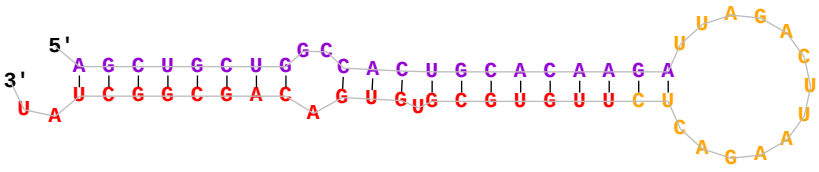
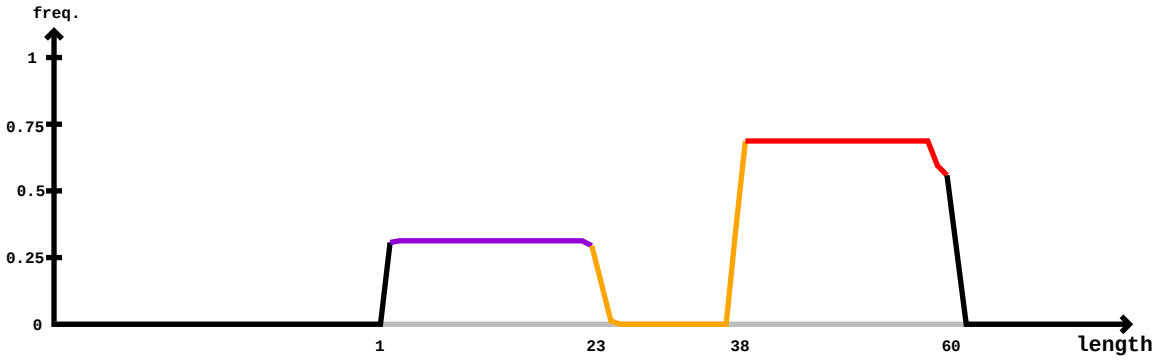
Provisional ID	:	X_dna:chromosome_chromosome:BDGP6:X:1:23542271:1_REF_27959
Score total	:	87.4
Score for star read(s)	:	3.9
Score for read counts	:	77.1
Score for mfe	:	1.8
Score for randfold	:	1.6
Score for cons. seed	:	3
Total read count	:	163
Mature read count	:	112
Loop read count	:	0
Star read count	:	51





Star Mature

```
5'- cgaccacguccggaaccgaaaggugcuuauugcagcugcuggccacugcacaagauuagacuuaagacucuugugcgugugacagcggcuauuguaagaggccauagaag  -3'   obs
    cgaccacguccggaaccgaaaggugcuuauugcagcugcuggccacugcacaagauuagacuuaagacucuugugcgugugacagcggcuauuguaagaggccauagaag        exp
    cgaccacguccggaaccgaaaggugcuuauugcagcugcuggccacugcacaagauuagacuuaagacucuugugcgugugacagcggcuauuguaagaggccauagaag        known
    .......(.((...(((....))).(((((...((((((((..((((((((((((.............)))))))))).))).))))))))...))))).)).)....... reads   mm   sample
    ................................agcugcuggccacugcacaag.................................................... 3       0    seq
    ................................agcugcuAgccacugcacaaga................................................... 4       1    seq
    ................................agcugcuggccacugcacaaga................................................... 26      0    seq
    ................................agcCgcuggccacugcacaaga................................................... 1       1    seq
    ................................agcugcuggccacugcacaagau................................................... 13      0    seq
    ................................agcugcuggccacugcacaagaA................................................... 1       1    seq
    ................................agcugcuggccacugcacaagauu.................................................. 1       0    seq
    ................................agcugcuggccacugcacaagauA................................................... 1       1    seq
    ................................gcugcuggccacugcacaaga.................................................... 1       0    seq
    ...................................................................cuugugcgugugacagcggcu.................... 9       0    seq
    ...................................................................cuugugcgugugacagcggcua................... 6       0    seq
    ...................................................................cuugugcgugugacagcggcuaA.................. 2       1    seq
    ...................................................................cuugugcgugugacagcggcuau.................. 35      0    seq
    ...................................................................cuugugcgugugacagcggcuaC.................. 2       1    seq
    ...................................................................cuugugUgugugacagcggcuau.................. 1       1    seq
    ...................................................................cuugugcgugugacagcggcuauu................. 1       0    seq
    ...................................................................cuugugcgugugacagcggcuauA................. 2       1    seq
    ...................................................................uugugcgugugacagcggcu.................... 6       0    seq
    ...................................................................uugugcgugugacagcggcuaA.................. 1       1    seq
    ...................................................................uugugcgugugacagcggcuau.................. 46      0    seq
    ...................................................................uugugcgugugacagcUgcuau.................. 1       1    seq
```