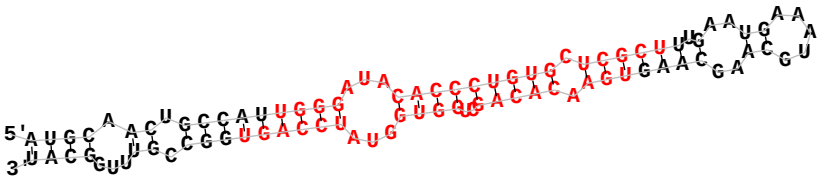
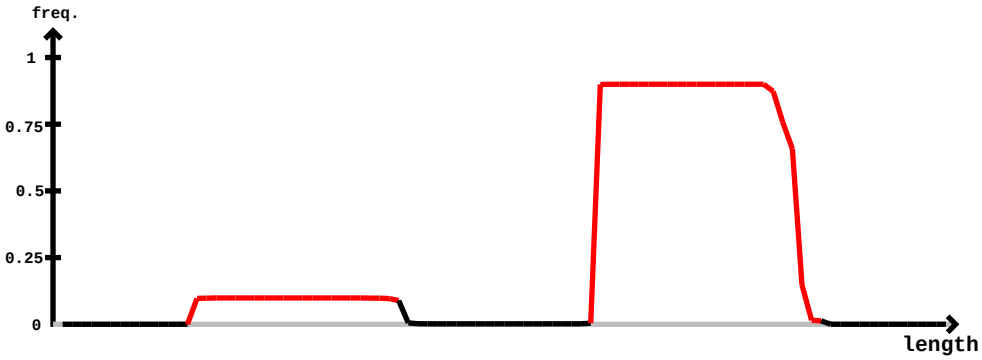
miRBase precursor : dme-mir-317_MI0000429
Total read count : 12108
dme-miR-317-5p_MIMAT0020843 read count
dme-miR-317-3p_MIMAT0000409 read count
remaining reads : 20





dme-miR-317-3p_MIMAT0000409

dme-miR-317-5p_MIMAT0020843

```
5'- augcaacugccauugggauacacccugugcucgcuuugaaugaaaugcaagcaagugaacacagcuggugguauccaguggccguuuggcau  -3'   exp
    ((((.((.((((((((((...((((((((((.((((((.(..((.....)).)).)))))).)))))..))))...)))))))))).))...))))   reads     mm        sample
    ............uugggauacacccugugcucgc.....................................................................   1         0         seq
    ............ugggauacacccugugcu.........................................................................   4         0         seq
    ............ugggauacacccugAgcu.........................................................................   1         1         seq
    ............ugggauacacccuUugcu.........................................................................   1         1         seq
    ............ugggauacacccAgugcu.........................................................................   1         1         seq
    ............ugggauacacccugugcuc........................................................................   2         0         seq
    ............ugggauacacccugugcucg.......................................................................   18        0         seq
    ............ugggauacacccugAgcucgc.....................................................................   1         1         seq
    ............ugggauacacccugugUucgc.....................................................................   1         1         seq
    ............ugggauacacccugugcucgc.....................................................................   77        0         seq
    ............ugggauacacccuguCcucgc.....................................................................   1         1         seq
    ............ugggauacacccugugcCcgc.....................................................................   1         1         seq
    ............ugggauacacccAgugcucgc.....................................................................   1         1         seq
    ............ugggauacacccuCugcucgc.....................................................................   1         1         seq
    ............ugggauacacccCgugcucgc.....................................................................   2         1         seq
    ............ugggauacacUcugugcucgcu....................................................................   1         1         seq
    ............ugggauacacccugugcucgUu....................................................................   2         1         seq
    ............ugggauacacccuAugcucgcu....................................................................   1         1         seq
    ............ugggauacacccuguAcucgcu....................................................................   17        1         seq
    ............ugggauacacccugugcucgcA....................................................................   3         1         seq
    ............ugggauacacccugAgcucgcu....................................................................   3         1         seq
    ............ugggauacacccugugcCcgcu....................................................................   3         1         seq
    ............ugggauacacccugugUucgcu....................................................................   10        1         seq
    ............Cgggauacacccugugcucgcu....................................................................   3         1         seq
    ............ugggauacacccuUugcucgcu....................................................................   2         1         seq
    ............ugggauaUacccugugcucgcu....................................................................   1         1         seq
    ............ugggauacacccugugcucAcu....................................................................   1         1         seq
    ............ugggauacacccugGgcucgcu....................................................................   1         1         seq
    ............ugggauacacccAgugcucgcu....................................................................   5         1         seq
    ............ugggauacacccugugcucUcu....................................................................   2         1         seq
    ............ugggauacacccuguCcucgcu....................................................................   5         1         seq
    ............ugggauGcacccugugcucgcu....................................................................   3         1         seq
    ............ugggauacacccugugcuUgcu....................................................................   3         1         seq
    ............ugggauacaccUugugcucgcu....................................................................   2         1         seq
```

dme-miR-317-3p_MIMAT0000409

dme-miR-317-5p_MIMAT0020843

augcaacugccauugggauacacccugugcucgcuuugaaugaaaugcaagcaagugaacacagcuggugguauccaguggccguuuggcau

.....ugggauacacccugugcuAgcu.....	2	1	seq
.....uAggauacacccugugcucgcu.....	3	1	seq
.....ugggauacacccugugcucgcu.....	900	0	seq
.....ugAgauacacccugugcucgcu.....	2	1	seq
.....ugggaGacacccugugcucgcu.....	2	1	seq
.....ugggauacacccuCugcucgcu.....	3	1	seq
.....uUggauacacccugugcucgcu.....	1	1	seq
.....ugggaAacacccugugcucgcu.....	6	1	seq
.....ugggauacacccugugcucgcG.....	2	1	seq
.....ugggauacacccugugcucgcC.....	6	1	seq
.....Agggauacacccugugcucgcu.....	2	1	seq
.....ugggauacGcccugugcucgcu.....	1	1	seq
.....ugggauacacccCgugcucgcu.....	9	1	seq
.....ugggauacacccugugcAcgcu.....	3	1	seq
.....ugggaCacacccugugcucgcu.....	2	1	seq
.....uggAauacacccugugcucgcu.....	2	1	seq
.....ugggauacacccGgugcucgcu.....	2	1	seq
.....ugggauacacccugCgcucgcu.....	4	1	seq
.....ugggauGcacccugugcucgcuu.....	3	1	seq
.....ugggauacacccuguAcucgcuu.....	1	1	seq
.....ugggauacaccUugugcucgcuu.....	1	1	seq
.....ugggauacacccugugcucgcuu.....	29	0	seq
.....ugggaAacacccugugcucgcuu.....	1	1	seq
.....ugggauacacccugugcucgcuA.....	2	1	seq
.....ugAgauacacccugugcucgcuu.....	1	1	seq
.....ugggauacacccugugcucgcuuu.....	3	0	seq
.....ugggauacacccuCugcucgcuuu.....	1	1	seq
.....ugggauacacccugugcucgcuuug.....	1	0	seq
.....gggauacacccugugcucgc.....	3	0	seq
.....gggauacacccugugcucgcu.....	5	0	seq
.....ggauacacccugugcucg.....	1	0	seq
.....ggauacacccugugcucgc.....	2	0	seq
.....ggauacacccugugcucgcu.....	7	0	seq
.....uugaaugaaaugcaagca.....	1	0	seq
.....uugaaugaaaugcaagcaag.....	17	0	seq
.....uugaaugaaaugcaagcaagA.....	1	1	seq
.....gcaagugaacacagcuggugguauc.....	1	0	seq
.....aAugaacacagcuggugguaucc.....	1	1	seq
.....gugaacacagcuggugguggua.....	1	0	seq
.....Augaacacagcuggugguggua.....	1	1	seq
.....Uugaacacagcuggugguggua.....	1	1	seq
.....gugaacacagcuggugguguau.....	7	0	seq
.....gugaacacagcuggugguggCau.....	1	1	seq
.....Augaacacagcuggugguguau.....	1	1	seq
.....gugaacacagcuggCgguau.....	1	1	seq
.....gugaacacagcuggugAuau.....	1	1	seq
.....Augaacacagcuggugguauc.....	1	1	seq
.....gugaacacagcugguAguauc.....	1	1	seq
.....gugaacacagcuggugguauc.....	7	0	seq
.....Cugaacacagcuggugguauc.....	2	1	seq
.....Augaacacagcuggugguaucc.....	6	1	seq
.....ugaacacagcugguggua.....	288	0	seq
.....ugaaUacagcugguggua.....	2	1	seq
.....ugaacacagcuggAggua.....	1	1	seq
.....ugaacacagcuggugguG.....	2	1	seq
.....uUaacacagcugguggua.....	1	1	seq
.....ugaacacagcugguAgua.....	1	1	seq
.....ugaacacagcCgguggua.....	1	1	seq
.....Cgaacacagcugguggua.....	1	1	seq
.....ugaacacagcugguUgua.....	1	1	seq
.....ugaacGcagcugguggua.....	1	1	seq
.....ugaacacagcuggugAua.....	2	1	seq
.....ugaacacagcuggugguU.....	9	1	seq
.....ugaacacagcugguggGa.....	2	1	seq
.....ugaacacagcuAguggua.....	1	1	seq
.....ugaacacaAcugguggua.....	1	1	seq
.....ugGacacagcuggugguau.....	1	1	seq
.....ugaacacagcuUgugguau.....	1	1	seq
.....ugaacacagcuggGgguau.....	3	1	seq

dme-miR-317-3p_MIMAT0000409

dme-miR-317-5p_MIMAT0020843

augcaacugccauugggauacacccugugcucgcuuugaaugaaaugcaagcaagugaacacagcuggugguauccaguggccguuuggcau

.....ugaacaUagcuggugguau.....	2	1	seq
.....ugaacacagcugAugguau.....	1	1	seq
.....ugaacacagcuggugguUu.....	7	1	seq
.....ugaacacagcuggugguaA.....	11	1	seq
.....ugaacacagcuggugCuau.....	1	1	seq
.....ugaacacaAcuggugguau.....	3	1	seq
.....ugaacacagcCggugguau.....	2	1	seq
.....ugaaUacagcuggugguau.....	1	1	seq
.....Cgaacacagcuggugguau.....	5	1	seq
.....ugaGcacagcuggugguau.....	9	1	seq
.....ugaacacagcuggugUuau.....	5	1	seq
.....ugaacaAagcuggugguau.....	2	1	seq
.....ugaacacGgcuggugguau.....	4	1	seq
.....ugaacacagcuAgugguau.....	5	1	seq
.....ugaacGcagcuggugguau.....	8	1	seq
.....ugaacacagcugguggCau.....	4	1	seq
.....ugaacacagcugCugguau.....	2	1	seq
.....ugaacacagcugguggAau.....	9	1	seq
.....ugaacacagcAggugguau.....	8	1	seq
.....ugaacUcagcuggugguau.....	2	1	seq
.....ugaacacagcuggAgguau.....	7	1	seq
.....ugaacacagcugguggGau.....	3	1	seq
.....ugaacacagcuggugguau.....	1260	0	seq
.....uAaacacagcuggugguau.....	10	1	seq
.....ugaacacagUuggugguau.....	1	1	seq
.....ugaacacagcuggugguGu.....	2	1	seq
.....ugaacacagcuggugguaC.....	2	1	seq
.....ugaacacagcuggCgguau.....	10	1	seq
.....uUaacacagcuggugguau.....	1	1	seq
.....ugaacacagcuggugAuau.....	3	1	seq
.....ugaacacagcuggugguaG.....	3	1	seq
.....ugaacacagcuAgugguauc.....	5	1	seq
.....Agaacacagcuggugguauc.....	1	1	seq
.....ugaacacagcuggugguGuc.....	4	1	seq
.....ugaUcacagcuggugguauc.....	1	1	seq
.....ugaacacagcuggugguauG.....	1	1	seq
.....ugaacacaAcuggugguauc.....	6	1	seq
.....ugaacacagcugguggGauc.....	4	1	seq
.....ugaacacagcuggugAuauc.....	11	1	seq
.....ugaCcacagcuggugguauc.....	1	1	seq
.....ugaacacagcugAugguauc.....	3	1	seq
.....ugaacacagcuggugguauU.....	35	1	seq
.....ugaacacagcuggCgguauc.....	6	1	seq
.....ugaacacagcuggugguauA.....	4	1	seq
.....ugaacacagcugguUguauc.....	1	1	seq
.....ugaacGcagcuggugguauc.....	3	1	seq
.....ugaacacagcuggugguCuc.....	1	1	seq
.....ugaacacaCcuggugguauc.....	1	1	seq
.....ugaacacagcuggugguaAc.....	5	1	seq
.....ugaacacagcGggugguauc.....	2	1	seq
.....ugGacacagcuggugguauc.....	4	1	seq
.....ugaacacagcuggugguaCc.....	2	1	seq
.....ugaacacagcuggugCuauc.....	3	1	seq
.....ugaacacagcuggugguaGc.....	1	1	seq
.....ugaaUacagcuggugguauc.....	3	1	seq
.....ugaacacaUcuggugguauc.....	1	1	seq
.....ugaacacagcugguggAauc.....	17	1	seq
.....Cgaacacagcuggugguauc.....	1	1	seq
.....ugaacacagcuggGgguauc.....	2	1	seq
.....uUaacacagcuggugguauc.....	1	1	seq
.....ugaacacagcuggAgguauc.....	6	1	seq
.....ugaacacagcugguggCauc.....	8	1	seq
.....ugaacacagcuggugguUuc.....	2	1	seq
.....ugaacacagcCggugguauc.....	2	1	seq
.....ugaacacagcuCgugguauc.....	1	1	seq
.....ugaacaUagcuggugguauc.....	1	1	seq
.....ugaacacagcAggugguauc.....	4	1	seq
.....ugaacacagcugguAguauc.....	3	1	seq
.....ugaacacagcuggugguauc.....	1035	0	seq

dme-miR-317-3p_MIMAT0000409

dme-miR-317-5p_MIMAT0020843

augcaacugccauugggauacacccugugcucgcuuugaaugaaaugcaagcaagugaacacagcuggugguauccaguggccguuuggcau

.....ugaacacagcugguCguauc.....	2	1	seq
.....uUaacacagcuggugguaucc.....	8	1	seq
.....ugaacacagcuggugguaCcc.....	4	1	seq
.....ugaacacGgcuggugguaucc.....	7	1	seq
.....ugaacacagcugAugguaucc.....	7	1	seq
.....ugaacacagcuggCgguaucc.....	39	1	seq
.....ugaacaAagcuggugguaucc.....	4	1	seq
.....ugaacacagcugguggAaucc.....	81	1	seq
.....ugaacacagcuCgugguaucc.....	6	1	seq
.....ugaacacagcuggugguUucc.....	11	1	seq
.....ugaacaUagcuggugguaucc.....	5	1	seq
.....ugaacacCgcuggugguaucc.....	2	1	seq
.....ugaacacagcAggugguaucc.....	33	1	seq
.....ugaacacagcuggugguaucc.....	5445	0	seq
.....ugaacacagcuggugguauGc.....	4	1	seq
.....ugaacacagcugguggCaucc.....	44	1	seq
.....ugaacacagcugCugguaucc.....	4	1	seq
.....ugaacacagcugguAguaucc.....	8	1	seq
.....ugaacacagcuggugguauUc.....	16	1	seq
.....ugaacacagcuggugguCucc.....	6	1	seq
.....ugaacacagcuggugguauAc.....	3	1	seq
.....ugaacacagcuggGgguaucc.....	13	1	seq
.....ugaacacagcuggugAuaucc.....	57	1	seq
.....ugaaUacagcuggugguaucc.....	4	1	seq
.....ugaacacagcuAgugguaucc.....	8	1	seq
.....ugaacacaCcuggugguaucc.....	5	1	seq
.....ugaGcacagcuggugguaucc.....	10	1	seq
.....uCaacacagcuggugguaucc.....	4	1	seq
.....ugaacacagcuggugguaucG.....	11	1	seq
.....Cgaacacagcuggugguaucc.....	2	1	seq
.....ugaacacaUcuggugguaucc.....	2	1	seq
.....ugaacacagcuUguggguaucc.....	3	1	seq
.....ugaacacagcuggugguaAcc.....	29	1	seq
.....ugaacacagcugUugguaucc.....	3	1	seq
.....ugaaAacagcuggugguaucc.....	3	1	seq
.....ugaacacagcGggugguaucc.....	2	1	seq
.....ugaacacagUuggugguaucc.....	3	1	seq
.....ugaacacUgcuggugguaucc.....	2	1	seq
.....ugaacacagcuggugguaucU.....	49	1	seq
.....ugaacUcagcuggugguaucc.....	4	1	seq
.....ugaacacagcuggAgguaucc.....	40	1	seq
.....ugaacGcagcuggugguaucc.....	24	1	seq
.....ugaacacagcuggugUuaucc.....	4	1	seq
.....ugaacacagcugguUguaucc.....	2	1	seq
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.....ugaacacagcugguggGaucc.....	13	1	seq
.....ugaacacagcugguCguaucc.....	5	1	seq
.....uAaacacagcuggugguaucc.....	14	1	seq
.....ugaacacagcuggugguaGcc.....	2	1	seq
.....Agaacacagcuggugguaucc.....	2	1	seq
.....ugGacacagcuggugguaucc.....	14	1	seq
.....ugUacacagcuggugguaucc.....	1	1	seq
.....ugaacacagcuggugguGucc.....	8	1	seq
.....ugaacaGagcuggugguaucc.....	1	1	seq
.....ugaacacagcuggugCuaucc.....	14	1	seq
.....ugaacacaAcuggugguaucc.....	12	1	seq
.....ugaacacagcuggugguaucA.....	36	1	seq
.....ugaacacagcuggugUuaucca.....	2	1	seq
.....ugaacacagcuggugAuaucca.....	8	1	seq
.....ugaacacagcuggugguaucGa.....	2	1	seq
.....ugaacacagcuggugguaAcca.....	4	1	seq
.....ugaacacagcuggugguaucUa.....	3	1	seq
.....ugaacacagcugguAguaucca.....	3	1	seq
.....ugaacacagcuggugguaCcca.....	1	1	seq
.....ugaacaAagcuggugguaucca.....	1	1	seq
.....ugaacacagcuCguggguaucca.....	1	1	seq
.....ugaacacagcuggugguGucca.....	1	1	seq
.....ugaacacaCcuggugguaucca.....	1	1	seq
.....ugaacacagcuggugguaucca.....	1426	0	seq

dme-miR-317-3p_MIMAT0000409

dme-miR-317-5p_MIMAT0020843

augcaacugccauugggauacacccugugcucgcuuugaaugaaaugcaagcaagugaacacagcuggugguauccaguggccguuuggcau

.....ugaacacagcuggugguaGcca.....	1	1	seq
.....ugaacacagcuggugguAaucca.....	15	1	seq
.....ugaacacGgcuggugguaucca.....	1	1	seq
.....ugaacaUagcuggugguaucca.....	2	1	seq
.....ugaacacagcuggugguCaucca.....	6	1	seq
.....ugaacacagcuggugguauccG.....	4	1	seq
.....uCaacacagcuggugguaucca.....	1	1	seq
.....ugaacacagcuggAgguaucca.....	4	1	seq
.....ugaacacagcAggugguaucca.....	4	1	seq
.....ugCacacagcuggugguaucca.....	1	1	seq
.....ugaacacagcuggugguGaucca.....	4	1	seq
.....ugaacUcagcuggugguaucca.....	1	1	seq
.....ugaacacagcuUgugguaucca.....	3	1	seq
.....ugaacacagcugCugguaucca.....	1	1	seq
.....uUaacacagcuggugguaucca.....	4	1	seq
.....ugaacacagcuggGgguaucca.....	6	1	seq
.....ugaacacagcuAgugguaucca.....	3	1	seq
.....ugaacGcagcuggugguaucca.....	7	1	seq
.....ugaacacagcugAugguaucca.....	5	1	seq
.....ugaacacagcuggugguauUca.....	3	1	seq
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.....ugGacacagcuggugguaucca.....	2	1	seq
.....ugaacacagcuggugguauccU.....	27	1	seq
.....Cgaacacagcuggugguaucca.....	1	1	seq
.....ugaacacagUuggugguaucca.....	1	1	seq
.....ugaacacagcCggugguaucca.....	4	1	seq
.....ugaacacagcugUugguaucca.....	4	1	seq
.....ugaacacaAcuggugguaucca.....	2	1	seq
.....ugaacacagcuggugguauccC.....	11	1	seq
.....uAaacacagcuggugguaucca.....	4	1	seq
.....ugaacacagcuggugguauGca.....	1	1	seq
.....ugaacacagcuggCgguaucca.....	5	1	seq
.....ugaGcacagcuggugguaucca.....	5	1	seq
.....ugaUcacagcuggugguaucca.....	1	1	seq
.....ugaacacagcuggugCuaucca.....	1	1	seq
.....ugaacacagcuggugguauccag.....	15	0	seq
.....ugaacacagcuggugguauccaC.....	1	1	seq
.....ugaacacagcuggugguauccaA.....	3	1	seq
.....ugaacacagcugAugguauccag.....	1	1	seq
.....ugaacacaCcuggugguauccag.....	1	1	seq
.....ugaacacagcuggugguauccGg.....	1	1	seq
.....ugaacacagcuggugguaAccag.....	1	1	seq
.....ugaacacagcuggCgguauccag.....	1	1	seq
.....ugaacacagcuggugUuauccag.....	1	1	seq
.....ugaacacagcuggugguauccaU.....	4	1	seq
.....ugaacacagcuggugguauccaCu.....	1	1	seq
.....ugaacacagcuggugguauccagC.....	1	1	seq
.....ugaacacagcuggGgguauccagu.....	1	1	seq
.....ugaacacagcuggugguauccagG.....	1	1	seq
.....ugaacacagcuggugguGuccagu.....	1	1	seq
.....ugaacacagcuggugguauccagu.....	121	0	seq
.....ugaacacagUuggugguauccagu.....	1	1	seq
.....ugaacacagcuggugGgauccagu.....	4	1	seq
.....ugaacacagcuggugguauccaUu.....	2	1	seq
.....ugaacacagcuggugguAccagu.....	1	1	seq
.....ugaGcacagcuggugguauccagu.....	1	1	seq
.....ugaacacagcuggugguaucGagu.....	1	1	seq
.....ugaacacagcuggugguauccagA.....	7	1	seq
.....ugaacacaUcuggugguauccagu.....	1	1	seq
.....ugaacacagcCggugguauccagu.....	2	1	seq
.....ugaacacagcuggugguauccCgu.....	1	1	seq
.....ugaacacagcuggugAuauccagu.....	2	1	seq
.....ugaacacagcuggAgguauccagu.....	1	1	seq
.....ugaacacagcugguAguauccagu.....	1	1	seq
.....ugaacacagcuggugguauccaguU.....	1	1	seq
.....ugaacacagcuggugguauccaguC.....	1	1	seq
.....ugaacacagcuggugguauccaguA.....	1	1	seq
.....Uaacacagcuggugguaucc.....	1	1	seq
.....gaacacagcuggugguaucc.....	6	0	seq

dme-miR-317-3p_MIMAT0000409

dme-miR-317-5p_MIMAT0020843

augcaacugccauugggauacacccugugcucgcuuugaaugaaaugcaagcaagugaacacagcuggugguauccaguggccguuuggcau

.....gaacacagcuggugguGucc.....	1	1	seq
.....gaacacagcuggugguaucca.....	1	0	seq
.....aacacagcuggugguaucca.....	1	0	seq
.....acacagcuggugguaucc.....	1	0	seq