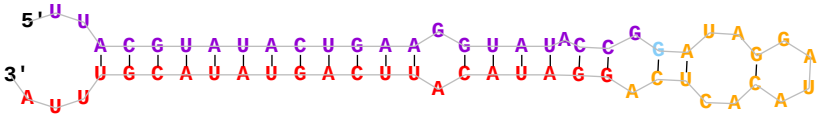
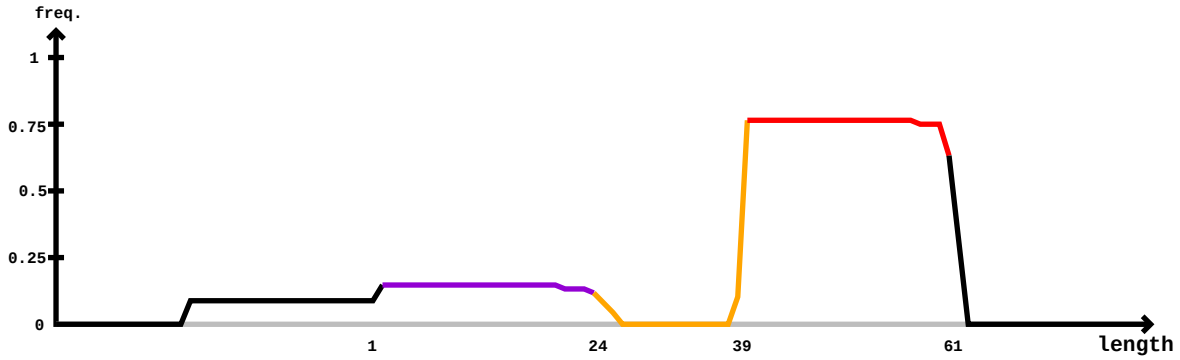
Provisional ID	:	3R_dna:chromosome_chromosome:BDGP6:3R:1:32079331:1_REF_25308
Score total	:	32.2
Score for star read(s)	:	3.9
Score for read counts	:	25.6
Score for mfe	:	1.7
Score for randfold	:	1.6
Score for cons. seed	:	-0.6
Total read count	:	62
Mature read count	:	52
Loop read count	:	0
Star read count	:	10





Star **Mature**

```
5'- uucacaucccacguuguuagugcucucucgguuacguauacugaagguauaccggauaggauacacucaggauacauucaguauacguuuacgaggucgcacagaaucgcc  -3'   obs
    uucacaucccacguuguuagugcucucucgguuacguauacugaagguauaccggauaggauacacucaggauacauucaguauacguuuacgaggucgcacagaaucgcc        exp
    uucacaucccacguuguuagugcucucucgguuacguauacugaagguauaccggauaggauacacucaggauacauucaguauacguuuacgaggucgcacagaaucgcc        known
    ...........((...((.((((..(((((((..(((((((((((.((((.((.((..(....)..)).)))))).)))))))))))).)))))))..)))).))..))..  reads  mm  sample
    ...........guuguuagugcucucucgug.....................................................................................  6      0   seq
    ............................uuacgCauacugaagguau.....................................................................  1      1   seq
    ............................uuacguauacugaagguauacc..................................................................  1      0   seq
    ............................uuacguauacugaagguauaccg.................................................................  5      0   seq
    ............................Auacguauacugaagguauaccgga...............................................................  1      1   seq
    ............................uuacguauacugaagguauaAcgga...............................................................  1      1   seq
    ............................uuacguauacugaagguauaccggU...............................................................  1      1   seq
    ................................................................aggauacaCucaguauacguuu..........................  1      1   seq
    ................................................................aggauacauucaguauacguuu..........................  2      0   seq
    ................................................................aggauacauucaguauacguuua.........................  4      0   seq
    ..................................................................ggauacauucaguauacg............................  1      0   seq
    ..................................................................ggauacaCucaguauacguuu..........................  1      1   seq
    ..................................................................ggauacauucaguauacguuu..........................  4      0   seq
    ..................................................................ggaGacauucaguauacguuua.........................  1      1   seq
    ..................................................................ggauacauucaguauacguuua.........................  35     0   seq
    ..................................................................ggauacauucaguauacguuuaA........................  3      1   seq
```