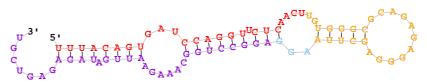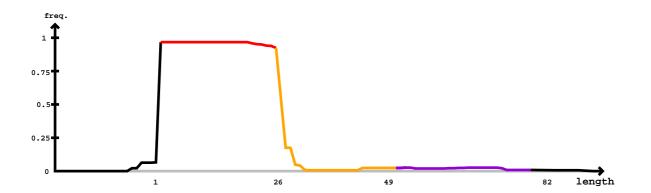| Provisional ID	: NC_058078.1_Acropora_millepora_isolate_JS-1_chromosome_13_Amil_v2.1_whole_genome_shotgun_sequence_485838
Score total	: 0
Score for star read(s)	: -1.3
Score for read counts	: 0
Score for mfe	: -1.6
Score for randfold	: 0
Score for cons. seed	: 3
Total read count	: 416
Mature read count	: 412
Loop read count	: 3
Star read count	: 1


```
        Mature                              Star
5'- ucguggauggccagagaugauuuacagugauccagguucucaacuuguggggcgcagagagggagcuuaaggaggccuggcaaagaauugauagagagucguuacaaagagau  -3'   obs
    ucguggauggccagagaugauuuacagugauccagguucucaacuuguggggcgcagagagggagcuuaaggaggccuggcaaagaauugauagagagucguuacaaagagau       exp
    ...(((....)))..(((..((((((((((...((((((.(((......(((((...........)))))..))))))))))......))))..))))..)))............     reads   mm      sample
```

	reads	mm	sample
.............Ggaugauuuacagugauccagguuc...	1	1	seq
.............agaugauuuacagugauccagguuc...	1	0	seq
.............agaugauuuacagugauccagguucuc...	1	0	seq
.............agaugauuuacagugauccagguucuca..	1	0	seq
.............agaugauuuacagugauccagguucucaaU..	2	1	seq
.............agaugauuuacagugauccagguucucaac..	2	0	seq
.............agaugauuuacagugauccagguucucaacC...	1	1	seq
.............agaugauuuacagugauccagguucucaacu...	1	0	seq
...............augauuuacagugauccagguuc...	2	0	seq
...............augauuuacagugauccagguucC..	1	1	seq
...............augauuuacagugauccagguucA..	2	1	seq
...............augauuuacagugauccagguucucaacu...	5	0	seq
...............augauuuacagugauccagguucucaacA...	1	1	seq
...............augauuuacagugauccagguucucaacuA..	1	1	seq
...............augauuuacagugauccagguucucaacuu..	7	0	seq
.................auuuacagugauccagguucucaacCu...	1	1	seq
..................uuuacagugauccagguucuca...	3	0	seq
..................uuuacagugauccagguucucaa..	1	0	seq
..................uuuacagugauccagguucucaaU...	1	1	seq
..................uuuacagugauccagguucucaac...	2	0	seq
..................uuuacagugauccagguucucaacC..	37	1	seq
..................uuuacagugauccagguucucaacG..	3	1	seq
..................uuuacagugauccaggCucucaacu..	2	1	seq
..................uuuacagugauccagguucucaGcu..	2	1	seq
..................uuuacagugauccagguucucaacu..	194	0	seq
..................uuuacagugauccagguucucaacA..	23	1	seq
..................uAuacagugauccagguucucaacu..	1	1	seq
..................Auuacagugauccagguucucaacu..	1	1	seq
..................Cuuacagugauccagguucucaacu..	1	1	seq
..................uuuacagugaCccagguucucaacu..	3	1	seq
..................uuuacagugauccagguucucGacu..	1	1	seq
..................uuuacagugauccagguucucGacuu...	1	1	seq
..................uuuacagugauccagguucucaacuu...	40	0	seq

Mature   **Star**

```
ucguggauggccagagaugauuuacagugauccagguucucaacuuguggcgcagagagggagcuuaaggaggccuggcaaagaauugauagagagucguuacaaagagau
```

Sequence	Count		
`....................uuuacagugauccagguucucaacuC............................................................`	6	1	seq
`....................uuuacagugauccagguCcucaacuu............................................................`	3	1	seq
`....................uuuacGgugauccagguucucaacuu............................................................`	1	1	seq
`....................uuuacagugauccagguucucaacuA............................................................`	4	1	seq
`....................uuuacagugauccagguucucaacuG............................................................`	2	1	seq
`....................uuuacagugauccagguucucaacuuU...........................................................`	2	1	seq
`....................uuuacagugauccagguucucaacuug...........................................................`	1	0	seq
`....................uuuacagugauccagguucucaacCugu..........................................................`	1	1	seq
`....................uuuacagugauccGgguucucaacuugu..........................................................`	1	1	seq
`....................uuuacagugauccagguucucaacuugA..........................................................`	1	1	seq
`....................uuuacagugauccagguucucaacuugC..........................................................`	8	1	seq
`....................uuuacagugauccagguucucaacuugu..........................................................`	46	0	seq
`....................uuuacagugauccagguucucaacuugG..........................................................`	1	1	seq
`....................uuuacagugauccagguucucaacuugug.........................................................`	3	0	seq
`....................uuuacagugaCccagguucucaacuugugg........................................................`	1	1	seq
`....................uuuacagugauccagguucucaacuugugC........................................................`	1	1	seq
`....................uuuacagugauccagguucucaacuugugg........................................................`	9	0	seq
`....................uuuacagugauccagguucucaacuugugA........................................................`	4	1	seq
`....................uuuacagugauccagguucucaacuuguggA.......................................................`	1	1	seq
`.................................................uugggcgcagagagggagcuuaaggag..............................`	1	0	seq
`.................................................uugggcgcagagagggagcuuaaggaA..............................`	2	1	seq
`.......................................................agcuuaaggaggccuggcaaagaauugau.....................`	2	0	seq
`.......................................................agcuuaaggaggccuggcaaagaauugaua....................`	4	0	seq
`.......................................................agcuuaaggaggccuggcaaagaauugaCa....................`	2	1	seq
`...................................................................aggccuggcaaagaauugauagagagucgG...........`	1	1	seq
`...........................................................................aaagaauugauagagagucguuacaaaga...`	1	0	seq
`.............................................................................agaauugauagagagucguuacaGagag..`	1	1	seq
`..............................................................................aauugauagagagucguuacaaag....`	1	0	seq