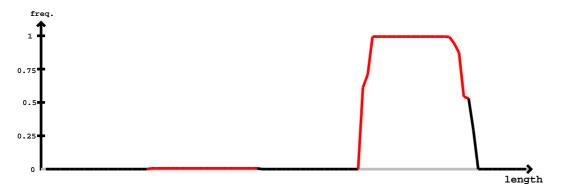miRBase precursor   : hsa-mir-3605
Total read count   : 330
hsa-miR-3605-5p read count 2
hsa-miR-3605-3p read count 328
remaining reads   : 0

hsa-miR-3605-3p

hsa-miR-3605-5p

```
5'- acuuuauacguguaauuugaugaggauggauagcaaggaagccgcucccaccugacccucacggccuccguguuaccuguccucagguggggacgcucg  -3'      exp
     .........(((((.........(((((((((.(((((.(((.((((.................))))).))).))))).))))))))).........))))...   reads      mm       sample
     ...................ugaggauggauagcaaggaagcc..........................................................      2          0        seq
     .....................................................ccuccguguuaccuguccc...............              1          0        seq
     .....................................................ccuccguguuaccuguccu...............              16         0        seq
     .....................................................ccuccguguuaccuguccuc..............              21         0        seq
     .....................................................Ucuccguguuaccuguccucu.............              1          1        seq
     .....................................................ccuccguguuaccuguccucu.............              100        0        seq
     .....................................................ccuccgAguuaccuguccucu.............              1          1        seq
     .....................................................ccuccguguuaccuguccucua............              3          0        seq
     .....................................................ccuccgugCuaccuguccucuag...........              1          1        seq
     .....................................................ccuccguguuaccuguccucuaU...........              1          1        seq
     .....................................................ccuccguguuaccuguccucuag...........              39         0        seq
     .....................................................ccuccguguuaccuguccucuagA..........              13         1        seq
     .....................................................ccuccguguuaccuguccucuagU..........              5          1        seq
     ......................................................cuccguguuaccuguccuc...............              2          0        seq
     ......................................................cuccguguuaccuguccucu.............              7          0        seq
     ......................................................cuccguguuaccuguccucua............              1          0        seq
     ......................................................cuccguguuaccuguccucuag...........              13         0        seq
     ......................................................cuccguguuaccuguccucuagA..........              6          1        seq
     ......................................................cuccguguuaccuguccucuagU..........              3          1        seq
     .......................................................uccgCguuaccuguccucuag...........              1          1        seq
     .......................................................uccgguuaccuguccucuag............              24         0        seq
     .......................................................uccgguuaccuguccucuagU...........              15         1        seq
     .......................................................uccgguuaccuguccucuagA...........              54         1        seq
```