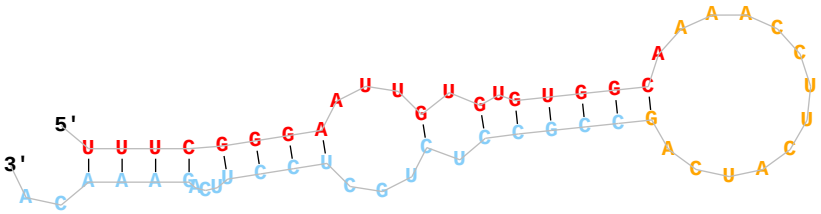
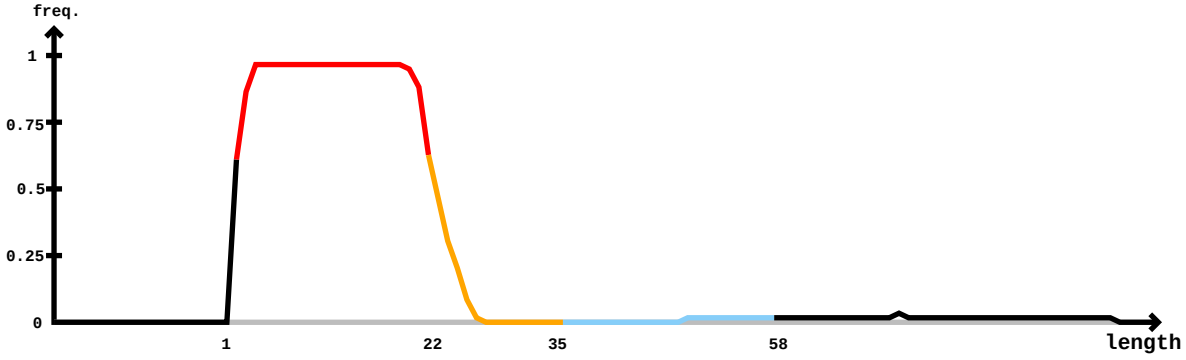
Provisional ID	:	NW_025322618.1_Acropora_millepora_isolate_JS-1_unplaced_genomic_scaffold_Amil_v2.1_Sc0000038_whole_genome_shotgun_sequen
Score total	:	1.1
Score for star read(s)	:	-1.3
Score for read counts	:	0
Score for mfe	:	-0.5
Score for randfold	:	0
Score for cons. seed	:	3
Total read count	:	57
Mature read count	:	57
Loop read count	:	0
Star read count	:	0





Mature Star

```
5'- aguugaaaacucgcauuuuucgggaauugugugugggcaaaaccuucaucagccgccucugcuccuucagaaacagugccaggcgagucauucucaucuacguugccagcagcc  -3'    exp
    .((((..(((..((((((((((((...(.(.(((((.............)))))).)...))))...)))).)))))).((..(((.....)))..))..)))..)))).....   reads     mm        sample
    ................uuucgggaauugugugugu.A.......................................................................   1         1         seq
    ................uuucgggaauugugugugg.A.......................................................................   4         1         seq
    ................uuucgggaauugugugugg.U.......................................................................   1         1         seq
    ................uuucgggaauugugugugg.c.......................................................................   3         0         seq
    ................uuucgggaauugugugugg.A.......................................................................   1         1         seq
    ................uuucgggaauugugugugg.cG......................................................................   1         1         seq
    ................uuucgggaauugugugugg.ca......................................................................   9         0         seq
    ................uuucgggaauugugugugg.caa.....................................................................   3         0         seq
    ................uuucgggaauugugugugg.caaa....................................................................   4         0         seq
    ................uuucgggaauugugugugg.caaaa...................................................................   4         0         seq
    ................uuucgggaauugugugugg.caaaac..................................................................   4         0         seq
    ................uuucgggaauugugugugg.caaaacG.................................................................   1         1         seq
    ...............uucgggaauugugugugg.U.........................................................................   1         1         seq
    ...............uucgggaauugugugugg.c.........................................................................   7         0         seq
    ...............uucgggaauugugugugg.ca........................................................................   3         0         seq
    ...............uucgggaauugugugugg.caaC.....................................................................   1         1         seq
    ...............uucgggaauugugugugg.caaa......................................................................   1         0         seq
    ...............uucgggaauugugugugg.caaaa.....................................................................   2         0         seq
    ..............ucgggaauugugugugg.U...........................................................................   1         1         seq
    ..............ucgggaauugugugugg.c...........................................................................   1         0         seq
    ..............ucgggaauugugugugg.ca..........................................................................   3         0         seq
    ..............ucgggaauugugugugg.caaaa.......................................................................   1         0         seq
    ...........................................................uucaAaaacagugccaggcgagu.........................   1         1         seq
    ...........................................................................ucauuAucaucuacguugccagc....       1         1         seq
```