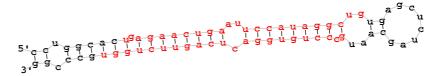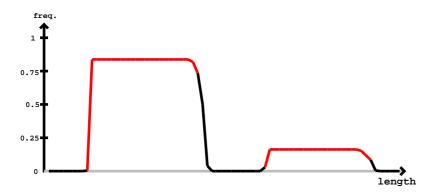miRBase precursor          : hsa-mir-146b
Total read count           : 203
hsa-miR-146b-5p read count 170
hsa-miR-146b-3p read count 33
remaining reads            : 0





```
                                                                    hsa-miR-146b-3p

                     hsa-miR-146b-5p
5'- ccuggcacugagaacugaauuccauaggcugugagcucuagcaaugcccuguggacucaguucgggugcccgg   -3'    exp
    ((.(((((((..((((((((((..((((((((((..((.........))..).))))))))).)))))))))))).))    reads    mm         sample
    .......Uugagaacugaauuccauaggcu.............................................    1        1          seq
    .......ugagaacugaauuccauaggc...............................................    3        0          seq
    .......ugagCacugaauuccauaggc...............................................    1        1          seq
    .......ugagCacugaauuccauaggcu..............................................    1        1          seq
    .......Cgagaacugaauuccauaggcu..............................................    1        1          seq
    .......ugagaacugaauuccauaggcA..............................................    2        1          seq
    .......ugagaacugaauuccauaggcu..............................................    11       0          seq
    .......ugagaacugaauuccUuaggcu..............................................    1        1          seq
    .......ugagaacugaauuccauaggcCg.............................................    1        1          seq
    .......ugagaacugaauuccauaggcug.............................................    46       0          seq
    .......ugaAaacugaauuccauaggcugu............................................    1        1          seq
    .......ugagaacugaauuccauaggcugG............................................    19       1          seq
    .......ugagaacugaauuccauaggcugu............................................    71       0          seq
    .......ugagaacugaauuccauaggcugA............................................    2        1          seq
    .......ugagCacugaauuccauaggcugu............................................    1        1          seq
    .......ugagaacugaauuccauaggcuguU...........................................    8        1          seq
    ..................................................ugcccuguggacucaguucug........    1        0          seq
    ..................................................ugcccuguggacucaguucgg.......    3        0          seq
    ..................................................ugcccuguggacucaguucggu......    1        0          seq
    ..................................................ugcccuCuggacucaguucggu......    1        1          seq
    ..................................................gcccuguggacucaguucu.........    2        0          seq
    ..................................................gcccuguggacucaguucg.........    6        0          seq
    ..................................................gcccuguggacucaguucgg........    4        0          seq
    ..................................................gcccuguggacucaguucggA.......    6        1          seq
    ..................................................gcccuguggacucaguucggu.......    8        0          seq
    ..................................................gcccuguggacucaguucgguU......    1        1          seq
```