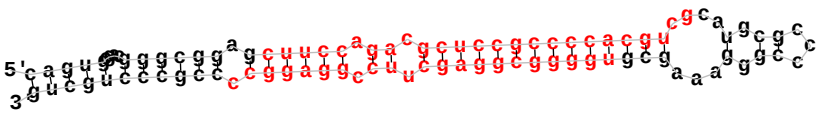
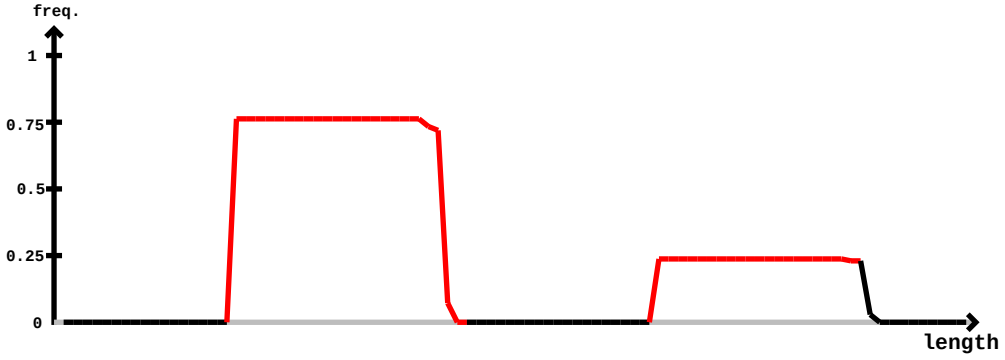
miRBase precursor : hsa-mir-3180-1
Total read count : 172
hsa-miR-3180-5p read count 106
hsa-miR-3180-3p read count 33
hsa-miR-3180 read count : 33
remaining reads : 0





hsa-miR-3180-3p

hsa-miR-3180-5p

hsa-miR-3180

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

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

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

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

hsa-miR-3180-3p

hsa-miR-3180-5p

cagugcgacgggcggagcuuccagacgcuccgccccacgucgcaugcgccccgggaaagcgugggucggagcuccggaggcccccgcccugcug

.....cuuccagacgcuccgccccacA.....	1	1	GDM
.....cuuccagacgcuccgccccacg.....	40	0	GDM
.....cuuccagacgcuccgccccacgA.....	4	1	GDM
.....cuuccagacgcuccgccccacgA.....	4	1	GDM
.....uggggcggagcuuccggaggcU.....	3	1	GDM
.....uggggcggagcuuccggaggcA.....	3	1	GDM
.....uggggcggagcuuccggaggcc.....	1	0	GDM
.....uggggcggagcuuccggagAcc.....	2	1	GDM