| | | |
|---|---|---|
| Provisional ID | : | NW_025322633.1_Acropora_millepora_isolate_JS-1_unplaced_genomic_scaffold_Amil_v2.1_Sc0000098_whole_genome_shotgun_sequer |
| Score total | : | 0.5 |
| Score for star read(s) | : | -1.3 |
| Score for read counts | : | 0 |
| Score for mfe | : | 2.4 |
| Score for randfold | : | 0 |
| Score for cons. seed | : | -0.6 |
| Total read count | : | 30 |
| Mature read count | : | 29 |
| Loop read count | : | 0 |
| Star read count | : | 1 |





```
            Star                              Mature
5'- ugucgaucccaguugauucaaacgucgagcuuugcaugugccgaaccuaauuauuagguucgacacauguaaaguucgacguuugaaccggaucuuaauccuuuauuag  -3'   obs
    ugucgaucccaguugauucaaacgucgagcuuugcaugugccgaaccuaauuauuagguucgacacauguaaaguucgacguuugaaccggaucuuaauccuuuauuag        exp
    ....(((((.....(.((((((((((((((((((((((((.((((((((.....)))))))).)))))))))))))))))))))))))).))))).............    reads   mm      sample
    ...................aacgucgagcuuugcaugu..................................................................            1      0        seq
    .......................................agguucgGcacauguaaagu............................................            1      1        seq
    ...............................................cacauguaaaguucgacguC....................................            1      1        seq
    ...............................................cacauguaaaguucgacgCu....................................            1      1        seq
    ...............................................cacauguaaaguucgacguu....................................           21      0        seq
    ................................................acauguaaaguucgacguu....................................            1      0        seq
    ................................................acauguaaaguucgacguAu...................................            1      1        seq
    .................................................cauguaaaguucgacguuga..................................            1      0        seq
    ....................................................uguaaaguucgacguuugU.................................            1      1        seq
    ....................................................uguaaaguucgacguuuga.................................            2      0        seq
    .......................................................aguucgacguuugaaccggGu...........................            1      1        seq
```