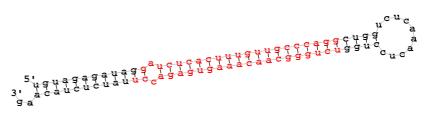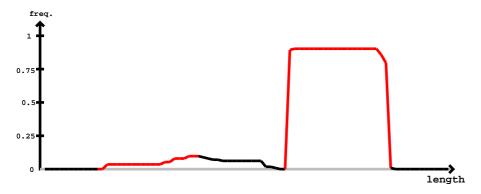miRBase precursor : hsa-mir-1285-1
Total read count : 113
hsa-miR-1285-5p read count 4
hsa-miR-1285-3p read count 102
remaining reads : 7

freq.

1

0.75

0.5

0.25

0

length

hsa-miR-1285-3p

hsa-miR-1285-5p

```
5'- uguagagauaggaucucacuuuguugcccaggcuggucucaaacuccuggucugggcaacaaagugagaccuuaucucuacaag    -3'    exp
     (((((((((((.((((((((((((((((((((((..........)))))))))))))))))))))).)))))))))))..      reads       mm        sample
     ...........aucucacuuuguugcccagg.....................................................     1          0         seq
     ...........aucucacuuuguugcccaggc....................................................     1          0         seq
     ...........aucucacuuuguugcccaggcu...................................................     1          0         seq
     ...........aucucacuuuguugcccaggcuAg.................................................     1          1         seq
     ....................ugcccaggcugCucucaaacu...........................................     2          1         seq
     ......................cccaggcuggucucaaacu...........................................     1          0         seq
     ......................cccaggcuggucucGaacu...........................................     2          1         seq
     ........................aggcuggucucGaacucc.........................................      1          1         seq
     ........................aggcuggucucaaacuccu.........................................     1          0         seq
     ...............................................ucugggcaacaaagugaga..................     5          0         seq
     ...............................................ucugggcaacaaagugagac.................     7          0         seq
     ...............................................ucugggcaacaaagugagacc................    83          0         seq
     ...............................................ucugggcaCcaaagugagacc................     1          1         seq
     ...............................................ucugggcaacaaagugagacU................     1          1         seq
     ...............................................ucCgggcaacaaagugagacc................     1          1         seq
     ...............................................ucugggcaaUaaagugagacc................     1          1         seq
     ...............................................ucugggcaacaaUgugagacc................     1          1         seq
     ...............................................ucugggcaacaaagugagaccu...............     1          0         seq
     ...............................................cugggcaacaUagugagacc.................     1          1         seq
```