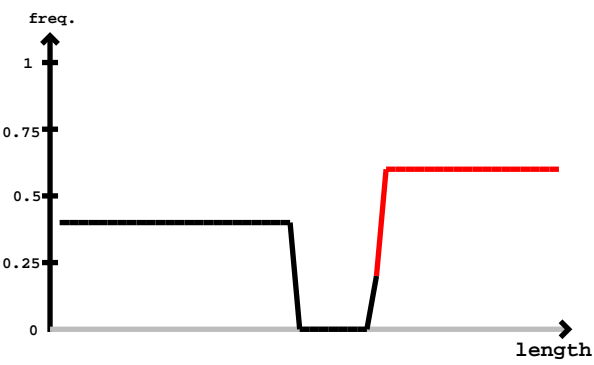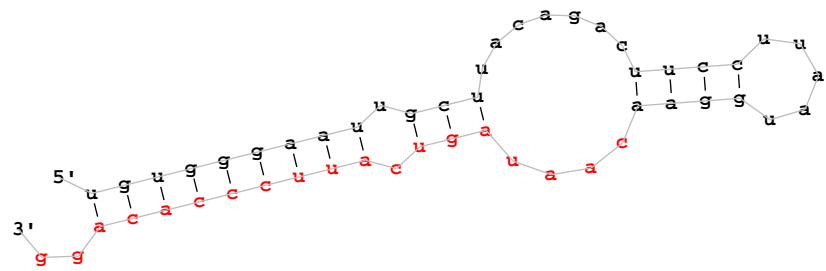miRBase precursor   : chromosome_2_5012
Total read count   : 5
chromosome_2_5012 read count : 3
remaining reads   : 2

```
5'- ugugggaauugcuuacagacuuccuuaauggaacaauagucauucccacagg  -3'    exp
    ((((((((((.(((.......(((((.....)))).....))).))))))))))..    reads     mm         sample
    ugugggaauugcuuacagacuucA..........................    2         1          seq
    ...............................acaauagucauucccacagg    1         0          seq
    ...............................caauagucauucccacagg     2         0          seq
```