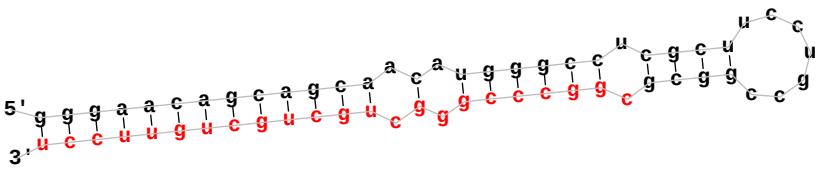
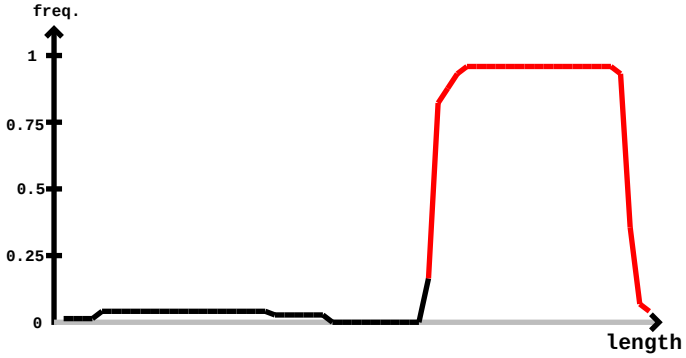
| | |
|---|---|
| miRBase precursor | : hsa-mir-1538 |
| Total read count | : 73 |
| hsa-miR-1538 read count | : 70 |
| remaining reads | : 3 |




```
                                        hsa-miR-1538
5'- gggaacagcagcaacaugggccucgcuuccugccggcgcggcccgggcugcugcuguuccu  -3'    exp
    ((((((((((((.(.((((((.((((.......)))).)))))).).)))))))))))))   reads      mm         sample
    ...aacagcagcaacaugggccucgUu...............................      2          1          Z1_
    ....................................gcggcccgggcugcugcuguu...      3          0          Z1_
    ....................................cggcccgggcugcugcuguC...      1          1          Z1_
    ....................................cggcccgggcugcugcuguu...      8          0          Z1_
    ....................................cggcccgggcugcugcuguA...      1          1          Z1_
    ....................................cggcccgggcugcugcuguuU..      3          1          Z1_
    ....................................cggcccgggcugcugcuguuccu      1          0          Z1_

    ....................................gcggcccgggcugcugcugGu...      2          1          GDM
    ....................................gcggcccgggcugcugcuguuU..      2          1          GDM
    ....................................cggcccgggcugcugcugu....      1          0          GDM
    ....................................cggcccgggcugcugcuguu...      15         0          GDM
    ....................................cggcccgggcugcugcuguu...      15         0          GDM
    ....................................cggcccgggcugcugcuguC...      3          1          GDM
    ....................................cggcccgggcugcugcuguC...      3          1          GDM
    ....................................cggcccgggcugcugcuguu...      15         0          GDM
    ....................................cggcccgggcugcugcuguuA..      2          1          GDM
    ....................................cggcccgggcugcugcuguuc..      2          0          GDM
    ....................................cggcccgggcugcugcuguuU..      3          1          GDM
    .....................................gcccgggcugcugcuguu...      2          0          GDM
    .....................................cccgggcugcugcuguucc.      2          0          GDM

    gggaacagcagcaacaugggc.....................................      1          0          Z2_
    ....................................gcggcccgggcugcugcuguC...      1          1          Z2_
    ....................................gcggcccgggcugcugcuguuU..      4          1          Z2_
    ....................................cggcccgggcugcugcugu....      1          0          Z2_
    ....................................cggcccgggcugcugcuguu...      3          0          Z2_
    ....................................cggcccgggcCgcugcuguu...      1          1          Z2_
    ....................................cggcccgggcugcugcuguuccu      1          0          Z2_
    .....................................ggcccgggcugcugcuguuU..      4          1          Z2_
    .....................................gcccgggcugcugcuguu...      1          0          Z2_
    ......................................gcccgggcugcugcuguuccu      1          0          Z2_

    ....................................cggcccgggcugcugcuguu...      1          0          Z3_
    ....................................cggcccgggcugcugcuguuc..      1          0          Z3_
```

**hsa-miR-1538**

**gggaacagcagcaacaugggccucgcuuccugccggcgcggcccgggcugcugcuguuccu**