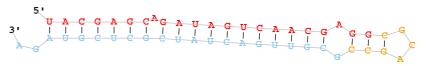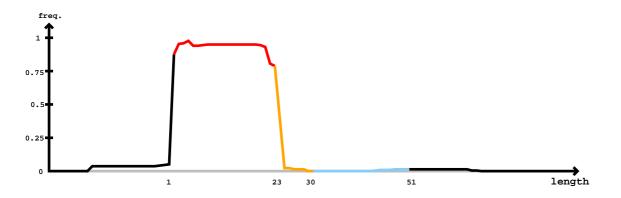| Provisional ID	:	NC_058075.1_Acropora_millepora_isolate_JS-1_chromosome_10_Amil_v2.1_whole_genome_shotgun_sequence_402538
Score total	:	4.2
Score for star read(s)	:	-1.3
Score for read counts	:	0
Score for mfe	:	2.5
Score for randfold	:	0
Score for cons. seed	:	3
Total read count	:	203
Mature read count	:	203
Loop read count	:	0
Star read count	:	0

Mature   **Star**

```
5'- auuagacuacgagcacgaguuuucuacgagcagauagucaacgaggcgcagccgcguugacuaucgcucguagaaaacgagggcguguagucuaauuguuuuaguaua  -3'   exp
    (((((((((((.((.(..(((((((((((.(((((((((((.(((...))).))))))))))))))))))))))))..).)).)))))))))............     reads   mm      sample
    ......uacgagcUcgaguuuucuacg.....................................................................            8       1       seq
    ....................uucuacgagcagauagucaacgaggcgcagc............................................            1       0       seq
    .................ucuacgagcagauagucaacgaggU...................................................            1       1       seq
    ..................cuacgagcagauagucaacgaCg....................................................            1       1       seq
    ..................uacgagcagauagucaac........................................................            1       0       seq
    ..................uacgagcagauagucaacg.......................................................            3       0       seq
    ..................uacgagcagauagCcaacga......................................................            1       1       seq
    ..................uacgagcagauagucaacga......................................................            22      0       seq
    ..................uacgagcagauagucaacgG......................................................            1       1       seq
    ..................uacgagcagauagucaacgaC.....................................................            1       1       seq
    ..................uUcgagcagauagucaacgag.....................................................            1       1       seq
    ..................uacgagcagauagucaacgag.....................................................            2       0       seq
    ..................uacgagcagauagucaacgaCg....................................................            28      1       seq
    ..................uaUgagcagauagucaacgagg....................................................            1       1       seq
    ..................uacgagcagauagucaacgagA....................................................            25      1       seq
    ..................uacgagUagauagucaacgagg....................................................            1       1       seq
    ..................uacgagcagauagucaacgGgg....................................................            1       1       seq
    ..................uacgagcagauagCcaacgagg....................................................            1       1       seq
    ..................uacgagcagauagucaacgagg....................................................            84      0       seq
    ..................uacgagcagauagucaacgagC....................................................            1       1       seq
    ..................uGcgagcagauagucaacgagg....................................................            2       1       seq
    ..................uacgagcagauagucaacgaggA...................................................            2       1       seq
    ..................uacgagcagauagucaacgaggcgcagc..............................................            1       0       seq
    ...................acgagcagauagucaacga......................................................            3       0       seq
    ...................acgagcagauagucaacgaC.....................................................            1       1       seq
    ...................acgagcagauagucaacgaCg....................................................            6       1       seq
    ...................acgagcagauagucaacgaUg....................................................            1       1       seq
    ...................acgagcagauagucaacgagA....................................................            1       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
.............................agauagucaacgaggcgcagU...	1	1	seq
....cucguagaaaacAagggcgu......................	1	1	seq
....ucguagaaaacgagggcgu......................	1	0	seq
....uagaaaacgagggcgugu....................	1	0	seq