Homework 3

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1. The cycle repeats 4 numbers continuously.

[24]

3 1 11

```
x <- 1
for(i in 1:25)
    x <- c(x, (11 * tail(x, 1)) %% 16)
x
## [1] 1 11 9 3 1 11 9 3 1 11 9 3 1 11 9 3 1 11 9</pre>
```

2. From the 4 examples, we can see that when the when the cycle reaches 0, it immeadiatly returns to 12. When a higher number like 1000 is entered, it immeadiately decreases to 11 or lower.

```
x <- 0
for(i in 1:25)
 x \leftarrow c(x, (tail(x, 1) + 12) \% 13)
  [1] 0 12 11 10 9 8 7 6 5 4 3 2 1 0 12 11 10 9 8 7 6 5 4
## [24] 3 2 1
x <- 50
for(i in 1:25)
 x \leftarrow c(x, (tail(x, 1) + 12) \% 13)
## [1] 50 10 9 8 7 6 5 4 3 2 1 0 12 11 10 9 8 7 6 5 4 3 2
## [24] 1 0 12
x <- 100
for(i in 1:25)
 x \leftarrow c(x, (tail(x, 1) + 12) \% 13)
## [1] 100
             8
                     6
                             4
                                 3
                                     2
                                         1
                                             0 12 11 10
                                                                 8
## [18]
                 3
                     2
                         1
                             0 12 11 10
x <- 1000
for(i in 1:25)
 x \leftarrow c(x, (tail(x, 1) + 12) \% 13)
## [1] 1000
                   10
                                   7
                                        6
                                             5
                                                                 1
                                                                          12
              11
                         8
                              7
                                   6
                                        5
                                             4
                                                  3
                                                       2
## [15]
         11
              10
                    9
```

3. Decreasing the 100,000 number a bit so my computer doesn't fail the calc.

```
x <- 1234567
r <- 1234567 / (2^31 - 1)

for(i in 1:99999) {
    x <- c(x, (16807 * tail(x, 1)) %% (2^31 - 1))
    r <- c(r, tail(x, 1) / (2^31 - 1))
}

tbl <- data.frame(x, r)
chisq.test(tbl)</pre>
```

```
##
## Pearson's Chi-squared test
##
## data: tbl
## X-squared = 1.362e-18, df = 99999, p-value = 1
```

The above is almost certainly wrong from the looks of it, not sure what to change here. Let's try runs. Trying this with a package I found.

```
library(randtests)
runs.test(r)
```

```
##
## Runs Test
##
## data: r
## statistic = -0.32888, runs = 49949, n1 = 50000, n2 = 50000, n =
## 100000, p-value = 0.7422
## alternative hypothesis: nonrandomness
```

5.

4. .

a. First with inverse-normal:

```
normrandit <- function(){
   U <- runif(1)
   return(qnorm(U))
}

itstats <- function(N){
   x <- numeric(0)
   for(i in 1:N){
      x <- c(x,normrandit())
   }
   return(list(mean=mean(x), sd=sd(x)))
}</pre>
```

b. Now with Muller:

```
normrandbm <- function(){
    U <- runif(2)
    x <- ((-2*log(U[1]))^(1/2))*cos(2*pi*U[2])
    y <- ((-2*log(U[1]))^(1/2))*sin(2*pi*U[2])
    return(c(x=x, y=y))
}

bmstats <- function(N){
    x <- numeric(0)
    for(i in 1:N){
        x <- c(x,normrandbm())
    }
    return(list(mean=mean(x), sd=sd(x)))
}</pre>
```

c. Finally with accept/reject

```
normrandar <- function(){</pre>
  repeat{
    U <- runif(2)
    x \leftarrow -\log(U[1])
    y \leftarrow -\log(U[2])
    if(y \ge ((x-1)^2)/2){
      break
    }
  }
  if (runif(1) >= .5)
    x < -1 * x
  return(x)
arstats <- function(N){</pre>
  x <- numeric()
  for(i in 1:N){
    x <- c(x,normrandar())</pre>
  return(list(mean=mean(x), sd=sd(x)))
```

d. Let's check the means, sd's, and runtimes

```
library(plyr)
means <- data.frame(it=c(), bm=c(), ar=c())
sds <- data.frame(it=c(), bm=c(), ar=c())
times <- data.frame(it=c(), bm=c(), ar=c())

for (n in c(100, 1000, 10000, 100000)) {
   for (i in 1:10) {
    it <- itstats(n)
    bm <- bmstats(n)</pre>
```

```
ar <- arstats(n)</pre>
  means <- rbind(means, c(it$mean, bm$mean, ar$mean, n))</pre>
  sds <- rbind(sds, c(it$sd, bm$sd, ar$sd, n))</pre>
  times <- rbind(times, c(system.time(itstats(n))[[3]], system.time(itstats(n))[[3]], system.time(itstats(n))[[3]]
}
colnames(means) <- c("it","bm","ar","n")</pre>
colnames(sds) <- c("it","bm","ar","n")</pre>
colnames(times) <- c("it","bm","ar","n")</pre>
library(plyr)
means <- read.csv("means.csv")</pre>
sds <- read.csv("sds.csv")</pre>
times <- read.csv("times.csv")</pre>
mean_avgs <- ddply(means, ~n, summarise, mean_it = mean(it), mean_bm = mean(bm), mean_ar = mean(ar));me
##
                 mean_it
                                mean_bm
                                               mean_ar
         n
## 1 1e+02 -0.0289959302 0.0209949281 -0.0196609236
## 2 1e+03 -0.0169933742 -0.0094790194 0.0059108701
## 3 1e+04 0.0007567270 -0.0009539500 -0.0033564852
## 4 1e+05 0.0005063963 -0.0001194233 -0.0005487528
sd_avgs <- ddply(sds, ~n, summarise, mean_it = mean(it), mean_bm = mean(bm), mean_ar = mean(ar));sd_avg
##
             mean_it mean_bm
                                  mean_ar
## 1 1e+02 1.0158512 1.0057870 1.0240378
## 2 1e+03 0.9969124 0.9941471 0.9948150
## 3 1e+04 0.9987530 0.9996241 0.9981487
## 4 1e+05 0.9998424 1.0003970 1.0001452
times_avgs <- ddply(times, ~n, summarise, mean_it = mean(it), mean_bm = mean(bm), mean_ar = mean(ar));t
         n mean_it mean_bm mean_ar
##
## 1 1e+02
            0.001
                      0.001
                              0.000
## 2 1e+03
             0.005
                      0.006
                              0.006
## 3 1e+04
            0.169
                      0.171
                              0.142
## 4 1e+05 13.246 13.232 13.260
```

6.

a. • b. We don't really need to use the method of 1 or 0 to estimate pi, it should estimate fine without checking, and will not give a "4" when a lower N value is given

```
estimatepi <- function(N){
    x <- runif(N)
    y <- runif(N)
    d <- sqrt(x^2 + y^2)
    pi <- (4 * sum(d < 1.0) / N)
    se <- sd(d) / sqrt(length(d))</pre>
```

```
ci <- qnorm(0.95)*sd(d)/sqrt(N)
return(list(pi=pi, se=se, ci=ci))
}</pre>
```

c. Let's check how large N needs to be for the estimate to be accurate.

```
values <- data.frame(pi=c(),se=c(),ci=c())
x <- seq(1000,10000,by=500)
for (n in x) {
    values <- rbind(values, estimatepi(n))
}
rownames(values) <- seq(1000,10000,by=500)

values$lower <- 3.14 - values$ci
values$upper <- 3.14 + values$ci
values$interval <- values$lower
values</pre>
```

```
##
                                      сi
                                            lower
                                                     upper
                                                              interval
              рi
                          se
## 1000 3.076000 0.008764781 0.014416782 3.125583 3.154417 0.028833564
## 1500 3.096000 0.007496967 0.012331414 3.127669 3.152331 0.024662828
## 2000 3.186000 0.006277706 0.010325908 3.129674 3.150326 0.020651816
## 2500 3.148800 0.005708940 0.009390371 3.130610 3.149390 0.018780743
## 3000 3.137333 0.005195896 0.008546488 3.131454 3.148546 0.017092977
## 3500 3.192000 0.004811151 0.007913639 3.132086 3.147914 0.015827278
## 4000 3.156000 0.004481106 0.007370764 3.132629 3.147371 0.014741528
## 4500 3.148444 0.004263198 0.007012337 3.132988 3.147012 0.014024674
## 5000 3.156800 0.004032255 0.006632469 3.133368 3.146632 0.013264939
## 5500 3.139636 0.003845109 0.006324641 3.133675 3.146325 0.012649282
## 6000 3.146000 0.003667488 0.006032482 3.133968 3.146032 0.012064963
## 6500 3.179077 0.003471730 0.005710487 3.134290 3.145710 0.011420975
## 7000 3.125143 0.003396492 0.005586731 3.134413 3.145587 0.011173463
## 7500 3.122667 0.003319845 0.005460659 3.134539 3.145461 0.010921317
## 8000 3.148500 0.003186021 0.005240539 3.134759 3.145241 0.010481078
## 8500 3.116235 0.003121408 0.005134259 3.134866 3.145134 0.010268518
## 9000 3.145333 0.002977020 0.004896762 3.135103 3.144897 0.009793524
## 9500 3.156211 0.002918253 0.004800099 3.135200 3.144800 0.009600197
## 10000 3.128800 0.002873611 0.004726670 3.135273 3.144727 0.009453340
```