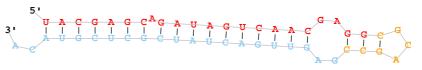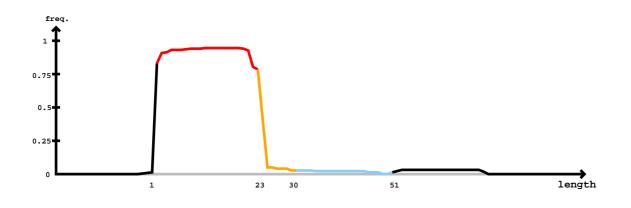|		
Provisional ID	:	NW_025322628.1_Acropora_millepora_isolate_JS-1_unplaced_genomic_scaffold_Amil_v2.1_Sc0000086_whole_genome_shotgun_sequer
Score total	:	0.1
Score for star read(s)	:	-1.3
Score for read counts	:	0
Score for mfe	:	2
Score for randfold	:	0
Score for cons. seed	:	-0.6
Total read count	:	203
Mature read count	:	203
Loop read count	:	0
Star read count	:	0

Mature   **Star**

```
5'- acuaagagcgcgaguuuucuacgagcagauagucaacgaggcgcagccgaguugacuaucgcucguacaaaacgagggcgaguagucaauuguuuuaguauaaauuua  -3'   exp
    (((((...((.(..(((((.((((((((.((((((((((..(((...)))..)))))))))))))))))))).)))))..).))...))))....................   reads    mm        sample
    ...............uucuacgagcagauagucaacgaggcgcagc.............................................................       1      0        seq
    ................ucuacgagcagauagucaacgaggU.................................................................       1      1        seq
    .................cuacgagcagauagucaacgaCg.................................................................       1      1        seq
    .................uacgagcagauagucaac......................................................................       1      0        seq
    .................uacgagcagauagucaacg.....................................................................       3      0        seq
    .................uacgagcagauagucaacga....................................................................      22      0        seq
    .................uacgagcagauagCcaacga....................................................................       1      1        seq
    .................uacgagcagauagucaacgG....................................................................       1      1        seq
    .................uUcgagcagauagucaacgag...................................................................       1      1        seq
    .................uacgagcagauagucaacgag...................................................................       2      0        seq
    .................uacgagcagauagucaacgaC..................................................................       1      1        seq
    .................uacgagcagauagucaacgaCg.................................................................      28      1        seq
    .................uacgagcagauagucaacgagC.................................................................       1      1        seq
    .................uaUgagcagauagucaacgagg.................................................................       1      1        seq
    .................uGcgagcagauagucaacgagg.................................................................       2      1        seq
    .................uacgagcagauagCcaacgagg.................................................................       1      1        seq
    .................uacgagUagauagucaacgagg.................................................................       1      1        seq
    .................uacgagcagauagucaacgagA.................................................................      25      1        seq
    .................uacgagcagauagucaacgGgg.................................................................       1      1        seq
    .................uacgagcagauagucaacgagg.................................................................      84      0        seq
    .................uacgagcagauagucaacgaggA................................................................       2      1        seq
    .................uacgagcagauagucaacgaggcgcagc...........................................................       1      0        seq
    ..................acgagcagauagucaacga...................................................................       3      0        seq
    ..................acgagcagauagucaacgaUg.................................................................       1      1        seq
    ..................acgagcagauagucaacgagA.................................................................       1      1        seq
    ..................acgagcagauagucaacgagC.................................................................       1      1        seq
    ..................acgagcagauagucaacgaCg.................................................................       6      1        seq
    ..................acgagcagauagucaacgagg.................................................................       4      0        seq
    ..................acgagcagauagucaacgaggU................................................................       1      1        seq
    ...................cgagcagauagucaacgagg................................................................       1      0        seq
    ....................gagcagauagucaacgagg...............................................................       1      0        seq
    ....................gagcagauagucaacgagA...............................................................       2      1        seq
    ....................gagcagauagucaacgaggcgc............................................................       1      0        seq
    ......................cagauagucaacgaggcgU............................................................       1      1        seq
```

acuaagagcgcgaguuuucuacgagcagauagucaacgaggcgcagccgaguugacuaucgcucguacaaaacgagggcgaguagucaauuguuuuaguauaaauuua

..........................agauagucaacgaggcgcagU..	1	1	seq
........................uagucaacgaggcgcagccgagu..	1	0	seq
...............................cgcagccgaguugacuaucgcC...	2	1	seq
..............................cgcagccgaguugacuaucgcucgu...	2	0	seq
..............................cgcagccgaguugGcuaucgcucgu...	1	1	seq
....aaaacgagggcgaguaguc....................................	1	0	seq
....aaaacgagggUgaguaguca...................................	2	1	seq
....aaacgagggcgaguaguU....................................	1	1	seq
....aaacgagggcgaguaguc....................................	1	0	seq
....aacgagggcgaguaguca...................................	2	0	seq