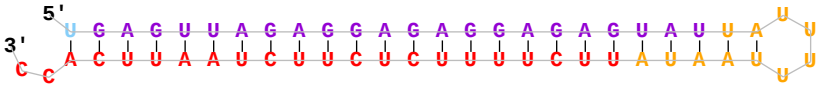
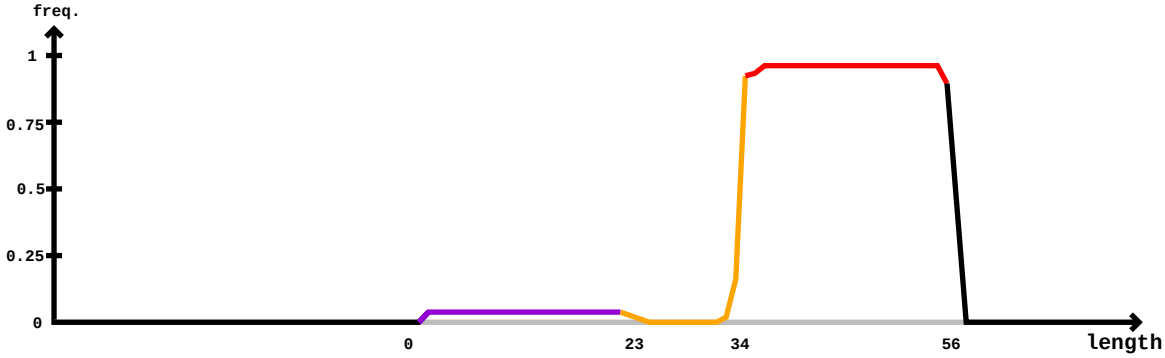
Provisional ID	:	NW_025322763.1_Acropora_millepora_isolate_JS-1_unplaced_genomic_scaffold_Amil_v2.1_xfSc0000033_whole_genome_shotgun_sequ
Score total	:	56.3
Score for star read(s)	:	3.9
Score for read counts	:	47.5
Score for mfe	:	1.8
Score for randfold	:	0
Score for cons. seed	:	3
Total read count	:	105
Mature read count	:	101
Loop read count	:	0
Star read count	:	4






```
                                   Star                                   Mature
5'- ucaaaggagacuaaugggaacuaugacauuccaugaugaguuagaggagaggagaguauuauuuuuaauauucuuuucucuucuaauucaccauuguaugguaauuguuaa  -3'   obs
    ucaaaggagacuaaugggaacuaugacauuccaugaugaguuagaggagaggagaguauuauuuuuaauauucuuuucucuucuaauucaccauuguaugguaauuguuaa        exp
    ....((....)).........(((((..(((((.(((((((((((((((((((((((....)))))))))))))))))))))))).....)))))....)))))).   reads   mm   sample
    ....................................gaguuagaggagaggagagua.................................................   2       0    seq
    ....................................gaguuagaggagaggagaguaC................................................   2       1    seq
    ....................................................................uauucuuuucucuucuaauucacA...................   2       1    seq
    ....................................................................auucuuuucucuucuaauucacU....................   11      1    seq
    ....................................................................auucuuuucucuucuaauucacA....................   4       1    seq
    ....................................................................uucuuuucucuucuaauucaU.....................   1       1    seq
    ....................................................................uucuuuucucuucuaauucac.....................   3       0    seq
    ....................................................................uucuuuucucuucuaauucacA....................   4       1    seq
    ....................................................................uucuuuucucuucuaauucacc....................   11      0    seq
    ....................................................................uucuuuucucuucuaauucacG....................   1       1    seq
    ....................................................................uucuuuucucuucuaauucacU....................   60      1    seq
    ....................................................................ucuuuucucuucuaauucacU.....................   1       1    seq
    ......................................................................cuuuucucuucuaauucac.....................   2       0    seq
    ......................................................................Uuuuucucuucuaauucac.....................   1       1    seq
```