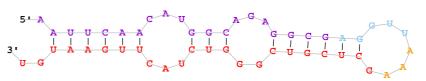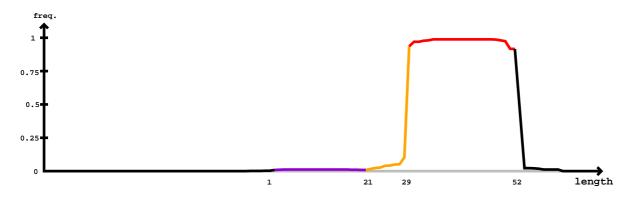|		
Provisional ID	:	NC_058066.1_Acropora_millepora_isolate_JS-1_chromosome_1_Amil_v2.1_whole_genome_shotgun_sequence_14067
Score total	:	318
Score for star read(s)	:	3.9
Score for read counts	:	311.6
Score for mfe	:	-0.5
Score for randfold	:	0
Score for cons. seed	:	3
Total read count	:	623
Mature read count	:	617
Loop read count	:	0
Star read count	:	6


```
Star          Mature
5'- guacauagaggccacuuuauuuauaagggaaggagaacucuuuucgaauucaacauggcagaggcgagguuaaagcucgucgggucuacuugaauguucauucaguaauaacag  -3'   obs
    guacauagaggccacuuuauuuauaagggaaggagaacucuuuucgaauucaacauggcagaggcgagguuaaagcucgucgggucuacuugaauguucauucaguaauaacag         exp
    (((.(((((((....))))))).)))...(((((((....)))))))..(((((...(((...(((((((.......))))))..)))...)))))).............       reads   mm   sample
    ................................................uuucgaauucaacGuggcaga.................................................   1    1    seq
    ................................................cgaauucaacauggcagaggcgaggu............................................   1    0    seq
    ................................................aauucaacauggcagagg.....................................................   1    0    seq
    ................................................aauucaacauggcagaggcg...................................................   2    0    seq
    ................................................aauucaacauggcagaggcgU..................................................   1    1    seq
    .................................................auucaacauggcagaggcgaggu...............................................   1    0    seq
    .................................................uucaacauggcagaggcgU...................................................   1    1    seq
    ......................................................agguuaaagcucgucgggucuacuugaau....................................   1    0    seq
    ......................................................agguuaaagcucgucgggucuacuugaaugu..................................   3    0    seq
    ......................................................aggCuaaagcucgucgggucuacuugaaugu..................................   1    1    seq
    ......................................................agguuaaagcucgucgggucuacuugaaugC..................................   3    1    seq
    .......................................................gguuaaagcuUgcgggucuacuugaaugu...................................   1    1    seq
    .......................................................gguuaaagcucgucgggucuacuugaaugu...................................  5    0    seq
    ........................................................guuaaagcucgucgggucuacuugaaugu...................................  2    0    seq
    .........................................................uuaaagcucgucgggucuacuugaaugu...................................  5    0    seq
    .........................................................uuaaagcucgucgggucuacuugaaugC...................................  2    1    seq
    .........................................................uuaaagcucgucgCgucuacuugaaugu...................................  1    1    seq
    .........................................................uuaaUgcucgucgggucuacuugaaugu...................................  1    1    seq
    ..........................................................uaaagcucgucgggucuacuugaaC.....................................  1    1    seq
    ..........................................................Caaagcucgucgggucuacuugaaugu...................................  2    1    seq
    ...........................................................aaagcucgucCggucuacuugaau.....................................  1    1    seq
    ...........................................................aaagcucgucgggucuacuugaaC.....................................  1    1    seq
    ...........................................................aaagcucgucgggucuacuugaaugu...................................  2    0    seq
    ...........................................................Gaagcucgucgggucuacuugaaugu...................................  1    1    seq
    ............................................................aagcucgucgggucuacCugaaugu...................................  1    1    seq
    .............................................................agcucgucgggucuacuug.......................................  1    0    seq
    .............................................................agcucgucgggucuacuuga......................................  1    0    seq
    .............................................................agcucgucgggucuacuugaa.....................................  3    0    seq
    .............................................................agcucgucgggucuacuugaU.....................................  1    1    seq
    .............................................................agcucgucgggucuacuugaau....................................  2    0    seq
    .............................................................agcucgucgggucuacuugaaugu..................................  23   0    seq
    .............................................................agcucgucggAucuacuugaaugu..................................  2    1    seq
    .............................................................agcucgucgggucuacuugaaugA.................................  1    1    seq
```

guacauagaggccacuuuauuuauaagggaaggagaacucuuuucgaauucaacauggcagaggcgagguuaaagcucgucgggucuacuugaauguucauucaguaauaacag

Sequence	Count	Mismatch	Type
....agcucgucgggucuacuugaaugG................	1	1	seq
....gcucgucgggucuacuugU....................	3	1	seq
....gcucgucgggucuacuugaaC...................	4	1	seq
....gcucAucgggucuacuugaau...................	2	1	seq
....gcucgucgggucuacuugaUu...................	1	1	seq
....gcucgucgggucuacuugaau...................	24	0	seq
....gcucgucgggucuacuugaauC..................	1	1	seq
....Acucgucgggucuacuugaaugu.................	9	1	seq
....gcucgucgggucuacuGgaaugu.................	1	1	seq
....gcucgucgggucuacuugaaGgu.................	2	1	seq
....gcucgucgggucuacAugaaugu.................	3	1	seq
....gcGcgucgggucuacuugaaugu.................	4	1	seq
....gcucguUgggucuacuugaaugu.................	1	1	seq
....gcucgucgggucuacuugaUugu.................	6	1	seq
....gcucgucgggucuacuugaaCgu.................	1	1	seq
....gcucgucgggucuacuCgaaugu.................	2	1	seq
....gcucgucgCgucuacuugaaugu.................	1	1	seq
....gcucgucgggucuacuugaaugG.................	12	1	seq
....gcucgucggAucuacuugaaugu.................	21	1	seq
....gcucgucgggucuacuugaaugA.................	17	1	seq
....gcucgucgggucuacuugaaugu.................	430	0	seq
....gcucgucgggucCacuugaaugu.................	1	1	seq
....gcucgucgggucuacuugGaugu.................	2	1	seq
....gcucgucgggucuacuugaauguu................	1	0	seq
....gcucgucgggucuacuugaauguuca..............	1	0	seq
....gcucgucgggucuacuugaauguucaC.............	1	1	seq
....gcucgucgggucuacuugaauguucau.............	1	0	seq
....gcucgucgggucuacuugaauguucUuu............	1	1	seq
....gcucgucgggucuacuugaauguucauu............	1	0	seq
....cucgucgggucuacuugaau...................	1	0	seq
....cucgucgggucuacuugaaugu.................	14	0	seq
....cucgucggAucuacuugaaugu.................	1	1	seq
....cucgucgggucuacuugaauguucauucagu........	7	0	seq
....cgucggAucuacuugaaugu.................	5	1	seq
....Aucgggucuacuugaaugu................	1	1	seq
....gucgggucuacuugaauguucauucagC.......	1	1	seq
....ucgggucuacuugaaugu................	2	0	seq
....ucgggucuacuugaauguu..............	1	0	seq
....ucgggucuacuugaauguucauu..........	2	0	seq