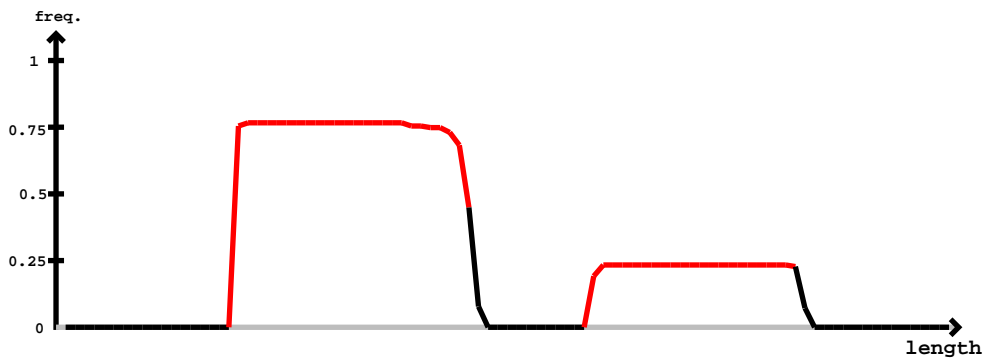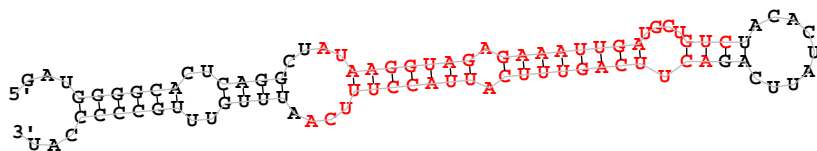miRBase precursor   : dme-mir-962_MI0005817
Total read count   : 167
dme-miR-962-3p_MIMAT0020857 read count
dme-miR-962-5p_MIMAT0005476 read count
remaining reads   : 0


```
                      dme-miR-962-5p_MIMAT0005476

                                    dme-miR-962-3p_MIMAT0020857

5'- gaugggcacucaggcuauaagguagagaaauugaugcugucuacacuauucagacuucaguuucauuaccuucaauuuguuugcccccau  -3'   exp
    ...(((((((..(((((....(((((((((.(((((((((....(((.........))))-.))))))))).)))))))...))))..)))))))...  reads   mm    sample
    ................auaagguagagaaauuga.............................................................    2      0     seq
    ................auaagguagagaaauugaug...........................................................    1      0     seq
    ................auaagguagagaaauugaugcu.........................................................    3      0     seq
    ................auaagguagagaaauugaugcug........................................................    8      0     seq
    ................auaagguagagaaauugaugcugu.......................................................   35      0     seq
    ................auaaggGagagaaauugaugcugu.......................................................    1      1     seq
    ................auaagguagagaaauugaCgcugu.......................................................    1      1     seq
    ................auaagguagagaaauugaugcGgu.......................................................    1      1     seq
    ................auGagguagagaaauugaugcugu.......................................................    1      1     seq
    ................auaagguagagaaauugaugcuguA......................................................    1      1     seq
    ................auaagguagagaaauugaGgcuguc......................................................    1      1     seq
    ................auaagguagagaaauugaCgcuguc......................................................    1      1     seq
    ................auaagguagagaaauuUaugcuguc......................................................    1      1     seq
    ................auaagguagagaaauugauCcuguc......................................................    1      1     seq
    ................auaagguagagaaauugaugcuguU......................................................    1      1     seq
    ................auaagguagagaaauugaugcuguG......................................................    1      1     seq
    ................aGaagguagagaaauugaugcuguc......................................................    1      1     seq
    ................auaUgguagagaaauugaugcuguc......................................................    1      1     seq
    ................auaagguagagaaauuAaugcuguc......................................................    1      1     seq
    ................auaagguagagaaauugaugcuguc......................................................   52      0     seq
    ................auaagguagagaaauugaugcugucu.....................................................   11      0     seq
    .................uaagguagagaaauugaugcugucu.....................................................    2      0     seq
    ...................................................acuucaguuucauuaccuuuc.......................    1      0     seq
    ...................................................acuucaguuucauuaccCuuca......................    1      1     seq
    ...................................................acuucaguuucauuaccuuuca......................   24      0     seq
    ...................................................Ucuucaguuucauuaccuuuca......................    1      1     seq
    ...................................................acuucaguuucauuaccuuucaa.....................    5      0     seq
    ....................................................cuucaguuucauCaccuuucaa.....................    1      1     seq
    ....................................................cuucaguuucauuaccuuucaa.....................    6      0     seq
```