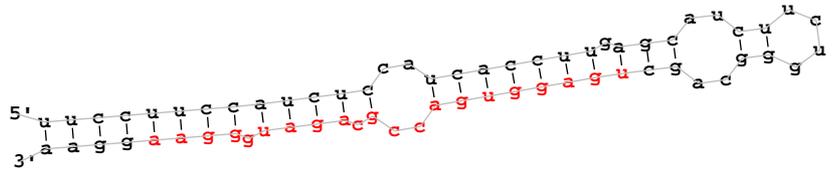miRBase precursor : hsa-mir-7854
Total read count : 404
hsa-miR-7854-3p read count 14
remaining reads : 390


```
                              hsa-miR-7854-3p
5'- uuccuuccaucuccaucaccuugagcaucuucugggcagcugaggugaccgcagaugggaaggaa  -3'    exp
    .((((((((((((((..(((((((.(((..((....))..))))))))))..).))))).)))))))).  reads    mm      sample
    ...cuuccaucuccaucaccuug.........................................       1        0       Z3_
    ...cuuccaucuccaucaccuuga........................................       3        0       Z3_
    ...cuuccaucuccaucaccCuga........................................       1        1       Z3_
    ...cuuccaucuccaucaccuugaU.......................................       2        1       Z3_
    ...cCuccaucuccaucaccuugag.......................................       1        1       Z3_
    ...cuuccaucuccaucaccuugag.......................................       6        0       Z3_
    ...cuuccaucuccaucaccuugagc......................................       7        0       Z3_
    ...cuuccaucuccaucaccuugagU......................................       2        1       Z3_
    ...cuuccaucuccaucaccuugagcU.....................................       1        1       Z3_
    ...cuuccaucuccaucaccuugagcC.....................................       1        1       Z3_
    ...................................ugaggugaccgcagauggga.....           2        0       Z3_

    ..ccuuccaucuccaucaccuugag.......................................       3        0       Z1_
    ...cuuccaucuccaucaccuug.........................................       6        0       Z1_
    ...cuCccaucuccaucaccuuga........................................       1        1       Z1_
    ...cuuccaucuGcaucaccuuga........................................       1        1       Z1_
    ...cuuccaucuccaucaccuuga........................................       22       0       Z1_
    ...cuuccaucuccaCcaccuugag.......................................       2        1       Z1_
    ...cuuccaucCccaucaccuugag.......................................       2        1       Z1_
    ...cuuccaucuccaucaccuugaA.......................................       1        1       Z1_
    ...cuuccaucuccaucaccuugag.......................................       33       0       Z1_
    ...cuuccaucuccaucaccuugagc......................................       22       0       Z1_
    ...cuuccaucuccaucaccuugagA......................................       1        1       Z1_
    ...cuuccaucuccaucacUuugagc......................................       1        1       Z1_
    ...cuuccaucuccaucaccuugagU......................................       1        1       Z1_
    ...cuuccaucuccaucaccAugagc......................................       1        1       Z1_
    ...cuuccaucuccaucaccuugagcau....................................       2        0       Z1_
    ...................................ugaggugGccgcagauggga.....           2        1       Z1_
    ...................................ugaggugaccgcagauggga.....           2        0       Z1_
    ...................................ugaggugacUgcagauggga.....           1        1       Z1_

    ...cuuccaucuccaucaccuug.........................................       10       0       GDM
    ...cuuccaucuccaucaccuGg.........................................       1        1       GDM
    ...cuuccaucuccaucaccuuA.........................................       2        1       GDM
    ...cuuccaucuccaucaccuug.........................................       10       0       GDM
```

hsa-miR-7854-3p

uuccuuccaucuccaucaccuugagcaucuucugggcagcugaggugaccgcagauggggaaggaa

Sequence	Count	MM	Group
...cuuccaucuccaucaccuug..	10	0	GDM
...cuuccaucuccaucaccuuA..	2	1	GDM
...cuuccaucuccaucGccuuga.......................................	3	1	GDM
...cuuccaucAccaucaccuuga.......................................	1	1	GDM
...cuuccaucuccaucaccuuga.......................................	61	0	GDM
...cuuccaucuccaucaccCuga.......................................	1	1	GDM
...cuuccaucuccaucaccuugU.......................................	2	1	GDM
...cuuccaucuccaucaccuuga.......................................	61	0	GDM
...cuuccaucuccaucaccuuga.......................................	61	0	GDM
...cuuccaucuccaucaccuugag.....................................	84	0	GDM
...cuCccaucuccaucaccuugag.....................................	1	1	GDM
...cuuccaucuccaucaccuugag.....................................	84	0	GDM
...cuuccaucuccaucaccuugaA.....................................	2	1	GDM
...cuuccaucuccaucGccuugag.....................................	1	1	GDM
...cuuccaucuccaucaccuugaA.....................................	2	1	GDM
...cuuccaucuccaucaccuugaC.....................................	1	1	GDM
...cuuccaucAccaucaccuugag.....................................	1	1	GDM
...cuuccaucuccaucaccuugag.....................................	84	0	GDM
...cuuccaucuccaucaccuugagc....................................	25	0	GDM
...cuucUaucuccaucaccuugagc....................................	1	1	GDM
...cuuccaucuccaucaccuugagA....................................	9	1	GDM
...cuuccaucuccaucaccuugagc....................................	25	0	GDM
...cuuccaucuccaucaccCugagc....................................	2	1	GDM
...cuuccaucuccaucaccuugagA....................................	9	1	GDM
...cuuccaucuccaucaccuugagc....................................	25	0	GDM
...cuuccaucuccaucaccuugagA....................................	9	1	GDM
...cuuccaucuccaucaccuugagU....................................	2	1	GDM
...cuuccaucuccaucaccuugagU....................................	2	1	GDM
....Cuccaucuccaucaccuugagc....................................	1	1	GDM
....uuccaucuccaucaccuugagc....................................	2	0	GDM
....uuccaucuccaucaccuugagc....................................	2	0	GDM
.....uccaucuccaucaccuuga......................................	2	0	GDM
.....uccaucuccaucaccuuga......................................	2	0	GDM
.....uccaucuccaucaccuugagU....................................	1	1	GDM
.....uccaucuccaucaccuugagc....................................	1	0	GDM
....ugaggugaccgcagauggga.....	3	0	GDM
....ugaggugGccgcagauggga.....	2	1	GDM
....ugaggugaccgcagaugggaaU...	1	1	GDM
....ugaggugaccgcGgaugggaag...	1	1	GDM
..ccuuccaucuccaucaccuuga......................................	1	0	Z2_
...cuuccaucuccaucaccuuga......................................	13	0	Z2_
...cuuccaucuccaucacAuugag.....................................	1	1	Z2_
...cuuccaucuccaucaccuugag.....................................	19	0	Z2_
...cuuccauUuccaucaccuugagc....................................	1	1	Z2_
...cuuccaucuccaucaccuugagU....................................	3	1	Z2_
...cuuccaucuccaucaccuugagc....................................	11	0	Z2_