hsa-miR-941

```
5'- ugugcacaugugcccagggcccgggacagcgccacggaagaggacgcacccggcugugugcacaugugccca  -3'    exp
    ((.(((((((((((.((.(((((.(((...(((((.........)).))).))))))).)).)))))))))).))    reads    mm    sample
    .........................................Ccacccggcugugugcacaugugc...          1        1     seq
    .........................................cacccggcugugugcaca........           13       0     seq
    .........................................cacccggcugugugcacau.......           3        0     seq
    .........................................cacccggcugugugcaUaug......           1        1     seq
    .........................................Uacccggcugugugcacaug......           1        1     seq
    .........................................cCcccggcugugugcacaug......           1        1     seq
    .........................................cacccggcugugugcacaug......           129      0     seq
    .........................................caUccggcugugugcacaug......           1        1     seq
    .........................................cacccggcugugugcacaugA.....           4        1     seq
    .........................................Aacccggcugugugcacaugu.....           1        1     seq
    .........................................cacccggcugugugUacaugu.....           1        1     seq
    .........................................cacccggcugugugcacaugu.....           65       0     seq
    .........................................caccUggcugugugcacaugug....           1        1     seq
    .........................................cacccggcugugugcacauguA....           7        1     seq
    .........................................cacccggcugugugcacaugug....           55       0     seq
    .........................................cacccggcugugugcacauguU....           1        1     seq
    .........................................cacccggcuguguAcacaugug....           1        1     seq
    .........................................cacccggcugugugcacaugugU...           79       1     seq
    .........................................cacccggcugugugcacauAugc...           1        1     seq
    .........................................cUcccggcugugugcacaugugc...           1        1     seq
    .........................................cacccggcugugugUacaugugc...           1        1     seq
    .........................................cacccggcugugugcacaugugc...           208      0     seq
    .........................................cacccggcuAugugcacaugugc...           1        1     seq
    .........................................cacccggcugCgugcacaugugc...           1        1     seq
    .........................................cacccggcugugugcacaugugG...           21       1     seq
    .........................................cacccggcugugugGacaugugc...           1        1     seq
    .........................................cacccggcugugugcacaugugA...           60       1     seq
    .........................................cacccggcugugugcacaugGgc...           1        1     seq
    .........................................cacccggcugugugcacauUugc...           1        1     seq
    .........................................cacccggcugugugcacaugugcU..           47       1     seq
    .........................................acccggcugugugcacaug......            1        0     seq
    .........................................cccggcugugugcacaugug....            1        0     seq
```