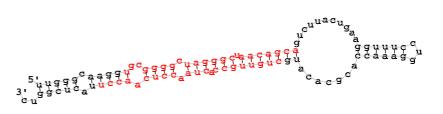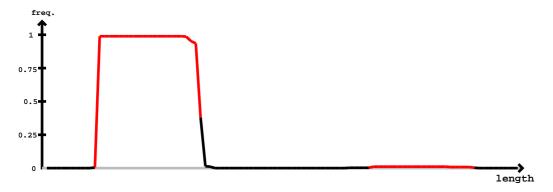miRBase precursor : hsa-mir-744
Total read count : 356
hsa-miR-744-5p read count : 352
hsa-miR-744-3p read count : 3
remaining reads : 1

freq.

length

hsa-miR-744-5p

hsa-miR-744-3p

```
5'- uugggcaaggugcggggcuagggcuaacagcagucuuacugaagguuuccuggaaaccacgcacaugcguugccacuaaccucaaccuuacucgguc  -3'   exp
    (((((.(((((..(((((.(((((((.((((((((...........((((((...))))))......)))))))))).))).))))).))))).)))))..    reads    mm    sample
    ........Uugcggggcuagggcuaacagc...............................................................         1        1     seq
    ........Uugcggggcuagggcuaacagca..............................................................         1        1     seq
    ........ugcggggcuagggcuaac...................................................................         1        0     seq
    ........ugcggggcuagggcuaaca..................................................................        10        0     seq
    ........ugcggggcuagggcuaacU..................................................................         2        1     seq
    ........ugcggggcuagggcuaacag.................................................................         6        0     seq
    ........ugcggggcuGgggcuaacagc................................................................         1        1     seq
    ........ugcggggcuagggcuaacagc................................................................       180        0     seq
    ........ugcggggcuagggcuaacagA................................................................        11        1     seq
    ........ugcggggcuagggcuaacaCc................................................................         1        1     seq
    ........ugcggggcuagggcuaacagU................................................................         4        1     seq
    ........ugcggggcuagggcuaacagAa...............................................................         3        1     seq
    ........ugcggggcuagggAuaacagca...............................................................         1        1     seq
    ........ugcggggcuagggcuaacagCU..............................................................        11        1     seq
    ........ugcggggcuagggcuaacagca..............................................................       114        0     seq
    ........ugcggggcuagggcuaacagcaU.............................................................         1        1     seq
    ........ugcggggcuagggcuaacagcaAu............................................................         2        1     seq
    ........ugcggggcuagggcuaacagcaCu............................................................         1        1     seq
    ........ugcggggcuagggcuaacagcaUu............................................................         1        1     seq
    .........................................................acaugcguugccacuaaccu...............         1        0     seq
    ..............................................................cguugccacuaaccucaacc..........         2        0     seq
    ..............................................................cguugccacuaaccucaaccu.........         1        0     seq
```