

# Communicating with Data via R Markdown

## Reproducible Reports

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These slides are viewed best by Chrome and occasionally need to be refreshed if elements did not load properly. See here for PDF [\[PDF\]](#).

# In a nutshell

R Markdown integrates **text + code** in one source document with ability to knit to many output formats (via Pandoc).



# Text in Markdown

```
# Header 1
## Header 2

- Unordered list 1
- Unordered list 2

1. Ordered list 1
1. Ordered list 2

This is italic. *This too.*
This is bold. **This too.**
*This is bold & italic.*
```

## Output

# Header 1

## Header 2

- Unordered list 1
- Unordered list 2

1. Ordered list 1
2. Ordered list 2

*This is italic. This too. This is bold.*  
**This too. This is bold & italic.**

# Shortcut for inserting code chunk

In RStudio .Rmd  press

- Mac:  +  + 
- PC:  +  + 

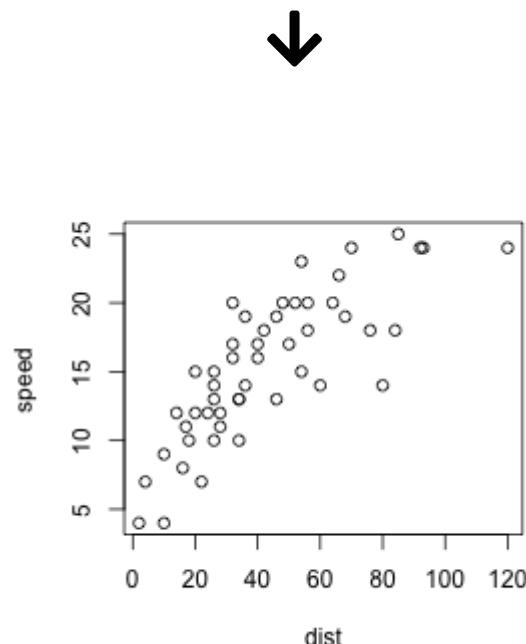
to insert a chunk of R code

```
```{r}  
...  
...
```



# Chunk options: echo & eval

```
```{r, echo = FALSE}  
plot(speed ~ dist, cars)  
```
```



```
```{r, eval = FALSE}  
plot(speed ~ dist, cars)  
```
```

↓

```
plot(speed ~ dist, cars)
```

There are many more **chunk** options.

Can you name 5 other ones?

Hint: <https://yihui.name/knitr/options/>

(We'll explore some later.)



# Valid chunk options

- Chunk options must be written in **one line**, i.e. no line break.
- All option values must be **valid R expressions**. Exception is the chunk name. E.g.
  - `fig.path = figures/` is not valid but  
`fig.path = "figures/"` is valid
  - `eval = true` is not valid but  
`eval = runif(1) > 0.5` is valid



# Chunk names (or labels)

The chunk below is called `plot1`.

```
```{r plot1}
ggplot(cars, aes(dist, speed)) + geom_point()
```
```

All chunks have a label regardless of whether it is explicitly supplied or not.

**⚠** Do not include spaces, "\_" or punctuation marks in your chunk name!



# Inline R Commands

```
Today's date is `r Sys.Date()`.
```

Today's date is 2019-10-03.

```
The value of $pi$ is `r pi`.
```

The value of  $\pi$  is 3.1415927.

- Note: the inline command needs to be R commands.
- Inline command does *not* echo and always evaluates.

# Go through

- challenge-02.Rmd
- challenge-03.Rmd
- challenge-04.Rmd
- challenge-05.Rmd
- challenge-06.Rmd



25 : 00

# R Markdown is not just for R

```
```{python, echo = FALSE}
a = [1, 2, 3]
a[0]
```
```

```
```{bash, echo = FALSE}
date +%B
```
```



```
## 1
```

```
## October
```



Can you name some other engines?

Hint:

<https://yihui.name/knitr/demo/engines/>



# YAML - YAML Ain't Markup Language

Basic format

```
---
```

**key: value**

```
---
```

Example

```
---
```

**title: "Communicating with Data via R Markdown"**

**subtitle: "Reproducible Reports"**

**author: "Emi Tanaka"**

**date: "`r Sys.Date()`"**

**output: html\_document**

```
---
```

There must be a space after ":"!

# Metadata

All YAML data are stored in `rmarkdown::metadata` as list.

```
rmarkdown::metadata$title
```

```
## [1] "Communicating with Data via R markdown"
```

```
rmarkdown::metadata$author
```

```
## [1] "Emi Tanaka"
```

# Default (minimal) html output

```
<!DOCTYPE html>
<html xmlns="http://www.w3.org/1999/xhtml">

<head>
<meta name="author" content="Emi Tanaka" />
<meta name="date" content="2019-10-04" />
<title>Communicating with Data via R Markdown</title>
</head>

<body>
<h1 class="title toc-ignore">Communicating with Data via R Markdown</h1>
<h3 class="subtitle">Reproducible Reports</h3>
<h4 class="author">Emi Tanaka</h4>
<h4 class="date">2019-10-04</h4>
</body>
</html>
```

output

Communicating  
with Data via R  
Markdown

Reproducible Reports

Emi Tanaka

2019-09-23

html meta data

Default html template add special YAML  
key values to file automatically

# YAML structure

- White spaces indicate structure in YAML - don't use tabs though!
- Same as R, you can comment lines by starting with #.
- YAML is case sensitive.
- A key can hold multiple values.

```
key:  
  - value 1  
  - value 2
```

```
key: [value 1, value 2]
```

# YAML with multiple key values

```
---
```

```
title: "Communicating with Data via R Markdown"
author:
  - "Emi Tanaka"
  - "Accomplice"
output: html_document
---
```

output

Communicating  
with Data via R  
Markdown

Emi Tanaka

Accomplice

```
<body>
<h1 class="title toc-ignore">Communicating with Data via R Markdown</h1>
<h4 class="author">Emi Tanaka</h4>
<h4 class="author">Accomplice</h4>
</body>
```

# key can contain keys

```
---
```

```
output:  
  html_document:  
    toc: true  
    toc_float: true
```

```
---
```



What does this do?

(Note: white space is important)

# Values spanning multiple lines

## output

```
---
```

```
title: >
  this is a\
  single line\

abstract: |
  this value spans\
  many lines and\
  appears as it is\

output: pdf_document
---
```

```
```r rmarkdown::metadata$title```
```r rmarkdown::metadata$abstract```



this is a single line



Abstract



this value spans  
  many lines and  
  appears as it is



this is a single line



this value spans  
  many lines and  
  appears as it is


```

# Go through challenge-07.Rmd



10 : 00

# Parametrized Report

```
---
```

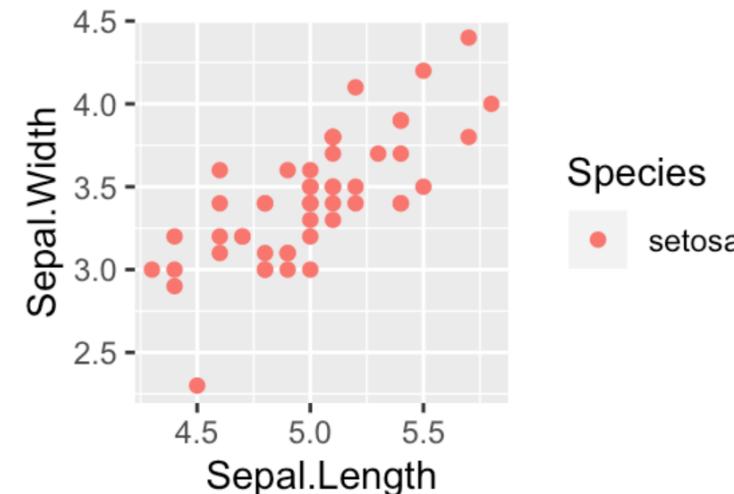
```
title: "Parameterized Report"
params:
  species: setosa
output: html_document
---

```{r, message = FALSE, fig.dim = c(3,2)}
library(tidyverse)
iris %>%
  filter(Species==params$species) %>%
  ggplot(aes(Sepal.Length, Sepal.Width)) +
  geom_point(aes(color=Species))
```
```
```

output

# Parameterized Report

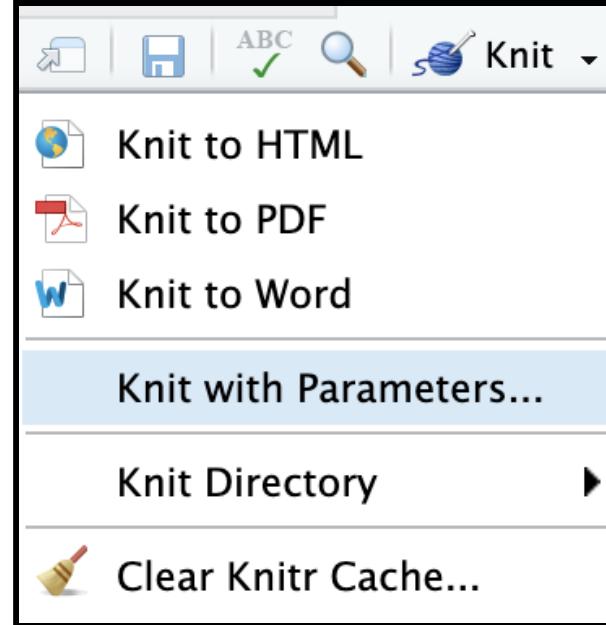
```
library(tidyverse)
iris %>%
  filter(Species==params$species) %>%
  ggplot(aes(Sepal.Length, Sepal.Width)) +
  geom_point(aes(color=Species))
```



# Knit with Parameters

```
---
```

```
title: "Parameterized Report"
params:
  species:
    label: "Species"
    value: setosa
    input: select
    choices: [setosa, versicolor, virginica]
  color: red
  max:
    label: "Maximum Sepal Width"
    value: 4
    input: slider
    min: 4
    max: 5
    step: 0.1
output: html_document
---
```

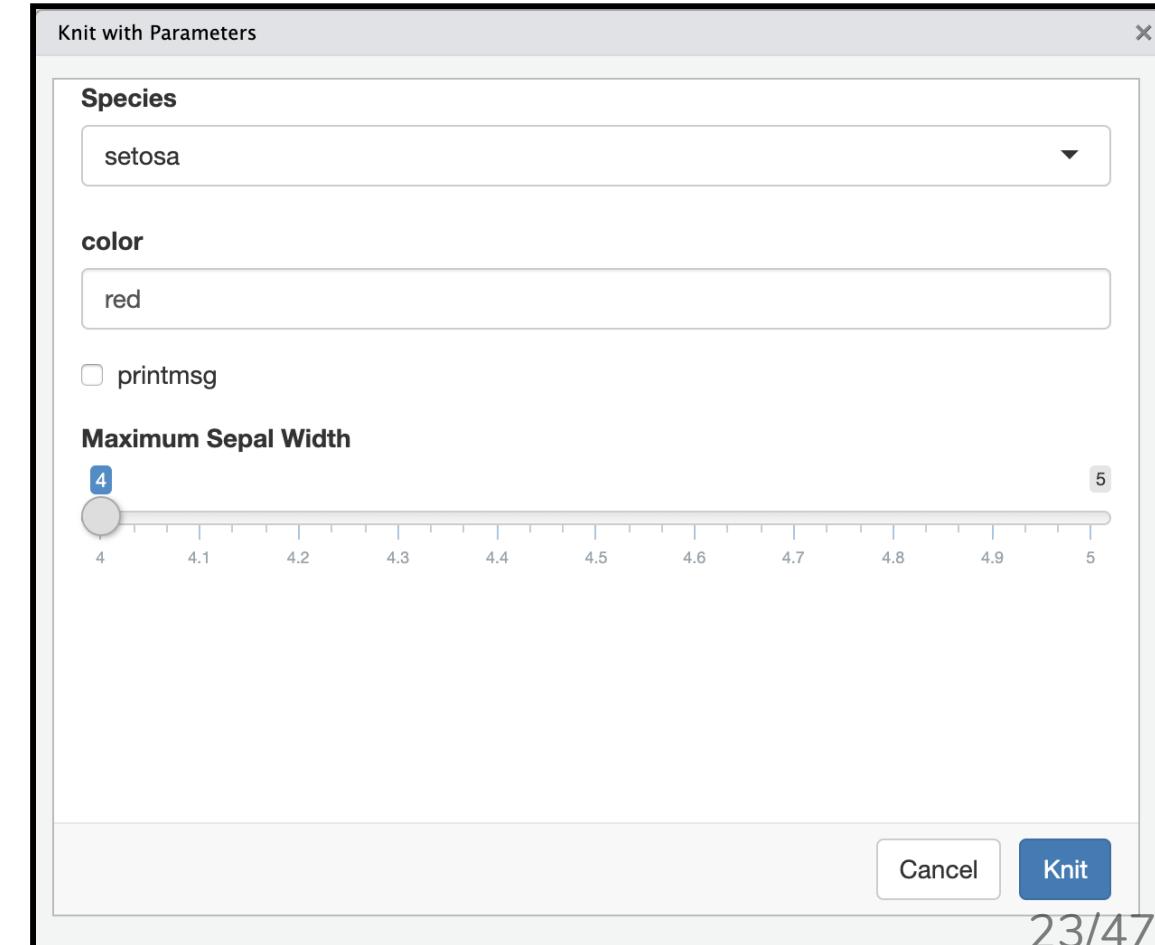


```
```{r, message = params$printmsg, fig.dim = c(
  library(tidyverse)
  iris %>%
    filter(Species==params$species) %>%
    filter(Sepal.Width < params$max) %>%
    ggplot(aes(Sepal.Length, Sepal.Width)) +
    geom_point(aes(color=Species),
               color = params$color)
  ``
```

# Shiny Report Generator

```
---
```

```
title: "Parameterized Report"
params:
  species:
    label: "Species"
    value: setosa
    input: select
    choices: [setosa, versicolor, virginica]
  color: red
  max:
    label: "Maximum Sepal Width"
    value: 5
    input: slider
    min: 4
    max: 5
    step: 0.05
  output: html_document
---
```



# R Markdown via Command Line

demo-render.Rmd

```
---
```

```
title: "Parameterized Report"
params:
  species: setosa
output: html_document
---  
  
```{r, message = FALSE, fig.dim = c(3,2)}
library(tidyverse)
iris %>%
  filter(Species==params$species) %>%
  ggplot(aes(Sepal.Length, Sepal.Width)) +
  geom_point(aes(color=Species))
```
```

You can knit this file via R command by using `render` function:

```
library(rmarkdown)
render("demo-render.Rmd")
```

You can overwrite the YAML values by supplying arguments to `render`:

```
library(rmarkdown)
render("demo-render.Rmd",
       output_format = "pdf_document",
       params = list(species = "virginica"))
```

**Go through**  
challenge-08.Rmd and  
challenge-09.Rmd



10 : 00

# Themes: html\_document

You can change the look of the html document by specifying themes:

- default
- cerulean
- journal
- flatly
- darkly
- readable
- spacelab
- united
- cosmo
- lumen
- paper
- sandstone
- simplex
- yeti
- NULL

```
output:  
html_document:  
theme: cerulean
```

These [bootswatch](#) themes attach the whole bootstrap library which makes your html file size larger.

# prettydoc

prettydoc 📦 is a community contributed theme that is light-weight:

- cayman 
- tactile 
- architect 
- leonids 
- hpstr 

```
output:  
prettydoc::html_pretty:  
theme: cayman
```

See more about it below:

<https://prettydoc.statr.me/>

# rmdformats

rmdformats 📦 contains four built-in html formats:

- `readthedown` 
- `html_clean` 
- `html_docco` 
- `material` 

You can use these formats by simply specifying the output in YAML as below:

```
output: rmdformats::readthedown
```

See more about it below:

<https://github.com/juba/rmdformats>

# rticles - LaTeX Journal Article Templates

- acm 
- acs 
- aea 
- agu 
- amq 
- ams 
- asa 
- biometrics 
- copernicus 
- elsevier 
- frontiers 
- ieee 
- jss 
- mdpi 
- mnras 
- peerj 
- plos 
- pnas 
- rjournal 
- rsos 
- rss 
- sage 
- sim 
- springer 
- tf 

Go to RStudio > File > New File > R Markdown ... > From Template

# External Files in Templating

- When using `rticles`, each journal usually require external files (e.g. `cls` or image files).
- These external components are stored within the package.
- If you are drafting an Rmd template with external components then you need to extract these to your folder first.

## GUI

- RStudio > File > New File > R Markdown ... > From Template

## Command line

```
rmarkdown::draft("file.Rmd",
  template = "biometrics_article",
  package = "rticles")
```

# More customisation needed?

Default templates for many output are found at

<https://github.com/jgm/pandoc-templates>

We'll go through the latex template.



I found this nice latex template online.

You can see it at `main.pdf`.

It was compiled from `main.tex`.

```
%%%%%%%%
% Wenneker Article
% LaTeX Template
% Version 2.0 (28/2/17)
%
% This template was downloaded from:
% http://www.LaTeXTemplates.com
%
% Authors:
% Vel (vel@LaTeXTemplates.com)
% Frits Wenneker
%
% License:
% CC BY-NC-SA 3.0 (http://creativecommons.org/licenses/by-nc-sa/3.0/)
%
%%%%%%%%
%
%-----%
%
%      PACKAGES AND OTHER DOCUMENT CONFIGURATIONS
%-----%
%
\documentclass[10pt, a4paper, twocolumn]{article} % 10pt font size (11 and 12 also possible), A4 paper (letterpaper for US letter) and two column layout (remove for one column)
\input{structure.tex} % Specifies the document structure and loads requires packages
%
%-----%
```

Find `main.tex` and `main.pdf` in `demo` folder.

How do I use this template so that I can  
write contents from an Rmd file  
instead?



# Templating

We will use

```
---
```

```
output:
```

```
  pdf_document:
```

```
    template: main.tex
```

```
---
```

But nothing written in the body shows up in the output!

You need to add `$body$` in the latex template file where you want the body of the md file to appear.

# Templating: few more tweaks

- R Markdown needs a few more special tweaks before `\begin{document}` in latex template:

```
\IfFileExists{bookmark.sty}{\usepackage{bookmark}}{\usepackage{hyperref}}
$if(highlighting-macros)$
$highlighting-macros$
$endif$
```

- These are *minimum* tweaks needed for a LaTeX template.
- You can find common tweaks (including for beamer) at  
<https://github.com/jgm/pandoc-templates>
- You can define your own tweaks but it is better practice to use the ones defined in pandoc template rather than trying to reinvent the wheel.

# How pandoc template works: key

Rmd 

```
---
```

```
title: "COMBINE 2019"
author: "Emi Tanaka"
output:
  pdf_document:
    template: "template.tex"
---
```

YAML meta data can be used by surrounding key with \$.

template.tex 

```
\documentclass{article}
\title{$title$}
\author{$author$}
\date{}

\begin{document}

\maketitle

\end{document}
```

COMBINE 2019  
Emi Tanaka

# How pandoc template works: if statements

Rmd 

template.tex 

```
---
```

```
title: "COMBINE 2019"
author: "Emi Tanaka"
output:
  pdf_document:
    template: "template.tex"
---
```

Simple "if null statements".

```
\documentclass[
  $if(fontsize)$
  $fontsize$,
  $endif$
] {article}
\title{$title$}
\author{$author$}
\date{ }

\begin{document}

\maketitle

\end{document}
```

# How pandoc template works: accessing list

Rmd 

template.tex 

```
---
```

```
title: "COMBINE 2019"
author:
  - name: "Rachel Wang"
    email: "rachel.wang@sydney.edu.au"
  - name: "Connor Smith"
    email: "connor.smith@sydney.edu.au"
output:
  pdf_document:
    template: "template.tex"
---
```

Here it will become

```
\author{Rachel Wang \and Connor Smith}
```

```
\documentclass{article}
\title{$title$}
\author{
$for(author)$
$author.name$$sep$ \and
$endfor$
}
\date{}

\begin{document}

\maketitle

\end{document}
```

# Go through challenge-10.Rmd



05 : 00

# Cross Reference

- When you make a header via Rmd

```
# Some Header
```

an id is created automatically.

- The id is created by replacing **space with –** and making it **all lower case**.
- Now you can link to this header by [some text] (#some-header).
- Cross references work for both pdf and html outputs.

## Demo: header cross-references

```
library(tidyverse)  
library(knitr)
```

### A look at iris

Let's have a look at the `iris` data set. The dataset contains 150 observations. This is cool chicken

### Count

```
iris %>%  
  group_by(Species) %>%  
  count(name = "Count")
```

| Species    | Count |
|------------|-------|
| setosa     | 50    |
| versicolor | 50    |

# Direct Reference for html

- For html output, you can also give a link directly to the relevant section.
- E.g. open `demo-header.html` in the `demo` folder in a web browser.
- Append say `#chicken-data` to the url. It should look like

`demo-header.html#chicken-`  
`data`

- It should have taken you straight to the corresponding header.

# User-defined id

- You can define your own id by appending {#your-id}.

```
# Some header {#header1}
```

- Now you can link to this header with the id header1.
- Note there should be no space in the id name!

# Bibliography

- BibTeX citation style format is used to store references in .bib files.
- Remember that you can get most BibTeX citation for R packages `citation` function. (Scroll below to see the BibTeX citation).

```
citation("xaringan")
```

```
##  
## To cite package 'xaringan' in publications use:  
##  
##   Yihui Xie (2019). xaringan: Presentation Ninja. R package  
##   version 0.9. https://CRAN.R-project.org/package=xaringan  
##  
## A BibTeX entry for LaTeX users is  
##  
##   @Manual{,  
##     title = {xaringan: Presentation Ninja},
```

# Citations

- You can include BibTeX by specifying the `bib` file at YAML as:

```
bibliography: bibliography.bib
```

`[@bibtex-key] → (Author et al. 2019)`

or

`@bibtex-key → Author et al. 2019`

- See `demo-citation.Rmd` in the demo folder.

R Markdown is such an *indispensable* tool for making documents, especially if you have plan to *include statistical output.*

How do you use (or plan to use)  
R Markdown?



# People that made R Markdown possible

The development of R Markdown is largely thanks to

- Yihui Xie  
*Software Engineer at RStudio*  
for `knitr`
- John MacFarlane  
*Professor of Philosophy at UC Berkeley*  
for `pandoc`
- and many contributors behind the development of these tools.



