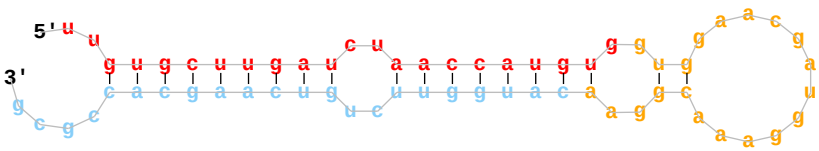
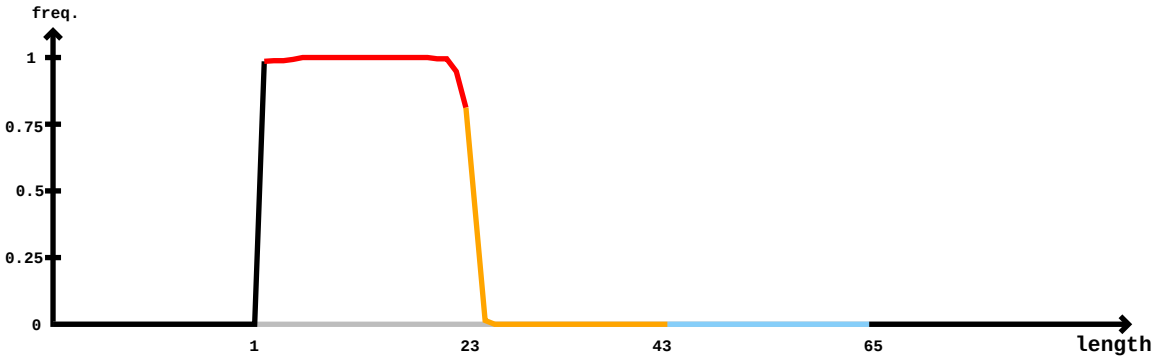
| | | |
|---|---|---|
| Provisional ID | : | RAE_1732_20314 |
| Score total | : | 2.4 |
| Score for star read(s) | : | -1.3 |
| Score for read counts | : | 0 |
| Score for mfe | : | 2.1 |
| Score for randfold | : | 1.6 |
| Score for cons. seed | : | |
| Total read count | : | 420 |
| Mature read count | : | 420 |
| Loop read count | : | 0 |
| Star read count | : | 0 |




**Mature** **Star**

```
5'- ggucgccgcggggcuuuccuuugugcuugaucuaaccaugugguggaacgauggaaacggaacaugguucugucaagcaccgcggaaagcaccgcguucuccugcagcau  -3'   exp
    ((....(((((.(((((((...(((((((((..(((((((((..((............))..))))))))).))))))))).))))))).)))))....)).........  reads     mm        sample
    ....................uugugcuugaucuaacca.....................................................................  2         0         seq
    ....................uugugcuugaucuaaccaug...................................................................  20        0         seq
    ....................uugugcuugaucuaaccaugu..................................................................  54        0         seq
    ....................uugugcuugaucuaCccaugu..................................................................  2         1         seq
    ....................uugugcuugaucuaaccaugAg.................................................................  1         1         seq
    ....................uugugcuugaucuaacAaugug.................................................................  1         1         seq
    ....................uugugcuugaucuaaccauguA.................................................................  7         1         seq
    ....................uugugcuugaucuaaccaugug.................................................................  274       0         seq
    ....................uugugcuugaucuaCccaugug.................................................................  13        1         seq
    ....................uugugcuugaucuaaccGugug.................................................................  1         1         seq
    ....................uugugcuugaucuaaccaugugA................................................................  8         1         seq
    ....................uugugcuugaucuaaccaugugC................................................................  1         1         seq
    ....................uugugcuugaucuaaccaugugg................................................................  6         0         seq
    ....................uugugcuugaucuaaccaugugU................................................................  18        1         seq
    ....................uugugcuugaucuaaccaugugCu...............................................................  1         1         seq
    ....................uugugcuugaucuaaccaugugUu...............................................................  1         1         seq
    ....................uugugcuugaucuaaccaugugAu...............................................................  4         1         seq
    ...................ugugcuugaucuaaccaugugC..................................................................  1         1         seq
    .....................ugcuugaucuaaccaugu.....................................................................  1         0         seq
    .....................ugcuugaucuaaccaugugC...................................................................  1         1         seq
    ......................gcuugaucuaaccauguA.....................................................................  2         1         seq
    ......................gcuugaucuaaccaugug.....................................................................  1         0         seq
```