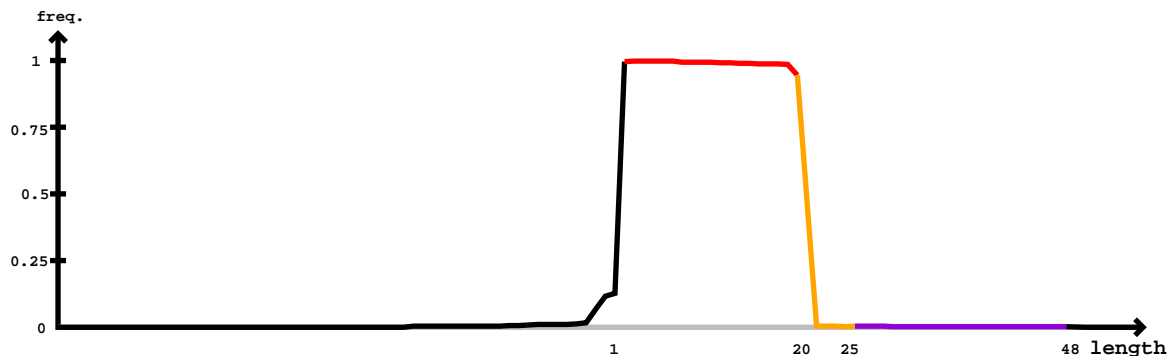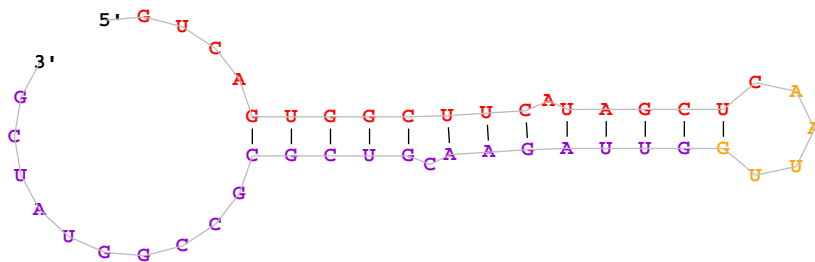|		
Provisional ID	:	NC_058074.1_Acropora_millepora_isolate_JS-1_chromosome_9_Amil_v2.1_whole_genome_shotgun_sequence_373046
Score total	:	223
Score for star read(s)	:	3.9
Score for read counts	:	217.8
Score for mfe	:	-1.6
Score for randfold	:	0
Score for cons. seed	:	3
Total read count	:	439
Mature read count	:	438
Loop read count	:	0
Star read count	:	1


```
                                                                                                                      obs
5'- cacuucacauucauuccucacgggcccauugcaacucacaaaugaccagcucccaacgucaguggcuucauagcucaauugguuagaacgucgcgccgguaucgcgagguc  -3'
    cacuucacauucauuccucacgggcccauugcaacucacaaaugaccagcucccaacgucaguggcuucauagcucaauugguuagaacgucgcgccgguaucgcgagguc       exp
    ...............(((((.((((..((...((...(((....)))...))..........(((((((((.(((((.....))))))))).)))))...))..))).))))).. reads   mm   sample
    .....................................ucacaaaugaccagcuGccaacgucagu............................................... 2       1    seq
    ............................................cUagcucccaacgucaguggcuucauagcuc.................................... 1       1    seq
    ....................................................agUucccaacgucaguggcu...................................... 1       1    seq
    ....................................................Ccucccaacgucaguggcuuc..................................... 1       1    seq
    .......................................................ccaacgucagugAcuucauagcuc.............................. 1       1    seq
    ........................................................caacgucaguggcuucauagcuA............................. 2       1    seq
    .........................................................aacgucaguggcuucau................................. 1       0    seq
    .........................................................Gacgucaguggcuucauagc.............................. 1       1    seq
    .........................................................aacgucagugAcuucauagcu.............................. 2       1    seq
    .........................................................aacgucaguggcuucauagcuA............................ 3       1    seq
    .........................................................aacgucaguggcuucauagcuc............................ 13      0    seq
    .........................................................aacgucaguggcuucauagUuc............................ 1       1    seq
    .........................................................aacgucaguggcuucauagcuca........................... 3       0    seq
    ..........................................................acgucaguggcuucauagcuA............................ 1       1    seq
    ..........................................................acgucagugAcuucauagcuc............................ 2       1    seq
    ..........................................................Gcgucaguggcuucauagcuc............................ 2       1    seq
    ..........................................................acgucaguggcuucauagcuc............................ 12      0    seq
    ..........................................................aGgucaguggcuucauagcuc............................ 2       1    seq
    ..........................................................acgucaguggcuucauagcucU........................... 1       1    seq
    ..........................................................acgucaguggcuucauagcuca........................... 3       0    seq
    ...........................................................cgucaguggcuucauagcu............................ 2       0    seq
    ...........................................................cgucagugAcuucauagcu............................ 1       1    seq
    ...........................................................cgucaguggcuucauagcuA........................... 1       1    seq
    ...........................................................Ggucaguggcuucauagcuc........................... 1       1    seq
    ...........................................................gucaguggcuucauagcu............................. 14      0    seq
    ...........................................................gucaguggcuucGuagcuc............................ 2       1    seq
    ...........................................................gCcaguggcuucauagcuc............................ 1       1    seq
    ...........................................................gucaguggcuucauaUcuc............................ 1       1    seq
    ...........................................................gucagugAcuucauagcucU........................... 32      1    seq
    ...........................................................gucaguggcuucauagUuc............................ 10      1    seq
    ...........................................................gucaguggcuucauagcuc............................ 285     0    seq
    ...........................................................gucaguggcuucaGagcuc............................ 3       1    seq
    ...........................................................gucaguggcuCcauagcuc............................ 5       1    seq
```

cacuucacauucauuccucacgggcccauugcaacucacaaaugaccagcucccaac**gucaguggcuucauagcuc**aauug**guuagaacgucgcgccgguaucg**cgagguc

....gucagCggcuucauagcuc.....................................	3	1	seq
....gAcaguggcuucauagcuc.....................................	10	1	seq
....gucaguggcCucauagcuc.....................................	2	1	seq
....gucaguggcuucauagcuA...................................	32	1	seq
....gucaguggcuucauagcuca..................................	8	0	seq
....gucaguggcuucauagcucaauu.............................	1	0	seq
....gucaguggcuucauagcucaCuugguua.........................	1	1	seq
....ucagugAcuucauagcuc......................................	1	1	seq
....guuagaacgucgcAccgguaucg.......	1	1	seq