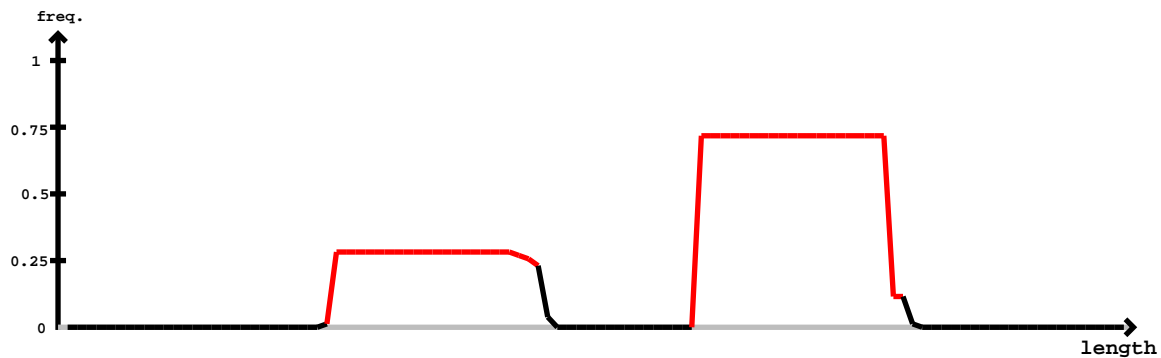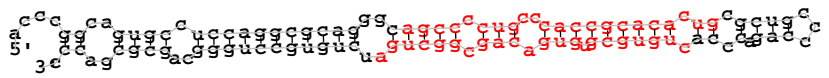miRBase precursor : hsa-mir-210
Total read count : 78
hsa-miR-210-5p read count : 22
hsa-miR-210-3p read count : 56
remaining reads : 0

hsa-miR-210-5p

hsa-miR-210-3p

```
5'- acccggcagugccuccaggcgcagggcagcccugcccaccgcacacugcgcugccccagacccacugugcgugugacagcggcugaucugugccugggcagcgcgaccc  -3'    exp
     ....((..(((((.(((((((((((..(((((.(((..(((((((((((.((.((((...))).).)).)))))).)))).))).)))))..)))))))))))..)))))..)).   reads      mm        sample
     .........................cagcccugcccaccgcacacu........................................................................    1         0          seq
     .........................agcccugcccaccgcaca..........................................................................    1         0          seq
     .........................Cgcccugcccaccgcacac.........................................................................    1         1          seq
     .........................agcccugcccaccgcacacu........................................................................    1         0          seq
     .........................agcccugcccaccgcacacug.......................................................................   14         0          seq
     .........................Cgcccugcccaccgcacacug.......................................................................    1         1          seq
     .........................agcccugcccaccgcacacugA......................................................................    2         1          seq
     .........................agcccugcccaccgcacacugU......................................................................    1         1          seq
     ...................................................cugugcgugugacagcggcu...............................................   44         0          seq
     ...................................................cugGgcgugugacagcggcu...............................................    1         1          seq
     ...................................................cugugcgugugacagcggcC...............................................    1         1          seq
     ...................................................cugugcgugugacagcggcA...............................................    1         1          seq
     ...................................................cugugcgugugacagcggcuga.............................................    8         0          seq
     ...................................................cugugcgugugacagcggcugau............................................    1         0          seq
```