# A Minimal Book Example

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# Chapter 1

# About

This is a *sample* book written in **Markdown**. You can use anything that Pandoc's Markdown supports; for example, a math equation  $a^2 + b^2 = c^2$ .

### 1.1 Usage

Each **bookdown** chapter is an .Rmd file, and each .Rmd file can contain one (and only one) chapter. A chapter *must* start with a first-level heading: # A good chapter, and can contain one (and only one) first-level heading.

Use second-level and higher headings within chapters like: ## A short section or ### An even shorter section.

The index.Rmd file is required, and is also your first book chapter. It will be the homepage when you render the book.

#### 1.2 Render book

You can render the HTML version of this example book without changing anything:

- 1. Find the Build pane in the RStudio IDE, and
- 2. Click on **Build Book**, then select your output format, or select "All formats" if you'd like to use multiple formats from the same book source files.

Or build the book from the R console:

bookdown::render\_book()

To render this example to PDF as a bookdown::pdf\_book, you'll need to install XeLaTeX. You are recommended to install TinyTeX (which includes XeLaTeX): https://yihui.org/tinytex/.

### 1.3 Preview book

As you work, you may start a local server to live preview this HTML book. This preview will update as you edit the book when you save individual .Rmd files. You can start the server in a work session by using the RStudio add-in "Preview book", or from the R console:

bookdown::serve\_book()

# Chapter 2

# Rstudio

## 2.1 Rstudio shortcuts

keyboard combination	function
opt + _	insert assignment operator <-
ESC  or  ctrl + C	exit + prompt
ctrl + shift + m	add pipe operator "%>%"
ctrl + [/]	indent or unindent
cmd + D	delete one row
cmd + 1	move cursor to console window
cmd + 2	move cursor to editor window
ctrl + shift + S	add 80 hyphens to signal a new
	chapter (Addin)
ctrl + shift + =	add $80$ equals $===$ to signal a new
	Chapter (Addin)
shift + cmd + N	new R script
$\operatorname{cmd} + \uparrow / \downarrow$	in console, get a list of command
	history
$shift + \uparrow / \downarrow$	select one line up/down
fn + F2	view() an object, don't select the
	object
cmd + shift + 1	activate X11() window
ctrl (+ shift) + tab	next (last) tab in scriptor (this applies
	to all apps); hit ctrl first, then shift if
	necessary, last tab

#### Source

keyboard combination	function
cmd + return	Run current line/selection
opt + return	Run current line/selection (retain cursor position)

#### Rmd related

keyboard	
combination	function
cmd + shift + K	Knit rmd
cmd + opt + C	run current code chunk in Rmd
cmd + opt + I	insert code chunks in Rmd, i.e., ```{r} and ```

Q: How to print output in console rather than inline in Rmd?

A: Choose the gear in the editor toolbar and choose "Chunk Output in Console".

#### Set working directory

```
dir_folder <- dirname(rstudioapi::getSourceEditorContext()$path) # get the dir name of
setwd(dir folder) # set as working dir</pre>
```

RStudio projects are associated with R working directories. You can create an RStudio project:

- In a brand new directory
- In an existing directory where you already have R code and data
- By cloning a version control (Git or Subversion) repository

Why using R projects:

- 1. I don't need to use **setwd** at the start of each script, and if I move the base project folder it will still work.
- 2. I have a personal package with a custom project, which creates my folders just the way I like them. This makes it so that the basic locations for data, outputs and analysis is the same across my work.

Double-click on a .Rproj file to open a fresh instance of RStudio, with the working directory and file browser pointed at the project folder.

Q: What is an **R session**? And when do I use it?

A: Multiple concurrent sessions can be useful when you want to:

- Run multiple analyses in parallel
- Keep multiple sessions open indefinitely
- Participate in one or more shared projects

#### Lauch a new project-less RStudio session

```
# run in console
rstudioapi::terminalExecute("open -n /Applications/RStudio.app", show = FALSE)
```

-n Open a new instance of the application(s) even if one is already running.

rstudioapi::terminalExecute(command, workingDir = NULL, env = character(), show = TRUE) tells R to run the system command in quotes.

- command System command to be invoked, as a character string.
- workingDir Working directory for command
- env Vector of name=value strings to set environment variables
- show If FALSE, terminal won't be brought to front

The rstudioapi package provides an interface for interacting with the RStudio IDE with R code. Usingrstudioapi, you can:

- Examine, manipulate, and save the contents of documents currently open in RStudio,
- Create, open, or re-open RStudio projects,
- Prompt the user with different kinds of dialogs (e.g. for selecting a file or folder, or requesting a password from the user),
- Interact with RStudio terminals,
- Interact with the R session associated with the current RStudio instance.

#### Set up Development Tools

https://cran.r-project.org/bin/macosx/tools/

• install Xcode command line tools

```
sudo xcode-select --install
```

• install GNU Fortran compiler

Using Apple silicon (aka arm64, aarch64, M1) Macs Fortran compiler

- Go to https://www.xquartz.org/, download the .dmg and run the installer.
- Verify that build tools are installed and available by opening an R console and running

```
install.packages("pkgbuild")
pkgbuild::check_build_tools()
```

#### Insert Code Session

To insert a new code section you can use the **Code** -> **Insert Section** command. Alternatively, any comment line which includes at least four trailing dashes (-), equal signs (=), or pound signs (#) automatically creates a code section.

#### Define your own shortcuts

https://www.statworx.com/ch/blog/defining-your-own-shortcut-in-rstudio/https://www.r-bloggers.com/2020/03/defining-your-own-shortcut-in-rstudio/Install the shortcut packages.

Add code session separators, --- or ===.

```
install.packages(
    # same path as above
"~/Downloads/shoRtcut_0.1.0.tar.gz",
# indicate it is a local file
repos = NULL)
install.packages(
    # same path as above
"~/Downloads/shoRtcut2_0.1.0.tar.gz",
# indicate it is a local file
repos = NULL)
```

Now go to Tools > Modify Keyboard Shortcuts and search for "dashes". Here you can define the keyboard combination by clicking inside the empty Shortcut field and pressing the desired key-combination on your keyboard. Click Apply, and that's it!

#### 2.2 Dark Theme

https://community.rstudio.com/t/fvaleature-req-word-background-highlight-color-in-find-and-spellcheck/18578/3

https://rstudio.github.io/rstudio-extensions/rstudio-theme-creation.html

https://docs.posit.co/ide/user/ide/guide/ui/appearance.html#creating-custom-themes-for-rstudio

.ace\_marker-layer .ace\_selection Changes the color and style of the highlighting for the currently selected line or block of lines.

.ace\_marker-layer .ace\_bracket Changes the color and style of the high-lighting on matching brackets.

Recommended highlight color: rgba(255, 0, 0, 0.47)

RStudio editor theme directory on Mac:

right click RStudio.app, "Show Package Contents" to navigate to the application folder.

/Applications/RStudio.app/Contents/Resources/resources/themes/ambiance.rstheme

Custom theme (user-defined) folder:

- ~/.config/rstudio/themes/idle\_fingers\_2.rstheme on mac
- viridis-theme

```
/* yaml tag */
.ace_meta.ace_tag {
  color: #2499DA;
}
/* quoted by $...$ and code chunk options */
.ace_support.ace_function {
  color: #55C667;
}
```

### 2.3 Packages Management

#### Install R pakages from source

```
# From local tarball
install.packages(
    # indicate path of the package source file
    "~/Documents/R/UserPackages/shoRtcut2_0.1.0.tar.gz",
    # indicate it is a local file
    repos = NULL)

# From github
install.packages("Rcpp", repos="https://rcppcore.github.io/drat")

Check insatlled packages

# print all installed packages
rownames(installed.packages())
# check if 'ggplot2' is installed
    "ggplot2" %in% rownames(installed.packages())

Update packages

packageVersion("ggplot2") # check package version
```

## update all installed packages in a stated library location, default to `.libPaths()`

Which will ask you for every package if you want to update, to just say yes to everything use ask = FAISE.

install.packages("ggplot2") # update one specific package

update.packages(lib.loc = .libPaths())

```
update.packages(ask = FALSE)
```

Unfortunately this won't update packages installed by devtools::install\_github()

#### Updating all Packages after R update

R packages are missing after updating. So have to save the installed packages and re-install them after updating.

```
## get packages installed
packs <- as.data.frame(installed.packages(.libPaths()[1]), stringsAsFactors = F)
# Save to local
f_name <- "~/Documents/R/packages.csv"
rownames(packs)
write.csv(packs, f_name, row.names = FALSE)
packs <- read_csv(f_name)
packs
## Re-install packages using install.packages() after updating R
install.packages(packs$Package)</pre>
```

R library path /Library/Frameworks/R.framework/Versions/4.2-arm64/Resources/library

- use find.package("ggplot2") to find the location of the source file.
- alternatively, .libPaths()
  - returns the directory within which packages are looked for.

library(package) returns an error if the package doesn't exist.

require(package) returns FALSE if the package doesn't exist. require is designed for use inside other functions.

#### 2.3.1 Put your R package on GitHub

https://jennybc.github.io/2014-05-12-ubc/ubc-r/session2.4\_github.html

- Change to the package directory
- Initialize the repository with git init
- Add and commit everything with
  - 1. git add . stage changes;
  - 2. git status optional check staged changes, but yet to submit;
  - 3. and git commit submit staged changes.
- Create a new repository on GitHub
- Connect your local repository to the GitHub one

git remote add origin https://github.com/username/reponame

• Push everything to github

```
git branch -M main git push -u origin main
```

## Chapter 3

## Knit Rmd

R Markdown is a powerful tool for combining analysis and reporting into the same document. R Markdown has grown substantially from a package that supports a few output formats, to an extensive and diverse ecosystem that supports the creation of books, blogs, scientific articles, websites, and even resumes.

#### Nice documentations

- R markdown: The definitive guide. provides detailed references
- R markdown cookbook concise and covers essential functions, with examples.

#### Quick takeaways:

- Can still use horizontal separator ctrl + shift + S for dashed lines and ctrl + shift + = for equals
- Headers must have one empty line above and below to separate it from other text

#### YAML metadata

Q: What is YAML?

A: YAML is a human-friendly data serialization language for all programming languages.

Q: What does YAML do?

A: It is placed at the very beginning of the document and is read by each of Pandoc, **rmarkdown**, and **knitr**.

- Provide metadata of the document.
- located at the top of the file.

• adheres to the YAML format and is delimited by lines containing three three dashes (---).

It can set values of the template variables, such as title, author, and date of the document.

• The output field is used by rmarkdown to apply the output format function rmarkdown::html\_document() in the rendering process.

There are two types of output formats in the **rmarkdown** package: documents (e.g., pdf\_document), and presentations (e.g., beamer\_presentation).

Supported output format examples: html\_document, pdf\_document.

R Markdown documents (html\_documents) and R Notebook documents (html\_notebook) are very similar; in fact, an R Notebook document is a special type of R Markdown document. The main difference is using R Markdown document (html\_documents) you have to knit (render) the entire document each time you want to preview the document, even if you have made a minor change. However, using an R Notebook document (html\_notebook) you can view a preview of the final document without rendering the entire document.

• Many aspects of the LaTeX template used to create PDF documents can be customized using *top-level* YAML metadata (note that these options do not appear underneath the output section, but rather appear at the top level along with title, author, and so on). For example:

```
title: "Crop Analysis Q3 2013"
output: pdf_document
fontsize: 11pt
geometry: margin=1in
```

A few available metadata variables are displayed in the following (consult the Pandoc manual for the full list):

Variable	Description
lang	Document language code
fontsize	Font size (e.g., 10pt, 11pt, or 12pt)
documentclass	LaTeX document class (e.g., article)
classoption	Options for documentclass (e.g., oneside)
geometry	Options for geometry class (e.g., margin=1in)
mainfont, sansfont,	Document fonts (works only with
monofont, mathfont	xelatex and lualatex)

Variable	Description
linkcolor, urlcolor, citecolor	Color for internal links (cross references), external links (link to websites), and citation links (bibliography)
linestretch	Options for line spacing (e.g. 1, 1.5, 3).

- In PDFs, you can use code, typesetting commands (e.g., \vspace{12pt}), and specific packages from LaTeX.
  - 1. The header-includes option loads LaTeX packages.

```
output: pdf_document
header-includes:
- \usepackage{fancyhdr}
---
\pagestyle{fancy}
\fancyhead[LE,R0]{Holly Zaharchuk}
\fancyhead[L0,RE]{PSY 508}
# Problem Set 12
```

2. Alternatively, use extra\_dependencies to list a character vector of LaTeX packages. This is useful if you need to load multiple packages:

```
title: "Untitled"
output:
   pdf_document:
     extra_dependencies: ["bbm", "threeparttable"]
```

f you need to specify options when loading the package, you can add a second-level to the list and provide the options as a list:

```
title: "Untitled"
output:
  pdf_document:
    extra_dependencies:
      caption: ["labelfont={bf}"]
      hyperref: ["unicode=true", "breaklinks=true"]
      lmodern: null
```

Here are some examples of LaTeX packages you could consider using

within your report:

- \* pdfpages: Include full PDF pages from an external PDF document within your document.
- \* caption: Change the appearance of caption subtitles. For example, you can make the figure title italic or bold.
- \* fancyhdr: Change the style of running headers of all pages.
- Some options are passed to Pandoc, such as toc, toc\_depth, and number\_sections. You should consult the Pandoc documentation when in doubt.

```
output:

pdf_document:

toc: true

keep_tex: true
```

\* keep\_tex: true if you want to keep intermediate TeX. Easy to debug. Defaults to false.

We can include variables and R expressions in this header that can be referenced throughout our R Markdown document. For example, the following header defines start\_date and end\_date parameters, which will be reflected in a list called params later in the R Markdown document.

Thus, if we want to use these values in our R code, we can access them via params\$start\_date and params\$end\_date.

Should I use quotes to surround the values?

- Whenever applicable use the unquoted style since it is the most readable.
- Use quotes when the value can be misinterpreted as a data type or the value contains a :.

```
# values need quotes
foo: '{{ bar }}' # need quotes to avoid interpreting as `dict` object
foo: '123'  # need quote to avoid interpreting as `int` object
foo: 'yes'  # avoid interpreting as `boolean` object
foo: "bar:baz:bam" # has colon, can be misinterpreted as key

# values need not quotes
foo: bar1baz234
bar: 123baz
```

### 3.1 Chunk Options

If you want to set chunk options globally, call knitr::opts\_chunk\$set() in a code chunk (usually the first one in the document), e.g.,

```
\``{r, label="setup", include=FALSE}
knitr::opts_chunk$set(
  comment = "#>", echo = FALSE, fig.width = 6
)
```

Full list of chunk options: https://yihui.org/knitr/options/

Chunk options can customize nearly all components of code chunks, such as the source code, text output, plots, and the language of the chunk.

#### Other languages are supproted in Rmd

You can list the names of all available engines via:

```
names(knitr::knit_engines$get())
## [1] "awk"
                        "bash"
                                       "coffee"
## [4] "gawk"
                        "groovy"
                                       "haskell"
   [7] "lein"
                        "mysql"
                                       "node"
                        "perl"
## [10] "octave"
                                       "php"
## [13] "psql"
                        "Rscript"
                                       "ruby"
                        "scala"
                                       "sed"
## [16] "sas"
## [19] "sh"
                        "stata"
                                       "zsh"
## [22] "asis"
                        "asy"
                                       "block"
## [25] "block2"
                        "bslib"
                                       "c"
                                       "comment"
## [28] "cat"
                        "cc"
                                       "dot"
## [31] "css"
                        "ditaa"
## [34] "embed"
                       "eviews"
                                       "exec"
## [37] "fortran"
                        "fortran95"
                                       "go"
                        "is"
                                       "julia"
## [40] "highlight"
                                       "Rcpp"
## [43] "python"
                        "R."
                        "scss"
                                       "sql"
## [46] "sass"
## [49] "stan"
                        "targets"
                                       "tikz"
                        "theorem"
## [52] "verbatim"
                                       "lemma"
## [55] "corollary"
                        "proposition"
                                       "conjecture"
## [58] "definition"
                        "example"
                                       "exercise"
                        "proof"
                                       "remark"
## [61] "hypothesis"
## [64] "solution"
                        "marginfigure"
```

The engines from theorem to solution are only available when you use the **bookdown** package, and the rest are shipped with the **knitr** package.

To use a different language engine, you can change the language name in the chunk header from  ${\tt r}$  to the engine name, e.g.,

```
""
python
x = 'hello, python world!'
print(x.split(' '))
```

For engines that rely on external interpreters such as python, perl, and ruby, the default interpreters are obtained from Sys.which(), i.e., using the interpreter found via the environment variable PATH of the system. If you want to use an alternative interpreter, you may specify its path in the chunk option engine.path.

For example, you may want to use Python 3 instead of the default Python 2, and we assume Python 3 is at /usr/bin/python3

- All outputs support markdown syntax.
- If the output is html, you can write in html syntax.

The **chunk label** for each chunk is assumed to be unique within the document. This is especially important for cache and plot filenames, because these filenames are based on chunk labels. Chunks without labels will be assigned labels like unnamed-chunk-i, where i is an incremental number.

- Chunk label doesn't need a tag, i.e., you only provide the value.
- If you prefer the form tag=value, you could also use the chunk option label explicitly, e.g.,

```
'``{r, label='my-chunk'}
# one code chunk example
'``
```

You may use knitr::opts\_chunk\$set() to change the default values of chunk options in a document.

#### Commonly used chunk options

• Complete list here. Or ?opts\_chunk to get the help page.

Options	Definitions
echo=TRUE	Whether to display the <b>source code</b> in the output
	document. Use this when you want to show the output
	but not the code itself.
eval=TRUE	Whether to evaluate the code chunk.
include=TRUE	Whether to include the chunk <b>output</b> in the output
	document. If FALSE, nothing will be written into the
	output document, but the code is still evaluated and plot
	files are generated if there are any plots in the chunk, so
	you can manually insert figures later.
message=TRUE	Whether to preserve messages emitted by message()

Options	Definitions
warning=TRUE	Whether to show warnings in the output produced by
	warning().
results='markup'	Controls how to display the text results. When
	results='markup' that is to write text output as-is, i.e.,
	write the raw text results directly into the output
	document without any markups. Useful when priting
	stargazer tables.
comment='##'	The prefix to be added before each line of the text
	output. Set comment = '' remove the default ##.
fig.keep='high'	How plots in chunks should be kept. high: Only keep
0 1 0	high-level plots (merge low-level changes into high-level
	plots). none: Discard all plots. all: Keep all plots
	(low-level plot changes may produce new plots). first:
	Only keep the first plot. last: Only keep the last plot.
	If set to a numeric vector, the values are indices of
	(low-level) plots to keep. If you want to choose the second
	to the fourth plots, you could use fig.keep = 2:4 (or
	remove the first plot via fig.keep = -1).
fig.align="center	
fig.pos="H"	A character string for the figure position arrangement to
TIE. POD II	be used in \begin{figure}[].
fig can	Figure caption.
fig.cap	rigure caption.

results='markup' note plural form for results.

• markup: Default. Mark up text output with the appropriate environments depending on the output format. For example, for R Markdown, if the text output is a character string "[1] 1 2 3", the actual output that knitr produces will be:

```
[1] 1 2 3
```

In this case, results='markup' means to put the text output in fenced code blocks ("').

• asis: Write text output as-is, i.e., write the raw text results directly into the output document without any markups.

```
```{r, results='asis'}
cat("I'm raw **Markdown** content.\n")
```

Sometime, you encounter the following error messages when you have R codes within enumerate environment.

You can't use 'macro parameter character #' in horizontal mode.

By default, knitr prefixes R output with ##, which can't be present in your TeX file.

#### Solution:

- specify results="asis" in code chunks.
- hold: Hold all pieces of text output in a chunk and flush them to the end of the chunk.
- hide (or FALSE): Hide text output.