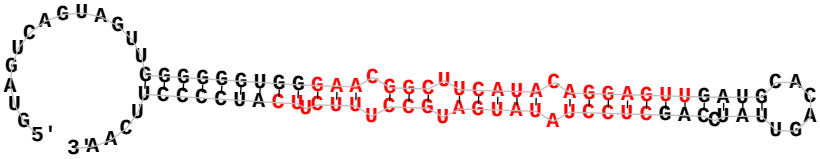
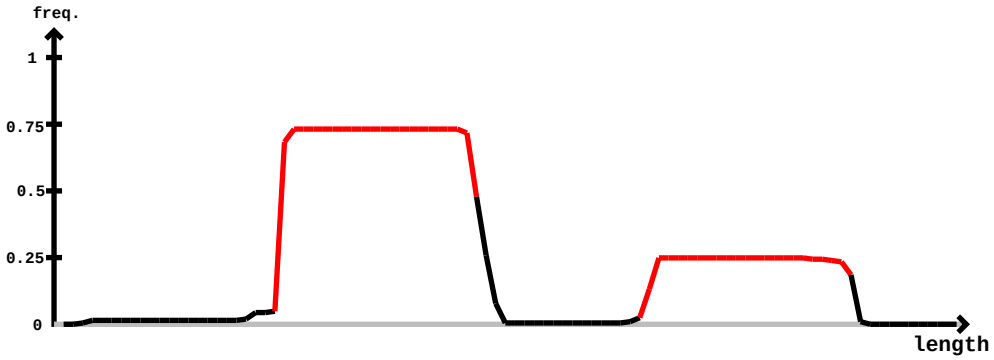
miRBase precursor : hsa-mir-337
Total read count : 205
hsa-miR-337-5p read count : 144
hsa-miR-337-3p read count : 51
remaining reads : 61





hsa-miR-337-5p

hsa-miR-337-3p

```
5'- guagucaguaguuggggggugggaacggcuucauacaggaguugaugcacaguuauccagcuccuauaugaugccuuucuucaucccuucaa  -3'   exp
    ..........(..(((((((((((.(((.((((((.((((((((((.....))).)))))))).)))))).))).))).)))))))).)..    reads      mm         sample
    .uagucaguaguuggggggugg.....................................................................    1          0          hsa
    ..agucaguaguuggggggugg.....................................................................    2          0          hsa
    ................gugggaacggcuucauaUaggagu...................................................    1          1          hsa
    ................ugggaacggcuucauacaggagu....................................................    3          0          hsa
    ................ugggaacggcuucauacaggaguuU..................................................    2          1          hsa
    ..................Ugaacggcuucauacaggagu....................................................    1          1          hsa
    ..................gaacggcuucauacaggag......................................................    3          0          hsa
    ..................gaacggcuucauacaggagu.....................................................    36         0          hsa
    ..................gaacggcuucauacCggagu.....................................................    1          1          hsa
    ..................gaCcggcuucauacaggagu.....................................................    1          1          hsa
    ..................gaacggcuucauacaggagC.....................................................    1          1          hsa
    ..................gaacggcuucauacagUagu.....................................................    1          1          hsa
    ..................gaacUgcuucauacaggagu.....................................................    1          1          hsa
    ..................gaacggcuucauacaggaUu.....................................................    3          1          hsa
    ..................gaacAgcuucauacaggagu.....................................................    1          1          hsa
    ..................gaacggcuucauacaggaUuu....................................................    4          1          hsa
    ..................Aaacggcuucauacaggaguu....................................................    1          1          hsa
    ..................gGacggcuucauacaggaguu....................................................    1          1          hsa
    ..................gaacUgcuucauacaggaguu....................................................    1          1          hsa
    ..................gaacggcuucauacaggaguu....................................................    37         0          hsa
    ..................gaacggcuucauacaggaguuC...................................................    1          1          hsa
    ..................gaacggcuucauacaggaguuA...................................................    13         1          hsa
    ..................gaacggcuucauacaggaguug...................................................    2          0          hsa
    ..................gaacggAuucauacaggaguug...................................................    1          1          hsa
    ..................gaacggcuucauacaggaguuU...................................................    16         1          hsa
    ..................gaacggcuucauacaggaguuAa..................................................    5          1          hsa
    ..................gaacggcuucauacaggaguugU..................................................    2          1          hsa
    ..................gaacggcuucauacaggaguuUa..................................................    1          1          hsa
    ....................aacggcuucauacaggaguu....................................................    1          0          hsa
    ....................aacggcuucauacaggaguug...................................................    1          0          hsa
    ....................aacggcuucauacaggaguuU...................................................    1          1          hsa
    ....................aacggcuucauacaggaguuga..................................................    7          0          hsa
    ..........................................ugaugcacaguuauccagc.....................................    1          0          hsa
    ..........................................................agcuccuauaugaugccuu.....................    1          0          hsa
```

hsa-miR-337-5p

hsa-miR-337-3p

guagucaguaguuggggggugggaacggcuucauacaggaguugaugcacaguuauccagcuccuauaugaugccuuucuucaucccccuucaa

.....gcuccuauaugaugccuuucuA.....	1	1	hsa
.....gcuccuauaugaugccuuucuu.....	1	0	hsa
.....gcuccuauaugaugccuuucuuc.....	1	0	hsa
.....cuccuauaugaugccuuucu.....	1	0	hsa
.....cuccuauaugaugccuuucuu.....	3	0	hsa
.....cuccuauaugaugccuuucuuc.....	17	0	hsa
.....cuccuGuaugaugccuuucuuc.....	1	1	hsa
.....uccuauaugaugccuuuc.....	1	0	hsa
.....uccuauaugaugccuuucuu.....	4	0	hsa
.....uccuauaugaugccuuCcuu.....	1	1	hsa
.....uccuauaugCugccuuucuuc.....	1	1	hsa
.....uccuauaugaugccuuucuuc.....	15	0	hsa
.....Accuauaugaugccuuucuuc.....	1	1	hsa
.....uccuauaugaugccuuucuucU.....	1	1	hsa
.....uccuauaugaugccuuucuuca.....	1	0	hsa