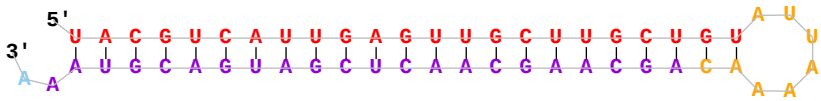
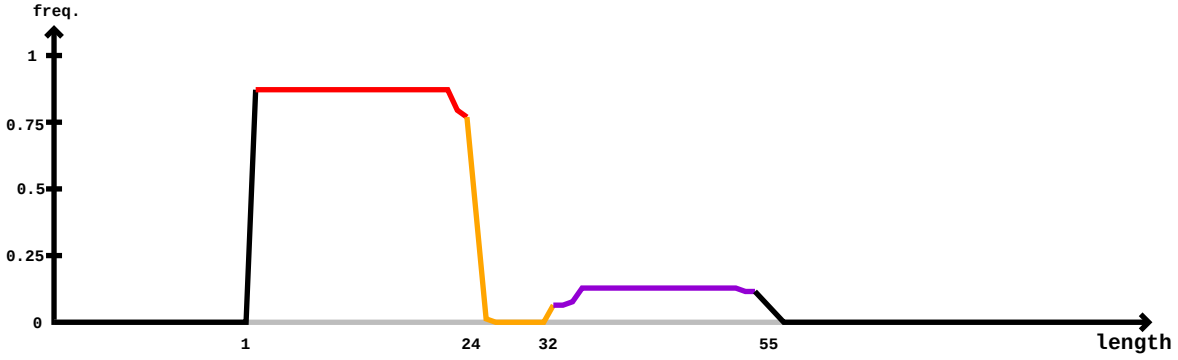
Provisional ID	:	NC_058078.1_Acropora_millepora_isolate_JS-1_chromosome_13_Amil_v2.1_whole_genome_shotgun_sequence_497039
Score total	:	0.5
Score for star read(s)	:	-1.3
Score for read counts	:	0
Score for mfe	:	2.4
Score for randfold	:	0
Score for cons. seed	:	-0.6
Total read count	:	78
Mature read count	:	68
Loop read count	:	0
Star read count	:	10






```
                    Mature                Star
5'- ccuuagucgaaaaagcuuuuacgucauugaguugcuugcuguauuaaaacagcaagcaacucgaugacguaaaagcuucaaauguucgaaaagacgcgcauauaaaagcgau  -3'    obs
    ccuuagucgaaaaagcuuuuacgucauugaguugcuugcuguauuaaaacagcaagcaacucgaugacguaaaagcuucaaauguucgaaaagacgcgcauauaaaagcgau         exp
    .(((..(((((.(((((((((((((((((((((((((((((.....)))))))))))))))))))))))))))))).....)))))..)))...(((........)))..   reads      mm         sample
    ..................uacgucauugaguugcuugcu.......................................................................   4          0          seq
    ..................uacgucauugaguugcuugcG.......................................................................   1          1          seq
    ..................uacgucauugaguugcuugcA.......................................................................   1          1          seq
    ..................uacgucauugaguugcuugcug......................................................................   2          0          seq
    ..................uacgucauugaguugcuugcugA.....................................................................   1          1          seq
    ..................uacgucauugaAuugcuugcugu.....................................................................   1          1          seq
    ..................uacgucauugagCugcuugcugu.....................................................................   1          1          seq
    ..................uacgucauugaguugcuugcugC.....................................................................   7          1          seq
    ..................uacgucauugaguugcuugcugG.....................................................................   1          1          seq
    ..................uacgucauugaguugcuugcugu......................................................................   43         0          seq
    ..................uacgucauugaguugcuugcuguU.....................................................................   4          1          seq
    ..................uacgucauugaguugcuugcugCa....................................................................   1          1          seq
    ..................uacgucauugaguugcuugcuguUu...................................................................   1          1          seq
    ...................................................agcaagcaacucgaugacguaU.....................................   4          1          seq
    ...................................................agcaagcaacucgaugacguaaa....................................   1          0          seq
    ....................................................caagcaacucgaugacgu........................................   1          0          seq
    ....................................................aagcaacucgaugacguaaa......................................   3          0          seq
    ....................................................aagcaacucgaugacguaaG.....................................   1          1          seq
```