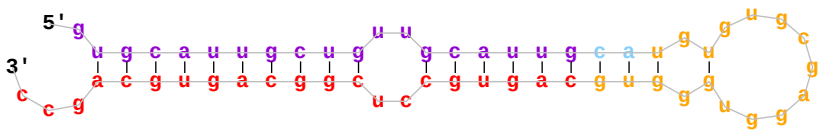
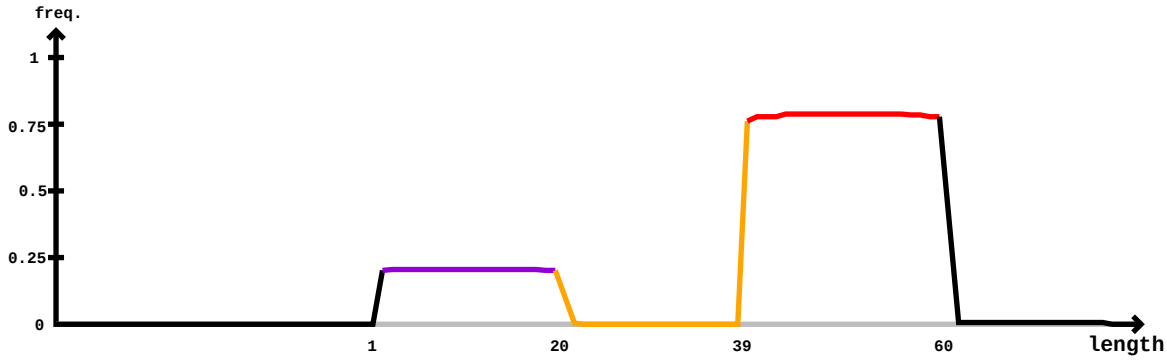
Provisional ID	:	MNA_323_7041
Score total	:	155
Score for star read(s)	:	3.9
Score for read counts	:	147
Score for mfe	:	2.6
Score for randfold	:	1.6
Score for cons. seed	:	
Total read count	:	300
Mature read count	:	238
Loop read count	:	0
Star read count	:	62





Star **Mature**

```
5'- gguggggggccuggagaggcgggcggccccgaggugcauugcuguugcauugcaugugugcgaggugggugcagugccucggcagugcagcccggagccggccccuggcacc  -3'   obs
    gguggggggccuggagaggcgggcggccccgaggugcauugcuguugcauugcaugugugcgaggugggugcagugccucggcagugcagcccggagccggccccuggcacc        exp
    .((.(((((((....)))))((.((((.(((.(((((((((((..(((((((((.(.........).)))))))))..)))))))))).)))))).))))).))...   reads      mm         sample
    ............................gugcauugcuguugcau.....................................................................   1          0          seq
    ............................gugcauugcuguugcauug....................................................................   56         0          seq
    ............................gugAauugcuguugcauug....................................................................   1          1          seq
    ............................gugcauugcuguugcCuug....................................................................   2          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
    ....................................................................cagugccucggcCgugcagcc..................   17         1          seq
    ....................................................................cUgugccucggcagugcagcc..................   1          1          seq
    ....................................................................cagugccucggcagugcCgcc..................   2          1          seq
    ....................................................................cagugccucggcagugcagcU..................   4          1          seq
    ....................................................................cagGgccucggcagugcagcc..................   1          1          seq
    ....................................................................cagugccucggcagugcagcc..................   180        0          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
```