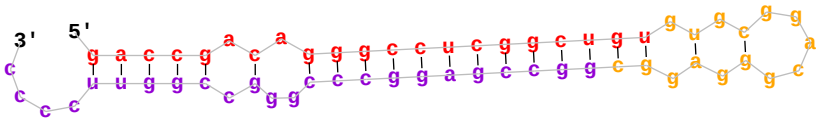
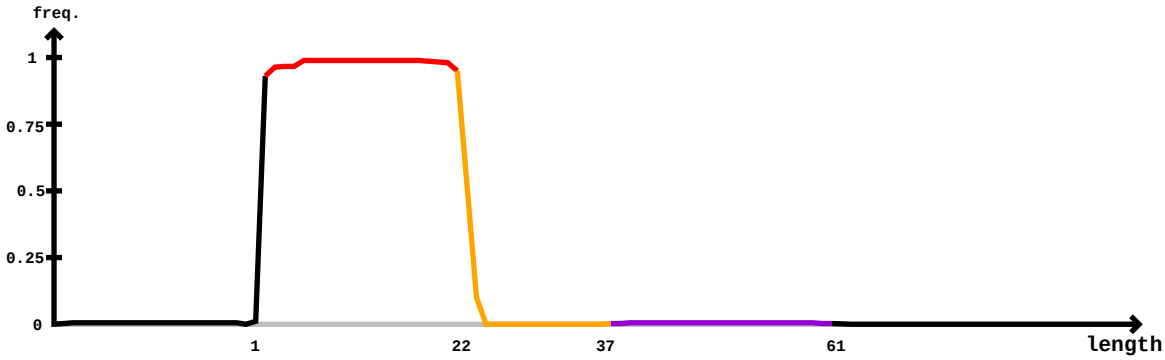
Provisional ID	:	RAE_779_9445
Score total	:	3.3
Score for star read(s)	:	-1.3
Score for read counts	:	0
Score for mfe	:	3
Score for randfold	:	1.6
Score for cons. seed	:	
Total read count	:	358
Mature read count	:	356
Loop read count	:	0
Star read count	:	2





Mature Star

```
5'- gccgaggggccgcggcgggggaccgacagggccucggcugugugcggacgggaggcggccgaggcccgggccgguuccccgaagcggcagcggcucccggcacuuccccg  -3'   obs
    gccgaggggccgcggcgggggaccgacagggccucggcugugugcggacgggaggcggccgaggcccgggccgguuccccgaagcggcagcggcucccggcacuuccccg        exp
    ((((.((((((((.((((((((((.(.((((((((((((.(.(.....).).))))))))))))..).))))))))).......)).))))))))))).........  reads   mm   sample
    gGcgaggggccgcggcgg.................................................................................  2       1    seq
    ...................ggaccgacagggccucggcug.......................................................  1       0    seq
    ...................ggaccgacagggccucggcuU......................................................  1       1    seq
    ...................ggaccgacagggccucggcugu.....................................................  2       0    seq
    ....................gaccgacagggccucgg.........................................................  1       0    seq
    ....................gaccgacagggccucggG........................................................  1       1    seq
    ....................gaccgacagggccucggcu.......................................................  1       0    seq
    ....................gaccgacagggccucggcug......................................................  8       0    seq
    ....................gaccgCcagggccucggcugu.....................................................  3       1    seq
    ....................gacGgacagggccucggcugu.....................................................  1       1    seq
    ....................gaccgacagggccucggcuUu.....................................................  1       1    seq
    ....................gaccgacagggccucggcugu.....................................................  232     0    seq
    ....................gaccgacagggccucggcugA.....................................................  2       1    seq
    ....................gaccgacagggccucggcuguU....................................................  49      1    seq
    ....................gaccgacagggccucggcuguAu...................................................  2       1    seq
    ....................gaccgacagggccucggcugugA....................................................  1       1    seq
    ....................gaccgacagggccucggcugugu....................................................  3       0    seq
    ....................gaccgacagggccucggcuguUu....................................................  26      1    seq
    .....................accgacagggccucggcug.......................................................  1       0    seq
    .....................accgacagggccucggcugu......................................................  7       0    seq
    .....................accgacagggccucggcuguAu....................................................  2       1    seq
    .....................accgacagggccucggcuguUu....................................................  2       1    seq
    ......................ccgacagggccucggcugu......................................................  1       0    seq
    .......................gacagggccucggcugu.......................................................  5       0    seq
    .......................gacagggccucggcugug......................................................  1       0    seq
    .......................gacagggccucggcuguU.....................................................  2       1    seq
    ..................................................ggccgaggcccgggccgguucccU.............................  1       1    seq
    ..................................................ccgaggcccgggccgguucA.................................  1       1    seq
```