

# Docker for Reproducible Research

Anastasiia Enne

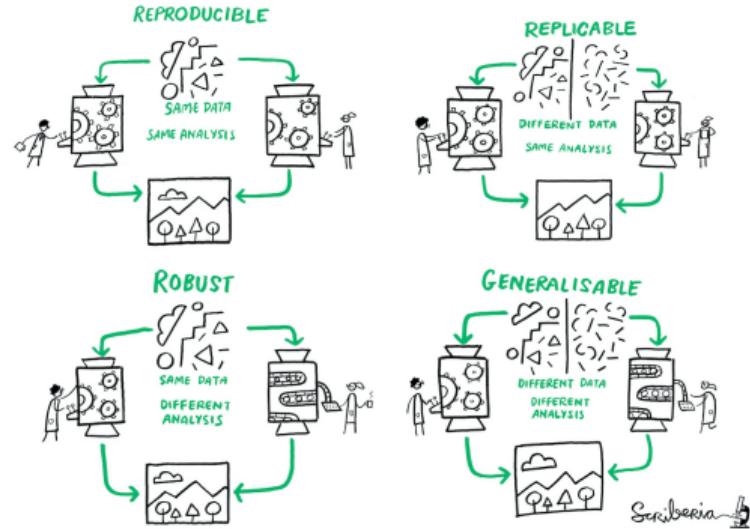
Group meeting, 14.01.2026

# Overview

- ① **Reproducibility crisis** — why just publishing your code is not enough
- ② **What is Docker** and how can it help us with reproducibility
- ③ **Dockerizing an existing project** — a step-by-step guide if you want to try Docker

# Reproducibility crisis

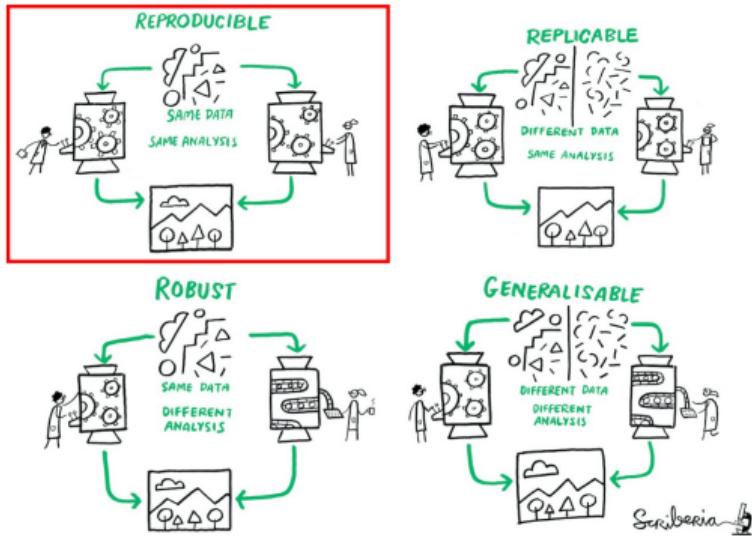
- Reminder: 4 definitions of reproducible research<sup>1</sup>



<sup>1</sup> The Turing Way Community. This illustration is created by Scriberia with The Turing Way community, used under a CC-BY 4.0 licence. DOI: [10.5281/zenodo.3332807](https://doi.org/10.5281/zenodo.3332807)

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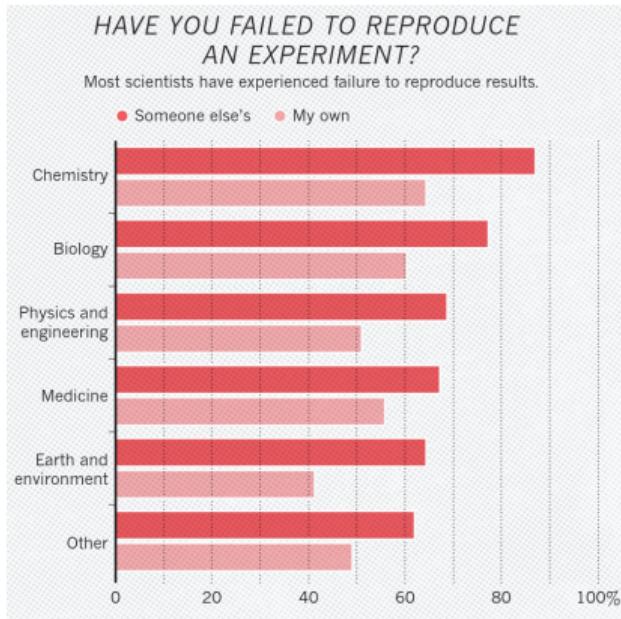
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- Our focus will be on REPRODUCIBILITY



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- Most scientists report that they have failed to reproduce an experiment<sup>2</sup>

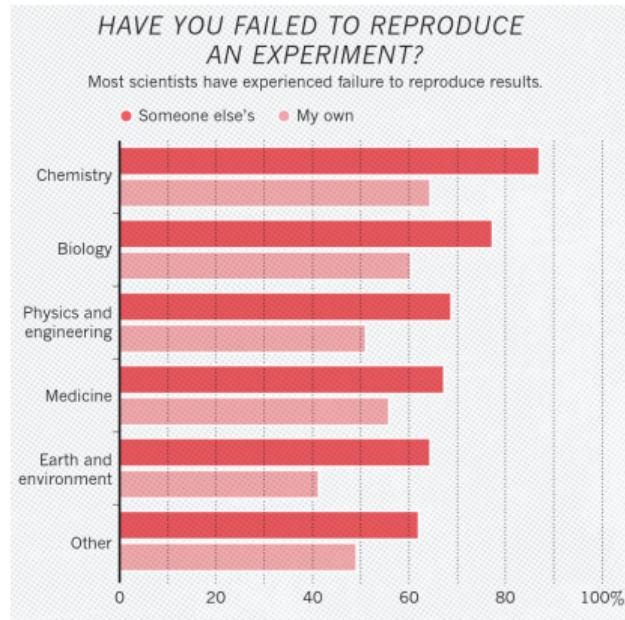


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# Reproducibility crisis

- Reminder: 4 definitions of reproducible research<sup>1</sup>
- Our focus will be on REPRODUCIBILITY
- Most scientists report that they have failed to reproduce an experiment<sup>2</sup>
- But what about modelling?



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# Reproducibility crisis in modelling studies?

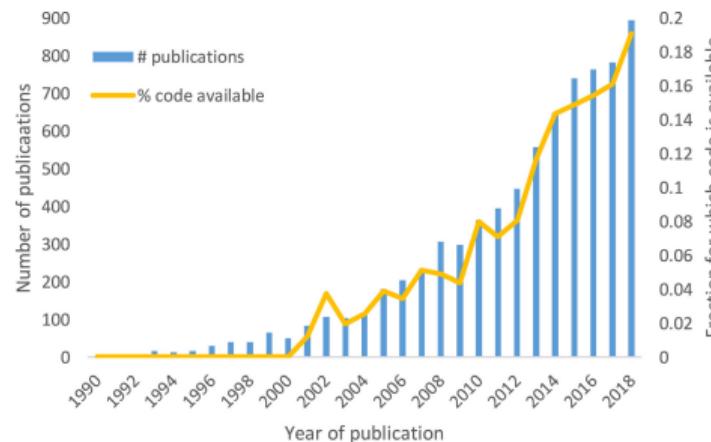
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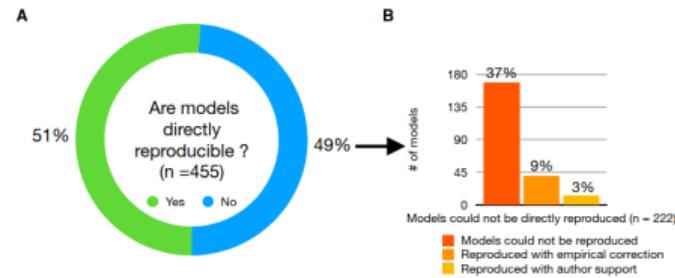
- How often do you manage to find and run the code for a paper without issues?
- Not many studies have investigated reproducibility in modelling papers...
  - Out of 7500 papers about individual-based and agent-based models only 11.2% provided their code<sup>1</sup>



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  - Out of 455 ODE models from [www.biomodels.org](http://www.biomodels.org) 49% are not directly reproducible<sup>2</sup>



<sup>1</sup> (Janssen, Pritchard, and Lee 2020)

<sup>2</sup> (Tiwari et al. 2021)

# Reproducibility crisis in modelling studies!

Main barriers to reproducibility<sup>1</sup>:

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- Computationally heavy and not scalable (hard to combine code from multiple studies)
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- A lot of proprietary formats
- Limited functionality

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- **Docker**

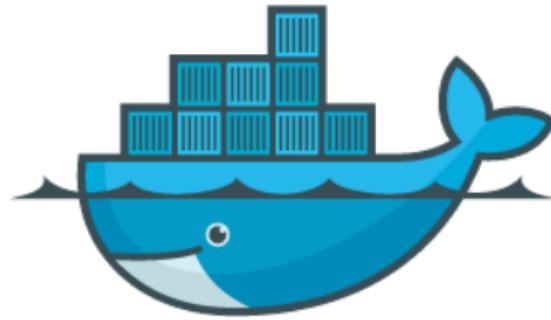
- Lightweight and easily scalable
- Clear list of dependencies
- Open source
- Linux functionality

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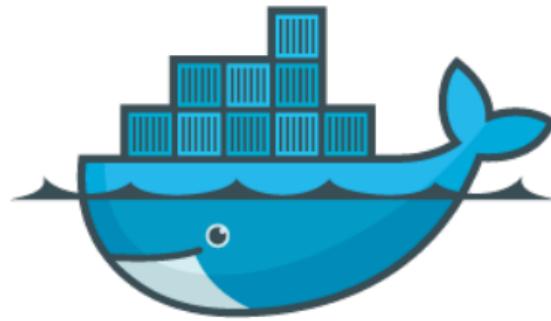
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- Docker becomes the new standard for reproducibility in science (e.g. Methods in Ecology and Evolution require the code to be Dockerized).



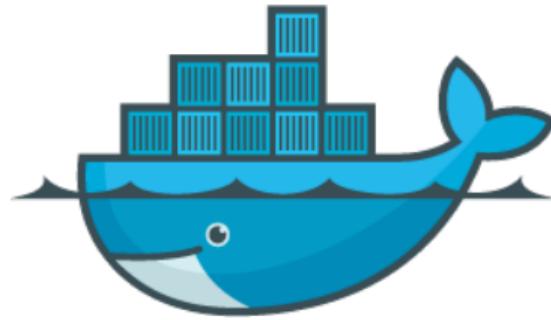
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  - **Understanding the basics of Docker and its components**
  - Being able to create and manage Docker containers



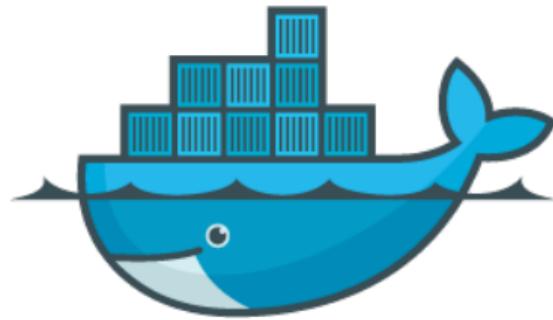
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  - Docker desktop app



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These slides are baked with Rmd and Dockerized! Go to [github.com/enne-anastasia/docker-for-reproducible-research](https://github.com/enne-anastasia/docker-for-reproducible-research) and see how I used Docker!

# What is Docker?

## Main idea

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- If Bob doesn't want to be bothered with using Docker he still can read the file and install all the dependencies on his own device manually.
- Alice is certain that she provided all the dependencies, because she tested this file with the Docker software.

# What is Docker?

docker build ⇒



Docker file

docker run ⇒



Docker image



Docker container

- Text file with "source code" of the image:
  - Instructions on how to build the image
  - Commands to run your project
- An executable snapshot of a container
- Includes all dependencies needed to run a container
- A running instance of the image
- Your "pocket Ubuntu"
- Self-contained

# What is Docker?

More about Docker containers



Docker container

- Containers stop existing after you exit them, meaning that all changes within the containers are lost.
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  - copy your files into the image (with a command in your Docker file), or
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- Consequently, you need to either
  - copy your files into the image (with a command in your Docker file), or
  - mount a folder to a container (during `docker run`)
- The rule of thumb is:
  - We MOUNT everything that takes plenty of space: e.g., data
  - We MOUNT everything that we want to be changed: e.g., figures
  - We COPY source code

# Dockerizing an existing project

What project are we going to Dockerize?

- We will Dockerize the R project that I used to prepare this presentation
- It has a very common structure for a code supplementary:
  - data — folder containing all the data used in this project (empty in this case)
  - docs — folder with Rmd file that generates these slides
  - figures — folder with all the figures I used
  - README file
  - run\_analysis.sh — one script to execute this project
  - src — source code (1 test R script in this case)

```
> tree -L 2
:
├── data
├── docker-for-reproducible-research.Rproj
└── docs
    ├── bibliography.bib
    ├── docker_slides.pdf
    ├── docker_slides.Rmd
    ├── docker_slides.tex
    └── preamble.tex
    └── figures
        └── downloaded
            └── screenshots
    └── README.html
    └── README.md
    └── run_analysis.sh
    └── src
        └── test.R
7 directories, 10 files
> █
```

# Dockerizing an existing project

## STEP 1: Installing Docker

- The official [docs.docker.com](https://docs.docker.com) provides full manuals on how to install Docker:
  - To work from the command line on Linux [install Docker Engine](#) (**this is what I did on Ubuntu**)

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- Verify that the installation is successful:

```
sudo systemctl status docker
sudo docker run hello-world
```

```
> sudo systemctl status docker
● docker.service - Docker Application Container Engine
   Loaded: loaded (/usr/lib/systemd/system/docker.service; enabled; preset: enabled)
   Active: active (running) since Tue 2026-01-13 11:24:15 CET; 28min ago
     TriggeredBy: ● docker.socket
       Docs: https://docs.docker.com
      Main PID: 18487 (dockerd)
        Tasks: 17
       Memory: 27.8M (peak: 33.0M)
         CPU: 1.155s
      CGroup: /system.slice/docker.service
              └─18487 /usr/bin/dockerd -H fd:// --containerd=/run/containerd/containerrd

Jan 13 11:24:15 ubio-23-25299 dockerd[18487]: time="2026-01-13T11:24:15.338176289+01:00"
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Jan 13 11:26:51 ubio-23-25299 dockerd[18487]: time="2026-01-13T11:26:51.667691881+01:00"
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[lines 1-22/22 (END)]
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> sudo docker run hello-world
```

Hello from Docker!

This message shows that your installation appears to be working correctly.

To generate this message, Docker took the following steps:

1. The Docker client contacted the Docker daemon.
2. The Docker daemon pulled the "hello-world" image from the Docker Hub. (amd64)
3. The Docker daemon created a new container from that image which runs the executable that produces the output you are currently reading.
4. The Docker daemon streamed that output to the Docker client, which sent it to your terminal.

To try something more ambitious, you can run an Ubuntu container with:

```
$ docker run -it ubuntu bash
```

Share images, automate workflows, and more with a free Docker ID:

<https://hub.docker.com/>

For more examples and ideas, visit:

<https://docs.docker.com/get-started/>

# Dockerizing an existing project

## STEP 2: Trying the basic Docker commands

- **Starting Docker:**

```
sudo systemctl start docker
```

*(Docker is running in the background  
waiting for you to use it)*

- **Stopping Docker:**

```
sudo systemctl stop docker
```

*(Docker is deactivated, but sometimes  
docker.socket remains active)*

- **Checking Docker status:**

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sudo systemctl status docker
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   Active: inactive (dead) since Tue 2026-01-13 12:05:12 CET; 14s ago
     Duration: 3min 53.508s
TriggeredBy: ● docker.socket
   Docs: https://docs.docker.com
   Process: 22747 ExecStart=/usr/bin/dockerd -H fd:// --containerd=/run/containerd/co...
 Main PID: 22747 (code=exited, status=0/SUCCESS)
    CPU: 656ms

Jan 13 12:01:18 ubio-23-25299 dockerd[22747]: time="2026-01-13T12:01:18.483714497+01:00"
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- To kill docker.socket use

```
sudo systemctl stop  
docker.socket
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STEP 2: Trying the basic Docker commands

- Listing all available Docker images:

```
sudo docker image ls
```

```
> sudo docker image ls
IMAGE          ID          DISK USAGE   CONTENT SIZE  EXTRA
hello-world:latest  d4aaab6242e0    25.9kB      9.52kB  U
>
```

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```

- Listing all containers that are currently running:

```
sudo docker container ls -a
```

```
> sudo docker container ls -a
CONTAINER ID IMAGE COMMAND CREATED STATUS PORTS NAME
S
73985d2e6242 hello-world "/hello" 4 hours ago Exited (0) 4 hours ago
efied_nightingale
stup
```

# Dockerizing an existing project

## STEP 2: Trying the basic Docker commands

- Listing all available Docker images:

```
sudo docker image ls
```

- Listing all containers that are currently running:

```
sudo docker container ls -a
```

- Deleting a container:

```
sudo docker container rm <ID>
```

```
sudo docker container rm <NAME>
```

```
> sudo docker container ls -a
CONTAINER ID IMAGE COMMAND CREATED STATUS PORTS NAME
S
73985d2e6242 hello-world "/hello" 4 hours ago Exited (0) 4 hours ago
efied_nightingale
```

# Dockerizing an existing project

## STEP 2: Trying the basic Docker commands

- Listing all available Docker images:

```
sudo docker image ls
```

- Listing all containers that are currently running:

```
sudo docker container ls -a
```

- Deleting a container:

```
sudo docker container rm <ID>
```

```
sudo docker container rm <NAME>
```

- Deleting an image:

```
sudo docker image rm <ID>
```

```
sudo docker image rm <IMAGE>
```

> sudo docker image ls					
IMAGE	ID	DISK USAGE	CONTENT SIZE	EXTRA	
hello-world:latest	d4aaab6242e0	25.9kB	9.52kB	U	
>					

# Dockerizing an existing project

## STEP 3: Writing the Docker file

- ① Create an empty Dockerfile in the root directory of your project

```
> tree -L 2
.
├── data
└── Dockerfile
    └── docker-for-reproducible-research.Rproj
        ├── docs
        │   ├── bibliography.bib
        │   ├── docker_slides.pdf
        │   ├── docker_slides.Rmd
        │   ├── docker_slides.tex
        │   └── preamble.tex
        ├── figures
        │   └── downloaded
        │       └── screenshots
        ├── README.html
        ├── README.md
        └── run_analysis.sh
    └── src
        └── test.R
7 directories, 11 files
>
```

# Dockerizing an existing project

## STEP 3: Writing the Docker file

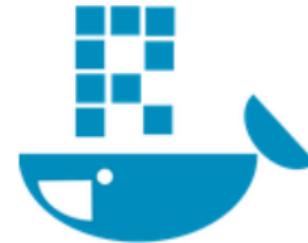
- ➊ Create an empty Dockerfile in the root directory of your project
- ➋ Pick the base image
  - Writing your Docker file from scratch is like setting a new work laptop from scratch
  - Thankfully, we do not have to do that because there are Docker images we can build upon!

```
> tree -L 2
.
├── data
└── Dockerfile
    └── docker-for-reproducible-research.Rproj
        ├── docs
        │   ├── bibliography.bib
        │   ├── docker_slides.pdf
        │   ├── docker_slides.Rmd
        │   ├── docker_slides.tex
        │   └── preamble.tex
        └── figures
            └── downloaded
                └── screenshots
        ├── README.html
        └── README.md
        └── run_analysis.sh
    └── src
        └── test.R
7 directories, 11 files
>
```

# Dockerizing an existing project

## STEP 3: Writing the Docker file

- ➊ Create an empty Dockerfile in the root directory of your project
- ➋ Pick the base image
  - Writing your Docker file from scratch is like setting a new work laptop from scratch
  - Thankfully, we do not have to do that because there are Docker images we can build upon!
  - For this project I will use rocker — Docker image with pre-installed R



## The Rocker Project

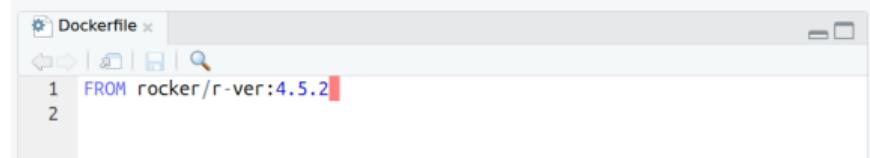
Docker Containers for the R Environment

# Dockerizing an existing project

## STEP 3: Writing the Docker file

- ③ Write the first line in your Dockerfile:

- We build our image starting from Docker image with pre-installed R
- I specify the version 4.5.2 of R since this is the version that I currently have on my laptop
- There can be only one FROM image!



A screenshot of a code editor window titled "Dockerfile x". The window contains the following code:

```
1 FROM rocker/r-ver:4.5.2
2
```

The line "FROM rocker/r-ver:4.5.2" is highlighted with a red rectangle.

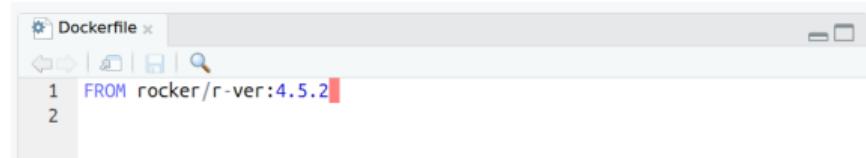
# Dockerizing an existing project

## STEP 3: Writing the Docker file

### ③ Write the first line in your Dockerfile:

- We build our image starting from Docker image with pre-installed R
- I specify the version 4.5.2 of R since this is the version that I currently have on my laptop
- There can be only one FROM image!

- Congratulations! This is the minimal Docker file we can use to build and run a Docker image (although, not very useful one)



```
Dockerfile x
1 FROM rocker/r-ver:4.5.2
2
```

# Dockerizing an existing project

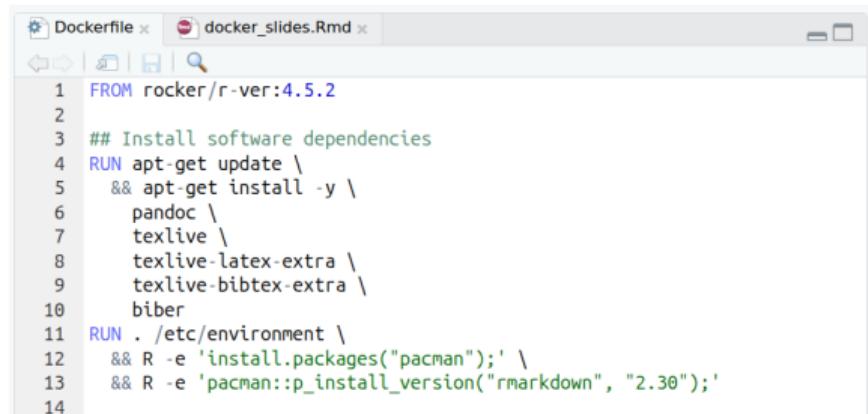
## STEP 3: Writing the Docker file

### ④ Specify software dependencies

- It is okay to do that in iterations.

When you think you got all the dependencies, go to the next step and see if you can run your project in the Docker container.

- Here I have 2 types of dependencies: OS-level and R-level.



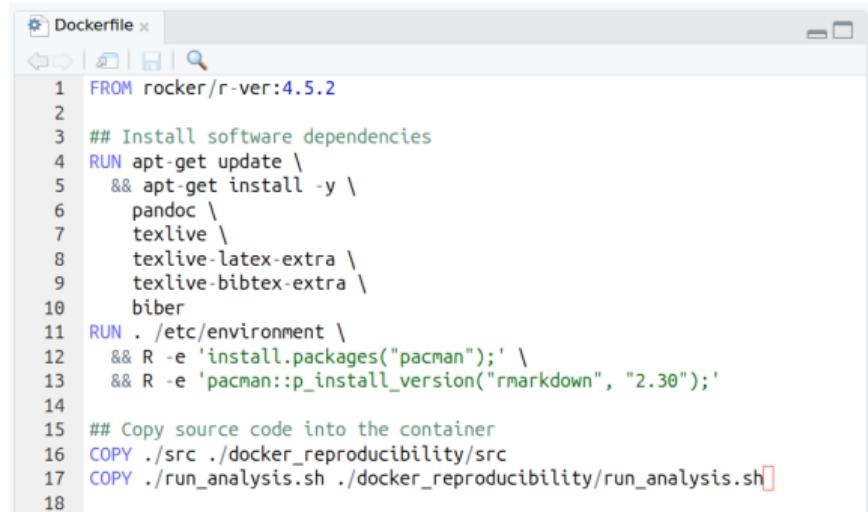
A screenshot of a code editor window titled "Dockerfile x docker\_slides.Rmd x". The editor shows a Dockerfile with the following content:

```
1 FROM rocker/r-ver:4.5.2
2
3 ## Install software dependencies
4 RUN apt-get update \
5     && apt-get install -y \
6         pandoc \
7         texlive \
8         texlive-latex-extra \
9         texlive-bibtex-extra \
10        biber
11 RUN . /etc/environment \
12     && R -e 'install.packages("pacman");' \
13     && R -e 'pacman::p_install_version("rmarkdown", "2.30");'
14
```

# Dockerizing an existing project

## STEP 3: Writing the Docker file

- ⑤ Copy the source code into the image
  - The inside of the container is an Ubuntu file system with installed dependencies, therefore I copy everything into a separate folder to keep things neat.



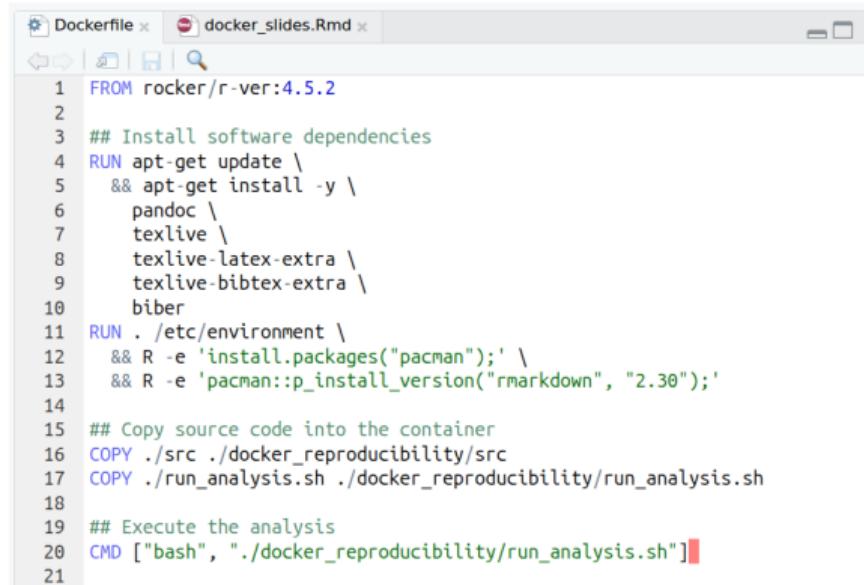
A screenshot of a code editor window titled "Dockerfile x". The editor shows a Dockerfile with the following content:

```
1 FROM rocker/r-ver:4.5.2
2
3 ## Install software dependencies
4 RUN apt-get update \
5     && apt-get install -y \
6         pandoc \
7         texlive \
8         texlive-latex-extra \
9         texlive-bibtex-extra \
10        biber
11 RUN . /etc/environment \
12     && R -e 'install.packages("pacman");' \
13     && R -e 'pacman::p_install_version("rmarkdown", "2.30");'
14
15 ## Copy source code into the container
16 COPY ./src ./docker_reproducibility/src
17 COPY ./run_analysis.sh ./docker_reproducibility/run_analysis.sh
18
```

# Dockerizing an existing project

## STEP 3: Writing the Docker file

- 5 Copy the source code into the image
  - The inside of the container is an Ubuntu file system with installed dependencies, therefore I copy everything into a separate folder to keep things neat.
- 6 Write the last line with the command to execute the analysis
  - There can be only one CMD line in your Docker file!



The screenshot shows a code editor window with a tab labeled "Dockerfile x" and another tab labeled "docker\_slides.Rmd x". The Dockerfile content is as follows:

```
1 FROM rocker/r-ver:4.5.2
2
3 ## Install software dependencies
4 RUN apt-get update \
5     && apt-get install -y \
6         pandoc \
7         texlive \
8         texlive-latex-extra \
9         texlive-bibtex-extra \
10        biber
11 RUN . /etc/environment \
12     && R -e 'install.packages("pacman");' \
13     && R -e 'pacman:::p_install_version("rmarkdown", "2.30");'
14
15 ## Copy source code into the container
16 COPY ./src ./docker_reproducibility/src
17 COPY ./run_analysis.sh ./docker_reproducibility/run_analysis.sh
18
19 ## Execute the analysis
20 CMD ["bash", "./docker_reproducibility/run_analysis.sh"]
```

# Dockerizing an existing project

## STEP 3: Writing the Docker file

- ➅ Copy the source code into the image
  - The inside of the container is an Ubuntu file system with installed dependencies, therefore I copy everything into a separate folder to keep things neat.
- ➆ Write the last line with the command to execute the analysis
  - There can be only one CMD line in your Docker file!



```
run_analysis.sh x
Run Run Script
1 cd docker_reproducibility
2 Rscript ./src/test.R
3 R -e 'library(rmarkdown); rmarkdown::render("./docs/docker_slides.Rmd")'
4
```

# Dockerizing an existing project

## STEP 4: Building the image

```
sudo docker build -t <TAG_NAME> ./
```

- It probably will take a while to install all dependencies
- When you change one layer of your image and build again, Docker will execute only starting from layers that were changed

```
> sudo docker build -t dore ./
[+] Building 14.8s (10/10) FINISHED
=> [internal] load build definition from Dockerfile
=> => transferring dockerfile: 589B
=> [internal] load metadata for docker.io/rocker/r-ver:4.5.2
=> [internal] load .dockerignore
=> => transferring context: 2B
=> [1/5] FROM docker.io/rocker/r-ver:4.5.2@sha256:76a8dec2998c79ceba36242e31db963ad15
=> => resolve docker.io/rocker/r-ver:4.5.2@sha256:76a8dec2998c79ceba36242e31db963ad15
=> [internal] load build context
=> => transferring context: 214B
=> CACHED [2/5] RUN apt-get update  && apt-get install -y      pandoc      texlive
=> CACHED [3/5] RUN . /etc/environment  && R -e 'install.packages("pacman");'  && R
=> CACHED [4/5] COPY ./src ./docker_reproducibility/src
=> [5/5] COPY ./run_analysis.sh ./docker_reproducibility/run_analysis.sh
=> exporting to image
=> => exporting layers
=> => exporting manifest sha256:160f570b294b814cc4a83c685771681dd473049bc4d44275afde1
=> => exporting config sha256:bff604ec4e5d344b0cdad97672ac880135cafe787002f92c2d3200a
=> => exporting attestation manifest sha256:19a27986c08162d902e9f501c16e83d96c740e5ca
=> => exporting manifest list sha256:704c22ad800cc602c4e940771696b74048c89ddcedbc76ce
=> => naming to docker.io/library/dore:latest
=> => unpacking to docker.io/library/dore:latest
12.9s
```

# Dockerizing an existing project

## STEP 4: Building the image

```
sudo docker build -t <TAG_NAME> ./
```

- It probably will take a while to install all dependencies
- When you change one layer of your image and build again, Docker will execute only starting from layers that were changed

IMAGE	ID	DISK USAGE	CONTENT SIZE	EXTRA
dore:latest	efa267ed23e7	3.53GB	984MB	
hello-world:latest	d4aaab6242e0	25.9kB	9.52kB	U

# Dockerizing an existing project

STEP 5: Testing your container

```
sudo docker run -v $(pwd)/figures:/docker_reproducibility/figures -v  
$(pwd)/docs:/docker_reproducibility/docs <TAG_NAME>
```

# Dockerizing an existing project

## STEP 5: Testing your container

```
> sudo docker run -v $(pwd)/figures:/docker_reproducibility/figures -v $(pwd)/docs:/docker_reproducibility/docs dore  
[1] 4
```

```
R version 4.5.2 (2025-10-31) -- "[Not] Part in a Rumble"  
Copyright (C) 2025 The R Foundation for Statistical Computing  
Platform: x86_64-pc-linux-gnu
```

```
R is free software and comes with ABSOLUTELY NO WARRANTY.  
You are welcome to redistribute it under certain conditions.  
Type 'license()' or 'licence()' for distribution details.
```

```
Natural language support but running in an English locale
```

```
R is a collaborative project with many contributors.  
Type 'contributors()' for more information and  
'citation()' on how to cite R or R packages in publications.
```

```
Type 'demo()' for some demos, 'help()' for on-line help, or  
'help.start()' for an HTML browser interface to help.  
Type 'q()' to quit R.
```

```
> library(rmarkdown); rmarkdown::render("./docs/docker_slides.Rmd")
```

```
processing file: docker_slides.Rmd  
1/3  
2/3 [setup]  
3/3  
output file: docker_slides.knit.md
```

```
/usr/bin/pandoc +RTS -K512m -RTS docker_slides.knit.md --to beamer --from markdown+autolink_bare_uris+tex_math_single_backslash --output docker_slides.tex --lua-filter /usr/local/lib/R/site-library/rmarkdown/rmarkdown/Lua/pagebreak.lua --lua-filter /usr/local/lib/R/site-library/rmarkdown/rmarkdown/Lua/latex-div.lua --variable theme=CambridgeUS --variable fonttheme=structurebold --highlight-style tango --pdf-engine pdflatex --biblatex --embed-resources --standalone --include-in-header /tmp/RtmpSm3bQ/rmarkdown-str1e2035fc3c.html
```

```
Output created: docker_slides.pdf  
>
```

-  Baker, Monya (2016). “1,500 scientists lift the lid on reproducibility”. In: *Nature* 533.7604, pp. 452–454. DOI: [10.1038/533452a](https://doi.org/10.1038/533452a).
-  Boettiger, Carl (2015). “An introduction to Docker for reproducible research”. In: *SIGOPS Oper. Syst. Rev.* 49.1, pp. 71–79. DOI: [10.1145/2723872.2723882](https://doi.org/10.1145/2723872.2723882).
-  Chan, Chung-hong and David Schoch (2023). “rang: Reconstructing reproducible R computational environments”. In: *PLOS ONE* 18.6, e0286761. DOI: [10.1371/journal.pone.0286761](https://doi.org/10.1371/journal.pone.0286761).
-  Janssen, Marco A., Calvin Pritchard, and Allen Lee (2020). “On code sharing and model documentation of published individual and agent-based models”. In: *Environmental Modelling & Software* 134, p. 104873. DOI: [10.1016/j.envsoft.2020.104873](https://doi.org/10.1016/j.envsoft.2020.104873).
-  Tiwari, Krishna et al. (2021). “Reproducibility in systems biology modelling”. In: *Molecular Systems Biology* 17.2, e9982. DOI: [10.15252/msb.20209982](https://doi.org/10.15252/msb.20209982).