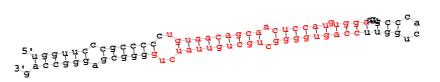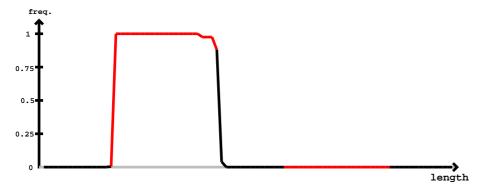miRBase precursor : hsa-mir-194-2
Total read count : 201
hsa-miR-194-5p read count : 201
hsa-miR-194-3p read count : 0
remaining reads : 0


```
                                                                    hsa-miR-194-3p

                                    hsa-miR-194-5p
5'- ugguucccgcccccuguaacagcaacuccauguggaagugcccacugguuccaguggggcugcuguuaucugggggcgagggccag  -3'      exp
    ((((((.((((((..((((((((((.((((((.(((...((((....))))))))))))))).)))))))))..))))))).)))))).       reads    mm       sample
    ............cuguaacagcaacuccaugugga...................................................          1        0        seq
    ............uguaacagcaacuccaug.......................................................          5        0        seq
    ............uguaCcagcaacuccaugugg....................................................          1        1        seq
    ............Cguaacagcaacuccaugugg....................................................          1        1        seq
    ............uguaacagcaacuccaugugg....................................................          17       0        seq
    ............uguaacagcaacuccaAgugga...................................................          1        1        seq
    ............Gguaacagcaacuccaugugga...................................................          1        1        seq
    ............uguaacagcaacuccauguggG...................................................          1        1        seq
    ............uguaacagcaacuccaugugga...................................................          155      0        seq
    ............uguaCcagcaacuccaugugga...................................................          10       1        seq
    ............uguaacagcaacuccauguggaa..................................................          6        0        seq
    ............uguaacagcaacucUauguggaa..................................................          1        1        seq
    ............uguaacagcaacuccauguggaU..................................................          1        1        seq
```