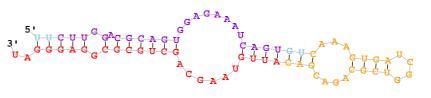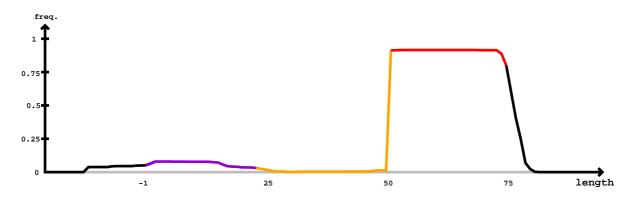Provisional ID   : NW_025322619.1_Acropora_millepora_isolate_JS-1_unplaced_genomic_scaffold_Amil_v2.1_Sc0000050_whole_genome_shotgun_sequer
Score total   :   3
Score for star read(s) :   -1.3
Score for read counts :   0
Score for mfe   :   1.3
Score for randfold :   0
Score for cons. seed :   3
Total read count   :   1333
Mature read count   :   1278
Loop read count   :   6
Star read count   :   49

Star   **Mature**

```
5'- ucucgaauagcgugguauuuucuuggacgcaguggagaaaucagugucaaagugaucggucgcagacgacauuguaagcagcugcgcggagggauggagguuccaugccaguua  -3'    obs
    ucucgaauagcgugguauuuucuuggacgcaguggagaaaucagugucaaagugaucggucgcagacgacauuguaagcagcugcgcggagggauggagguuccaugccaguua         exp
    .......((((.(((((.(..(((.(.((((((.........((((((...((((....))))....)))))))......)))))).)))..)..((.....)).))))))))   reads    mm        sample
    ......uagcgugguauuuucuuC.................................................................................................    2      1         seq
    ......uagcgugguauuuucuuggUc..............................................................................................    1      1         seq
    ......uagcgugguauuuucuuggacgcagu.........................................................................................    2      0         seq
    ......uagcgugguauuuucuuggacgcagC.........................................................................................    1      1         seq
    ......uaUcgugguauuuucuuggacgcagu.........................................................................................    1      1         seq
    ......uagcgugguauuuucuuggacgcagug........................................................................................    2      0         seq
    ......uagcgugguauuuucuuggacgcagugg.......................................................................................   13      0         seq
    ......uagcgugguauuuucuuggacgcagugA.......................................................................................    9      1         seq
    ......uagcgugguauuuucuuggacgcagugga......................................................................................   15      0         seq
    ......uagcguAguauuuucuuggacgcagugga......................................................................................    1      1         seq
    ......uagcgugguauuuucuuggacgcaguggU......................................................................................    1      1         seq
    ......uagcgugguauuuucuuggacgcaguggag.....................................................................................    2      0         seq
    ......uagcgugguauuuucuuggacgcaguggaU.....................................................................................    1      1         seq
    ......uagcgugguauuuucuuggacgcaguggaA.....................................................................................    1      1         seq
    ......uagcgugguauuuucuuggaUgcaguggag.....................................................................................    1      1         seq
    ........agcgugguauuuucuuggacgcagug......................................................................................    1      0         seq
    ...........ugguauuuucuuggacgcagC........................................................................................    1      1         seq
    ...........ugguauuCucuuggacgcaguggaga...................................................................................    1      1         seq
    ...........ugguauuuucuuggacgcaguggagaa..................................................................................    5      0         seq
    ...........ugguauuuucuuggacgcaguggagaaau................................................................................    1      0         seq
    ............gguauuuucuuggacgcaguggagaa..................................................................................    2      0         seq
    ...............uuuucuuggacgcaguggagaaauc................................................................................    1      0         seq
    ...............uuuucuuggacgcaguggagaaaucagu.............................................................................    3      0         seq
    ...............uuuucuuggacgcaguggagaaaucaguA............................................................................    1      1         seq
    ...............uuuucuuggacgcaguggagaaaucaguU............................................................................    1      1         seq
    ................uuucuuggacgcaguggagaaauc................................................................................    2      0         seq
    ................uuucuuggacgcaguggagaaaucaUuguca.........................................................................    1      1         seq
    .................uucuuggacgcaguggagaaauc................................................................................    2      0         seq
    ..................ucuuggacgcaguggagaaaucag..............................................................................    5      0         seq
    ..................ucuuggacgcaguggagaaaucagu.............................................................................    5      0         seq
    ..................ucuuggacgcaguggagaaaucaguA............................................................................    1      1         seq
    ..................ucuuggacgcaguggagaaaucagugu...........................................................................    1      0         seq
    ..................ucuuggacgcaguggagaaaucaguguc..........................................................................    1      0         seq
```

ucucgaauagcgugguauuuucuuggacgcaguggagaaaucagugucaaagugaucggucgcagacgacauuguaagcagcugcgcggagggauggagguuccaugccaguua

Sequence	Count 1	Count 2	
....................ucuuggacgcaguggagaaaucaguguUa..	1	1	seq
....................ucuuggacgcaguggagaaaucaUuguca...	2	1	seq
....................ucuuggacgcaguggagaaaucaguguca...	1	0	seq
...................cuuggacgcaguggagaaauc..	1	0	seq
...................cuuggacgcaguggagaaaucagC...	3	1	seq
...................cuuggacgcaguggagaaGucagu...	1	1	seq
...................cuuggacgcaguggagaaaucagu...	12	0	seq
...................cuuggacgcaguggagaaaucagA...	1	1	seq
...................cuuggacgcaguggagaaaucagug..	1	0	seq
...................cuuggacgcaguggagaaaucaguguca..	1	0	seq
.....................gacgcaguggUgaaaucagu..	1	1	seq
....agugaucggucgcagacgac...	3	0	seq
....agugaucggucgcagacgaU...	1	1	seq
....gugaucggucgcagacgac..	1	0	seq
....ugaucggucgcagacgac..	1	0	seq
....gcagacgacauuguaagcagcugUgcgg.....................................	1	1	seq
....cgacauuguaagcagcugUgcggaggg................................	2	1	seq
....cgacauuguaagcagcugUgcggagggau..............................	1	1	seq
....gacauuguaagcagcugUgcggaggga...............................	1	1	seq
....gacauuguaagcagcugUgcggagggau..............................	4	1	seq
....gacauuguaagcagcugUgcggagggaug.............................	3	1	seq
....cauuguaagcagcugUgcggagggauggag.........................	1	1	seq
....auuguaagcagcugcgcA......................................	1	1	seq
....auuguaagcagcugUgcggaggg................................	36	1	seq
....auuguaagcagcugcgcggaggga..............................	1	0	seq
....auuguaagcagcugUgcggaggga..............................	121	1	seq
....auuguaagcagcugcgcggagggau.............................	4	0	seq
....auuguaagcagcugUgcggagggau.............................	439	1	seq
....auuguaagcagcugUgcggagggaug............................	105	1	seq
....auuguaagcagcugcgcggagggaug............................	2	0	seq
....auuguaagcagcugUgcggagggaugg...........................	217	1	seq
....auuguaagcagcugcgcggagggaugC...........................	1	1	seq
....auuguaagcagcugcgcggaggAaugga..........................	1	1	seq
....auuguaagcagcugUgcggagggaugga..........................	255	1	seq
....auuguaagcagcugcgcggagggaugga..........................	1	0	seq
....auuguaagcagcugUgcggagggauggag.........................	64	1	seq
....auuguaagcagcugUgcggagggauggagg........................	29	1	seq
....uguaagcagcugUgcggagggauggaggu.......................	3	1	seq