PSTAT 5LS Lab 2

Professor Miller

July 6, 2023

Section 1

Learning Objectives

R Learning Objectives

- Learn how to make a frequency table and a two-way frequency table in R
- Learn how to make a bar chart in R
- Learn how to find the five-number summary of a variable, and find a specific numeric summary (statistic) in R
- Learn how to make a histogram in R
- 5 Learn how to make a boxplot in R
- Learn how to make side-by-side boxplots in R

Statistical Learning Objectives

- 1 Understand when to make a bar chart.
- Understand when to make a frequency/two-way frequency table.
- Be able to use these graphical summaries to discuss data.
- Understand when to make a bar chart versus a boxplot or a histogram.
- Understand when to make a side-by-side boxplot and how to use this type of comparison
- Understand when to make a frequency/two-way frequency table versus a number summary.
- Be able to use these graphical and numerical summaries to discuss data.

Functions covered in this lab

- 1 table()
- ② barplot()
- 3 summary()
- 4 hist()
- min(), mean(), median(), max(), sd(), IQR()
- 6 boxplot()

Section 2

Lab Tutorial

Penguins data set

We're back to hanging out with our penguin friends.

```
penguins <- read.csv("penguins.csv", stringsAsFactors = TRUE)</pre>
```

Notice that we added the additional argument stringsAsFactors = TRUE to the read.csv() function. This is important because the R default is to NOT do this. We want to be able to distinguish between something called a *string*, which is a phrase or word, and a **factor**, which represents the levels of a categorical variable. Basically, by setting stringsAsFactors = TRUE, we are letting R know to expect that any words or phrases in the data actually relate to categorical variables.

Go ahead and run the loadPenguins chunk of your lab2-notes.Rmd markdown file, and verify that the penguins data is in your environment in the top right corner of your RStudio project.

Frequency Tables

Let's start with the species variable. Is this a categorical or numeric variable? How do you know?

To make a frequency table of a categorical variable, we use the table() function. Try this code out on your lab2-notes.Rmd file in the tryit1 code chunk of your notes file. Recall that code goes in the middle of the chunk and not on the first or last line of the chunk.

```
table(penguins$species)
```

```
##
```

##	Adelie	Chinstrap	Gentoo
##	146	68	119

Frequency Tables continued

So, there are 119 Gentoo penguins in the data.

Notice that inside the table function, we have something that looks a little weird. We wrote penguins\$species. This is how we tell R to use the species variable **inside the data frame** penguins. The dollar sign (\$) tells R to look inside the data frame penguins for the column called species.

Frequency Table Common Error

It's very important that you tell R $\it which\ data\ frame$ the variable you're interested in is from. Let's see what happens when we don't:

```
table(species)
```

Error in eval(expr, envir, enclos): object 'species' not for

Notice that we get an error message here, stating that the "object 'species' not found".

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Two-Way Frequency Tables (Contingency Tables)

We can also make "two-way" frequency tables (sometimes called "contingency tables") to summarize counts for two categorical variables. Try this in the tryit2 code chunk in your notes file.

```
table(penguins$species, penguins$island)
```

```
## Biscoe Dream Torgersen
## Adelie 44 55 47
## Chinstrap 0 68 0
## Gentoo 119 0 0
```

Data is **R**eally **C**ool, so the first variable you give to table() is in the **r**ows of the table, and the second is in the **c**olumns.

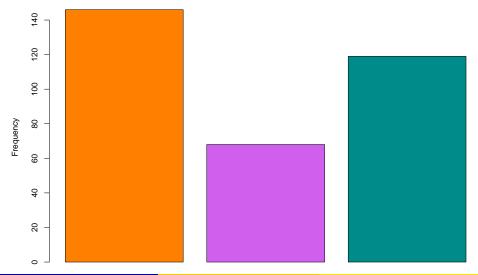
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Bar Charts in R

Let's explore our penguins data by making a plot that will help us visualize a categorical variable. We'll start by looking at the number of penguins observed of each species. We have provided to you the code in the tryit3 code chunk - all you need to do is run it!

Bar Chart of species in penguins

Bar Chart of Number of Penguins of Each Species
Observed



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Bar Chart Code for species in penguins

Bar Charts Code for species in penguins continued

Notice that we included the table from earlier in our code!

Also notice that we included some arguments, such as

- xlab, the label on the x (or horizontal) axis
- ylab, the label on the y (or vertical) axis
- main, the title of the graph, and
- a nice way to add some fun to an otherwise boring plot col for colors. We have three species of penguins, so we picked out three colors that we thought best represented each species. There are lots and lots of colors that you can try. There is a color palette "cheat sheet" from UCSB's own National Center for Ecological Analysis and Synthesis posted on Canvas if you want to check it out.

How to Find Help in R

R has built-in "documentation" for every function. If you want to find that documentation, you can Google it, but that takes too long. So it's better to use R's built in help! In the R console, just type a question mark? followed by the name of the function you want help with, then hit enter. For example, ?barplot will bring up the documentation for the barplot() function.

Let's try this in the tryit4 code chunk in your notes file.

At the end of the help file you may find an example of how to use the function. These examples are generally super helpful! You can directly run them using the example() function - e.g., example(barplot).

The most useful feature of help in R is a list of a function's arguments and a quick explanation of what each argument does. You may not be able to fully understand some of the terms in the documentation just yet, but try it out and your TA will be able to help!

Numerical Summaries

Let's start with the flipper_length_mm variable. Is this a categorical or quantitative variable? How do you know?

We can use R to summarize data numerically. We'll use the summary() function to do that for a given variable. Here, we'll summarize the flipper_length_mm variable, which is the length of the penguins' flippers (in millimeters). Try this code out on in the tryit5 code chunk.

summary(penguins\$flipper_length_mm)

```
## Min. 1st Qu. Median Mean 3rd Qu. Max.
## 172 190 197 201 213 231
```

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Numerical Summaries

[1] 14.01577

You might have noticed that the summary() function doesn't give you the standard deviation of the variable. To get the standard deviation, use the sd() function in addition to the summary() function.

Summarize the flipper_length_mm variable again adding the code to get the standard deviation as well. Try this code out on in the tryit6 code chunk.

```
summary(penguins$flipper_length_mm)

## Min. 1st Qu. Median Mean 3rd Qu. Max.
## 172 190 197 201 213 231

sd(penguins$flipper_length_mm)
```

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Number Summaries continued

You can always get just the one numerical summary you're looking for using the function for that specific summary. Try these out in the tryit7 code chunk.

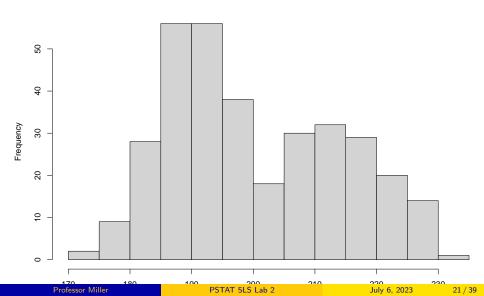
```
min(penguins$flipper_length_mm)
mean(penguins$flipper_length_mm)
median(penguins$flipper_length_mm)
max(penguins$flipper_length_mm)
sd(penguins$flipper_length_mm)
IQR(penguins$flipper_length_mm)
```

Histograms in R

One type of graphical display for a quantitative variable is a histogram. Histograms in R are also pretty easy – you just use the hist() function.

Histogram of Flipper Length of Penguins

Histogram of penguins\$flipper_length_mm



Don't Forget Labels and Titles!

hist(penguins\$flipper_length_mm)

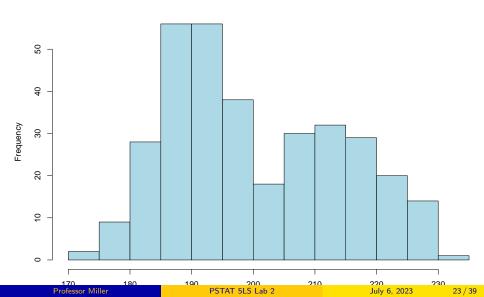
So here we've got a histogram. Notice that we didn't provide the main, xlab, and ylab arguments that we'd normally use for a plot title and an axis label, but R still gave us a title and labels. This is nice, but the labels are *horrible*: nobody (other than you) knows what penguin\$flipper_length_mm means, so we don't want to use that as a title or axis label.

The moral of the story is to always provide main, xlab, and ylab arguments when making a plot!

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That Same Histogram with Labels and a Title

Histogram of Penguin Flipper Length



Histogram Code

Try this code out in the tryit8 code chunk. Watch out for the dreaded typos! If you get an error message, try to debug it yourself before asking for help!

```
hist(penguins$flipper_length_mm,
    main = "Histogram of Penguin Flipper Length",
    xlab = "Flipper Length (mm)",
    col = "lightblue")
```

Recall from lecture that we describe distributions by addressing four aspects:

- Shape (number of modes + symmetry or lack thereof)
- Center
- Spread/Variability
- Outliers

A handy mnemonic to remember what to comment about is **SOCS**:

Shape Outliers Center Spread

Note: Be sure to mention whether there are or are not outliers. Not saying anything doesn't let us know that you know to check for outliers.

Describe the distribution of flipper lengths.

Do you think that the mean is the best measure of center? Why or why not?

Describe the spread (variability) of the distribution of flipper length.

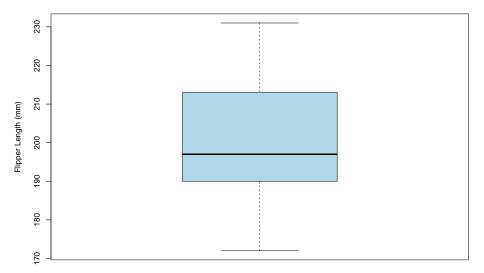
Are there any outliers or other unusual features that you'd like to mention about the distribution of flipper length?

Boxplots in R

Another type of graphical display for a quantitative variable is a boxplot. The command for making a boxplot in R is pretty simple: it's just boxplot(). To make a boxplot of a single variable, just give R the name of the data set, a dollar sign (\$), then the name of the variable. Also provide the arguments main and ylab for a plot title and y-axis label.

Boxplot of the Flipper Length of Penguins

Boxplot of Penguin Flipper Length



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Code for the Boxplot of the Flipper Length of Penguins

Try this code out yourself in the tryit9 code chunk. Watch out for the dreaded typos! If you get an error message, try to debug it yourself before asking for help!

Describing Boxplots

True or False:

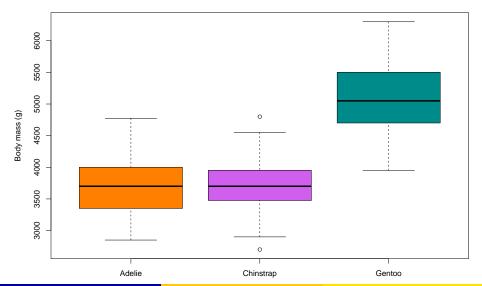
The boxplot of flipper lengths appears to be unimodal and symmetric.

Side-by-side Boxplots

Sometimes we're interested in comparing two or more groups using "side-by-side" boxplots. We can compare the different species of penguins' body masses in this way, still using the boxplot function. We have provided the code in the tryit10 code chunk.

Side-by-side Boxplots Continued

Boxplots of Penguin Body Mass by Species



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Penguin Body Mass By Species

Does it appear that a penguin's body mass is affected by its species, for the penguins in Palmer Archipelago? Why or why not?

Another Way to Code Side-by-side Boxplots

Another way to get the same side-by-side boxplots the to specify the variables themselves and adding in the code data = penguins.

It's up to you which of the two ways to specify the variables you use. We all have different things we prefer in coding. Try out some different options to find your style!

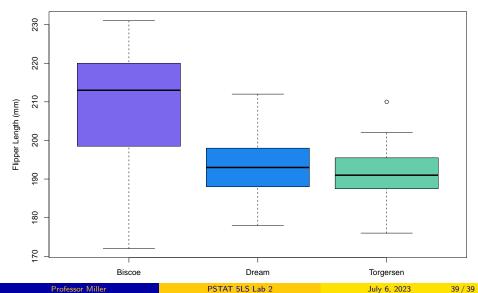
```
boxplot(body_mass_g ~ species, data = penguins,
    main = "Boxplots of Penguin Body Mass by Species",
    ylab = "Body mass (g)",
    xlab = "Species",
    col = c("darkorange1", "mediumorchid2", "darkcyan"))
```

Another Side-by-side Boxplot

Now, in the tryit11 code chunk, you will make a side-by-side boxplot of the numeric variable flipper_length_mm by the categorical variable island in the penguins data.

Another Side-by-side Boxplot

Boxplots of Penguin Flipper Length by Island



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