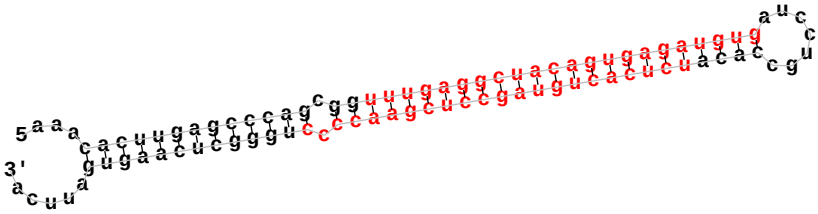
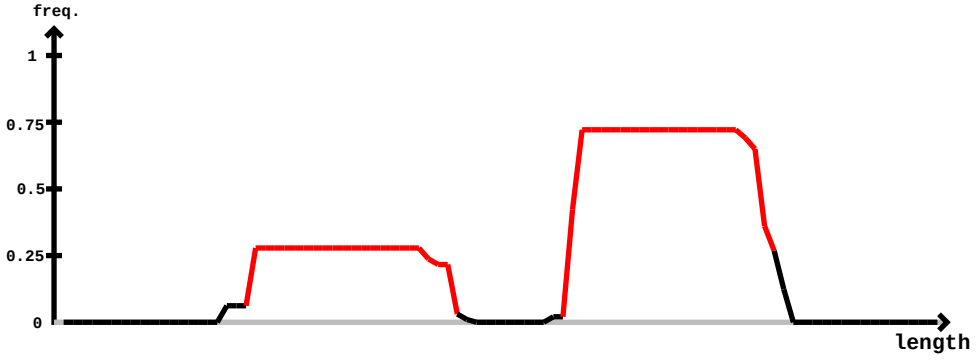
miRBase precursor : hsa-mir-1304
Total read count : 97
hsa-miR-1304-5p read count 21
hsa-miR-1304-3p read count 70
remaining reads : 6






```
                                   hsa-miR-1304-5p
                                                                   hsa-miR-1304-3p
5'- aaacacuugagcccagcggUUUGAGGCUACAGUGAGAUGUGauccugccacaucucacuguagccucgaacccccugggcucaagugauuca  -3'    exp
    ...(((((((((((((.((((((((((((((((((((((.......)))))))))))))))))))))))))))).))))))))))))))..... reads     mm        sample
    ...................................................caucucacuguagcAucgaa..................... 1         1         Z3_
    ...................................................ucucacuguagcAucgaacc...................... 4         1         Z3_
    ...................................................cucacuguagcAucgaacccc..................... 2         1         Z3_

    ...................................................ucucacuguagcAucgaacc...................... 8         1         Z1_
    ...................................................ucucacuguagcAucgaaccc..................... 3         1         Z1_
    ...................................................ucucacuguagcAucgaacccc.................... 2         1         Z1_
    ...................................................cucacuguagcAucgaacc...................... 3         1         Z1_
    ...................................................cucacuguagcAucgaaccc..................... 3         1         Z1_
    ...................................................cucacuguagcAucgaaccccu................... 2         1         Z1_

    ................cgguuugaggcuacagCgagau............................................... 1         1         Z2_
    ..................uuugaggcuacagCgagaugug.............................................. 1         1         Z2_
    ...................................................ucucacuguagcAucgaac...................... 3         1         Z2_
    ...................................................ucucaUuguagccucgaacc..................... 1         1         Z2_
    ...................................................ucucacuguagcAucgaacccc.................... 1         1         Z2_
    ...................................................cucacuguagcAucgaaccccu................... 5         1         Z2_

    ................cgguuugaggcuacGgugaga................................................ 3         1         GDM
    ................cgguuugaggcuacUgugaga................................................ 1         1         GDM
    ................cgguuugaggcuacGgugaga................................................ 3         1         GDM
    ................cgguuugagAcuacagugagau............................................... 1         1         GDM
    ..................uuugaggcuacagGgagaugu............................................... 18        1         GDM
    ..................uuugaggcuacagGgagaugu............................................... 18        1         GDM
    ..................uuugaggcuacagGgagaugug.............................................. 1         1         GDM
    ..................uuugaggcuacagGgagauguga............................................. 1         1         GDM
    ...................................................caucucacuguagcAucgaacc...................... 1         1         GDM
    ...................................................ucucacuguagcAucgaa....................... 2         1         GDM
    ...................................................ucucacuguagcAucgaac...................... 1         1         GDM
    ...................................................ucucacuguagcAucgaacc...................... 4         1         GDM
    ...................................................ucucacuguagcGucgaacc...................... 1         1         GDM
    ...................................................ucucacuguagcAucgaacc...................... 4         1         GDM
    ...................................................ucucacuguagcAucgaaccc..................... 1         1         GDM
    ...................................................ucucacuguagcAucgaacccc.................... 7         1         GDM
```

```
                    hsa-miR-1304-5p
                                                hsa-miR-1304-3p
aaacacuugagcccagcgguuugaggcuacagugagaugugauccugccacaucucacuguagccucgaaccccugggcucaagugauuca

...................................................ucucacuguagcAucgaacccc................   7          1          GDM
...................................................ucucacuguagcAucgaacccc................   7          1          GDM
...................................................ucucacuguagcAucgaaccccu...............   1          1          GDM
...................................................cucacuguagcAucgaacc...................   6          1          GDM
...................................................cucacuguagcAucgaaccc..................   2          1          GDM
...................................................cucacuguagcAucgaacccc.................   2          1          GDM
...................................................cucacuguagcAucgaaccccu................   4          1          GDM
...................................................cucacuguagcAucgaaccccu................   4          1          GDM
```