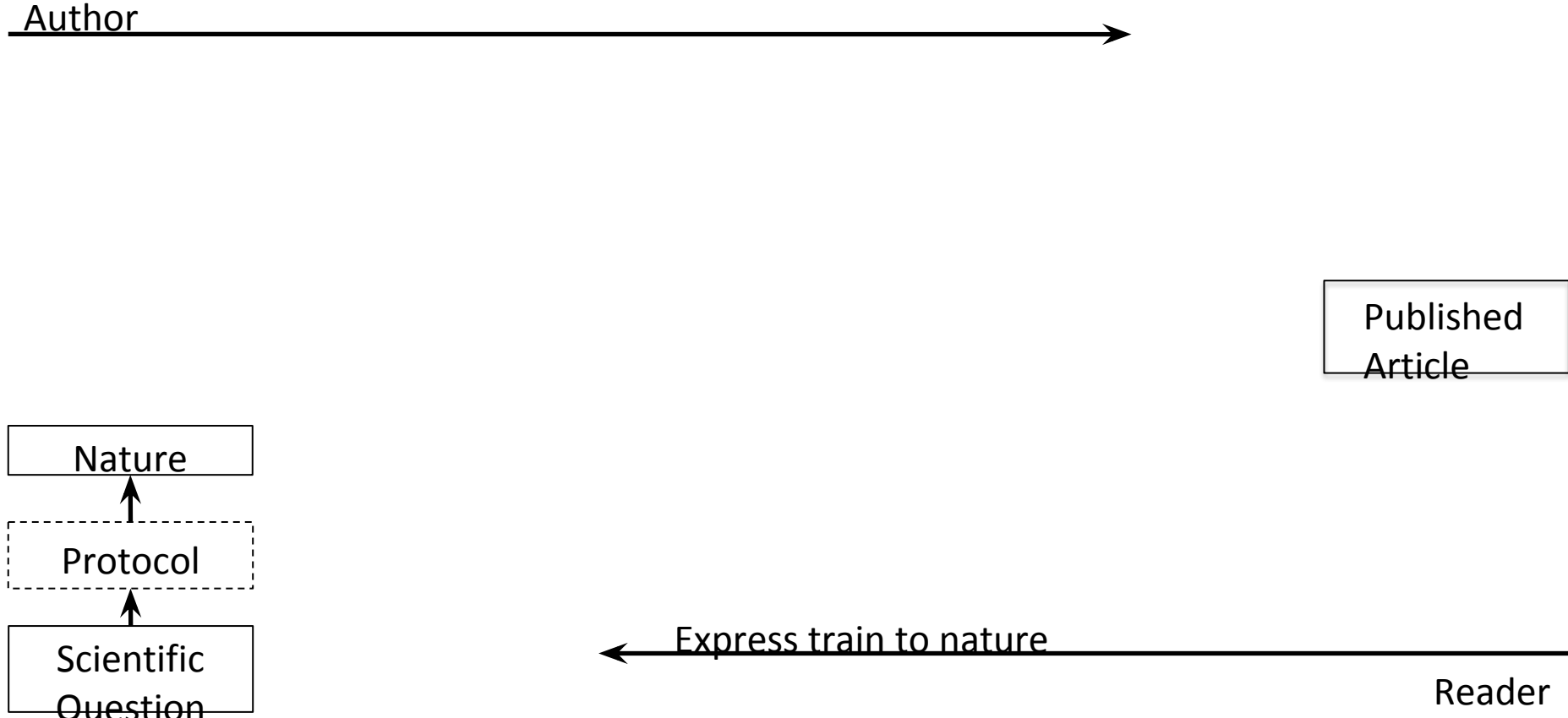


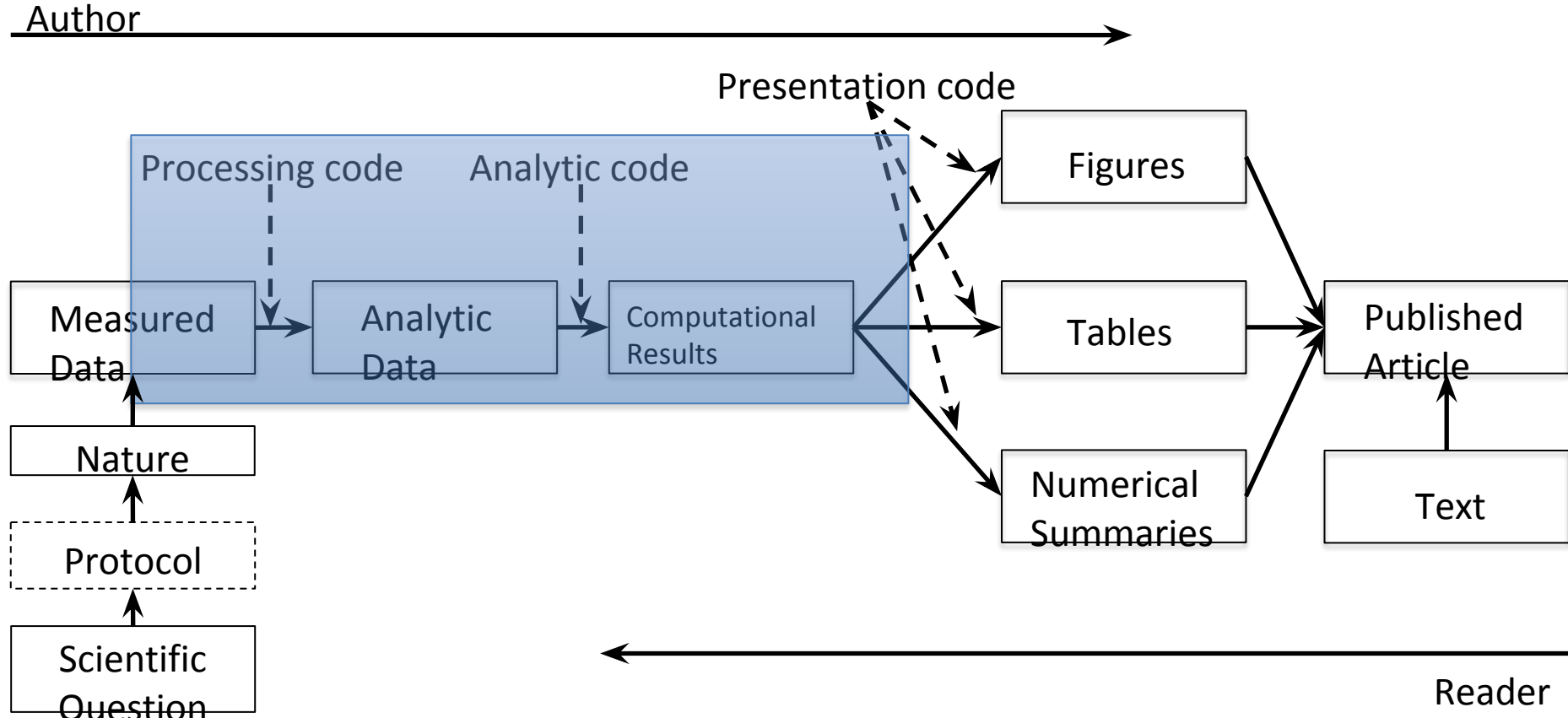
Data Wrangling in R

Reproducible Research

What is Reproducible Research?



What is Reproducible Research?



Uh....who cares?

FIXING SCIENCE

Most science research findings are false. Here's how we can change that



All

Images

Videos

News

Shopping

More

Settings

Tools

About 176,000,000 results (0.42 seconds)

Most Scientific Findings Are Wrong or Useless - Reason.com

reason.com/archives/2016/08/26/most-scientific-results-are-wrong-or-useless

Aug 26, 2016 - ScientistYanlevDreamstime Yanlev/Dreamstime"Science, the pride of modernity, our one source of objective knowledge, is in deep trouble.

PLOS Medicine: Why Most Published Research Findings Are False

journals.plos.org/plosmedicine/article?id=10.1371/journal.pmed.0020124

by JPA Ioannidis - 2005 - Cited by 4846 - Related articles

Aug 30, 2005 - Moreover, for many current scientific fields, claimed research findings ... Citation: Ioannidis JPA (2005) Why **Most** Published Research Findings Are False. what might have gone **wrong** with their data, analyses, and results.

Is Most Published Research Wrong? - YouTube

<https://www.youtube.com/watch?v=42QuXLucH3Q>

Aug 11, 2016 - Uploaded by Veritasium

Why **Most** Published Research Findings Are False: The problem with the approach to **science** is that ...





















Believe It Or Not, Most Published Research Findings Are Probably ...





bigthink.com/.../believe-it-or-not-most-published-research-findings-are-probably-false

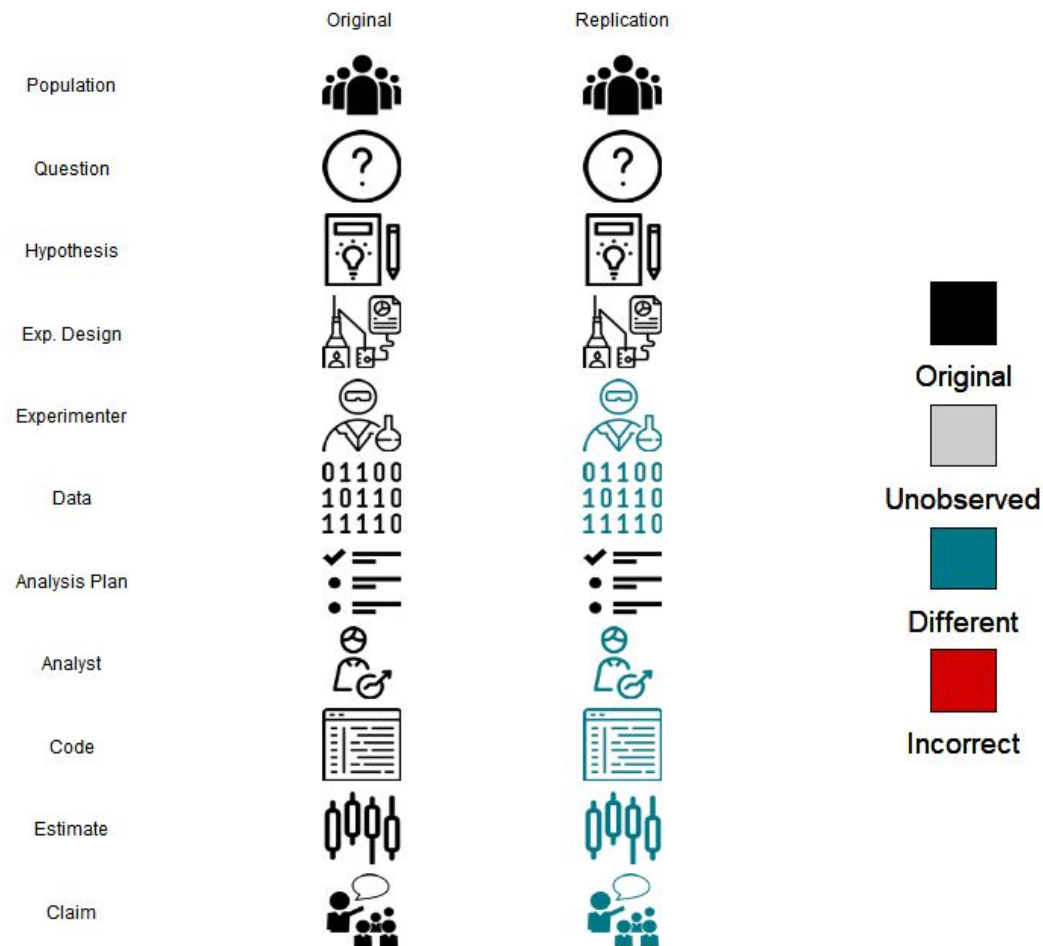
Ten years ago, a researcher claimed **most** published research findings are false; ... of the Internet has worked wonders for the public's access to **science**, but this ... the case, experiments are underpowered,

176,000,000!

Reproduce & replicate

	Original	Reproduction	
Population			
Question			
Hypothesis			
Exp. Design			
Experimenter			
Data	01100 10110 11110	01100 10110 11110	
Analysis Plan			
Analyst			
Code			
Estimate			
Claim			

 Original
 Unobserved
 Different
 Incorrect



When human harm isn't involved

A few things that would reduce stress around reproducibility/replicability in science

Jeff Leek 2017/11/21

I was listening to the Effort Report Episode on [The Messy Execution of Reproducible Research](#) where they were discussing the piece about [Amy Cuddy in the New York Times](#). I think both the article and the podcast did a good job of discussing the nuances of the importance of reproducibility and the challenges of the social interactions around this topic. After listening to the podcast I realized that I see a lot of posts about reproducibility/replicability, but many of them are focused on the technical side. So I started to think about compiling a list of more cultural things we can do to reduce the stress/pressure around the reproducibility crisis.

I'm sure others have pointed these out in other places but I am procrastinating writing something else so I'm writing these down while I'm thinking about them :).

1. **We can define what we mean by “reproduce” and “replicate”** Different fields have different definitions of the words *reproduce* and *replicate*. If you are publishing a new study we now have an [R package](#) that you can use to create figures that show what changed and what was the same between the original study and your new work. Defining concretely what was the same and different will reduce some of the miscommunication about what a reproducibility/replicability study means.

When human harm could happen

From the article:

Cancer trial errors revealed

2006 Anil Potti, a cancer geneticist at Duke University in Durham, North Carolina, and others file patent applications on the idea of using gene-expression data to predict sensitivity to cancer drugs. Potti is first author on a paper in *Nature Medicine*¹.

2007 Potti is last author on a paper in the *Journal of Clinical Oncology (JCO)*². Duke begins three clinical trials to test Potti's predictors in patients with breast or lung cancer.

SEPTEMBER 2009 Keith Baggerly and Kevin Coombes, statisticians at the University of Texas M. D. Anderson Cancer Centre in Houston, publish a paper in *Annals of Applied Statistics*³ stating that they could not replicate Potti's claims. Duke suspends the trials and asks a review panel to investigate.

NOVEMBER 2009 Potti places data underlying the *JCO* paper online. Baggerly writes to Sally Kornbluth, Duke vice-dean for research, and Michael Cuffe, Duke vice-president for medical affairs, to point out differences from raw data.

DECEMBER 2009 An unredacted copy of the report by Duke's review panel, later obtained by *Nature*, shows that the panel replicated Potti's claims using his data, but were unaware that those data contained discrepancies.

JANUARY 2010 Duke restarts clinical trials.

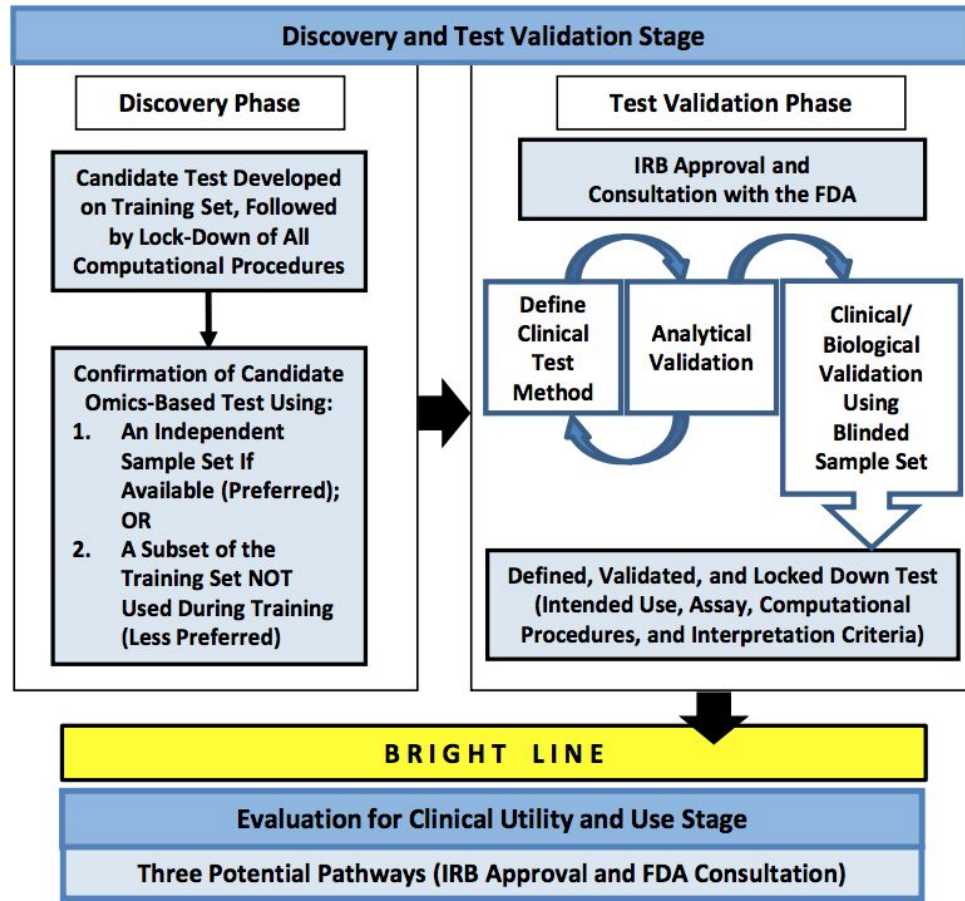
JULY 2010 *The Cancer Letter* reveals that Potti made false claims about his CV. Trials are suspended and an investigation begins. Harold Varmus, director of the National Cancer Institute in Bethesda, Maryland, asks the Institute of Medicine to review Duke's trials.

NOVEMBER 2010 *JCO* paper is retracted. Duke closes the trials permanently. Potti resigns.

DECEMBER 2010 Institute of Medicine study begins, but will now focus more generally on criteria for genomics predictor.

JANUARY 2011 *Nature Medicine* paper is retracted.

<http://www.nature.com/news/2011/110111/full/469139a/box/1.html>



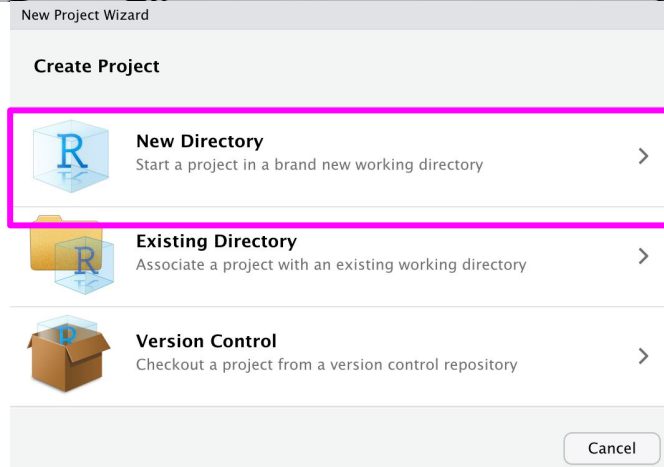
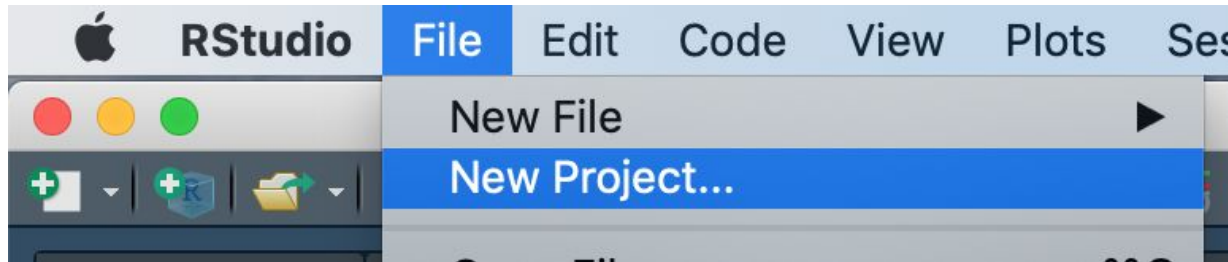
1. Code + documentation
2. Versions of software
3. Data provenance

Your closest collaborator is
you six months ago, but you
don't reply to emails

- Karl Broman

(http://kbroman.org/Tools4RR/assets/lectures/06_org_eda.pdf)

RStudio Projects



New Project Wizard

Back

Project Type



New Project



New Project Wizard

Back

Create New Project



Directory name:

SISBID

Create project as subdirectory of:

~/Desktop

Browse...

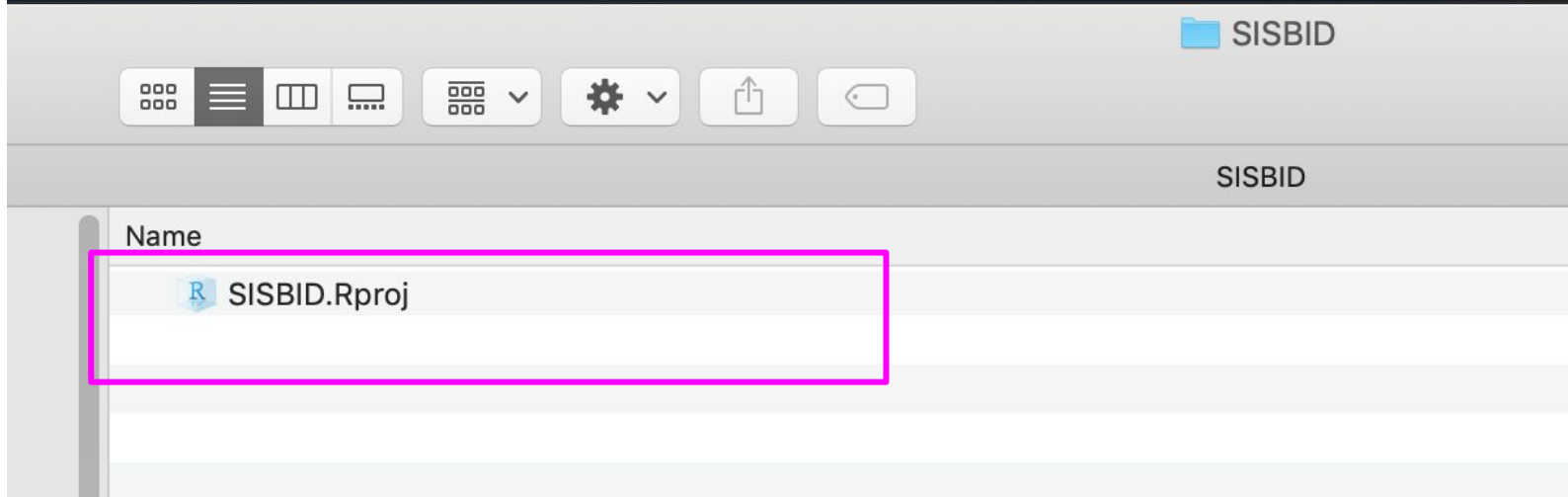
☒ Create a git repository

☐ Use renv with this project

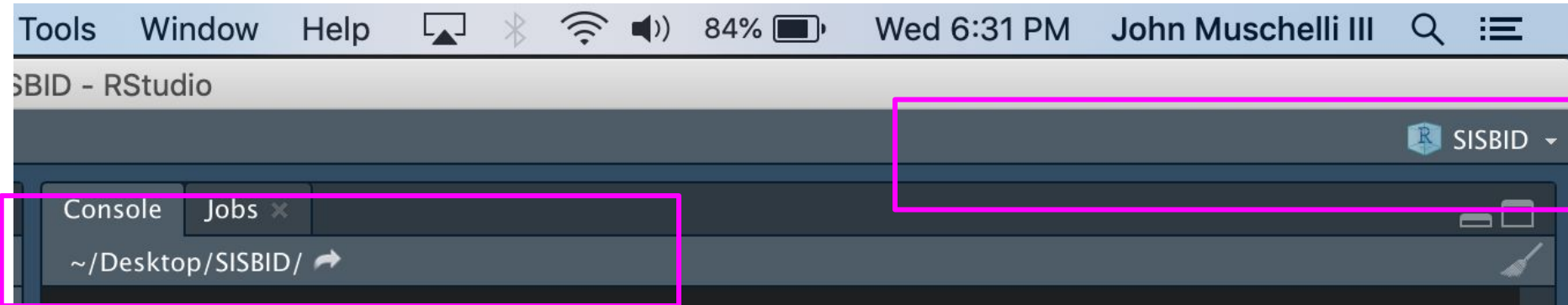
☐ Open in new session

Create Project

Cancel



Double click on the Rproj file - opens RStudio



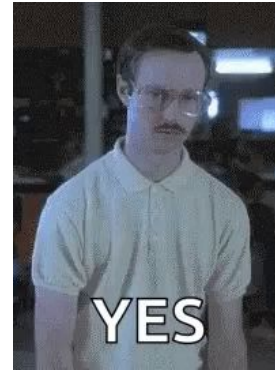
RStudio Projects

- Rproj - Open to directory of Project
- Everything is **relative** to working directory
- Zip up whole folder and send to someone else

```
read_csv("C:/terrible/path/john/blah.csv")
```



```
read_csv("data/blah.csv")
```



Could you just re-run that code
with the [latest/different/best]
parameters?

- Every collaborator/PI

The magic of Markdown

- bullets
- **bold**
- *italics*
- [links](https://google.com)
- or run inline `r code`



- bullets
- **bold**
- *italics*
- links
- or run inline `r code`

https://rmarkdown.rstudio.com/authoring_basics.html

Introduction

Here is some background you need to know:

Lorem ipsum dolor sit amet, consectetur adipiscing elit. Nullam iaculis enim ut enim viverra molestie. In lacinia aliquet urna, nec vulputate quam congue et. Maecenas porta mauris sem, nec laoreet sapien tincidunt non. Integer sit amet consequat neque, non iaculis ligula.

Hypothesis

Pellentesque molestie erat nec elit efficitur, sit amet sodales erat viverra. Mauris sed commodo eros, ac volutpat sem. Morbi convallis leo et dui cursus, eu suscipit turpis efficitur.

Section 1 code and results

First I will run this.

```
##{r}
print("Hello world")
print("Yup, this is important")
```

The output of which is consistent with my hypothesis.

Conclusion

I can move on to the next part of my project

.Rmd
document

Introduction

Here is some background you need to know:

Lorem ipsum dolor sit amet, consectetur adipiscing elit. Nullam iaculis enim ut enim viverra molestie. In lacinia aliquet urna, nec vulputate quam congue et. Maecenas porta mauris sem, nec laoreet sapien tincidunt non. Integer sit amet consequat neque, non iaculis ligula.

Hypothesis

Pellentesque molestie erat nec elit efficitur, sit amet sodales erat viverra. Mauris sed commodo eros, ac volutpat sem. Morbi convallis leo et dui cursus, eu suscipit turpis efficitur.

Section 1 code and results

First I will run this.

```
print("Hello world")
```

```
## [1] "Hello world"
```

```
print("Yup, this is important")
```

```
## [1] "Yup, this is important"
```

The output of which is consistent with my hypothesis.

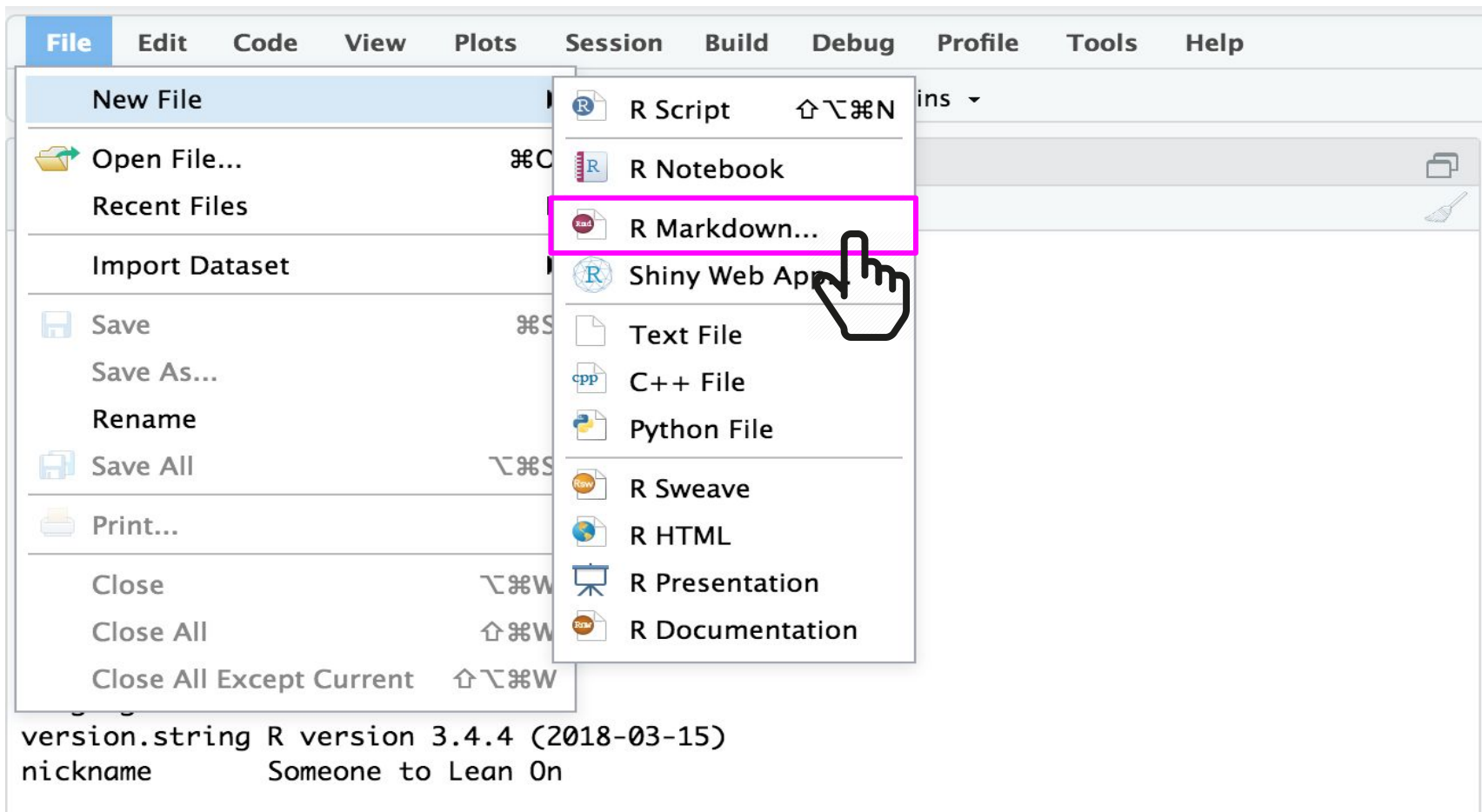
Conclusion

I can move on to the next part of my project

PDF, HTML or Word
document

Another major benefit to R Markdown is that since it is plain text, it works very well with version control systems. It is easy to track what character changes occur between commits; unlike other formats that aren't plain text. For example, in one version of this lesson, I may have forgotten to bold this word. When I catch my mistake, I can make the plain text changes to signal I would like that word bolded, and in the commit, you can see the exact character changes that occurred to now make the word bold.

Another major benefit to R Markdown is that since it is plain text, it works very well with version control systems. It is easy to track what character changes occur between commits; unlike other formats that aren't plain text. For example, in one version of this lesson, I may have forgotten to bold **this** word. When I catch my mistake, I can make the plain text changes to signal I would like that word bolded, and in the commit, you can see the exact character changes that occurred to now make the word bold.



Install Required Packages



Creating R Markdown documents requires updated versions of the following packages: evaluate, highr, markdown, yaml, htmltools, caTools, bitops, knitr, jsonlite, base64enc, rprojroot, rmarkdown.

Do you want to install these packages now?

Yes

No



New R Markdown



Document



Presentation



Shiny



From Template

Title:

Untitled

Author:

Default Output Format:

☒ HTML

Recommended format for authoring (you can switch to PDF or Word output anytime).

☐ PDF

PDF output requires TeX (MiKTeX on Windows, MacTeX 2013+ on OS X, TeX Live 2013+ on Linux).


☐ Word


Previewing Word documents requires an installation of MS Word (or Libre/Open Office on Linux).


OK


Cancel

New R Markdown

 Document

 Presentation

 Shiny

 From Template

Title:

Author:

Default Output Format:

☒ **HTML**
Recommended format for authoring (you can switch to PDF or Word output anytime).

☐ **PDF**
PDF output requires TeX (MiKTeX on Windows, MacTeX 2013+ on OS X, TeX Live 2013+ on Linux).

☐ **Word**
Previewing Word documents requires an installation of MS Word (or Libre/Open Office on Linux).

Untitled1 x

Insert | Run |

```
1 ---
2 title: "My First R Markdown Document!"
3 author: "Jane Doe"
4 date: "5/31/2018"
5 output: html_document
6 ---
7
8 ```{r setup, include=FALSE}
9 knitr::opts_chunk$set(echo = TRUE)
10 ```
11
12 ## R Markdown
13
14 This is an R Markdown document. Markdown is a simple formatting syntax for authoring HTML, PDF, and MS Word
15 documents. For more details on using R Markdown see <http://rmarkdown.rstudio.com>.
16
17 When you click the Knit button a document will be generated that includes both content as well as the
18 output of any embedded R code chunks within the document. You can embed an R code chunk like this:
19
20 ```{r cars}
21 summary(cars)
22 ```
23
24 ## Including Plots
25
26 You can also embed plots, for example:
27
28 ```{r pressure, echo=FALSE}
29 plot(pressure)
30 ```
31
32 2:1 | # My First R Markdown Document! | R Markdown
```

YAML - not markdown,
Spaces matter, usually goes
option: value

TEXT

CODE CHUNK

Untitled1 x

Knit

Insert

Run

```
1 ---
2 title: "My First R Markdown Document!"
3 author: "Jane Doe"
4 date: "5/31/2018"
5 output: html_document
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7
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19
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21 summary(cars)
22 ```
23
24 ## Including Plots
25
26 You can also embed plots, for example:
27
28 ```{r pressure, echo=FALSE}
29 plot(pressure)
30 ```
```

2:1 # My First R Markdown Document! R Markdown

Save File – Untitled1

File name:

> / > cloud > project



..



.Rhistory

0 B

May 30, 2018, 12:34 PM



project.Rproj

205 B

May 31, 2018, 3:35 PM

New Folder

Save

Cancel



Header rendered as the title

My First R Markdown Document!

Jane Doe

5/31/2018

R Markdown

Text section rendered as formatted text

This is an R Markdown document. Markdown is a simple formatting syntax for authoring HTML, PDF, and MS Word documents. For more details on using R Markdown see <http://rmarkdown.rstudio.com>.

When you click the **Knit** button a document will be generated that includes both content as well as the output of any embedded R code chunks within the document. You can embed an R code chunk like this:

```
summary(cars)
```

Code rendered as the input code AND the output of running the code chunk

```
##      speed      dist
##  Min.   : 4.0    Min.   : 2.00
## 1st Qu.:12.0    1st Qu.: 26.00
##  Median :15.0    Median : 36.00
##  Mean   :15.4    Mean    : 42.98
## 3rd Qu.:19.0    3rd Qu.: 56.00
##  Max.   :25.0    Max.    :120.00
```

Including Plots

You can also embed plots, for example:

test_document.Rmd x

Insert

Run

Knit

```
1 ---
2 title: "My First R Markdown Document!"
3 author: "Jane Doe"
4 date: "5/31/2018"
5 output: html_document
6 ---
7
8 ```{r setup, include=FALSE}
9 knitr::opts_chunk$set(echo = TRUE)
10 ```
11
12 ## R Markdown
13
14 This is an R Markdown document. Markdown is a simple formatting syntax for
15 authoring HTML, PDF, and MS Word documents. For more details on using R Markdown
16 see <http://rmarkdown.rstudio.com>.
17:1 R Markdown
```

Console Terminal x

/cloud/project/

> |

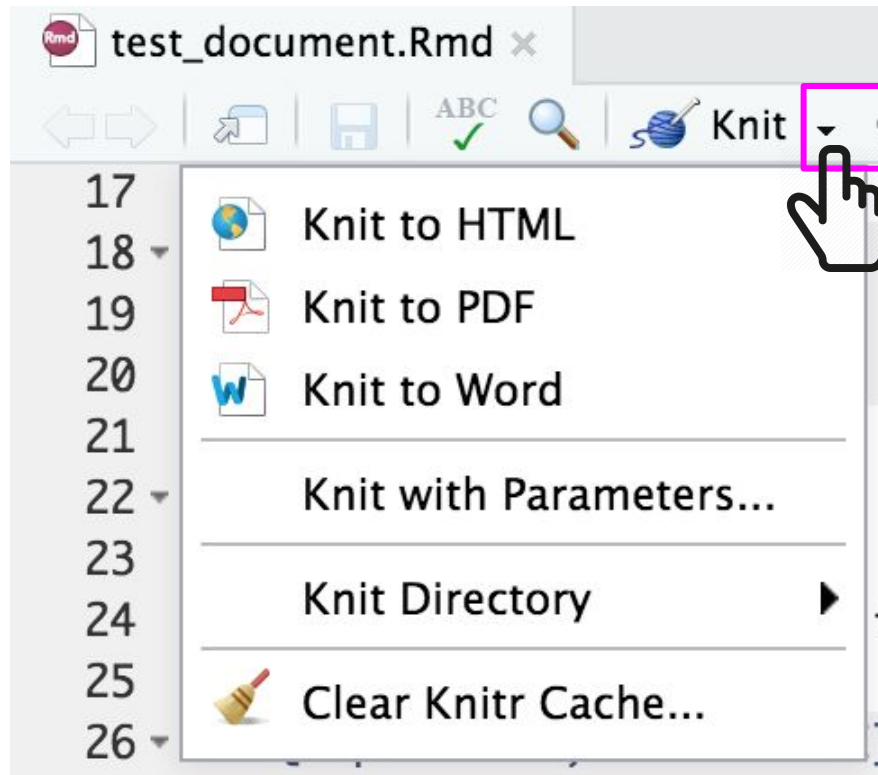
Environment History Connections

Files Plots Packages Help Viewer

New Folder Upload Delete Rename More

/ > / > cloud > project

	Name	Size	Modified
	..		
<input type="checkbox"/>	.Rhistory	0 B	May 30, 2018, 12:34 PM
<input type="checkbox"/>	project.Rproj	205 B	Jun 1, 2018, 8:33 AM
<input type="checkbox"/>	test_document.html	741.2 KB	May 31, 2018, 3:57 PM
<input type="checkbox"/>	test_document.Rmd	854 B	May 31, 2018, 3:57 PM



Rendering in R

```
library(rmarkdown)  
render("Untitled.Rmd")
```

Session information - what's loaded?

```
devtools::session_info()  
# comment-for specific package  
devtools::session_info("pkg")
```

Overview

Declararing Parameters

YAML Params Field

Parameter Types

Using Parameters

Accessing from R

Passing Parameters

Parameter User Interfaces

Parameterized Reports

Overview

R Markdown documents can optionally include one or more parameters. Parameters are useful when you want to re-render the same report with distinct values for various key inputs, for example:

1. Running a report specific to a department or geographic region.
2. Running a report that covers a specific period in time.
3. Running multiple versions of a report for distinct sets of core assumptions.

R Markdown parameter names, types, and default values are declared in the YAML section at the top of the document. To change these values for a given rendering you use the `params` argument to the `rmarkdown::render` function.

Note that parameterized reports are a new feature of R Markdown and therefore require very recent versions of the **knitr** (v1.11) and **rmarkdown** (v0.8) packages. You can install the most up to date versions with the following command:

http://rmarkdown.rstudio.com/developer_parameterized_reports.html

Set parameters in yaml

title: My Document

output: html_document

params:

region: east

Can set any R type if you use !r before expression

title: My Document

output: html_document

params:

start: !r as.Date("2020-01-01")

Accessing the parameters

```
params$region
```

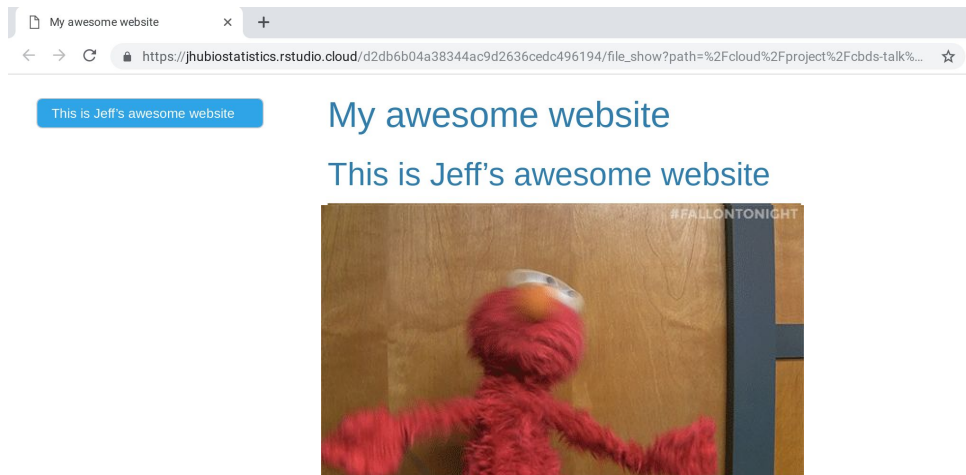
```
params$start
```

Barely scratching the surface

rmarkdown

```
---  
title: "My awesome website"  
output:  
  html_document:  
    toc: true  
    toc_float: true  
    theme: cerulean  
---  
# This is Jeff's awesome website  
  

```



flexdashboard

title: "How does your BMI measure up?"

output: flexdashboard::flex_dashboard

runtime: shiny

Inputs {.sidebar}

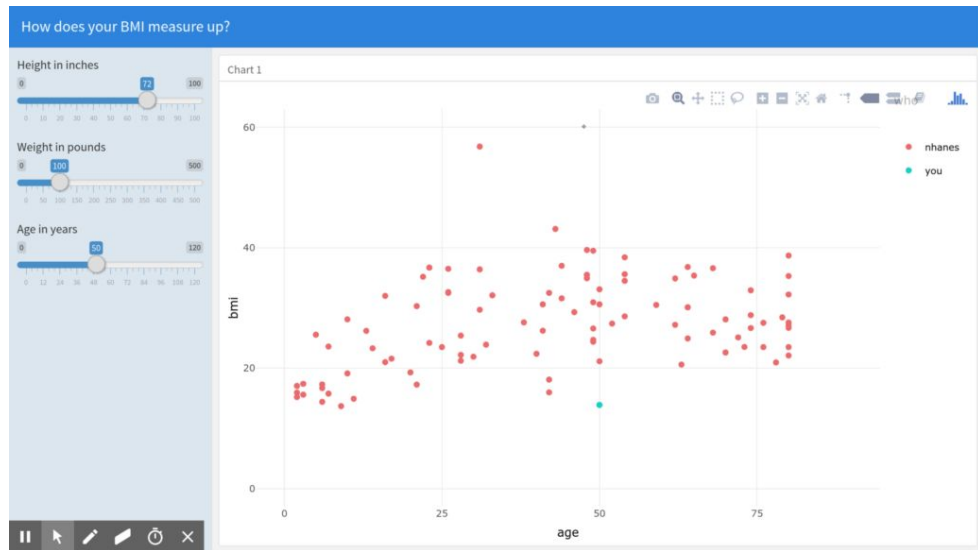
```
```{r}
library(flexdashboard); library(NHANES); library(plotly);library(dplyr)
sliderInput("height", "Height in inches",0,100,72)
sliderInput("weight", "Weight in pounds",0,500,100)
sliderInput("age", "Age in years",0,120,50)
```

...

Column

### Chart 1

```
```{r}
nhanes = sample_n(NHANES,100)
renderPlotly({
  df = data.frame(bmi = c(nhanes$BMI,input$weight*0.45/(input$height*0.025)^2),
    age = c(nhanes$Age,input$age),
    who = c(rep("nhanes",100),"you"))
  ggplotly(ggplot(df) +
    geom_point(aes(x=age,y=bmi,color=who)) +
    scale_x_continuous(limits=c(0,90)) +
    scale_y_continuous(limits=c(0,60)) +
    theme_minimal()
  )
})
```
```





Downloading data  
reproducibly

## Finding and creating files

```
getwd() # get working directory
```

```
setwd("data") # set
```

```
file.exists("data")
```

```
dir.create("data")
```

```
list.files("data")
```

Putting it together

```
if (!file.exists("data")) {
 dir.create("data")
}
```



## Finding and creating files

```
file.exists("data")
```

```
dir.create("data")
```

```
list.files("data")
```

```
fileUrl <-
"https://data.baltimorecity.gov/api/views/dz54-2aru/rows.csv?accessType=DOWNLOAD"

curl::curl_download(fileUrl,
 destfile="./data/cameras.csv")
list.files("./data")
dateDownloaded <- date()
dateDownloaded
```

# Rmarkdown lab

<https://bit.ly/1LRk3ds>

Download the file from

<https://github.com/SISBID/Module1/raw/gh-pages/abs/rmarkdown-lab.Rmd>