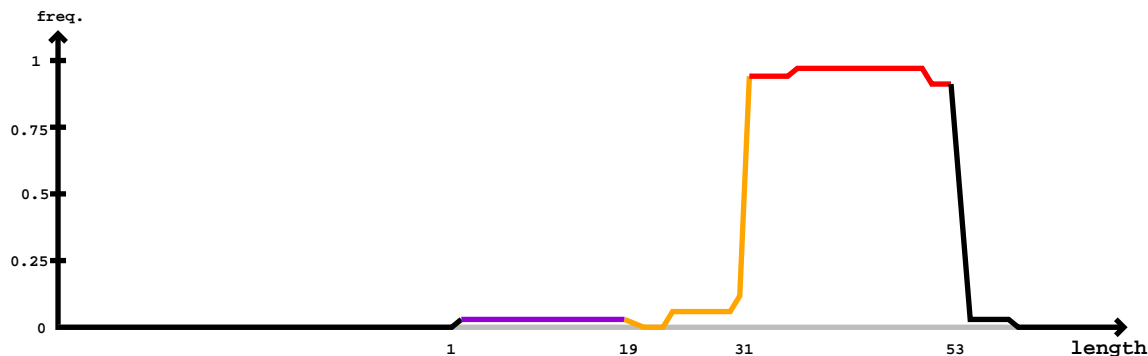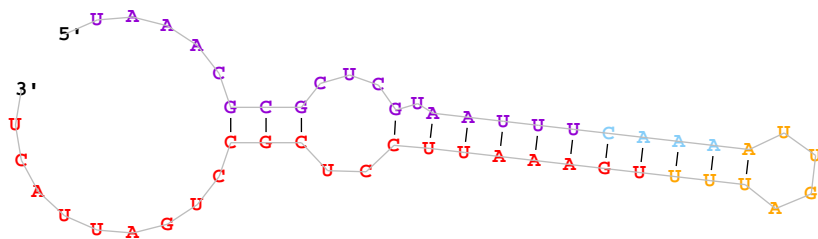| Provisional ID	: NW_025322636.1_Acropora_millepora_isolate_JS-1_unplaced_genomic_scaffold_Amil_v2.1_Sc0000107_whole_genome_shotgun_sequen
Score total	: 17.2
Score for star read(s)	: 3.9
Score for read counts	: 9.8
Score for mfe	: 0.5
Score for randfold	: 0
Score for cons. seed	: 3
Total read count	: 31
Mature read count	: 30
Loop read count	: 0
Star read count	: 1


```
                                                               Star                    Mature
5'- uuaauuaguaauuggacuucauguccuacaauucagggaauaaacgcgcucguaauuucaaaauugauuuugaaauuccucgccugauuacucccgaauuguacucca  -3'   obs
    uuaauuaguaauuggacuucauguccuacaauucagggaauaaacgcgcucguaauuucaaaauugauuuugaaauuccucgccugauuacucccgaauuguacucca        exp
    ............(((((.....)))).(((((((.((((.(((.((.((..(.(((((((((((....))))))))))).)..)).)).))).)))).))))))).....    reads   mm    sample
    ...................................uaaacgGgcucguaauuu.....................................................     1      1     seq
    ..................................................auugauuuugaaauuccucgccugauuacA...........................     2      1     seq
    ..................................................uugaaauuccucgccugauuacu.................................     2      0     seq
    ..................................................ugaaauuccucgccugauu.....................................     2      0     seq
    ..................................................ugaaauuccucgccugauuacG..................................     1      1     seq
    ..................................................ugaaauuccucgccugauuacA..................................     3      1     seq
    ..................................................ugGaauuccucgccugauuacu..................................     1      1     seq
    ..................................................ugaaauuccucgccugauuacC..................................     3      1     seq
    ..................................................ugaaauuccucgccugauuacu..................................    18      0     seq
    ....................................................uuAcucgccugauuacucccga...........                         1      1     seq
```