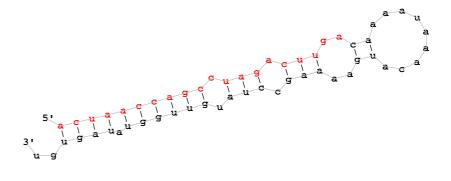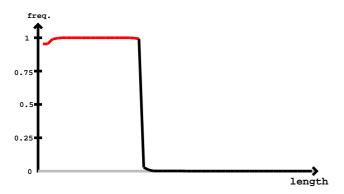miRBase precursor : chromosome_4_13569
Total read count : 800
chromosome_4_13569 read count : 799
remaining reads : 1

```
5'- acuaaccagccuagacuugacaaaauaaacaugaaaagccuauguugguauagugu -3'    exp
    (((((((((((.(((.(((..((..........))..))).))).)))))).))))..    reads    mm        sample
    acuaaccagccuagacuu.....................................    2        0         seq
    acuaaccagccuagacuug....................................    3        0         seq
    acuaaccagccuagacCug....................................    1        1         seq
    acuaaccGgccuagacuuga...................................    3        1         seq
    acuaaccagccuagCcuuga...................................    1        1         seq
    acuaaccagccuagGcuuga...................................    4        1         seq
    Ccuaaccagccuagacuuga...................................    2        1         seq
    acuaaccagccCagacuuga...................................    3        1         seq
    acuaaccagccuagUcuuga...................................    2        1         seq
    acuaacUagccuagacuuga...................................    2        1         seq
    acuaaccagccuagacuGga...................................    3        1         seq
    acuaaccagccuagaGuuga...................................    1        1         seq
    acuaGccagccuagacuuga...................................    4        1         seq
    acuaaccagccuagacuugC...................................    1        1         seq
    acGaaccagccuagacuuga...................................    2        1         seq
    Ucuaaccagccuagacuuga...................................    16       1         seq
    Gcuaaccagccuagacuuga...................................    59       1         seq
    acuaacAagccuagacuuga...................................    1        1         seq
    acuaaccagccuagacCuga...................................    2        1         seq
    acuaaccagccGagacuuga...................................    2        1         seq
    acuaaccagccuagacuAga...................................    1        1         seq
    acuaaAcagccuagacuuga...................................    2        1         seq
    acuaaccagccuagaUuuga...................................    2        1         seq
    acCaaccagccuagacuuga...................................    1        1         seq
    acuaaccagccuagacuCga...................................    2        1         seq
    acAaaccagccuagacuuga...................................    5        1         seq
    acuaaccagcGuagacuuga...................................    1        1         seq
    acuaaUcagccuagacuuga...................................    2        1         seq
    acuaaccagccuagacuuG....................................    5        1         seq
    acuaaccagccuaAacuuga...................................    1        1         seq
    acuaaGcagccuagacuuga...................................    1        1         seq
    acuaaccagccuGgacuuga...................................    6        1         seq
    acuaaccagcAuagacuuga...................................    1        1         seq
    acuaaccagUcuagacuuga...................................    1        1         seq
    acuaaccagAcuagacuuga...................................    1        1         seq
```

acuaaccagccuagacuugacaaaauaaacaugaaaagccuauguugguauagugu

```
acuaaccagccuagacuuga...................................     599     0     seq
acuaaccagccuagacuugaG..................................       4     1     seq
aAuaaccagccuagacuugac..................................       1     1     seq
acuaaccagccuagacuugac..................................       6     0     seq
acuaGccagccuagacuugaca.................................       1     1     seq
Ucuaaccagccuagacuugaca.................................       1     1     seq
acuaaccagccuagacuugaca.................................       2     0     seq
acGaaccagccuagacuugaca.................................       1     1     seq
acuaaccagccuagacGugaca.................................       1     1     seq
acuaaccagccuagacuugacaaaauaaa.........................       1     0     seq
.Auaaccagccuagacuuga...................................       3     1     seq
.Uuaaccagccuagacuuga...................................       2     1     seq
.cuaacGagccuagacuuga...................................       1     1     seq
.Guaaccagccuagacuuga...................................       1     1     seq
.cuaaccagccuagacuuga...................................      21     0     seq
.cuaaccagccuagacuugac.................................       1     0     seq
..uaaccagccuagacuuga...................................       5     0     seq
...Gaccagccuagacuugac.................................       2     1     seq
...aaccagccuagacuugaca.................................       1     0     seq
```