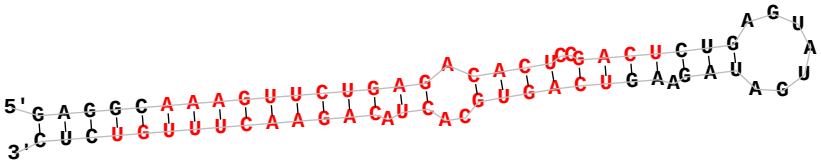
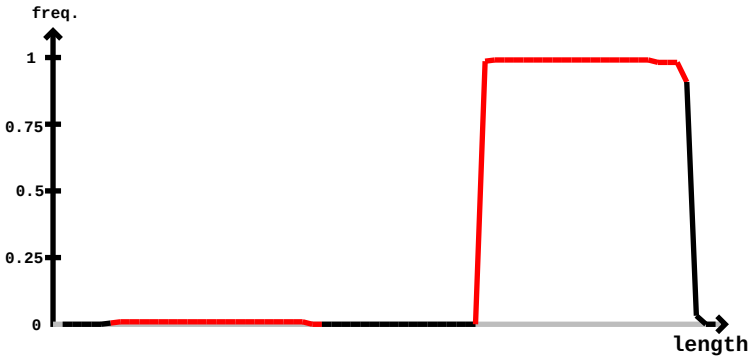
miRBase precursor : hsa-mir-148a
Total read count : 218
hsa-miR-148a-3p read count : 216
hsa-miR-148a-5p read count : 2
remaining reads : 216






hsa-miR-148a-3p

hsa-miR-148a-5p

```
5'- gaggcaaaguucugagacacuccgacucugaguaugauagaagucagugcacuacagaacuuugucuc  -3'    exp
    ((((((((((((((((.((((..(((((((.......))).)))))))..)).))))))))))))))    reads      mm         sample
    ....caaaguucugagacacuccga.....................................    1          0          hsa
    .....aaaguucugagacacuccga.....................................    1          0          hsa
    .....................................ucagugcacuacagaacA.......    1          1          hsa
    .....................................ucagugcaUuacagaacu.......    1          1          hsa
    .....................................ucagugcaUuacagaacuuug....    3          1          hsa
    .....................................ucagugcacuacagaacuuug....    12         0          hsa
    .....................................ucagugcacuacagaacuuGg....    1          1          hsa
    .....................................ucaUugcacuacagaacuuugu...    1          1          hsa
    .....................................ucagugcacuacagaacuuCgu...    1          1          hsa
    .....................................ucagugcacuacaAaacuuugu...    2          1          hsa
    .....................................ucagugcaUuacagaacuuugu...    3          1          hsa
    .....................................ucagugcacuacagaacuuugA...    2          1          hsa
    .....................................ucagugcacuacagaacuCugu...    1          1          hsa
    .....................................ucagugcacuacagaacuuuUu...    2          1          hsa
    .....................................ucagGgcacuacagaacuuugu...    1          1          hsa
    .....................................ucagugcacuacaUaacuuugu...    2          1          hsa
    .....................................ucagugcacGacagaacuuugu...    1          1          hsa
    .....................................ucagugcacuacagaacuuugu...    170        0          hsa
    .....................................ucagAgcacuacagaacuuugu...    1          1          hsa
    .....................................ucagugUacuacagaacuuugu...    1          1          hsa
    .....................................ucagCgcacuacagaacuuugu...    1          1          hsa
    .....................................ucCgugcacuacagaacuuugu...    1          1          hsa
    .....................................ucagugcacuacagaacuuuguU..    1          1          hsa
    .....................................ucagugcacuacagaacuuuguc..    6          0          hsa
    ........................................cagugcacuacagaacuuugu...    1          0          hsa
```