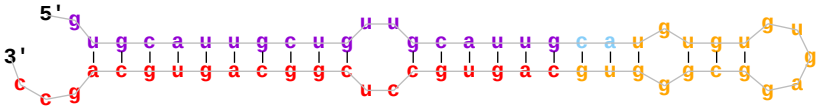
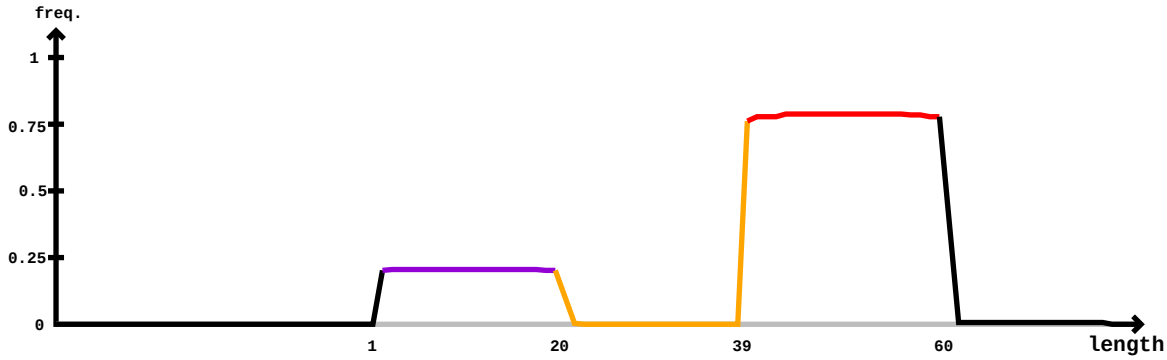
Provisional ID	:	RAE_1424_15437
Score total	:	155.1
Score for star read(s)	:	3.9
Score for read counts	:	147
Score for mfe	:	2.6
Score for randfold	:	1.6
Score for cons. seed	:	
Total read count	:	300
Mature read count	:	238
Loop read count	:	0
Star read count	:	62





Star **Mature**

```
5'- caugggggccuggagaggcgggcggccccgcggugcauugcuguugcauugcaugugugugaggcgggugcagugccucggcagugcagccCGGAGCCGGCCCCUGGCACC  -3'   obs
    caugggggccuggagaggcgggcggccccgcggugcauugcuguugcauugcaugugugugaggcgggugcagugccucggcagugcagccCGGAGCCGGCCCCUGGCACC        exp
    .....(((((.....(((.((.((((.(((.(((((((((((..(((((((((.(((.....))).)))))))))..)))))))))).)))))).))))).)))))))).))   reads    mm    sample
    ............................gugcauugcuguugcau.....................................................................   1        0     seq
    ............................gugcauugcuguugcCuug.....................................................................   2        1     seq
    ............................gugcauugcuguugcauug.....................................................................   56       0     seq
    ............................gugAauugcuguugcauug.....................................................................   1        1     seq
    ............................gugcauugcuguugcauugca...................................................................   1        0     seq
    ............................ugcauugcuguugcauug.....................................................................   1        0     seq
    ....................................................................cagugccucggcagugc.....................   1        0     seq
    ....................................................................cagugccucggcagugcag....................   2        0     seq
    ....................................................................cagugccucggcagugcagc...................   2        0     seq
    ....................................................................caguCccucggcagugcagcc..................   1        1     seq
    ....................................................................cagugccucggcagugcagcc..................   180      0     seq
    ....................................................................cagGgccucggcagugcagcc..................   1        1     seq
    ....................................................................cUgugccucggcagugcagcc..................   1        1     seq
    ....................................................................cagugccucggcagugcagcU..................   4        1     seq
    ....................................................................cagugccucggcagugcCgcc..................   2        1     seq
    ....................................................................cagugccucggcCgugcagcc..................   17       1     seq
    ....................................................................cagugccucggcagugcagccc.................   3        0     seq
    ....................................................................cagugccucggcagugcagccU.................   11       1     seq
    ....................................................................cagugccucggcagugcagccA.................   5        1     seq
    ....................................................................agugccucggcagugcagcc...................   3        0     seq
    ....................................................................agugccucggcagugcagccA..................   2        1     seq
    ..................................................................gccucggcagugcagcc.......................   2        0     seq
    ..................................................................gccucggcagugcagccA......................   1        1     seq
    ...................................................................................ccggagccggccccGggc...   2        1     seq
```