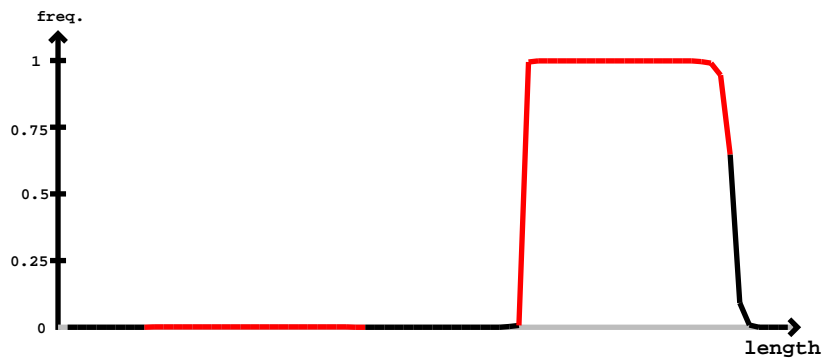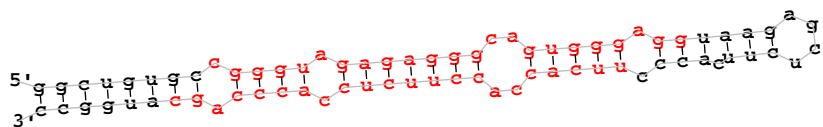miRBase precursor   : hsa-mir-197
Total read count   : 711
hsa-miR-197-5p read count : 1
hsa-miR-197-3p read count : 710
remaining reads   : 0

hsa-miR-197-3p

hsa-miR-197-5p

```
5'- ggcugugccggguagagagggcagugggagguaagagcucuucaccuuucaccaccuucuccacccagcauggcc   -3'    exp
    ((((((((.(((((.((((((((..(((((.((((((((....))).)))).)))))..))))))))).))))).))))))))    reads     mm       sample
    .......cggguagagagggcagugggga................................................     1        0        seq
    ..........................................ccuucaccaccuucuccacccag.......     1        0        seq
    ..........................................ccuucaccaccuucuccacccagA......     1        1        seq
    .........................................cuucaccaccuucuccacccag.......     1        0        seq
    .........................................Auucaccaccuucuccacccagc......     1        1        seq
    .........................................cuucaccaccuucuccacccagA......     1        1        seq
    ..........................................uucaccaccuucuccacc..........     2        0        seq
    ..........................................uucaccaccuucuccaccc.........     4        0        seq
    ..........................................uucaccaccuucuccaccca........     29       0        seq
    ..........................................uucCccaccuucuccaccca........     2        1        seq
    ..........................................uucaccaccuucuccacccGg.......     1        1        seq
    ..........................................uuAaccaccuucuccacccag.......     1        1        seq
    ..........................................uucaccaccuucuccacccCg.......     1        1        seq
    ..........................................uucCccaccuucuccacccag.......     1        1        seq
    ..........................................uucaccaccuucuccacccag.......     205      0        seq
    ..........................................uucaccaccuucuccacccagU......     32       1        seq
    ..........................................uucaccaccuucuccacccagA......     120      1        seq
    ..........................................uucaccaccuucuccacccagc......     232      0        seq
    ..........................................uucaccaccuucuccacccagG......     1        1        seq
    ..........................................Cucaccaccuucuccacccagc......     1        1        seq
    ..........................................uucaccaccuucuccaccAagc......     1        1        seq
    ..........................................uucCccaccuucuccacccagc......     3        1        seq
    ..........................................uucacUaccuucuccacccagc......     1        1        seq
    ..........................................uucaccaccuCuccacccagc......     1        1        seq
    ..........................................uucaccaccuucuccacccagcU.....     11       1        seq
    ..........................................uucCccaccuucuccacccagca.....     1        1        seq
    ..........................................uucaccaccuucuccacccagUa.....     2        1        seq
    ..........................................uucaccaccuucuccacccagca.....     36       0        seq
    ..........................................uucaccaccuucuccacccagGa.....     1        1        seq
    ..........................................uucaccaccuucuccacccagAa.....     7        1        seq
    ..........................................uucaccaccuucuccacccagcG.....     1        1        seq
    ..........................................uucaccaccuucuccacccagcUu....     2        1        seq
    ..........................................uucaccaccuucuccacccagcaG....     1        1        seq
    ..........................................uucaccaccuucuccacccagcaA....     2        1        seq
```

hsa-miR-197-3p

hsa-miR-197-5p

ggcugugccggguagagagggcagugggagguaagagcucuucacccuucaccaccuucuccacccagcauggcc

....ucaccaccuucuccaccca........   1   0   seq
....ucaccaccuucuccacccag.......   1   0   seq
....ucaccaccuucuccacccagcau....   1   0   seq