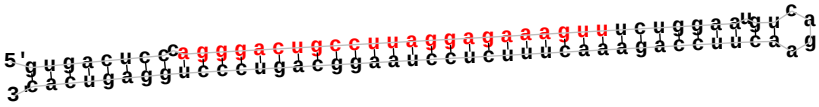
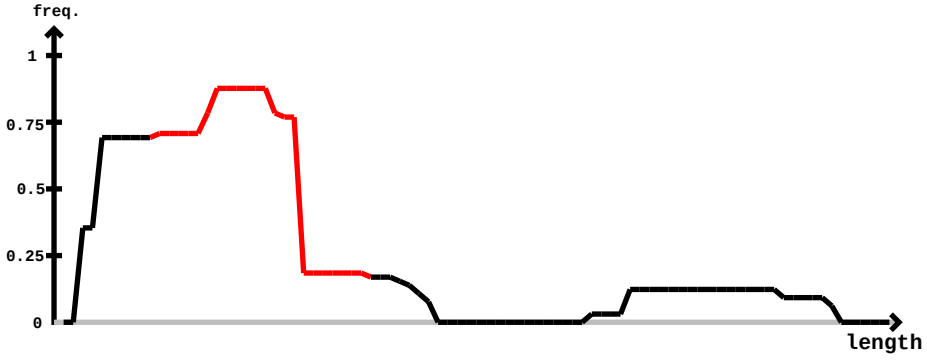
miRBase precursor	: hsa-mir-3199-2
Total read count	: 65
hsa-miR-3199 read count	: 7
remaining reads	: 58






```
                hsa-miR-3199
5'- gugacucccagggacugccuuaggagaaaguuucuggaaugucagaacuuccagaaacuuucuccuaaggcagucccuggagucac  -3'    exp
    (((((((.(((((((((((((((((((((((((((((((.((....)))))))))))))))))))))))))))))))))))))))))))   reads      mm         sample
    .ugacucccagggacugccuua.............................................................   1          0          Z1_
    .ugacucccagggacugccuuagg...........................................................   5          0          Z1_
    .ugacucccagggacCgccuuagg...........................................................   1          1          Z1_
    ...acucccagggacugccuuagg...........................................................   4          0          Z1_
    ...acucccagggacugccuuagA...........................................................   2          1          Z1_
    .........agggacugccuuaggagaaagu....................................................   1          0          Z1_
    ...............ugccuuaggagaaaguuucugg..............................................   2          0          Z1_
    ...............ugccuuaggagaaaguuucugga.............................................   3          0          Z1_

    .ugacucccagggacugccuu..............................................................   2          0          Z2_
    .ugacucccagggacugccuuagA...........................................................   1          1          Z2_
    .ugacucccagggacugccuuagg...........................................................   6          0          Z2_
    .............cugccuuaggagaaaguuucuU................................................   1          1          Z2_
    .............cugccuuaggagaaaguuucug................................................   1          0          Z2_
    .............cugccuuaggagaaaguuucugga..............................................   1          0          Z2_
    ......................................................aaacuuucuccuaaggcagu..........   2          0          Z2_

    .ugacucccagggacugccuu..............................................................   3          0          GDM
    .ugacucccagggacugccuC..............................................................   1          1          GDM
    .ugacucccagggacugccuuagg...........................................................   2          0          GDM
    .ugacucccagggacugccuuagg...........................................................   2          0          GDM
    .ugacucccagggacugccuuagA...........................................................   1          1          GDM
    ...acucccagggacugccuuagg...........................................................   9          0          GDM
    ...acucccagggacugccuuagA...........................................................   1          1          GDM
    ...acucccagggacugccuuagg...........................................................   9          0          GDM
    ...acucccagggGcugccuuagg...........................................................   3          1          GDM
    .............cugccuuaggagaaaguuuc.................................................   1          0          GDM
    .............cugccuuaggagaaagCuucu................................................   1          1          GDM
    ...............ugccuuaggagaaaguuucugga............................................   1          0          GDM
    ......................................................uuucuccuaaggcagucccuA.......   1          1          GDM
    ......................................................uuucuccuaaggcagucccug.......   1          0          GDM
    ......................................................uuucuccuaaggcagucccugg......   1          0          GDM

    ...acucccagggacugccuuagg...........................................................   3          0          Z3_
    ......................................................uuucuccuaaggcagucccugA......   1          1          Z3_
```

hsa-miR-3199

```
gugacucccagggacugccuuaggagaaaguuucuggaaugucagaacuuccagaaacuuucuccuaaggcagucccuggagucac
.............................................................uuucuccuaaggcagucccugg.....    2          0          Z3_
```