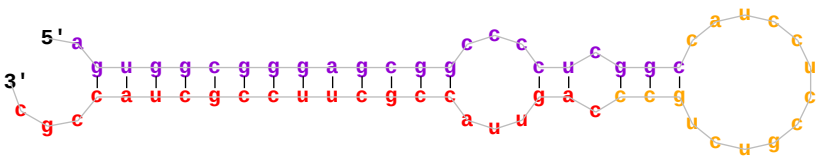
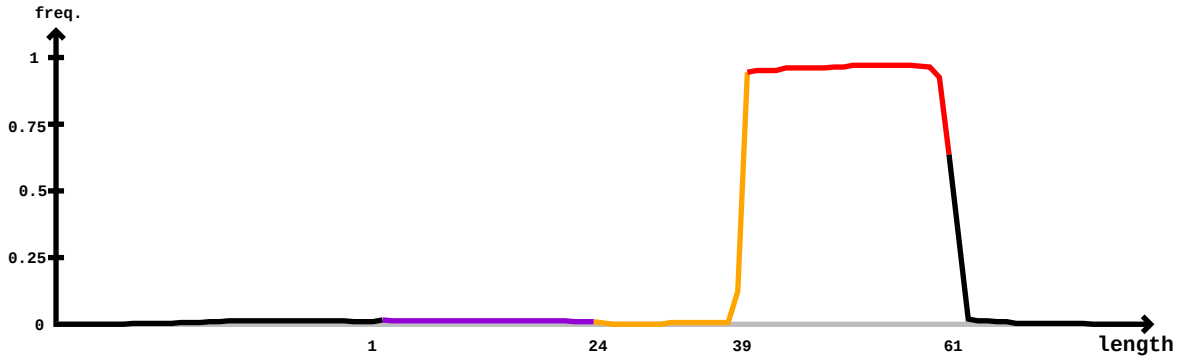
Provisional ID	:	RAE_2389_41090
Score total	:	154.5
Score for star read(s)	:	3.9
Score for read counts	:	146.4
Score for mfe	:	2.6
Score for randfold	:	1.6
Score for cons. seed	:	
Total read count	:	299
Mature read count	:	295
Loop read count	:	0
Star read count	:	4





Star Mature

```
5'- accugcucaaggccggcgggggcgcgggcggcaguggcgggagcggccccucggccauccuccgucugcccaguuaccgcuuccgcuaccgccgccgcucccgcucuagcuc  -3'    obs
    accugcucaaggccggcgggggcgcgggcggcaguggcgggagcggccccucggccauccuccgucugcccaguuaccgcuuccgcuaccgccgccgcucccgcucuagcuc         exp
    ..........(((.(((((((((...((((((.((((((((((((...((.(((............))).))...))))))))))))).))))))))))))))...))).   reads     mm        sample
    ......ucaaggccggcgggggcgcgggcggc.......................................................................................  1         0         seq
    ...........gcGggcgggggcgcgggc.........................................................................................  1         1         seq
    ..............ggcAggggcgcgggcggc.......................................................................................  1         1         seq
    ................cgggggcgcAggcggca.......................................................................................  1         1         seq
    ..............................aguggcgggagcggccccuc.......................................................................  1         0         seq
    ..............................aguggcgggagcggccccucggc....................................................................  2         0         seq
    ..............................aguggcgggagcggccccucggcA...................................................................  1         1         seq
    ...................................................................cgucugcccaguuaccgcuuccgcuaccg.....................  2         0         seq
    ...................................................................ccaguuaccgcuuccgcuacc.....................  3         0         seq
    ...................................................................ccaguuaccgcuuccgcuaccg.....................  6         0         seq
    ...................................................................ccaguuaccgcuuccgcuaccgA....................  2         1         seq
    ...................................................................ccaguuaccgcuuccgcuaccgU....................  1         1         seq
    ...................................................................ccaguuaccgcuuccgcuaccgc....................  24        0         seq
    ...................................................................caguuaccgcuuccgcua.......................  1         0         seq
    ...................................................................caguuaccgcuuccgcuac......................  2         0         seq
    ...................................................................caguuaccgcuuccgcuacc.....................  9         0         seq
    ...................................................................caguuCccgcuuccgcuaccg....................  1         1         seq
    ...................................................................caguuaccgcuuccgcuaccg....................  78        0         seq
    ...................................................................caguuaccgcuuccgcuaccgc....................  131       0         seq
    ...................................................................caguuaccgcuuccgcuaccgA....................  4         1         seq
    ...................................................................caguuaccgcuuccgcuaccgU....................  4         1         seq
    ...................................................................caguuaccgcuuccgcuCccgc....................  1         1         seq
    ...................................................................caguuaccgcuuccgcuaccgcA....................  6         1         seq
    ...................................................................caguuaccgcuuccgcuaccgcU....................  14        1         seq
    ...................................................................caguuaccgcuuccgcuaccgcUg...................  1         1         seq
    ...................................................................caguuaccgcuuccgcuaccgcAg...................  1         1         seq
    ...................................................................aguuaccgcuuccgcuaccg.....................  2         0         seq
    ..................................................................uaccgcuuccgcuaccgc....................  2         0         seq
    ..................................................................uaccgcuuccgcuaccgU....................  1         1         seq
    ...................................................................cuuccgcCaccgccgcc.................  1         1         seq
    ...................................................................uccgcuGccgccgccgc.................  2         1         seq
    ...................................................................cgccgccgcucccgcCc.......  1         1         seq
```