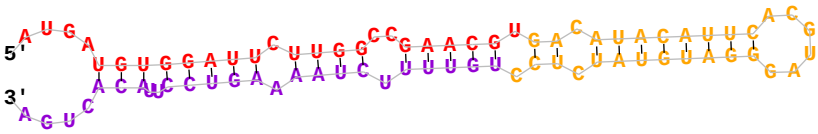
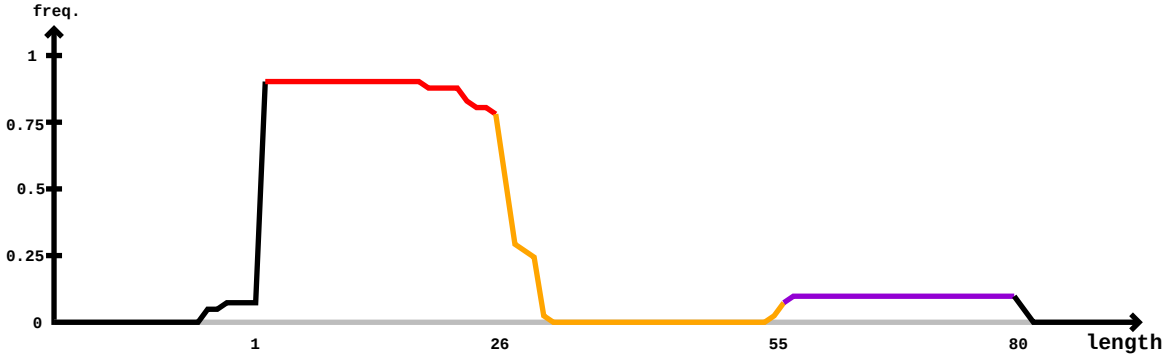
Provisional ID	:	NC_058079.1_Acropora_millepora_isolate_JS-1_chromosome_14_Amil_v2.1_whole_genome_shotgun_sequence_518700
Score total	:	20.4
Score for star read(s)	:	3.9
Score for read counts	:	13.4
Score for mfe	:	0.1
Score for randfold	:	0
Score for cons. seed	:	3
Total read count	:	38
Mature read count	:	34
Loop read count	:	0
Star read count	:	4





Mature **Star**

```
5'- uccgcgaaacaaguggacagaugauguggauucuuggccgaacgugacauacauucacguagggauguaucuccuguuuucuaaaaguccuuacacugaacaaguagccaa -3'   obs
    uccgcgaaacaaguggacagaugauguggauucuuggccgaacgugacauacauucacguagggauguaucuccuguuuucuaaaaguccuuacacugaacaaguagccaa        exp
    (((((.......)))))(((.((.((.(((((.((((..(((((.((.(((((((((.......)))))))).)).)))))).)))).)))))).)))))))............  reads   mm   sample
    ..............ggacagaugauguggauucuugg.......................................................................  1       0    seq
    ..............ggacagaugauguggauucuuggccgaa...................................................................  1       0    seq
    ................acagaugauguggauucuuggccgaacgu...............................................................  1       0    seq
    ....................augauguggauucuuggccga.....................................................................  2       0    seq
    ....................augauguggauucuuggccgaacA..................................................................  1       1    seq
    ....................augauguggauucuuggccgaacCu.................................................................  3       1    seq
    ....................augauguggauucuuggccgaacgA.................................................................  1       1    seq
    ....................augauguggauucuuggccgaacgu.................................................................  9       0    seq
    ....................augauguggauucuuggccgaacgC.................................................................  2       1    seq
    ....................augauguggauucuuggccgaacgug................................................................  3       0    seq
    ....................augauguggauucuuggccgaacguA................................................................  1       1    seq
    ....................augauguggauucuuggccgaacguga...............................................................  1       0    seq
    ....................augauguggauucuuggccgaacgugac..............................................................  1       0    seq
    ....................augauguggauucuuggccgaacgugacG.............................................................  1       1    seq
    ....................augauguggauucuCggccgaacgugaca.............................................................  1       1    seq
    ....................augauguggauucuuggccgaacgugaca..............................................................  7       0    seq
    ....................augauguggauucuuggccgaacgugacau.............................................................  1       0    seq
    ......................................................................cuguuuucuaaaaguccuuacacugaa..........  1       0    seq
    .......................................................................uguuuucuaaaaguccuuacacuga...........  2       0    seq
    ......................................................................guuuucuaaaaguccuuacacugaa..........  1       0    seq
```