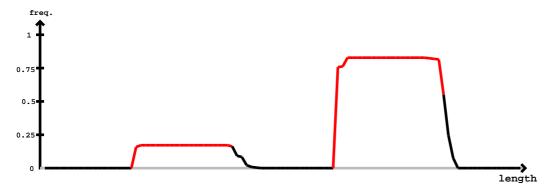miRBase precursor   : hsa-mir-342
Total read count   : 227
hsa-miR-342-5p read count : 39
hsa-miR-342-3p read count : 188
remaining reads   : 0


```
                                                                hsa-miR-342-3p

                                               hsa-miR-342-5p

5'- gaaacugggcucaaggugaggggugcuaucugugauugagggacaugguuaauggaauugucucacacagaaaucgcacccgucaccuuggccuacuua   -3'    exp
     .....(((((.(((((((((.((((((..(((((((..(((((.(((((.....))))....).)))))))))))....))))))).)))))))))))....    reads      mm          sample
.................aggggugcuaucugugauug........................................................................   2          0            seq
.................Cggggugcuaucugugauuga.......................................................................   2          1            seq
.................aggggugcuaucugugauAga.......................................................................   1          1            seq
.................aggggugcuaucugugauuga.......................................................................   13         0            seq
.................aggggugcuaucugugauugag......................................................................   2          0            seq
.................aggggugcuaucugugauugagg.....................................................................   5          0            seq
.................aggggugcuaucugugauugagA.....................................................................   7          1            seq
.................aggggugcuaucugugauugaggA....................................................................   2          1            seq
.................aggggugcuaucugugauugagggga..................................................................   1          0            seq
.................aggggugcuaucugugauugaggggaU.................................................................   1          1            seq
..................ggggugcuaucugugauugagg.....................................................................   2          0            seq
..................ggggugcuaucugugauugaggA....................................................................   1          1            seq
....................................................ucucacacagaaaucgcac.....................................   1          0            seq
....................................................ucucCcacagaaaucgcacc....................................   1          1            seq
....................................................ucucacacagaaaucgcaccc...................................   1          0            seq
....................................................ucucCcacagaaaucgcacccg..................................   5          1            seq
....................................................ucucacacagaaaucgcacccg..................................   48         0            seq
....................................................ucucacacagaaaucgcacUcg..................................   2          1            seq
....................................................ucucacacagaaaucgcaccUg..................................   2          1            seq
....................................................ucucacacagaaacgUacccg...................................   1          1            seq
....................................................ucucCcacagaaaucgcacccgu.................................   2          1            seq
....................................................ucucacacagaaaucgcacccAu.................................   1          1            seq
....................................................ucucacacagaaaucgcaccUgu.................................   1          1            seq
....................................................ucucacacagaaaucgcacccgu.................................   60         0            seq
....................................................ucucacacagaaaucgcacccgA.................................   3          1            seq
....................................................ucucacacagaaaucgcacccgC.................................   1          1            seq
....................................................uAucacacagaaaucgcacccguc................................   1          1            seq
....................................................ucucacacagaaaucgcacccguU................................   3          1            seq
....................................................ucucacacagaaaucgcacccguc................................   30         0            seq
....................................................ucucCcacagaaaucgcacccguca...............................   1          1            seq
....................................................ucucacacagaaaucgcacccguca...............................   4          0            seq
....................................................ucucacacagaaaucgcacccgucG...............................   1          1            seq
....................................................ucucacacagaaaucgcacccgucU...............................   3          1            seq
.....................................................cucacacagaaaucgcacccguc................................   1          0            seq
```

gaaacugggcucaaggugaggggugcuaucugugauugagggacaugguuaauggaauugucucacacagaaaucgcacccgucaccuuggccuacuua

```
..........................................................ucacacagaaaucgcacccg................    2      0      seq
..........................................................ucacacagaaaucgcacccgu...............    1      0      seq
..........................................................ucacacagaaaucgcacccguc..............    3      0      seq
..........................................................ucacacagaaaucgcacccguA..............    1      1      seq
..........................................................ucacacagaaaucgcacccguca.............    8      0      seq
```