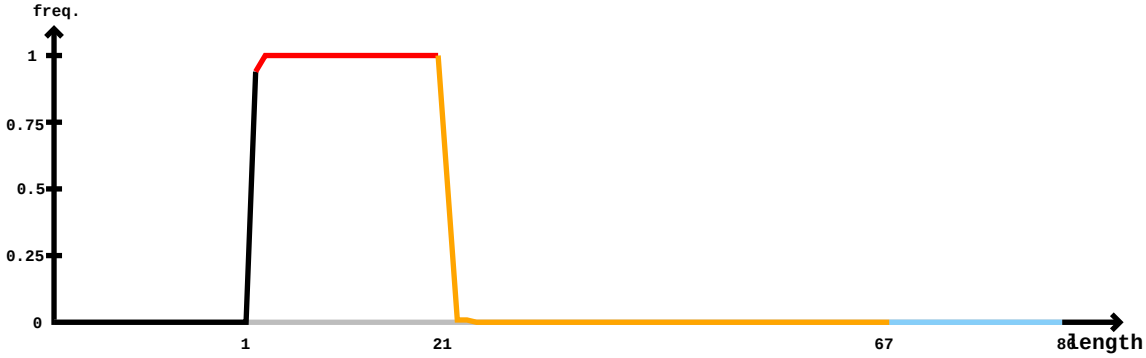
Provisional ID	:	Montipora_capitata_HIv3___Scaffold_14_879123
Score total	:	2.3
Score for star read(s)	:	-1.3
Score for read counts	:	0
Score for mfe	:	2
Score for randfold	:	1.6
Score for cons. seed	:	
Total read count	:	116
Mature read count	:	116
Loop read count	:	0
Star read count	:	0





Mature **Star**

```
5'- augauuaaaugauugaaugUAGGAGCGCGCGUAGCGUAGUGGGUAUUGCUUGCGGACUGCAUUCCAGCGCUCCGGGUUCGAGUCCCAGCUCGUGCGUUCCAAAGUUCAC  -3'   exp
    .............(((((...((((((((((.(((....((((..(((..(.((((.(((......))).)))).)..)))..))))))))))))))))))...)))))).  reads   mm   sample
    ...................AAGGAGCGCGCGUAGCGUAG.....................................................................  10      1    s23
    ...................AGGAGCGCGCGUAGCCUAG.....................................................................  2       1    s23

    ...................AAGGAGCGCGCGUAGCGUAG.....................................................................  24      1    s13

    ...................AAGGAGCGCGCGUAGCGUAG.....................................................................  8       1    s72

    ...................AAGGAGCGCGCGUAGCGUAG.....................................................................  1       1    s85

    ...................AAGGAGCGCGCGUAGCGUAG.....................................................................  16      1    s35
    ...................AGGAGCGCGCGUAGCGUAG......................................................................  1       0    s35

    ...................AAGGAGCGCGCGUAGCGUAG.....................................................................  24      1    s60
    ...................AGGAGCGCGCGUAGCGUAG......................................................................  3       0    s60

    ...................AAGGAGCGCGCGUAGCGUAG.....................................................................  8       1    s09
    ...................AGGAGCGCGCGUAGCGUAG......................................................................  1       0    s09

    ...................AAGGAGCGCGCGUAGCGUAG.....................................................................  17      1    s52
    ...................AAGGAGCGCGCGUAGCGUAGUGG..................................................................  1       1    s52
```