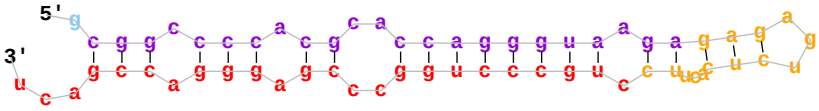
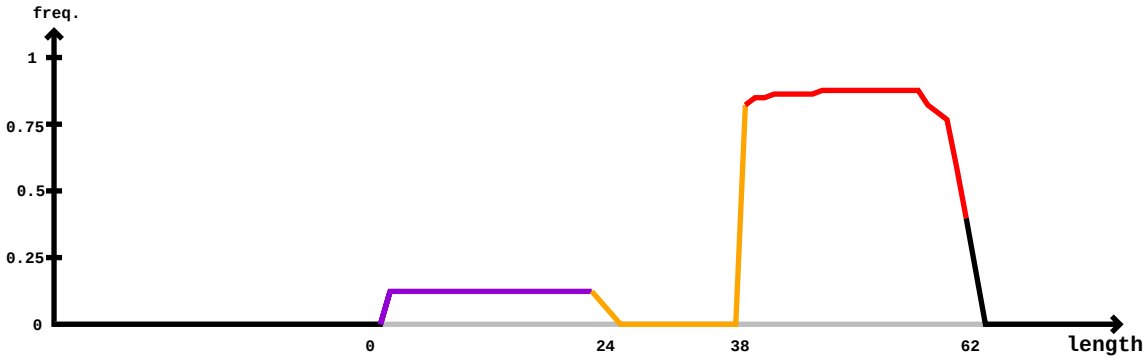
Provisional ID	:	RAE_1912_27457
Score total	:	34.7
Score for star read(s)	:	3.9
Score for read counts	:	31.2
Score for mfe	:	1.8
Score for randfold	:	-2.2
Score for cons. seed	:	
Total read count	:	73
Mature read count	:	64
Loop read count	:	0
Star read count	:	9





Star **Mature**

```
5'- gcuccuucggaugaauuuuugguguuagcccugcggccccacgcaccaggguaagagagagucucacuuccugcccuggcccgagggaccgacuggcugggccugccuu  -3'   obs
    gcuccuucggaugaauuuuugguguuagcccugcggccccacgcaccaggguaagagagagucucacuuccugcccuggcccgagggaccgacuggcugggccugccuu        exp
    ..(((...))).........(((..((((((...(((.(((.((..(((((((((.(((((...))))...)).)))))))))..)).)))).)))...)))))).)))......  reads   mm   sample
    ................................cggccccacgcaccaggguaag.....................................................  2       0    seq
    ................................cggccccacgcaccaggguaagU.....................................................  3       1    seq
    ................................cggccccacgcaccaggguaaga.....................................................  4       0    seq
    ....................................................................cugcccuggcccgagggac.....................  4       0    seq
    ....................................................................cugcccuggcccgagggacA....................  2       1    seq
    ....................................................................cugcccuggcccgagggaccg...................  2       0    seq
    ....................................................................cugcccuggAccgagggaccga..................  1       1    seq
    ....................................................................cugcccuggcccgagggaccga..................  9       0    seq
    ....................................................................cugcccuggcccgagggaccgU..................  1       1    seq
    ....................................................................cugcccuggcccgagggaccgaU.................  3       1    seq
    ....................................................................cugcccuggcccgagggaccgaA.................  1       1    seq
    ....................................................................cugcccuggcccgagggaccgac.................  9       0    seq
    ....................................................................cugcccuggcccgagggaccgaAu................  2       1    seq
    ....................................................................cugcccuggcccgCgggaccgacu................  1       1    seq
    ....................................................................cugcccuggccAgagggaccgacu................  1       1    seq
    ....................................................................cugcccuggcccgagggaccgaUu................  2       1    seq
    ....................................................................cugcccuggcccgagggaccgacu................  15      0    seq
    ....................................................................cugcccuggcccgagggaccgaGu................  2       1    seq
    ....................................................................cugcccuggcccgagggaccgacC................  3       1    seq
    ....................................................................cugcccuggcccgagggaccgacA................  2       1    seq
    ....................................................................ugcccuggcccgagggaccga...................  2       0    seq
    ....................................................................cccuggcccgagggaccgac....................  1       0    seq
    ....................................................................gccUgagggaccgacug.......................  1       1    seq
```