miRBase precursor : hsa-mir-191
Total read count : 30235
hsa-miR-191-5p read count : 30195
hsa-miR-191-3p read count : 40
remaining reads : 40





hsa-miR-191-3p

hsa-miR-191-5p

```
5'- cggcuggacagcgggcaacggaaucccaaaagcagcuguugucuccagagcauuccagcugcgcuuggauuucgucccugcucuccugccu  -3'   exp
    .(((.(((.((((((..((((((((((.((..(((((((..((((....)).)).)..)))))))..))))))))))..))))))).))).   reads     mm        sample
    ..............Acaacggaaucccaaaagcagcug.....................................................   4         1         hsa
    ..............Ucaacggaaucccaaaagcagcug.....................................................   1         1         hsa
    ..............Ccaacggaaucccaaaagcagcugu....................................................   1         1         hsa
    ..............caacggaaucccaaaagc..........................................................   5         0         hsa
    ..............caacggaaCcccaaaagca........................................................   1         1         hsa
    ..............caacggaaucccaaaaUca........................................................   1         1         hsa
    ..............caacggaaucccaaaagca........................................................   12        0         hsa
    ..............caacggaaucccaaaCgcag.......................................................   2         1         hsa
    ..............caaUggaaucccaaaagcag.......................................................   1         1         hsa
    ..............cUacggaaucccaaaagcag.......................................................   1         1         hsa
    ..............caacgAaaucccaaaagcag.......................................................   1         1         hsa
    ..............caacggaaucccaaaagcag.......................................................   50        0         hsa
    ..............caacggaaucccaaUagcagc......................................................   1         1         hsa
    ..............caacggaaucccCaaagcagc......................................................   1         1         hsa
    ..............cCacggaaucccaaaagcagc......................................................   3         1         hsa
    ..............caacggaaucUcaaaagcagc......................................................   1         1         hsa
    ..............caacggaaucccaaaagcagc......................................................   180       0         hsa
    ..............caacggaaucccaaaagcCgc......................................................   2         1         hsa
    ..............caacUgaaucccaaaagcagc......................................................   2         1         hsa
    ..............caacggaaucccaaGagcagc......................................................   1         1         hsa
    ..............caacggaaucccaaaaUcagc......................................................   1         1         hsa
    ..............caacggCaucccaaaagcagc......................................................   1         1         hsa
    ..............caacggaaucccaaaagcaUc......................................................   2         1         hsa
    ..............caGcggaaucccaaaagcagc......................................................   1         1         hsa
    ..............caacggaaucccaaaagUagc......................................................   1         1         hsa
    ..............caacggaauUccaaaagcagc......................................................   2         1         hsa
    ..............Uaacggaaucccaaaagcagc......................................................   1         1         hsa
    ..............caacggaaucccaaaGgcagcu.....................................................   1         1         hsa
    ..............caacggaaucccaaaagUagcu.....................................................   6         1         hsa
    ..............caacggaaucccaaaagcCgcu.....................................................   4         1         hsa
    ..............caacggaaucccaaaaUcagcu.....................................................   13        1         hsa
    ..............Uaacggaaucccaaaagcagcu.....................................................   6         1         hsa
    ..............caacggaaucccaaaagcGgcu.....................................................   1         1         hsa
    ..............caacggaaucccaaaagGagcu.....................................................   1         1         hsa
```

hsa-miR-191-3p

hsa-miR-191-5p

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.....caacgUaaucccaaaagcagcu.....	2	1	hsa
.....caacgCaaucccaaaagcagcu.....	2	1	hsa
.....cCacggaaucccaaaagcagcu.....	15	1	hsa
.....caacggaauAccaaaagcagcu.....	2	1	hsa
.....caacggaaucccaaaagcaUcu.....	14	1	hsa
.....caacggaaucccaaaCgcagcu.....	15	1	hsa
.....caacAgaaucccaaaagcagcu.....	4	1	hsa
.....caacggaaucccaaaaAcagcu.....	3	1	hsa
.....caacggCaucccaaaagcagcu.....	10	1	hsa
.....caacggaGucccaaaagcagcu.....	1	1	hsa
.....caacggaaucUcaaaagcagcu.....	8	1	hsa
.....caacggaauGccaaaagcagcu.....	1	1	hsa
.....cGacggaaucccaaaagcagcu.....	2	1	hsa
.....caacggaCucccaaaagcagcu.....	4	1	hsa
.....caacggaaucccaaaagcagcu.....	1878	0	hsa
.....caacggaaucccaaaagcagcC.....	1	1	hsa
.....caaUggaaucccaaaagcagcu.....	1	1	hsa
.....caacggUaucccaaaagcagcu.....	1	1	hsa
.....caacggaaucccaaaaCcagcu.....	1	1	hsa
.....caacggaaucccaaaagcagUu.....	6	1	hsa
.....caacggaauccUaaaagcagcu.....	3	1	hsa
.....caacgAaaucccaaaagcagcu.....	4	1	hsa
.....caacggaaucccaaGagcagcu.....	4	1	hsa
.....caCcggaaucccaaaagcagcu.....	1	1	hsa
.....caacggaaucccaaaagcUgcu.....	1	1	hsa
.....caacggaauccAaaaagcagcu.....	1	1	hsa
.....caacggaaucccaaUagcagcu.....	3	1	hsa
.....Gaacggaaucccaaaagcagcu.....	1	1	hsa
.....caacggaaucccaGaagcagcu.....	1	1	hsa
.....caacggaaucccaaCagcagcu.....	6	1	hsa
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.....caacggaaucAcaaaagcagcu.....	2	1	hsa
.....caacggaaucccaaaagcaCcu.....	2	1	hsa
.....caacggaaucccaaaUgcagcu.....	1	1	hsa
.....caaAggaaucccaaaagcagcu.....	1	1	hsa
.....caGcggaaucccaaaagcagcu.....	1	1	hsa
.....caacUgaaucccaaaagcagcu.....	3	1	hsa
.....caacggaUucccaaaagcagcu.....	2	1	hsa
.....caacggaaucccaaaagcagcA.....	4	1	hsa
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.....caacggaaCcccaaaagcagcu.....	3	1	hsa
.....caacggaaucccaaaagcagcG.....	2	1	hsa
.....caacggaauUccaaaagcagcu.....	8	1	hsa
.....Aaacggaaucccaaaagcagcug.....	6	1	hsa
.....caacggaaucccaaaagcaCcug.....	11	1	hsa
.....caacggaauccUaaaagcagcug.....	42	1	hsa
.....caacggaaucccaaaaUcagcug.....	93	1	hsa
.....cCacggaaucccaaaagcagcug.....	50	1	hsa
.....caacggaaucccaaaagcCgcug.....	32	1	hsa
.....caacggGaucccaaaagcagcug.....	14	1	hsa
.....caacggaaucccaaaagcagAug.....	5	1	hsa
.....caacggaaucccaaaGgcagcug.....	5	1	hsa
.....caacggaaucccaGaagcagcug.....	6	1	hsa
.....caaGggaaucccaaaagcagcug.....	8	1	hsa
.....caaAggaaucccaaaagcagcug.....	13	1	hsa
.....caacggaaucccaaCagcagcug.....	39	1	hsa
.....caacggaUucccaaaagcagcug.....	3	1	hsa
.....Uaacggaaucccaaaagcagcug.....	47	1	hsa
.....caacggaaucccUaaagcagcug.....	3	1	hsa
.....caacggaaucccaaaagAagcug.....	6	1	hsa
.....cGacggaaucccaaaagcagcug.....	11	1	hsa
.....caaUggaaucccaaaagcagcug.....	27	1	hsa
.....caacggaauUccaaaagcagcug.....	60	1	hsa
.....caacggaaGcccaaaagcagcug.....	1	1	hsa
.....caacggCaucccaaaagcagcug.....	52	1	hsa
.....caacggaaucccaaaagcagcGg.....	4	1	hsa
.....caacggaaucccaaaagcUgcug.....	8	1	hsa
.....caacggaGucccaaaagcagcug.....	9	1	hsa
.....caacggaaucAcaaaagcagcug.....	8	1	hsa

hsa-miR-191-3p

hsa-miR-191-5p

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.....caacggaaucccaaaCgcagcug.....	76	1	hsa
.....caacCgaaucccaaaagcagcug.....	6	1	hsa
.....caacggaaucccaaGagcagcug.....	15	1	hsa
.....caacAgaaucccaaaagcagcug.....	11	1	hsa
.....caacggaauGccaaaagcagcug.....	29	1	hsa
.....caacggaaucccaaaagcagcuA.....	100	1	hsa
.....caacggaauccAaaaagcagcug.....	11	1	hsa
.....caacggaaucccaaaagcagGug.....	3	1	hsa
.....caacgUaaucccaaaagcagcug.....	28	1	hsa
.....caacggaaucccaaaagcaAcug.....	8	1	hsa
.....caacggaaucccaaaagcagcug.....	13350	0	hsa
.....caacggaaCcccaaaagcagcug.....	9	1	hsa
.....caacggaaucUcaaaagcagcug.....	53	1	hsa
.....caacggaaucccaaaUgcagcug.....	5	1	hsa
.....cUacggaaucccaaaagcagcug.....	2	1	hsa
.....caacggaCucccaaaagcagcug.....	30	1	hsa
.....caacggaauccGaaaagcagcug.....	3	1	hsa
.....caacggUaucccaaaagcagcug.....	6	1	hsa
.....caacggaaucccaCaagcagcug.....	21	1	hsa
.....caacggaaucccaUaagcagcug.....	6	1	hsa
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.....caacggaaucccaaaagcagcuU.....	99	1	hsa
.....caacgAaaucccaaaagcagcug.....	5	1	hsa
.....caacggaaucccaaaagUagcug.....	12	1	hsa
.....caCcggaaucccaaaagcagcug.....	33	1	hsa
.....caacggaaucccCaaagcagcug.....	15	1	hsa
.....caacggaaucccaaaagcagcCg.....	20	1	hsa
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.....caacgCaaucccaaaagcagcug.....	5	1	hsa
.....caacggaauAccaaaagcagcug.....	34	1	hsa
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.....caGcggaaucccaaaagcagcug.....	3	1	hsa
.....caacggaaucGcaaaagcagcug.....	5	1	hsa
.....caacggaaucccaaaagcagcAg.....	9	1	hsa
.....caacggaaAcccaaaagcagcug.....	7	1	hsa
.....caacggaaucccaaaagcagcuC.....	10	1	hsa
.....caacggaaucccaaaagcaUcug.....	113	1	hsa
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.....caacggaaucccaaaagAagcugu.....	1	1	hsa
.....caacggaaucccaaaagcaCcugu.....	1	1	hsa
.....caacgCaaucccaaaagcagcugu.....	2	1	hsa
.....caacggaCucccaaaagcagcugu.....	8	1	hsa
.....caGcggaaucccaaaagcagcugu.....	5	1	hsa
.....caacggaaucccaUaagcagcugu.....	2	1	hsa
.....caacggaaucccaCaagcagcugu.....	8	1	hsa
.....caacggaaCcccaaaagcagcugu.....	5	1	hsa
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.....caacggaaGcccaaaagcagcugu.....	2	1	hsa
.....Aaacggaaucccaaaagcagcugu.....	2	1	hsa
.....caacggaaucccaaUagcagcugu.....	9	1	hsa
.....caacggaaucccGaaagcagcugu.....	5	1	hsa
.....caacggaaucccaaaagGagcugu.....	3	1	hsa
.....cCacggaaucccaaaagcagcugu.....	39	1	hsa
.....caacggaaucccaaaagcagcAgu.....	4	1	hsa
.....caacggaaucccaaaagcagcugG.....	26	1	hsa

hsa-miR-191-3p

hsa-miR-191-5p

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.....caacggaUucccaaaagcagcugu.....	4	1	hsa
.....caacUgaaucccaaaagcagcugu.....	15	1	hsa
.....caacggaaucccaaaagcCgcugu.....	15	1	hsa
.....caaUggaaucccaaaagcagcugu.....	13	1	hsa
.....caacggUaucccaaaagcagcugu.....	8	1	hsa
.....caacggaaucccaaaagcagcugu.....	7592	0	hsa
.....caacggaaucccaaaaAcagcugu.....	7	1	hsa
.....caacggaaucccaaaagUagcugu.....	7	1	hsa
.....caacggaaucAcaaaagcagcugu.....	3	1	hsa
.....caacggaaAcccaaaagcagcugu.....	2	1	hsa
.....caacCgaaucccaaaagcagcugu.....	4	1	hsa
.....caacggaaucccaaaagcagcCgu.....	10	1	hsa
.....caacggaaucccUaaagcagcugu.....	2	1	hsa
.....caacggGaucccaaaagcagcugu.....	7	1	hsa
.....caacggaauAccaaaagcagcugu.....	31	1	hsa
.....caacggaaucccaaaagcUgcugu.....	6	1	hsa
.....caacggaaucccaaaGgcagcugu.....	1	1	hsa
.....caacggaaucccaaaUgcagcugu.....	5	1	hsa
.....caacgUaaucccaaaagcagcugu.....	21	1	hsa
.....caacggaaucGcaaaagcagcugu.....	2	1	hsa
.....caacggaaucccaaaagcaUcugu.....	76	1	hsa
.....caacggaaucUcaaaagcagcugu.....	25	1	hsa
.....caacggaaucccaaaagcagUugu.....	35	1	hsa
.....caacggaaucccaaaagcagcugA.....	1895	1	hsa
.....caaGggaaucccaaaagcagcugu.....	6	1	hsa
.....caacggaaucccaGaagcagcugu.....	4	1	hsa
.....cUacggaaucccaaaagcagcugu.....	2	1	hsa
.....caacgAaaucccaaaagcagcugu.....	9	1	hsa
.....caacggaaucccaaaagcGgcugu.....	5	1	hsa
.....caacAgaaucccaaaagcagcugu.....	14	1	hsa
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.....caacggaaucccaaaCgcagcugu.....	38	1	hsa
.....caacggaaucccaaaagcagcuCu.....	3	1	hsa
.....caacggaaucccaaaagcagcugC.....	53	1	hsa
.....Uaacggaaucccaaaagcagcugu.....	17	1	hsa
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.....caaAggaaucccaaaagcagcugu.....	7	1	hsa
.....caacggCaucccaaaagcagcugu.....	33	1	hsa
.....caacggaauccGaaaagcagcugu.....	7	1	hsa
.....caCcggaaucccaaaagcagcugu.....	14	1	hsa
.....caacggaaucccaaaagcagGugu.....	3	1	hsa
.....caacggaauUccaaaagcagcugu.....	32	1	hsa
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.....caacggaaucccaaaagcagcuUu.....	34	1	hsa
.....caacggaaucccaaaaUcagcugu.....	38	1	hsa
.....caacggaaucccaaaagcagcuguA.....	202	1	hsa
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.....caacggaaucccaaaagcagcugGu.....	2	1	hsa
.....caacggaaucccaaaagcagcuUuu.....	2	1	hsa
.....Aaacggaaucccaaaagcagcuguu.....	1	1	hsa
.....caacggaaucccaaaagcagcugCu.....	1	1	hsa
.....caacggaaucccaaaagcagcuguG.....	7	1	hsa
.....caacggaaucccaaaaUcagcuguu.....	2	1	hsa
.....caacggaaucccaaaagcagUuguu.....	2	1	hsa
.....caacggaaucccaaaagcagcuguC.....	3	1	hsa
.....caacggCaucccaaaagcagcuguu.....	1	1	hsa
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.....caacggaaucccaaaagcagcugAu.....	62	1	hsa
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.....caacggaaucccaaaagcaUcuguu.....	2	1	hsa
.....caacggaaucUcaaaagcagcuguu.....	1	1	hsa
.....caacggaaucccaaaaAcagcuguu.....	1	1	hsa
.....caacggaaucccaaaagcagcuguuA.....	9	1	hsa
.....caacggaaucccaaaagcagcuguAg.....	8	1	hsa

hsa-miR-191-3p

hsa-miR-191-5p

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.....caacggaaucccaaaagcagcuguuU.....	19	1	hsa
.....caacggaaucccaaaagcagcuguAgu.....	1	1	hsa
.....caacggaaucccaaaagcagcuguuUu.....	1	1	hsa
.....caacggaaucccaaaagcagcuguugu.....	1	0	hsa
.....caacggaaucccaaaagcagcuguuAu.....	1	1	hsa
.....caacggaaucccaaaagcagcuguCgu.....	1	1	hsa
.....aacggaaucccaaaagca.....	1	0	hsa
.....aacggaaucccaaaagcag.....	3	0	hsa
.....aacggaaucUcaaaagcagc.....	1	1	hsa
.....aacggaaucccaaaagcagc.....	7	0	hsa
.....aCcggaaucccaaaagcagc.....	1	1	hsa
.....aacgUaaucccaaaagcagcu.....	1	1	hsa
.....aacggaaucccaaaagcagcu.....	67	0	hsa
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.....aacggGaucccaaaagcagcug.....	1	1	hsa
.....aacggaaucccaaaagcCgcug.....	1	1	hsa
.....aaAggaaucccaaaagcagcug.....	3	1	hsa
.....aacggaaucccaaaagcagcuU.....	7	1	hsa
.....Cacggaaucccaaaagcagcug.....	1	1	hsa
.....aacUgaaucccaaaagcagcug.....	8	1	hsa
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.....Gacggaaucccaaaagcagcug.....	1	1	hsa
.....aacgUaaucccaaaagcagcug.....	2	1	hsa
.....aacggaaucGcaaaagcagcug.....	1	1	hsa
.....aacggaauccUaaaagcagcug.....	6	1	hsa
.....aacggaauUccaaaagcagcug.....	2	1	hsa
.....aacggaaucccaaaagcGgcug.....	1	1	hsa
.....aacggaaucccaaaagcagcCg.....	1	1	hsa
.....aacggaaucccUaaagcagcug.....	2	1	hsa
.....aaUggaaucccaaaagcagcug.....	3	1	hsa
.....aacggaCucccaaaagcagcug.....	4	1	hsa
.....aacggaaucAcaaaagcagcug.....	1	1	hsa
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.....aacggaaucccaaaagcagUug.....	5	1	hsa
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.....aacggaaucccaCaagcagcug.....	3	1	hsa
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.....aacggaaucccaaaUgcagcug.....	1	1	hsa
.....aCcggaaucccaaaagcagcug.....	7	1	hsa
.....aacAgaaucccaaaagcagcug.....	1	1	hsa
.....aacggaauccAaaaagcagcug.....	1	1	hsa
.....aacggaaucccaaaagcagGug.....	1	1	hsa
.....aacggaaucccaaaagcagcuC.....	1	1	hsa
.....aacggaaucccaaaagcaCcug.....	1	1	hsa
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.....aacggaaGcccaaaagcagcugu.....	1	1	hsa
.....aacggCaucccaaaagcagcugu.....	1	1	hsa
.....aaUggaaucccaaaagcagcugu.....	2	1	hsa
.....aacggaaucUcaaaagcagcugu.....	4	1	hsa
.....aacggaaucccaaaagcagcugG.....	1	1	hsa
.....aacggaaucccaaaagcagcugC.....	2	1	hsa
.....aacggaaAcccaaaagcagcugu.....	1	1	hsa
.....aacggaaucccaaaagcagcugA.....	265	1	hsa
.....aacggaaucccaGaagcagcugu.....	1	1	hsa
.....aacUgaaucccaaaagcagcugu.....	3	1	hsa
.....aacggaaucAcaaaagcagcugu.....	1	1	hsa
.....aacggaGucccaaaagcagcugu.....	1	1	hsa
.....aacggaaucccaaaagcagcuAu.....	1	1	hsa
.....aacggaaucccaaaagcagcuUu.....	1	1	hsa

hsa-miR-191-3p

hsa-miR-191-5p

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.....aacggaauAccaaaagcagcugu.....	1	1	hsa
.....Cacggaaucccaaaagcagcugu.....	3	1	hsa
.....aacggaCucccaaaagcagcugu.....	2	1	hsa
.....aacggaaucccaaaCgcagcugu.....	3	1	hsa
.....aacggaaucccaaaagcagcugu.....	609	0	hsa
.....aacggaaucccaaaaUcagcugu.....	1	1	hsa
.....aacggaaucccaaaagcagcCgu.....	1	1	hsa
.....aacAgaaucccaaaagcagcugu.....	3	1	hsa
.....aacggaaucccaaaagUagcugu.....	2	1	hsa
.....aacggaaucccaaaagcagAugu.....	1	1	hsa
.....aacCgaaucccaaaagcagcugu.....	1	1	hsa
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