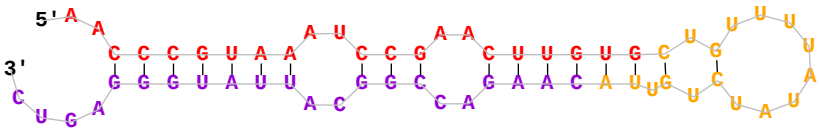
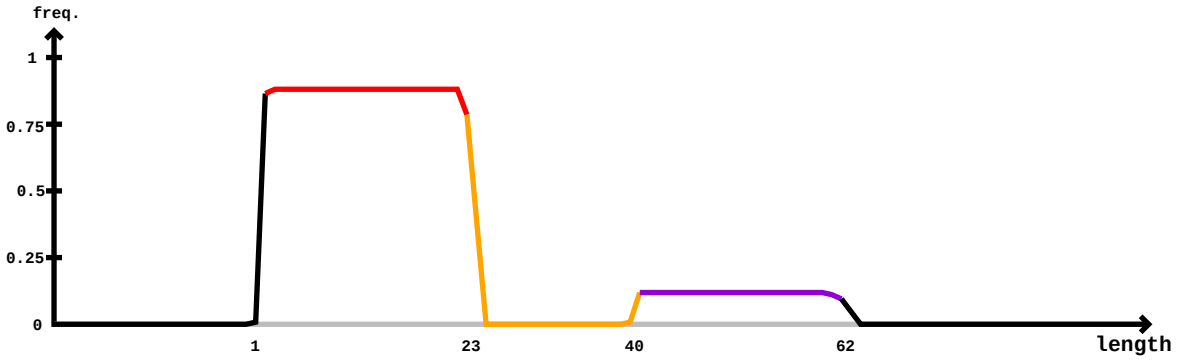
Provisional ID	:	2L_dna:chromosome_chromosome:BDGP6:2L:1:23513712:1_REF_638
Score total	:	66.9
Score for star read(s)	:	3.9
Score for read counts	:	58.2
Score for mfe	:	0.2
Score for randfold	:	1.6
Score for cons. seed	:	3
Total read count	:	126
Mature read count	:	111
Loop read count	:	0
Star read count	:	15





Mature **Star**

```
5'- acauauggaccauuaacagaaacccguaaauccgaacuugugcuguuuuauaucuguuacaagaccggcauuaugggagucugucaaugcaaacaacuggguuuuuggcaaca  -3'    obs
    acauauggaccauuaacagaaacccguaaauccgaacuugugcuguuuuauaucuguuacaagaccggcauuaugggagucugucaaugcaaacaacuggguuuuuggcaaca        exp
    acauauggaccauuaacagaaacccguaaauccgaacuugugcuguuuuauaucuguuacaagaccggcauuaugggagucugucaaugcaaacaacuggguuuuuggcaaca        known
    .((..((...((((.(((((..(((((((..(((..(((((((.(........).).)))))))..)))..)))))))..)))))).))))....))..))).....((....))  reads   mm   sample
    ...................aaacccguaaauccgaacuugug.....................................................................  1       0    seq
    ...................aacccguaaauccgaacuugu......................................................................  8       0    seq
    ...................aacccguaGauccgaacuugu......................................................................  1       1    seq
    ...................aacccguaaauccgaaUuugu......................................................................  1       1    seq
    ...................aacccguaaaCccgaacuugu......................................................................  1       1    seq
    ...................aacccgAaaauccgaacuugu......................................................................  1       1    seq
    ...................aacccguaaauccAaacuugug.....................................................................  2       1    seq
    ...................aacccguaaauccgaacuugGg.....................................................................  1       1    seq
    ...................aacccguaaauccgaacuGgug.....................................................................  2       1    seq
    ...................aacccguaaauccgaaUuugug.....................................................................  1       1    seq
    ...................Gacccguaaauccgaacuugug.....................................................................  1       1    seq
    ...................aacccguaaaucGgaacuugug.....................................................................  1       1    seq
    ...................aacccguaGauccgaacuugug.....................................................................  3       1    seq
    ...................aacccguaaaCccgaacuugug.....................................................................  1       1    seq
    ...................aacccguaaauccgaacuugug.....................................................................  78      0    seq
    ...................aacccguaaauccgaacAugug.....................................................................  1       1    seq
    ...................aaccUguaaauccgaacuugug.....................................................................  1       1    seq
    ...................aacccguaGauccgaacuugugc....................................................................  3       1    seq
    ...................aacccguaaauccgaacuugugU....................................................................  1       1    seq
    ..................acccguaaauccgaacuugug......................................................................  2       0    seq
    ..........................................................Ccaagaccggcauuaugggagu...............................  1       1    seq
    ..........................................................caagaccggcauuaugggag................................  1       0    seq
    ..........................................................caagaccggcauuaugggagu...............................  1       0    seq
    ..........................................................caagGccggcauuaugggaguc..............................  1       1    seq
    ..........................................................caagaccggcauuaugggaguc..............................  11      0    seq
```