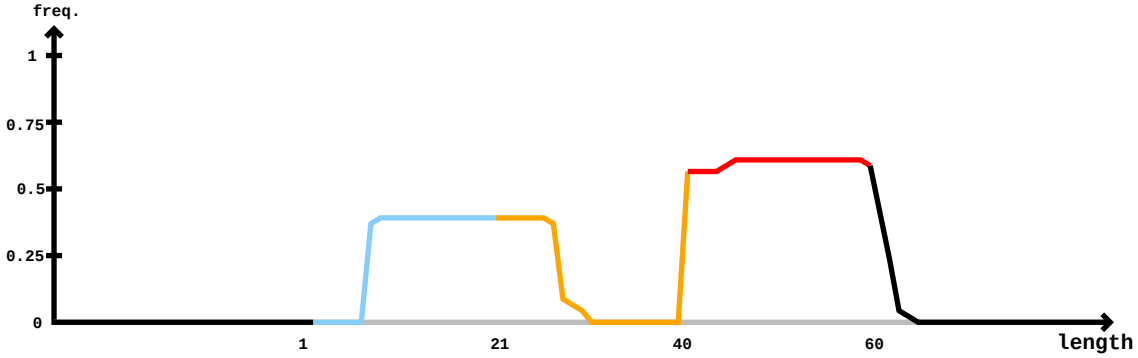
Provisional ID	:	NC_058067.1_Acropora_millepora_isolate_JS-1_chromosome_2_Amil_v2.1_whole_genome_shotgun_sequence_100232
Score total	:	0.7
Score for star read(s)	:	-1.3
Score for read counts	:	0
Score for mfe	:	2.6
Score for randfold	:	0
Score for cons. seed	:	-0.6
Total read count	:	28
Mature read count	:	28
Loop read count	:	0
Star read count	:	0



Star Mature

```
5'- ggggggguccaaauccgcgaagggggguccaaauccgcuagcggauauggaccggggggguccaaauccgcuagcggauauggaccaggggguccaaauccgcgggggguc  -3'    exp
    ...((..((....((.....)).((((((.((((((((((((((.((((((.....)))))).)))))))))))))).)))))).))..)).)..((((.....)))).   reads      mm         sample
    ..............................uccgcuagcggauauggac.................................................................   2          0          seq
    ..............................uccgcuagcggauauggacc................................................................   16         0          seq
    ..............................uccgcuagcggauauggacA................................................................   10         1          seq
    ..............................uccgcuagcggauauggaccU...............................................................   2          1          seq
    ..............................uccgcuagcggauauggaccgg..............................................................   1          0          seq
    ..............................uccgcuagcggauauggaccggU.............................................................   1          1          seq
    ..............................cUgcuagcggauauggaccggg..............................................................   1          1          seq
    ...............................................................uccgcuagcggauauggac.........................   2          0          seq
    ...............................................................uccgcuagcggauauggacA........................   10         1          seq
    ...............................................................uccgcuagcggauauggacc........................   16         0          seq
    ...............................................................uccgcuagcggauauggacca.......................   2          0          seq
    ...............................................................uccgcuagcggauauggaccU.......................   2          1          seq
    ...............................................................uccgcuagcggauauggaccaU......................   7          1          seq
    ...............................................................uccgcuagcggauauggaccaA......................   1          1          seq
    ...............................................................uccgcuagcggauauggaccaC......................   1          1          seq
    ..................................................................cuagcggauauggaccagU.......................   1          1          seq
    ...................................................................uagcggauauggaccaggU......................   1          1          seq
```