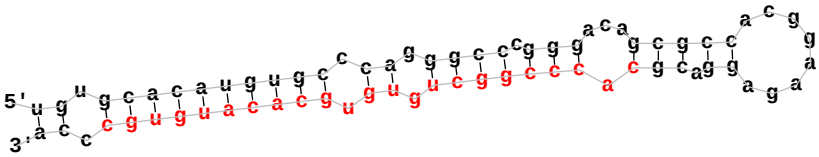
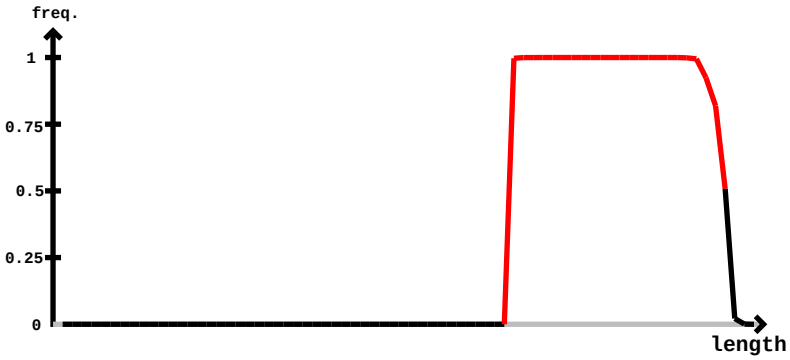
miRBase precursor : hsa-mir-941-4
Total read count : 14188
hsa-miR-941 read count : 14188
remaining reads : 0





hsa-miR-941

```
5'- ugugcacaugugcccagggcccgggacagcgccacggaagaggacgcacccggcugugugcacaugugccca  -3'    exp
    ((.((((((((((.((.((((.(((...(((((........)).))).)))))))).)).))))))))))).))   reads      mm         sample
    ..............................................cacccggcugugugcaca........   1          0          Z3_
    ..............................................cacccggcugugugcacau.......   1          0          Z3_
    ..............................................cacccggcugugugcacaC.......   1          1          Z3_
    ..............................................caUccggcugugugcacau.......   1          1          Z3_
    ..............................................Uacccggcugugugcacau.......   1          1          Z3_
    ..............................................cacccggcugugGgcacaug......   1          1          Z3_
    ..............................................cGcccggcugugugcacaug......   2          1          Z3_
    ..............................................cacccggcugugugcacauA......   4          1          Z3_
    ..............................................cacccggcugugCgcacaug......   1          1          Z3_
    ..............................................cacccggcugugugcacaug......   67         0          Z3_
    ..............................................cacccggcugCgugcacaug......   1          1          Z3_
    ..............................................cacccggcugugugcacaugG.....   2          1          Z3_
    ..............................................cacccggcuAugugcacaugu.....   2          1          Z3_
    ..............................................cUcccggcugugugcacaugu.....   1          1          Z3_
    ..............................................cacccggcugugugGacaugu.....   5          1          Z3_
    ..............................................cacccggcugugugcacaugA.....   1          1          Z3_
    ..............................................cacccggcugugugcacaugu.....   106        0          Z3_
    ..............................................cacccggcAgugugcacaugu.....   1          1          Z3_
    ..............................................cacccggcugugugcacaugC.....   10         1          Z3_
    ..............................................cGcccggcugugugcacaugu.....   1          1          Z3_
    ..............................................cacccggcugugugcGcaugu.....   1          1          Z3_
    ..............................................cacccAgcugugugcacaugug....   1          1          Z3_
    ..............................................cacccggcugugugcacauguU....   3          1          Z3_
    ..............................................cacccggcugugugcUcaugug....   1          1          Z3_
    ..............................................cacccggcCgugugcacaugug....   1          1          Z3_
    ..............................................cacccggcugugugcacaugCg....   3          1          Z3_
    ..............................................cacccggcugugugcacauAug....   1          1          Z3_
    ..............................................cacccggcugugugcacauguC....   1          1          Z3_
    ..............................................cacccggcugugugcacaugug....   402        0          Z3_
    ..............................................Uacccggcugugugcacaugug....   1          1          Z3_
    ..............................................cacccggcugugugcacGugug....   1          1          Z3_
    ..............................................cacccggcugugugGacaugug....   20         1          Z3_
    ..............................................cacccggcugugugcacUugug....   1          1          Z3_
    ..............................................cacccggcugugAgcacaugug....   1          1          Z3_
    ..............................................cacccggUugugugcacaugug....   2          1          Z3_
```

hsa-miR-941

ugugcacaugugcccagggcccgggacagcgccacggaagaggacgcacccggcugugugcacaugugccca

.....cacccggcugugugcacauguA....	23	1	Z3_
.....cacccggcugCgugcacaugug....	2	1	Z3_
.....cacccggcugugugcacaAgug....	2	1	Z3_
.....cacccggcugugugcGcaugug....	3	1	Z3_
.....cGcccggcugugugcacaugug....	1	1	Z3_
.....cacccggcuguguUcacaugug....	1	1	Z3_
.....Aacccggcugugugcacaugug....	1	1	Z3_
.....cacccggcugugCgcacaugug....	2	1	Z3_
.....cacccggcugugugcacaCgug....	1	1	Z3_
.....cacccggcugugugcacauAugc...	1	1	Z3_
.....cacccggcugugugcacaugugU...	245	1	Z3_
.....cacccggcugugugcUcaugugc...	1	1	Z3_
.....cacccggcAgugugcacaugugc...	2	1	Z3_
.....caccUggcugugugcacaugugc...	1	1	Z3_
.....cacccggcugugugcaUaugugc...	1	1	Z3_
.....cacccggcCgugugcacaugugc...	5	1	Z3_
.....cacccggcugugugcacaCgugc...	1	1	Z3_
.....cacccggcugugugcacaugugA...	106	1	Z3_
.....caUccggcugugugcacaugugc...	1	1	Z3_
.....cacccggcugugCgcacaugugc...	8	1	Z3_
.....cacccggcugugugcacauguAc...	2	1	Z3_
.....cacccggcugugugcGcaugugc...	2	1	Z3_
.....cacccggcugugugcacGugugc...	3	1	Z3_
.....cacccggcugCgugcacaugugc...	7	1	Z3_
.....cacccggcugugugGacaugugc...	7	1	Z3_
.....cGcccggcugugugcacaugugc...	1	1	Z3_
.....cacUcggcugugugcacaugugc...	1	1	Z3_
.....Uacccggcugugugcacaugugc...	4	1	Z3_
.....cacccggcuAugugcacaugugc...	3	1	Z3_
.....cacccggcugugugUacaugugc...	1	1	Z3_
.....Aacccggcugugugcacaugugc...	2	1	Z3_
.....cacccggcugugugcacaugugc...	477	0	Z3_
.....cacccggcugugugcacaugAgc...	1	1	Z3_
.....cUcccggcugugugcacaugugc...	3	1	Z3_
.....cacccgCcugugugcacaugugc...	1	1	Z3_
.....cacccggcuguAugcacaugugc...	1	1	Z3_
.....cacccggcugugugcacaugugG...	3	1	Z3_
.....cacccggcugugugcacUugugc...	2	1	Z3_
.....Gacccggcugugugcacaugugc...	1	1	Z3_
.....cacccggcugugugcacaugugcU..	28	1	Z3_
.....cacccggcugugugGacaugugcc..	2	1	Z3_
.....cacccggcugugugcacaugugcA..	2	1	Z3_
.....cacccggcugugugcacaugugcc..	7	0	Z3_
.....cacccggcugugugcacaugugcAc.	2	1	Z3_
.....acccggcugugugcacaugug....	1	0	Z3_
.....acccggcugugugcacaugugc...	2	0	Z3_
.....acccggcCgugugcacaugugc...	1	1	Z3_
.....Ccccggcugugugcacaugugc...	1	1	Z3_

.....gcacccggcugugugGacaugug....	1	1	Z1_
.....caccGggcugugugcaca........	2	1	Z1_
.....cacccggcugugugcaca........	5	0	Z1_
.....cacccggcugugugcacau.......	4	0	Z1_
.....cacccggcugugugcacaC.......	1	1	Z1_
.....cacccggcugugugcacauA......	13	1	Z1_
.....cacccggcCgugugcacaug......	3	1	Z1_
.....cacccggcugugugcacaug......	300	0	Z1_
.....cacUcggcugugugcacaug......	1	1	Z1_
.....Uacccggcugugugcacaug......	3	1	Z1_
.....cacccggcuguCugcacaug......	1	1	Z1_
.....cacccggcugugCgcacaug......	1	1	Z1_
.....cacccggcugugugcacGug......	1	1	Z1_
.....cacccggcugugugGacaug......	1	1	Z1_
.....cacccggcuguAugcacaug......	1	1	Z1_
.....cUcccggcugugugcacaug......	1	1	Z1_
.....cacccggcugCgugcacaug......	1	1	Z1_
.....cacccggcugugAgcacaug......	1	1	Z1_
.....cacccggcugugugcacauU......	3	1	Z1_
.....cacAcggcugugugcacaugu.....	1	1	Z1_
.....cacccggcugugugcacaugC.....	44	1	Z1_

hsa-miR-941

ugugcacaugugcccagggcccgggacagcgccacggaagaggacgcacccggcugugugcacaugugccca

.....cacccggcugugugcGcaugu.....	2	1	Z1_
.....cacUcggcugugugcacaugu.....	1	1	Z1_
.....cacccggcugCgugcacaugu.....	3	1	Z1_
.....cacccAgcugugugcacaugu.....	1	1	Z1_
.....cacccggcugugugcacaugG.....	2	1	Z1_
.....Aacccggcugugugcacaugu.....	1	1	Z1_
.....cacccggcugugugcacaugu.....	405	0	Z1_
.....cacccggcugugugcacGugu.....	1	1	Z1_
.....caUccggcugugugcacaugu.....	1	1	Z1_
.....cacccggcugugCgcacaugu.....	3	1	Z1_
.....cacccggcugugugcacaugA.....	7	1	Z1_
.....cGcccggcugugugcacaugu.....	4	1	Z1_
.....cacccggcugugugcacUugu.....	2	1	Z1_
.....cacccggcCgugugcacaugu.....	4	1	Z1_
.....cacccggcugAgugcacaugu.....	1	1	Z1_
.....cacccggcuguguAcacaugu.....	1	1	Z1_
.....cacccggcuguguCcacaugu.....	1	1	Z1_
.....cacccggAugugugcacaugu.....	1	1	Z1_
.....cacccggcugugGgcacaugu.....	1	1	Z1_
.....cacccggcugugugAacaugug....	1	1	Z1_
.....cacccggcugugCgcacaugug....	10	1	Z1_
.....cacccggcugugugcacaCgug....	5	1	Z1_
.....cacccggcugugugcacauguU....	6	1	Z1_
.....cUcccggcugugugcacaugug....	1	1	Z1_
.....cacccggcugugugGacaugug....	17	1	Z1_
.....cacccggcuguguAcacaugug....	3	1	Z1_
.....cacccggcugugugcacauAug....	3	1	Z1_
.....Uacccggcugugugcacaugug....	4	1	Z1_
.....cacccggcugugugcacaugug....	1221	0	Z1_
.....cacccggcCgugugcacaugug....	5	1	Z1_
.....cacccggcuguAugcacaugug....	3	1	Z1_
.....cCcccggcugugugcacaugug....	3	1	Z1_
.....cacccggcAgugugcacaugug....	2	1	Z1_
.....cacccggcugugugcacGugug....	2	1	Z1_
.....cacccggcugugugcacUugug....	1	1	Z1_
.....cacccggcugugugcGcaugug....	4	1	Z1_
.....cacccggcugugugcaUaugug....	1	1	Z1_
.....cacccggcugugugcUcaugug....	3	1	Z1_
.....cacccggcugAgugcacaugug....	1	1	Z1_
.....cacccggcugugugcacaugGg....	1	1	Z1_
.....cacccggcugugugcacauguC....	3	1	Z1_
.....cacccggcugugugUacaugug....	2	1	Z1_
.....cacccggcugugugcacaugAg....	1	1	Z1_
.....cacccgAcugugugcacaugug....	1	1	Z1_
.....cacccggcuAugugcacaugug....	2	1	Z1_
.....cacAcggcugugugcacaugug....	3	1	Z1_
.....cacccggcugugugcacaugCg....	3	1	Z1_
.....cacccggcugCgugcacaugug....	5	1	Z1_
.....cacccggcugugugcacauguA....	50	1	Z1_
.....cacccggcugugugcacaGgug....	3	1	Z1_
.....cGcccggcugugugcacaugug....	9	1	Z1_
.....caUccggcugugugcacaugug....	1	1	Z1_
.....caGccggcugugugcacaugug....	1	1	Z1_
.....cacccggcugugugcacCugug....	2	1	Z1_
.....cacUcggcugugugcacaugug....	2	1	Z1_
.....cacccggcugCgugcacaugugc...	10	1	Z1_
.....cacccggcGgugugcacaugugc...	1	1	Z1_
.....cacccggcugugugcacGugugc...	9	1	Z1_
.....cacccggcugugugcGcaugugc...	11	1	Z1_
.....cacccggcugugugGacaugugc...	7	1	Z1_
.....cacUcggcugugugcacaugugc...	4	1	Z1_
.....cacccggcugugGgcacaugugc...	1	1	Z1_
.....cacccggcuguguAcacaugugc...	1	1	Z1_
.....cacccggcugugAgcacaugugc...	3	1	Z1_
.....Gacccggcugugugcacaugugc...	1	1	Z1_
.....cacccggcugugugcacaugugU...	570	1	Z1_
.....cUcccggcugugugcacaugugc...	1	1	Z1_
.....cCcccggcugugugcacaugugc...	1	1	Z1_
.....cacccggcugugugcacUugugc...	2	1	Z1_
.....Aacccggcugugugcacaugugc...	4	1	Z1_

hsa-miR-941

ugugcacaugugcccagggcccgggacagcgccacggaagaggacgcacccggcugugugcacaugugccca

.....cGcccggcugugugcacaugugc...	6	1	Z1_
.....cacccggcugugugUacaugugc...	2	1	Z1_
.....cacccggcugugugAacaugugc...	4	1	Z1_
.....cacccggcugugugCcaugugc...	2	1	Z1_
.....caccUggcugugugcacaugugc...	4	1	Z1_
.....cacccggcugugugcacaugugA...	216	1	Z1_
.....cacccggcugugugcacaCgugc...	1	1	Z1_
.....Uacccggcugugugcacaugugc...	7	1	Z1_
.....cacccggcugugugcacaugugc...	1550	0	Z1_
.....cacccggcugugugcacaugCgc...	13	1	Z1_
.....cacccggcuguCugcacaugugc...	2	1	Z1_
.....cacccgCcugugugcacaugugc...	2	1	Z1_
.....cacccgAcugugugcacaugugc...	1	1	Z1_
.....cacccggcugugugcacaugAgc...	2	1	Z1_
.....cacccggcugAgugcacaugugc...	1	1	Z1_
.....cacccggcugugugcaUaugugc...	2	1	Z1_
.....cacccggcugugugcacaugugG...	6	1	Z1_
.....cacccggcugugugcacauguAc...	2	1	Z1_
.....cacccggcugugCgcacaugugc...	6	1	Z1_
.....cacccggcugugugcUcaugugc...	3	1	Z1_
.....caccAggcugugugcacaugugc...	1	1	Z1_
.....caUccggcugugugcacaugugc...	2	1	Z1_
.....cacccggcugugugcacaugugcA..	3	1	Z1_
.....cacccggcugugugcacaugugAc..	3	1	Z1_
.....cacccggcugugugcacaugugUc..	1	1	Z1_
.....cacccggcugugugcacaugugcc..	9	0	Z1_
.....cacccggcugugugcGcaugugcc..	1	1	Z1_
.....cacccggcugugugcacaugugcG..	1	1	Z1_
.....cacccggcugugugGacaugugcc..	2	1	Z1_
.....cacccggcugCgugcacaugugcc..	2	1	Z1_
.....cacccggcugugugcacaugugcU..	91	1	Z1_
.....cacccggcugugugcacaugugccU.	1	1	Z1_
.....cacccggcugugugcacaugugcAc.	4	1	Z1_
.....acccggcugugugcacaugu.....	1	0	Z1_
.....acccggcugugugcacaugug....	3	0	Z1_
.....acccggcugugugcacaugugc...	7	0	Z1_

.....cacccggcugugAgcacaug.....	1	1	Z2_
.....cacccggcugugugcacaug.....	40	0	Z2_
.....cacccggcCgugugcacaug.....	1	1	Z2_
.....cacccgAcugugugcacaugu.....	1	1	Z2_
.....cacccggcugugugcacGugu.....	1	1	Z2_
.....cacccggcugugugcacaugG.....	1	1	Z2_
.....cacccggcugugAgcacaugu.....	1	1	Z2_
.....cacccggcugugugcacaugu.....	80	0	Z2_
.....cacccggcugugugUacaugu.....	2	1	Z2_
.....Gacccggcugugugcacaugu.....	2	1	Z2_
.....cacccggcugugugcacaugC.....	7	1	Z2_
.....cacccggAugugugcacaugug....	1	1	Z2_
.....cacccggcugugugcacaugug....	277	0	Z2_
.....cacccggcugugugcacauguA....	14	1	Z2_
.....cacccggcugugugCcaugug....	1	1	Z2_
.....Uacccggcugugugcacaugug....	1	1	Z2_
.....cUcccggcugugugcacaugug....	1	1	Z2_
.....cacccggcugugugUacaugug....	1	1	Z2_
.....cacccggUugugugcacaugug....	1	1	Z2_
.....cacccggcuguguAcacaugug....	1	1	Z2_
.....cacccggcugugugcacaugCg....	1	1	Z2_
.....cacccggcugugugcacaugAg....	2	1	Z2_
.....caccUgcugugugcacaugug....	1	1	Z2_
.....cacccggcugugugcacauguU....	3	1	Z2_
.....caccAgcugugugcacaugug....	1	1	Z2_
.....cacccggcugugCgcacaugug....	1	1	Z2_
.....cacccggcugugGgcacaugug....	1	1	Z2_
.....caccUggcugugugcacaugug....	1	1	Z2_
.....cacccggcugugugcacCugug....	1	1	Z2_
.....cacccggcugugugGacaugug....	11	1	Z2_
.....cacccggcCgugugcacaugug....	1	1	Z2_
.....cGcccggcugugugcacaugug....	1	1	Z2_
.....cacccggcuAugugcacaugug....	1	1	Z2_

hsa-miR-941

ugugcacaugugcccagggcccgggacagcgccacggaagaggacgcacccggcugugugcacaugugccca

.....cacccggcugCgugcacaugug....	2	1	Z2_
.....cacccggcugugAgcacaugugc...	1	1	Z2_
.....cacccggcugugugcacaugugc...	324	0	Z2_
.....cacccggcugugugcacaugugU...	153	1	Z2_
.....cacccggcugugugcacGugugc...	3	1	Z2_
.....cacccgAcugugugcacaugugc...	3	1	Z2_
.....cacccggcugGgugcacaugugc...	1	1	Z2_
.....cacccggcugAgugcacaugugc...	3	1	Z2_
.....cacccggcugugugcacaugAgc...	2	1	Z2_
.....cacccggcugCgugcacaugugc...	2	1	Z2_
.....cacccggcugugugcacauguUc...	1	1	Z2_
.....cacccggcuguguCcacaugugc...	1	1	Z2_
.....cacccggcugugugcacaugugA...	46	1	Z2_
.....cacccggcugugugcacaugugG...	1	1	Z2_
.....cacccggcugugugcacauguAc...	1	1	Z2_
.....Uacccggcugugugcacaugugc...	3	1	Z2_
.....cacccAgcugugugcacaugugc...	1	1	Z2_
.....cacAcggcugugugcacaugugc...	1	1	Z2_
.....cacccggcugugugGacaugugc...	14	1	Z2_
.....cacccggcuguguAcacaugugc...	1	1	Z2_
.....cacccggcugugugcGcaugugc...	3	1	Z2_
.....cacccggcCgugugcacaugugc...	2	1	Z2_
.....cacUcggcugugugcacaugugc...	1	1	Z2_
.....cacccggcugugugcacaugCgc...	1	1	Z2_
.....cacccggcugugugcacaugugcc..	3	0	Z2_
.....cacccggcugugugcacaugugcG..	2	1	Z2_
.....cacccggcugugugcacaugugAc..	4	1	Z2_
.....cacccggcugugugcacaugugcU..	17	1	Z2_
.....cacccggcugugugcacaugugcAc.	1	1	Z2_
.....acccggcugugugcacaug......	1	0	Z2_

.....cacccggcugugugcaca........	8	0	GDM
.....cacGcggcugugugcaca........	1	1	GDM
.....cacccggcugugugcaca........	8	0	GDM
.....cacccggcugugugcaca........	8	0	GDM
.....cacccggcugugugcacau.......	44	0	GDM
.....cacccggcugugugcacau.......	44	0	GDM
.....cacccggcugugugcacau.......	44	0	GDM
.....cacccggcugugugcacauU......	2	1	GDM
.....cacccggcugugugcGcaug......	3	1	GDM
.....caccUggcugugugcacaug......	1	1	GDM
.....cacccggcugugugcaUaug......	2	1	GDM
.....caccAggcugugugcacaug......	2	1	GDM
.....cacccggcugugugcacGug......	3	1	GDM
.....cacccggcugugugcacauA......	24	1	GDM
.....cacccggcugugugcacauA......	24	1	GDM
.....cacccggcugugugGacaug......	4	1	GDM
.....cacccggcuguguCcacaug......	4	1	GDM
.....cacccggcuAugugcacaug......	1	1	GDM
.....cacccggcugugugcacaug......	474	0	GDM
.....cacccggcCgugugcacaug......	2	1	GDM
.....cacccggcugugCgcacaug......	5	1	GDM
.....cacccggcuguguAcacaug......	1	1	GDM
.....cacccggcugugCgcacaug......	5	1	GDM
.....cacccggcugugugcacaCg......	3	1	GDM
.....cacccggcugugugcacaCg......	3	1	GDM
.....cacccggcugugugcaUaug......	2	1	GDM
.....cacccggcugugugcacGug......	3	1	GDM
.....cacccggcugugugcacauA......	24	1	GDM
.....cacccggcugugugcacaug......	474	0	GDM
.....cacccggcugCgugcacaug......	5	1	GDM
.....cacccggcugugugcGcaug......	3	1	GDM
.....Aacccggcugugugcacaug......	2	1	GDM
.....cacccggcugCgugcacaug......	5	1	GDM
.....Uacccggcugugugcacaug......	1	1	GDM
.....cacccggcugugCgcacaug......	5	1	GDM
.....cacccAgcugugugcacaug......	2	1	GDM
.....cacccggcuguguCcacaug......	4	1	GDM
.....cacccggcugugugcacaGg......	10	1	GDM
.....Aacccggcugugugcacaug......	2	1	GDM

hsa-miR-941

ugugcacaugugcccagggcccgggacagcgccacggaagaggacgcacccggcugugugcacaugugccca

.....caUccggcugugugcacaug.....	1	1	GDM
.....cacccggcugugugcacaug.....	474	0	GDM
.....cacccggcugugugcacauC.....	1	1	GDM
.....cacccggcuAugugcacaugu.....	2	1	GDM
.....cacccggcuguguCcacaugu.....	1	1	GDM
.....cacccggcugugugcUcaugu.....	3	1	GDM
.....cacccggcugugugcCcaugu.....	1	1	GDM
.....cacccggcugugugcacaugu.....	643	0	GDM
.....cacccggcugAgugcacaugu.....	1	1	GDM
.....cacccggcugugugcaUaugu.....	2	1	GDM
.....cacccggcugugugcacauUu.....	2	1	GDM
.....cacccggcugugugcacaugA.....	28	1	GDM
.....cGcccggcugugugcacaugu.....	3	1	GDM
.....cacccggcugugugcacaugu.....	643	0	GDM
.....cacccggcugugugcacaCgu.....	5	1	GDM
.....cacccggcuguAugcacaugu.....	1	1	GDM
.....cacccggcCgugugcacaugu.....	2	1	GDM
.....cUcccggcugugugcacaugu.....	1	1	GDM
.....cacccggcugugCgcacaugu.....	3	1	GDM
.....cacccggcugugugcacaugC.....	51	1	GDM
.....cacccggcugugugcaUaugu.....	2	1	GDM
.....cacccAgcugugugcacaugu.....	1	1	GDM
.....cacccggcugugugcacaugC.....	51	1	GDM
.....caUccggcugugugcacaugu.....	1	1	GDM
.....cacUcggcugugugcacaugu.....	1	1	GDM
.....cacccggcugugugUacaugu.....	4	1	GDM
.....Aacccggcugugugcacaugu.....	1	1	GDM
.....cGcccggcugugugcacaugu.....	3	1	GDM
.....cacccggcugugugcacGugu.....	1	1	GDM
.....cacccggcugugugcacaugA.....	28	1	GDM
.....cacccggcugugugcacaugG.....	1	1	GDM
.....cacccggcugCgugcacaugu.....	6	1	GDM
.....cacccggcuguguAcacaugu.....	1	1	GDM
.....cacccggcugugugGacaugu.....	7	1	GDM
.....cacccggcugugugcUcaugu.....	3	1	GDM
.....cacccggcugugugcacaugu.....	643	0	GDM
.....cacccggcugCgugcacaugu.....	6	1	GDM
.....cacccggcugugugcGcaugu.....	5	1	GDM
.....cacccggcugCgugcacaugu.....	6	1	GDM
.....cacAcggcugugugcacaugu.....	1	1	GDM
.....cacccggcugugugUacaugu.....	4	1	GDM
.....cacccggcugugugcacaugC.....	51	1	GDM
.....cacccggcugugugcacauAu.....	1	1	GDM
.....cacccggcugugugcacaAgu.....	1	1	GDM
.....cacccggcugAgugcacaugug....	4	1	GDM
.....cacccUgcugugugcacaugug....	3	1	GDM
.....cGcccggcugugugcacaugug....	9	1	GDM
.....cacccggcuguguAcacaugug....	5	1	GDM
.....caccUggcugugugcacaugug....	2	1	GDM
.....cacccggcugugugcacaugAg....	2	1	GDM
.....cacccggcugugugcacaugAg....	2	1	GDM
.....cacccgAcugugugcacaugug....	2	1	GDM
.....cacccggcugCgugcacaugug....	10	1	GDM
.....cUcccggcugugugcacaugug....	8	1	GDM
.....cacccUgcugugugcacaugug....	3	1	GDM
.....cacccggcuguguAcacaugug....	5	1	GDM
.....cacccggcugugugGacaugug....	45	1	GDM
.....cacccggcuCugugcacaugug....	1	1	GDM
.....cacccggcugugugcacaAgug....	1	1	GDM
.....caAccggcugugugcacaugug....	1	1	GDM
.....cacccggcugugugGacaugug....	45	1	GDM
.....Uacccggcugugugcacaugug....	7	1	GDM
.....cacccggcugugugcUcaugug....	1	1	GDM
.....cacccggcugugugcGcaugug....	17	1	GDM
.....cacccggcuguguUcacaugug....	1	1	GDM
.....cacccggcugugugcacaugug....	1826	0	GDM
.....cacccggcugugugcacauguA....	113	1	GDM
.....cacccggcAgugugcacaugug....	3	1	GDM
.....cacccggcugugugcacaugCg....	5	1	GDM
.....cacccggcugugugcacaugug....	1826	0	GDM

hsa-miR-941

ugugcacaugugcccagggcccgggacagcgccacggaagaggacgcacccggcugugugcacaugugccca

.....cacccggcugugugcacaugGg....	71	1	GDM
.....cacccggcugugugcaAaugug....	1	1	GDM
.....cacccggcuAugugcacaugug....	2	1	GDM
.....cacccggcugugugcCcaugug....	1	1	GDM
.....cacccggUugugugcacaugug....	5	1	GDM
.....cacAcggcugugugcacaugug....	3	1	GDM
.....cacccggUugugugcacaugug....	5	1	GDM
.....cacccggcugugugcacaCgug....	9	1	GDM
.....Uacccggcugugugcacaugug....	7	1	GDM
.....cacccCgcugugugcacaugug....	1	1	GDM
.....cacccAgcugugugcacaugug....	4	1	GDM
.....cacccggcuguUugcacaugug....	1	1	GDM
.....cacccggcugugugGacaugug....	45	1	GDM
.....Aacccggcugugugcacaugug....	2	1	GDM
.....cacccggcugugugUacaugug....	7	1	GDM
.....cacccggcugugugcacaugug....	1826	0	GDM
.....cacccggcugugugcacUugug....	1	1	GDM
.....cacccggcugCgugcacaugug....	10	1	GDM
.....cacccggcAgugugcacaugug....	3	1	GDM
.....cacccggcugugCgcacaugug....	7	1	GDM
.....cacccggcuguAugcacaugug....	1	1	GDM
.....caUccggcugugugcacaugug....	5	1	GDM
.....cacccggcugugugcacGugug....	6	1	GDM
.....caUccggcugugugcacaugug....	5	1	GDM
.....cacccggcugugugcGcaugug....	17	1	GDM
.....Uacccggcugugugcacaugug....	7	1	GDM
.....cacccggcugugugcacauAug....	2	1	GDM
.....cacccggcugugugcacaugCg....	5	1	GDM
.....cacccggcugugAgcacaugug....	3	1	GDM
.....cacAcggcugugugcacaugug....	3	1	GDM
.....cacccggcCgugugcacaugug....	5	1	GDM
.....cacccggcugCgugcacaugug....	10	1	GDM
.....cacccggcugugugcacaugGg....	71	1	GDM
.....cacccggcugugugcacauguA....	113	1	GDM
.....cacccggcugugugcacGugug....	6	1	GDM
.....cacccggcGgugugcacaugug....	1	1	GDM
.....cacccggcuguguAcacaugug....	5	1	GDM
.....cacccggcugugugcGcaugug....	17	1	GDM
.....cacccggcugugugcacGugug....	6	1	GDM
.....cacccggcugugugcacauguU....	7	1	GDM
.....cacccggcugugugcacauAug....	2	1	GDM
.....cacccggcugugugcacaCgug....	9	1	GDM
.....cacccggcugugugcacaCgug....	9	1	GDM
.....cacccggcugugugcacauguU....	7	1	GDM
.....cacccggcugugugAacaugug....	1	1	GDM
.....cacccggcugAgugcacaugug....	4	1	GDM
.....cCcccggcugugugcacaugug....	1	1	GDM
.....cacccggcugGgugcacaugug....	2	1	GDM
.....cacccggcugugugUacaugug....	7	1	GDM
.....caccUggcugugugcacaugug....	2	1	GDM
.....cacccggUugugugcacaugug....	5	1	GDM
.....cacccggcugugugcacauguC....	4	1	GDM
.....cacccggcugugCgcacaugug....	7	1	GDM
.....cacccggcugugugcacaugCg....	5	1	GDM
.....cacccggcugugugcacauguA....	113	1	GDM
.....cacccggcuAugugcacaugug....	2	1	GDM
.....Aacccggcugugugcacaugug....	2	1	GDM
.....cacccAgcugugugcacaugug....	4	1	GDM
.....cGcccggcugugugcacaugug....	9	1	GDM
.....cacccgUcugugugcacaugug....	1	1	GDM
.....cacccggcugugugcacauguC....	4	1	GDM
.....cacUcggcugugugcacaugug....	3	1	GDM
.....cacccggcugugugcacaugGg....	71	1	GDM
.....cacccggcCgugugcacaugug....	5	1	GDM
.....cacccggcugugugcacaugugc...	1538	0	GDM
.....cacccggcugCgugcacaugugc...	4	1	GDM
.....cacccggcugugugcacaCgugc...	5	1	GDM
.....cacccggcugugugcacGugugc...	11	1	GDM
.....cacccggcugugugcacaugugG...	11	1	GDM
.....cacccggcuAugugcacaugugc...	2	1	GDM

hsa-miR-941

ugugcacaugugcccagggcccgggacagcgccacggaagaggacgcacccggcugugugcacaugugccca

.....Aacccggcugugugcacaugugc...	4	1	GDM
.....cacccggcugugugGacaugugc...	28	1	GDM
.....cacccggcugugugcacGugugc...	11	1	GDM
.....cacccCgcugugugcacaugugc...	1	1	GDM
.....cacccggcugugugcUcaugugc...	1	1	GDM
.....cUcccggcugugugcacaugugc...	2	1	GDM
.....cGcccggcugugugcacaugugc...	9	1	GDM
.....cacccggcugCgugcacaugugc...	4	1	GDM
.....cacccggcugugCgcacaugugc...	10	1	GDM
.....cacccggUugugugcacaugugc...	2	1	GDM
.....caccGggcugugugcacaugugc...	2	1	GDM
.....cacccggcugugugcGcaugugc...	5	1	GDM
.....cacccggcugugugcacaugugA...	407	1	GDM
.....cacccggcCgugugcacaugugc...	11	1	GDM
.....cacccggcugugugcacaugugU...	842	1	GDM
.....cacccUgcugugugcacaugugc...	1	1	GDM
.....cacccggcugugugUacaugugc...	4	1	GDM
.....Gacccggcugugugcacaugugc...	1	1	GDM
.....cacccggcuguAugcacaugugc...	3	1	GDM
.....cacccggcugugugcacaugugG...	11	1	GDM
.....cacccggcugugugUacaugugc...	4	1	GDM
.....cacccAgcugugugcacaugugc...	2	1	GDM
.....cacccAgcugugugcacaugugc...	2	1	GDM
.....cacccggcugugugcacaugGgc...	1	1	GDM
.....cacccggcugugugcacauUugc...	1	1	GDM
.....caccGggcugugugcacaugugc...	2	1	GDM
.....Aacccggcugugugcacaugugc...	4	1	GDM
.....cacccggcugugGgcacaugugc...	1	1	GDM
.....cacccggcugugugcacaugugc...	1538	0	GDM
.....cacccggcugugCgcacaugugc...	10	1	GDM
.....cacccggcugGgugcacaugugc...	1	1	GDM
.....cacccggcugugugcacaugCgc...	6	1	GDM
.....Uacccggcugugugcacaugugc...	6	1	GDM
.....cacccggcugugugcacaugCgc...	6	1	GDM
.....cGcccggcugugugcacaugugc...	9	1	GDM
.....cacccggcugugugcacCugugc...	1	1	GDM
.....Uacccggcugugugcacaugugc...	6	1	GDM
.....cacccggcugugugGacaugugc...	28	1	GDM
.....cacGcggcugugugcacaugugc...	3	1	GDM
.....cacccggcugugugcacUugugc...	3	1	GDM
.....cacccggcugugugcGcaugugc...	5	1	GDM
.....cacccggcuAugugcacaugugc...	2	1	GDM
.....cacccggcugugugcacaugugU...	842	1	GDM
.....cacccggcuguAugcacaugugc...	3	1	GDM
.....cacccggcugugugcacGugugc...	11	1	GDM
.....cacAcggcugugugcacaugugc...	3	1	GDM
.....cacAcggcugugugcacaugugc...	3	1	GDM
.....cacccggcugugugcaUaugugc...	4	1	GDM
.....cacUcggcugugugcacaugugc...	1	1	GDM
.....cacccggcCgugugcacaugugc...	11	1	GDM
.....cacccggcugugugcacaugugG...	11	1	GDM
.....cacccggcugugugcacaugugA...	407	1	GDM
.....cacccggcugugugcacaugugA...	407	1	GDM
.....cacccggcugugCgcacaugugc...	10	1	GDM
.....cacccggcugugugcacaugugU...	842	1	GDM
.....cacccggcugugugcacauguAc...	3	1	GDM
.....caUccggcugugugcacaugugc...	3	1	GDM
.....cacGcggcugugugcacaugugc...	3	1	GDM
.....cacccggcugugugcacaCgugc...	5	1	GDM
.....cacccggcugugugcGcaugugc...	5	1	GDM
.....caUccggcugugugcacaugugc...	3	1	GDM
.....cGcccggcugugugcacaugugc...	9	1	GDM
.....cacccggcugugugGacaugugc...	28	1	GDM
.....cacccggcugugugcaUaugugc...	4	1	GDM
.....cacccggcugugugcacaugugc...	1538	0	GDM
.....cacccggcugugugcacaugugcc..	6	0	GDM
.....cacccggcugugugcacaugugUc..	2	1	GDM
.....cacccggcugugugcacaugugAc..	1	1	GDM
.....cacccggcugugugcacaugugcU..	101	1	GDM
.....cacccggcugugugcacaugugcA..	9	1	GDM

hsa-miR-941

```
ugugcacaugugcccagggcccgggacagcgccacggaagaggacgcacccggcugugugcacaugugccca
.............................................cacccggcugugugcacaugugcA..    9          1          GDM
.............................................cacccggcugugugcacaugugcU..    101        1          GDM
.............................................cacccggcugugugcacaugugcc..    6          0          GDM
.............................................cacccggcugugugcacaugugcU..    101        1          GDM
.............................................cacccggcugugugcacaugugUc..    2          1          GDM
.............................................cacccggcugugugcacaugugcG..    1          1          GDM
.............................................acccggcugugugcacaug......    1          0          GDM
.............................................acccggcugugugcacaugu.....    1          0          GDM
.............................................acccggcugugugGacaugug....    2          1          GDM
.............................................acccggcugugugcacaugug....    3          0          GDM
.............................................acccggcugugugcacaugugc...    9          0          GDM
.............................................acccggcugugugcacaugugc...    9          0          GDM
.............................................acccggcugugugcacaugugU...    3          1          GDM
.............................................acccggcugugugcacaugugc...    9          0          GDM
.............................................acccggcugugugcacaugugU...    3          1          GDM
.............................................acccggcuguguAcacaugugc...    1          1          GDM
.............................................acccggcugugugcacaugugU...    3          1          GDM
..............................................ccggcugugugcacaugugc...    3          0          GDM
```