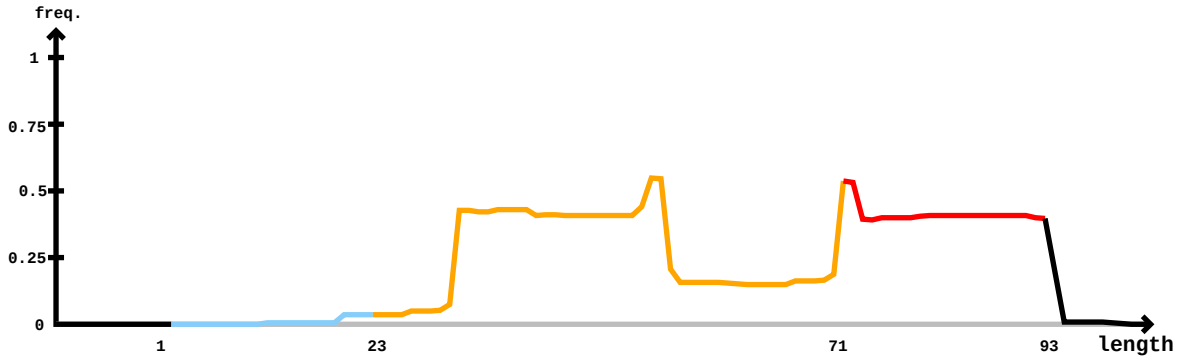
Provisional ID	:	NC_058079.1_Acropora_millepora_isolate_JS-1_chromosome_14_Amil_v2.1_whole_genome_shotgun_sequence_517732
Score total	:	1
Score for star read(s)	:	-1.3
Score for read counts	:	0
Score for mfe	:	2.9
Score for randfold	:	0
Score for cons. seed	:	-0.6
Total read count	:	194
Mature read count	:	140
Loop read count	:	54
Star read count	:	0



Star Mature

```
5'- ugcgauggcaaguuauaacuugcgauugcgacuuauaacuugcgaucgcaacuuauaacuugcgaucgcaaguuauaacuugcgaucgcaacuuauaacuugcgaucgcaac  -3'   exp
    ((((((.(((((((((((.((((((((((((.(((((((((((((((((((.........)))))))))))))))))))).))))))))))))).))))))))))).))))))..  reads   mm   sample
    ....................ugcgauugcgacuuauaacuug.......................................................................  2       0    seq
    ............................cgacuuauaacuugcgaucU.......................................................................  1       1    seq
    ............................cgacuuauaacuugcgaucA.......................................................................  9       1    seq
    ............................cgacuuauaacuugcgaucgcaU....................................................................  1       1    seq
    ..................................uaacuugcgaucgcaacuuauaacuug.............................................................  4       0    seq
    ..................................uaacuugcgaucgcaacuuauaacuugc............................................................  6       0    seq
    ....................................cuugcgaucgcaacuuauaacuA...............................................................  2       1    seq
    ....................................uugcgaucgcaacuuauGacuug...............................................................  4       1    seq
    ....................................uugcgaucgcaacuuauaacuug...............................................................  8       0    seq
    ....................................uugcgaucgcaacuuauaacuugc..............................................................  4       0    seq
    ....................................ugcgaucgcaacuuauaacu..................................................................  6       0    seq
    ....................................ugcgaucgcaacuuauaacuuU................................................................  6       1    seq
    ....................................ugcgaucgcaacuuauaacAug................................................................  2       1    seq
    ....................................ugcgaucgcaacuuauaacuAg................................................................  6       1    seq
    ....................................ugcgaucgcaacuuauaacuug................................................................  206     0    seq
    ....................................ugcgaucgcaacuuauaacuCg................................................................  2       1    seq
    ....................................Cgcgaucgcaacuuauaacuug................................................................  2       1    seq
    ....................................ugcgaucgcaacuuauaacuugc...............................................................  26      0    seq
    ...................................aucgcaacuuauaacuug.....................................................................  6       0    seq
    ..................................................caacuuauaacuugcgaucA.....................................................  2       1    seq
    ..................................................caacuuauaacuugcgaucgc....................................................  2       0    seq
    ..................................................aacuuauaacuugcgaucgca....................................................  2       0    seq
    ..................................................................uugcgaucgcaaguuauaacuu...............................  2       0    seq
    ..................................................................uCgcgaucgcaaguuauaacuug..............................  2       1    seq
    ..................................................................uugcgaucgcaaguuauaacuuA..............................  2       1    seq
    ..................................................................uugcgaucgcaaguuauaacuCg..............................  1       1    seq
    ..................................................................uugcgaucgcaaguuauaacuug..............................  3       0    seq
    ..................................................................uugcgaucgcaaguuauaacuuC..............................  1       1    seq
    ..................................................................uugcgaucgcaaguuauaacuugc.............................  1       0    seq
    ..................................................................ugcgaucgcaaguuauaacA..................................  1       1    seq
    ..................................................................ugcgaucgcaaguuGuaacuug................................  2       1    seq
    ..................................................................ugcgaucgcaaguuauaacuug................................  28      0    seq
    ..................................................................ugcgaucgcaaguuauGacuug................................  1       1    seq
    ..................................................................ugcgaucgcaaguuauaacuuA................................  6       1    seq
```

Star Mature

ugcgauggcaaguuauaacuugcgauugcgacuuauaacuugcgaucgcaacuuauaacuugcgaucgcaaguuauaacuugcgaucgcaacuuauaacuugcgaucgcaac

..........ugcgaucgcaaguuauaacuuU..........	3	1	seq
..........uCcgaucgcaaguuauaacuug..........	1	1	seq
..........uaacuugcgaucgcaacuuauaacuug.....	4	0	seq
..........uaacuugcgaucgcaacuuauaacuugc....	6	0	seq
..........cuugcgaucgcaacuuauaacuA.........	2	1	seq
..........uugcgaucgcaacuuauaacuug.........	8	0	seq
..........uugcgaucgcaacuuauGacuug.........	4	1	seq
..........uugcgaucgcaacuuauaacuugc........	4	0	seq
..........ugcgaucgcaacuuauaacu............	6	0	seq
..........ugcgaucgcaacuuauaacuCg.........	2	1	seq
..........ugcgaucgcaacuuauaacuug.........	206	0	seq
..........ugcgaucgcaacuuauaacuuU.........	6	1	seq
..........ugcgaucgcaacuuauaacAug.........	2	1	seq
..........ugcgaucgcaacuuauaacuAg.........	6	1	seq
..........Cgcgaucgcaacuuauaacuug.........	2	1	seq
..........ugcgaucgcaacuuauaacuugc........	26	0	seq
..........aucgcaacuuauaacuug.............	6	0	seq
..........caacuuauaacuugcgaucA....	2	1	seq
..........caacuuauaacuugcgaucgc...	2	0	seq
..........aacuuauaacuugcgaucgca..	2	0	seq