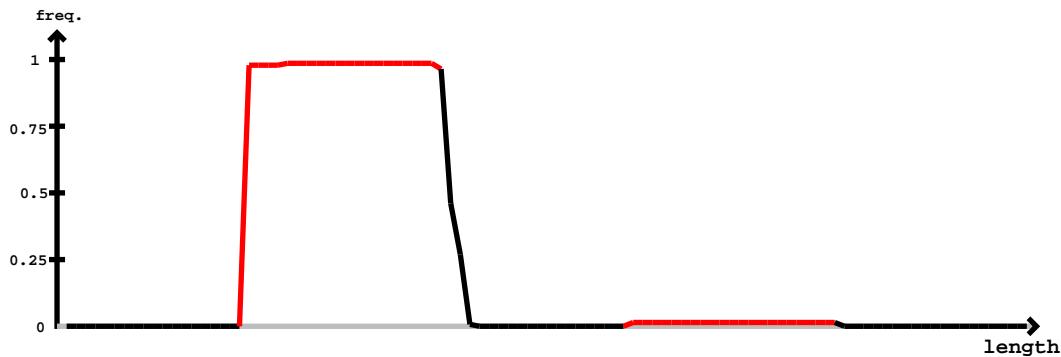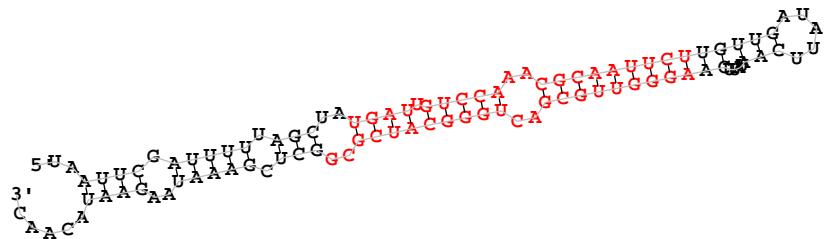miRBase precursor : dme-mir-219_MI0000358
Total read count : 141
dme-miR-219-5p_MIMAT0000335 read count : 139
dme-miR-219-3p_MIMAT0020803 read count : 2
remaining reads : 0

freq.

length

```
5'- uaauucgauuuuuagcuaugauuguccaaacgcaauucuuguugauauucaauauucaaggguugcgacugggcaucgcggcucgaaauaagaauacaac  -3'   exp
     ..(((((.(((((.(((..(((((.(((((..(((((((((((((((.....)))....)))))))))))))..)))))))))..))).))))))..)))))....   reads   mm        sample
     .................ugauuguccaaacgcaauuc..........................................................................   3      0         seq
     .................Agauuguccaaacgcaauucu.........................................................................   2      1         seq
     .................ugauuguccaaacgcaauuUu.........................................................................   1      1         seq
     .................ugauuguccaaacgcaauCcu.........................................................................   1      1         seq
     .................Cgauuguccaaacgcaauucu.........................................................................   1      1         seq
     .................ugauuguccaaacgcaaCucu.........................................................................   2      1         seq
     .................ugauuguccaaacAcaauucu.........................................................................   1      1         seq
     .................ugauuguccaaacgcaauucu.........................................................................   56     0         seq
     .................ugaAuguccaaacgcaauucu.........................................................................   1      1         seq
     .................ugauuguccaaacCcaauucu.........................................................................   4      1         seq
     .................ugauuguccaaacgcaUuucu.........................................................................   1      1         seq
     .................ugauuguccaaacgcCauucu.........................................................................   1      1         seq
     .................ugauuguccaaacgcaauucuu........................................................................   27     0         seq
     .................ugauugCccaaacgcaauucuug.......................................................................   1      1         seq
     .................ugauuguccaaacgcaauucuug.......................................................................   33     0         seq
     .................ugauuguccaaacgcaaCucuug.......................................................................   1      1         seq
     .................ugauuguccaGacgcaauucuug.......................................................................   1      1         seq
     .................ugauuguccaaacgcaauucuugu......................................................................   1      0         seq
     ....................uguccaaacgcaauuUuug........................................................................   1      1         seq
     .....................................................aggguugcgacugggcaucgcg....................................   2      0         seq
```