Provisional ID	: NW_025322620.1_Acropora_millepora_isolate_JS-1_unplaced_genomic_scaffold_Amil_v2.1_Sc0000055_whole_genome_shotgun_sequer
Score total	: 3.2
Score for star read(s)	: -1.3
Score for read counts	: 0
Score for mfe	: 1.5
Score for randfold	: 0
Score for cons. seed	: 3
Total read count	: 270
Mature read count	: 270
Loop read count	: 0
Star read count	: 0

Star   **Mature**

```
5'- ggccggcgccgcuuaugcauugucugcucacaacggcagccacgucgcagauggccagccugcacagacucagauugcuggcuuuugauguuaugguaauguucgcugguaaga  -3'    exp
    .(((((((((((.....(((.....)))......))))).(((((...(((((((.((((((((((((.......)))...))))))))))).))...)))).......))))))....     reads    mm    sample
    ....................ucugcucacaacggcagccacgucgcagaug........................................................              2        0     seq
    ....................ugcucacaacggcagccacgucgcagaug..........................................................            3        0     seq
    ..........................ucacaacggcagccacgucgcagauggcU...................................................          1        1     seq
    .................................................................acucagauugcuggAuuuugauguuau..................        1        1     seq
    .................................................................ucagauugcuggAuuuugaugu.......................      1        1     seq
    .................................................................ucagauugcuggcuuuugauguuaA...................       2        1     seq
    .................................................................ucagauugcuggcuuuugauguuaC...................       2        1     seq
    .................................................................ucagauugcuggAuuuugauguuau...................       240      1     seq
    .................................................................ucagauugcuggGuuuugauguuau...................       1        1     seq
    .................................................................ucagauugcuggcuuuugauguuau...................       18       0     seq
    .................................................................ucagauugcuggcuuuugauguuauA..................       2        1     seq
    .................................................................ucagauugcuggcuuuugauguuauAgua...............      1        1     seq
    ............................................................................ugcuggAuuuugauguuau...............       2        1     seq
```