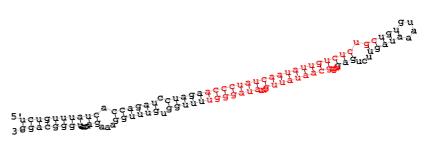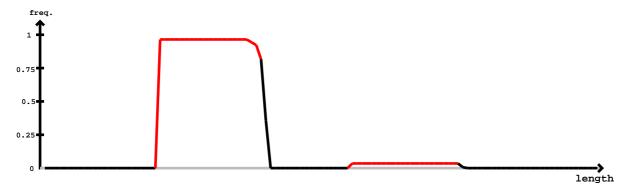miRBase precursor   : hsa-mir-454
Total read count   : 142
hsa-miR-454-5p read count : 137
hsa-miR-454-3p read count : 5
remaining reads   : 0

hsa-miR-454-5p

hsa-miR-454-3p

```
5'- ucuguuuaucaccagauccagaaacccuaucaauauugucucugcuguguaaauaguucgaguagugcaauauugcuuauagggguuuugguguuuggaaagaacaaugggcagg  -3'    exp
     .(((((((((.(((((((.((((((((((((((((((((((((.(((((....))))))...))....)))))))))))...))))))))))))).)))))...)))....))))))).   reads     mm      sample
     ....................acccuaucaauauugucuc.................................................................................     3        0       seq
     ....................acccuaucaauauugucucu................................................................................     3        0       seq
     ....................acccuaucaaCauugucucug...............................................................................     1        1       seq
     ....................Ccccuaucaauauugucucug...............................................................................     1        1       seq
     ....................acccuaucaauauugucucug...............................................................................    13        0       seq
     ....................acccuaucaauauugucucugU..............................................................................    38        1       seq
     ....................acccuaucaauauugucucugc..............................................................................     7        0       seq
     ....................acccuaucaauauugucucugA..............................................................................    19        1       seq
     ....................Ccccuaucaauauugucucugc..............................................................................     1        1       seq
     ....................Ccccuaucaauauugucucugcu.............................................................................     1        1       seq
     ....................acccuaucaauauugucucugUu.............................................................................     1        1       seq
     ....................Ucccuaucaauauugucucugcu.............................................................................     1        1       seq
     ....................acccuaucaauauugucucugcu.............................................................................    44        0       seq
     ....................acccuaucaauauugucucugcA.............................................................................     4        1       seq
     ...............................................................uagugcaauauugcuuauaggggu..............................     4        0       seq
     ...............................................................uagugcaauauugcuuauaggguu..............................     1        0       seq
```