

# Setting up R projects

Applied Data Science using R

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# Goals for today

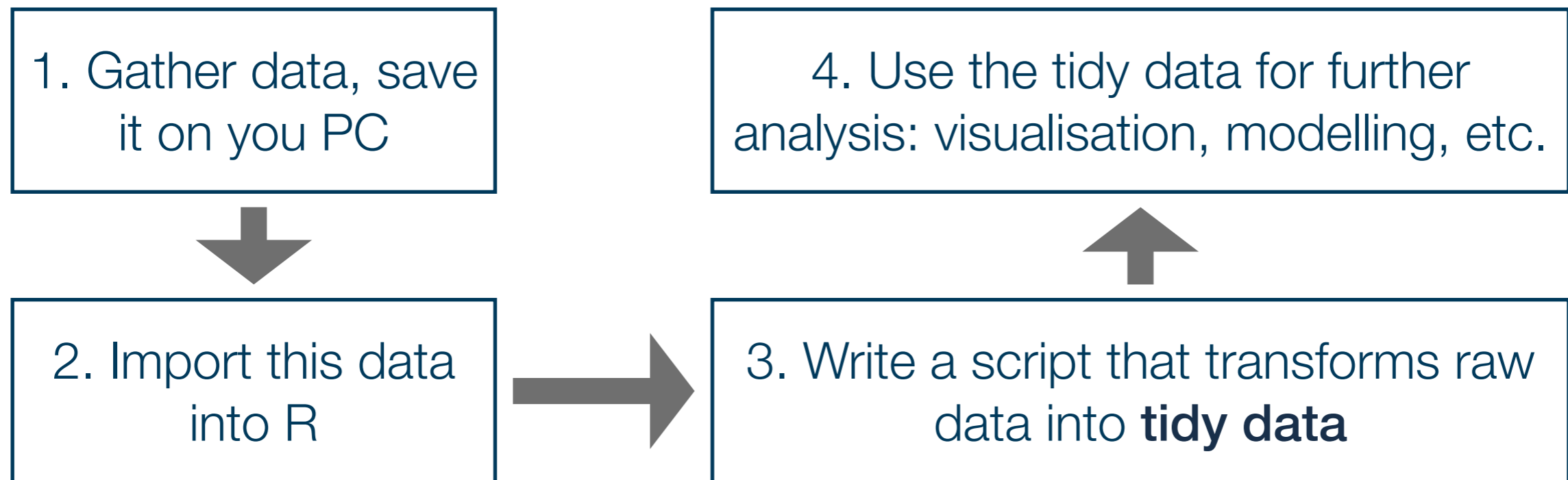
- I. Learn how to set up an R project
- II. Learn about the difference between absolute and relative paths
- III. Learn how to use the here package

# Our goal

- Learn about a **default directory structure** and a general way to **document everything** you do in your project
  - Facilitates the collaboration with future-you considerably
  - 👉 Nothing is worse than hating your past-you for not documenting correctly where data came from, or how it has been prepared 🤔
- Introduce general workflow to avoid most editable problems in the context of project management
- Central idea: all results must be **reproducible** from the raw data at any time
  - This implies that you **must not manipulate your raw data** at any cost
  - Raw data = what you download from the internet, gather through an experiment, or code yourself
  - Focus here: organization of your overall project

# How to keep your work transparent

- Raw data must not be changed, but is usually not in a state we can work with 😬



- Saving the scripts in steps 2 & 3 makes your work **fully reproducible**
- By looking into the script you will always know what you did to your raw data → you can also heal basically every mistake you made, not harm done!

# Outlook

Set up you project environment

This is done only once per project

Import data

Transform raw data into tidy data

This might be done several times

Save data

# Outlook

**Set up you project environment**

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Import data

Transform raw data into tidy data

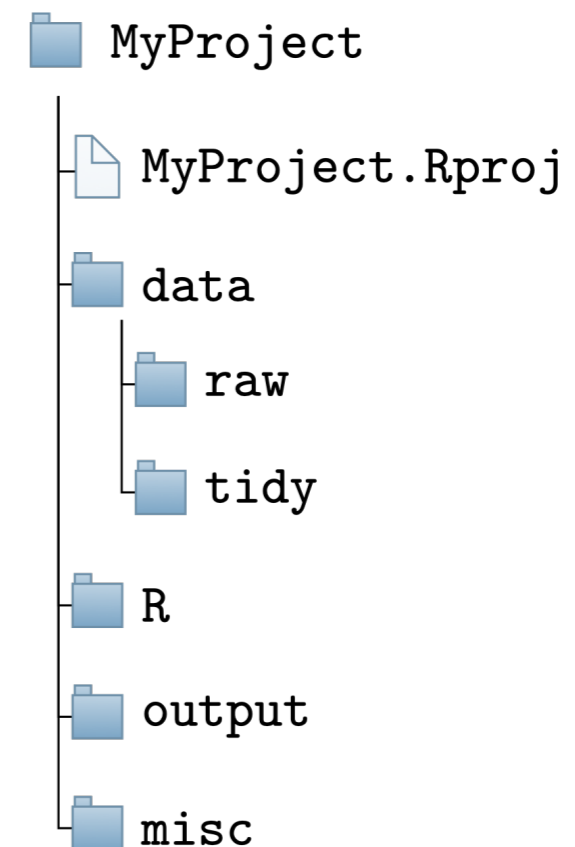
This might be done several times

Save data

# Set up your R project

# Setting up your working environment

- Before we talk about importing raw data we need to discuss where the raw data should actually be saved
- A prerequisite for a transparent, reproducible, and easy-to-work-with project is the right **directory structure**
- Thus, for every task in R you should set up your project like this:
- All the relevant steps to set this up, and the rationality for this structure are described in the respective tutorial



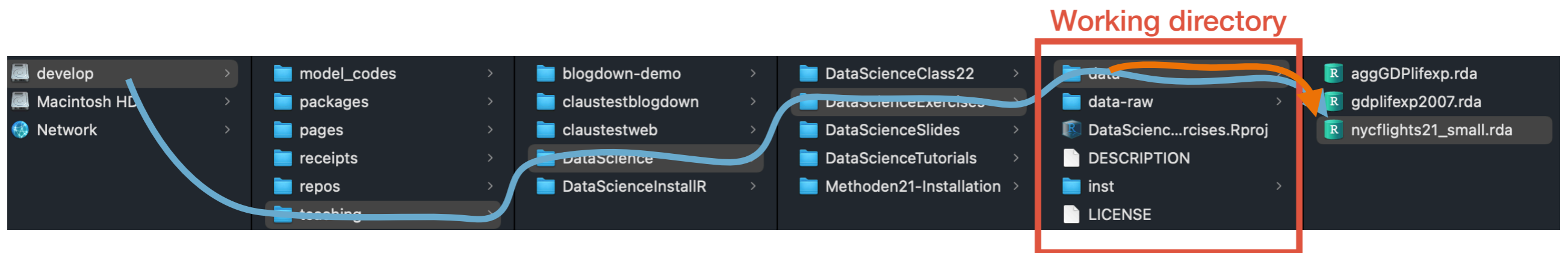


# Creating an R project

See the associated tutorial and video for the documentation of the relevant steps (slides focus on selected background concepts only)

# Paths and the here-package

- There are two ways in which you tell your computer where a certain file is located:
  - Via an absolute path: description starts at the root directory 🌲
  - Via a relative path: description starts at your current position in the file system



- Assuming we are 'located' in the folder `DataScienceExercises`: and want to point to the file `nycflights21_small.rda`:
  - `"/Volumes/develop/teaching/DataScience/DataScienceExercises/data/nycflights21_small.rda"`
  - `"data/nycflights21_small.rda"`

# Relative paths and `setwd()`

- The relative path seems nicer...
  - Its shorter 😊 and you can share code without forcing others to adjust the path
- Problem: how to set our location to the directory `DataScienceExercises`?
- We can do this using `setwd()`, providing the absolute path to `DataScienceExercises` as an argument:
  - `setwd("/Volumes/develop/teaching/DataScience/DataScienceExercises")`
  - Then we can use `"data/nycflights21_small.rda"`
- Many people put `setwd()` at the top of their scripts
  - **BUT YOU MUST NEVER EVER DO THIS!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!**



# Why setwd() is evil and not to be used

- You should never ever use `setwd()` in your scripts
- First, it does not help avoiding absolute paths because you have to provide an absolute path to `setwd()` 🤪
- Second, it makes people hate you:

Abby writes amazing\_script.R 🧑💻

```
setwd("/Volumes/Macintosh HD/Users/AbbysUserName/  
PathToFolderThatOnlyExistsHere/ProjectName")  
data_file <- data.table::fread("data/file.csv")
```


Sends file to Ellie 📧

Ellie opens file and executes it 😊

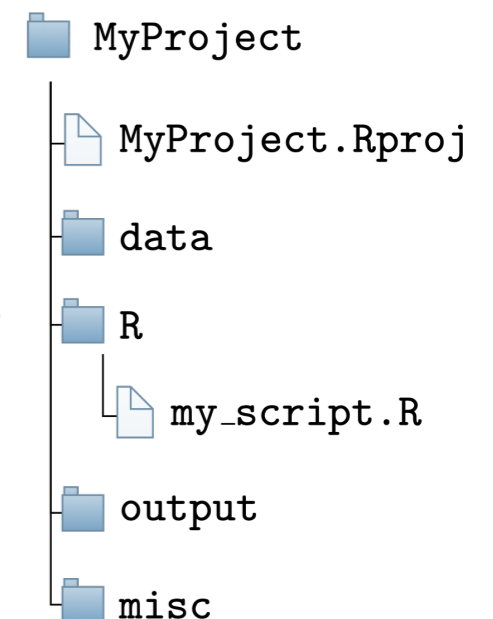


```
> setwd("/Volumes/Macintosh HD/Users/AbbysUserName/PathToFolderThatOnlyExistsHere/ProjectName/file.txt")  
Error in setwd("/Volumes/Macintosh HD/Users/AbbysUserName/PathToFolderThatOnlyExistsHere/ProjectName/file.txt") :  
cannot change working directory
```

# The better alternative to `setwd()` is here

- Thankfully, there is a very simple solution: the package [here](#)
- It allows you to set an anchor  in your project directory
- Then you can create paths relative to this anchor using the function `here::here()`
  - These commands will always work on every machine
- Always put `here::i_am()` into the first line of your scripts
  - As an argument, provide the location of the script relative to the project root
- From now on, only provide paths relative to this root using `here::here()`

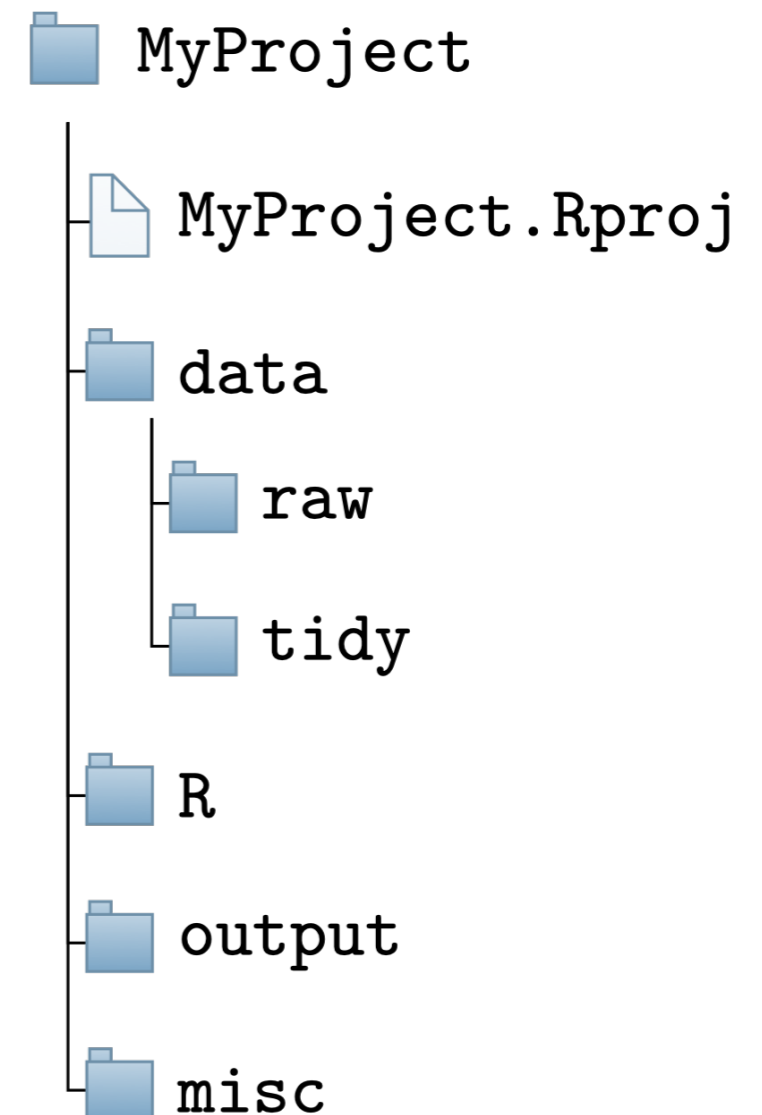
```
1 here::i_am("R/my_script.R")
2 library(here)
3 library(ggplot2)
4 # Script content
5
```



# Your turn: final exercise

- Create a new R-Project on your computer
- Create all the required folders
- Write an R script, put it into the right directory, and make it usable for the `here`-package
- Check out what the function `here::here()` returns and experiment with its use

```
1 here::i_am("R/my_script.R")
2 library(here)
3 library(ggplot2)
4 # Script content
5
```



# Summary and outlook

- We now know how to organise our **working directory**
- Important difference between **absolute and relative paths**
- Challenge of using code on different machines can be addressed using the **here package**
  - Better alternative than using `setwd()`
- Project management essential but often under-appreciated!
- Further topics:
  - Using a version control system (such as Git)
  - Using virtual programming environments (e.g. via the package **renv**)

