

## STAT 515 hw 4

*Binomial, Normal probabilities/quantiles, Q-Q plots*

*Attach a sheet with the R plots and R code printed on it. You may write out your other answers by hand if you want. Just try to make it to grade!!*

1. Each visitor entering a museum must pass through one of five turnstiles: turnstiles A, B, C, D, and E. Suppose 10 visitors enter the museum and that each chooses a turnstile at random, independently of the others.

- (a) Find  $P(\text{None pass through turnstile A})$ .

The probability that a visitor selects turnstile A is  $1/5$ , so if  $X$  is the number of the ten visitors that pass through turnstile A, then  $X \sim \text{Binomial}(10, 1/5)$ . So we have

$$P(X = 0) = \binom{10}{0} (1/5)^0 (1 - 1/5)^{10} = (4/5)^{10} = 0.1073742.$$

- (b) Find  $P(\text{Exactly three pass through turnstile A})$

According to the same binomial distribution, we have

$$P(X = 3) = \binom{10}{3} (1/5)^3 (1 - 1/5)^{10-3} = 0.2013266.$$

- (c) Find  $P(\text{All pass through turnstiles C, D, and E})$

If  $Y$  is the number of visitors entering through turnstiles C, D, and E, then  $Y \sim \text{Binomial}(10, 3/5)$ . So we have

$$P(Y = 10) = \binom{10}{10} (3/5)^{10} (1 - 3/5)^0 = (3/5)^{10} = 0.006046618.$$

- (d) Find  $P(\text{Fewer than half of the visitors enter through turnstiles C,D, and E})$

We have

$$P(Y < 5) = P(Y \leq 4) = \sum_{y=0}^4 \binom{10}{y} (3/5)^y (1 - 3/5)^{10-y} = \text{pbinom}(4, 10, 3/5) = 0.1662386.$$

2. Your company makes bars of soap sold with labels claiming that they weigh 100 grams. A consumer advocacy group has said they will sue you if more than 5% of your bars weigh less than the stated 100 grams.

- (a) If your current production process makes bars with weights having the Normal( $\mu = 105, \sigma^2 = 5^2$ ) distribution, find the percentage of bars which will weigh less than 100 grams.

Let  $X \sim \text{Normal}(105, 25)$ . Then

$$P(X \leq 100) = P(Z \leq (100 - 105)/5) = P(Z \leq -1) = \text{pnorm}(-1) = 0.1586553.$$

- (b) You decide you must alter your production process: If your process makes bars with weights having the Normal( $\mu = 105, \sigma^2$ ) distribution, find the largest value of  $\sigma$  such that no more than 5% of your bars will weigh less than the stated 100 grams.

Let  $X \sim \text{Normal}(105, \sigma^2)$  and set  $P(X \leq 100) = 0.05$ . Then we have

$$\begin{aligned} P(X \leq 100) = 0.05 &\iff P(Z \leq (100 - 105)/\sigma) = 0.05 \\ &\iff q_{0.05}^Z = (100 - 105)/\sigma \\ &\iff \sigma = (100 - 105)/q_{0.05}^Z. \end{aligned}$$

Plugging in  $q_{0.05}^Z = \text{qnorm}(0.05) = -1.644854$ , we obtain

$$\sigma = 3.039784.$$

- (c) Your process engineer claims it is impossible to make  $\sigma$  any smaller than 4. If your process makes bars with weights having the Normal( $\mu, \sigma^2 = 4^2$ ) distribution, find the smallest  $\mu$  such that no more than 5% of your bars will weigh less than the stated 100 grams.

Let  $X \sim \text{Normal}(\mu, \sigma^2 = 4^2)$  and set  $P(X \leq 100) = 0.05$ . Then

$$\begin{aligned} P(X \leq 100) = 0.05 &\iff P(Z \leq (100 - \mu)/4) = 0.05 \\ &\iff q_{0.05}^Z = (100 - \mu)/4 \\ &\iff \mu = 100 - 4q_{0.05}^Z. \end{aligned}$$

Plugging in  $q_{0.05}^Z = \text{qnorm}(0.05) = -1.644854$ , we obtain

$$\mu = 106.5794.$$

3. Suppose a population has values that are Normally distributed. Then:

- (a) How many standard deviations should you go above and below the mean to in order to capture 1/3 of the values?

The middle 1/3 of the values lie between the 1/3 and the 2/3 quantiles. The 2/3 quantile of the standard Normal distribution is  $q_{2/3}^Z = 0.4307$ . So if we go 0.4307 standard deviations

above and below the mean, we capture 1/3 of the values.

- (b) The top 10% of the values are all at least how many standard deviations above the mean?

The 0.90 quantile of the standard Normal distribution is  $q_{0.90}^Z = 1.2820$ , so the top 10% of the values are all at least 1.2820 standard deviations above the mean.

- (c) What proportion of the values lie within 3 standard deviations of the mean?

This is given by  $P(-3 < Z < 3)$ , where  $Z \sim \text{Normal}(0, 1)$ . We have  $P(-3 < Z < 3) = 0.9973$ .

- (d) What is the distance between the median and the 75th percentile in terms of the number of standard deviations?

The 0.75 quantile of the standard Normal distribution is  $q_{0.75}^Z = 0.6745$ , and the median is equal to 0. So the 75th percentile lies 0.6745 standard deviations above the median.

- (e) At what percentile is a value that is 2 standard deviations above the mean?

This is given by  $P(Z < 2)$ , where  $Z \sim \text{Normal}(0, 1)$ . This is equal to 0.9772, so a value that is 2 standard deviations above the mean is at the 97.72 percentile.

- (f) The bottom 5% of values lie at least how many standard deviations below the mean?

The 0.05 quantile of the standard Normal distribution, which is  $q_{0.05}^Z = -1.645$ , so the bottom 5% of values lie at least 1.645 standard deviations below the mean.

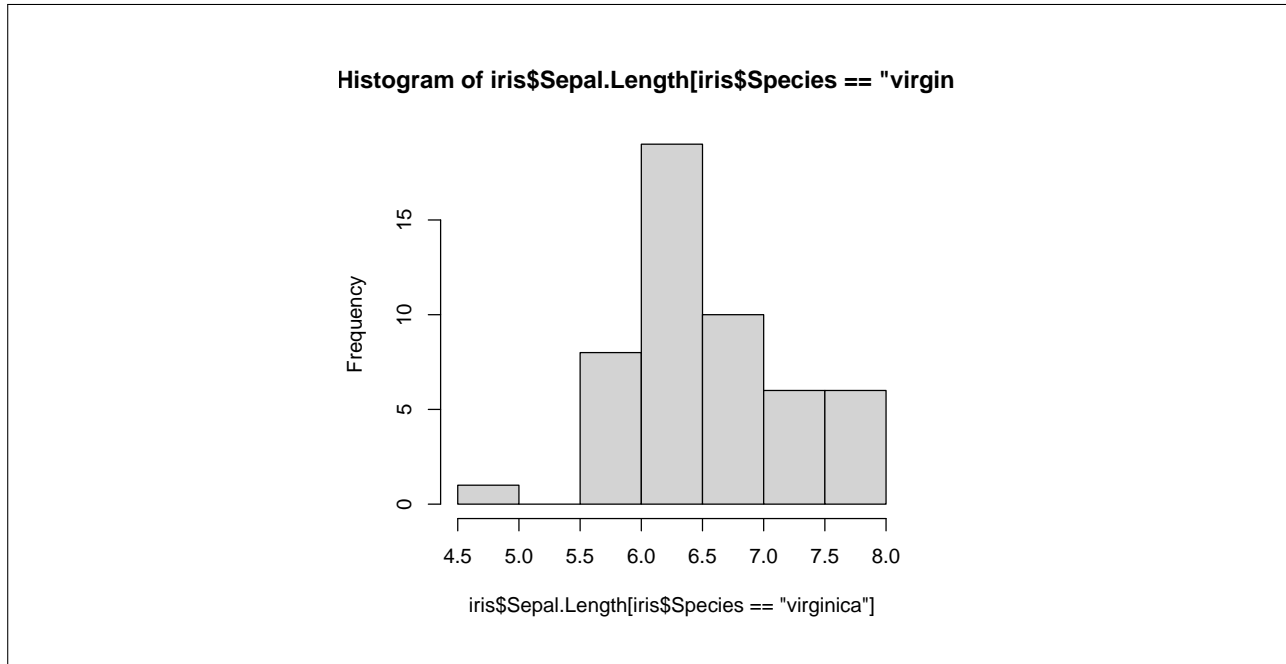
4. Open R and enter `data()` into the console. A window opens showing you a bunch of data sets. You may close this window. This command just imported these standard R datasets into the workspace so you can play with them. Now type `iris` and press enter. You will see a data set printed out. Find a description of this data set at [https://en.wikipedia.org/wiki/Iris\\_flower\\_data\\_set](https://en.wikipedia.org/wiki/Iris_flower_data_set). Now type the command

```
hist(iris$Sepal.Length[iris$Species=="virginica"])
```

You can increase the number of bars in the histogram with

```
hist(iris$Sepal.Length[iris$Species=="virginica"], breaks=20)
```

- (a) Choose one of these plots to hand in. It doesn't matter which one.



- (b) Do you think that the sepal length of virginica irises follows a Normal distribution? Say why or why not.

There is no wrong answer for this one; we can't really be conclusive when we only have a histogram.

- (c) Compute the mean  $\bar{X}$  and the sample standard deviation  $s$  for the this species of iris. I suggest storing the sepal lengths in a variable:

```
x <- iris$Sepal.Length[iris$Species=="virginica"]
```

Then to compute the mean  $\bar{X}$ , you can simply type `mean(x)` and for the standard deviation, you can type `sd(x)`.

We get  $\bar{X} = 6.588$  and  $s = 0.6358796$ .

- (d) What proportion of the sepal lengths of virginica irises are less than 6.25? Hint: You can type `mean(x < 6.25)`. Note that if you type `x < 6.25`, you will get a vector of 0s and 1s, a 1 whenever  $x < 6.25$  is true. Taking the mean of this vector is like counting the number of 1s and dividing by  $n$ , which is the same as computing a proportion.

We get 0.26, so that 26% of the sepal lengths are less than 6.25.

- (e) If the sepal lengths of virginica irises followed a Normal distribution with a mean  $\mu$  equal to `mean(x)` and a standard deviation  $\sigma$  equal to `sd(x)`, then what proportion of sepal lengths would you expect to fall below 6.25?

We have

$$P(X < 6.25) = P((X - 6.588)/0.6358796 < (6.25 - 6.588)/0.6358796) = P(Z < -0.5315472),$$

where  $Z$  has the standard Normal distribution. We get

$$P(Z < -0.5315472) = \text{pnorm}(-0.5315472) = 0.2975198.$$

- (f) What is the 0.90 quantile of the 50 sepal lengths of the virginica irises? Hint: Sort the values using `sort(x)` and take the 45th value. This is the value exceeded by only 10% of the sample values.

The 0.90 quantile is 7.6.

- (g) What is the 0.90 quantile of a Normal distribution with mean  $\mu$  equal to `mean(x)` and a standard deviation  $\sigma$  equal to `sd(x)`?

We have

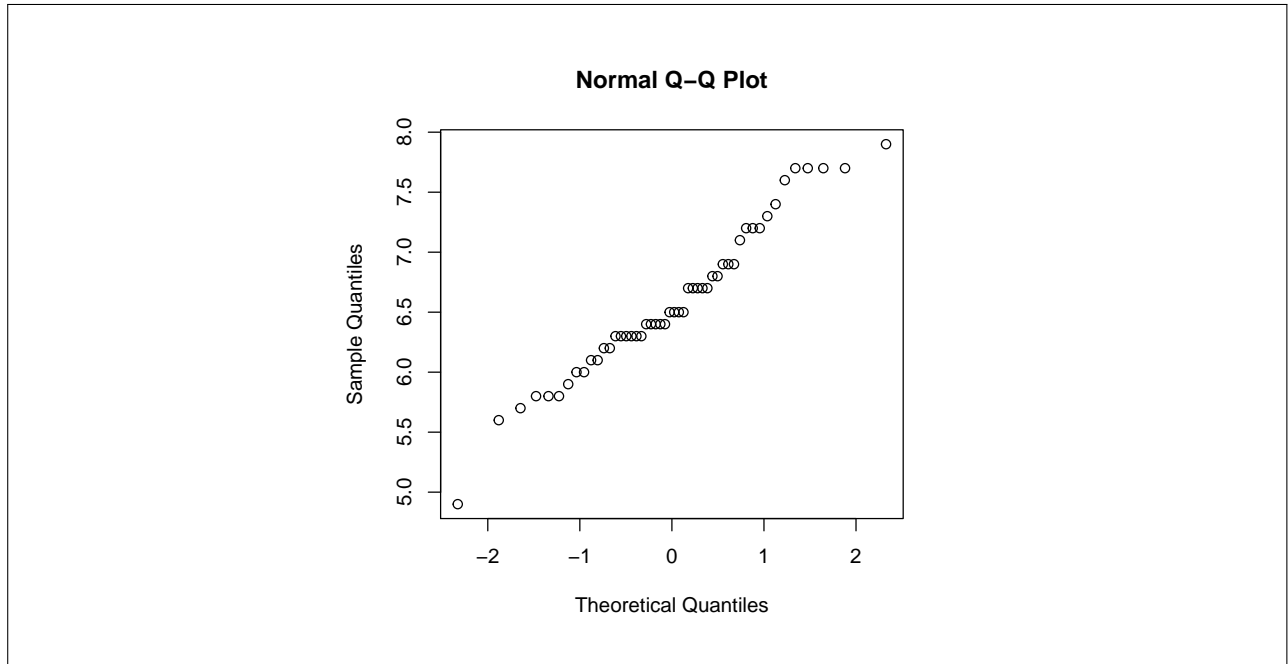
$$6.588 + 0.6358796(1.281552) = 7.402913.$$

- (h) Would the answers from parts (d)–(g) support or not support the claim that the distribution of sepal lengths of virginica irises is Normal?

The answers seem to suggest that the data come from a Normal distribution. However, this is not a rigorous statistical test.

- (i) Imagine doing parts (f) and (g) for many different quantiles, not just the 0.90 quantile. If the population distribution of sepal lengths of virginica irises is Normal, then the quantiles of the sample should agree with the quantiles of a Normal distribution. We can perform a visual test of this by producing a Normal Quantile-Quantile or Normal QQ plot. This plots many quantiles of the sample against the corresponding quantiles of the standard Normal distribution. *If the points in the Normal QQ plot fall more or less on a straight line*, then we say that the distribution is Normal. If the points do not fall on a straight line, we say that the distribution is not Normal.

Make a Normal QQ plot of the 50 sepal lengths of the virginica iris by using the command `qqnorm(x)`. Turn in this plot.



- (j) Do you think that the distribution of sepal lengths of the virginica iris follow a Normal distribution? Why or why not?

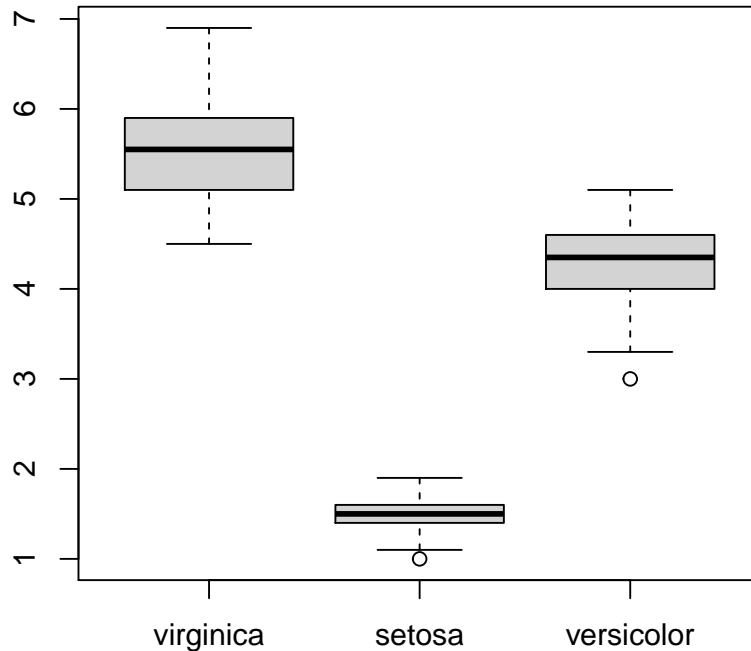
It is just important to argue correctly: points along a line are supportive of Normality, points not along a line are not supportive of Normality. The points may look different to different people. This suggests the need for a more rigorous test.

5. The purpose of this question is to make you familiar with boxplots. Enter the following commands to store the petal lengths of the three species of irises in the objects `vir.pl`, `set.pl`, and `ver.pl`:

```
vir.pl <- iris$Petal.Length[iris$Species == "virginica"]
set.pl <- iris$Petal.Length[iris$Species == "setosa"]
ver.pl <- iris$Petal.Length[iris$Species == "versicolor"]
```

Now use the following command to create a plot which has three boxplots side by side which are labeled with the names of the species.

```
boxplot(vir.pl,set.pl,ver.pl, names=c("virginica","setosa","versicolor"))
```



(a) For which species does the sample contain an outlying petal length value?

For setosa and versicolor.

(b) Which species appears to have the greatest average petal length?

Virginica.

(c) Do the petal lengths look symmetrically distributed or skewed?

They appear to be quite symmetric.

(d) Order the species according to what you believe the sample variances  $s^2$  are: which has the smallest, the second-smallest, and the largest sample variance (do this by just looking at the plot)?

Setosa, Versicolor, Virginica, by the widths of the rectangle parts of the boxplots.

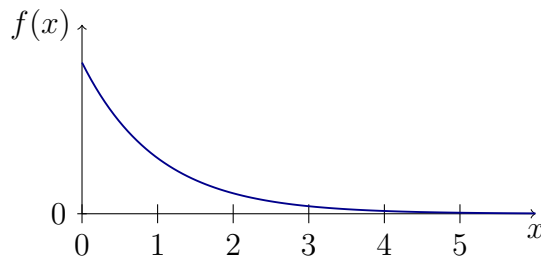
- (e) Now compute the sample variances  $s^2$  of the petal lengths for each species using the commands `var(vir.pl)`, `var(set.pl)`, and `var(ver.pl)`.

We get `var(vir.pl) = 0.3045878`, `var(set.pl) = 0.03015918`, and `var(ver.pl) = 0.2208163`.

- (f) If you have two boxplots, how do you judge which one (probably) has the larger sample variance  $s^2$ ?

A wider rectangle part and longer “whiskers” are indicative of a greater sample variance.

6. The purpose of this exercise is to show you what is meant by the terms “right-skewed” and “left-skewed” and what histograms and boxplots look like for right- and left-skewed data. Use R to generate 1000 random values of a random variable with a probability distribution having a probability density function which looks like



by using the command

```
simulated.data <- rexp(1000,.01)
```

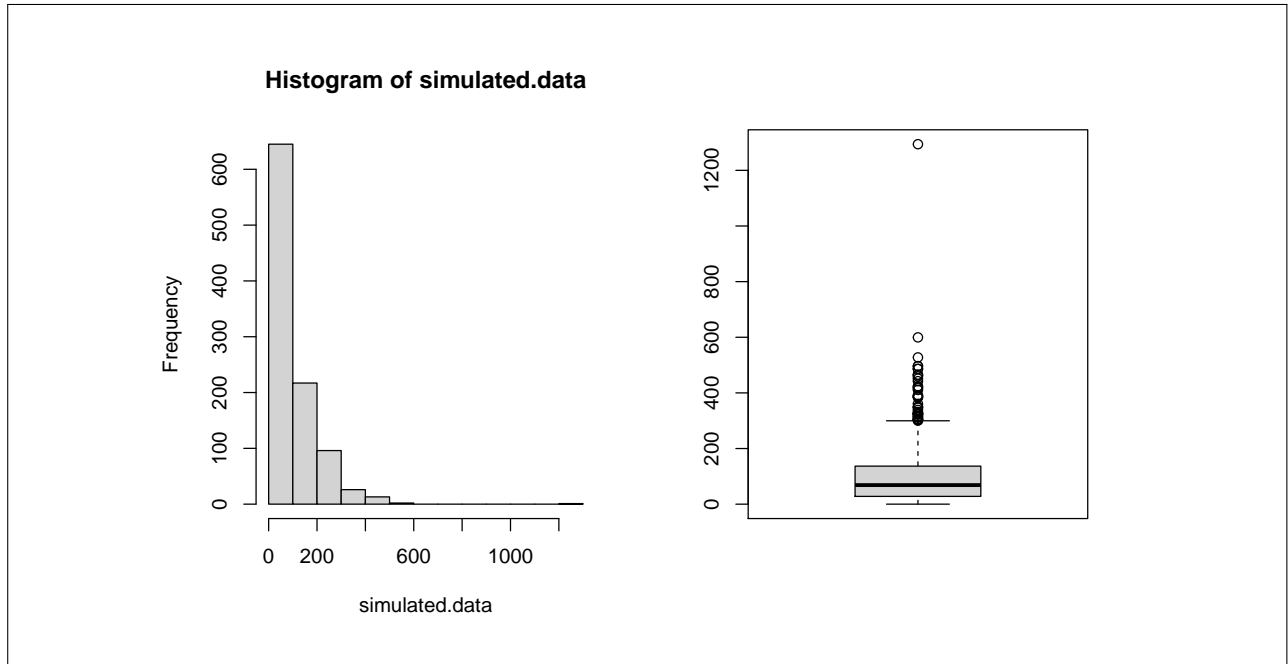
(`rexp()` generates random values from a distribution called the exponential distribution, about which we will later learn more. You may type `?rexp` into the console to open the help page for the function `rexp()` if you like).

- (a) Make a figure with two panels, one showing the histogram and the other showing a boxplot of the 1000 random values:

```
par(mfrow=c(1,2)) # this command puts the next two plots side by side
hist(simulated.data)
boxplot(simulated.data)
```

Turn in this two-paneled figure.





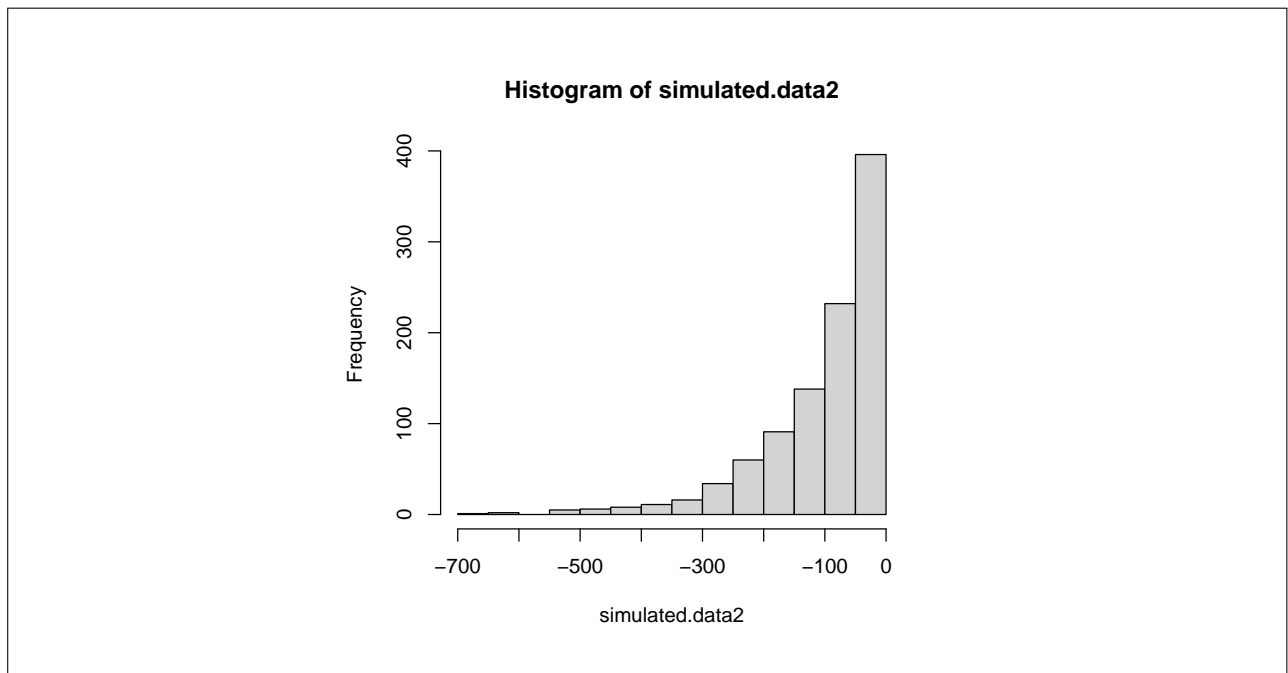
(b) Now use the command

```
simulated.data2 <- - rexp(1000,.01)
```

(notice the minus sign!) to generate some more data and make a histogram with the command

```
hist(simulated.data2)
```

Turn in this plot.



(c) Sketch, by hand, what you think the boxplot would look like for `simulated.data2`.

This plot will be like the mirror image of the boxplot from part (a).

(d) Which data, `simulated.data` or `simulated.data2` is right-skewed and which is left-skewed (For a review on skewness look at [these slides](#), pages 8 and 9)?

`simulated.data` is right-skewed and `simulated.data2` is left-skewed.