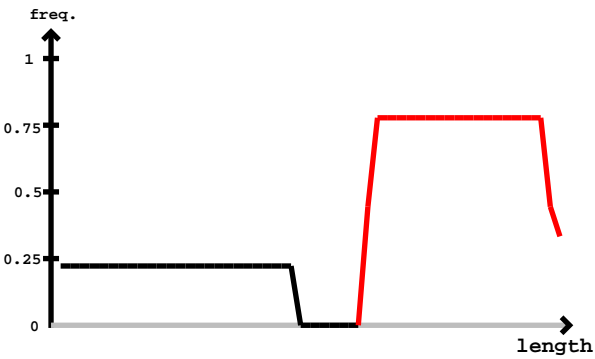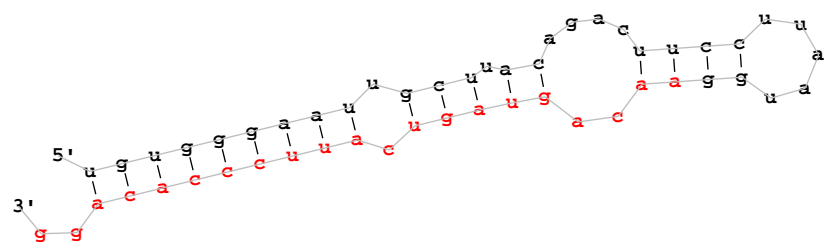miRBase precursor : chromosome_2_5007
Total read count : 9
chromosome_2_5007 read count : 7
remaining reads : 2

chromosome_2_5007

```
5'- uguggggaauugcuuacagacuuccuuaauggaacaguagucauucccacagg  -3'    exp
     ((((((((((.(((.((....(((.....)))).)))))).)))))))))..    reads    mm         sample
     uguggggaauugcuuacagacuucA...........................      2      1            seq
     ...............................aacaguagucauucccaca..      1      0            seq
     ...............................aacaguagucauucccacagg      3      0            seq
     ...............................acaguagucauucccaca..       2      0            seq
     ...............................acaguagucauucccacag.       1      0            seq
```