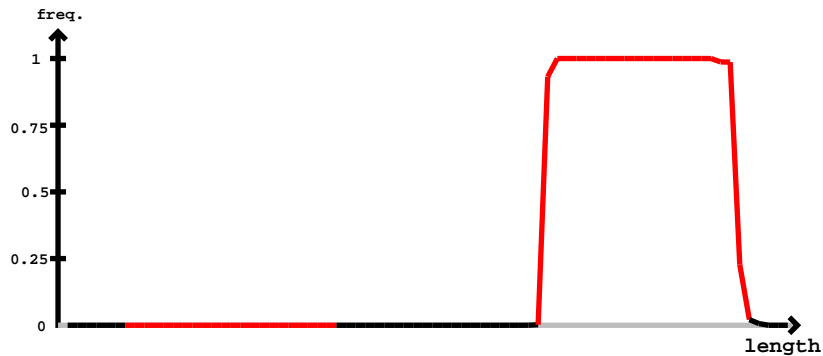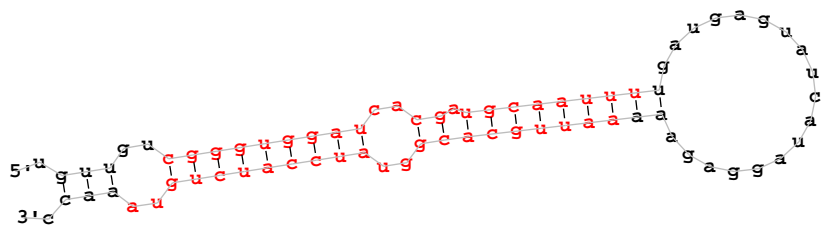miRBase precursor : hsa-mir-363
Total read count : 4436
hsa-miR-363-5p read count : 0
hsa-miR-363-3p read count : 4436
remaining reads : 0

hsa-miR-363-5p

hsa-miR-363-3p

```
5'- uguugucggguggaucacgaugcaauuuugaugaguaucauaggagaaaaauugcacgguauccaucuguaaacc  -3'   exp
     .(((..((((((((((..((.((((((((((....................))))))))))..))))))))))..))).   reads   mm   sample
     ................................................Caaauugcacgguauccaucug......    1      1    seq
     ................................................aaaauugcacgguauccaucug......    6      0    seq
     ................................................aaaauugcacgguauccaucugu.....    2      0    seq
     ................................................aauugcacgguauccauc........    51     0    seq
     ...............................................Cauugcacgguauccauc........     6      1    seq
     ...............................................aauugcacgguauccaucu.......     3      0    seq
     ...............................................aauugAacgguauccaucug......     5      1    seq
     ...............................................aauugcUcgguauccaucug......     1      1    seq
     ...............................................aauugcacgguauccaucGg......     9      1    seq
     ...............................................aauugcacgguauUcaucug......     1      1    seq
     ...............................................aauugcacgCuauccaucug......     1      1    seq
     ...............................................aauugcacgguauccaucuA......     1      1    seq
     ...............................................Gauugcacgguauccaucug......     1      1    seq
     ...............................................aGuugcacgguauccaucug......     1      1    seq
     ...............................................aauCgcacgguauccaucug......     2      1    seq
     ...............................................aauugUacgguauccaucug......     4      1    seq
     ...............................................Uauugcacgguauccaucug......     1      1    seq
     ...............................................aauugcacgguauccaucCg......     1      1    seq
     ...............................................Cauugcacgguauccaucug......     229    1    seq
     ...............................................aauugcacAguauccaucug......     2      1    seq
     ...............................................aauugcGcgguauccaucug......     2      1    seq
     ...............................................aauugcacgguaucUaucug......     2      1    seq
     ...............................................aauugcacgguUuccaucug......     3      1    seq
     ...............................................aauugcacgguaucGaucug......     1      1    seq
     ...............................................aaAugcacgguauccaucug......     3      1    seq
     ...............................................aauugcacgguauccaAcug......     2      1    seq
     ...............................................aauuAcacgguauccaucug......     1      1    seq
     ...............................................aauugcacgguauGcaucug......     2      1    seq
     ...............................................aauugcacggCauccaucug......     1      1    seq
     ...............................................aauugcacgguaAccaucug......     2      1    seq
     ...............................................aauugcCcgguauccaucug......     1      1    seq
     ...............................................aauugcaAgguauccaucug......     1      1    seq
     ...............................................aauugcacgUuauccaucug......     1      1    seq
     ...............................................aauugcacgguauccauUug......     1      1    seq
```

hsa-miR-363-5p

hsa-miR-363-3p

uguugucggguggaucacgaugcaauuuugaugaguaucauaggagaaaaauugcacgguauccaucguaaacc

....aauugcacgAuauccaucug......	4	1	seq
....aaCugcacgguauccaucug......	2	1	seq
....aCuugcacgguauccaucug......	2	1	seq
....aauugcacggAauccaucug......	8	1	seq
....aauugcacggGauccaucug......	1	1	seq
....aauugcacgguauccaucug......	2857	0	seq
....aauugGacgguauccaucug......	1	1	seq
....aauugcacgguaucAaucug......	1	1	seq
....aauCgcacgguauccaucugu.....	1	1	seq
....aauugcacggCauccaucugu.....	1	1	seq
....aauugcacgguauGcaucugu.....	1	1	seq
....aauugcacgCuauccaucugu.....	1	1	seq
....Gauugcacgguauccaucugu.....	1	1	seq
....aauugcacgguauccaucCgu.....	1	1	seq
....aauugcacgguauccaucugA.....	4	1	seq
....aauugcacgguauccaucugG.....	3	1	seq
....aauugcacgguauccaucugu.....	757	0	seq
....Uauugcacgguauccaucugu.....	1	1	seq
....aauugcacgguCuccaucugu.....	1	1	seq
....aauugAacgguauccaucugu.....	2	1	seq
....aauugcacCguauccaucugu.....	1	1	seq
....aauugcacAguauccaucugu.....	2	1	seq
....aauugcacgguauccauGugu.....	2	1	seq
....Cauugcacgguauccaucugu.....	52	1	seq
....aaCugcacgguauccaucugu.....	1	1	seq
....aauugcacgguauccaucugua....	47	0	seq
....aauugcacgguauccaucuAua....	1	1	seq
....aauugcacgguauccaucuguU....	1	1	seq
....Cauugcacgguauccaucugua....	6	1	seq
....aauugcacgguauccaucuguaG...	5	1	seq
....aauugcacgguauccaucugAaa...	1	1	seq
....aauugcacgguauccaucuguaU...	5	1	seq
....aauugcacgguauccaucuguaa...	10	0	seq
....auugcaGgguauccaucug......	1	1	seq
....auugGacgguauccaucug......	1	1	seq
....Cuugcacgguauccaucug......	3	1	seq
....auugcacgguauccaCcug......	1	1	seq
....auugcacgguauccauGug......	1	1	seq
....auugcacgguauccaucug......	196	0	seq
....auugcacgguauccaucugu.....	80	0	seq
....auugcacggGauccaucugu.....	1	1	seq
....Cuugcacgguauccaucugua....	1	1	seq
....auugcacgguauccaucugAa....	1	1	seq
....auugcacgguauccaucugua....	8	0	seq
....auugcacgguauccaucuguaa...	5	0	seq
....auugcacgguauccaucuguaG...	1	1	seq
....auugcacgguauccaucuguaU...	1	1	seq