The University of British Columbia I.K. Barber Faculty of Science

COSC 405/DATA 405/DATA 505/COSC 505 Modelling and Simulation Practice Term Test 1 Solutions

1	(4	marks	Write	out	the	R	code	required	to
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```
## [1] 27 48 72 101 98 37 22 55 41 79 58 44 61

x <- c(27, 48, 72, 101, 98, 37, 22, 55, 41, 79, 58, 44, 61)

sd(x)
```

(b) produce a boxplot of the sample above.

```
boxplot(x)
```

(c) produce a normal QQ-plot of the sample above with reference line.

```
qqnorm(x)
qqline(x)
```

(d) test whether the mean of the sample differs from 50. and constructs a 99% confidence interval for the mean of the population from which the sample was taken.

```
t.test(x, mu = 50, conf.level=.99)
```

2. (10 marks) Consider the following simulated data and analysis. The simulation is to emulate a paper airplane throwing experiment involving a number of throws for each of a number of different types of paper.

```
papergroup <- factor(rep(1:5, 7))
distances <- rnorm(35, mean = 9, sd = 3)</pre>
```

- (a) How many different types of paper are being simulated?

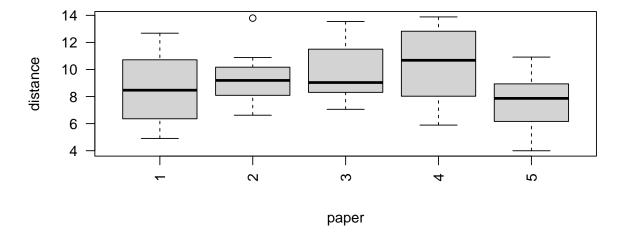
 Since there are 5 levels in the paper factor, there must be 5 different types paper.
- (b) Is there a true difference in the mean distance travelled by paper airplanes in the different groups?

The mean for all 35 measurements is 9; therefore, all simulated paper airplanes in all of the groups have the same mean. There is no difference.

(c) Provide the R code to obtain the following plot

```
boxplot(distances ~ papergroup, las = 2, xlab = "paper", ylab = "distance")
title("Paper Airplane Experiment Data")
```

Paper Airplane Experiment Data



(d) Fill in the blanks in the code below in order to get the following output.

```
_____(lm(_____~___))
```

(e) Based on the output from part (d), what can you conclude about the simulated paper airplanes? Briefly support your conclusions.

The p-value is .3316 which is not small. Therefore, there is no evidence that the mean distance travelled by the simulated paper airplanes are different for the different groups.

- 3. (6 marks) Consider the gas mileage data in table.b3 of the MPV package.
 - (a) Fit a multiple regression model to estimate mean gas mileage y for cars with x_7 number of transmission speeds and having weight x_{10} .

The fitted model is

$$\hat{y} = 30.375 + 2.048x_7 - 0.005x_{10}$$

where the error has mean 0 and an estimated standard deviation of 3.16.

(b) Use the model to estimate mean gas mileage for cars having weight 5000 pounds and 4 transmission speeds.

```
predict(b3.lm, newdata = data.frame(x7 = 4, x10 = 5000),
    interval="confidence")

## fit lwr upr
## 1 14.86889 10.83812 18.89965
```

The 95% confidence interval for gas mileage for such cars is (10.8, 18.9).

4. (5 marks) Consider a data frame, such as the women object built into R, for which the heights could be taken as x values and the weights could be taken as y values.

Write an R function called TukeySmooth which outputs a new data frame consisting of a column of equally spaced x values and a column of corresponding local medians, and which takes the following arguments

- \mathbf{x} : the vector of x values
- \bullet y: the vector of y values
- x.min: a constant which specifies the left boundary of the plotted curve
- x.max: a constant which specifies the right boundary of the plotted curve
- window: a constant which specifies the range of the x values used to calculate each of the moving medians.

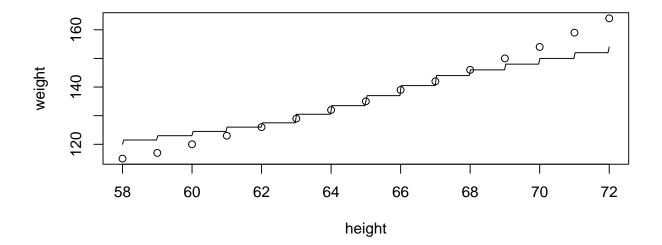
The output for this function will be a data frame with 2 columns: \mathbf{x} and \mathbf{y} , which will correspond to the y-medians and the corresponding x locations where the medians are taken.

This is very similar to the **smoother()** function described in the textbook, with median in place of mean.

```
TukeySmooth <- function(x, y, x.min, x.max, window=1) {
    xpoints <- seq(x.min, x.max, len=401)
    ymedians<- numeric(401)
    for (i in 1:length(xpoints)) {
        indices <- which(abs(x - xpoints[i]) < window)
        if (length(indices) < 1) {
            stop("Your choice of window width is too small.")
        } else {
            ymedians[i] <- median(y[indices])
        }
    }
    data.frame(x = xpoints, y = ymedians)
}</pre>
```

5. (2 marks) Apply the function obtained in the previous question, using a window width of 5, to the data in women, plotting the data, and overlaying the smooth curve.

```
women.TS <- TukeySmooth(women$height, women$weight, x.min = 58, x.max=72, window=5)
plot(weight ~ height, data = women)
lines(women.TS)</pre>
```



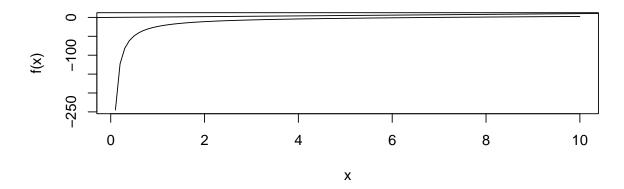
6. (5 marks) Consider the following iteration scheme:

$$x_{n+1} = f(x_n) := \frac{x_n}{2} - \frac{24.5}{x_n}$$

where x_0 is the initial value, say, $x_0 = 0.5$.

(a) Plot the graph of the function f(x), for $x \in [0.1, 10]$ and overlay the straight line with intercept 0 and slope 1. Does f(x) have a fixed point?

```
f <- function(x) x/2 - 24.5/x
curve(f(x), 0.1, 10)
abline(0,1)</pre>
```



Since the function and the line do not intersect, the function does not have a fixed point.

(b) Using a for loop in R, run 10 steps of the proposed scheme, printing out the value of x_n at each step.

```
x <- 25
for (i in 1:10) {
    x \leftarrow f(x)
    print(x)
}
##
   [1] 11.52
##
   [1] 3.633264
## [1] -4.926616
  [1] 2.509679
##
   [1] -8.507364
## [1] -1.373824
## [1] 17.14652
## [1] 7.144399
## [1] 0.1429396
## [1] -171.3296
```

- (c) Based on your analysis, could the proposed scheme lead to a useful pseudorandom number generator?
 - On the basis of the analysis so far, we would not rule out this function as a possible starting point to produce a useful generator, but a lot of additional testing would be needed to ensure that it produced approximately independent and uniformly distributed numbers.

- 7. (4 marks) Which of the following linear congruential pseudorandom number generators have maximal cycle length?
 - (a) $a = 1025, c = 27, m = 2^{31} Yes$
 - (b) $a = 1025, c = 54, m = 2^{31}$ No, since c and m are both multiples of 2.
 - (c) $a = 1025, c = 375, m = 2^{31} Yes$
 - (d) $a = 10241, c = 375, m = 2^{31} Yes$

8. (8 marks) Write a function called myrbinom() to compute N independent binomial random variates which have parameter n and p. The function should use the runif() function in a single for() loop to simulate N vectors of n Bernoulli(p) random variates which are then added to obtain the binomial numbers:

$$X = \sum_{i=1}^{n} B_i$$

is a binomial (n, p) random variable if B_1, \ldots, B_n are independent Bernoulli (p) random variables.

Following the instructions of the question directly, we have:

```
myrbinom <- function(N, n, p) {
    X <- numeric(N)
    for (j in 1:N) {
        X[j] <- sum(runif(n) <= p)
    }
    X
}</pre>
```