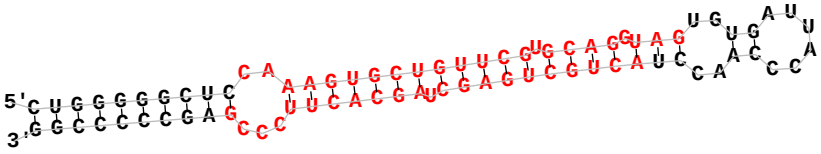
miRBase precursor	: hsa-mir-93
Total read count	: 3821
hsa-miR-93-5p read count	: 3773
hsa-miR-93-3p read count	: 47
remaining reads	: 48





hsa-miR-93-5p

hsa-miR-93-3p

```
5'- cuggggcuccaaagugcuguucgugcagguagugugauuacccaaccuacugcugagcuagcacuucccgagccccgg  -3'    exp
    ((((((((((..((((((((((.((((.(((..((......)).)))))))))))).)))))))...))))))))))   reads      mm         sample
    .........ccaaagugcuguucgugcagguag.....................................................   1          0          hsa
    .........Acaaagugcuguucgugcagguag.....................................................   1          1          hsa
    .........caaagugcuguucgugca...........................................................   5          0          hsa
    .........caaagugcuguucgugcag..........................................................   6          0          hsa
    .........caaagugcuguucgugcagg.........................................................   18         0          hsa
    .........cCaagugcuguucgugcagg.........................................................   1          1          hsa
    .........caaagugcuguucgugcagUu........................................................   1          1          hsa
    .........caaagugcuguucguUcaggu........................................................   1          1          hsa
    .........caaagugcuguucgugAaggu........................................................   1          1          hsa
    .........caaagugcuguucgugcaggu........................................................   41         0          hsa
    .........caaagugcuguucgugcaCgua.......................................................   1          1          hsa
    .........caaCgugcuguucgugcaggua.......................................................   1          1          hsa
    .........caaagugcugAucgugcaggua.......................................................   1          1          hsa
    .........caaagugcuguCcgugcaggua.......................................................   1          1          hsa
    .........caaagugcuguucgugcaggua.......................................................   162        0          hsa
    .........cCaagugcuguucgugcaggua.......................................................   2          1          hsa
    .........caaagugcuguucgugcagUua.......................................................   2          1          hsa
    .........caaagugcuguucgugcCggua.......................................................   1          1          hsa
    .........caaagugcuguucgugcaggCa.......................................................   1          1          hsa
    .........caaagugcuguucgugcagAua.......................................................   6          1          hsa
    .........caaagugcugGucgugcaggua.......................................................   1          1          hsa
    .........caaagugcuguucgugcagguC.......................................................   2          1          hsa
    .........caaagugcuguuUgugcaggua.......................................................   2          1          hsa
    .........caaagugcuUuucgugcaggua.......................................................   2          1          hsa
    .........Gaaagugcuguucgugcaggua.......................................................   1          1          hsa
    .........caaagugcuguucgugcaUgua.......................................................   2          1          hsa
    .........caaagGgcuguucgugcagguag......................................................   1          1          hsa
    .........caaagugcuguAcgugcagguag......................................................   1          1          hsa
    .........caaagugcuguucgCgcagguag......................................................   4          1          hsa
    .........caaagugcuguucguCcagguag......................................................   3          1          hsa
    .........caaagugcuguucCugcagguag......................................................   1          1          hsa
    .........caaagugcuguucgugcCgguag......................................................   1          1          hsa
    .........caaagugcuguucgugcagUuag......................................................   24         1          hsa
    .........caaagugcuguucgugGagguag......................................................   1          1          hsa
```

hsa-miR-93-5p

hsa-miR-93-3p

cuggggcuccaaagugcuguucgugcagguagugugauuacccaaccuacugcugagcuagcacuucccgagccccgg

.....caaGgugcuguucgugcagguag.....	4	1	hsa
.....caaagugcuguucgugAagguag.....	1	1	hsa
.....caaagugAuguucgugcagguag.....	1	1	hsa
.....Aaaagugcuguucgugcagguag.....	2	1	hsa
.....caaagugcuguucgugcagCuag.....	2	1	hsa
.....caaagugcuCuucgugcagguag.....	5	1	hsa
.....caaUgugcuguucgugcagguag.....	1	1	hsa
.....caaagugcuAuucgugcagguag.....	13	1	hsa
.....Uaaagugcuguucgugcagguag.....	3	1	hsa
.....caaagugcuguucgugcagguUg.....	3	1	hsa
.....caaagugcugGucgugcagguag.....	2	1	hsa
.....caGagugcuguucgugcagguag.....	1	1	hsa
.....caaagugcugCucgugcagguag.....	4	1	hsa
.....caaagugcuguucgugcagguaU.....	40	1	hsa
.....caaCgugcuguucgugcagguag.....	15	1	hsa
.....cCaagugcuguucgugcagguag.....	13	1	hsa
.....caaagugcuguucgugUagguag.....	5	1	hsa
.....cUaagugcuguucgugcagguag.....	2	1	hsa
.....caaagugcuguucUugcagguag.....	15	1	hsa
.....caaaguAcuguucgugcagguag.....	2	1	hsa
.....caaagugcCguucgugcagguag.....	1	1	hsa
.....caCagugcuguucgugcagguag.....	9	1	hsa
.....caaagugcuguucgugcagguaC.....	2	1	hsa
.....caaagugcuguucgugcagguag.....	2549	0	hsa
.....caaagugcuguucgugcaggCag.....	4	1	hsa
.....caaagugGuguucgugcagguag.....	1	1	hsa
.....caaaAugcuguucgugcagguag.....	2	1	hsa
.....caaagugcuguucguAcagguag.....	1	1	hsa
.....caaagugcugAucgugcagguag.....	2	1	hsa
.....caaagAgcuguucgugcagguag.....	6	1	hsa
.....Gaaagugcuguucgugcagguag.....	1	1	hsa
.....caaagugcuguucgGgcagguag.....	4	1	hsa
.....caaagugcuguGcgugcagguag.....	3	1	hsa
.....caaagugcuguucgugcaggGag.....	5	1	hsa
.....caaagugcuguCcgugcagguag.....	2	1	hsa
.....caaagugcuguucgugcaAguag.....	2	1	hsa
.....caaagugcuguucgugcGgguag.....	2	1	hsa
.....caaagugcuguucAugcagguag.....	3	1	hsa
.....caaagugcuguucgugcaUguag.....	14	1	hsa
.....caaaCugcuguucgugcagguag.....	1	1	hsa
.....caaagugcuguuUgugcagguag.....	1	1	hsa
.....caaagugcuguucgugcagguCg.....	12	1	hsa
.....caaaUugcuguucgugcagguag.....	2	1	hsa
.....caaagugcuguucguUcagguag.....	8	1	hsa
.....caaagugcuguucgAgcagguag.....	1	1	hsa
.....caaagugcuguucgugcagguaA.....	43	1	hsa
.....caaagugcAguucgugcagguag.....	1	1	hsa
.....caaagugcuguucgugcaggAag.....	11	1	hsa
.....caaaguCcuguucgugcagguag.....	3	1	hsa
.....caaagugcuUuucgugcagguag.....	14	1	hsa
.....caaagCgcuguucgugcagguag.....	4	1	hsa
.....caaagugcuguucgugcaCguag.....	3	1	hsa
.....caaagugUuguucgugcagguag.....	4	1	hsa
.....caaaguUcuguucgugcagguag.....	3	1	hsa
.....cGaagugcuguucgugcagguag.....	1	1	hsa
.....caaagugcuguucgugcaUguagu.....	1	1	hsa
.....caaagugcuUuucgugcagguagu.....	2	1	hsa
.....caaUgugcuguucgugcagguagu.....	1	1	hsa
.....caaagugcuguucgugcagguagA.....	403	1	hsa
.....caaagugcuguucgugcagguaUu.....	1	1	hsa
.....caaagugcuguucgugcagguagu.....	129	0	hsa
.....caaagugcuguucgugcagguaCu.....	1	1	hsa
.....caaagugcuguucCugcagguagu.....	1	1	hsa
.....caaagugcugCucgugcagguagu.....	1	1	hsa
.....caaagugcuguucgugcaggGagu.....	1	1	hsa
.....caaagugcuCuucgugcagguagu.....	2	1	hsa
.....caaagugcuguucgugcagguagC.....	5	1	hsa
.....caaaguUcuguucgugcagguagu.....	1	1	hsa
.....caaagugcuguucgugcagguagG.....	3	1	hsa

hsa-miR-93-5p

hsa-miR-93-3p

cuggggcucCaaagugcuguucgugcagguagugugauuacccaaccuacugcugagcuagcacuucccgagccccgg

.....caaagugcuAuucgugcagguagu.....	1	1	hsa
.....caaagugcuguucUugcagguagu.....	2	1	hsa
.....caaagugcuguucgugcagguCgu.....	1	1	hsa
.....caaagugcuguucgugcagguaguU.....	1	1	hsa
.....caaagugcuguucgugcagguagAg.....	4	1	hsa
.....caaagugcuguucgugcagguaguA.....	4	1	hsa
.....caaagugcuguucgugcagguagugA.....	2	1	hsa
.....caaagugcuguucgugcagguagugu.....	1	0	hsa
.....aCagugcuguucgugcaggua.....	1	1	hsa
.....aaagugcuguucgugcagguU.....	1	1	hsa
.....aaagugcuguucgugcaggua.....	2	0	hsa
.....aaagugcuguuUgugcagguag.....	1	1	hsa
.....aaagugcuguucgugcagguaA.....	2	1	hsa
.....aaagugcuguucgugcagguag.....	21	0	hsa
.....aaagugcuguucgugcagguagu.....	2	0	hsa
.....aaagugcuguucgugcagguagA.....	7	1	hsa
.....aaagugcuguucgugcagguagugu.....	1	0	hsa
.....aagugcuguucgugcagguag.....	1	0	hsa
.....gugcuguucgugcagguag.....	4	0	hsa
.....gugcuguucgugcagguagA.....	2	1	hsa
.....gugcuguucgugcagguagu.....	1	0	hsa
.....ccuacugcugagcuagcacuucc.....	1	0	hsa
.....cuacugcugagcuagcac.....	1	0	hsa
.....cuacugcugagcuagcacuucc.....	1	0	hsa
.....uacugcugagcuagcacuucU.....	1	1	hsa
.....uacugcugagcuagcacuucc.....	2	0	hsa
.....uacugcugagcuagcacuucccga.....	1	0	hsa
.....uacugcugagcuagcacuucccgaU.....	1	1	hsa
.....acugcugagcuagcacuuccc.....	1	0	hsa
.....acugcugagcuagcacuucccA.....	1	1	hsa
.....acugcugagcuagcacuucUcg.....	1	1	hsa
.....acugcugagcuagcacuucccg.....	10	0	hsa
.....acugcugCgcuagcacuucccga.....	1	1	hsa
.....acugcugagcuagcacuucccgU.....	8	1	hsa
.....acugcugagcuagcacuucccga.....	12	0	hsa
.....acugcugagcuagcacuucUcga.....	1	1	hsa
.....acugcugagcuagcacuucccgG.....	2	1	hsa
.....acugcugagcuagcacuucccgag.....	1	0	hsa
.....acugcugagcuagcacuucccgaU.....	1	1	hsa
.....cugcugagcuagcacuucccga.....	1	0	hsa