# Homework 3

## DATA604 Simulation and Modeling

Daniel Dittenhafer
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```
##
## Attaching package: 'gplots'
## The following object is masked from 'package:stats':
##
##
       lowess
library(knitcitations)
library(RefManageR)
cleanbib()
cite_options(style="markdown")
bibPkgTseries <- bibentry(bibtype="Misc",</pre>
                 author=personList(person(family="Trapletti", given="Adrian"),
                                    person(family="Hornik", given="Kurt")),
                 journal="Annual Review of Sociology",
                 title="tseries: Time Series Analysis and Computational Finance",
                 year=2015,
                 note="R package version 0.10-34",
                 url="http://CRAN.R-project.org/package=tseries")
```

#### 1

Starting with  $X_0 = 1$ , write down the entire cycle for  $X_i = 11X_{i-1} \mod(16)$ 

```
fn1 <- function(x0)</pre>
{
  df <- data.frame(X=c(), R=c())</pre>
  x <- x0
  continue <- TRUE
  while(continue)
    xi <- (11 * x) %% 16
    df <- rbind(df, data.frame(X=x, R=xi))</pre>
    x <- xi
    if(xi == x0)
      break
    }
  }
  return(df)
}
res <- fn1(1)
```

| X  | R  |
|----|----|
| 1  | 11 |
| 11 | 9  |
| 9  | 3  |
| 3  | 1  |
|    |    |

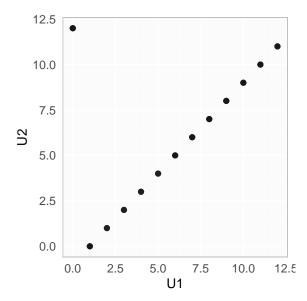
## 2

\*Using the LCG provided below:  $X_i = (X_{i-1} + 12) mod(13)$ , plot the pairs  $(U_1.U_2), (U_2, U_3), ...$  and observe the lattice structure obtained. Discuss what you observe.

```
fn2 <- function(x0, max=100)
{
    df <- data.frame(U1=c(), U2=c())
    x <- x0

    for(i in 1:max)
    {
        xi <- (x + 12) %% 13
        df <- rbind(df, data.frame(U1=x, U2=xi))
        x <- xi
    }

    return(df)
}
# Call the function starting at x0=1
res <- fn2(1)</pre>
```



The chart above suggests there are only 13 points, but actually the LCG cycle period is 13 and numbers are repeating.

| U1 | U2 |
|----|----|
| 1  | 0  |
| 0  | 12 |
| 12 | 11 |
| 11 | 10 |
| 10 | 9  |
| 9  | 8  |
| 8  | 7  |
| 7  | 6  |
|    |    |

| U1 | U2 |
|----|----|
| 6  | 5  |
| 5  | 4  |
| 4  | 3  |
| 3  | 2  |
| 2  | 1  |
| 1  | 0  |
|    |    |

3

Implement the pseudo-random number generator:

$$X_i = 16807X_{i-1} \bmod (2^{31} - 1)$$

Using the seed 1234567, run the generator for 100,000 observations. Perform a chi-square goodness-of-fit test on the resulting PRN's. Use 20 equal-probability intervals and level  $\alpha=0.05$ . Now perform a runs up-and-down test with  $\alpha=0.05$  on the observations to see if they are independent.

```
fnLCG3 <- function(seed = 1, n = 1)
{
  rands <- rep(NA, n)
  x <- seed
  modVal <- (2^31 - 1)

  for(i in 1:n)
  {
    xi <- (16807 * x) %% (modVal)
    rands[i] <- xi
    x <- xi
  }

  return(rands)
}

n=100000
rn <- fnLCG3(1234567, n)</pre>
```

## Chi-Square Test

```
intervals <- 20
maxRn <- max(rn)
minRn <- min(rn)
intWidth <- (maxRn - minRn) / intervals
lwr <- minRn
dfCounts <- data.frame(intID=c(), count=c())
# Bin the data ourselves, I'd guess there
# is an easier way, but this will do.</pre>
```

```
for(i in 1:intervals)
{
  upr <- lwr + intWidth</pre>
  inRange <- rn[lwr <= rn & rn < upr]</pre>
  dfCounts <- rbind(dfCounts, data.frame(intID=i, count=length(inRange)))</pre>
  # setup for next interval range
  lwr <- upr
}
# Do our own Chi-Squared test
Expected <- (100000 / intervals)</pre>
chi2 <- sum((dfCounts$count - Expected)^2 / Expected)</pre>
## [1] 14.7762
# Use built-in function to compare
chiTest <- chisq.test(dfCounts$count)</pre>
chiTest
##
```

The p-value = 0.7367029 is not less than  $\alpha = 0.05$ , therefore we doesn't reject the null hypothesis that the distribution is uniform.

| intID | count |
|-------|-------|
| 1     | 5069  |
| 2     | 5028  |
| 3     | 5044  |
| 4     | 5087  |
| 5     | 4948  |
| 6     | 4953  |
| 7     | 4937  |
| 8     | 4933  |
| 9     | 4900  |
| 10    | 4957  |
| 11    | 5088  |
| 12    | 4994  |
| 13    | 5076  |
| 14    | 5019  |
| 15    | 5002  |
| 16    | 5067  |
| 17    | 4981  |
| 18    | 4914  |
| 19    | 5062  |
| 20    | 4940  |
|       |       |

#### Runs Up-and-Down Test

##

##

## data: dfCounts\$count

Chi-squared test for given probabilities

## X-squared = 14.776, df = 19, p-value = 0.7367

Using the tseries package, we execute the Runs test (Trapletti and Hornik, 2015). First we have to construct the +/- vector. Here we simply convert to boolean.

```
s <- rep(NA, n - 1)
for(i in 1:n - 1)
{
    s[i] <- rn[i] < rn[i + 1]
}
runsTest <- runs.test(as.factor(s))
runsTest</pre>
```

```
##
## Runs Test
##
## data: as.factor(s)
## Standard Normal = 105.84, p-value < 2.2e-16
## alternative hypothesis: two.sided</pre>
```

Based on the p-value < 0.05, we reject the null hypothesis and conclude there is not evidence to support independence in the psuedo-random numbers.

#### 4.

 $Give\ inverse-transforms,\ composition,\ and\ acceptance-rejectino\ algorithms\ for\ generating\ from t\ he\ following\ density:$ 

$$f(x) = \begin{cases} \frac{3x^2}{2} & -1 \le x \le 1\\ 0 & otherwise \end{cases}$$

### References

Trapletti, A. and K. Hornik. tseries: Time Series Analysis and Computational Finance. R package version 0.10-34. 2015. URL: http://CRAN.R-project.org/package=tseries.