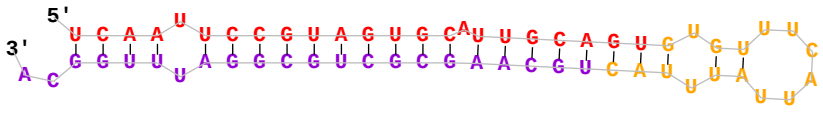
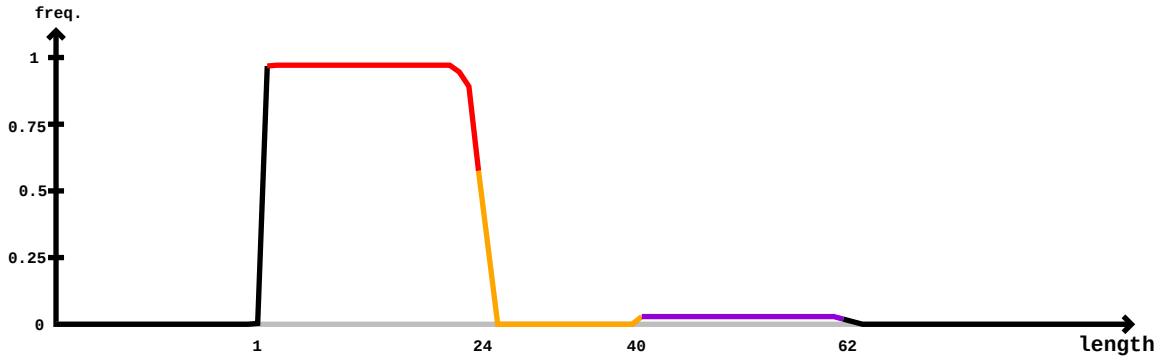
Provisional ID	:	2L_dna:chromosome_chromosome:BDGP6:2L:1:23513712:1_REF_276
Score total	:	164.1
Score for star read(s)	:	3.9
Score for read counts	:	153.6
Score for mfe	:	2
Score for randfold	:	1.6
Score for cons. seed	:	3
Total read count	:	313
Mature read count	:	304
Loop read count	:	0
Star read count	:	9





Mature Star

```
5'- gaguauugguuuugaaguuuucaauuccguagugcauugcagugugguucauuauuuacugcaagcgcugcggauuuggcaacuuugacgaccuucaucaucugcuagag  -3'   obs
    gaguauugguuuugaaguuuucaauuccguagugcauugcagugugguucauuauuuacugcaagcgcugcggauuuggcaacuuugacgaccuucaucaucugcuagag        exp
    gaguauugguuuugaaguuuucaauuccguagugcauugcagugugguucauuauuuacugcaagcgcugcggauuuggcaacuuugacgaccuucaucaucugcuagag        known
    ((.....(((((..(((((.((((.(((((((((.(((((((.((......)).))))))))))))))))).)))).)))))..).))))....))........... reads   mm   sample
    ...................Aucaauuccguagugcauugcag.................................................................... 1       1    seq
    ...................ucaauuccguaguAcauugc........................................................................ 1       1    seq
    ...................ucaauuccguagugcauugc........................................................................ 7       0    seq
    ...................ucaauuccguagugcauugca....................................................................... 17      0    seq
    ...................ucaauuccgGagugcauugcag...................................................................... 1       1    seq
    ...................ucaGuuccguagugcauugcag...................................................................... 2       1    seq
    ...................ucaauuccguagugcauugcag...................................................................... 90      0    seq
    ...................ucaauAccguagugcauugcag...................................................................... 2       1    seq
    ...................ucaauuccguagugcauugcaA...................................................................... 1       1    seq
    ...................ucaauuccguagugcauAgcag...................................................................... 1       1    seq
    ...................ucaauuccguagugcauCgcagu..................................................................... 1       1    seq
    ...................ucaauuccguagugcauugcagu..................................................................... 164     0    seq
    ...................ucaauuccguagugcauugcaCu..................................................................... 1       1    seq
    ...................ucaauuccguaAugcauugcagu..................................................................... 1       1    seq
    ...................ucaauuccguagugcauAgcagu..................................................................... 1       1    seq
    ...................ucaauuccguagugcGuugcagu..................................................................... 1       1    seq
    ...................ucaauuccguagCgcauugcagu..................................................................... 2       1    seq
    ...................ucaauuccguagugcauugcGgu..................................................................... 1       1    seq
    ...................ucaauuccguagugcauugcagN..................................................................... 1       1    seq
    ...................ucaauuccguagugAauugcagu..................................................................... 1       1    seq
    ...................ucaauuccguagugcUuugcagu..................................................................... 1       1    seq
    ...................ucaauuccguagugcauugcagC..................................................................... 1       1    seq
    ...................ucaauuccguagugcaAugcagu..................................................................... 1       1    seq
    ...................ucaauuccguagugcauugcaguU.................................................................... 3       1    seq
    ....................caauuccguagugcauugcag...................................................................... 1       0    seq
    ..................................................ugcaagcgcugcggauuuggc.......................................... 3       0    seq
    ..................................................uAcaagcgcugcggauuuggca.......................................... 1       1    seq
    ..................................................ugcaagcgcugcggauuuggca.......................................... 5       0    seq
```