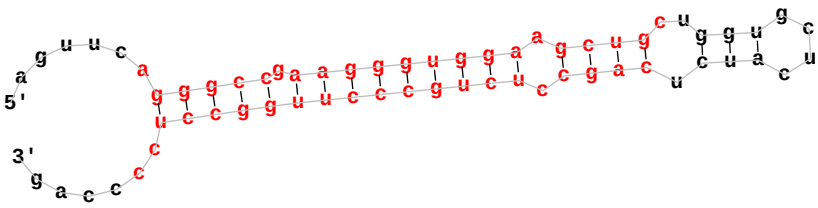
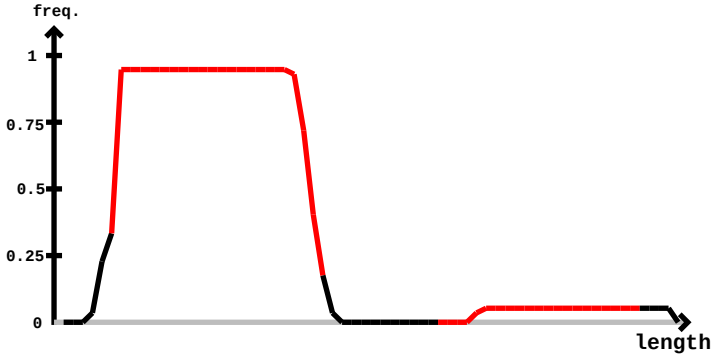
miRBase precursor : hsa-mir-6877
Total read count : 57
hsa-miR-6877-5p read count 52
hsa-miR-6877-3p read count 3
remaining reads : 2






```
                                                  hsa-miR-6877-3p
          hsa-miR-6877-5p
5'- aguucagggccgaagggugggaagcugcuggugcucaucucagccucugcccuuggccucccccag  -3'    exp
    ......(((((.(((((((((.((((..(((....))).)))).)))))))))))))).....   reads      mm         sample
    ...ucagggccgaagggugggaaA.......................................   1          1          Z1_
    ...ucagggccgaagggugggaagU......................................   2          1          Z1_
    ....cagggccgaagggugggaagcu.....................................   4          0          Z1_
    .....agggccgaagggugggaagcC.....................................   1          1          Z1_
    .....agggccgaagggugggaagcu.....................................   3          0          Z1_
    .....agggccgaagggugggaagcug....................................   7          0          Z1_
    .....agggccgaagggugggaagcugcu..................................   1          0          Z1_
    ........................................cucugcccuuggccuccccU.   1          1          Z1_

    ....cagggccgaagggugggaagcugU...................................   2          1          Z2_
    .....agggccgaagggugggaagcug....................................   2          0          Z2_
    .....agggccgaagggugggaagcugA...................................   3          1          Z2_
    .....agggccgaagggugggaagcugU...................................   2          1          Z2_

    ..uucagggccgaagggugggaagA......................................   2          1          GDM
    ...ucagggccgaagggugggaagc......................................   3          0          GDM
    ...ucagggccgaagggugggaagU......................................   4          1          GDM
    ...ucagggccgaagggugggaagc......................................   3          0          GDM
    ...ucGgggccgaagggugggaagc......................................   1          1          GDM
    .....agggccgaagggugggaagcu.....................................   10         0          GDM
    .....agggccgaagggugggaagcu.....................................   10         0          GDM
    .....agggccgaagggugggaagcug....................................   1          0          GDM
    .....agggccgGagggugggaagcug....................................   2          1          GDM
    .....agggccgaagggugggaagcGg....................................   1          1          GDM
    .....agggccgaagggugggaagcugU...................................   1          1          GDM
    .....agggccgaagggugggaagcugcu..................................   1          0          GDM
    ........................................ccucugcccuuggccucccca.   1          0          GDM
    ........................................ccucugcccuuggccucccccG.   1          1          GDM
```