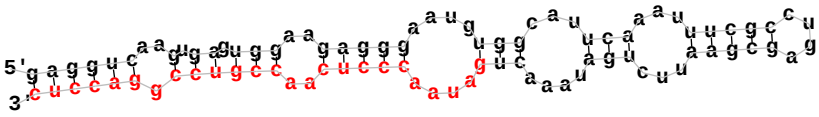
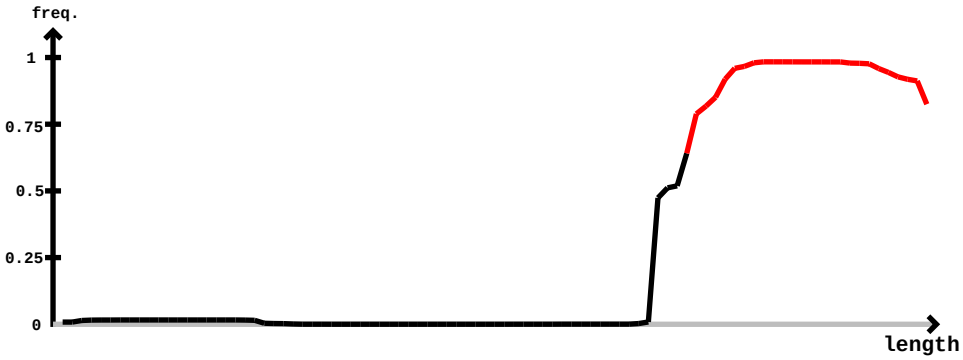
miRBase precursor : chromosome_14_73226
Total read count : 8219
chromosome_14_73226 read count : 3883
remaining reads : 4336





chromosome_14_73226

```
5'- gaggucaagugaguggaagagggaauguggcauucaaauuucgccugagcgaauucugauaaacugauaacccucaaccguccggaccuc  -3'   exp
    ((((((..(.((.(((..((((((....(((...(((...((((((....))))))...)))....)))....))))))..))))))).))))))   reads     mm        sample
    Aaggucaagugaguggaa.....................................................................................   1         1         seq
    Uaggucaagugaguggaa.....................................................................................   1         1         seq
    gaggucaagugaguggaa.....................................................................................   1         0         seq
    Uaggucaagugaguggaag....................................................................................   2         1         seq
    gUggucaagugaguggaag....................................................................................   1         1         seq
    Aaggucaagugaguggaag....................................................................................   3         1         seq
    gUggucaagugaguggaaga...................................................................................   2         1         seq
    gaggucaaguUaguggaaga...................................................................................   1         1         seq
    Aaggucaagugaguggaaga...................................................................................   16        1         seq
    gaggucaagugaguggaaga...................................................................................   12        0         seq
    Uaggucaagugaguggaaga...................................................................................   19        1         seq
    gaggucaagugaguggaagagg.................................................................................   2         0         seq
    Aaggucaagugaguggaagaggg................................................................................   1         1         seq
    gaggucaaguUaguggaagaggg................................................................................   4         1         seq
    gaggucaagugaguggaagagggG...............................................................................   1         1         seq
    .aggucaagugaguggaag....................................................................................   1         0         seq
    .aggucaaCugaguggaag....................................................................................   1         1         seq
    .aggucaaguUaguggaaga...................................................................................   8         1         seq
    .Uggucaagugaguggaaga...................................................................................   15        1         seq
    .Cggucaagugaguggaaga...................................................................................   2         1         seq
    .Gggucaagugaguggaaga...................................................................................   2         1         seq
    .aggucaagugaguggaaga...................................................................................   14        0         seq
    .Uggucaagugaguggaagag..................................................................................   1         1         seq
    .aggucaagugaguggaagag..................................................................................   1         0         seq
    .Gggucaagugaguggaagag..................................................................................   2         1         seq
    .aggucaaguUaguggaagagg.................................................................................   1         1         seq
    .aggucaaguUaguggaagaggg................................................................................   1         1         seq
    .aggucaagugaguggaagagggG...............................................................................   1         1         seq
    ..Ugucaagugaguggaagag..................................................................................   1         1         seq
    ..ggucaaguUaguggaagag..................................................................................   1         1         seq
    ..ggucaagugaguggaagag..................................................................................   1         0         seq
    ..Agucaagugaguggaagaggg................................................................................   1         1         seq
    ..Ugucaagugaguggaagaggg................................................................................   2         1         seq
    ..ggucaagugaguggaagaggg................................................................................   3         0         seq
    ..ggucaagugaguggaagagggG...............................................................................   1         1         seq
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chromosome_14_73226

gaggucaagugaguggaagagggaauguggcauucaaauuucgccugagcgaauucugauaaacugauaaacccucaaccguccggaccuc

..ggucaagugaguggaagagggaaug.....	1	0	seq
...gucaaguUaguggaagag.....	1	1	seq
...gucaagugaguggaagagggG.....	2	1	seq
.....caagugaguggaagagggG.....	1	1	seq
.....agggaauguggcauucaaauuucgccugag.....	1	0	seq
.....gagcgaauucugauaaacugauaacccuca.....	1	0	seq
.....cgaauucugauaaacugauaacccucaacc.....	1	0	seq
.....Gauucugauaaacugauaacccuc.....	1	1	seq
.....uaaaUugauaacccucaaccguccggaccu.	19	1	seq
.....aaaUugauaacccucaaccguccg.....	1	1	seq
.....aaaUugauaacccucaaccguccgga....	4	1	seq
.....aaaUugauaacccucaaccguccggac...	1	1	seq
.....aaaUugauaacccucaaccguccggaccu.	9	1	seq
.....aGacugauaacccucaaccguccggaccuc	1	1	seq
.....aaaUugauaacccucaaccguccggaccuc	32	1	seq
.....aaGcugauaacccucaaccguccggaccuc	1	1	seq
.....aaUugauaacccucaaccgucc.....	1	1	seq
.....aaUugauaacccucaaccguccg.....	25	1	seq
.....aaUugauaacccucaaccguccgg.....	18	1	seq
.....aaUugauaacccucaaccguccgga....	3	1	seq
.....aaUugauaacccucaaccguccggac...	11	1	seq
.....aaUugauaacccucaaccguccggacc..	1	1	seq
.....aGcugauaacccucaaccguccggaccu.	1	1	seq
.....aacGgauaacccucaaccguccggaccu.	1	1	seq
.....aaUugauaacccucaaccguccggaccu.	162	1	seq
.....aacGgauaacccucaaccguccggaccuc	12	1	seq
.....aacAgauaacccucaaccguccggaccuc	1	1	seq
.....aacugauaacccucaaccguccggaccuc	16	0	seq
.....Gacugauaacccucaaccguccggaccuc	5	1	seq
.....aaAugauaacccucaaccguccggaccuc	1	1	seq
.....aaUugauaacccucaaccguccggaccuc	3558	1	seq
.....aacugauaacccCcaaccguccggaccuc	1	1	seq
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.....aUugauaacccucaaccgu.....	16	1	seq
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.....aUugauaacccucaaccguccggac...	8	1	seq
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.....acugGuaacccucaaccguccggaccuc	2	1	seq
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.....Uugauaacccucaaccgucc.....	1	1	seq
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.....cGgauaacccucaaccguccgg.....	2	1	seq
.....Uugauaacccucaaccguccggacc..	1	1	seq
.....Uugauaacccucaaccguccggaccu.	10	1	seq
.....cGgauaacccucaaccguccggaccuc	1	1	seq
.....Uugauaacccucaaccguccggaccuc	22	1	seq
.....cAgauaacccucaaccguccggaccuc	2	1	seq
.....Augauaacccucaaccguccggaccuc	1	1	seq
.....ugauaacccucaaccguc.....	3	0	seq
.....Agauaacccucaaccguc.....	1	1	seq
.....Ggauaacccucaaccguc.....	1	1	seq
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.....Ggauaacccucaaccgucc.....	4	1	seq
.....Agauaacccucaaccgucc.....	1	1	seq
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.....ugauaacccucUaccguccg.....	1	1	seq
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.....Ggauaacccucaaccguccg.....	17	1	seq
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chromosome_14_73226

gaggucaagugaguggaagagggaauguggcauucaaauuucgccugagcgaauucugauaaacugauaacccucaaccguccggaccuc

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.....AgauaacccucaaccguccG.....	11	1	seq
.....ugauaacccucaaccgCccG.....	1	1	seq
.....Agauaacccucaaccguccgg.....	3	1	seq
.....ugGuaacccucaaccguccgg.....	2	1	seq
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.....ugauaacccucaaccguccggaccA.	1	1	seq
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.....uAauaacccucaaccguccggaccuc	4	1	seq
.....ugauaacccucaaccguccggaUcuc	1	1	seq
.....ugauaaAccucaaccguccggaccuc	1	1	seq
.....ugauaacccucaaccguccggGccuc	1	1	seq
.....ugauaacccucaaccgCccggaccuc	3	1	seq
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.....ugauaacUcucaaccguccggaccuc	1	1	seq
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.....gCuaacccucaaccguccG.....	1	1	seq
.....AauaacccucaaccguccG.....	8	1	seq
.....gGuaacccucaaccguccgg.....	4	1	seq
.....gauaaGccucaaccguccgg.....	1	1	seq
.....Aauaacccucaaccguccgg.....	2	1	seq
.....gauaacccucaaccguccgg.....	21	0	seq
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chromosome_14_73226

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.....gauaacUcucaaccguccgga....	1	1	seq
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.....gGuaacccucaaccguccggaccu.	15	1	seq
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.....Uauaacccucaaccguccggaccu.	4	1	seq
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.....gauaacccucaaccguccggaccuU	7	1	seq
.....gauaaGccucaaccguccggaccuc	1	1	seq
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.....gauaacccucaaccguccggaccuG	2	1	seq
.....gauaacccucaacGguccggaccuc	2	1	seq
.....gauaacccucaaccguccggaccGc	1	1	seq
.....gaGaacccucaaccguccggaccuc	1	1	seq
.....gauaacccucaaccguccgAaccuc	1	1	seq
.....gauaacccGcaaccguccggaccuc	2	1	seq
.....gauGacccucaaccguccggaccuc	2	1	seq
.....gauaacccuUaaccguccggaccuc	3	1	seq
.....Uauaacccucaaccguccggaccuc	8	1	seq
.....gauaacccucaaccgCccggaccuc	1	1	seq
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.....gauaacccucaaccgucGggaccuc	1	1	seq
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.....gauaacccucUaccguccggaccuc	3	1	seq
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.....gauaaAccucaaccguccggaccuc	4	1	seq
.....gauaacccucaGccguccggaccuc	2	1	seq
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.....gauaacccCcaaccguccggaccuc	5	1	seq
.....gauaacccucaaccguccggaccAc	2	1	seq
.....gauaacccucaaccguUcggaccuc	1	1	seq
.....gauaacccucaaccguccggaccCc	5	1	seq
.....gauaacccucaaccUuccggaccuc	1	1	seq
.....gauaacUcucaaccguccggaccuc	1	1	seq
.....gauaacccucaaccguccggaccuc	778	0	seq
.....gauCacccucaaccguccggaccuc	2	1	seq
.....gauaacccucGaccguccggaccuc	6	1	seq
.....gauaaccGucaaccguccggaccuc	1	1	seq
.....gauaacccucaUccguccggaccuc	1	1	seq
.....gauaacccucaaUcguccggaccuc	3	1	seq
.....gauaacccucaaccguccggaccuA	3	1	seq
.....gauaacAcucaaccguccggaccuc	1	1	seq
.....gauaacccucaacUguccggaccuc	1	1	seq
.....auaacccucaaccguccg......	11	0	seq
.....Guaacccucaaccguccg......	2	1	seq
.....Uuaacccucaaccguccg......	1	1	seq

chromosome_14_73226

gaggucaagugaguggaagagggaauguggcauucaaauuucgccugagcgaauucugauaaacugauaacccucaaccguccggaccuc

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.....auaacccucaaccgucUgg.....	2	1	seq
.....auaacccucaaccguccgg.....	9	0	seq
.....auaacccucaaccguccgA.....	1	1	seq
.....auaacUcucaaccguccgga....	1	1	seq
.....auaacAcucaaccguccgga....	2	1	seq
.....auaacccucaaccguccgga....	27	0	seq
.....Guaacccucaaccguccgga....	1	1	seq
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.....auaacccucaaccguccggacc..	1	0	seq
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.....Guaacccucaaccguccggaccu.	6	1	seq
.....auaacUcucaaccguccggaccu.	2	1	seq
.....auaacccucaaccgCccggaccuc	1	1	seq
.....auaacccucaaccguccggaccCc	1	1	seq
.....auaacccucaaUcguccggaccuc	1	1	seq
.....Cuaacccucaaccguccggaccuc	1	1	seq
.....aAaacccucaaccguccggaccuc	8	1	seq
.....auaacccucaaccguccggaccuc	108	0	seq
.....aCaacccucaaccguccggaccuc	4	1	seq
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.....Uuaacccucaaccguccggaccuc	2	1	seq
.....Guaacccucaaccguccggaccuc	13	1	seq
.....auaGcccucaaccguccggaccuc	1	1	seq
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.....uaacccucaaccguccgg.....	3	0	seq
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.....uaacccucaaccguccggacc..	2	0	seq
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.....Caacccucaaccguccggaccu.	1	1	seq
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.....uaacccucaaccguccggaccu.	42	0	seq
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.....Gaacccucaaccguccggaccu.	1	1	seq
.....uGacccucaaccguccggaccuc	1	1	seq
.....Gaacccucaaccguccggaccuc	3	1	seq
.....uaacccucaaccguccggaccCc	1	1	seq
.....uaacccuAaaccguccggaccuc	1	1	seq
.....Aaacccucaaccguccggaccuc	33	1	seq
.....uUacccucaaccguccggaccuc	2	1	seq
.....uaacccucaaccguccggaAcuc	1	1	seq
.....uaacccucaaccguccggGccuc	3	1	seq
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.....uaacccucaGccguccggaccuc	2	1	seq
.....uaacccucGaccguccggaccuc	1	1	seq
.....uaacccucaaccguccggaccuU	1	1	seq
.....uaacccucaaccguccggaccuc	151	0	seq
.....uaacccucaaccguccggaccAc	1	1	seq
.....uaacccucaacAguccggaccuc	1	1	seq
.....Gacccucaaccguccgga....	4	1	seq
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.....aacccucaaccguccggC....	1	1	seq
.....aacccucaaccAuccgga....	2	1	seq
.....aacccucaacAguccggac...	1	1	seq
.....aUcccucaaccguccggac...	1	1	seq
.....aacccucaacUguccggac...	1	1	seq
.....aacccucaaccguccggac...	7	0	seq
.....Gacccucaaccguccggac...	1	1	seq
.....aacccucaaccguccggacc..	9	0	seq
.....Gacccucaaccguccggaccu.	6	1	seq

chromosome_14_73226

gaggucaagugaguggaagagggaauguggcauucaaauuucgccugagcgaauucugauaaacugauaacccucaaccguccggaccuc

.....aacccucaaccAuccggaccu.	2	1	seq
.....Uacccucaaccguccggaccu.	1	1	seq
.....aacccucGaccguccggaccu.	2	1	seq
.....aCcccucaaccguccggaccu.	1	1	seq
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.....aacccucaaAcguccggaccu.	1	1	seq
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.....aacccucaaccguGcggaccuc	1	1	seq
.....aCcccucaaccguccggaccuc	1	1	seq
.....Uacccucaaccguccggaccuc	6	1	seq
.....aacccucaaccgCccggaccuc	1	1	seq
.....aacccucaCccguccggaccuc	1	1	seq
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.....aacUcucaaccguccggaccuc	1	1	seq
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.....aacccucaaccguccAgaccuc	1	1	seq
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.....aacccucaaccguccggacUuc	2	1	seq
.....aacccucaacGguccggaccuc	3	1	seq
.....aacccucaaccguccggGccuc	4	1	seq
.....aacccucaaccguccggaccuU	3	1	seq
.....aaccUucaaccguccggaccuc	1	1	seq
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.....aacccucaaccUuccggaccuc	1	1	seq
.....aacccCcaaccguccggaccuc	3	1	seq
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.....aacccucGaccguccggaccuc	2	1	seq
.....aacccucaaccguccggaccuG	1	1	seq
.....aacccucaaccguccggaccuc	325	0	seq
.....aacccGcaaccguccggaccuc	2	1	seq
.....aacccucaGccguccggaccuc	2	1	seq
.....aacccucaaccAuccggaccuc	1	1	seq
.....aacccucaacAguccggaccuc	1	1	seq
.....acccucaaccguccggac...	15	0	seq
.....acccucaaccguccggacc..	3	0	seq
.....acccucaaGcguccggacc..	1	1	seq
.....acccucaaccguccAgaccu.	1	1	seq
.....Gcccucaaccguccggaccu.	1	1	seq
.....acccucaaccguccggaCUu.	1	1	seq
.....acccucaaccguccggaccu.	21	0	seq
.....acccucaaccguccggGccu.	1	1	seq
.....acccAcaaccguccggaccu.	1	1	seq
.....acccucaacUguccggaccuc	2	1	seq
.....acccucaaccguccggaccuU	1	1	seq
.....acccucaaccguccggaAcuc	1	1	seq
.....aUccucaaccguccggaccuc	1	1	seq
.....acccucaGccguccggaccuc	2	1	seq
.....acccucaaccguccggaccuc	230	0	seq
.....acccucaaAcguccggaccuc	1	1	seq
.....aAccucaaccguccggaccuc	1	1	seq
.....acccucaaccguccggacAuc	1	1	seq
.....acccucaaccAuccggaccuc	2	1	seq
.....Ucccucaaccguccggaccuc	6	1	seq
.....Gcccucaaccguccggaccuc	34	1	seq
.....acAcucaaccguccggaccuc	1	1	seq
.....acccucaaccguccggaccAc	1	1	seq
.....acccucaaccguccggCccuc	1	1	seq
.....Ccccucaaccguccggaccuc	3	1	seq
.....acccucaaccguccAgaccuc	1	1	seq
.....cccucaaccguccggacc..	1	0	seq
.....cccucaaccguAcggacc..	1	1	seq
.....Accucaaccguccggaccu.	1	1	seq
.....cccucaaccguccggaccu.	4	0	seq

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gaggucaagugaguggaagagggaauguggcauucaaauuucgccugagcgaauucugauaaacugauaacccucaaccguccggaccuc

.....Uccucaaccguccggaccuc	7	1	seq
.....cccucaaAcguccggaccuc	1	1	seq
.....cccucaaccguccggaccuc	42	0	seq
.....cccucaaccguccggaccuU	1	1	seq
.....Gccucaaccguccggaccuc	1	1	seq
.....Ucucaaccguccggaccu.	1	1	seq
.....ccucaaccAuccggaccu.	2	1	seq
.....ccucaaccguccggaccu.	21	0	seq
.....ccucaaccguccggGccuc	1	1	seq
.....Acucaaccguccggaccuc	2	1	seq
.....Ucucaaccguccggaccuc	9	1	seq
.....ccucaaccguccggaccuc	65	0	seq
.....ccucaaccguccgAaccuc	2	1	seq
.....ccucaaccCuccggaccuc	1	1	seq
.....ccucaaccUuccggaccuc	1	1	seq
.....Gcucaaccguccggaccuc	2	1	seq
.....cUucaaccguccggaccuc	1	1	seq
.....ccucaaccAuccggaccuc	2	1	seq
.....ccucaaccguccggaccCc	1	1	seq
.....cucaaccguccggaccuU	1	1	seq
.....cucaaccguAcggaccuc	1	1	seq
.....cucaaccguccggaccuc	27	0	seq
.....Uucaaccguccggaccuc	1	1	seq