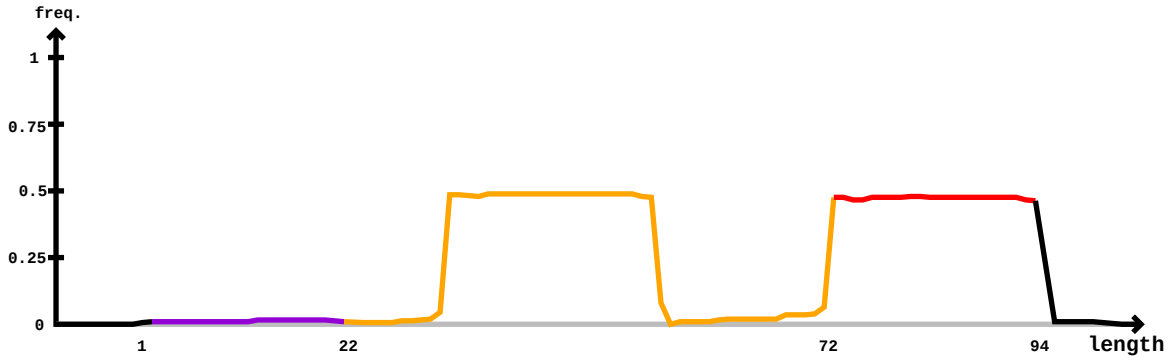
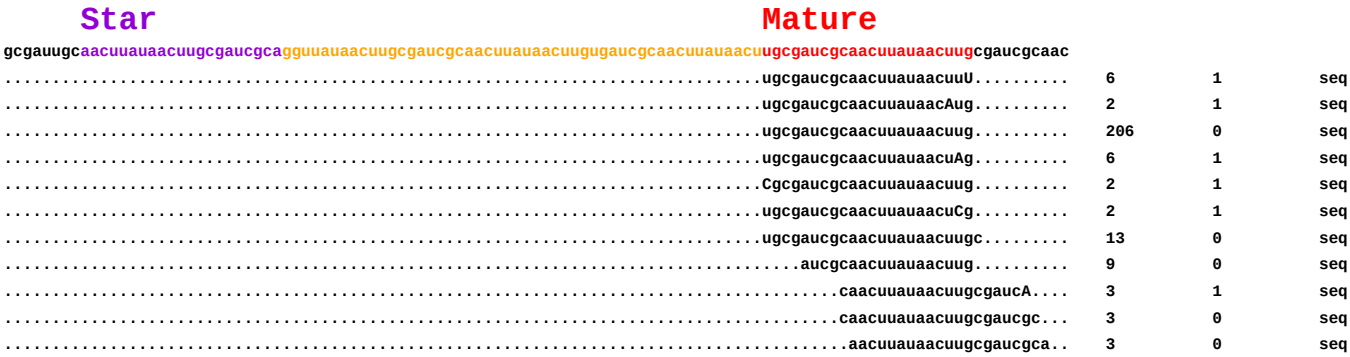
| | | |
|---|---|---|
| Provisional ID | : | NW_025322629.1_Acropora_millepora_isolate_JS-1_unplaced_genomic_scaffold_Amil_v2.1_Sc0000089_whole_genome_shotgun_sequer |
| Score total | : | 84.7 |
| Score for star read(s) | : | 3.9 |
| Score for read counts | : | 78.6 |
| Score for mfe | : | 2.7 |
| Score for randfold | : | 0 |
| Score for cons. seed | : | -0.6 |
| Total read count | : | 166 |
| Mature read count | : | 140 |
| Loop read count | : | 25 |
| Star read count | : | 1 |




**Star** **Mature**

```
5'- gcgauugcaacuuauaacuugcgaucgcagguuauaacuugcgaucgcaacuuauaacuugugaucgcaacuuauaacuugcgaucgcaacuuauaacuugcgaucgcaac  -3'   obs
    gcgauugcaacuuauaacuugcgaucgcagguuauaacuugcgaucgcaacuuauaacuugugaucgcaacuuauaacuugcgaucgcaacuuauaacuugcgaucgcaac        exp
    ((((((((((.((((((.((((((((((((((((((.((((((((((((.........))))))))))))).))))))))))))))))))).)))))).))))))))))...   reads   mm   sample
    .......caacuuauaacuugcgaucA.................................................................................   3       1    seq
    .......caacuuauaacuugcgaucgc................................................................................   3       0    seq
    ........aacuuauaacuugcgaucgca...............................................................................   3       0    seq
    ...................ugcgaucgcagguuauaacuuA.................................................................   1       1    seq
    ...................ugcgaucgcGgguuauaacuugc................................................................   1       1    seq
    ................................uaacuugcgaucgcaacuuauaacuug...................................................   4       0    seq
    ...................................acuugcgaucgcaacuuauaacuugu...................................................   1       0    seq
    ..................................cuugcgaucgcaacuuauaacuA.......................................................   2       1    seq
    ....................................uugcgaucgcaacuuauaacuug.....................................................   8       0    seq
    ....................................uugcgaucgcaacuuauGacuug.....................................................   4       1    seq
    ....................................uugcgaucgcaacuuauaacuugu....................................................   2       0    seq
    .....................................ugcgaucgcaacuuauaacu.......................................................   6       0    seq
    .....................................ugcgaucgcaacuuauaacuug.....................................................   206     0    seq
    .....................................Cgcgaucgcaacuuauaacuug.....................................................   2       1    seq
    .....................................ugcgaucgcaacuuauaacuAg.....................................................   6       1    seq
    .....................................ugcgaucgcaacuuauaacAug.....................................................   2       1    seq
    .....................................ugcgaucgcaacuuauaacuuU.....................................................   6       1    seq
    .....................................ugcgaucgcaacuuauaacuCg.....................................................   2       1    seq
    .....................................ugcgaucgcGacuuauaacuugu....................................................   14      1    seq
    .....................................ugcgaucgcaacuuauaacuuCu....................................................   1       1    seq
    .....................................ugcgaucgcaacuuauaacuugu....................................................   7       0    seq
    ........................................aucgcaacuuauaacuug.......................................................   9       0    seq
    ..................................................................aucgcaacuuauaacuug.........................   9       0    seq
    ...................................................................caacuuauaacuugcgaucA.......................   3       1    seq
    ...................................................................caacuuauaacuugcgaucgc......................   3       0    seq
    ...................................................................aacuuauaacuugcgaucgca......................   3       0    seq
    ..................................................................uaacuugcgaucgcaacuuauaacuug..........   4       0    seq
    ..................................................................uaacuugcgaucgcaacuuauaacuugc.........   3       0    seq
    ...................................................................cuugcgaucgcaacuuauaacuA...........   2       1    seq
    ...................................................................uugcgaucgcaacuuauaacuug..........   8       0    seq
    ...................................................................uugcgaucgcaacuuauGacuug..........   4       1    seq
    ...................................................................uugcgaucgcaacuuauaacuugc.........   2       0    seq
    ...................................................................ugcgaucgcaacuuauaacu............   6       0    seq
```


Star
Mature
gcgauugcaacuuauaacuugcgaucgcaggguuauaacuugcgaucgcaacuuauaacuugugaucgcaacuuauaacuugcgaucgcaacuuauaacuugcgaucgcaac
.....ugcgaucgcaacuuauaacuuU..... 6 1 seq
.....ugcgaucgcaacuuauaacAug..... 2 1 seq
.....ugcgaucgcaacuuauaacuug..... 206 0 seq
.....ugcgaucgcaacuuauaacuAg..... 6 1 seq
.....Cgcgaucgcaacuuauaacuug..... 2 1 seq
.....ugcgaucgcaacuuauaacuCg..... 2 1 seq
.....ugcgaucgcaacuuauaacuugc..... 13 0 seq
.....aucgcaacuuauaacuug..... 9 0 seq
.....caacuuauaacuugcgaucA.... 3 1 seq
.....caacuuauaacuugcgaucgc... 3 0 seq
.....aacuuauaacuugcgaucgca.. 3 0 seq