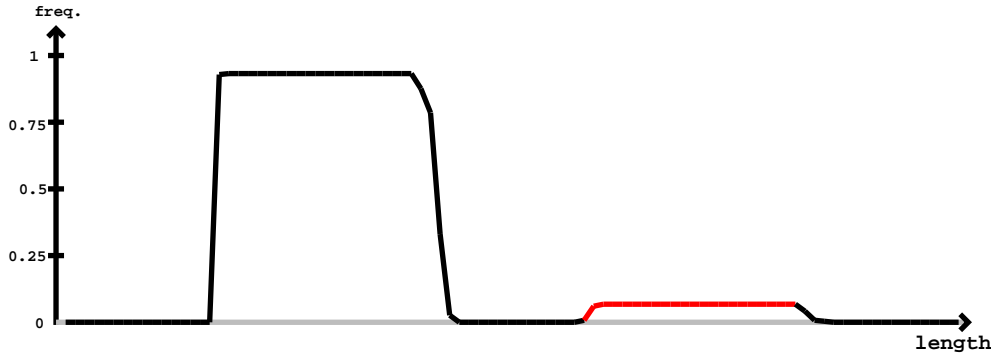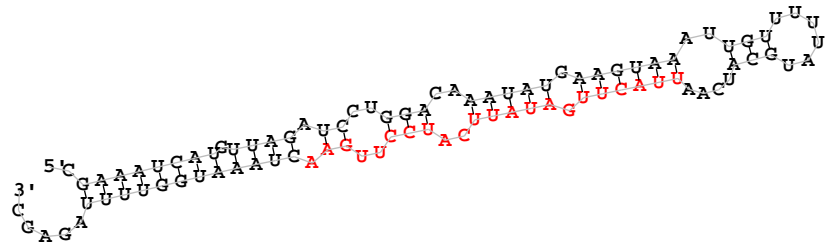miRBase precursor   : dme-mir-982_MI0005839
Total read count   : 266
dme-miR-982-3p_MIMAT0020879 read count   : 18
remaining reads   : 248

<div align="center">dme-miR-982-3p_MIMAT0020879</div>

```
5'- cgaaaucauguuagauccuggacaaauaugaaguaaauuguuuuuaugcaucaauuacuugauauucauccuugaacuaaaugguuuuagagc  -3'    exp
    .((((((((.((((.((..(((..(((((.((((((.((.(((......)))....))))))).)))))..))).))).))))))))))....    reads    mm    sample
    ..............uccuggacaaauaugaaguaa....................................................          14       0     seq
    ..............uccuggacaaauaCgaaguaa....................................................          1        1     seq
    ..............uccuCgacaaauaugaaguaaa...................................................          2        1     seq
    ..............uccuggacaaauaugaaguaaa...................................................          22       0     seq
    ..............uccuggacaaauaugaaguCaau..................................................          1        1     seq
    ..............uccuggacaaauaugaGguaaau..................................................          1        1     seq
    ..............uccuUgacaaauaugaaguaaau..................................................          1        1     seq
    ..............uccuggacaaauauAaaguaaau..................................................          1        1     seq
    ..............uccuggacaaauaugaagGaaau..................................................          1        1     seq
    ..............uccuggacaaauaugaaguaaau.................................................           113      0     seq
    ..............uccuggacaaauaugaaguaaaA.................................................           1        1     seq
    ..............uccuggacaGaugaaguaaau...................................................           1        1     seq
    ..............uccuggacaaauaugaaguaaauu................................................           74       0     seq
    ..............Cccuggacaaauaugaaguaaauu................................................           1        1     seq
    ..............uccuggacaaauaugaaguaaauA................................................           1        1     seq
    ..............uccuggacaaauaugaGguaaauu................................................           1        1     seq
    ..............Accuggacaaauaugaaguaaauu................................................           1        1     seq
    ..............uccuggacGaaauaugaaguaaauu...............................................           2        1     seq
    ..............uccuggaGaaauaugaaguaaauu................................................           1        1     seq
    ..............uccuggacaaauaugaaguaaauuA...............................................           7        1     seq
    ...............ccuggacaaauaugaaguaaau.................................................           1        0     seq
    ...........................................................auuacuugauauucauccuugaa...............   1    0     seq
    ...........................................................auuacuugauauucauccuugaac..............   1    0     seq
    ............................................................uuacuugauauucauccuugaa...............   6    0     seq
    ............................................................uuacuugauauucauccuugaac..............   6    0     seq
    ............................................................uuacuugauauucauccuugaacu.............   1    0     seq
    ............................................................uuacuugauauucauccuugaacAa............   1    1     seq
    .............................................................uacuugauauucauccuugaac..............   2    0     seq
```