Reproducibility in R

Sharing interactive environments with Binder

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Hello!

Here I would like to share with you this code I am working on. Let me know if you have any problems.



Issue 1

Absolute path

Issue 2

Package version of tidyr 0.8.3 does not include pivot_longer() and pivot_wider() functions added in tidyr version 1.0.0

Use sessionInfo() to check this

Would I have reproducible work only sharing the code and the data?

Reproducible environments

There are several tools to capture computational environments

- Package management systems (packrat renv)
- Binder
- Virtual machines
- Containers

Package management systems

renv package



Reproducible Environments -RStudio

renv package

- 1. Parenv::init() works by creating a new library. A library stores installed packages.
- 2. The file contains all the information you need to communicate your project's dependencies at the moment you call snapshot.
- 3. Frenv::restore() recreates the environment!

^{*} Read more about reny here

Binder

Jupyter Notebooks

Shiny

RStudio

Tutorial 1 - Ines Montani

Tutorial 2 - LearnR

Binder

Binder is an open source web service that lets users create sharable, interactive, reproducible environments in the cloud.



Advantages

- Easy to use
- You can access the infomation with one click
- It is free

Limitations

- Computational power
- Security/privacy (using mybinder.org BinderHub)
- Also no FTP for connecting to some data.



Binder's goal is to lower the barrier to interactivity, and to allow users to utilize code that is hosted in repository providers such as GitHub

Binder 2.0 - Reproducible, interactive, sharable environments for science at scale

mybinder.org

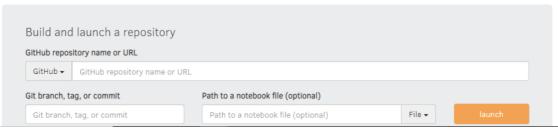
A free, public BinderHub. Because it is public, you should not use it if your project requires personal or confidential information (such as passwords).



Turn a Git repo into a collection of interactive notebooks

Have a repository full of Jupyter notebooks? With Binder, open those notebooks in an executable environment, making your code immediately reproducible by anyone, anywhere.

New to Binder? Get started with a Zero-to-Binder tutorial in Julia, Python or R.



"Binderizing" your project (9)

- 1- Specify the computational environment
- + intall.R
- + runtime.txt
- 2- Upload the project files to a publicly available repository hosting service, such as GitHub / GitLab
- 3- "Binderize" the project (mybinder.org)
- 4- Use the correct URL @



Demo

+ install.R

This file should have listed all of the packages to be installed

```
install.packages("ggplot2")
install.packages("shiny")
```

What is MRAN?

Since September 17th, 2014, the checkpoint server has been taking a daily snapshot at precisely midnight UTC of the entire CRAN repository and storing it on Microsoft R Archived Network (MRAN)

⚠ Non-CRAN packages, such as those available on GitHub, are not part of the snapshot process.

EXTRA: checkpoint package

© checkpoint package allows you to install packages as they existed on CRAN on a specific snapshot date as if you had a CRAN time machine.

```
library(checkpoint)
checkpoint("YYYY-MM-DD")
```



Specify the R and package versions used

For this you must choose a date where the versions of your packages are captured in MRAN.

r-version-<YYYY>-<MM>-<DD>

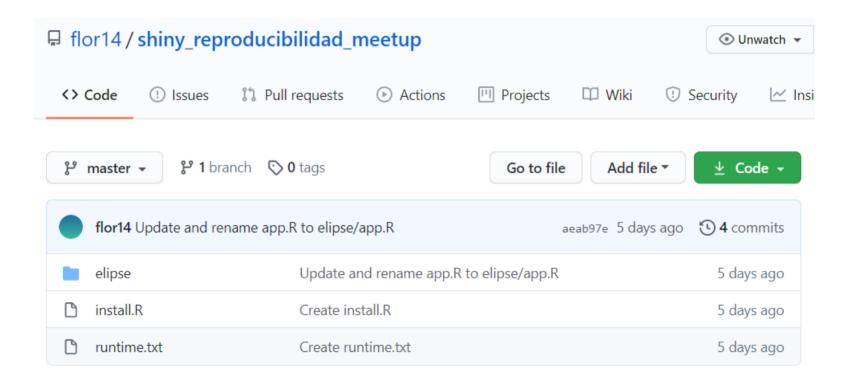
READ HERE - Important about R versions

+ I runtime.txt

r-3.6-2020-08-20 #r-version-<YYYY>-<MM>-<DD>



2. Upload your code to the repository



3. "Binderize" your project @

- a. Go to https://mybinder.org
- b. Paste the repository URL https://github.com/<your-username>/<your-repository>
- c. Finally, click the Launch button.

Patience! This could take a while \(\mathbb{T} \)

RStudio IDE URL

+?urlpath=rstudio

You should call the binderized project using this template link

https://mybinder.org/v2/gh/<user>/<repository>/<branch>?urlpath?rstudio

Example **@**:

http://mybinder.org/v2/gh/flor14/shiny_reproducibilidad_meetup/master?urlpath=rstudio

Ejemplos en el repositorio de Binder

Shiny app URL \$\lambda

+?urlpath=shiny/<folder>/

You should call the binderized project using this template link

https://mybinder.org/v2/gh/<user>/<repository>/<branch>? urlpath=shiny/<folder>/

Example **@**:

https://mybinder.org/v2/gh/flor14/shiny_reproducibilidad_meetup/master?urlpath=shiny/elipse/

Ejemplos en el repositorio de Binder

- Ines Montani framwework uses Binder
- Interactive Tutorial with learnr and Binder Sang-Yun Oh blog post

Others

- ✓ Holepunch Package
- ✓ Faster installation

r-conda

- ✓ More info about Binder
- Changes to Docker terms of service on November 1. Lack of activity for 6 months could leave links inactive.
- **P**discourse.jupyter.org

Practice 6

Exercise

Could you modify the code from the first exercise to make it work?



Links @

- Binder 2.0 Reproducible, interactive, sharable environments for science at scale
- Reproducibility in Production Webinar
- The Turing Way Book
- Reproducible Environments RStudio
- renv: Project Environments with R RStudio blog
- Putting the R into Reproducible Research Anna Krystalli
- Demo renv package



iThank you!

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