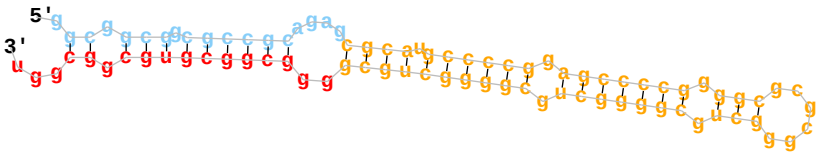
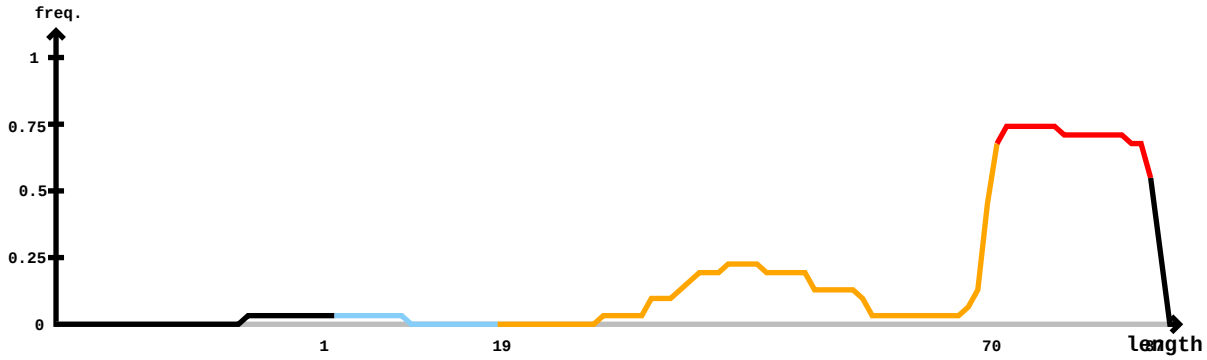
Provisional ID	:	RAE_1569_16342
Score total	:	3.3
Score for star read(s)	:	-1.3
Score for read counts	:	0
Score for mfe	:	3
Score for randfold	:	1.6
Score for cons. seed	:	
Total read count	:	28
Mature read count	:	21
Loop read count	:	7
Star read count	:	0





Star Mature

```
5'- aacucugccgcgggagggggugggggagggcggcggcgccgcagagcgcaugccccggagccccggggcgcgcgggcugcggggcugcggggcugcggggcggcgugcggcggugg  -3'   exp
    ..(((((((.(.....).)))))))...((.(((.((((((....((((.((((((.(((((((.(((.......))).)))))))).))))))))))..)))))))))).))..... reads   mm   sample
    ..................gCuggggagggcggcgg.......................................................................................  1      1    seq
    ..................................................ggagccccggggcgcgU.......................................................  1      1    seq
    ..................................................................cccgggCcgcgcgggcu.......................................  2      1    seq
    ..................................................................ggggcgcgcgggcugcgggC...................................  1      1    seq
    ..................................................................gggcgcgcgggcugcggC.....................................  1      1    seq
    ..................................................................ggGgcgcgggcugcgggg.....................................  1      1    seq
    ..................................................................Ccgcgggcugcggggcu.......................................  1      1    seq
    ..................................................................................gcggggcugcAgggcggc.............  1      1    seq
    ..................................................................................gcggggcggcgugGggc.....  1      1    seq
    ..................................................................................cggggcggcguCcggcggu..  2      1    seq
    ..................................................................................ggggcggcguCcggcgg...  4      1    seq
    ..................................................................................ggggcggcguCcggcggu..  4      1    seq
    ..................................................................................ggggcggcguCcggcggug.  2      1    seq
    ..................................................................................gggcggcguCcggcggu..  7      1    seq
    ..................................................................................ggcggcguCcggcggug.  2      1    seq
```