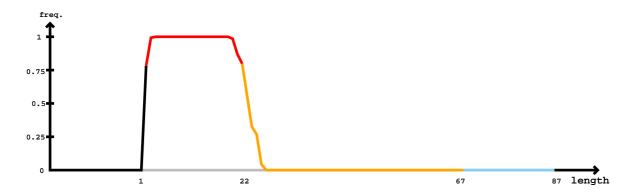|		
Provisional ID	:	NC_058067.1_Acropora_millepora_isolate_JS-1_chromosome_2_Amil_v2.1_whole_genome_shotgun_sequence_107295
Score total	:	0.3
Score for star read(s)	:	-1.3
Score for read counts	:	0
Score for mfe	:	2.2
Score for randfold	:	0
Score for cons. seed	:	-0.6
Total read count	:	138
Mature read count	:	138
Loop read count	:	0
Star read count	:	0

Mature   **Star**

```
5'- acugaguggaguaaaauucagggaguaaucggacgagugauuucaaaacagacgaacgcgaagcgugaggcugauuugaaauuacuagcacgauuaccccugaauuguacga -3'     exp
          .........(((.(((((((((.((((((((..(.(((((((((((((.(((.(.((((...)))).).))).)))))))))))).)..))))))))))))).)))..     reads   mm   sample
.................cagggaguaaucggacga.............................................................................       1       0    seq
.................cagggaguaaucggacgU.............................................................................       1       1    seq
.................cagggaguaaucggacgag............................................................................       9       0    seq
.................cagggaguaaucggacgaA............................................................................       3       1    seq
.................cagggaguaaucggacgaU............................................................................       1       1    seq
.................cagggaguaaucggacgaUu...........................................................................       3       1    seq
.................cagggaguaaucggacgagu...........................................................................       7       0    seq
.................cagggaguaaucggGcgagug..........................................................................       19      1    seq
.................cagggaguaaucggacgagug..........................................................................       25      0    seq
.................cagggaguaaucggacgaguU..........................................................................       1       1    seq
.................cagggaguaaucggacgaguA..........................................................................       4       1    seq
.................cagggaguaaucggacgagugC.........................................................................       5       1    seq
.................cagggaguaaucggacgagugU.........................................................................       8       1    seq
.................cagggaguaaucggacgaguga.........................................................................       2       0    seq
.................cagggaguaaucggGcgaguga.........................................................................       1       1    seq
.................cagggaguaaucggacgagugau........................................................................       4       0    seq
.................cagggaguaaucggacgagugUu........................................................................       1       1    seq
.................cagggaguaaucggGcgagugauu.......................................................................       1       1    seq
.................cagggaguaaucggacgagugauu.......................................................................       6       0    seq
.................cagggaguaaucggacgagugauA.......................................................................       1       1    seq
.................cagggaguaaucggacgagugauuu......................................................................       5       0    seq
.................agggaguaaucggacgaA.............................................................................       2       1    seq
.................agggaguaaucggacgag.............................................................................       1       0    seq
.................agggaguaaucggacgagugau.........................................................................       1       0    seq
.................agggaguaaucggGcgagugau.........................................................................       1       1    seq
.................agggaguaaucggacgagugaC.........................................................................       1       1    seq
.................agggaguaaucggacgagugauu........................................................................       12      0    seq
.................agggaguaaucggacgagugaCu........................................................................       1       1    seq
.................agggaguaaucggacgagugauC........................................................................       2       1    seq
.................agggaguaaucggGcgagugauu........................................................................       7       1    seq
.................agggaguaaucggGcgagugauuu.......................................................................       1       1    seq
.................gggaguaaucggGcgagugauu.........................................................................       1       1    seq
```