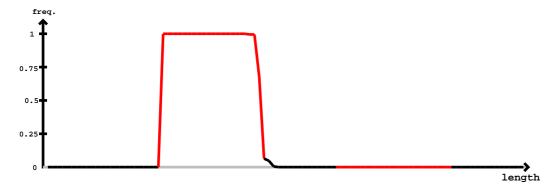miRBase precursor : hsa-mir-589
Total read count : 188
hsa-miR-589-5p read count : 188
hsa-miR-589-3p read count : 0
remaining reads : 0

hsa-miR-589-5p

hsa-miR-589-3p

```
5'- uccagccugugcccagcagccccugagaaccacgucugcucugagcuggguacugccuguucagaacaaaugccgguucccagacgcugccagcuggcc  -3'     exp
     .(((((.((......(((((..(((.(((((((.(((.((.(((((((((.((......)).)))))))).)).))).))))).))).))))))))))))..    reads      mm        sample
     .....................ugagaaccacgucugcuG............................................................       1          1         seq
     .....................ugagaaccacgucugcucug..........................................................       57         0         seq
     .....................ugagaaccacgucugcuUug..........................................................       1          1         seq
     .....................ugagCaccacgucugcucug..........................................................       2          1         seq
     .....................ugagaaccacgucugcCcuga.........................................................       1          1         seq
     .....................ugagaaccacgucugcucuga.........................................................       111        0         seq
     .....................ugagaaccacCucugcucuga.........................................................       1          1         seq
     .....................ugGgaaccacgucugcucuga.........................................................       1          1         seq
     .....................ugagaaccaAgucugcucuga.........................................................       1          1         seq
     .....................ugagaaccacgucugcucugaA........................................................       1          1         seq
     .....................ugagaaccacgucugcucugag........................................................       2          0         seq
     .....................ugagCaccacgucugcucugagc.......................................................       1          1         seq
     .....................ugagaaccacgucugcucugagc.......................................................       4          0         seq
     .....................ugagaaccacgucugcucugagA.......................................................       3          1         seq
     .....................ugagaaccacgucugcucugagcu......................................................       1          0         seq
```