Summary statistics

Put your name here
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Introduction

In this module, we'll learn about summary statistics, which are numerical summaries calculated from data.

Instructions

Presumably, you have already created a new project and downloaded this file into it. Please knit the document and work back and forth between this R Markdown file and the PDF output as you work through this module.

When you are finished with the assignment, knit to PDF one last time, proofread the PDF file **carefully**, export the PDF file to your computer, and then submit your assignment.

Sometimes you will be asked to add your own R code. That will appear in this document as a code chunk with a request for you to add your own code, like so:

```
## Add code here to [do some task]...
```

Be sure to remove the line ## Add code here to [do some task]... when you have added your own code. Sometimes you will be asked to type up your thoughts. That will appear in the document as follows:

Please write up your answer here.

Again, please be sure to remove the line "Please write up your answer here" when you have written up your answer. In these areas of the assignment, please use contextually meaningful full sentences/paragraphs (unless otherwise indicated) and proper spelling, grammar, punctuation, etc. This is not R code, but rather a free response section where you talk about your analysis and conclusions. If you need to use some R code as well, you can use inline R code inside the block between \begin{answer} and \end{answer}, or if you need an R code chunk, please go outside the answer block and start a new code chunk.

Load Packages

We load the mosaic package as well as the MASS package so that we can work with the birth weight data.

```
library(mosaic)
library(MASS)
```

Mean and standard deviance

Summary statistics come in two general flavors: measures of center and measures of spread.

The first pair we'll consider is the mean and the standard deviation. The mean—denoted \bar{y} —of a variable y is calculated by summing all the values of the variable, and dividing by the total number of observations. In formula form, this is

$$\bar{y} = \frac{\sum y}{n}$$
.

This is a measure of center since it estimates the "middle" of a set of numbers.

It is calculated in R using the mean command. For example, if we want to calculate the mean weight in pounds of the mother at the last menstual period, we type the following:

mean(birthwt\$lwt)

[1] 129.8148

The corresponding measure of spread is the $standard\ deviation$. Usually this is called s and is calculated using a much more complicated formula:

$$s = \sqrt{\frac{\sum (y - \bar{y})^2}{n - 1}}.$$

This is a measure of spread because the $(y - \bar{y})$ term measures the how far away each data point is from the mean.

In R, this is calculated with the sd command.

sd(birthwt\$lwt)

[1] 30.57938

The mean and the standard deviation should always be reported together.

Another related measurement is the *variance*, but this is nothing more than the standard deviation squared:

$$s^2 = \frac{\sum (y - \bar{y})^2}{n - 1}.$$

(Compare this formula to the one for the standard devation. Nothing has changed except for the removal of the square root.) We rarely use the variance in an introductory stats class because it's not as interpretable as the standard deviation.¹ If you need to do this in R, the command is var.

var(birthwt\$lwt)

[1] 935.0985

(You can check and see that the number above really is just 30.5793804 squared.)

Median and IQR

Another choice for measuring the center and spread of a data set is the median and the IQR. The median is just the middle value if the list of values is ordered. In R, it is calculated using the median command.

¹The main reason for this is units. If the data variable is the mother's weight in pounds, then both the mean and the standard deviation are also reported in pounds. The variance has units of "pounds squared", but what does that even mean?

median(birthwt\$lwt)

```
## [1] 121
```

You can check that this is correct. If we print out the entire lwt variable, you can see all 189 values, and if we're clever about it, we can see them in order.

sort(birthwt\$lwt)

```
##
                90
                  90
                     90
                          92
                             94
                               95
                                  95
                                     95
   Γ11
           85
             89
                        91
                                       95
                                          95
##
        98 100 100 100 100 100 101 102 102 103 103 103 105 105 105 105
  ##
  [103] 124 125 125 125 127 128 128 129 130 130 130 130 130 130 130 130 130
 [120] 130 130 130 130 131 132 132 132 133 133 134 134 134 135 135 135 135
 [137] 137 138 138 140 140 140 141 142 142 147 147 148 150 150 150 150 150
 [154] 153 154 154 155 155 155 158 158 160 160 165 167 168 169 169 170 170
## [171] 170 170 175 182 184 185 186 187 187 189 190 190 200 202 215 229 235
 [188] 241 250
```

Exercise

If there are 189 mothers in this data set, in which position in the list will the median value appear? (Be careful! You can't just divide 189 by 2!) Verify that the median value 121 does appear in the position you calcuated above.

Please write up your answer here.

Calculating the *interquartile range*—or IQR—requires first the calculation of the first and third quartiles, denoted Q1 and Q3. If the median is the 50% mark in the sorted data, the first and third quartiles are the 25% and the 75% marks, respectively.² Then the IQR is just

$$IQR = Q3 - Q1$$

In R, you can get the IQR by using—are you ready for this?—the IQR command.

IQR(birthwt\$lwt)

[1] 30

The IQR is a measure of spread because the distance between Q1 and Q3 measures the span of the "middle 50%" of the data.

²One way to compute these is to calculate the median of the lower and upper halves of the data separately. Then again, it's hard to know how to split the data set into halves if there are an odd number of observations. There are many different methods for computing percentiles in general, but you don't need to worry too much about the particular implementation in R.

Exercise

A general function for computing any percentile in R is the quantile function. For example, since Q1 is the 25th percentile, you can compute it as follows:

```
Q1 <- quantile(birthwt$lwt, 0.25)
unname(Q1)
```

```
## [1] 110
```

(The unname command just helps clean up the output here. Don't worry too much about it.)

Now compute Q3. Also check that the IQR calculated above matches the value you get from subtracting Q3 minus Q1.

```
## Add code here to compute and print out Q3.
## You should use the unname command as in the above chunk.
```

```
## Add code here to compute Q3 - Q1
```

The median and the IQR should always be reported together.

Five-number summary

A five-number summary is the minimum, Q1, median, Q3, and maximum of a set of numbers.

The summary command in R gives you the five-number summary, and throws the mean in for good measure.

```
summary(birthwt$lwt)
```

```
## Min. 1st Qu. Median Mean 3rd Qu. Max.
## 80.0 110.0 121.0 129.8 140.0 250.0
```

You can, of course, isolate the various pieces of this. You already know most of the commands below.

```
min(birthwt$lwt)
```

```
## [1] 80
```

```
median(birthwt$lwt)
```

[1] 121

```
max(birthwt$lwt)
```

[1] 250

```
mean(birthwt$lwt)
```

[1] 129.8148

quantile(birthwt\$lwt)

```
## 0% 25% 50% 75% 100%
## 80 110 121 140 250
```

Exercise

What is the difference between the way quantile was used in a previous exercise versus the way it was used here? How did that change the output?

Please write up your answer here.

The mosaic package also has a summary command called favstats that has a little more information, including the standard deviation, the sample size, and a count of any cases that are missing data. This command is much more useful than the standard summary command.

favstats(birthwt\$lwt)

```
## min Q1 median Q3 max mean sd n missing
## 80 110 121 140 250 129.8148 30.57938 189 0
```

Also, don't forget about the trick for using R commands inline. If you need to mention a statistic in the middle of a sentence, there is no need to break the sentence and display a code chunk. Be sure you're looking at the R Markdown document to note that the numbers in the next sentence are not manually entered, but are calculated on the fly:

There are 189 births represented in this data and the median weight of the women as of their last menstrual period is 121 pounds.

Your turn

Type a full sentence using inline R code (as above) summarizing the minimum and maximum baby weights (in grams) in our data set.

Please write up your answer here.

Summary statistics by group

Using base R, it's not so easy to get summary statistics for each group separately. Fortunately, the mosaic package comes to the rescuem, allowing for more flexibility. For example:

favstats(~ birthwt\$lwt | birthwt\$race)

```
## birthwt$race min Q1 median Q3 max mean sd n missing
## 1 1 90 112 129.5 143.25 235 132.0521 29.09381 96 0
## 2 2 98 120 129.0 179.00 241 146.8077 39.63939 26 0
## 3 3 80 105 119.0 130.00 250 120.0149 25.13026 67 0
```

The notation is a little weird. Don't worry about the tilde for now. Just learn that it needs to be there. The important part is the birthwt\$lwt | birthwt\$race. This says, "Look at the numerical variable birthwt\$lwt broken down by birthwt\$race." Indeed, the output has three lines, one for each race in the data. This extended notation works for lots of commands, like mean, sd median, IQR, quantile, etc. as long as the mosaic package is loaded.

Your turn

Choose one numerical variable and one categorical variable from the birthwt data set. Find the five-number summary of your numerical variable grouped by your categorical variable.

Add code here to compute the five-number summary of your ## numerical variable grouped by your categorical variable.

Conclusion

. . .