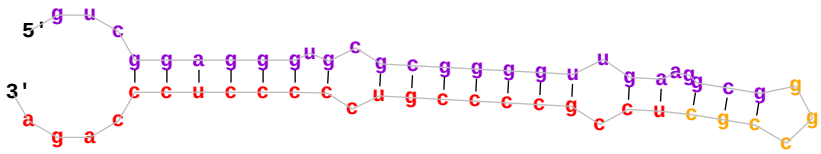
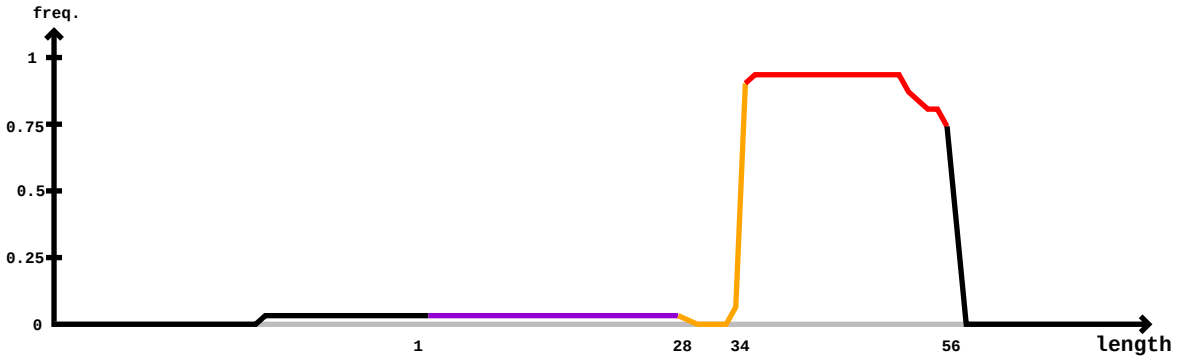
Provisional ID	:	DRO_27851_37293
Score total	:	13.5
Score for star read(s)	:	3.9
Score for read counts	:	9.3
Score for mfe	:	2.5
Score for randfold	:	-2.2
Score for cons. seed	:	
Total read count	:	30
Mature read count	:	29
Loop read count	:	0
Star read count	:	1





Star Mature

```
5'- cgccucgccggcccccggggcgguuagcgcuuccggggucggagggugcgcggggguugaaggcgggccgcuccgccccgucccccucccagaccagcagaggcagcagccgg  -3'    obs
    cgccucgccggcccccggggcgguuagcgcuuccggggucggagggugcgcggggguugaaggcgggccgcuccgccccgucccccucccagaccagcagaggcagcagccgg         exp
    .(((((((.((.((((((((((.....))).)))))))..((((((.(.(((((((.((..(((...))))).)))))))).)))))))....)).)).)))))........   reads     mm        sample
    ....................cgguuagcgcuuUcggg.................................................................................   1         1         seq
    ....................................gucggagggugcgcggggguugaaAgcg.............................................................   1         1         seq
    ..................................................................cuccgccccgucccccucccaUa...................   1         1         seq
    ..................................................................cuccgccccgucccccucccagU...................   1         1         seq
    ..................................................................uccgccccgucccccCc........................   2         1         seq
    ..................................................................uccgccccgucccccCcc.......................   1         1         seq
    ..................................................................uccgccccgucccccCccc......................   1         1         seq
    ..................................................................uccgccccgucccccucccag....................   2         0         seq
    ..................................................................uccgccccgucccccucccaga...................   14        0         seq
    ..................................................................uccgccccgucccccucccagU...................   1         1         seq
    ..................................................................uccgccccgucccccucccagaU..................   2         1         seq
    ..................................................................uccgccccgucccccucccagaA..................   3         1         seq
    ..................................................................ccgccccgucccccucccagU...................   1         1         seq
```