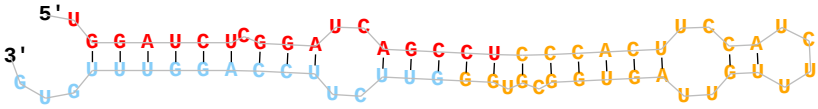
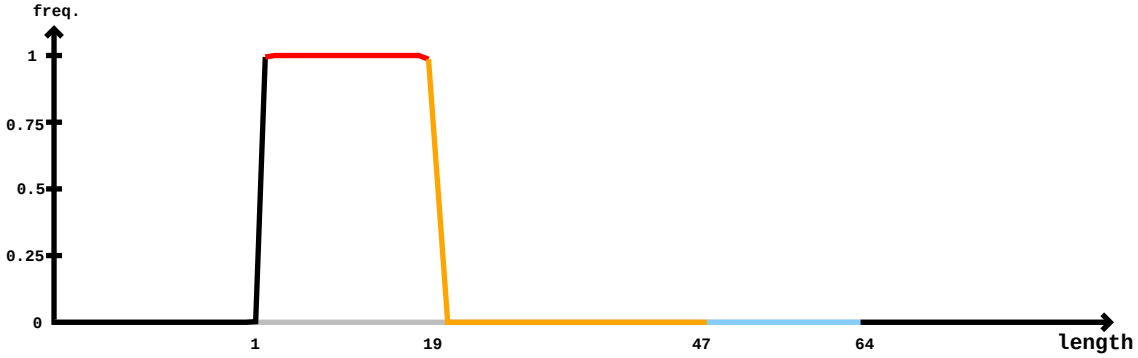
Provisional ID	:	NC_058067.1_Acropora_millepora_isolate_JS-1_chromosome_2_Amil_v2.1_whole_genome_shotgun_sequence_72848
Score total	:	1.4
Score for star read(s)	:	-1.3
Score for read counts	:	0
Score for mfe	:	-0.2
Score for randfold	:	0
Score for cons. seed	:	3
Total read count	:	1292
Mature read count	:	1292
Loop read count	:	0
Star read count	:	0





Mature Star

```
5'- guaaccauaauagccguucuuggaucucggaucagccucccacuuccaucuuuguuaguggcgugggguucuuccagguuugugggaccaauauccgucuuuacccaaa  -3'    exp
    ((((..((.(((...((((((..((((.(((..((((((((((((..((....))..)))))).).)))))..)))))))..).)))))...)))..))..))))..... reads      mm         sample
    ...................Guggaucucggaucagcc..................................................................... 1          1          seq
    ...................uuggaCcucggaucagcc..................................................................... 1          1          seq
    ...................uuggaCcucggaucagccu.................................................................... 1          1          seq
    ....................Cggaucucggaucagcc..................................................................... 1          1          seq
    ....................uggaucucggaucagcc..................................................................... 1          0          seq
    ....................uggaCcucggaucagcc..................................................................... 14         1          seq
    ....................uggaucucggaucagccu.................................................................... 2          0          seq
    ....................Gggaucucggaucagccu.................................................................... 1          1          seq
    ....................uggaCcucggaucagccu.................................................................... 1172       1          seq
    ....................Cggaucucggaucagccu.................................................................... 70         1          seq
    ....................uggaucucggaucagccuA................................................................... 1          1          seq
    ....................uggaucucggaucagccuG................................................................... 3          1          seq
    ....................uggaCcucggaucagccuc................................................................... 17         1          seq
    .....................ggaCcucggaucagccuc................................................................... 7          1          seq
```