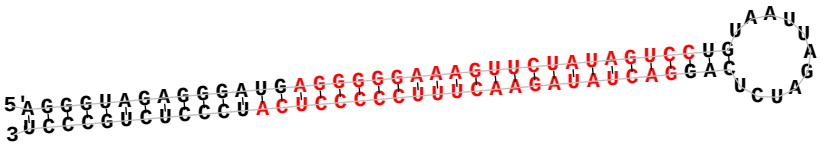
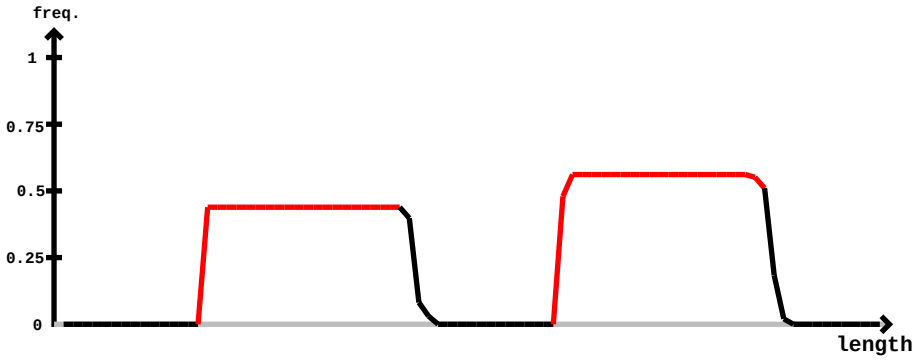
miRBase precursor	: hsa-mir-625
Total read count	: 98
hsa-miR-625-3p read count	: 55
hsa-miR-625-5p read count	: 43
remaining reads	: 55





hsa-miR-625-5p

hsa-miR-625-3p

```
5'- aggguagagggaugagggggaaaguucuauaguccuguaauuagaucucaggacuauagaacuuuccccccucaucccucugcccu  -3'    exp
    ((((((((((((((((((((((((((((((((((((((.....))))))))))))))))))))))))))))))))))))))))))   reads      mm         sample
    ..............agggggaaaguucuauagucU.....................................................   1          1          hsa
    ..............agggggaaaguucuauagucc.....................................................   3          0          hsa
    ..............agggggCaaguucuauaguccu....................................................   1          1          hsa
    ..............agggggaaaguucuauaguccu....................................................   28         0          hsa
    ..............aggggGaaguucuauaguccu.....................................................   1          1          hsa
    ..............agggggaaaguucuUuaguccu....................................................   1          1          hsa
    ..............agggggaaaguucuauaguccuU...................................................   3          1          hsa
    ..............agggggaaaguucuauaguccuA...................................................   2          1          hsa
    ..............agggggaaaguucuauaguccugu..................................................   1          0          hsa
    ..............agggggaaaguucuauaguccugA...................................................   2          1          hsa
    ..................................................gacuauGgaacuuuccccu.....................   1          1          hsa
    ..................................................gacuauagCacuuuccccuc....................   1          1          hsa
    ..................................................gacuauagaacuuuccccuc....................   3          0          hsa
    ..................................................gacuauagaacuuuccccuca...................   26         0          hsa
    ..................................................gacuauagaacuuuccccucau..................   2          0          hsa
    ..................................................gacuauagaacuuuccccucaA..................   12         1          hsa
    ..................................................gGcuauagaacuuuccccucau..................   1          1          hsa
    ..................................................gacuauagaacuuuccccucaAc.................   1          1          hsa
    ..................................................acuauagaacuuuccccuca....................   6          0          hsa
    ..................................................acuGuagaacuuuccccucau...................   1          1          hsa
    ..................................................acuauagaacuuuccccucaAc..................   1          1          hsa
```