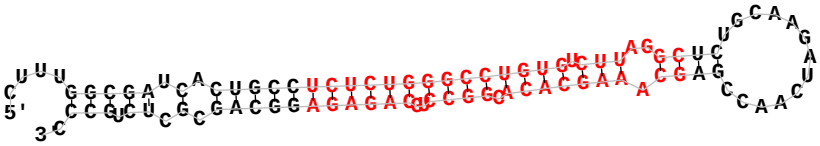
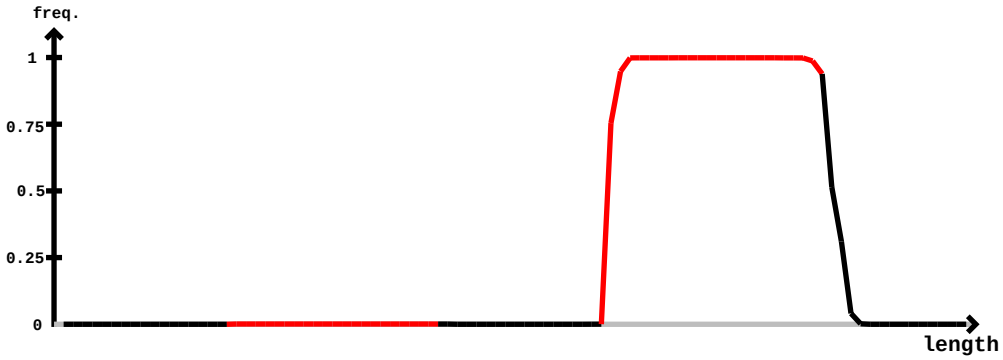
miRBase precursor : hsa-mir-330
Total read count : 4916
hsa-miR-330-3p read count : 4911
hsa-miR-330-5p read count : 5
remaining reads : 4911





hsa-miR-330-5p

hsa-miR-330-3p

```
5'- cuuuggcgaucacugccucucugggccugugucuuaggcucugcaagaucaaccgagcaaagcacacggccugcagagaggcagcgcucugccc  -3'    exp
    ....(((((.(.((((((((((((((((((((.(((..((((.............)))).)))))))..))))..)))))))))))).).)).))).   reads      mm         sample
    .................ucucugggccugugucuuagg.....................................................................   1          0          hsa
    .................ucucugggccugugucuuaggcA...................................................................   1          1          hsa
    .................ucucugggccugugucuuaggcu..................................................................   3          0          hsa
    ....................................................Aagcaaagcacacggccugcagaga..............................   1          1          hsa
    ....................................................agcaaagcacacggccugcaga.................................   1          0          hsa
    ....................................................gcaaagcacacggccugc.....................................   2          0          hsa
    ....................................................gcaaagcacacggccugcag....................................   1          0          hsa
    ....................................................gAaaagcacacggccugcaga...................................   1          1          hsa
    ....................................................gcaaagcacacggccugcaga...................................   47         0          hsa
    ....................................................gcaaagcacacggccugcCga...................................   1          1          hsa
    ....................................................gcaaagcacacggccugcaUa...................................   1          1          hsa
    ....................................................gcaaagAacacggccugcaga...................................   1          1          hsa
    ....................................................gcaaagcacacggccugcagag..................................   191        0          hsa
    ....................................................gcaaagUacacggccugcagag..................................   1          1          hsa
    ....................................................gcaaaCcacacggccugcagag..................................   1          1          hsa
    ....................................................gcaaagcacacgAccugcagag..................................   1          1          hsa
    ....................................................gcaaagcacacggccugcagaA..................................   5          1          hsa
    ....................................................gcaaagcacacggccugcaCag..................................   1          1          hsa
    ....................................................gcaaagcCcacggccugcagag..................................   1          1          hsa
    ....................................................gcaaagcacacggccugcaUag..................................   3          1          hsa
    ....................................................gcaCagcacacggccugcagag..................................   1          1          hsa
    ....................................................gcaaagcacacUgccugcagag..................................   2          1          hsa
    ....................................................gcaaCgcacacggccugcagag..................................   3          1          hsa
    ....................................................gcaaagcacacggccugcCgag..................................   1          1          hsa
    ....................................................gcaaagcacacggcGugcagag..................................   1          1          hsa
    ....................................................gcaaagcacacggccugcagaU..................................   1          1          hsa
    ....................................................gcaaagcacacggccugcagCg..................................   2          1          hsa
    ....................................................gcaaagGacacggccugcagag..................................   1          1          hsa
    ....................................................gcaaagcacacggccugcagaUa.................................   11         1          hsa
    ....................................................gcaaagcacacggccugcaAaga.................................   3          1          hsa
    ....................................................gcaaagcacacggccugcagagC.................................   3          1          hsa
    ....................................................gcUaagcacacggccugcagaga.................................   2          1          hsa
    ....................................................gcaCagcacacggccugcagaga.................................   8          1          hsa
    ....................................................gcaaagcaUacggccugcagaga.................................   5          1          hsa
```

hsa-miR-330-5p

hsa-miR-330-3p

cuuuggcgaucacugccucucugggccugugucuuaggcucugcaagaucaaccgagcaaagcacacggccugcagagaggcagcgcucugccc

.....gcaaagcacacggccuAcagaga.....	2	1	hsa
.....gcaaagcacacggUcugcagaga.....	2	1	hsa
.....gcaaagcacacggcAugcagaga.....	1	1	hsa
.....gcaaagcacacggccugcagagG.....	6	1	hsa
.....gcaaagcacacggccCgcagaga.....	3	1	hsa
.....gcaaagcacacggccugcagUga.....	4	1	hsa
.....gAaaagcacacggccugcagaga.....	1	1	hsa
.....gcaaagcacacggAcugcagaga.....	2	1	hsa
.....gcaaagcacacggccAgcagaga.....	1	1	hsa
.....gcaaagcacacCgccugcagaga.....	2	1	hsa
.....gcaaagcacacggccugcUgaga.....	4	1	hsa
.....gcaaagcacacggccugUagaga.....	1	1	hsa
.....gUaaagcacacggccugcagaga.....	2	1	hsa
.....gcGaagcacacggccugcagaga.....	1	1	hsa
.....gcaaagcacacggccuUcagaga.....	9	1	hsa
.....gcCaagcacacggccugcagaga.....	3	1	hsa
.....gcaaagcacacggGcugcagaga.....	1	1	hsa
.....gcaaagcGcacggccugcagaga.....	1	1	hsa
.....gcaaagcacacggccugcagaga.....	1564	0	hsa
.....gcaaagcacacgAccugcagaga.....	2	1	hsa
.....gcaaagcacacggccugcagagU.....	140	1	hsa
.....gcaaaUcacacggccugcagaga.....	1	1	hsa
.....gcaaagcacacggccugcagaCa.....	2	1	hsa
.....gcaaagcacacggcUugcagaga.....	2	1	hsa
.....gcaaagAacacggccugcagaga.....	3	1	hsa
.....gcaaagcacacgUccugcagaga.....	2	1	hsa
.....gcaaagcacGcggccugcagaga.....	4	1	hsa
.....gcaaagcacacggccugAagaga.....	1	1	hsa
.....gcaaCgcacacggccugcagaga.....	8	1	hsa
.....gcaaagcacacggccugcagaAa.....	5	1	hsa
.....gcaaagcacacggccugcaUaga.....	18	1	hsa
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.....gcaaagcacacggccugcagagaC.....	6	1	hsa
.....gcaaCgcacacggccugcagagag.....	2	1	hsa
.....gcaaagcacUcggccugcagagag.....	2	1	hsa
.....gcaaagcacacggccuUcagagag.....	7	1	hsa
.....gcaaaCcacacggccugcagagag.....	1	1	hsa
.....gcaaagcacacggccugcagagaU.....	137	1	hsa
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.....gcaaagcacacggccugcagGgag.....	1	1	hsa
.....gcCaagcacacggccugcagagag.....	3	1	hsa

hsa-miR-330-5p

hsa-miR-330-3p

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.....gcaaagcacacggccugcagagagA.....	40	1	hsa
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.....caaagcacacggccugcagagU.....	5	1	hsa
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.....caaagcacacggccugcagagaA.....	5	1	hsa
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hsa-miR-330-5p

hsa-miR-330-3p

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.....caaagcacacgUccugcagagagg.....	3	1	hsa
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.....aaagcacacggccugcagagagA.....	10	1	hsa
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.....aaagcacacggccugcagagaggUa.....	1	1	hsa
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