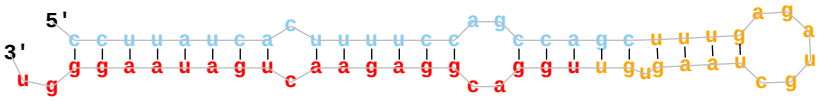
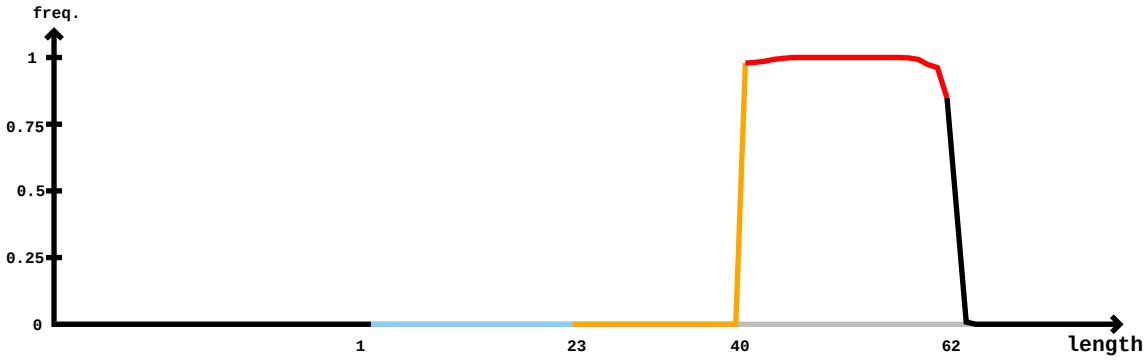
Provisional ID	:	RAE_2235_37451
Score total	:	2.2
Score for star read(s)	:	-1.3
Score for read counts	:	0
Score for mfe	:	1.9
Score for randfold	:	1.6
Score for cons. seed	:	
Total read count	:	716
Mature read count	:	716
Loop read count	:	0
Star read count	:	0





Star Mature

```
5'- gucaaagccgcuucagcagccagucacaucccccuuaucacuuuuccagccagcuuugagaugcuaagugu uggacggagaacugauaagggu aggugacugacacagcc  -3'   exp
    ......((.((....)).))(((((((...((((((((.((((((..((((((((((......)))).)))))..))))))).))))))))...))))))).....  reads   mm   sample
    ...................................................................uggacggagaacugaua.....................  1       0    seq
    ...................................................................uggacggagaacugauaa....................  4       0    seq
    ...................................................................uggacggagaaUugauaag...................  1       1    seq
    ...................................................................uggacggagaacugauaag...................  13      0    seq
    ...................................................................uggacggagaacugCuaagg..................  1       1    seq
    ...................................................................uggacggagaacugauaagg..................  7       0    seq
    ...................................................................uggacggagaacugauaaggU.................  1       1    seq
    ...................................................................uggacggagaacugauaaggA.................  4       1    seq
    ...................................................................uggacggagaacugauaaggg.................  71      0    seq
    ...................................................................uggacggagaacugCuaaggg.................  5       1    seq
    ...................................................................uggacggagaacugCuaagggu................  1       1    seq
    ...................................................................uggacggagaacugauaagggC................  476     1    seq
    ...................................................................uggacggagaacugauaagggu................  11      0    seq
    ...................................................................uggacggagaacugauaagggA................  39      1    seq
    ...................................................................uggacggagaacugauaaggguU................  1       1    seq
    ...................................................................uggacggagaacugauaagggAa................  7       1    seq
    ...................................................................uggacggagaacugauaagggCa................  52      1    seq
    ...................................................................uggacggagaacugauaagggCag...............  4       1    seq
    ...................................................................uggacggagaacugauaaggguaA...............  1       1    seq
    ...................................................................uggacggagaacugauaaggguaU...............  1       1    seq
    ...................................................................ggacggagaacugauaagggC.................  1       1    seq
    ...................................................................ggacggagaacugauaagggCa................  1       1    seq
    ...................................................................gacggagaacugauaagggC.................  2       1    seq
    ...................................................................gacggagaacugauaagggCa................  1       1    seq
    ...................................................................acggagaacugauaaggg...................  1       0    seq
    ...................................................................acggagaacugauaagggC.................  4       1    seq
    ...................................................................cggagaacugauaagggA.................  1       1    seq
    ...................................................................cggagaacugauaagggC.................  1       1    seq
    ...................................................................cggagaacugauaagggCa................  1       1    seq
    ...................................................................ggagaacugauaagggC.................  2       1    seq
```