chromosome_7_30929

```
5'- acccguagauccgaacuugugggauuuuucuccacaaguucggcuccauggucc   -3'     exp
    ..(((((.((.(((((((((((((((....)).)))))))))))))).)).)))...    reads    mm       sample
    Ccccguagauccgaacuu.................................          1        1        seq
    acccguagauccgaacuC.................................          13       1        seq
    acccguagauccgaacuA.................................          4        1        seq
    acUcguagauccgaacuu.................................          1        1        seq
    acccAuagauccgaacuu.................................          1        1        seq
    acccgCagauccgaacuu.................................          2        1        seq
    acccguagauccgaacuu.................................          216      0        seq
    acccguagauccgaacCu.................................          1        1        seq
    acAcguagauccgaacuu.................................          1        1        seq
    acccguGgauccgaacuu.................................          2        1        seq
    acccguagauUcgaacuu.................................          1        1        seq
    aAccguagauccgaacuu.................................          1        1        seq
    acccguagauccgaaGuu.................................          1        1        seq
    acccguagauccgaacGu.................................          2        1        seq
    acccguagGuccgaacuu.................................          2        1        seq
    acccguagauccgaacAug................................          1        1        seq
    accUguagauccgaacuug................................          1        1        seq
    acccguagauccAaacuug................................          1        1        seq
    acccguagaCccgaacuug................................          1        1        seq
    acccgCagauccgaacuug................................          1        1        seq
    acccguCgauccgaacuug................................          1        1        seq
    Ccccguagauccgaacuug................................          1        1        seq
    acccguagauccgaaUuug................................          2        1        seq
    Gcccguagauccgaacuug................................          4        1        seq
    aAccguagauccgaacuug................................          5        1        seq
    acccguagauccgaCcuug................................          1        1        seq
    acccguagauccgaaGuug................................          2        1        seq
    acccguagGuccgaacuug................................          7        1        seq
    acccguagauccgaacuug................................          1012     0        seq
    acccguagauccgaacuuA................................          163      1        seq
    acccguagaucUgaacuug................................          2        1        seq
    acccguagaAccgaacuug................................          1        1        seq
    acccguagauccgaGcuug................................          12       1        seq
    acccguagauccCaacuug................................          1        1        seq
    acccguagauccgaacGug................................          1        1        seq
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acccguagauccgaacuugugggauuuuucuccacaaguucggcuccaugguccc

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acccguagauccgaacuuC.................................    1      1        seq
acccguagauccgaacuCg.................................    1      1        seq
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acccguaAauccgaacuug.................................    2      1        seq
acccguagauccgGacuug.................................    2      1        seq
acccguagauccgCacuug.................................    3      1        seq
acccguGgauccgaacuug.................................    4      1        seq
acGcguagauccgaacuug.................................    1      1        seq
acccguagauccgaacuGg.................................    2      1        seq
acccguagauccgaacuAg.................................    2      1        seq
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acUcguagauccgaacuug.................................    3      1        seq
acccguagauccgaacuuU.................................    20     1        seq
acccguagauccUaacuug.................................    1      1        seq
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acccgAagauccgaacuugu................................    4      1        seq
acccguagauAcgaacuugu................................    1      1        seq
acccguagauccgaaGuugu................................    4      1        seq
acccguagauccgaacuAu.................................    24     1        seq
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acccguagauccgUacuugu................................    1      1        seq
Ucccguagauccgaacuugu................................    11     1        seq
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Gcccguagauccgaacuugu................................    14     1        seq
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acccguagauccgaacuuUu................................    5      1        seq
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acccguagauccgCacuugu................................    12     1        seq
acccguaAauccgaacuugu................................    4      1        seq
acccguaCauccgaacuugu................................    2      1        seq
```

chromosome_7_30929

acccguagauccgaacuugugggauuuuucuccacaaguucggcuccauggucc

acccguagauccgaacuuCu.................................	3	1	seq
acccguagauUcgaacuugug...............................	2	1	seq
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.cccguagauccgaacuuA..................................	2	1	seq
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.cccguagauccgaacuugC.................................	10	1	seq

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.cccguagauccgaacuugugg.............................     7      0     seq
..ccguagauccgaaUuugu..............................      1      1     seq
...cguagauccgaacuugug.............................      2      0     seq
.....uagauccgaacuuguggga...........................     2      0     seq
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.................................acaaguucggcuccaugg...  34     0     seq
.................................acaaguucggcCccaugg...  1      1     seq
.................................acaaguucggcuccaugA...  8      1     seq
.................................Gcaaguucggcuccaugg...  1      1     seq
.................................acaaguucggcuccaugU...  3      1     seq
.................................acaaguucggcuccauggu..  7      0     seq
.................................acaaguucggcuccauggA..  2      1     seq
.................................acaaguucggcuccaugAu..  1      1     seq
.................................acaaguucggcuccauggC..  2      1     seq
.................................acaaguucggcuccaugguA.  1      1     seq
.................................acaaguucgUcuccaugguc.  1      1     seq
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.................................acaaguucggcuccaugguU.  2      1     seq
.................................acaaguucgAcuccaugguc.  1      1     seq
.................................acaaguucggcuccaugguc.  25     0     seq
.................................acaaguucggcuccauggucU  13     1     seq
.................................acaaguucggcuAcauggucc  1      1     seq
.................................acaaguucggcuccauggucG  1      1     seq
.................................acaaguucggcuccauggucA  23     1     seq
.................................acaaguucggcuccauggucc  16     0     seq
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