

Lab Session 2: Vectors and Data Types

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We expect you to review the `class 2` material, [here](#) prior to lab.

1 Introduction to Rmds

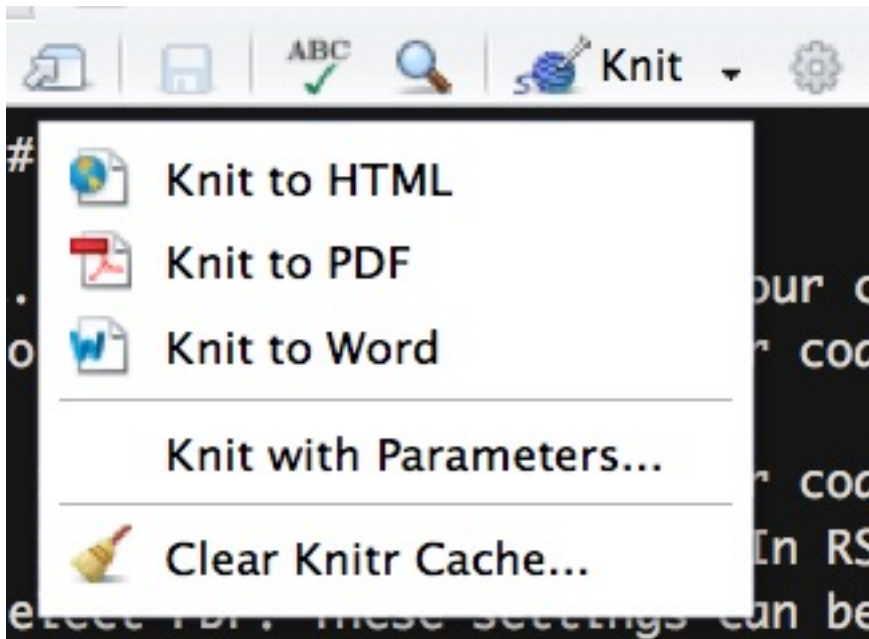
1.1 Intro to Rmds (R Markdown documents). (To be covered during QA)

1. Before getting started, run the following code in the console. This gives R the necessary tools to make pdfs from your Rmds.

```
install.packages("tinytex")
tinytex::install_tinytex()
```

Note: You never want to include code that installs packages or use `View()` in a code chunk that is evaluated in an Rmd. When knitting you will get an error or worse create unusual behavior without an error.

1. Create an Rmd file. (In RStudio's menu `File > New File > R Markdown`). Name the document and select PDF. These settings can be changed later.
2. Save the Rmd in your coding lab folder as `lab_3.Rmd`.
3. New Rmds come with some example code. Read the document and then knit to pdf by clicking the knit button. If pdf doesn't work, try html.



Pay attention to the syntax and ask your group or TA about anything you don't understand. Rmds start with meta information which provides instructions to `knitr` on how to knit. After that, there's a normal code chunk which runs, but you won't see because they of the `include=FALSE` bit at start of the code chunk.

```
1 ---
2 title: "Lab Session 1: Read and manipulate data"
3 author: "Ari Anisfeld"
4 date: "8/26/2020"
5 output: pdf_document
6 ---
7
8 ```{r setup, include=FALSE}
9 knitr::opts_chunk$set(echo = TRUE)
10 ```
```

Keep the part shown in the image above and erase the rest of the code and text in the document. This is how we start our own Rmd code.

4. Make a new code chunk by pressing `Ctrl + Alt + I` (`Cmd + Option + I` on macOS).¹ Then, add code to load the `tidyverse` in that chunk.

2 Warm-up: Vector creation.

1. Create an R script or Rmd to capture your work. Load the `tidyverse` libraries.
2. In the lecture, we covered `c()`, `:`, `rep()`, `seq()`, `rnorm()`, `runif()` among other ways to create vectors. Use each of these functions once as you create the vectors required below.
 - a. Create an integer vector from seven to seventy.
 - b. Create a numeric vector with 60 draws from the random uniform distribution
 - c. Create a character vector with the letter "x" repeated 1980 times.
 - d. Create a character vector of length 5 with the items "Nothing" "works" "unless" "you" "do". Call this vector `angelou_quote` using `<-`.
 - e. Create a numeric vector with $1e4$ draws² from a standard normal distribution.
 - f. Create an integer vector with the numbers 0, 2, 4, ... 20.
3. Run this code and explain why we get an error. (Make sure you did question 1.d above first!)

```
# make sure you followed direction in part d above.
sum(angelou_quote)
```

4. If we want `angelou_quote` to be a single string, we can use `paste0`.

```
paste0(angelou_quote, collapse = " ")
```

- a. We gave `collapse` the argument `" "` i.e. a character string that is a blank space. Try a different character string.

¹You can also type three tick marks with `{r}` and then another three tick marks.

²This is scientific notation. Try `1e4 - 1 + 1` in the console.

5. Try these lines of code using `paste0` (or its tidyverse synonym `str_c`)³.

```
paste0(angelou_quote, ".com")
paste0(angelou_quote, c("!", "!", "?", " :(", "!!!"))
```

- a. Explain to your partner what `paste0` is doing.
6. Common error alert. Run the following code and explain why it throws an error.

```
c(1, 2) + c(1 2)
```

This is an example where the error is not so helpful. I get this one a lot, because I forget to put a comma where it should be.

3 Calculating Mean and Standard Deviation with vectors

3.1 Is the coin fair?

In this exercise, we will calculate the mean of a vector of random numbers. To get started, we'll generate some fake data using built-in random⁴ sampling functions. Let's start by flipping coins.

```
(coin_flips <- sample(c("Heads", "Tails"), 10, replace = TRUE))
```

```
## [1] "Tails" "Heads" "Heads" "Heads" "Heads" "Heads" "Heads" "Heads" "Tails"
## [10] "Heads"
```

`sample()` is a function that requires two arguments.

- In the first position, we have a vector of any type. We sample *from* this vector.
- In the second position, we have `size` which is the number of items to choose.

If we want to have independent draws from our sampling vector, we say `replace = TRUE`. By default `replace` is `FALSE`.

1. We hope the following code will give us 100 independent die rolls (i.e. random numbers between 1 and 6), but we get an error. Run the code to reproduce the error.
- a. Interpret the error. I.e. why does the code fail?
- b. Adjust the code so that you simulate 100 independent die rolls.

```
die_rolls <- sample(c(1, 2, 3, 4, 5, 6), 100)
```

2. In my coin-toss simulation above, I sample from a character vector. Doing so, makes it easier to interpret the outcome, but difficult to do stuff with the results. Replace the characters with 1 and 0. Now, you'll be able to do math, but the results are more abstract. You can choose whether 1 represents heads or tails, just be consistent. Collect samples of size 10, 1000 and 1000000.⁵

```
# replace ... with suitable code
ten_flips <- ...
thousand_flips <- ...
million_flips <- ...
```

- a. What data type are your `xxx_rolls` vectors?
- b. Use `sum()` on your vectors. What does this represent?
- c. Use `length()` on your vectors to verify the vectors are the right length. What does this represent?

³tidyverse synonyms are often preferable since they have ironed out quirky behaviors. For example, try `str_c(c("bob", NA, "maya"), "@gmail.com")` vs `paste0(c("bob", NA, "maya"), "@gmail.com")`

⁴Technically, "pseudo-random", but who's asking.

⁵Note: you can use scientific notation `1e6` is short for 1 with 6 zeros.

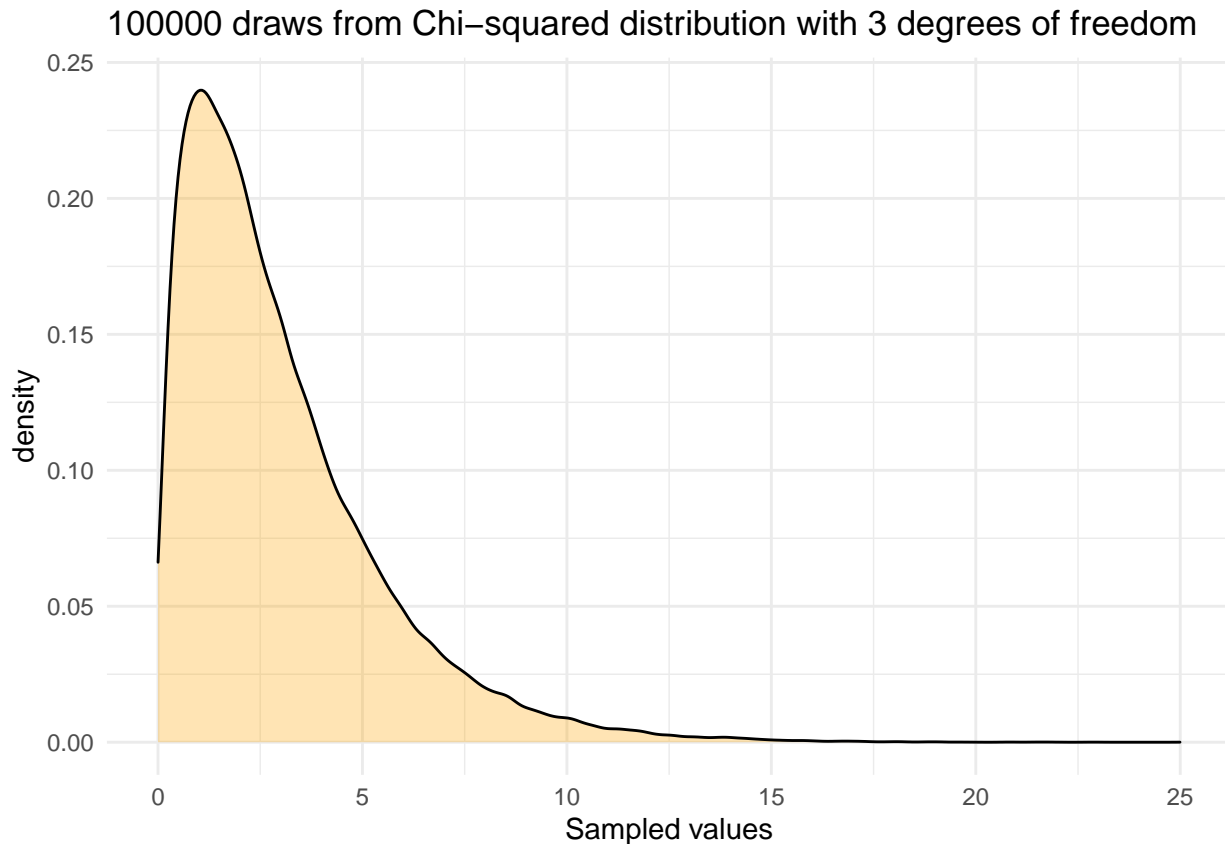
3. A fair coin assigns equal probability to heads and tails. Thus, the probability of heads or tails is 50 percent or .5. We can run experiments or simulations to see if our “coins” are fair. In particular, we can calculate an estimate of the probability of heads by computing estimated probability heads = $\hat{p}(\text{heads}) = \frac{\text{n_heads}}{\text{n_flips}}$. The estimated probability is often called \hat{p} . Use the starter code to calculate the estimated probability of heads from your `ten_flips` sample.

```
n_heads <- ...  
n_flips <- ...  
p_hat_ten <- n_heads / n_flips
```

4. Repeat the code from part 3 to find the estimated probability of heads from your `thousand_flips` sample and `million_flips` sample.⁶
 - a. Re-run all the code from parts 2 through 4 a few times. Notice that the random number generator will give a different sequence of flips each time.
 - b. What do you notice about the estimated probabilities as the sample size gets larger? (This is an example of the “Law of Large Numbers”)
5. We had you calculate the estimated probability with `sum() / length()`. R also has a function `mean()` built in. Simplify the computation for `p_hat_xxx` by using `mean()`.

3.2 A new distribution.

Now we are going to take random samples from a chi-squared distribution with 3 degrees of freedom. Do not worry about what the distribution’s name means, but be aware of that it looks something like the picture below. It’s possible—but highly unlikely—to get values up to `Inf`, which is R for infinity.



⁶In the fall, we’ll discuss ways to write this type of code more efficiently without copy paste.

We are going to calculate the mean, variance and standard deviation of the distribution using vectors in three different ways.

1. First, we'll do it "by hand". The formula for sample variance is $Var(x) = \frac{\sum(x-\bar{x})^2}{n-1}$. where
 - \bar{x} is the sample mean.
 - n is the sample size and
 - \sum means we add up

```
# fill in the ... with appropriate code.
x <- rchisq(100000, 3)

# this one should be straight forward!
# (See what we did with coin flips)
x_bar <- ...
n <- ...
# The formula in R will be exactly the same as the
# fomula in math thanks to vectorization!
# If you aren't sure the code will work the way you want
# try with a simpler x. x <- c(1, 0, 1, 1)
var_x <- sum(...) / ...
```

2. Standard deviation is the square root of Variance, i.e. $sd(x) = \sqrt{Var(x)}$. Calculate the standard deviation.⁷
3. Now, we'll check your work using built in R functions. To calculate variance use `var()`. To calculate standard deviation use `sd()`. Try them out. If you disagree with your previous results, it's most likely a coding error in the definition of `var_x`.⁸
4. Finally, we can do this in a `tibble` setting and use `summarize`. You may need to load a package.⁹ Using a `tibble` provides two services 1) the results print as an organized table. 2) We can do further `tidyverse` processing with it.

```
# replace the ... with suitable code.
tibble(x = ... ) %>%
  summarize(mean = mean(x),
            variance = ...,
            `standard deviation` = ...)
```

5. Copy your code from the previous problem, but replace `summarize` with `mutate`. Explain the result to your group.

3.3 Challenge problems

1. Run the code below. The resulting graph shows three chi-sq distributions determined by their degrees of freedom.

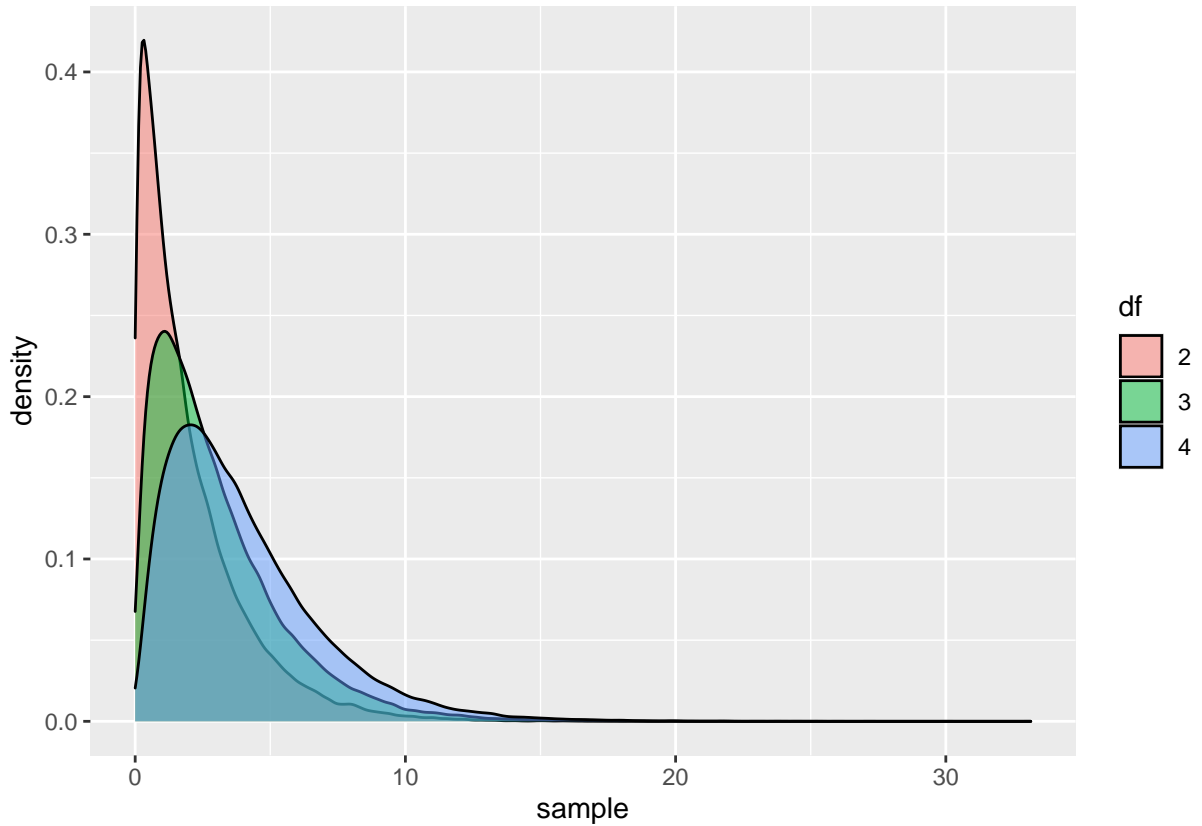
```
chi_sq_samples <-
  tibble(x = c(rchisq(100000, 1) + rchisq(100000, 1),
              rchisq(100000, 3),
              rchisq(100000, 4)),
        df = rep(c("2", "3", "4"), each = 1e5))
```

⁷Hint: we have the function `sqrt()`

⁸The most common errors are about where you put your parentheses. The second most common error is where you put the power i.e. `^`.

⁹Hint: it's the `tidyverse`.

```
chi_sq_samples %>%
  ggplot(aes(x = x, group = df, fill =df)) +
  geom_density( alpha = .5) +
  labs(fill = "df", x = "sample")
```



2. How many rows are in the tibble? Explain how the code that defines `x` and the code that defines `df` make vectors that are the right length.
3. Temporarily delete `each = (keep 1e5)`. Explain why the resulting graph looks the way it does. Make sure you put `each =` back.

Want to improve this tutorial? Report any suggestions/bugs/improvements on [here!](#) We're interested in learning from you how we can make this tutorial better.