Provisional ID	: NW_025322770.1_Acropora_millepora_isolate_JS-1_unplaced_genomic_scaffold_Amil_v2.1_xfSc0000077_whole_genome_shotgun_sequ
Score total	: 0.2
Score for star read(s)	: -1.3
Score for read counts	: 0
Score for mfe	: 2.1
Score for randfold	: 0
Score for cons. seed	: -0.6
Total read count	: 355
Mature read count	: 353
Loop read count	: 0
Star read count	: 2


```
5'- auguaguacaauucagggaguaaucaggcucguaauuucaaaucauuucugaaauuacucgcccgauuacucccugaauuguucuccacucgguccaauuacuacuacu -3'   obs
    auguaguacaauucagggaguaaucaggcucguaauuucaaaucauuucugaaauuacucgcccgauuacucccugaauuguucuccacucgguccaauuacuacuacu       exp
    ..(((((((((((((((((((((((.(((..(((((((((.........)))))))))..))).)))))))))))))))))))))....((((..))......)))))).......   reads   mm   sample
    .....uacaauucagggaguaaAca.....................................................................................   1    1    seq
    .................uaaucUggcucguaauuuca.........................................................................   1    1    seq
    .................uaaucUggcucguaauuucaa........................................................................   1    1    seq
    .................uaaucUggcucguaauuucaaauca....................................................................   1    1    seq
    ..................ucGggcucguaauuucaaauca......................................................................   3    1    seq
    ...........................................ugaaauuacucgcccgauuacu.............................................   10   0    seq
    ...........................................ugaaauuacucgcccgauuacC.............................................   2    1    seq
    ............................................aaauuacucgcccgauuacu..............................................   1    0    seq
    ............................................aaauuacucgcccgauuacucA............................................   1    1    seq
    ............................................aaUuuacucgcccgauuacucc............................................   1    1    seq
    ............................................aaauuacucgcccgauuacucc............................................   1    0    seq
    ............................................aaauuacucgcccgauuacuccU...........................................   1    1    seq
    .............................................aauuacucgcccgauuacucA...........................................   2    1    seq
    .............................................aauuacucgcccgauuacuccc..........................................   1    0    seq
    ..............................................auuacucgcccgauuacucA..........................................   1    1    seq
    ...............................................uuacucgcccgauuacuccA.........................................   1    1    seq
    ...............................................uuacucgcccgauuacuccc.........................................   1    0    seq
    ...............................................uuacucgcccgauuacucccC........................................   3    1    seq
    ...............................................uuacucgcccgauuacucccu........................................   2    0    seq
    ...............................................uuacucgcccgauuacucccug.......................................   13   0    seq
    ...............................................uuacucgcccgauuacucccuga......................................   22   0    seq
    ...............................................uuacucgcccgauuacucccugG......................................   1    1    seq
    ...............................................uuacucgcccgauuacucccugU......................................   52   1    seq
    ...............................................uuacucgcccgauuacucccugC......................................   9    1    seq
    ...............................................uuacucgcccgauuacucccugaU.....................................   9    1    seq
    ...............................................uuacucgcccgauuacucccugaUu....................................   3    1    seq
    ...............................................uuacucgcccgauuacucccugaauu...................................   1    0    seq
    ................................................uacucgcccgauuacucc.........................................   1    0    seq
    ................................................uacucgcccgauuacuccc........................................   2    0    seq
    ................................................uacucgcccgauuacuccU........................................   1    1    seq
    ................................................uacucgcccgauuacucccu.......................................   1    0    seq
    ................................................uacucgcccgauuacucccA.......................................   1    1    seq
    ................................................uacucgcccgauuacucccCg......................................   2    1    seq
```

auguaguacaauucagggaguaaucaggcucguaauuucaaaucauuucugaaauuacucgcccgauuacucccugaauuguucuccacucgguccaauuacuacuacuacu

Sequence	Count	Mismatch	Type
....uacucgcccgauCacucccug.....................................	1	1	seq
....uacucgcccgauuacucccug.....................................	11	0	seq
....uacucgcccgauuacucccuC.....................................	3	1	seq
....uacucgcccgauuacucccugG....................................	1	1	seq
....uacucgcccgauuacuccUuga...................................	2	1	seq
....uacucgcccgauuacucccugU....................................	8	1	seq
....uacucgcccgauuacucccuga....................................	38	0	seq
....uacucgcccgauuacucccugaa...................................	14	0	seq
....uacucgcccgauuacucccugaU...................................	11	1	seq
....uacucgcccgauuacucccugaC...................................	3	1	seq
....uacucgcccgauuacucccugaUu..................................	4	1	seq
....uacucgcccgauCacucccugaau..................................	1	1	seq
....uacucgcccgauuacucccUUgaau.................................	1	1	seq
....uacucgcccgauuacucccugaCu..................................	1	1	seq
....uacucgcccgauuacucccugaaA..................................	2	1	seq
....uacucgcccgauuacucccugaau..................................	60	0	seq
....uacucgcccgauuacucccugaauu.................................	16	0	seq
....uacucgcccgauuacucccugaauA.................................	1	1	seq
....uacucgcccgauuacucccugaauC.................................	1	1	seq
....acucgcccgauuacuccUug...................................	2	1	seq
....acucgcccgauuacuccUuga..................................	1	1	seq
....acucgcccgauuacuccUugaa.................................	1	1	seq
....acucgcccgauuacuccUugaau................................	3	1	seq
....cuUgcccgauuacucccuga................................	3	1	seq
....cucgcccgauuacucccuga................................	1	0	seq
....cuUgcccgauuacucccugaa...............................	1	1	seq
....cucgcccgauuacucccugaC...............................	1	1	seq
....cucgcccgauuacucccugaau..............................	7	0	seq
....cucgcccgauuacuccUugaau..............................	2	1	seq
....cuUgcccgauuacucccugaau..............................	3	1	seq
....cuUgcccgauuacucccugaauu.............................	1	1	seq
....cucgcccgauuacucccugaauu.............................	2	0	seq
....ucgcccgauuacucccug.................................	1	0	seq
....ucgcccgauuacuccUugaau..............................	1	1	seq
....ucgcccgauuacucccugaauugG...........................	1	1	seq
....Ugcccgauuacucccugaau.............................	2	1	seq
....Ugcccgauuacucccugaauu............................	3	1	seq
....cccgauuacucccugaauugu.........................	3	0	seq
....ccgauuacucccugaauugu.......................	9	0	seq