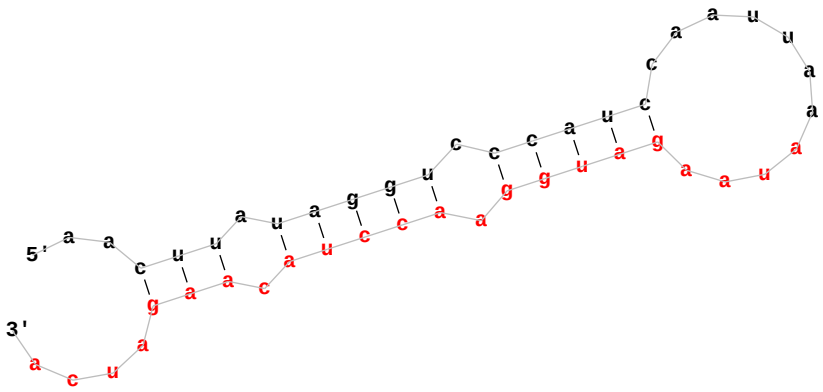
miRBase precursor : chromosome_3_9496
Total read count : 1514
chromosome_3_9496 read count : 1514
remaining reads : 0





chromosome_3_9496

```
5'- aacuuauaggucccauccaauuaaauaagauggaaccuacaagauca  -3'    exp
    ..(((.(((((.(((((..........))))).)))))).))....   reads      mm         sample
    ......................Uaauaagauggaaccuacaagauca   6          1          seq
    ......................aUauaagauggaaccuacaagauca   3          1          seq
    ......................aaauaagauggaaccuacaagauca   2          0          seq
    ......................aauaagauggaaccuacaagau..    1          0          seq
    ......................aauaagauggaaccuacaagauc.    2          0          seq
    ......................aUuaagauggaaccuacaagauca   1          1          seq
    ......................aCuaagauggaaccuacaagauca   2          1          seq
    ......................aauaagauggaacAuacaagauca   1          1          seq
    ......................aGuaagauggaaccuacaagauca   1          1          seq
    ......................aauaagauggaaccuacUagauca   1          1          seq
    ......................aauaagauggaaccuacaagauca   89         0          seq
    ......................aauaagauggaaccuacaagaCca   1          1          seq
    ......................aauaagauggaaccuacaCgauca   1          1          seq
    ......................Uauaagauggaaccuacaagauca   16         1          seq
    ......................aauaagauggaaccuacaagCuca   1          1          seq
    ......................Gauaagauggaaccuacaagauca   10         1          seq
    ......................auaagauggaaccuacaagauc.    1          0          seq
    ......................auaagauggGaccuacaagauc.    1          1          seq
    ......................auaagauggaaccuacaagaucG    6          1          seq
    ......................auaagauggaaccuacaagaCca    2          1          seq
    ......................auaagauggaaccuacaCgauca    2          1          seq
    ......................auaagauggaaccuacaaUauca    1          1          seq
    ......................auaagauggaaccCacaagauca    4          1          seq
    ......................auaagauggaaccuacGagauca    4          1          seq
    ......................auaagauggaaccuGcaagauca    1          1          seq
    ......................auaagauggaacAuacaagauca    1          1          seq
    ......................auaagauggaaccuacaagCuca    2          1          seq
    ......................auaagauAgaaccuacaagauca    2          1          seq
    ......................auaagauCgaaccuacaagauca    2          1          seq
    ......................auaagauggaGccuacaagauca    10         1          seq
    ......................auaCgauggaaccuacaagauca    1          1          seq
    ......................auaagauggaaccuacaagauca    731        0          seq
    ......................auaagaCggaaccuacaagauca    5          1          seq
    ......................Cuaagauggaaccuacaagauca    7          1          seq
    ......................Uuaagauggaaccuacaagauca    13         1          seq
```

chromosome_3_9496

aacuuauaggucccauccaauuaaauaagauggaaccuacaagauca

| | | | |
|---|---|---|---|
| .....auaagauggaaAcuacaagauca | 3 | 1 | seq |
| .....auaagauggaaccuacaagaucU | 1 | 1 | seq |
| .....auaUgauggaaccuacaagauca | 1 | 1 | seq |
| .....auaagauggaaccuacaaCauca | 2 | 1 | seq |
| .....auCagauggaaccuacaagauca | 1 | 1 | seq |
| .....auaagauggaacUuacaagauca | 2 | 1 | seq |
| .....auaagauggaaccuacaagauAa | 1 | 1 | seq |
| .....auaagaugUaaccuacaagauca | 2 | 1 | seq |
| .....auaaUauggaaccuacaagauca | 2 | 1 | seq |
| .....auaagauggaaccuacaGgauca | 2 | 1 | seq |
| .....auaagaGggaaccuacaagauca | 1 | 1 | seq |
| .....auaagauggaCccuacaagauca | 1 | 1 | seq |
| .....aAaagauggaaccuacaagauca | 2 | 1 | seq |
| .....auaagauggUaccuacaagauca | 1 | 1 | seq |
| .....auaagauggGaccuacaagauca | 5 | 1 | seq |
| .....auaagauggaaGcuacaagauca | 1 | 1 | seq |
| .....auaagauggaaccuacaagGuca | 4 | 1 | seq |
| .....auaaAauggaaccuacaagauca | 1 | 1 | seq |
| .....auaagauggaaccuCcaagauca | 3 | 1 | seq |
| .....auaagauggaaccuacaagaucC | 1 | 1 | seq |
| .....Guaagauggaaccuacaagauca | 69 | 1 | seq |
| .....auaagauggaaccuacaagauUa | 2 | 1 | seq |
| .....aCaagauggaaccuacaagauca | 5 | 1 | seq |
| .....auaagauggaaUcuacaagauca | 1 | 1 | seq |
| .....auaagauUgaaccuacaagauca | 2 | 1 | seq |
| .....auaagauggaaccuacUagauca | 1 | 1 | seq |
| .....auaGgauggaaccuacaagauca | 1 | 1 | seq |
| .....auaagGuggaaccuacaagauca | 7 | 1 | seq |
| .....uaagauggaaccuacaagauc. | 1 | 0 | seq |
| .....uaagauggaacUuacaagauca | 2 | 1 | seq |
| .....uaagaCggaaccuacaagauca | 1 | 1 | seq |
| .....uaagauggaaAcuacaagauca | 2 | 1 | seq |
| .....uaagauggaGccuacaagauca | 3 | 1 | seq |
| .....Caagauggaaccuacaagauca | 1 | 1 | seq |
| .....uaagGuggaaccuacaagauca | 2 | 1 | seq |
| .....uaGgauggaaccuacaagauca | 1 | 1 | seq |
| .....uaagauggaaccuacaagauUa | 1 | 1 | seq |
| .....uCagauggaaccuacaagauca | 3 | 1 | seq |
| .....uaagauggaacGuacaagauca | 2 | 1 | seq |
| .....uaagauggaaccuacaaAauca | 1 | 1 | seq |
| .....uaagauggaaccuacaagauca | 322 | 0 | seq |
| .....uaagauggaaccuacaagCuca | 1 | 1 | seq |
| .....uaagaugCaaccuacaagauca | 1 | 1 | seq |
| .....uaagauUgaaccuacaagauca | 1 | 1 | seq |
| .....uaagauggaaccuacaUgauca | 1 | 1 | seq |
| .....uaagauCgaaccuacaagauca | 1 | 1 | seq |
| .....uaagauggaaccuacaaUauca | 3 | 1 | seq |
| .....uaagauggaaccuacGagauca | 2 | 1 | seq |
| .....uaagauggaaccuacaagaucG | 1 | 1 | seq |
| .....uaagauggaaccCacaagauca | 2 | 1 | seq |
| .....uaagauggaaccuacaGgauca | 2 | 1 | seq |
| .....uaagaugGaccuacaagauca | 1 | 1 | seq |
| .....Aaagauggaaccuacaagauca | 13 | 1 | seq |
| .....uaagauggaaccuacaagaCca | 1 | 1 | seq |
| .....uaagauggaaGcuacaagauca | 2 | 1 | seq |
| .....Gagauggaaccuacaagauca | 2 | 1 | seq |
| .....aagauggaaccCacaagauca | 1 | 1 | seq |
| .....aagauggaaccuacaagauca | 38 | 0 | seq |
| .....Ugauggaaccuacaagau.. | 1 | 1 | seq |
| .....Ggauggaaccuacaagauca | 1 | 1 | seq |
| .....Cgauggaaccuacaagauca | 2 | 1 | seq |
| .....Ugauggaaccuacaagauca | 17 | 1 | seq |
| .....agauggaaccuacaagauca | 8 | 0 | seq |
| .....agaCggaaccuacaagauca | 1 | 1 | seq |
| .....Uauggaaccuacaagauca | 4 | 1 | seq |
| .....Aauggaaccuacaagauca | 5 | 1 | seq |
| .....auggaaccuacaagauca | 1 | 0 | seq |