

Center for Conservation Biology | UC Riverside  
Installation Guide

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# Contents

<b>Informational Resource for Software and Application Installation Utilized by CCB Faculty &amp; Staff</b>	<b>5</b>
<b>1 Introduction</b>	<b>7</b>
1.1 Computer requirements . . . . .	7
1.2 Software . . . . .	7
<b>2 R and RStudio Installation</b>	<b>9</b>
2.1 Install or Update R . . . . .	9
2.2 Install or Update R Studio . . . . .	9
2.3 Install Quarto . . . . .	10
2.4 Learn How to Use R & RStudio . . . . .	11
<b>3 Bookdown Guide</b>	<b>13</b>
3.1 Primary Reference Resources . . . . .	13
3.2 Formatting . . . . .	13
3.3 Citations . . . . .	14
3.4 Alt Text for Accessibility . . . . .	14
3.5 Rendering Bookdown to Build & Publish . . . . .	14
<b>4 Git Installation &amp; GitHub Account</b>	<b>17</b>
4.1 Git . . . . .	17
4.2 GitHub . . . . .	17
4.3 Configure Git . . . . .	18
4.4 Personal Access Token . . . . .	18
<b>5 Project Management Tools</b>	<b>25</b>
5.1 Trello . . . . .	25
5.2 Slack . . . . .	25
5.3 Google Apps . . . . .	26
5.4 Zotero . . . . .	26



# Informational Resource for Software and Application Installation Utilized by CCB Faculty & Staff



Chino Canyon Wildflowers, Coachella Valley, California. **Image Credit:**  
Coachella Valley Mountains Conservancy: *Bill Havert*



# Chapter 1

## Introduction

This is an installation and account set up reference guide for the Center for Conservation Biology team. Team members may contribute to this reference file as we expand the tools utilized during research efforts.

### 1.1 Computer requirements

Work computers (laptop or desktop) operating systems should either be Windows or macOS. Please note that Windows based machine it required to run ESRI ArcDesktop and ArcPro software. Both Windows and macOS may utilize ESRI ArcOnline tools.

For **Mac** users, update macOS to the newest supported version. Navigate to System Preferences → Software Update.

For **PC** users, ensure you have Windows 10 or 11 installed. If not, request a Windows key from UCR IT at UC Riverside ServiceLink

### 1.2 Software

Software covered in this reference guide includes:

- git
- GitHub
- Google Apps
- R
- RStudio
  - Quarto
  - Bookdown
- Slack
- Trello

- Trello 4 Slack
- Zotero

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Thank you to UC Santa Barbara's Bren School of Environmental Science & Management and National Center for Ecological Analysis and Synthesis (NCEAS) staff for providing many of the resources listed in this reference guide. Information was made available on the UCSB-MEDS GitHub page.



## Chapter 2

# R and RStudio Installation

### 2.1 Install or Update R

**R** is a programming language and environment used for statistical computing and graphics. For more information, please visit [What is R](#).

To install R, visit [cloud.r-project.org](http://cloud.r-project.org) to download the most recent version for your operating system.

### 2.2 Install or Update R Studio

RStudio is a software (considered an Integrated Development Environment, or IDE) that provides R programmers with an easy-to-use interface for coding in R.

**Note:** RStudio will not work without R installed, and you won't particularly enjoy using R without having RStudio installed. Be sure to install both!

- **New install:** To install RStudio, visit [rstudio.com/products/rstudio/](http://rstudio.com/products/rstudio/). Download the free ("Open Source Edition") Desktop version for your operating system.
- **Update:** If you already have RStudio and need to update: Open RStudio, and under 'Help' in the top menu, choose 'Check for updates.' If you have the most recent release, it will return 'No update available. You are running the most recent version of RStudio.' Otherwise, you should follow the instructions to install an updated version.

Open RStudio (logo you'll click on shown below). **If you are prompted to install Command Line Tools, do it.**



### Mac Users

There may be a need to install command line tools and XQuartz:

- To install command line tools (if you're not automatically prompted), run in the Terminal tab in RStudio: `xcode-select --install`
- Visit [xquartz.org](http://xquartz.org) to download & install XQuartz.

## 2.3 Install Quarto

*This is an optional tool within R Studio that is extremely powerful, but not required for your job*

Quarto is a scientific publishing tool built on Pandoc that allows R, Python, Julia, and ObservableJS users to create dynamic documents, websites, books

and more.

As of *July 2022*, Quarto comes pre-installed in R Studio (v2022.07). If you haven't updated your R Studio IDE (and concerned about doing so), you can install Quarto separately.

- Download Quarto [here](#) and install
- To use Quarto through the RStudio IDE, be sure to have at least version v2022.02 installed (see directions in step 2, above)

Learn more about Quarto [here](#).

## 2.4 Learn How to Use R & RStudio

There are a lot of resources out there to help you learn how to use R and the RStudio IDE (YouTube, Googling, StackOverflow, etc...). This is a short list of primary resources for you to reference. It is highly recommended to join R community Slack channels, R User community groups (such as R-Ladies), and UC Riverside Data Science clubs!

- ACM at UCR - UC Riverside community dedicated to technical, professional, and personal development in the context of computer science.
- R for Data Science
- R Markdown Cookbook
- Cookbook for R
- Advanced R
- R Studio (posit) Book Catalog
- R for Data Science Online Learning Community
- R OpenSci
- RStudio Cloud - interactive tutorials to learn data science basics.
- R Studio Cheatsheets - invaluable tool to learn how to use various R packages.
- Swirl - R package that is a built in tutorial.



## Chapter 3

# Bookdown Guide

The first step to edit and add to this bookdown is to install the **bookdown** package from CRAN or Github. In the RStudio console, run the following:

```
install.packages("bookdown")  
# or the development version  
# devtools::install_github('rstudio/bookdown')
```

### 3.1 Primary Reference Resources

Here is a list of resources to learn how to use and edit in bookdown

- Bookdown Package Documentation
- Authoring Books with R Markdown
- R Markdown Cookbook
- R Markdown: The Definitive Guide

---

The following information is directly taken from the *bookdown* package (Xie, 2022).

### 3.2 Formatting

You can use anything that Pandoc's Markdown supports, e.g., a math equation  $a^2 + b^2 = c^2$ .

Remember each Rmd file contains one and only one chapter, and a **chapter** is defined by the first-level heading #.

You can label chapter and section titles using `{#label}` after them, e.g., we can reference Chapter 1. If you do not manually label them, there will be automatic

labels anyway, e.g., Chapter ??.

Figures and tables with captions will be placed in `figure` and `table` environments, respectively.

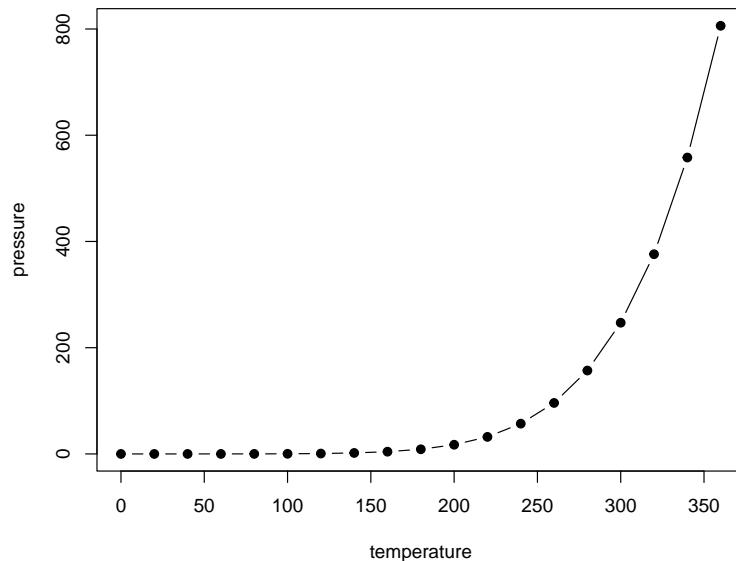


Figure 3.1: Here is a nice figure!

Reference a figure by its code chunk label with the `fig:` prefix, e.g., see Figure 3.1. Similarly, you can reference tables generated from `knitr::kable()`, e.g., see Table 3.1.

### 3.3 Citations

You can easily write citations using `.bib` files within this repository formatted using BibTEX. For example, the **bookdown** package (Xie, 2022) in this reference book, which was built on top of R Markdown and **knitr** (Xie, 2015).

### 3.4 Alt Text for Accessibility

Use the `knitr` package to add alt text to graphics in R Markdown files

### 3.5 Rendering Bookdown to Build & Publish

In your Console, type either of these commands depending on which type of render you prefer:

Table 3.1: Here is a nice table!

Sepal.Length	Sepal.Width	Petal.Length	Petal.Width	Species
5.1	3.5	1.4	0.2	setosa
4.9	3.0	1.4	0.2	setosa
4.7	3.2	1.3	0.2	setosa
4.6	3.1	1.5	0.2	setosa
5.0	3.6	1.4	0.2	setosa
5.4	3.9	1.7	0.4	setosa
4.6	3.4	1.4	0.3	setosa
5.0	3.4	1.5	0.2	setosa
4.4	2.9	1.4	0.2	setosa
4.9	3.1	1.5	0.1	setosa
5.4	3.7	1.5	0.2	setosa
4.8	3.4	1.6	0.2	setosa
4.8	3.0	1.4	0.1	setosa
4.3	3.0	1.1	0.1	setosa
5.8	4.0	1.2	0.2	setosa
5.7	4.4	1.5	0.4	setosa
5.4	3.9	1.3	0.4	setosa
5.1	3.5	1.4	0.3	setosa
5.7	3.8	1.7	0.3	setosa
5.1	3.8	1.5	0.3	setosa

```
bookdown::render_book("index.Rmd", "bookdown::gitbook") bookdown::render_book("index.Rmd",  
"bookdown::pdf_book")
```

To compile to PDF, you need XeLaTeX. It is recommended to install TinyTeX (which includes XeLaTeX): <https://yihui.name/tinytex/>.



## Chapter 4

# Git Installation & GitHub Account

### 4.1 Git

Check to see if your computer already has git:

- Open RStudio
- In the terminal, run the following command:

```
which git
```

If `git` is already installed, the return to the above command should return a filepath (ex: `/usr/local/bin/git`).

- If there is no response, download and install `git` from here: [git-scm.com/downloads](https://git-scm.com/downloads). Select the default settings within the prompts **except** the default to **master** branch. This branch is being phased out, so select the option that let's you select alternative branches (ex: `main`).

### 4.2 GitHub

GitHub is a internet based code hosting platform for collaboration and version control. GitHub lets you (and others) work together on projects.

Navigate to [github.com](https://github.com), and create an account! Please use either your work or personal email account. You can add several emails to your account, and assign a particular email as the primary email for the account.

Review this article on choosing a GitHub username: [happygitwithr.com/github-acct.html](https://happygitwithr.com/github-acct.html).

### 4.3 Configure Git

Once Git is installed and your GitHub account has been set up, Git needs to be configured on the computer. Configuring Git is required to push & pull commits to GitHub.

- In RStudio, open the terminal.
- Run the following commands separately, pressing **Enter** after each line. Replace username and email with your GitHub account username and email. Make sure to keep the quotes around the username in the code below:

```
- git config --global user.name "Jane Doe"
- git config --global user.email janedoe@example.com
```

- Once you have entered the above command line code, check that the configuration was set correctly. In the RStudio Terminal, type the following command and hit enter:

```
- git config --list --global
```

In the terminal, it should show code similar to this in the Terminal:

```
user.name=janedoe user.email=janedoe@ucr.edu core.excludesfile=/Users/janedoe/.gitignore
```

If when installing or updating Git, the default branch was not set to **main** (it is defaulting to the old **master** branch), you can change this setting globally. In the Terminal again, enter the line below:

```
git config --global init.defaultBranch main
```

For more information on configuring Git: check out this [Git reference](#)

### 4.4 Personal Access Token

Once Git has been configured to commit to your GitHub account, a **Personal Access Token (PAT)** must be created for **each computer you intend to use**. A PAT is an alternative password authentication method for Git to access GitHub accounts.

- In the RStudio Console, install the **usethis** package in R by running the following code:

```
install.packages("usethis")
```

If the **usethis** package is installed correctly, at the end of the stream of text there should be a message similar to the image below:

```
The downloaded binary packages are in  
      /var/folders/ (...random gibberish here...) /downloaded_packages  
> |
```




- Once the **usethis** package is installed, run the following in the RStudio Console:

```
usethis::create_github_token()
```

- Enter your GitHub password when prompted.

This should take you to the **Settings/Developer settings** section of your GitHub account:

[Settings](#) / Developer settings

-  GitHub Apps
-  OAuth Apps
-  Personal access tokens

## New personal access token

Personal access tokens function like OAuth tokens, but can be used over HTTPS, or can be used to [authenticate](#) requests.

### Note

CCB\_PC\_WorkComputer

What's this token for?

### Expiration \*

Custom...

01/31/2023

### Select scopes

Scopes define the access for personal access tokens.

- |                                     |                 |   |
|-------------------------------------|-----------------|---|
| <input checked="" type="checkbox"/> | repo            | Full control of repository                |
| <input checked="" type="checkbox"/> | repo:status     | Access to repository status               |
| <input checked="" type="checkbox"/> | repo_deployment | Access to repository deployment           |
| <input checked="" type="checkbox"/> | public_repo     | Access to public repository               |
| <input checked="" type="checkbox"/> | repo:invite     | Access to repository invite               |
| <input checked="" type="checkbox"/> | security_events | Read access to repository security events |
| <input checked="" type="checkbox"/> | workflow        | Upload and run workflows                  |
| <input checked="" type="checkbox"/> | write:packages  | Upload packages                           |
| <input checked="" type="checkbox"/> | read:packages   | Download packages                         |

- **Note Field** Change the PAT name to a meaningful reference (see image for an example). You may end up creating multiple PATs, so you want to ensure that you know which PAT is designated for each computer | server.
- **Expiration Field**

This is to select a set expiration timeframe for your PAT. Setting an expiration is highly recommended, and GitHub will send you *SEVERAL* emails prior to it expiring to remind you to renew it. Use the drop down to select a set time frame (7 days to 90 days) or create a custom expiration time frame (exactly a year or particular date).

- **Select Scopes Field**

Define access for the Personal Access Token being generated.

It is recommended to select at least the following scopes: - repo - workflow - write:packages - notifications - delete repo - write:discussions - project

To learn more about Scopes, visit the GitHub Scopes for OAuth Apps page.

- Once Scopes are selected, click on the green **Generate token** button:

<input type="checkbox"/> <b>admin:gpg_key</b>	Full control of public user GPG keys
<input type="checkbox"/> <b>write:gpg_key</b>	Write public user GPG keys
<input type="checkbox"/> <b>read:gpg_key</b>	Read public user GPG keys

**Generate token** Cancel

- Copy the generated PAT to your clipboard.
- Paste and Save this PAT in a text file in a secure folder that will NEVER be accessed by other users or the internet. You can create a **private** folder on your personal computer to store these files.
- Return to RStudio Console and run the following command:

```
gitcreds::gitcreds_set().
```

You will be prompted to paste the PAT into the console:

```
> gitcreds::gitcreds_set()

? Enter password or token: REDACTED
-> Adding new credentials...
-> Removing credetials from cache...
-> Done.
```

Paste the PAT at the end of the line `Enter password or token:` and press enter.

- In the console, run:

```
usethis::git_sitrep()
```

This command should return your GitHub account information (see example below).

```
> usethis::git_sitrep()
Git config (global)
• Name: 'juliaparish'
• Email: 'jparish@bren.ucsb.edu'
• Global (user-level) gitignore file: '/Users/julia/.gitignore'
• Vaccinated: FALSE
i See `?git_vaccinate` to learn more
i Defaulting to 'https' Git protocol
• Default Git protocol: 'https'
• Default initial branch name: <unset>
GitHub
• Default GitHub host: 'https://github.com'
• Personal access token for 'https://github.com': '<discarded>'
• GitHub user: 'juliaparish'
```

For more information on PATs, check out GitHub's PAT information page.





## Chapter 5

# Project Management Tools

### 5.1 Trello



Trello is a project management app tool that provides teams the opportunity to create task lists, reference & resource lists, and communicate via tagging! To connect with CCB's Trello Workspace, send a request to CCB's PI. The main board is **CCB Team Projects**.

### 5.2 Slack



Slack is a messaging app that connects teams via channels and direct messages (DMs). To join the CCB Slack workspace, follow this link.

There is a Trello app incorporated in the CCB UCR slack workspace. You can create cards in Slack for the CCB Team Projects Trello workspace. To learn how, follow this Slack connect tool.

Other cool Slack workspaces to join:

- EcoDataScience - an environmental data science study group that started at UC Santa Barbara, but now has an international following!
- R4DS Online Learning Community
- Society for Open, Reliable, and Transparent Ecology and Evolutionary Biology (SORTEE)

### 5.3 Google Apps

The CCB team utilizes a number of Google applications for project management, data & file sharing, and communicating with colleagues.

**Google Calendar:** There is a lab Google calendar, *UCR CCB Palm Desert*. This calendar maintains lab mates work and leave schedules as well as any team meetings. Please make sure this calendar is shared with you once you have obtained your UCR NetId.

**Google Drive:** CCB utilizes *Shared Drive* folders as a file sharing application. You must request access to the shared drive folder from the lab P.I.

**Google Sites:** There is a CCB Google Site that is in draft form at the moment. This site may serve as an intranet or provide public facing information in the future. If you would like edit access, please submit a request to the CCB PI.

### 5.4 Zotero

The Center for Conservation Biology shares research references through Zotero software. Zotero is a citation management tool and allows groups to collect and share references easily. The Zotero browser extension makes it easy to copy citations from webpages.

To install Zotero and the extension, please follow UC Riverside's installation instructions here. *Note:* These instructions are for a **Windows** machine, but it is a similar process for Macs.

## Install Zotero

If you have never installed Zotero, or if you are transitioning from the Firefox-embedded version,

1. Go to [Zotero.com](https://www.zotero.com) and click on

A red rectangular button with the word "Download" in white text.

3. Download **Zotero for Windows**. Select the blue download button, and save the file (if necessary).



4. Run the install executable wherever it downloaded to.

### Install Connectors

For each browser you use, you would want to install a connector, so that you can e

For each browser (all at once or when you are next using it):

1. Go to [Zotero.com](https://www.zotero.com)



8. When prompted click **add extension, or allow or whatever prompting your b**

The next step is to create a Zotero account and enable syncing to enable extension use and to be able to join the CCB Zotero Reference Group. Make sure to use your UCR email address when setting up the Zotero account.

### Syncing Your Account

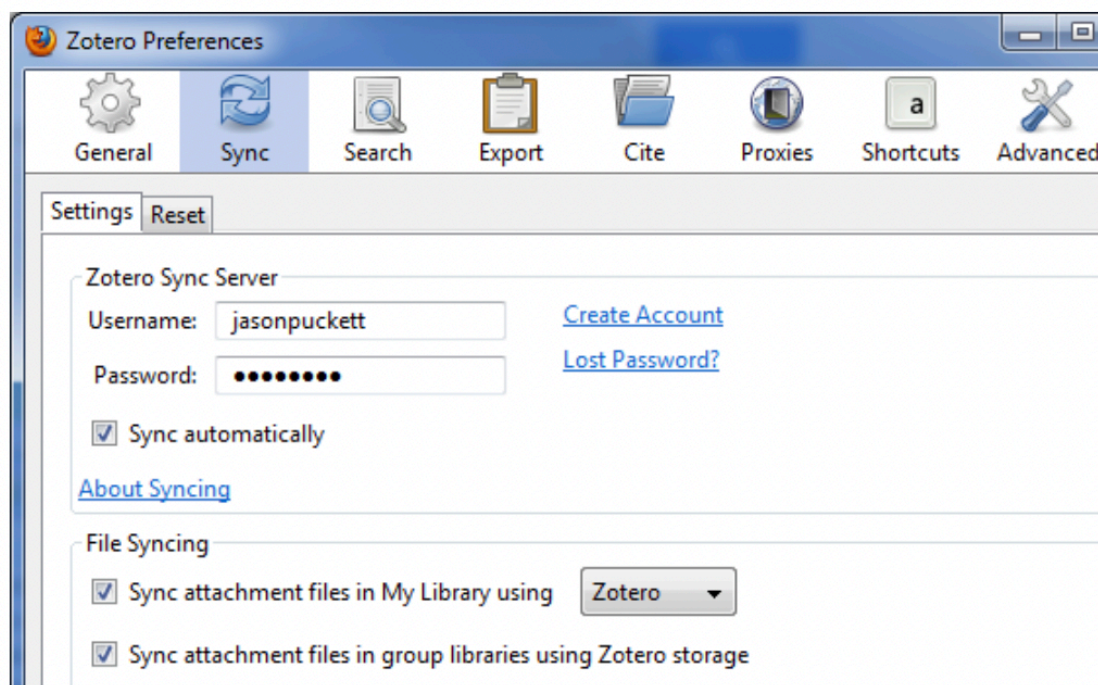
If you're regularly using more than one computer in your research, Zotero's [sync feature](#) can keep your library up to date on all copy of your library on the Zotero.org server and check it for updates whenever you open your library on a different computer. running the same version of Zotero.

First, set up a (free, of course) [Zotero.org user account](#). Then:

- Open Zotero preferences (via the gear menu) and select the Sync tab.
- Enter your Zotero user name and password.
- Check the "sync automatically" box.
- Check both boxes under File Syncing and choose Zotero storage for My Library. This will sync your PDF attachments a
- Click the green circular arrow button at the top right corner of the Zotero window.
- Zotero will upload your library to the server.

Repeat this configuration on each of your computers. Any updates you make on one of your computers will be reflected on the synchronize your library among Windows, Mac and Linux computers.

For more details and help troubleshooting sync problems, check the [Zotero site](#).



To connect to the CCB Zotero Reference Group, navigate to this link. If you do not have access to the Group page, please send a request that you be added to the CCB PI.

For more information on how to use Zotero, check out these links:

- UC Riverside Zotero Tips & Tricks
- Zotero Quick Start Guide
- Zotero Groups
- RStudio Citations - add BibTex citation format in Zotero then quickly use .bib files in R to create references in markdown.

# Bibliography

Xie, Y. (2015). *Dynamic Documents with R and knitr*. Chapman and Hall/CRC, Boca Raton, Florida, 2nd edition. ISBN 978-1498716963.

Xie, Y. (2022). *bookdown: Authoring Books and Technical Documents with R Markdown*. R package version 0.27.