

Statistical Computing and Simulation: HW1

Department of Statistics, NCCU

葉佐晨

112354016@nccu.edu.tw

高崇哲

112354@nccu.edu.tw

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Statistical Computing and Simulation

Assignment 1, Due March 15/2024

- (a) Write a computer program using the Mid-Square Method using 6 digits to generate 10,000 random numbers ranging over $[0, 999999]$. Use the Kolmogorov-Smirnov Goodness-of-fit test to see if the random numbers that you create are uniformly distributed. (Note: You must notify the initial seed number used, and you may adapt 0.05 as the α value. Also, you may find warning messages for conducting the Goodness-of-fit test, and comment on the Goodness-of-fit test.)
- (b) Consider the combination of 3 multiplicative congruential generators, i.e.,

$$u_i = \frac{x_i}{30269} + \frac{y_i}{30307} + \frac{z_i}{30323} \pmod{1}$$

with $x_i = 171x_{i-1} \pmod{30269}$, $y_i = 172y_{i-1} \pmod{30307}$, $z_i = 170z_{i-1} \pmod{30323}$. Compare the results in (a) and (b), and discuss your findings.

```
library(magrittr)
library(extraDistr)
```

```
midSquare <- function(size = 1){
  initSample <- function(){sample(0:999999, size = 1)}

  num <- initSample()
```

```

nums <- c(num)

for (i in seq(size)){
  num <- num^2 %>%
    format(scientific = FALSE) %>%
    substr(start = 4, stop = 9) %>%
    ifelse(. == "", initSample(), .) %>%
    as.integer()

  num <- ifelse(num == nums[length(nums)], initSample(), num)
  nums <- c(nums, num)
}

return(nums)
}

```

```

randSeq <- midSquare(size = 10000)
ks.test(randSeq, "pdunif", 0, 999999)

```

```

## Warning in ks.test.default(randSeq, "pdunif", 0, 999999): ties should not be
## present for the Kolmogorov-Smirnov test

```

```

##
## Asymptotic one-sample Kolmogorov-Smirnov test
##
## data:  randSeq
## D = 0.13611, p-value < 2.2e-16
## alternative hypothesis: two-sided

```

```

LCG <- function(size, mod, mul, seed = NULL){
  nums <- c()
  if (!is.null(seed)){set.seed(seed)}

  xyz <- sapply(mod, function(x){sample(0:x, size = 1)})

  for (i in seq(size)){
    u <- sum(xyz / mod)
    nums <- c(nums, u)
    xyz <- (xyz * mul) %% mod
  }
}

```

```

}

return(nums)
}

mod <- c(30269, 30307, 30323)
mul <- c(172, 171, 170)
randomSeq <- LCG(size = 10000, mod = mod, mul = mul)

ks.test(randomSeq, "punif", 0, 3)

##
## Asymptotic one-sample Kolmogorov-Smirnov test
##
## data: randomSeq
## D = 0.18528, p-value < 2.2e-16
## alternative hypothesis: two-sided

```

3. There are several ways for checking the goodness-of-fit for empirical data. In specific, there are a lot of normality tests available in R. Generate a random sample of size 10, 50, and 100 from $N(0,1)$ and t-distribution (with degrees 10 and 20) in R. You may treat testing random numbers from t-distribution as the power. For a level of significance $\alpha = 0.05$ test, choose at least four normality tests in R (“nortest” module) to check if this sample is from $N(0,1)$. Tests used can include the Kolmogorov-Smirnov test and the Cramer-von Mises test. Note that you need to compare the differences among the tests you choose.

```

library(nortest)

norTest <- function(testList, n, seed = NULL){
  if (!is.null(seed)) {set.seed(seed)}

  res <- rbind(
    sapply(testList, function(test){test(rnorm(n))["p.value"]}),
    sapply(testList, function(test){test(rt(n, df = 10))["p.value"]}),
    sapply(testList, function(test){test(rt(n, df = 20))["p.value"]})
  )

  colnames(res) <- c(c("ad", "cvm", "lillie", "pearson", "sf"))
  rownames(res) <- c("N(0,1)", "t(df=10)", "t(df=20)")
}

```

```

    return(res)
}

countReject <- function(n_trial, n, alpha){
  replicate(n = n_trial,
            norTest(testList, n = n) <= alpha) %>%
    apply(MARGIN = c(1, 2), sum)
}

```

```

n <- c(10, 50, 100)
testList <- c(ad.test, cvm.test, lillie.test, pearson.test, sf.test)
n_trial <- 1000
alpha <- 0.05

```

```

rejList <- lapply(n, function(n){countReject(n_trial = 1000, n = n, alpha = 0.05)})
names(rejList) <- paste0("n=", n)
print(rejList)

```

```

## $'n=10'
##           ad cvm lillie pearson sf
## N(0,1)    57  48     48     71 63
## t(df=10)  75  61     72     71 82
## t(df=20)  58  56     60     72 79
##
## $'n=50'
##           ad cvm lillie pearson  sf
## N(0,1)     47  47     45     56 68
## t(df=10)  126 100     71     62 195
## t(df=20)   75  82     68     65 105
##
## $'n=100'
##           ad cvm lillie pearson  sf
## N(0,1)     54  51     54     59 56
## t(df=10)  187 165    110     63 289
## t(df=20)   89  74     83     60 147

```

5. (a) Use the search engine to download the first one million digits of pi (for example, <https://www.piday.org/million/>) and check via graphic tools if the numbers violate the assumption of random numbers.

- (b) Apply the appropriate tools to test if the random numbers from
(a) satisfy the assumption of random numbers.

```
library(randtoolbox)
```

```
## Loading required package: rngWELL
```

```
## This is randtoolbox. For an overview, type 'help("randtoolbox")'.
```

```
pi1M <- readLines("Pi1MDP.txt", warn = FALSE) %>%  
  strsplit(split = "") %>%  
  unlist() %>%  
  as.integer()
```

```
isRandom <- function(seq, alpha = 0.05){  
  tb <- table(pi1M)  
  testChi <- tb %>% chisq.test  
  print(testChi)  
  isUnif <- ifelse(testChi[["p.value"]] < alpha, FALSE, TRUE)  
  
  cat(rep("*", 60), sep = "")  
  
  r <- diff(range(seq))  
  isIndept <- ifelse(gap.test(seq/r, echo = TRUE)[["p.value"]] < alpha,  
                    FALSE, TRUE)  
  
  return(isUnif & isIndept)  
}
```

```
isRandom(pi1M, alpha = 0.05)
```

```
##  
## Chi-squared test for given probabilities  
##  
## data: .  
## X-squared = 5.5091, df = 9, p-value = 0.7879  
##  
## *****  
## Gap test
```

```
##
## chisq stat = 44, df = 20, p-value = 0.0013
##
##      (sample size : 1000000)
##
## length  observed freq      theoretical freq
## 1          125596          125000
## 2          62325          62500
## 3          31348          31250
## 4          15788          15625
## 5           7900          7812
## 6          3857          3906
## 7          1888          1953
## 8           925          977
## 9           444          488
## 10          232          244
## 11          148          122
## 12           60           61
## 13           20           31
## 14           18           15
## 15            6          7.6
## 16            9          3.8
## 17            3          1.9
## 18            0          0.95
## 19            0          0.48
## 20            1          0.24
## 21            1          0.12

## [1] FALSE
```