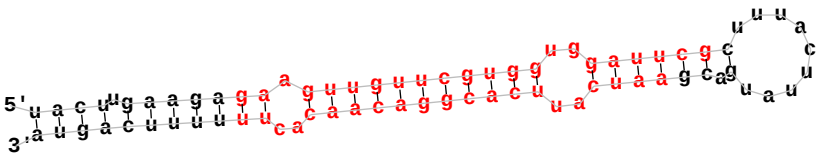
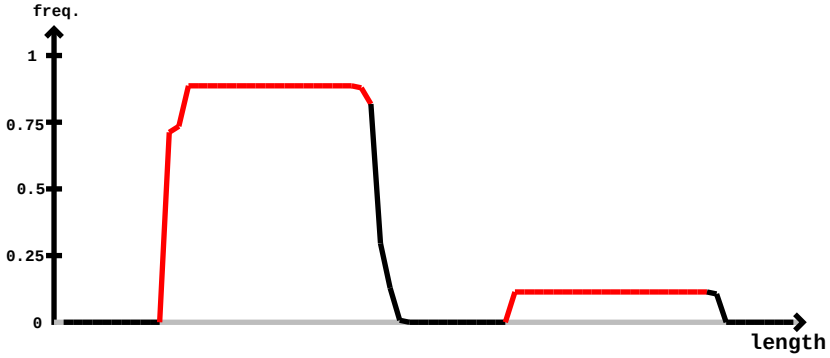
miRBase precursor : hsa-mir-382
Total read count : 132
hsa-miR-382-5p read count : 117
hsa-miR-382-3p read count : 15
remaining reads : 0





hsa-miR-382-3p

hsa-miR-382-5p

```
5'- uacuugaagagaagauuguucgugguggauucgcuuuacuuaugacgaaucauucacggacaacacuuuuucagua  -3'    exp
    ((((.(((((((.(((((((((((..(((((((.........).)))))))..)))))))))))..)))))))))))   reads     mm        sample
    ..........gaCguuguucgugguggauuc.............................................   1         1         Z1_
    ..........gaaguuguucgugguggauuc.............................................   6         0         Z1_
    ..........gaaguuguucguggAggauucg............................................   1         1         Z1_
    ..........gaaguuguucgugguggauucA............................................   3         1         Z1_
    ..........gaaguuguucguggCggauucg............................................   1         1         Z1_
    ..........gaaguuguucgugguggauucg............................................   21        0         Z1_
    ..........gaaguuguucgugguggauucgA...........................................   5         1         Z1_
    ............aguuguucgugguggauucg............................................   1         0         Z1_
    ............aguuguucgugguggauucA............................................   1         1         Z1_
    ...............................................aaucauucacggacaacacuu........   1         0         Z1_
    ...............................................aaucauucacggacaacacuuu.......   5         0         Z1_

    ..........gGaguuguucgugguggauuc.............................................   1         1         Z2_
    ..........gaaguuguucgugguggauucg............................................   8         0         Z2_
    ..........gaaguuguucgugguggauucU............................................   4         1         Z2_
    ...........aaguuguucgugguggauucg............................................   2         0         Z2_
    ............aguuguucgugguggauucgc...........................................   1         0         Z2_

    ..........gaUguuguucgugguggauu..............................................   1         1         GDM
    ..........gaaguuguucgugguggauucg............................................   20        0         GDM
    ..........gaaguuguucgugguggauucg............................................   20        0         GDM
    ..........gaaguuguucgugguggauucU............................................   1         1         GDM
    ..........gaaguuguucgCgguggauucg............................................   1         1         GDM
    ..........gaaguuguucgugguggauucA............................................   2         1         GDM
    ..........gaaguuguucgugguggauucg............................................   20        0         GDM
    ..........gaaguuguucgugguggauucgc...........................................   1         0         GDM
    ..........gaaguuguucgugguggauucgU...........................................   9         1         GDM
    ..........gaaguuguucgugguggauucgA...........................................   6         1         GDM
    ..........gaaguuguucgugguggauucgU...........................................   9         1         GDM
    ..........gaaguuguucgugguggauucgU...........................................   9         1         GDM
    ..........gaaguuguucgugguggauucgA...........................................   6         1         GDM
    ............aguuguucgugguggauucgcu.........................................   15        0         GDM
    ............aguuguucgugguggauucgcC.........................................   1         1         GDM
    ............aguuguucgugguggauucgcu.........................................   15        0         GDM
```

```
                                                  hsa-miR-382-3p
          hsa-miR-382-5p
uacuugaagagaagauguucgugguggauucgcuuuacuuaugacgaaucauucacggacaacacuuuuuucagua

............aguuguucgugguggauucgcu.........................................   15        0        GDM
............aguugCucgugguggauucgcuu........................................   1         1        GDM
...............................................aaucauucacggacaacacuuA........   2         1        GDM
...............................................aaucauucacggacaacacCuu........   2         1        GDM
...............................................aaucauucacggacaacacuuu........   5         0        GDM
...............................................aaucauucacggacaacacuuu........   5         0        GDM

..........gaaguuguucgugguggauucg...........................................   2         0        Z3_
...........aaguuguucgugguggauucg...........................................   1         0        Z3_
```