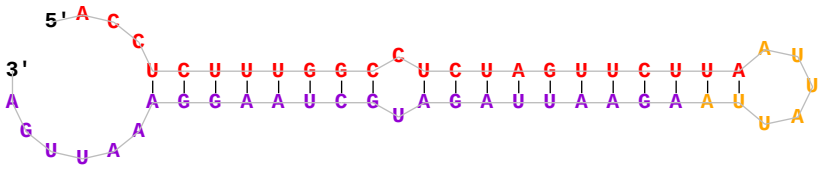
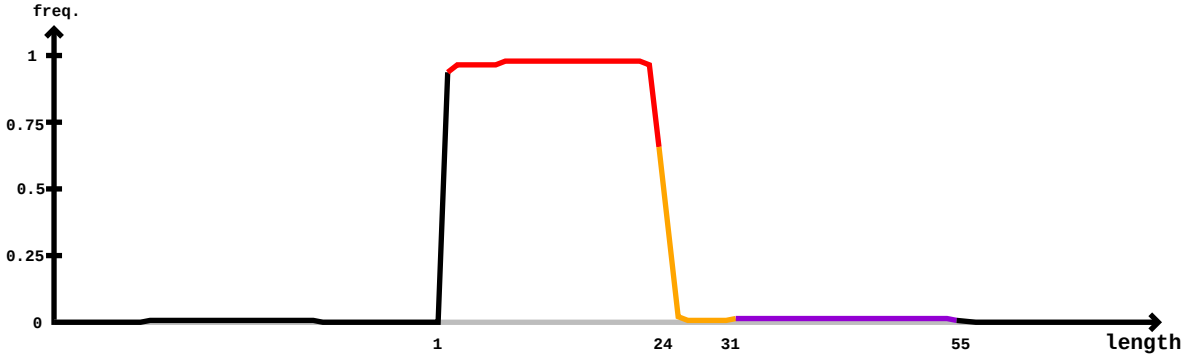
Provisional ID	:	NC_058066.1_Acropora_millepora_isolate_JS-1_chromosome_1_Amil_v2.1_whole_genome_shotgun_sequence_66778
Score total	:	70.2
Score for star read(s)	:	3.9
Score for read counts	:	65.9
Score for mfe	:	1.1
Score for randfold	:	0
Score for cons. seed	:	-0.6
Total read count	:	141
Mature read count	:	139
Loop read count	:	0
Star read count	:	2





Mature Star

```
5'- aaguguaauuacaugaugaauagauuacuauuuuucucaaccucuuuggccucuaguucuuaauuauuaagaauuagaugcuaaggaaauugagaaucaaacagcaaagacuc  -3'   obs
    aaguguaauuacaugaugaauagauuacuauuuuucucaaccucuuuggccucuaguucuuaauuauuagaauuagaugcuaaggaaauugagaaucaaacagcaaagacuc        exp
    ..((((....))))...((((((....))))))(((((((..((((((((.((((((((((.....)))))))))))).)))))))).))))))).................   reads   mm   sample
    ........uuGcaugaugaauagauu.......................................................................................   1       1    seq
    ......................................accucuuuggccucuaguucu.......................................................   1       0    seq
    ......................................accucuuuggccucAaguucu.......................................................   1       1    seq
    ......................................accucuuuggccucAaguucuu......................................................   19      1    seq
    ......................................accucuuuggccucuaguucuC......................................................   5       1    seq
    ......................................accucuuuggccucuaguucuu......................................................   19      0    seq
    ......................................accucuuuggccucuaguucuCa.....................................................   1       1    seq
    ......................................accucuuuggccucuaguucuua.....................................................   39      0    seq
    ......................................accucuuugAccucuaguucuua.....................................................   1       1    seq
    ......................................accucuuuggccucAaguucuua.....................................................   36      1    seq
    ......................................accucuuuggccucuaguucuuU.....................................................   1       1    seq
    ......................................accucuuuggccucuaguucuuaC....................................................   1       1    seq
    ......................................accucuuuggccucuaguucuuaU....................................................   9       1    seq
    ......................................accucuuuggccucuaguucuuaUu...................................................   1       1    seq
    .....................................ccucuuuggccucAaguucuu........................................................   1       1    seq
    .....................................ccucuuuggccucuaguucuuaU.....................................................   2       1    seq
    .....................................ccucuuuggccucuaguucuuaC.....................................................   1       1    seq
    ...........................................uuggccucuaguucuuaau.....................................................   1       0    seq
    ...........................................uuggccucuaguucuuaauuauua................................................   1       0    seq
    ..............................................................agaauuagaugcuaaggaaauug..........................   1       0    seq
    ..............................................................agaauuagaugcuaaggaaauugU.........................   1       1    seq
```