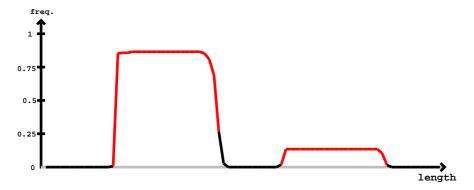miRBase precursor   : hsa-mir-185
Total read count   : 252
hsa-miR-185-5p read count : 218
hsa-miR-185-3p read count : 34
remaining reads   : 0


```
                                                          hsa-miR-185-3p

                             hsa-miR-185-5p

5'- aggggcgagggauuggagagaaaggcaguuccugauggucccuccccagggggcuggcuuuccucugguccuucccucccca   -3'      exp
    .(((((.(((((((((((((.(((((.((((((((((..((.......)).)))))))))).))))))))))))))).))))).      reads        mm        sample
    ............uuggagagaaaggcaguuccGg.................................................      1            1         seq
    ............uuggagagaaaggcaguuccuga...............................................      1            0         seq
    .............uggagagaaaggcaguuc...................................................      3            0         seq
    .............uggagagaaaggcaguuUc..................................................      1            1         seq
    .............uggagagaaaggcaguucc..................................................      10           0         seq
    .............uggagagaaaggcaguuccu.................................................      29           0         seq
    .............uggagCgaaaggcaguuccu.................................................      1            1         seq
    .............uggagagaaaggcCguuccug................................................      1            1         seq
    .............uggagagaaaggcaguuccug................................................      102          0         seq
    .............uggagagaaaggcaguuccugU...............................................      3            1         seq
    .............uggagagaaaggcaguuccugG...............................................      1            1         seq
    .............uggagagaaaggcaguuccuga...............................................      54           0         seq
    .............uggagagaaaggcaguuUcuga...............................................      1            1         seq
    .............uggagagaaaggcaguuccugaG..............................................      4            1         seq
    .............uggagagaaaggcaguuccugaA..............................................      2            1         seq
    .............uggagagaaaggcaguuccugau..............................................      1            0         seq
    ..............Ugagagaaaggcaguuccug................................................      1            1         seq
    ...............agagaaaggcaguuccug.................................................      2            0         seq
    ..................................................cagggcuggcuuCccucuggu...........      1            1         seq
    ..................................................cagggcuggcuuuccucuggu...........      3            0         seq
    ..................................................cagggcuggcuuuccucuggA...........      1            1         seq
    ...................................................agggcuggcuuuccucugg............      7            0         seq
    ...................................................aAgggcuggcuuuccucugg...........      1            1         seq
    ...................................................Cgggcuggcuuuccucuggu...........      3            1         seq
    ...................................................agggcuggcuuuccucuggu...........      13           0         seq
    ...................................................agggcuggcuuuccucuggA...........      1            1         seq
    ...................................................agggcuggcuuuccucugguU..........      1            1         seq
    ...................................................agggcuggcuuuccucugguA..........      3            1         seq
```