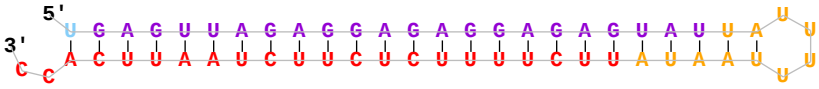
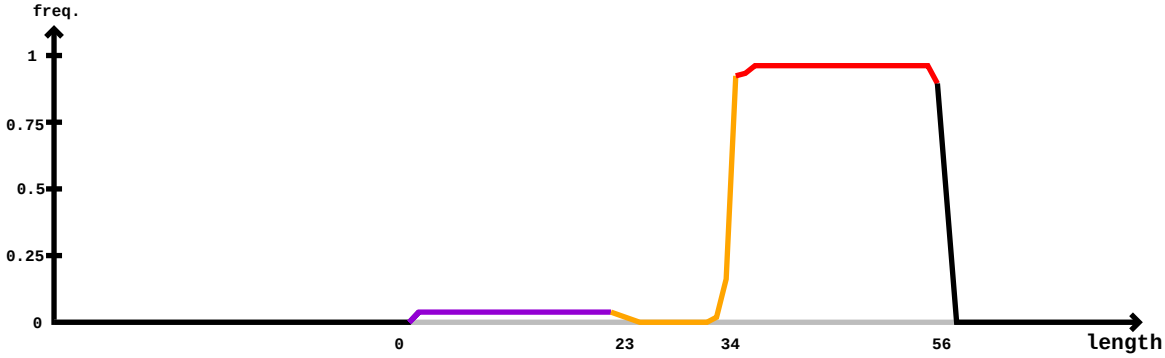
Provisional ID	:	NW_025322864.1_Acropora_millepora_isolate_JS-1_unplaced_genomic_scaffold_Amil_v2.1_xfSc0000213_whole_genome_shotgun_sequ
Score total	:	52.7
Score for star read(s)	:	3.9
Score for read counts	:	47.5
Score for mfe	:	1.8
Score for randfold	:	0
Score for cons. seed	:	-0.6
Total read count	:	105
Mature read count	:	101
Loop read count	:	0
Star read count	:	4






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