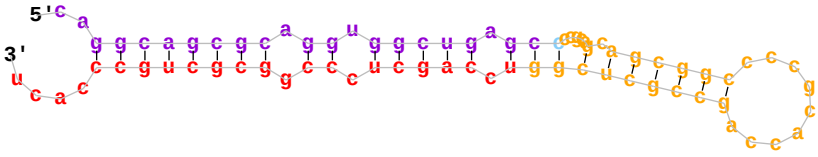
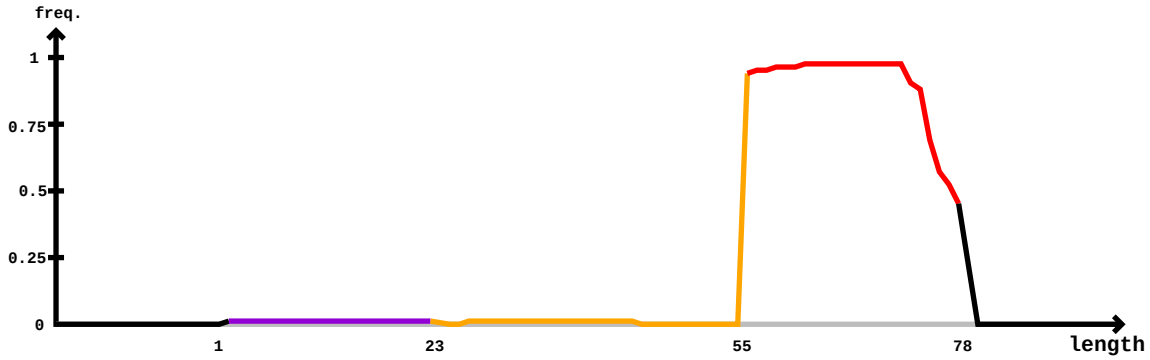
Provisional ID	:	RAE_1708_19325
Score total	:	44.6
Score for star read(s)	:	3.9
Score for read counts	:	36.8
Score for mfe	:	2.2
Score for randfold	:	1.6
Score for cons. seed	:	
Total read count	:	84
Mature read count	:	82
Loop read count	:	1
Star read count	:	1





Star **Mature**

```
5'- ucugcccgccccccuggcaggcagcgcagguggcugagccccgcagcagcggccccgcaccagccgcucgguccagcucccggcgcugcccacuggguacggaucaacaa  -3'   obs
    ucugcccgccccccuggcaggcagcgcagguggcugagccccgcagcagcggccccgcaccagccgcucgguccagcucccggcgcugcccacuggguacggaucaacaa        exp
    .....(((..(((.((..(((((((((.((.((((((.(((.....(.((((((.........)))))))))).)))))).)).)))))))..)))))..)))........  reads    mm       sample
    ...............caggcagcgcaggCggcugagc.....................................................................................  1        1        seq
    ........................................gcagcagcggccccgGac.....................................................................  1        1        seq
    ..............................................................................uccagcucccggcgcug.....................  6        0        seq
    ..............................................................................uccagcucccggcgcugc.....................  1        0        seq
    ..............................................................................uccagcucccggcgcugA.....................  1        1        seq
    ..............................................................................uccagcucccggcgcugcc.....................  16       0        seq
    ..............................................................................uccagcucccggcgcugccU.....................  1        1        seq
    ..............................................................................uccagcucccggcgcugccA.....................  1        1        seq
    ..............................................................................uccagcucccggcgcugccc.....................  8        0        seq
    ..............................................................................uccagcucccggcgcugccca.....................  4        0        seq
    ..............................................................................uccagcucccggcgcugcccac.....................  2        0        seq
    ..............................................................................uccagcucccggcgcugcccaU.....................  4        1        seq
    ..............................................................................uccagcucccggcgcugcccaAu.....................  1        1        seq
    ..............................................................................uccagcucccggcgcugcccaUu.....................  7        1        seq
    ..............................................................................uccagcucccggcgcugcccacu.....................  24       0        seq
    ..............................................................................uccagcucccggcgcugcccacuU.....................  3        1        seq
    ..........................................................................ccagcucccggcgcugcccacu.....................  1        0        seq
    ......................................................................................agcuGccggcgcugcccacu.....................  1        1        seq
    ..............................................................................ucccggcgcugcccaAu.....................  1        1        seq
```