

R Markdown Workshop



Creation, organisation, storage, and collaboration

Anne-Kathrin Kleine | University of Groningen

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...taking the pain away

- ✍ efficient management of code and results

- 📘 easy conversion to pdf, html, word

- 🧘 guaranteed reproducibility, less mistakes

Workshop overview

Day 1:

1. R Markdown basics
2. Writing text
3. Reference management
4. Writing code
5. Figures, tables, and plots
6. R project organization

Day 1:

1. R Markdown basics

GOAL 1 💪 Write your analysis scripts in R Markdown

2. Writing text

GOAL 2 💪 Convert the output into a format of your choice (word, html, pdf)

3. Reference management

GOAL 3 💪 Add text, figures/tables, and references

4. Writing code

GOAL 4 💪 Learn proper project structuring and using templates

5. Figures, tables, and plots

6. R project organization

Day 2:

1. Git & GitHub basics
2. Collaborating using Git & GitHub
3. Osf integration
4. Blogdown - build your project website
5. Xaringan, bookdown & Co.

Day 2:

1. Git & GitHub basics

2. Collaborating using Git & GitHub

3. Osf integration

4. Blogdown - build your project website

5. Xaringan, bookdown & Co.

GOAL 1 💪 Learn how to use Git & GitHub for version control

GOAL 2 💪 Learn how to collaborate using Git & GitHub

GOAL 3 💪 Share your code and data through osf

GOAL 4 💪 Build your own data analysis project website

GOAL 5 💪 Learn how to use R & Markdown to produce a multitude of outputs (books, presentations, interactive applications)

General organization:

Presentation, then practice

You will work in breakout rooms, groups of 3 to 4

? ? Questions ? ?

1. Ask your group 🧑

2. Google (together) 🧑 🧑 🧑

3. Ask me 🤟



Find me at...

[LinkedIn](#)

[GitHub](#)

[AnneKathrinKleine.com](#)

a.kleine@rug.nl

Or find me ...



Or find me ...



Or find me ...



How familiar are you with using R for data analysis?

Go to <https://www.menti.com/yzy8sx3mb6>  (and enter the code 6959 8626)

[See results](#)

Major resources:

[Advanced R Markdown Workshop R Conf 2019](#)

[R Markdown: The Definitive Guide](#)

R Markdown Workshop by [Resul Umit](#)

Detailed Outline

- Did you do the setup?
- Creating an R project
- The YAML header
- Writing text 
- Reference management
- Writing code
- Figures
- Plots - ggplot2
- Structure your project
-  Yey, you're all set! 
- Your turn! 

Did you do the

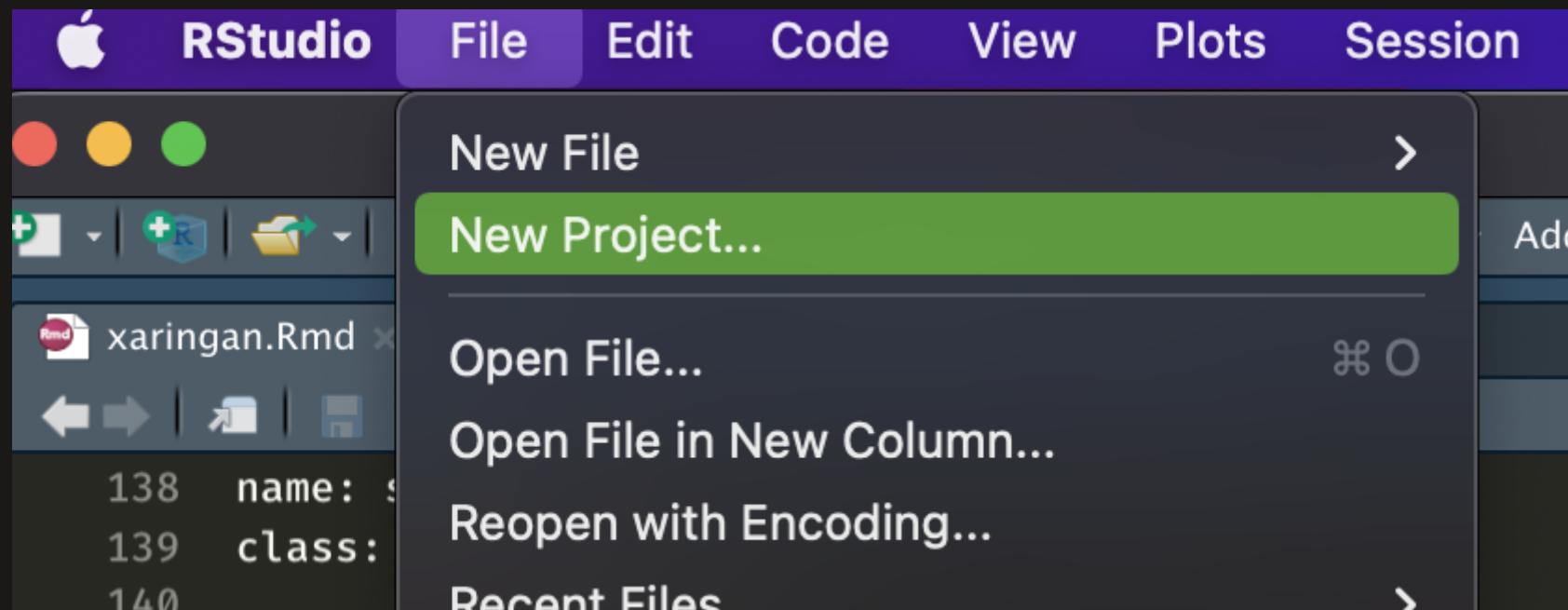
setup?

[Link to setup](#)

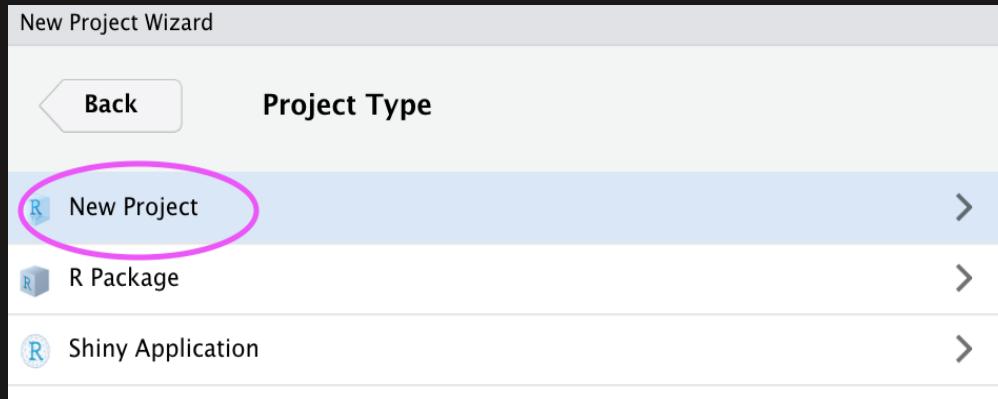
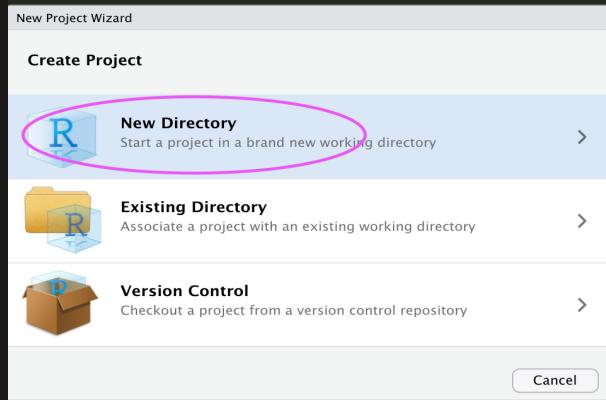
Creating an R project

Creating an R project

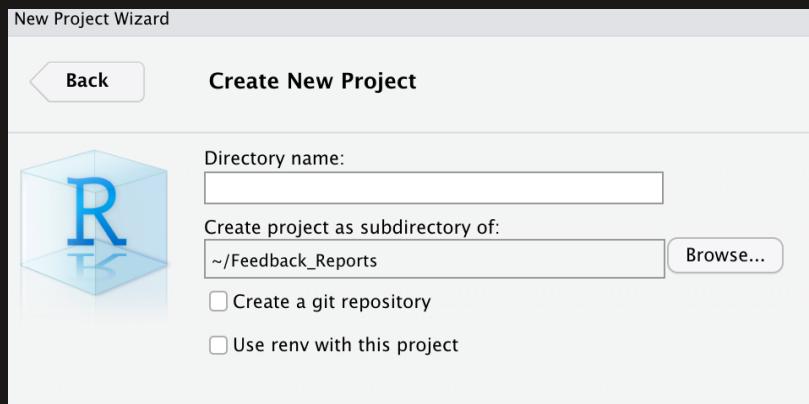
Create a new R Project from within R Studio



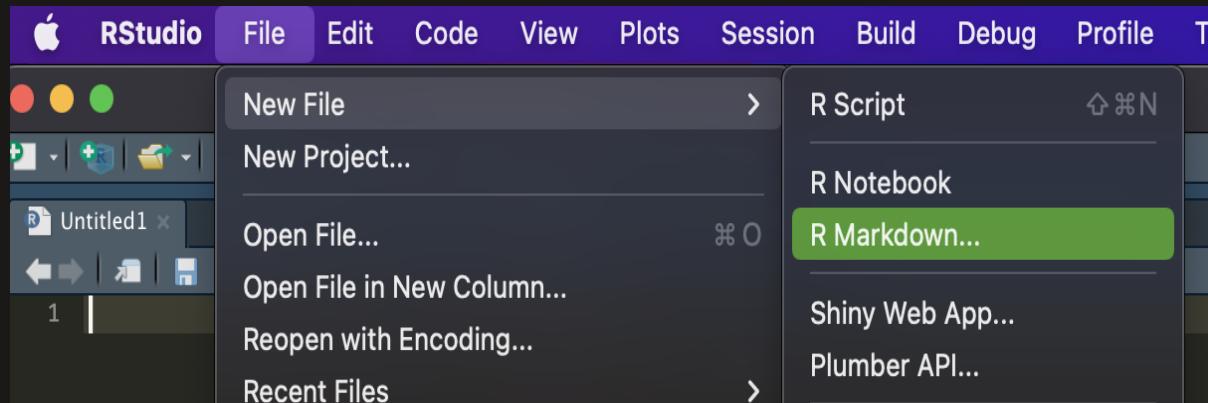
Save in new directory



Choose the folder where you want to store your R Markdown file



In that new folder, create an R Markdown file



When you knit, the following happens:



knit executes the code if there is any, converts the resulting document from .Rmd (R Markdown) into .md (Markdown)

pandoc transforms the .md document into your preferred output format(s) (e.g., word, pdf, html)

The YAML header

YAML — General

YAML includes the metadata variables

- e.g., title, output format
- written between a pair of three hyphens -

```
---
```

```
title:
```

```
output:
```

```
---
```

YAML — Variables

- see [Pandoc User's Guide](#) and [R Markdown Cheat Sheet](#) for documentation and help
- Typical YAML variables for an research paper are as follows:

```
---
```

```
title:
author:
date:
bibliography:
csl:
output:
---
```

YAML – Variables

Variables can be provided as **strings**

```
---  
title: "My very funny and descriptive article title"  
output:  
---
```

YAML – Variables

Variables can be provided as strings, options

```
---
```

```
title: "My very funny and descriptive article title"
output: pdf_document
---
```

YAML — Variables

Variables can be provided as strings, options, and sub-options

```
---
```

```
title: "My very funny and descriptive article title"
output:
  pdf_document:
    keep_tex: true
---
```

YAML — Variables

Variables can be provided as strings, options, sub-options, and code

```
---
```

```
title: "My very funny and descriptive article title"
output:
  pdf_document:
    keep_tex: true
date: "\`r format(Sys.Date(), '%d %B %Y')`"
---
```

YAML — Variables — Output Formats

Documents as output formats include

- **HTML**

```
---
```

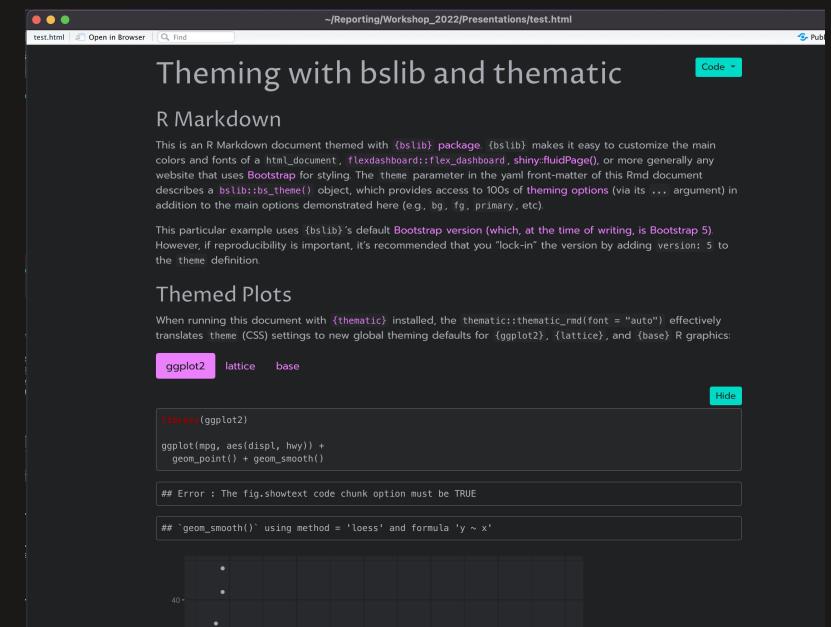
```
title: "Theming with bslib and thematic"
```

```
output:
```

```
  html_document:
```

```
  ...
```

```
---
```



YAML — Variables — Output Formats

Documents as output formats include

- HTML
- **LaTeX**

```
---
```

title: "Theming with bslib and thematic"

output: latex_document

```
---
```

```
2  \PassOptionsToPackage{unicode}{hyperref}
3  \PassOptionsToPackage{hyphens}{url}
4  %
5  \documentclass[
6  ]{article}
7  \usepackage{amsmath,amssymb}
8  \usepackage{lmodern}
9  \usepackage{ifxetex}
10 \ifxetex
11   \usepackage[1]{fontenc}
12   \usepackage[utf8]{inputenc}
13   \usepackage{textcomp} % provide euro and other symbols
14 \else % if luatex or xetex
15   \usepackage{unicode-math}
16   \defaultfontfeatures{Scale=MatchLowercase}
17   \defaultfontfeatures[\rmfamily]{Ligatures=TeX,Scale=1}
18 \fi
19 % Use upquote if available, for straight quotes in verbatim environments
20 \IfFileExists{upquote.sty}{\usepackage{upquote}}{}
21 \IfFileExists{microtype.sty}{% use microtype if available
22   \usepackage[microtype]
23   \usepackageSet[protrusion]{basicmath} % disable protrusion for tt fonts
24 }
25 \makeatletter
26 \ifundefined{KOMAClassName}{% if non-KOMA class
27   \IfFileExists{parskip.sty}{%
28     \usepackage{parskip}
29   }%
30   \setlength{\parindent}{0pt}
31   \setlength{\parskip}{6pt plus 2pt minus 1pt}
32 }% if KOMA class
33 \KOMAoptions{parskip=half}
34 \makeatother
35 \usepackage{xcolor}
36 \IfFileExists{xurl.sty}{\usepackage{xurl}}{} % add URL line breaks if available
37 \IfFileExists{bookmark.sty}{\usepackage{bookmark}}{\usepackage{hyperref}}
38 \hypersetup{
39   pdftitle=(Theming with bslib and thematic),
40   hidelinks,
41   pdfcreator={LaTeX via pandoc}
42   \urlstyle{same} % disable monospaced font for URLs
43   \usepackage[margin=1in]{geometry}
44   \usepackage{color}
45   \usepackage{fancyvrb}
46   \newcommand{\VerbBar}{|}
47   \newcommand{\VERB}{\VerbBar\Verbatim\VerbBar}
48   \DefineVerbatimEnvironment{Highlighting}{Verbatim}{commandchars=\\\{\}}
49   % Add more commandchars (e.g. {, })
50   \usepackage{framed}
51   \definecolor{shadecolor}{RGB}{248, 248, 248}
52   \newenvironment{Shaded}{\begin{snugshade}}{\end{snugshade}}
53   \newcommand{\AlertTok}[1]{\textcolor{rgb}{0.94, 0.16, 0.16}{\#1}}
54   \newcommand{\AnnotationTok}[1]{\textcolor{rgb}{0.56, 0.35, 0.01}{\textbf{\textit{#1}}}}
55   \newcommand{\AttributeTok}[1]{\textcolor{rgb}{0.77, 0.63, 0.00}{\#1}}
56   \newcommand{\BaseNTok}[1]{\textcolor{rgb}{0.00, 0.00, 0.81}{\#1}}
57   \newcommand{\BuiltInTok}[1]{\textcolor{black}{#1}}
58   \newcommand{\ColorTok}[1]{\textcolor{black}{#1}}
59   \newcommand{\DocumentationTok}[1]{\textcolor{black}{#1}}
60   \newcommand{\ErrorTok}[1]{\textcolor{black}{#1}}
61   \newcommand{\GeneralTok}[1]{\textcolor{black}{#1}}
```

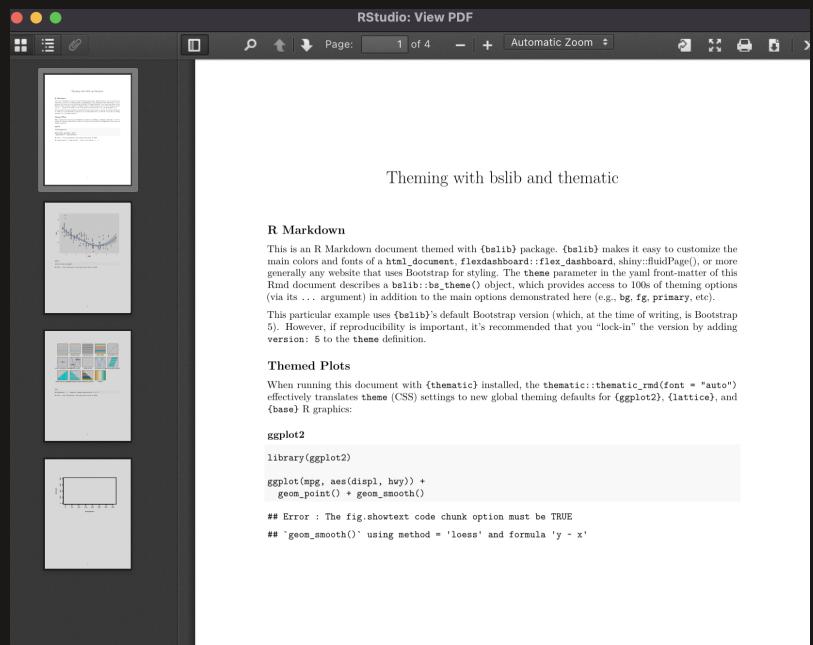
YAML — Variables — Output Formats

Documents as output formats include

- HTML
- LaTeX
- PDF

```
---
```

```
title: "Theming with bslib and thematic"
output: pdf_document
---
```



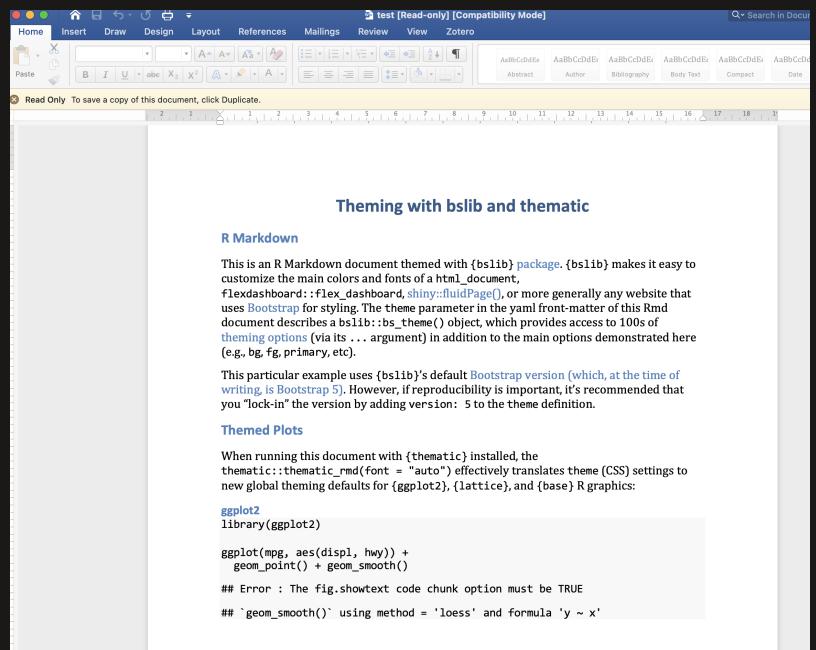
YAML — Variables — Output Formats

Documents as output formats include

- HTML
- LaTeX
- PDF
- Word

```
---
```

```
title: "Theming with bslib and thematic"
output: word_document
---
```



YAML — External Files

```
---
title: "Journal title"
author: "Jane Doe^[Department of Science, University of Random. Email: jane.doe@random.edu. Website: http://www.jane
date: 4 March 2022
bibliography: references.bib
csl: apa.csl
output: pdf_document
---
```

YAML — Strings — External Files

```
---
```

...
bibliography: references/ref_library.bib
csl: "../../paperstyle/csl/apa.csl" # two dots (...) mean moving one folder up
...

- ! ! locations of files re specified as **relative to the working directory ! !**
- for reproducibility reasons, hard-coded strings should be avoided 🙏
 - e.g.,  "C:/Users/Dropbox/styles/apa.csl" 

Writing text



Lines

Multiple spaces on a given line are reduced to one

This is a sentence followed by four spaces. This is another sentence on the same line.

This is a sentence followed by four spaces. This is another sentence on the same line.

Line endings with fewer than two spaces are ignored

This is a sentence followed by one space.
This is another sentence on a new line.

This is a sentence followed by one space. This is another sentence on a new line.

Hard Breaks

Two or more spaces at the end of lines introduce hard breaks, forcing a new line

This is a sentence followed by two spaces.
This is another sentence on a new line.

This is a sentence followed by two spaces.
This is another sentence on a new line.

Line Blocks

Spaces on lines that start with a vertical line | are kept

```
| a one-space indent  
|   a five-space indent  
|     a ten-space indent
```

```
a one-space indent  
a five-space indent  
a ten-space indent
```

Block Quotes

Lines starting with the greater-than sign > introduce block quotes*

```
> In God, we trust. All others must bring data.  
>  
> --- Anonymous
```

In God, we trust. All others must bring data.

— Anonymous

Paragraphs

One or more blank lines introduce a new paragraph

This is the first sentence of a paragraph as it is preceded by a blank line. This is the second sentence of that paragraph, which is followed by a blank line.

This is the first sentence of a *new paragraph* as it is preceded by a blank line.

This is the first sentence of a paragraph as it is preceded by a blank line. This is the second sentence of that paragraph, which is followed by a blank line.

This is the first sentence of a *new paragraph* as it is preceded by a blank line.

Comments

Text with the syntax `<!-- comments -->` is omitted from output

<!-- This paragraph needs re-writing -->

This is the first sentence of a paragraph as it is preceded by a blank line.

This is the first sentence of a new paragraph *<!-- I've removed italics -->* as it is preceded by a blank line.

This is the first sentence of a paragraph as it is preceded by a blank line.

This is the first sentence of a new paragraph as it is preceded by a blank line.

Headers

The number sign # introduces headers; lower levels are created with additional signs – up to total five levels

Introduction becomes

Introduction

1. Introduction becomes

Introduction

3.1 Introduction becomes

Introduction

Introduction becomes

Introduction

Introduction becomes

Introduction

Emphases

A pair of single asterisk `*` or underscores `_` introduces italics

`*italics*` becomes `italics`

`_italics_` becomes `italics` as well

A pair of double asterisk or underscores introduces bold

`**bold**` becomes `bold`

`--bold--` becomes `bold` as well

These two rules can be combined

`**_bolditalics_**` becomes `bolditalics`

`_**bolditalics**_` becomes `bolditalics` as well

Strikethrough

A pair of double tildes ~ introduces strikethrough

~~**strikethrough**~~ becomes **strikethrough**

Strikethrough can be combined with italics or bold

~~strikebold**~~** or __~~**strikebold**~~__, they both become **strikebold**

~~****strikebold****~~ or ~~__**strikebold**__~~, they both become **strikebold** as well

*~~**strikeitalics**~~* or __~~**strikeitalics**~~__, they both become **strikeitalics**

~~***strikeitalics***~~ or ~~__**strikeitalics**__~~, they both become **strikeitalics** as well

External links

You can link text to URLs

[visit my website](<https://anneokk.netlify.app/>) becomes [visit my website](#)

You can also link text to an email address

[email me](mailto:a.k.kleine@rug.nl) becomes [email me](mailto:a.k.kleine@rug.nl)

<a.k.kleine@rug.nl> becomes a.k.kleine@rug.nl

Equations

Inline equations go between a pair of single dollar signs "\$" with no space between the signs and the equation itself

`$E = mc^2$` becomes `E = mc2`

Block equations go in between a pair of double dollar signs

`$$E = mc^2$$` becomes

$$E = mc^2$$

Inline Notes

For inline footnotes, use the `^[footnote]` syntax

`Jane Doe1[Corresponding author.] becomes Jane Doe1`

¹ Corresponding author.

Lists

Lines starting with asterisk * as well as plus + or minus – signs introduce lists

- books
- articles
- reports

- books
- articles
- reports

Lists

Lists can be nested within each other, with indentation

```
+ books
+ articles
  - published
  - under review
    + revised and resubmitted
  - work in progress
```

- books
- articles
 - published
 - under review
 - revised and resubmitted
 - work in progress

Lists

List items can be numbered

1. books
2. articles
 - published
 - under review
 - + revised and resubmitted
 - work in progress

1. books
2. articles
 - published
 - under review
 - revised and resubmitted
 - work in progress

Dashes

Two hyphens grouped together introduce an en-dash

-- becomes –

Three hyphens grouped together introduce an em-dash

--- becomes —

Subscripts and Superscripts

A pair of tildes introduces subscript

C \sim 2 \sim becomes CO₂

A pair of carets introduces superscript

R \wedge 2 \wedge becomes R²

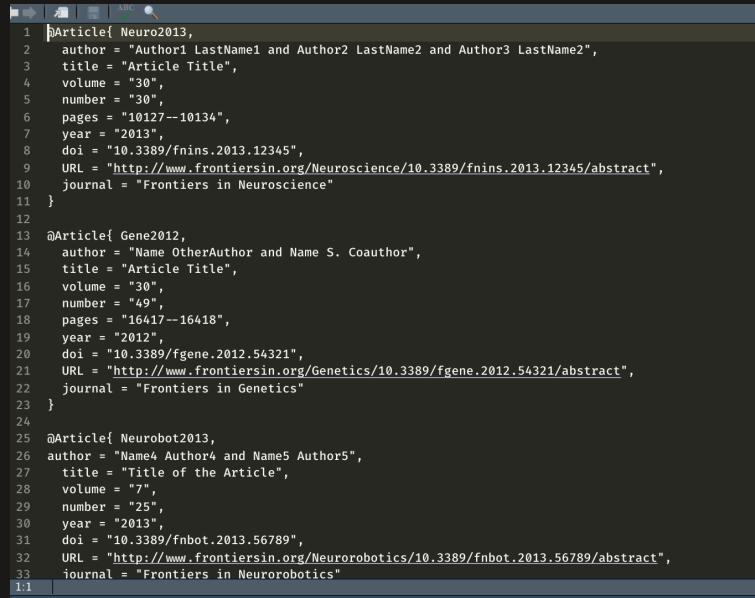
Reference management

References — Bibliography Database

- References are defined in .bib files
 - they follow the BibTeX format
- pandoc looks for a .bib file, and for the definitions therein, to process citations

```
---  
...  
bibliography: refs.bib  
...  
---
```

- .bib files are specified with the bibliography variable in YAML



```
1 @Article{ Neuro2013,  
2   author = "Author1 LastName1 and Author2 LastName2 and Author3 LastName2",  
3   title = "Article Title",  
4   volume = "30",  
5   number = "30",  
6   pages = "10127--10134",  
7   year = "2013",  
8   doi = "10.3389/fnins.2013.12345",  
9   URL = "http://www.frontiersin.org/Neuroscience/10.3389/fnins.2013.12345/abstract",  
10  journal = "Frontiers in Neuroscience"  
11 }  
12  
13 @Article{ Gene2012,  
14   author = "Name OtherAuthor and Name S. Coauthor",  
15   title = "Article Title",  
16   volume = "30",  
17   number = "49",  
18   pages = "16417--16418",  
19   year = "2012",  
20   doi = "10.3389/fgene.2012.54321",  
21   URL = "http://www.frontiersin.org/Genetics/10.3389/fgene.2012.54321/abstract",  
22   journal = "Frontiers in Genetics"  
23 }  
24  
25 @Article{ Neurobot2013,  
26   author = "Name4 Author4 and Name5 Author5",  
27   title = "Title of the Article",  
28   volume = "",  
29   number = "25",  
30   year = "2013",  
31   doi = "10.3389/fnbot.2013.56789",  
32   URL = "http://www.frontiersin.org/Neurorobotics/10.3389/fnbot.2013.56789/abstract",  
33   journal = "Frontiers in Neurorobotics"  
34 }  
35
```

- A BibTeX entry consists of three elements
 - a type (e.g., @Article)
 - a citation-key (e.g., Neuro2013)
 - a number of tags (e.g., title, volume, ...)

References — Bibliography Database — Entries

Get information via **Google Scholar**:

follow **cite** -> **BibTex** and copy

paste into .bib, edit if necessary, and save

References — Bibliography Database — Entries

Get information via **Google Scholar**:

follow **cite** -> **BibTex** and copy

paste into .bib, edit if necessary, and save

Better options: Zotero, Mendeley

★ Useful article on how to use Zotero with R Markdown ★

References — Style

- Reference styles are defined in `.csl` files
- files for different styles (e.g., APA) are available at <https://www.zotero.org/styles>

```
---
```

```
...
csl: "../apa.csl"
...
---
```

- `.csl` files are specified with the `csl` variable in YAML

References — In-text Citation Syntax — Author-Date Styles (APA 7)

[@bennett2015] becomes (Bennett, 2015)

@bennett2015 becomes Bennett (2015)

[‐@bennett2015] becomes (2015)

-@bennett2015 becomes 2015

[@bennett2015 35] becomes (Bennett, 2015, p. 35)

[@bennett2015 33‐35] becomes (Bennett, 2015,
pp. 33‐35)

[@bennett2015, ch. 1] becomes (Bennett, 2015,
ch. 1)

[@bennett2015; @gilbert2019] becomes
(Bennett, 2015; Gilbert, 2019)

[see @bennett2015, for details] becomes
(see Bennett, 2015, for details)

@bennett2015 [33‐35] becomes Bennett (2015,
pp. 33‐35)

Citations — Reference List

This is the last sentence of an APA style manuscript.
References

This is the last sentence of an APA style manuscript.

References

Bennett, S. (2015). Peanut butter and jelly. *Journal of Bone*, 1(12), 3–35.

Gilbert, T. (2019). Turning wine into water. In M. Albert (Ed.), *The book of ground* (pp. 124–142). Antman.

Writing code

Data: [Superhero Dataset](#)

Code, in and outside chunks

Code chunks:

```
library(readxl)
library(dplyr)
df <- read_excel("data/heroes_information.xlsx") %>%
  na_if(., -99) %>%
  mutate(bmi = round(Weight/((Height/100)**2)), 2) %>%
  rename(., Water_allergy = Gender) %>%
  mutate(Water_allergy = recode(Water_allergy, Male = "Yes", Female = "No")) %>%
  select(., -...1, -`2`)
```

Inline:

The average height of a superhero is `r round(mean(df$Height, na.rm = T), 2)` centimeter
becomes:

The average height of a superhero is 186.73 centimeter.

Code Chunks

- Code chunks are delimited spaces between a pair of three backticks `````
 - their output, if there is any, appears in the output document
-
- in curly brackets `{}`, code chunks take
 - a language engine (e.g., `r`)
 - a label (e.g., `setup`)
 - options (e.g., `echo = FALSE`)
 - The complete list of options is available at
<https://yihui.org/knitr/options>

```
```{r, setup, echo=FALSE}
```

# Chunk options

`echo = FALSE` to exclude code

```
```{r, show_df, echo = FALSE}
head(df)
```

```

```
A tibble: 6 × 11
name Water_allergy `Eye color` Race `Hair color` Height Publisher `Skin color` Align
<chr> <chr> <chr> <chr> <chr> <dbl> <chr> <chr> <chr>
1 A-Bomb Yes yellow Human No Hair 203 Marvel Comics - good
2 Abe Sapien Yes blue Ichyo Sapien No Hair 191 Dark Horse Com... blue good
3 Abin Sur Yes blue Ungaran No Hair 185 DC Comics red good
4 Abomination Yes green Human / Radiation No Hair 203 Marvel Comics - bad
5 Abraxas Yes blue Cosmic Entity Black NA Marvel Comics - bad
6 Absorbing Man Yes blue Human No Hair 193 Marvel Comics - bad
```

# Chunk options

`echo = TRUE` to include code and output

```
```{r, show_df_1, echo = TRUE}
head(df)
```

```
head(df)
```

```
## # A tibble: 6 × 11
##   name      Water_allergy `Eye color` Race      `Hair color` Height Publisher      `Skin color` Align
##   <chr>     <chr>        <chr>       <chr>      <chr>       <dbl> <chr>        <chr>       <chr>
## 1 A-Bomb    Yes          yellow     Human      No Hair      203  Marvel Comics -       good
## 2 Abe Sapien Yes          blue      Ichyo Sapien No Hair      191  Dark Horse Com... blue    good
## 3 Abin Sur  Yes          blue      Ungaran    No Hair      185  DC Comics    red    good
## 4 Abomination Yes         green     Human / Radiation No Hair      203  Marvel Comics -       bad
## 5 Abraxas   Yes          blue      Cosmic Entity Black     NA   Marvel Comics -       bad
## 6 Absorbing Man Yes         blue     Human      No Hair      193  Marvel Comics -       bad
```

Chunk options

include = FALSE to exclude entire chunk content (code and output)

```
```{r, show_df_2, include = FALSE}
head(df)
```

# Chunk options

**results = "hide"** to include only code (no output)

```
```{r, show_df_3, results = "hide"}  
head(df)
```

```
head(df)
```

Chunk options

Cache results, useful for complex analyses:

```
```{r ... cache=TRUE}
```

Prevent evaluation:

```
```{r ... eval=FALSE}
```

Show results as produced by code (no transformation with **pandoc**, necessary for some packages (e.g., `stargazer`)):

```
```{r ... results="asis"}
```

Prevent showing warnings, messages, errors:

```
```{r ...error=FALSE, message=FALSE, warning=FALSE}
```

The Setup Chunk

It is recommended to use the first code chunk for general setup, where you can

- define your own defaults for chunk options, with `knitr::opts_chunk$set()`
- load the necessary packages
- import raw data

```
```{r, setup, include=FALSE}
chunk option defaults
knitr::opts_chunk$set(echo=FALSE, message=FALSE)
packages
library(dplyr)
library(ggplot2)
library(stargazer)
library(readxl)
```

# The Data Chunk

Second chunk for the main operations on raw data

- cleaning, transformations

```
```{r, data, ...}
df <- read_excel("data/heroes_information.xlsx") %>%
  na_if(., -99) %>%
  mutate(bmi = round(Weight/((Height/100)**2)), 2) %>%
  rename(., Water_allergy = Gender) %>%
  mutate(Water_allergy = recode(Water_allergy, Male = "Yes", Female = "No")) %>%
  select(., -...1, -`2`)
```
```

# Inline Code

If we multiply `_pi_` by `5`, we get ``r pi * 5``.

If we multiply `pi` by `5`, we get `15.7079633`.

The average height of a superhero `in` the dataset is ``r mean(df$Height, na.rm = T)``, which would round to ``r round(mean(df$Height, na.rm = T), digits = 1)``.

The average height of a superhero in the dataset is `186.7263056`, which would round to `186.7`.

Only ``r nrow(subset(df, Height < 100))`` superheros `in` the dataset are Smallings.

Only **9** superheros in the dataset are Smallings.

# *Figures*

# Figures — Markdown Syntax

The syntax `![Figure Caption](figure.extension)` embeds images, and/or figures into .Rmd documents

```
![A superhero's best friend](img/superdog.jpg)
```



Figure 1: A superhero's best friend.

# Figures — `knitr`

The `knitr` package offers a capable alternative with the `include_graphics()` function

```
```{r, superdog, echo=FALSE, fig.cap="A superhero's best friend."}
knitr::include_graphics("img/superdog.jpg")
```
```



Figure 1: A superhero's best friend.

# Figures — knitr

Size is defined with the chunk options `out.width` or `out.height`

```
```{r ... out.width="15%"}  
knitr::include_graphics("img/superdog.jpg")
```



```
```{r ... out.width="40%"}  
knitr::include_graphics("img/superdog.jpg")
```



# Plots - *ggplot2*

# ggplot2

- 1) The `ggplot` function and the `data` argument

```
ggplot(data = df)
```

# ggplot2

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2) The mapping aesthetics, or `aes`; most importantly, the variable(s) that we want to plot

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ggplot(data = df,
 mapping = aes(x = Height, y = Weight, color = Alignment))
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```
ggplot(data = df,
 mapping = aes(x = Height, y = Weight, color = Alignment))
```

3) The geometric objects, or `geom`; the visual representations

```
ggplot(data = df,
 mapping = aes(x = Height, y = Weight, color = Alignment)) +
 geom_point()
```

# ggplot2

Put the code in a chunk, and give it a caption

```
```{r, scatterplot, `fig.cap = "Superheros size"`}
ggplot(data = df,
       mapping = aes(x = Height, y = Weight, color = Alignment)) +
  geom_point()
...````
```

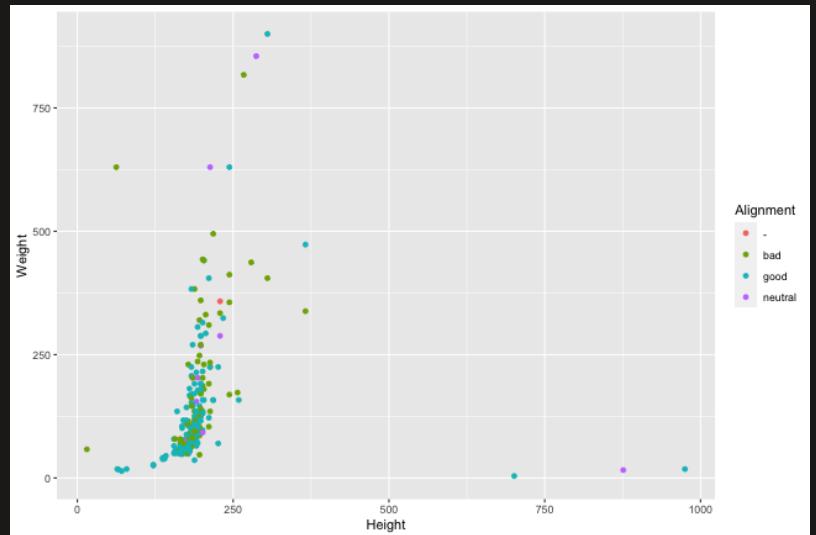


Figure 1. A scatterplot of superheros' height and weight.

ggplot2

Add facets for subgroups, e.g., branch

```
```{r, scatterplot, fig.cap = "A scatterplot of superheros' height  
ggplot(data = df,
 mapping = aes(x = Height, y = Weight, color = Alignment)) +
 geom_point() +
 facet_wrap(. ~ Water_allergy)
...````
```

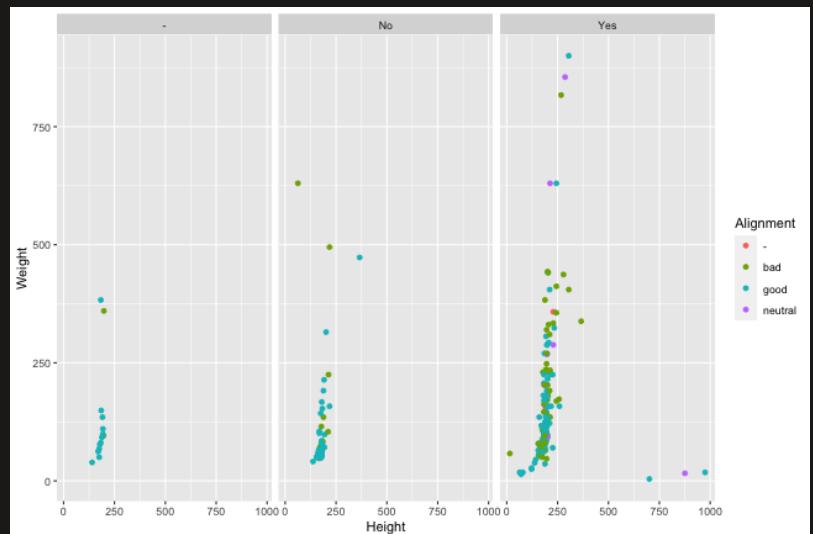


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# ggplot2

Add facets for subgroups, e.g., branch

```
```{r, scatterplot, fig.cap = "A scatterplot of superheros' height  
ggplot(data = df,  
        mapping = aes(x = Height, y = Weight, color = Alignment)) +  
        geom_point() +  
        facet_wrap(. ~ Water_allergy)  
...````
```

Every plot you can think of - you can create it in R! See [R Graph Gallery!](#)

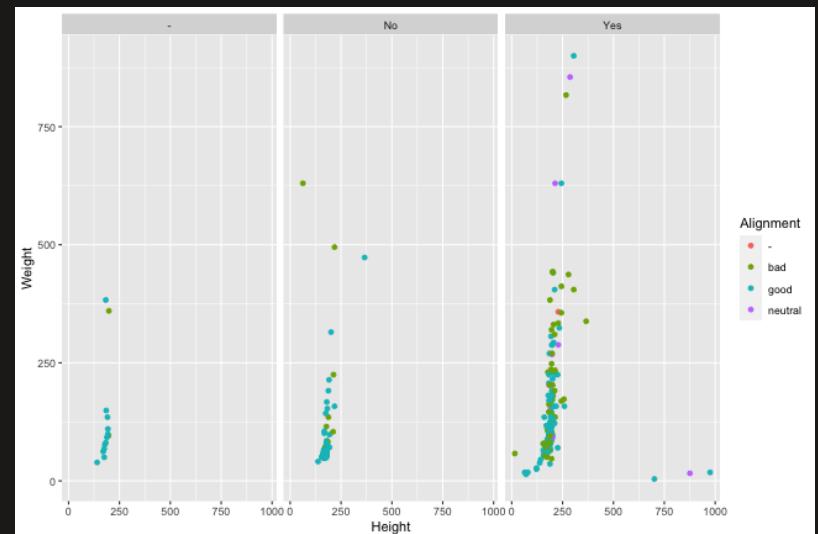


Figure 1. A scatterplot of superheros' height and weight.

ggplot2

Add facets for subgroups, e.g., branch

```
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...````
```

Every plot you can think of - you can create it in R! See [R Graph Gallery!](#)  
Also have a look at the [plotly graphical library](#).

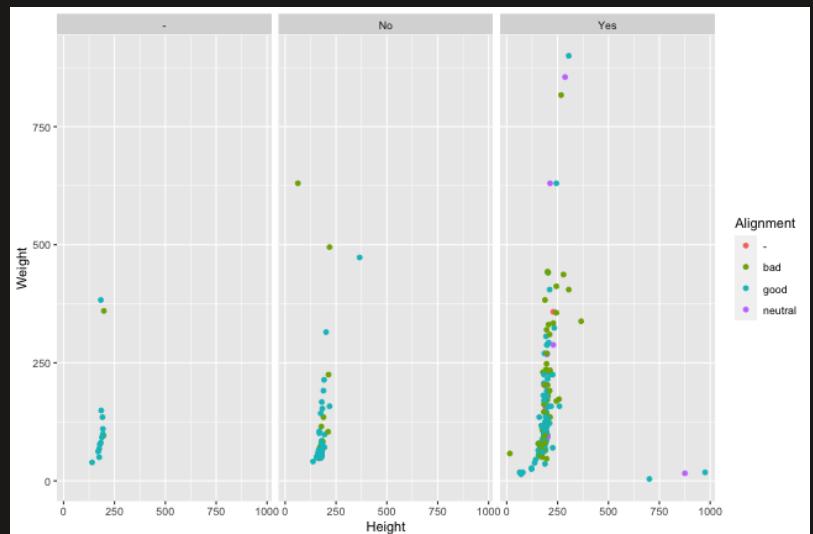


Figure 1. A scatterplot of superheros' height and weight.

# Tables

# Tables with **stargazer**

- A capable package for creating at least three kinds of tables
  - raw data, in columns and rows
  - descriptive/summary statistics
  - regression models

# Tables with `stargazer`

- The `stargazer` package requires specific settings
  - in the chunk options
  - and, in the `type` argument of the `stargazer()` function
- These settings depend on the desired output format,<sup>\*</sup> as shown below

Output	Chunk Option	Type Argument
LaTex / PDF	<code>results="asis"</code>	<code>latex</code>
HTML	<code>results="asis"</code>	<code>html</code>
Word	<code>comment=""</code>	<code>text</code>

<sup>\*</sup> The following slides use the setting for LaTex and PDF outputs.

# Tables with `stargazer`

- `stargazer` tables look slightly different in different output formats
- workarounds for Word:
  - `knit` to HTML as well as Word, copy the tables from HTML to Word
  - `knit` to PDF, open the PDF in Word
  - use a different package to create tables, such as `huxtable`

# Example data table with **stargazer**

Table of first three rows of the dataset

```
```{r, data_table, echo=FALSE, results="asis"}  
stargazer(data = head(df, n = 3), type = "latex", summary = FALSE)
```

Example data table with **stargazer**

Table of first three rows of the dataset

```
```{r, data_table, echo=FALSE, results="asis"}  
stargazer(data = head(df, n = 3), type = "latex", summary = FALSE)
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Notice the options of the chunk and the arguments of the function

# Example data table with **stargazer**

Table of first three rows of the dataset

```
```{r, data_table, echo=FALSE, results="asis"}  
stargazer(data = head(df, n = 3), type = "latex", summary = FALSE)
```

Notice the options of the chunk and the arguments of the function

- with `results="asis"`, `knitr` will pass through results without reformatting them
- they should remain LaTeX (`type = "latex"`) because our outcome document is PDF, converted from LaTeX
- with `summary = FALSE`, the table will present the data, not its descriptive statistics

Example data table with **stargazer**

Table of the first three rows of the dataset

```
```{r, data_table, echo=FALSE, results="asis"}  
stargazer(data = head(df, n = 3), type = "latex", summary = FALSE)
```

Table 1:

	name	Water_allergy	Eye color	Race	Hair color	Height	Publisher	Skin color	Alignment	Weight	bmi
1	A-Bomb	Yes	yellow	Human	No Hair	203	Marvel Comics	-	good	441	107
2	Abe Sapien	Yes	blue	Icthyo Sapien	No Hair	191	Dark Horse Comics	blue	good	65	18
3	Abin Sur	Yes	blue	Ungaran	No Hair	185	DC Comics	red	good	90	26

# Example regression table with **stargazer**

Create a table of regression models instead

```
```{r, regression_table, echo=FALSE, results="asis"}  
stargazer(data = lm(Height ~ Weight, data = df),  
          type = "html", header = FALSE,  
          title = "Regression Results")  
...````
```

Regression Results	
<i>Dependent variable:</i>	
	Height
Weight	0.109*** (0.025)
Constant	174.865*** (3.843)
<hr/>	
Observations	490
R ²	0.037
Adjusted R ²	0.035
Residual Std. Error	57.935 (df = 488)
F Statistic	18.971 *** (df = 1; 488)
<hr/>	
Note:	* p<0.1; ** p<0.05; *** p<0.01

Summary statistics with `vtable`

```
library(vtable)
df %>% select(Alignment, bmi) %>% st(.)
```

Summary Statistics								
Variable	N	Mean	Std. Dev.	Min	Pctl. 25	Pctl. 75	Max	
Alignment	734							
... -	7	1%						
... bad	207	28.2%						
... good	496	67.6%						
... neutral	24	3.3%						
bmi	490	38.557	133.965	0	21	30	2510	

There are many alternatives for creating tables in R Markdown, see `knitr`, `kableExtra`, and `huxtable` packages.

Structure your project

Structure your project

Minimal example

```
: └── example_project
    ├── R
    ├── README.md
    └── analysis
        └── template.Rmd
    ├── data
    ├── docs
    ├── example_project.Rproj
    ├── output
    └── .gitignore
```

- **R** - Resuable R code (functions etc.)
- **analysis** - R Markdown analysis files
- **docs** - Rendered analysis reports
- **data** - (Raw) data used for analysis
- **output** - Output files (e.g., figures, plots...)
- **README.md** - project description (will be displayed if added to GitHub)
- **.gitignore** - The purpose of gitignore files is to ensure that certain files remain untracked (not added when your code is pushed to GitHub); e.g., data that is not anonymized, irrelevant files

Easy project structuring with cookiecutter

1. Install cookiecutter
2. Generate a new analysis directory: type **cookiecutter**
gh:lazappi/cookiecutter-r-analysis (in  terminal )

Useful folders to add

Subfolders for `/data`

```
└── data
    ├── raw
    └── processed
```

Subfolders for `/output`

```
└── output
    ├── figs
    └── ...
```

`/config` folder for csl and other custom styling

```
└── config
```

The **R** and **analysis** folders

Your .rmd file lives in the **/analysis** folder. This is where the final script is executed and the final output (e.g., a pdf) is created.

The **/R** folder is used for custom functions that you load in the main .rmd file via `source("remote-functions.R")` at the top of the file.

The **R** and **analysis** folders

Your .rmd file lives in the **/analysis** folder. This is where the final script is executed and the final output (e.g., a pdf) is created.

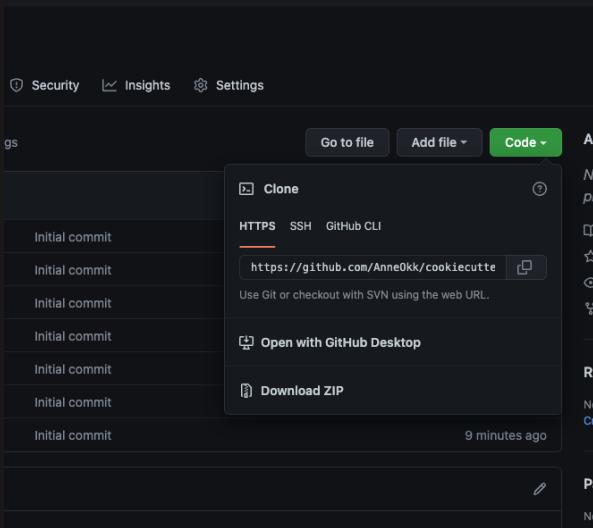
The **/R** folder is used for custom functions that you load in the main .rmd file via **source("remote-functions.R")** at the top of the file.

The setup chunk becomes:

```
```{r, setup, include=FALSE}
chunk option defaults
knitr::opts_chunk$set(echo=FALSE, message=FALSE)
packages
library(dplyr)
library(ggplot2)
library(stargazer)
library(readxl)
custom functions
source("remote-functions.R")
```
```

Alternative: Easy project structuring with templates from GitHub

- Search for an example project structure that serves the purpose, e.g., [here](#).
- Download the directory
- save at desired location



Next level stuff: R packages



Building R packages is beyond the scope of the workshop but definitely useful to get into. See [R Packages Book](#) for more information.

R Markdown templates

Setup for APA7 articles:

- install and load the papaja library

```
remotes::install_github("crsh/papaja@devel")
```

- Install the APA 7 (or another style) document class [here](#)
- Move csl into the correct folder (e.g., `/config`)
- Open an R Markdown APA template in your `/analysis` folder

Add this to the YAML header (for APA7 style, see also [here](#)):

```
header-includes:
  - |
    \makeatletter
    \renewcommand{\paragraph}{\@startsection{paragraph}{4}{\parindent}%
      {0\baselineskip \@plus 0.2ex \@minus 0.2ex}%
      {-1em}%
      {\normalfont\normalsize\bfseries\typesectitle}}
    \renewcommand{\ subparagraph}[1]{\@startsection{subparagraph}{5}{1em}%
      {0\baselineskip \@plus 0.2ex \@minus 0.2ex}%
      {-\z@\relax}%
      {\normalfont\normalsize\bfseries\itshape\hspace{\parindent}\#1\textit{\addperi}}{\relax}}
    \makeatother

csl           : "../config/apa.csl" #< path to csl file
documentclass : "apa7"
```

Add this to the YAML header (to knit output to `docs` folder):

```
knit: (function(inputFile, encoding) {
  rmarkdown::render(inputFile, encoding = encoding, output_dir = "../docs") })
```

Yey, you're all set!

Your turn!



Create an R project for your analysis (~ 45 min.)

1. Use `cookiecutter` to create a project from scratch or use a project template that serves the purpose, e.g., [The Example Project](#)
2. Create an R Markdown template that you would like to use (e.g., APA7 paper with `papaja` package)
3. If you already have an analysis script, move the files into the correct folders (e.g., raw data into `data/raw` subfolder, custom R functions into `R` folder) ! ! Do not forget to refer to the files using relative paths ! !
4. 🎨 KNIT to html or pdf