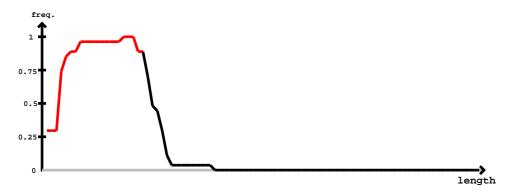miRBase precursor   : chromosome_14_91697
Total read count   : 27
chromosome_14_91697 read count : 26
remaining reads   : 1

chromosome_14_91697

```
5'- ggcaacguggccgagugguuagggcgccggacuugaaauccgaaggccccgggguucaagucccguucugaccacuaccuggguguuguuuc  -3'    exp
    .(((((...(((((((((((((((((..(((((((((.((((.......))))))))))))))))))))))))))))...)))))))))...  reads    mm    sample
    gAcaacguggccgagugg.......................................................................     3       1     seq
    gAcaacguggccgaggguu......................................................................     4       1     seq
    gAcaacguggccgaggguua.....................................................................     1       1     seq
    ..caacguggccgaggguu......................................................................     1       0     seq
    ..caacguggccgaggguua.....................................................................     3       0     seq
    ..cGacguggccgaggguua.....................................................................     2       1     seq
    ..caacguggccgaggguuaA....................................................................     1       1     seq
    ..caacguggccgaggguuaAg...................................................................     1       1     seq
    ..caacguggccgaggguuaAgg..................................................................     2       1     seq
    ..caacguggccgaggguuaAggc.................................................................     2       1     seq
    ...aacguggccgaggguuaAg...................................................................     2       1     seq
    ...aacguggccgaggguuaAgg..................................................................     1       1     seq
    ....acguggccgaggguuaAg...................................................................     1       1     seq
    ......guggccgaggguuaAgg..................................................................     2       1     seq
    ..............ugguuagAgcgccggacuu........................................................     1       1     seq
```