miRBase precursor   : hsa-mir-125a
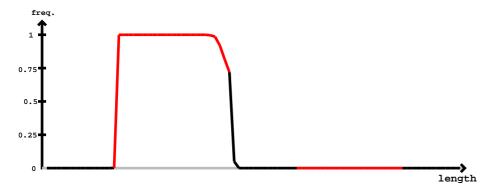Total read count   : 563
hsa-miR-125a-5p read count 563
hsa-miR-125a-3p read count 0
remaining reads   : 0


```
                    hsa-miR-125a-5p

                                          hsa-miR-125a-3p
5'- ugccagucucuaggucccugagacccuuuaaccugugaggacauccagggucacaggugagguucuugggagccuggcgucuggcc  -3'    exp
   .(((((.(.((((((.(((((((.((((..(((((((((((....))....)))))))))))).))))))).)))))).).))))).    reads      mm        sample
...........Uucccugagacccuuuaaccug...........................................................     1         1         seq
...............ucccugagacccuuuaacc...........................................................     2         0         seq
...............ucccugagacccuuuaaccu..........................................................     6         0         seq
...............ucccugagaUccuuuaaccug.........................................................     1         1         seq
...............ucccugagacccuuuaaccug.........................................................    35         0         seq
...............ucccugagacccuuuaaccugA........................................................     2         1         seq
...............ucccugagacccuuuaaccugG........................................................     1         1         seq
...............ucccugagacccuuuaaccugu........................................................    55         0         seq
...............ucccugagacccuuuaaccugug.......................................................    52         0         seq
...............ucccugagacccuuuaCccugug.......................................................     1         1         seq
...............uccccCgagacccuuuaaccugug......................................................     1         1         seq
...............ucccugagCccuuuaaccugug........................................................     1         1         seq
...............ucccugagacccuuuaaccugugU......................................................    14         1         seq
...............ucccugagacccuuuaaccuguga......................................................   287         0         seq
...............ucccugagacccuuuaaccCguga......................................................     1         1         seq
...............ucccugagCccuuuaaccuguga.......................................................     2         1         seq
...............ucccugagacccuuuCaccuguga......................................................     1         1         seq
...............ucccugagacccuuuaaccugugG......................................................    71         1         seq
...............ucccugagacUcuuuaaccuguga......................................................     1         1         seq
...............ucccugagacccuuuaaccugugaA.....................................................    26         1         seq
...............ucccugagacccuuuaaccugugaU.....................................................     2         1         seq
```