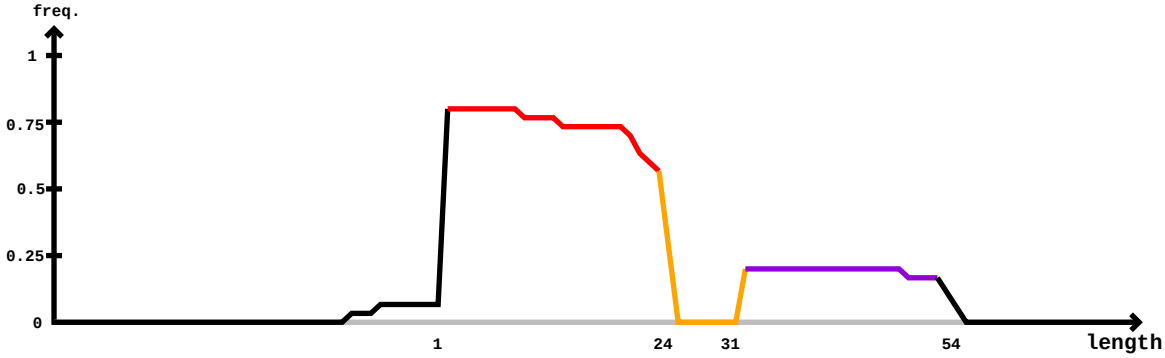
Provisional ID	:	NC_058067.1_Acropora_millepora_isolate_JS-1_chromosome_2_Amil_v2.1_whole_genome_shotgun_sequence_71083
Score total	:	17.5
Score for star read(s)	:	3.9
Score for read counts	:	8.3
Score for mfe	:	2.3
Score for randfold	:	0
Score for cons. seed	:	3
Total read count	:	28
Mature read count	:	22
Loop read count	:	0
Star read count	:	6



Mature Star

```
5'- uacuaaaacaauauuauuauuccucuagcccucauggccucugagucaauagcccauucggccuuuggccucgugggcuauugauucagggccccaugcauuugggcucaag  -3'   obs
    uacuaaaacaauauuauuauuccucuagcccucauggccucugagucaauagcccauucggccuuuggccucgugggcuauugauucagggccccaugcauuugggcucaag        exp
    ......................((..(((((.(((((.(((((((((((((((((((..((((...)))).))))))))))))))))))).)))))).....)))))..))   reads   mm   sample
    ............................ccucauggccucugaguc.......................................................................   1       0    seq
    ...............................cauggGcucugagucaaua....................................................................   1       1    seq
    ....................................ucugagucaauagcccauu...................................................................   1       0    seq
    ....................................ucuAagucaauagcccauuc..................................................................   2       1    seq
    ....................................ucugagucaauagcccauucU.................................................................   1       1    seq
    ....................................ucugagucaauagcccauucUg................................................................   1       1    seq
    ....................................ucugagucaauagcccauucggc...............................................................   8       0    seq
    ....................................ucugaguUaauagcccauucggc...............................................................   4       1    seq
    ....................................ucugagucaauagcccauucggU...............................................................   2       1    seq
    ....................................ucugaCucaauagcccauucggc...............................................................   2       1    seq
    ....................................ucugagucaauagcccauucggcU..............................................................   1       1    seq
    ......................................................................ucgugggcuauugauuc.......................   1       0    seq
    ......................................................................ucgugggcuauugauucaggg...................   1       0    seq
    ......................................................................ucgugggcuauugauucagggc..................   1       0    seq
    ......................................................................ucgugggcuauugauucagggU..................   3       1    seq
```