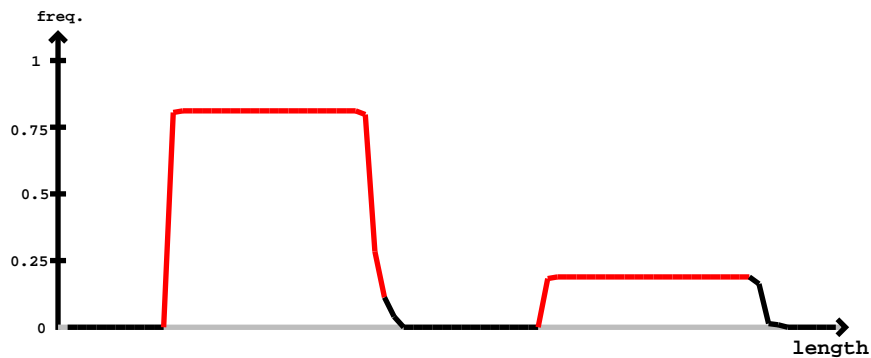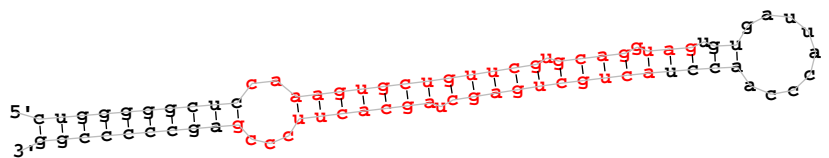miRBase precursor   : hsa-mir-93
Total read count   : 652
hsa-miR-93-5p read count : 529
hsa-miR-93-3p read count : 123
remaining reads   : 0

hsa-miR-93-5p

hsa-miR-93-3p

```
5'- cuggggcuccaaagugcguucgugcagguagugugauuacccaaccuacugcugagcuagcacuucccgagccccgg   -3'    exp
    ((((((((((..(((((((((((((.((((.(((.((.........))))))))))))))).)))))))...)))))))))    reads    mm      sample
    .........caaagugcguucgugcagg....................................................       9       0        seq
    .........caaagCgcguucgugcaggu...................................................       1       1        seq
    .........Uaaagugcguucgugcaggu...................................................       1       1        seq
    .........cUaagugcguucgugcaggu...................................................       1       1        seq
    .........caaagugcguucgugUaggu...................................................       1       1        seq
    .........caaagugcguucgugcaggA...................................................       2       1        seq
    .........caaagugGuucgugcaggu...................................................       1       1        seq
    .........caaagugcguucgugcaggu...................................................     319       0        seq
    .........caaagugcguuCcgugcaggu..................................................       1       1        seq
    .........caaagugcguucgugcaggG...................................................       1       1        seq
    .........caaCgugcguucgugcaggu...................................................       2       1        seq
    .........caaaAugcguucgugcaggu...................................................       1       1        seq
    .........caaagugcugAucgugcaggu..................................................       1       1        seq
    .........caaagugcguucgugUaggua..................................................       2       1        seq
    .........caaagugcguucgugcaggua.................................................     110       0        seq
    .........caaagugcguucgugcagguaU................................................       2       1        seq
    .........caaagugcguucgugcagguag...............................................      44       0        seq
    .........cCaagugcguucgugcagguag...............................................       1       1        seq
    .........caaagugcguucgugcagguagA..............................................      15       1        seq
    .........caaagugcguucgugcagguagu..............................................      10       0        seq
    .........aaagugcguucgugcaggu...................................................       2       0        seq
    .........Caagugcguucgugcaggu...................................................       1       1        seq
    .........aaagugcguucgugcagguagA...............................................       1       1        seq
    ..............................................Ccugcugagcuagcacuucccg.........       1       1        seq
    ..............................................acugcugagcuagcacuucccU.........       1       1        seq
    ..............................................acugcugagcuagcacuucccg.........      15       0        seq
    ..............................................acugcugagcuagcacuucccga.......      85       0        seq
    ..............................................Ccugcugagcuagcacuucccga.......       8       1        seq
    ..............................................acugcugagcuagcacuucccgaA......       3       1        seq
    ..............................................acugcugagcuagcacuucccgagA.....       6       1        seq
    ..............................................cugcugagcuagcacuucccga........       4       0        seq
```