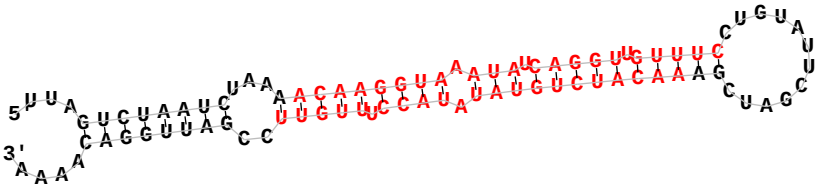
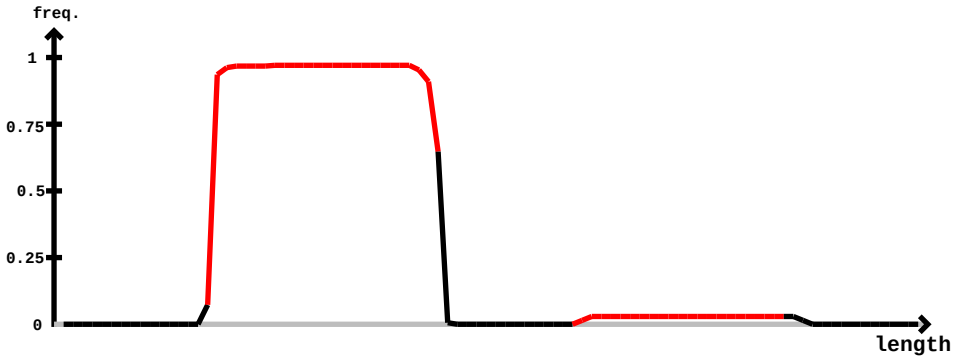
miRBase precursor : dme-mir-963_MI0005818
Total read count : 343
dme-miR-963-3p_MIMAT0020858 read count
dme-miR-963-5p_MIMAT0005477 read count
remaining reads : 0





dme-miR-963-5p_MIMAT0005477

dme-miR-963-3p_MIMAT0020858

```
5'- uuagucuaaucuaaaacaagguaaauaucagguuguuucuguauucgaucgaaacaucuguauauaccuuuguuccgauuggacaaaa  -3'    exp
   ...(((((((...(((((((((.(((.(((((.(((((.............)))))))))))))).)))).)))))..)))))))....    reads      mm         sample
   ..............aacaagguaaauaucagguuguu.....................................................    4          0          seq
   ..............aacaagguaaauaucaAguuguuu....................................................    1          1          seq
   ..............aacaagguaaauaucagguuguuu....................................................    17         0          seq
   ..............aacaagguaaauaucagguuguuuc...................................................    3          0          seq
   ..............acaagguaaauaucagUuugu.......................................................    1          1          seq
   ..............acaagguaaauaucagguugu.......................................................    5          0          seq
   ..............acaagguaaauaucagguugGu......................................................    1          1          seq
   ..............acaagguaaGuaucagguuguu.....................................................    1          1          seq
   ..............acaagguaaauaucagguuguu.....................................................    9          0          seq
   ..............acaagguaaauaucagguuguuu....................................................    65         0          seq
   ..............acaagguaaauaucaggGuguuu....................................................    1          1          seq
   ..............acaagAuaaauaucagguuguuu....................................................    1          1          seq
   ..............Gcaagguaaauaucagguuguuu....................................................    1          1          seq
   ..............acaagguaaauaucaAguuguuu....................................................    1          1          seq
   ..............acaagguaaauaucCguuguuuc....................................................    1          1          seq
   ..............Gcaagguaaauaucagguuguuuc...................................................    1          1          seq
   ..............acaagguaaauaucagguCguuuc...................................................    1          1          seq
   ..............acaagguaaauaAcagguuguuuc...................................................    1          1          seq
   ..............acaagguaaauaucaggAuguuuc...................................................    2          1          seq
   ..............acaagguaaauaucagguuguuuU...................................................    2          1          seq
   ..............acaagguaaaAaucagguuguuuc...................................................    1          1          seq
   ..............acaagguaaauaucaggCuguuuc...................................................    1          1          seq
   ..............acaagguaaauaucagUuuguuuc...................................................    2          1          seq
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   ...............caagguaaauaucagguuguuu....................................................    3          0          seq
   ...............caagguaaauaucagguuguuuc...................................................    4          0          seq
   ...............caagUuaaauaucagguuguuuc...................................................    1          1          seq
   ...............caagguaaauaucagguuguuucA..................................................    1          1          seq
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   ....................uaaauaucagguuguuuc.....................................................    1          0          seq
   ..........................................................aacaucuguauauaccuuuguuU.............    5          1          seq
   ..........................................................acaucuguauauaccuuuguucU.............    4          1          seq
```

dme-miR-963-5p_MIMAT0005477

dme-miR-963-3p_MIMAT0020858

uuagucuaaucuaaaacaagguaaauaucagguuguuuccuguauucgaucgaaacaucuguauauaccuuuguuccgauuggacaaaa

....acaucuguauauaccuuuguucc..... 1 0 seq