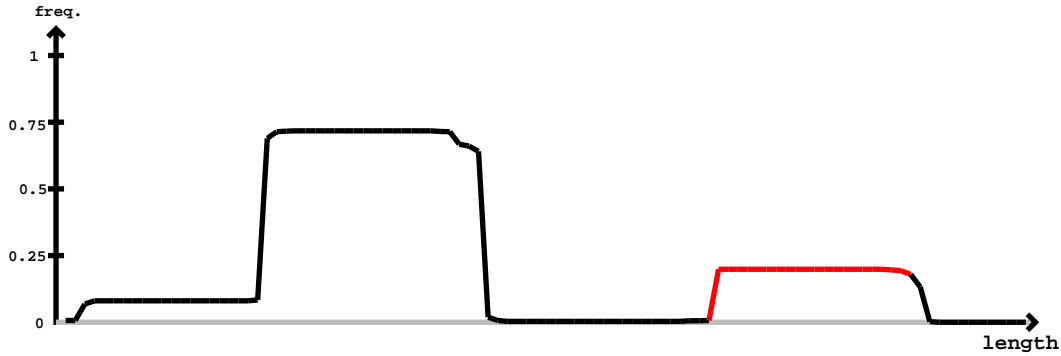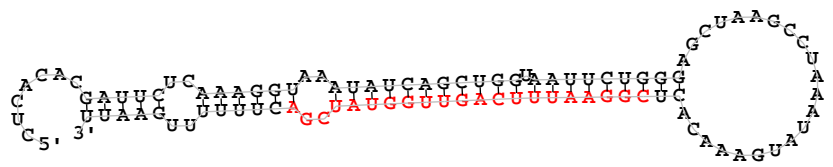miRBase precursor : dme-mir-283_MI0000368
Total read count : 896
dme-miR-283-3p_MIMAT0020810 read count
remaining reads : 720

dme-miR-283-3p_MIMAT0020810

```
5'- cucacacgauucucaaagguaaauaucagcugguaauucugggagcuaagccuaaauaugaaacacucggaauuucaguuggguaucgacuuuuuugaauu -3'    exp
     ......(((((..(((((((..((((((((((((.(((((((((....................)))))))))))))))))))))))))..))))))..)))))    reads   mm    sample
     cucacacgauucucaaaggu.................................................................................    5       0     seq
     Aucacacgauucucaaaggu.................................................................................    1       1     seq
     .ucacacgauucucaaaggG.................................................................................    1       1     seq
     .ucacacgauucucCaaggu.................................................................................    1       1     seq
     .Acacacgauucucaaaggu.................................................................................    4       1     seq
     .ucGcacgauucucaaaggu.................................................................................    2       1     seq
     .uUacacgauucucaaaggu.................................................................................    1       1     seq
     .ucacacgauucucaaaggu.................................................................................    46      0     seq
     ..cacacgauucucaaaggu.................................................................................    10      0     seq
     ..Aacacgauucucaaaggu.................................................................................    1       1     seq
     ....................uaaauaucagcugguaauu..............................................................    1       0     seq
     ....................uaaauaucagcugguaauucu............................................................    1       0     seq
     ....................Aaaauaucagcugguaauucuggg.........................................................    1       1     seq
     ....................aaauaucagcugguaauu...............................................................    1       0     seq
     ....................aaauaucagcugguaauuc..............................................................    1       0     seq
     ....................aaaCaucagcugguaauucu.............................................................    1       1     seq
     ....................aaauaucagcugguaauucu.............................................................    31      0     seq
     ....................aaauaucagcuggCaauucu.............................................................    1       1     seq
     ....................aaauaucagcAgguaauucu.............................................................    1       1     seq
     ....................aaauaucaAacugguaauucu............................................................    1       1     seq
     ....................aaauaCcagcugguaauucu.............................................................    1       1     seq
     ....................aaauaucagcugguaauAcu.............................................................    1       1     seq
     ....................aaauaucagUugguaauucu.............................................................    1       1     seq
     ....................aaauaucagcuUguaauucu.............................................................    2       1     seq
     ....................aaauaucagcugguaauucug............................................................    4       0     seq
     ....................aaauaucagcugguaauucuU............................................................    1       1     seq
     ....................aaauaucagcugguaauucuA............................................................    1       1     seq
     ....................aaauaucagcuggGaauucugg...........................................................    1       1     seq
     ....................aaauaucagcugguaauucugg...........................................................    17      0     seq
     ....................Caauaucagcugguaauucuggg..........................................................    4       1     seq
     ....................aaauaucagcuggAaauucuggg..........................................................    1       1     seq
     ....................aaauaucagcugguaauucugAg..........................................................    2       1     seq
     ....................aaauaucagcugguaauucuggU..........................................................    3       1     seq
     ....................aaauaucagcugguaauucuAgg..........................................................    1       1     seq
     ....................aaauaucagcugguaauGcuggg..........................................................    1       1     seq
```

dme-miR-283-3p_MIMAT0020810

```
cucacacgauucucaaagguaaaauaucagcugguaauucugggagcuaagccuaaaauaugaaacacucggaauuucaguugguaucgacuuuuuugaauu
```

Sequence	Count	Mis	Type
....................aaauaucagcugguaauucAggg..	1	1	seq
....................aaauaucagcuggGaauucuggg...	5	1	seq
....................aaauaucagcugguCauucuggg...	1	1	seq
....................aaauaucagcuggCaauucuggg...	1	1	seq
....................aaauaucagAugguaauucuggg...	1	1	seq
....................aaauaucagcugguaauucuggA...	1	1	seq
....................aaaCaucagcugguaauucuggg...	2	1	seq
....................aaCuaucagcugguaauucuggg...	1	1	seq
....................aaauaucagcugguaauAcuggg...	1	1	seq
....................aaauaCcagcugguaauucuggg...	1	1	seq
....................aCauaucagcugguaauucuggg...	2	1	seq
....................aaauaucagcugguaauucuUgg...	2	1	seq
....................aaaGaucagcugguaauucuggg...	2	1	seq
....................aaauaucagcugguaauucugUg...	4	1	seq
....................aaauaucagcugguaauucuggg...	493	0	seq
....................aaauaucagcugguaauucCggg...	1	1	seq
....................aaauauAagcugguaauucuggg...	1	1	seq
....................aaauaucagcugguaCuucuggg...	3	1	seq
....................aaauaucagcuggCaauucuggga..	1	1	seq
....................aaauaucagcugguaauucuggga..	3	0	seq
....................aaauaucagcugguaauucugggU..	7	1	seq
....................aaauaucagcugguaauucugggaC...	1	1	seq
....................aaauaucagcugguaauucugggaA...	2	1	seq
.....................aauaucagcugguaauucu...	1	0	seq
.....................aauaucaAcugguaauucuggg..	1	1	seq
.....................aauaucagcugguaauucuggg..	20	0	seq
......................auaucagcugguaauucuggg..	2	0	seq
.......................uaucagcugguaauucuggg..	1	0	seq
....agcuaagccuaaaauaugaaacacu.......................	3	0	seq
....acucggaauuucaguugguauc....................	2	0	seq
....ucggaauuucaAuugguauc..................	1	1	seq
....cggaauuucaguugguau..................	2	0	seq
....cggaauuGcaguugguaucg................	1	1	seq
....cggaauuucaguugguaucg................	11	0	seq
....cggaauuucaguugguaucga...............	41	0	seq
....cggaauuucGguugguaucga...............	1	1	seq
....Aggaauuucaguugguaucgac.............	12	1	seq
....cggaauuucaguugguaucgac.............	91	0	seq
....cggaauuucaguugguaucgaA.............	8	1	seq
....cggaauuucaguuAguaucgac.............	1	1	seq
....cggaauuucaguugguaucgaU.............	1	1	seq
....cgAaauuucaguugguaucgac.............	1	1	seq
....cggaauGcaguugguaucgac.............	1	1	seq
....cggaauuucaguugguaucgUc.............	2	1	seq
....cggaauuucaguugguaucgacA............	2	1	seq