

Practical 224

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Reproducibility

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Markdown

A essential tool used in creating these materials is [RMarkdown](#) That is an R library that allows you to create scripts that mix the Markdown mark-up language and R, to create dynamic documents. RMarkdown script can be compiled, at which point, the Markdown notation is interpreted to create the output files, while the R code is executed and the output incorporated in the document.

For instance the following markdown code

[This is a link to the University of Leicester](http://le.ac.uk) and **this is in bold**.
is rendered as

[This is a link to the University of Leicester](#) and **this is in bold**.

The core Markdown notation used in this session is presented below. A full RMarkdown *cheatsheet* is available [here](#).

```
# Header 1
## Header 2
### Header 3
#### Header 4
##### Header 5
```

```
**bold**
*italics*
```

```
[This is a link to the University of Leicester](http://le.ac.uk)
```

```
- Example list
  - Main folder
    - Analysis
    - Data
    - Utils
  - Other bullet point
- And so on
  - and so forth
```

```
1. These are
```

1. Numeric bullet points
2. Number two
2. Another number two
3. This is number three

R Markdown

R code can be embedded in RMarkdown documents as in the example below. That results in the code chunk be displayed within the document (as `echo=TRUE` is specified), followed by the output from the execution of the same code.

```
```{r, echo=TRUE}
a_number <- 0
a_number <- a_number + 1
a_number <- a_number + 1
a_number <- a_number + 1
a_number
```
```

```
a_number <- 0
a_number <- a_number + 1
a_number <- a_number + 1
a_number <- a_number + 1
a_number
```

```
## [1] 3
```

Exercise 224.1

Create a new R project named *Practical_224* as the directory name. Create an RMarkdown document in RStudio by selecting *File > New File > R Markdown ...* – this might prompt RStudio to update some packages. On the RMarkdown document creation menu, specify “Practical 05” as title and your name as the author, and select *PDF* as default output format.

The new document should contain the core document information, as in the example below, plus some additional content that simply explains how RMarkdown works.

```
---
title: "Practical 224"
author: "A. Student"
date: "7 October 2018"
output: pdf_document
---
```

Delete the contents below the document information and copy the following text below the document information.

```
# Pipe example
```

```
This is my first [RMarkdown](https://rmarkdown.rstudio.com/) document.
```

```
```{r, echo=TRUE}
library(tidyverse)
```
```

The code uses the pipe operator:

- takes 2 as input

- calculates the square root
- rounds the value
 - keeping only two digits

```

'''{r, echo=TRUE}
2 %>%
  sqrt() %>%
  round(digits = 2)
'''

```

The option `echo=TRUE` tells RStudio to include the code in the output document, along with the output of the computation. If `echo=FALSE` is specified, the code will be omitted. If the option `message=FALSE` and `warning=FALSE` are added, messages and warnings from R are not displayed in the output document.

Save the document by selecting *File > Save* from the main menu. Enter *Square_root* as file name and click *Save*. The file is saved using the *Rmd* (RMarkdown) extension.

Click on the *Knit* button on the bar above the editor panel (top-left area) in RStudio, on the left side. Check the resulting *pdf* document. Try adding some of your own code (e.g., using some of the examples above) and Markdown text, and compile the document again.

Exercise 224.2

Create an analysis document based on RMarkdown for each one of the two analyses seen in the practical sessions 3 and 4. For each of the two analyses, within their respective R projects, first, create an RMarkdown document. Then, add the code from the related R script. Finally add additional content such as title, subtitles, and most importantly, some text describing the data used, how the analysis has been done, and the result obtained. Make sure you add appropriate links to the data sources, as available in the practical session materials.

Git

Git is a free and opensource version control system. It is commonly used through a server, where a master copy of a project is kept, but it can also be used locally. Git allows storing versions of a project, thus providing file synchronisation, consistency, history browsing, and the creation of multiple branches. For a detailed introduction to Git, please refer to the [Pro Git book, written by Scott Chacon and Ben Straub](#).

As illustrated in the image below, when working with a git repository, the most common approach is to first check-out the latest version from the main repository before start working on any file. Once a series of edits have been made, the edits to stage are selected and then committed in a permanent snapshot. One or more commits can then be pushed to the main repository.

Git and RStudio

In RStudio Server, in the *Files* tab of the bottom-left panel, click on *Home* to make sure you are in your home folder – if you are working on your own computer, create a folder for *granolarr* wherever most convenient. Click on *New Folder* and enter *Repos* (short for repositories) in the prompt dialogue, to create a folder named *Repos*.

Create a GitHub account at github.com, if you don't have one, and create a new repository named *my-granolarr*, following the instructions available on the [GitHub help pages](#). Make sure you tick the box next to *Initialize this repository with a README*, which adds a `README.md` markdown file to your repository.

Once the repository has been created, GitHub will take you to the repository page. Copy the link to the repository `.git` file by clicking on the green *Clone or download* button and copying the `https` URL there available. Back to RStudio Server, select *File > New Project...* from the top menu and select *Version Control* and then *Git* from the *New Project* panel. Paste the copied URL in the *Repository URL* field,



Figure 1: by Scott Chacon and Ben Straub, licensed under CC BY-NC-SA 3.0

select the *Repos* folder created above as folder for *Create project as subdirectory of*, and click *Create Project*. RStudio might ask you for your GitHub username and password at this point.



Figure 2: Cloning my-granolarr as R project in RStudio

Before continuing, you need to record your identity with the Git system installed on the RStudio Server, for it to be able to communicate with the GitHub's server. Open the *Terminal* tab in RStudio (if not visible, select *Tools > Terminal > New Terminal* from the top menu). First, paste the command below substituting `you@example.com` with your university email (make sure to maintain the double quotes) and press the return button.

```
git config --global user.email "you@example.com"
```

Then, paste the command below substituting `Your Name` with your name (make sure to maintain the double quotes) and press the return button.

```
git config --global user.name "Your Name"
```

RStudio should have now switched to a new R project linked to your *my-granolarr* repository. In the RStudio *File* tab in the bottom-right panel, navigate to the file created for the exercises above, select the files and copy them in the folder of the new project (in the *File* tab, *More > Copy To...*).

Check the now available *Git* tab on the top-right panel, and you should see at least the newly copied files marked as untracked. Tick the checkboxes in the *Staged* column to stage the files, then click on the *Commit* button.

In the newly opened panel *Commit* window, the top-left section shows the files, and the bottom section shows the edits. Write **My first commit** in the *Commit message* section in the top-right, and click the *Commit* button. A pop-up should notify the completed commit. Close both the pop-up panel, and click the *Push* button on the top-right of the *Commit* window. Another pop-up panel should ask you for your GitHub username and password and then show the executed push operation. Close both the pop-up panel and the *Commit* window.

Congratulations, you have completed your first commit! Check the repository page on GitHub. If you reload the page, the top bar should show *2 commits* on the left and your files should now be visible in the file list below. If you click on *2 commits*, you can see the commit history, including both the initial commit that created the repository and the commit you just completed.

Exercise 224.3

Create a new GitHub repository named **GY7702_224_Exercise3**. Clone the repository to RStudio Server as a new R project from that repository. Create an RMarkdown document exploring the presence of the different living arrangements in Leicester among both the different categories of the 2011 Output Area Classification and deciles of Index of Multiple Deprivations, copying the required data into the project folder.

Your analysis should include:

- an introduction to the data and the aims of the project;
- a justification of the analysis methods;
- the code and related results;
- and a discussion of the results within the same document.

Cloning granolarr

You can follow the steps listed below to clone the granolarr repository.

1. Create a folder named **Repos** in your home directory. If you are working on RStudio Server, in the *Files* panel, click on the **Home** button (second bar, next to the house icon), then click on **New Folder**, enter the name **Repos** and click **Ok**.
2. In RStudio or RStudio Server select **File > New Project...**
3. Select **Version Control** and then **Git** (you might need to set up Git first if you are working on your own computer)
4. Copy <https://github.com/sdesabbata/granolarr.git> in the **Repository URL** field and select the **Repos** folder for the field **Create project as subdirectory of**, and click on **Create Project**.
5. Have a look around
6. Click on the project name **granolarr** on the top-left of the interface and select **Close Project** to close the project.

As granolarr is a public repository, you can clone it, edit it as you wish and push it to your own copy of the repository. However, contributing your edits to the original repository would require a few further steps. Check out the [GitHub help pages](#) if you are interested.