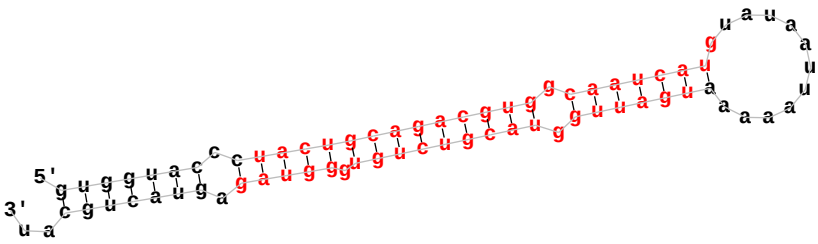
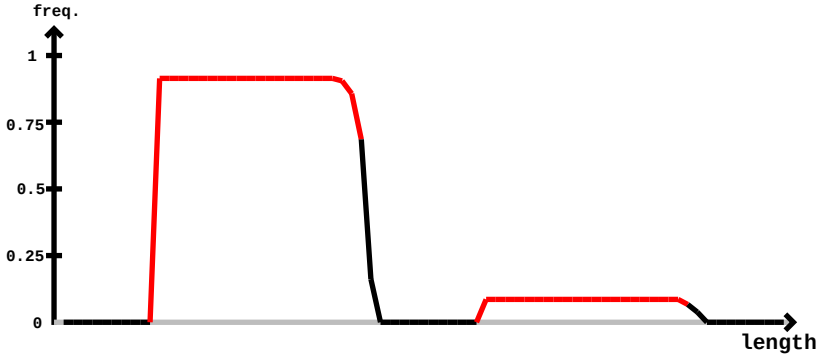
miRBase precursor : hsa-mir-509-3
Total read count : 105
hsa-miR-509-3p read count : 9
hsa-miR-509-3-5p read count96
remaining reads : 0





hsa-miR-509-3p

hsa-miR-509-3-5p

```
5'- gugguacccuacugcagacguggcaaucaugauaauuaaaaaugauugguacgucugugggugagaguacugcau  -3'    exp
    (((((((.(((((((((((((((.(((((((.............)))))))).)))))))))).)))))).)))))))..   reads      mm          sample
    .........uacugcagacguggcaaucaug.............................................   2          0           Z3_
    .........uacugcagacguggcaaucaugA............................................   1          1           Z3_

    .........uacugcagacguggcaaucaug.............................................   1          0           Z2_
    .........uacugcagacguggcaaucauA............................................   1          1           Z2_

    .........uacugcagacguggcaauc................................................   1          0           GDM
    .........uacugcagacguggcaauca...............................................   3          0           GDM
    .........uacugcagacguggcaauAa...............................................   1          1           GDM
    .........uacugcagacguggcaaucU...............................................   1          1           GDM
    .........uacugcagacguggcaauca...............................................   3          0           GDM
    .........uacugcagacguggcaaucau..............................................   14         0           GDM
    .........uacugcagacguggcaaucau..............................................   14         0           GDM
    .........uacugcagGcguggcaaucau..............................................   1          1           GDM
    .........uacugcagacguggcaaucau..............................................   14         0           GDM
    .........uacugcagacguggcaaucaC..............................................   1          1           GDM
    .........Aacugcagacguggcaaucaug.............................................   2          1           GDM
    .........uacugcagacguggcGaucaug.............................................   2          1           GDM
    .........uacugcagacguggcaaucaug.............................................   28         0           GDM
    .........uacugcagacguggcaaucaGg.............................................   5          1           GDM
    .........uacugcagacguggcaaucaug.............................................   28         0           GDM
    .........uacugcagacguggcaaucauA............................................   5          1           GDM
    .........uacugcagacguggcaaucauU............................................   1          1           GDM
    .........uacugcaUacguggcaaucaug.............................................   1          1           GDM
    .........uacugcagacguggcaaucaug.............................................   28         0           GDM
    .........uacugcagacguggcaaucauA............................................   5          1           GDM
    .........uacugcagacguggcaaucaugA............................................   10         1           GDM
    .........uacugcagacguggcaaucaugu............................................   4          0           GDM
    .........uacugcagacguggcaaucaugu............................................   4          0           GDM
    .........uacugcagacguggcaaucaugA............................................   10         1           GDM
    .........uacugcagacguggcaaucaugA............................................   10         1           GDM
    ........................................ugauugguacgucugugggua...............   2          0           GDM
    ........................................ugauugguacgucuguggguaA..............   1          1           GDM
    ........................................ugauugguacgucuguggguag..............   2          0           GDM
```

```
                                                hsa-miR-509-3p
          hsa-miR-509-3-5p
gugguacccuacugcagacguggcaaucauguauaauuaaaaaugauugguacgucugugggguagaguacugcau

.........................................ugauugguacgucugugggguaga.........    3          0          GDM
.........................................ugauugguacgucugugggguaga.........    3          0          GDM

.........uacugcagacguggcaaucau...........................................    2          0          Z1_
.........uacugcagacguggcaaucaug...........................................    6          0          Z1_
.........uacugcagacguggcaaucauA...........................................    1          1          Z1_
.........uacugcagacguggcaaucaugA..........................................    2          1          Z1_
.........................................ugauugguacgucugugggguaga.........    1          0          Z1_
```