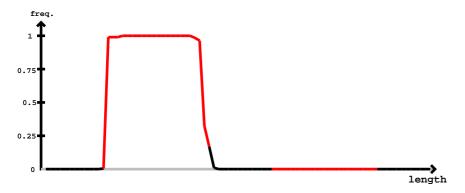miRBase precursor   : hsa-mir-100
Total read count   : 195
hsa-miR-100-5p read count : 195
hsa-miR-100-3p read count : 0
remaining reads   : 0

hsa-miR-100-5p

hsa-miR-100-3p

```
5'- ccguugccacaaacccguagauccgaacuugugguauuaguccgcacaagcuuguaucuauagguaugugucuguuagg  -3'    exp
    ((((....((((.(((.(((((((.(((.(((((((...........))))))).))).))))))).))).))))....))))  reads   mm      sample
    ..........aaacccguagauccgaacuug.............................................    1       0       seq
    ..........aacccguagauccgaacu...............................................    2       0       seq
    ..........aacccguagaucUgaacu...............................................    1       1       seq
    ..........aacccguagauccgaacuu..............................................    2       0       seq
    ..........aacccguagaucUgaacuu..............................................    2       1       seq
    ..........aacccguagaucUgaacuug.............................................    3       1       seq
    ..........aacccguagaCccgaacuug.............................................    1       1       seq
    ..........aacccguagauccgaacuuA.............................................    1       1       seq
    ..........Cacccguagauccgaacuug.............................................    8       1       seq
    ..........aacccguagauUcgaacuug.............................................    1       1       seq
    ..........aacccguagauccgaacuug.............................................    110     0       seq
    ..........Cacccguagauccgaacuugu............................................    2       1       seq
    ..........aacccguagaucUgaacuugu............................................    1       1       seq
    ..........aacccguagauccgaacuugu............................................    25      0       seq
    ..........aacccguagauccgaacuugG............................................    1       1       seq
    ..........aacccguagauccgaacuugGg...........................................    1       1       seq
    ..........Cacccguagauccgaacuugug...........................................    1       1       seq
    ..........aacccguagauccgaacuugug...........................................    28      0       seq
    ..........aacccguagauccgaacuuggg...........................................    1       0       seq
    ..........aacccguagauccgaacuugugA..........................................    1       1       seq
    ..............ccguagauccgaacuugu...........................................    1       0       seq
    ..............ccguCgauccgaacuugug..........................................    1       1       seq
```