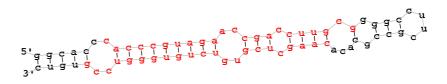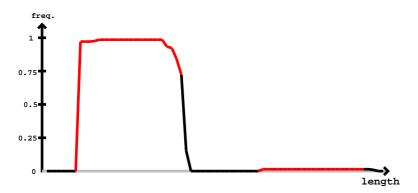miRBase precursor : hsa-mir-99b
Total read count : 209
hsa-miR-99b-5p read count : 206
hsa-miR-99b-3p read count : 3
remaining reads : 0

freq.

hsa-miR-99b-5p

hsa-miR-99b-3p

```
5'- ggcacccacccguagaaccgaccuugcggggccuucgccgcacacaagcucgugucuguggguccguguc  -3'     exp
    (((((..((((((((((..(((.(((((.(.(((....))).).)...))))).))).))))))))))..)))))     reads     mm          sample
    .....cacccguagaaccgaccC............................................              1         1           seq
    .....cacccguagaaccgaccu............................................              10        0           seq
    .....cacccguagaaccgaccuu...........................................              2         0           seq
    .....cacccguagaaccgaccuug..........................................              17        0           seq
    .....cacccguagaaccgaccuugA.........................................              1         1           seq
    .....cacccguagaaccgaccuugc.........................................              23        0           seq
    .....cacccguagaacAgaccuugc.........................................              1         1           seq
    .....cacccguagaaccgaccuugcg........................................              113       0           seq
    .....cacccguGgaaccgaccuugcg........................................              1         1           seq
    .....cacccguagaaccgaccuugcA........................................              1         1           seq
    .....cacccgCagaaccgaccuugcg........................................              1         1           seq
    .....cacccguagaaccgaccuugcgA.......................................              14        1           seq
    .....cacccguagaaccgaccuugcgU.......................................              18        1           seq
    .........ccguagaaccgaccuugcg.......................................              1         0           seq
    .........cguagaaccgaccuugcg.......................................              1         0           seq
    .........cguagCaccgaccuugcg.......................................              1         1           seq
    ........................................caagcucgugucuguggguccgA...              1         1           seq
    ........................................caagcucgugucuguggguccguU..              2         1           seq
```