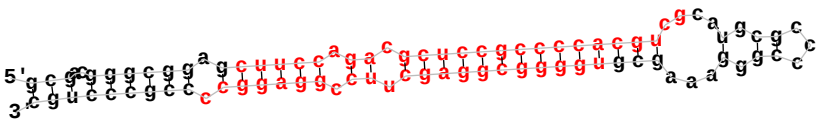
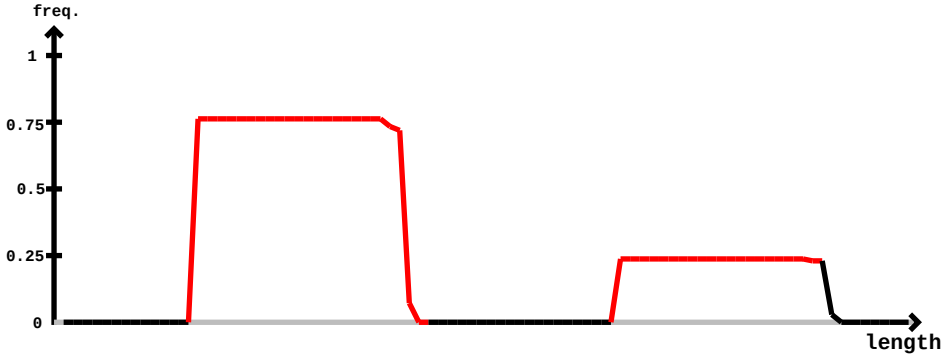
miRBase precursor : hsa-mir-3180-2
Total read count : 172
hsa-miR-3180-5p read count 106
hsa-miR-3180-3p read count 33
hsa-miR-3180 read count : 33
remaining reads : 0






hsa-miR-3180-3p
hsa-miR-3180
hsa-miR-3180-5p

```
5'- gcgacgggcggagcuuccagacgcuccgccccacgucgcaugcgccccgggaaagcgugggcggagcuuccggaggcccgcccugc  -3'    exp
    (((..((((((.((((((.((.(((((((((((((....(.((...)).)...)))))))))))))).)).)))))).)))))))))   reads    mm    sample
    ............cuuccagacgcuccgccccac..................................................   1        0     Z3_
    ............cuuccagacgcuccgccccacg.................................................   11       0     Z3_
    ............cuuccagacgcuccgccccacgA................................................   5        1     Z3_
    ...........................................................uggggcggagcuuccggaggcU.....   1        1     Z3_
    ...........................................................uggggcggagcuuccggaggcc.....   1        0     Z3_
    ...........................................................uggggcggagcuuccggaggccU....   3        1     Z3_

    ............cuuccagacgcuccgccccacg.................................................   10       0     Z2_
    ............cuuccagacgcuccgccccacgA................................................   1        1     Z2_
    ...........................................................uggggcggagcuuccggaggcU.....   1        1     Z2_
    ...........................................................uggggcggagcuuccggaggcA.....   1        1     Z2_

    ............cuuccagacgcuccgccccacC................................................   1        1     Z1_
    ............cuuccagacgcuccgccccacg.................................................   16       0     Z1_
    ............cuuccagacgcuccgccccacA................................................   1        1     Z1_
    ............cuuccagacgUuccgccccacg.................................................   2        1     Z1_
    ............cuuccagacgcuccgccccacU................................................   2        1     Z1_
    ............cCuccagacgcuccgccccacg................................................   2        1     Z1_
    ...........................................................uggggcggagcuuccggagA......   1        1     Z1_
    ...........................................................uggggcggagcuuccggaggcc.....   13       0     Z1_
    ..........................................................Cggggcggagcuuccggaggcc.....   1        1     Z1_
    ...........................................................uggggcggagcuuccggaggcU.....   1        1     Z1_
    ...........................................................uggggcggagcuuccggaggccA....   1        1     Z1_

    ............cuuccagacgcuccgcccca...................................................   4        0     GDM
    ............cuuccagacgcuccgcccca...................................................   4        0     GDM
    ............cuuccagacgcuccgcccca...................................................   4        0     GDM
    ............cuuccagacgcuccgccccac..................................................   1        0     GDM
    ............cuuccagacgcCccgccccacg.................................................   2        1     GDM
    ............cuuccagacgcuccgccccacg.................................................   40       0     GDM
    ............cuuccagacgUuccgccccacg.................................................   1        1     GDM
    ............cuuccagacgcuccgccccacA................................................   1        1     GDM
    ............cuuccagacgcuccgccccacg.................................................   40       0     GDM
```

hsa-miR-3180-3p

hsa-miR-3180

gcgacgggcggagcccccacgucgcaugcgccccgggaaagcguggggcggagcuuccggaggccccgcccugc

| | | | |
|---|---|---|---|
| .....cuuccagacgcuccgccccacC..... | 1 | 1 | GDM |
| .....cuuccagacgcuccgccccacg..... | 40 | 0 | GDM |
| .....cuuccagacgcuccgccccacgA..... | 4 | 1 | GDM |
| .....cuuccagacgcuccgccccacgA..... | 4 | 1 | GDM |
| .....uggggcggagcuuccggaggcU..... | 3 | 1 | GDM |
| .....uggggcggagcuuccggaggcA..... | 3 | 1 | GDM |
| .....uggggcggagcuuccggagAcc..... | 2 | 1 | GDM |
| .....uggggcggagcuuccggaggcc..... | 1 | 0 | GDM |