

The Alan Turing Institute

The Turing Way Workshop: Reproducible, Open and FAIR Research

Session 2

Emma Karoune & Malvika Sharan

Pronouns: she/her/hers



Assignment: Create a project repository

Add top-level files: **README** and **LICENSE**

- README to communicate about your work
- Also try to add a License - to allow others to use, modify, build upon your work
 - We will cover licenses in the next session in more detail.

Use The Turing Way chapter for README to guide your assignment

<https://the-turing-way.netlify.app/project-design/project-repo/project-repo-readme.html>

Create a GitHub Repository with README file

Create a new repository

A repository contains all the files for your project, including the revision history.

Initialize this repository with:

Skip this step if you're importing an existing repository.

Add a README file

This is where you can write a long description for your project. [Learn more.](#)

Add .gitignore

Choose which files not to track from a list of templates. [Learn more.](#)

Note

Three lessons about README

- Know your users and what they need
- Get users doing powerful things quickly
- Watch out for jargon!

Source: Hao Ye. (2021, March). Collaborations Workshop 2021 Mini-Workshop: README tips to make your project more approachable (Version v1.0.0). Zenodo.

<http://doi.org/10.5281/zenodo.4647391>

End of Part 1: Further Reading and Examples

- *The Turing Way* README
- Open Life Science README
- Purple Booth's README Template
- Thoughtbot's Blog on How to Write a Good README
- Matias Singer's curated List of Awesome READMEs

Assignment: Create a project repository

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Use The Turing Way chapter for README to guide your assignment

<https://the-turing-way.netlify.app/project-design/project-repo/project-repo-readme.html>

Owner *

 malvikasharan ▾

Repository name *

2022-03-project-example ✓

Great repository names are short and memorable. Need inspiration? How about [automatic-funicular?](#)

Description (optional)



Public

Anyone on the internet can see this repository. You choose who can commit.



Private

You choose who can see and commit to this repository.

Initialize this repository with:

Skip this step if you're importing an existing repository.



Add a README file

This is where you can write a long description for your project. [Learn more.](#)



Add .gitignore

Choose which files not to track from a list of templates. [Learn more.](#)



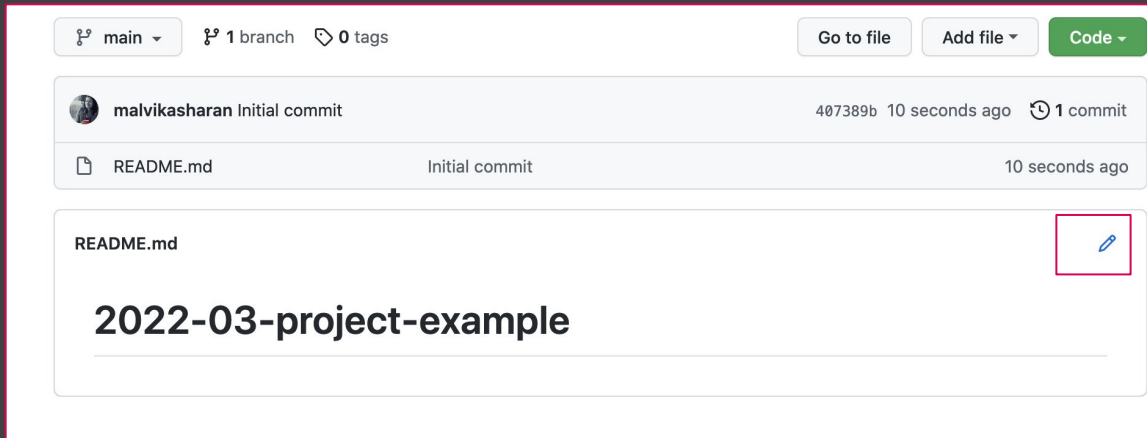
Choose a license

A license tells others what they can and can't do with your code. [Learn more.](#)

This will set  `main` as the default branch. Change the default name in your [settings](#).

Create repository

<https://github.com/malvikasharan/2022-03-project-example>



1. Copy Template:



https://raw.githubusercontent.com/ha0ye/CW21-README-tips/main/template_README.md

2. Add your project information

Bonus: Badges and Additional Information

README.md

The Turing Way



Contributing

🚧 This repository is always a work in progress and **everyone** is encouraged to help us build something that is useful to the many. 🚧

Everyone is asked to follow our [code of conduct](#) and to checkout our [contributing guidelines](#) for more information on how to get started.

If you are not familiar or confident contributing on GitHub, you can also contribute a case study and your tips and tricks via our [Google submission form](#).


Citing *The Turing Way*

You can reference *The Turing Way* through the project's Zenodo archive using DOI: [10.5281/zenodo.3233853](#). DOIs allow us to archive the repository and they are really valuable to ensure that the work is tracked in academic publications.

Custom Badges

The Shields service (at [shields.io](#)) provides a way to create custom badges for your projects. These are badges are very common and are frequently used to show status information about the project, or demonstrate tools that were used for the development of your project.

Example badge:



Static



Using dash "-" separator

```
https://img.shields.io/badge/<LABEL>-<MESSAGE>-<COLOR>
```

Dynamic



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Session 2

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Project communication, collaboration and sharing

- **License** to allow reuse of your research outputs in a way YOU want
- Make your work **citable**: Digital Object Identifiers, Zenodo
- Setting online repository for **collaboration**
 - CoC, Contributing guidelines, continuous integration
- **Reproducible environment** and sharing to aid reproducibility: **Binder**
- Revisiting how these contribute to implementing FAIR practices
- What more you can do.

Open Licenses

Disclaimer: We are not lawyers

Licensing Chapter in The Turing Way

Slides are CC-BY 4.0. Open Life Science. Contributors: OLS Team, Hao Ye, Christine Rogers, Josh Simmons, Mozilla Open Leadership team, Licensing chapter in The Turing Way:
<https://the-turing-way.netlify.app/reproducible-research/licensing.html>

Motivation:

Allow others to use, remix and share your work.

Method:

Add an open license for use, remixing and sharing.

Addressing Common Concerns

- Sharing something on GitHub does **NOT automatically make it reusable.**
- Sharing with a license does **NOT give away your rights** to publish, sell, etc.
- Work shared with an open license **NOT** to be used **without attribution.**
- *Not attributing to authors violates academic ethics.*

Truly Open Licenses: common elements



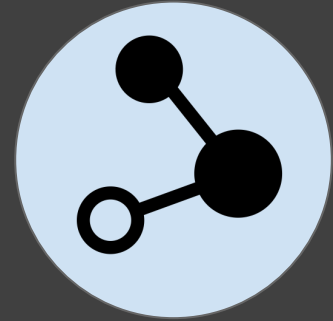
use

Anyone can use the work for any purpose



modify

Anyone can modify the work



share

Anyone can redistribute both the original and modified work

“Open source software is software that can be freely used, modified, and shared (in both modified and unmodified form) by anyone.”

- [GitHub Glossary, Open Source](#)

@openlifesci, CC-BY 4.0, The Turing Way, DOI: 10.5281/zenodo.6346555

Attribution

Most open licenses require crediting the authors of the work.

Non-copyleft: *Permissive*, non-reciprocal - CC BY, MIT, BSD

- **do not require** derivative works to be shared with the same license

Copyleft: *Viral*, reciprocal - CC BY-SA, GPLv3, MPL-2.0

- **require** derivative works to be shared with the same license

Exception: CC0 (public domain, no copyright holder)

How to Apply License



Place the **full text of the license** in a file named **LICENSE** in the root directory.

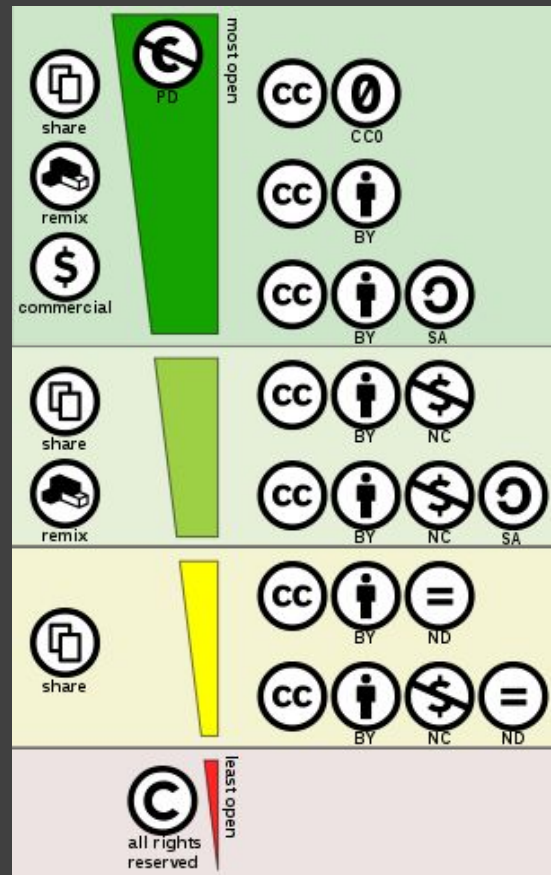
Apply licenses to all components, **content: CC-BY, software: MIT, data: CC0**

Describe in README which license applies to which parts of your work.

Assignment: Add a License

Use a **different license** for content, code and data. For example

- Writing/docs/images/metadata: **CC-BY**
- Code: **MIT License**
- Data will also have different license based on what you can/can't share!



GitHub can add a license for you

Create a new repository

A repository contains all the files for your project, including the revision history.

Initialize this repository with:

Skip this step if you're importing an existing repository.

Add a README file

This is where you can write a long description for your project. [Learn more.](#)

Add .gitignore

Choose which files not to track from a list of templates. [Learn more.](#)

Choose a license

A license tells others what they can and can't do with your code. [Learn more.](#)

License: None ▾

This repository will be licensed with the license you select. You can change the default name in your [settings](#).

License

Filter licenses...

✓ None

Apache License 2.0

GNU General Public License v3.0

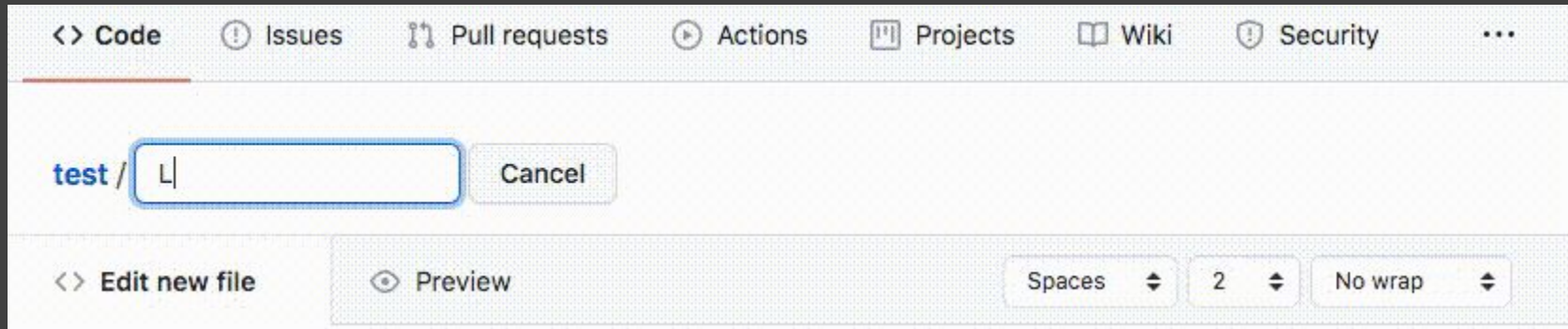
MIT License

BSD 2-Clause "Simplified" License

BSD 3-Clause "New" or "Revised"

GitHub can add a license for you, Part 2

when adding a new file named LICENSE...



Further Reading

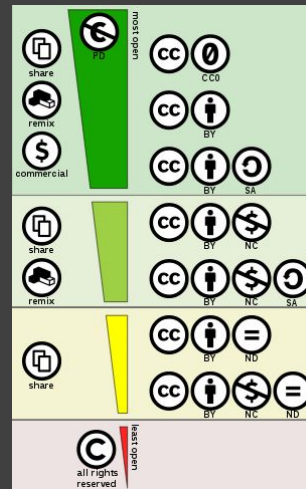
Licensing Chapter in *The Turing Way*

- [The Open Source Definition \(10 Criteria\)](#) | [opensource.org](#)
- [Legal Matters](#) | [producingoss.com](#)
- [Open Source Guide](#) | [opensource.guide](#)
- Software: [Choose an Open Source License](#) | [choosealicense.com](#)
- Content: [Choose a License](#) | [creativecommons.org](#)

Breakout

Make sure that you have done the following:

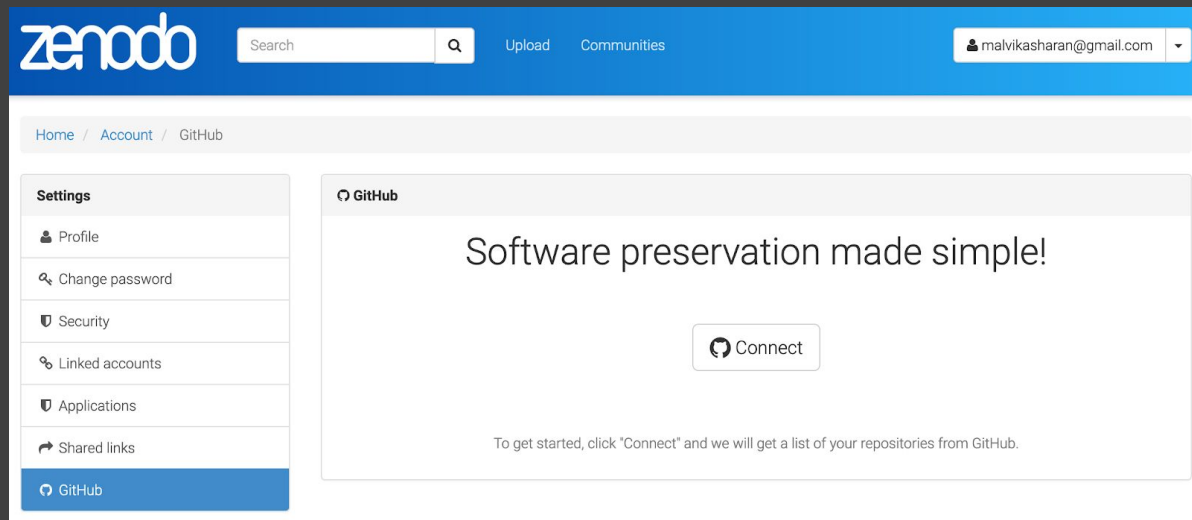
- Set a project repository
- Add a minimal README
- Add a License



You can use part of your project to use as example for this workshop!

You can fork: <https://github.com/malvikasharan/2022-03-project-example>

Making your work citable



The screenshot shows the Zenodo website interface. At the top, there is a blue header with the Zenodo logo, a search bar, and navigation links for 'Upload' and 'Communities'. A user profile dropdown menu is visible on the right, showing the email 'malvikasharan@gmail.com'. Below the header, the breadcrumb navigation reads 'Home / Account / GitHub'. On the left, a 'Settings' sidebar is open, listing options like Profile, Change password, Security, Linked accounts, Applications, Shared links, and GitHub (which is highlighted). The main content area is titled 'GitHub' and features the text 'Software preservation made simple!' and a 'Connect' button. Below the button, a message states: 'To get started, click "Connect" and we will get a list of your repositories from GitHub.'

(DOI) is a **persistent identifier** used to identify research objects uniquely.

- Zenodo, Figshare (research objects)
- Data Dryad (data)
- Preprint (manuscript)
- Docker (env)
- ORCID (researchers)

Zenodo

malvikasharan@gmail.com

- Profile
- Change password
- Security
- Linked accounts
- Applications
- Shared links
- GitHub
- Log out

malvikasharan/2022-03-project-example ON

DOI 10.5281/zenodo.6345410

GitHub

malvikasharan / 2022-03-project-example Public

<> Code Issues Pull requests Actions Projects

main 1 branch 1 tag

malvikasharan Create LICENSE.md

Releases Tags

There aren't any releases here

You can create a release to package software, along with release notes and links to binary files, for other people to use. Learn more about releases in our docs.

Create a new release

Releases Tags

Choose a tag Target: gh-pages

Choose a tag

Find or create a new tag

Create new tag: on publish

This is a pre-release We'll point out that this is a pre-release

Publish release

Zenodo

https://zenodo.org/account/settings/github/

zenodo Search Upload Communities

Drafts 3 Published 49 All versions

March 11, 2022 (v0.001) Software Open Access

malvikasharan/2022-03-project-example: test release

Created Mar 11, 2022 12:05:46 AM, modified Mar 11, 2022 12:05:51 AM

Publication date: March 11, 2022

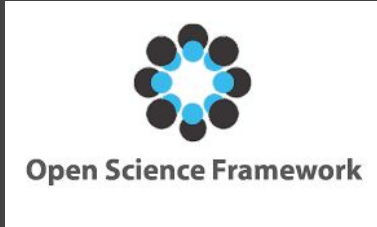
DOI: DOI 10.5281/zenodo.6345410

Related identifiers: Supplement to https://github.com/malvikasharan/2022-03-project-example/tree/v0.001

License (for files): Other (Open)

Citable DOI

Upload to free openly accessible online repositories



- Not a TDR
- cannot ensure long-term access





- Trustworthy digital repository (TDR)
- Allows to generate shareable DOI - Digital Object Identifier

Contribution Guideline

Contributing to *The Turing Way*

 Welcome to *The Turing Way* repository! 

 We're excited that you're here and want to contribute. 

We want to ensure that every user and contributor feels welcome, included and supported to participate in *The Turing Way* community. We hope that the information provided in this document will make it as easy as possible for you to get involved.

We welcome all contributions to this project via GitHub issues and pull requests. Please follow these guidelines to make sure your contributions can be easily integrated into the projects. As you start contributing to *The Turing Way*, don't forget that your ideas are more important than perfect pull requests. ❤️

If you have any questions that aren't discussed below, please let us know through one of the many ways to [get in touch](#).

Table of contents

Been here before? Already know what you're looking for in this guide? Jump to the following sections:

- [Joining the community](#)
- [Inclusivity](#)
- [Get in touch](#)
- [Contributing through GitHub](#)
- [Writing in Markdown](#)
- [Where to start: issues](#)

CONTRIBUTING.md

- how to file a bug report
- how to suggest a feature
- how to contribute changes
- roadmap & project vision
- how contributors should ask for help and guidance

Code of Conduct

Code of Conduct

We value the participation of every member of our community and want to ensure that every contributor has an enjoyable and fulfilling experience. Accordingly, everyone who participates in the Turing Way project is expected to show respect and courtesy to other community members at all times.

Kirstie Whitaker, as PI of this project, and all project members, are dedicated to a *harassment-free experience for everyone*, regardless of gender, gender identity and expression, sexual orientation, disability, physical appearance, body size, race, age or religion. **We do not tolerate harassment by and/or of members of our community in any form.**

We are particularly motivated to support new and/or anxious collaborators, people who are looking to learn and develop their skills, and anyone who has experienced discrimination in the past.

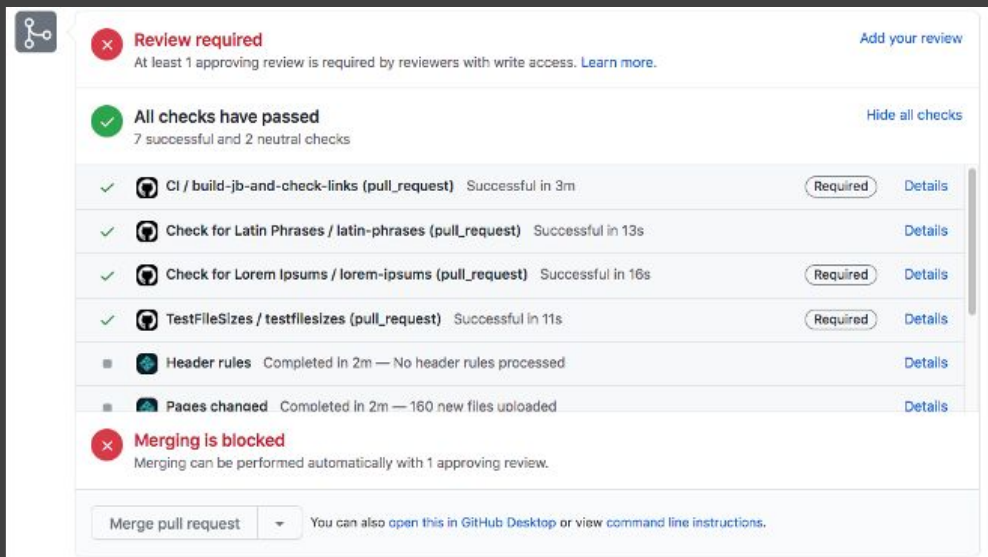
To make clear what is expected, we ask all members of the community to conform to the following Code of Conduct.

- [1 Introduction](#)
- [2 Code of Conduct](#)
 - [2.1 Expected Behaviour](#)
 - [2.2 Unacceptable Behaviour](#)
 - [2.3 Consequences of Unacceptable Behaviour](#)
 - [2.4 Feedback](#)
- [3 Incident Reporting Guidelines](#)
 - [3.1 Contact points](#)

CODE_OF_CONDUCT.md

- What is expected of the contributors
- What culture do we want to promote
- What if something should be reported

Continuous Integration



The screenshot displays a GitHub pull request interface. At the top, a red 'Review required' status is shown with a message: 'At least 1 approving review is required by reviewers with write access. Learn more.' and a link to 'Add your review'. Below this, a green 'All checks have passed' status is shown with a message: '7 successful and 2 neutral checks' and a link to 'Hide all checks'. A list of checks follows, including 'CI / build-jb-and-check-links (pull_request)' (Successful in 3m, Required), 'Check for Latin Phrases / latin-phrases (pull_request)' (Successful in 13s), 'Check for Lorem Ipsums / lorem-ipsums (pull_request)' (Successful in 16s, Required), and 'TestFileSizes / testfilesizes (pull_request)' (Successful in 11s, Required). Two other checks are shown as completed: 'Header rules' (Completed in 2m — No header rules processed) and 'Pages changed' (Completed in 2m — 160 new files uploaded). At the bottom, a red 'Merging is blocked' status is shown with a message: 'Merging can be performed automatically with 1 approving review.' and a 'Merge pull request' button.

Practice of integrating changes to a project made by individuals into a main, shared version -- frequently

All these aspects
enhance collaboration in
your project!

How to make your code
easy to test?



Reproducible computational environment

Reproducible research

same analysis steps on
the same dataset
produces same answer

		Data	
		Same	Different
Analysis	Same	(Reproducible)	Replicable
	Different	Robust	Generalisable

*“An article about computational science in a scientific publication is not the scholarship itself, it is merely advertising of the scholarship. The actual scholarship is the **complete software development environment** and the complete set of instructions which generated the figures.”*

— *Buckheit and Donoho (paraphrasing John Claerbout)*

WaveLab and Reproducible Research, 1995

(slide courtesy of Chris Holdgraf and the Jupyter Team)

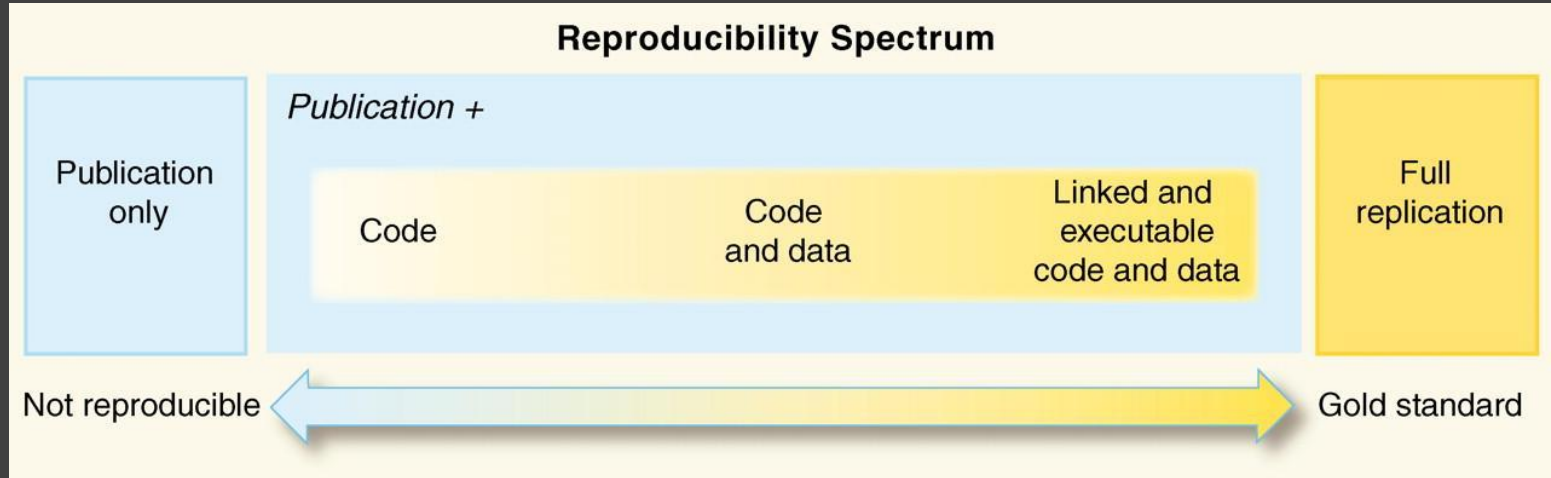
@turingway, CC-BY 4.0, The Turing Way, DOI: 10.5281/zenodo.6346555

Take home message

sharing your code and
data isn't enough



You need the computational environment too



You need the computational environment too

Publication
only

Not reproducible



Full
replication

Gold standard

Peng, 2011, doi: [10.1126/science.1213847](https://doi.org/10.1126/science.1213847)


@turingway, CC-BY 4.0, The Turing Way, DOI: [10.5281/zenodo.4609987](https://doi.org/10.5281/zenodo.4609987)

What is a computational environment?

- hardware (e.g. CPU)
- software
 - operating system
 - programming languages
 - packages

What is a computational environment?

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their versions
and their
configuration

What is a computational environment?

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- software
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 - packages

their versions
and their
configuration

and their
interaction

What is *Binder*?

What is Binder?



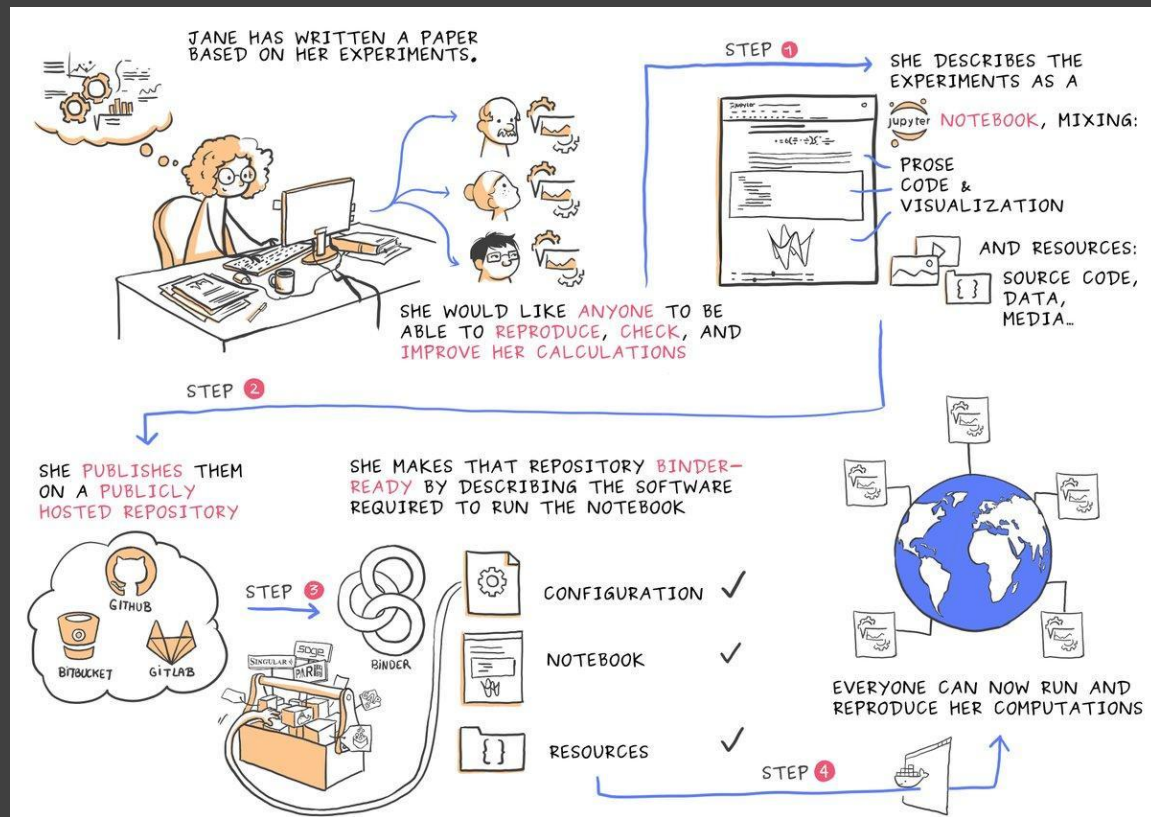
choldgraf Distinguished Contributor

3  Nov '18

The Binder Project helps you create one-click, sharable, live code environments from public code repositories that runs entirely in the cloud.

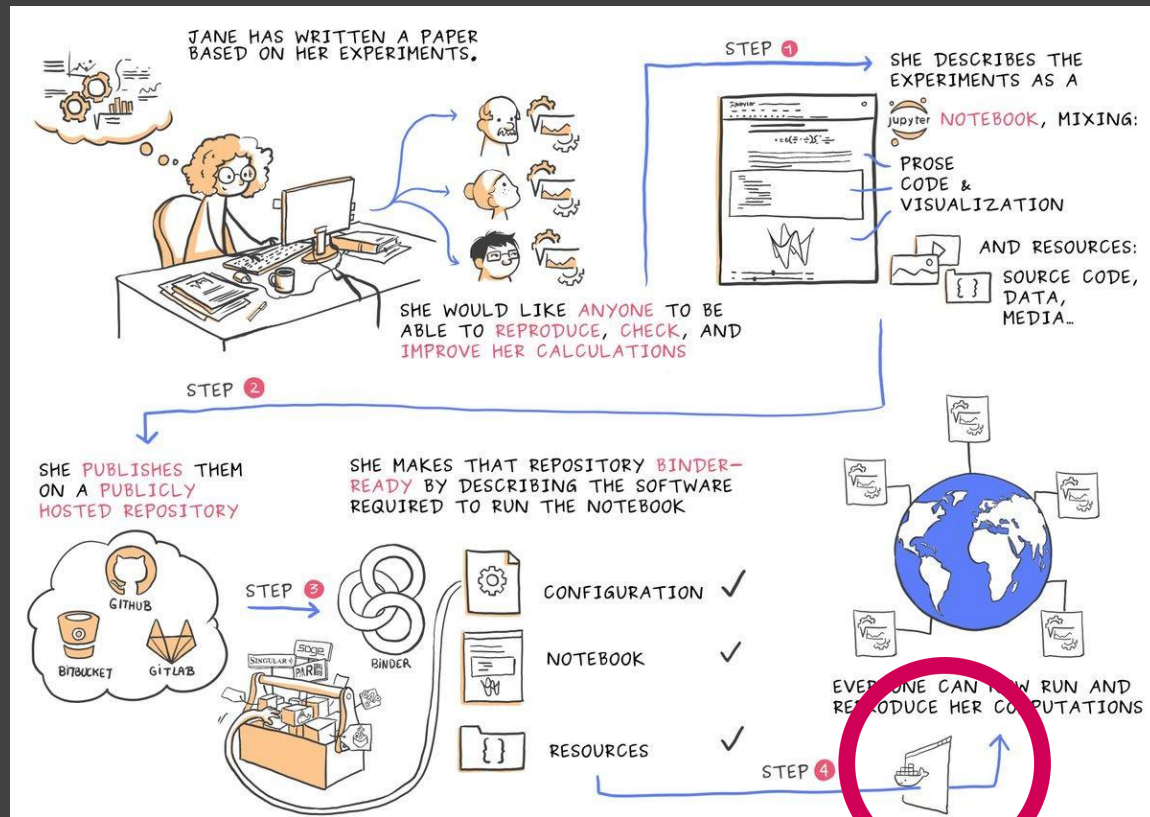
<https://discourse.jupyter.org/t/about-the-binder-category/200>

@turingway, CC-BY 4.0, The Turing Way, DOI: 10.5281/zenodo.6346555



Courtesy of Juliette Taka: <https://twitter.com/mybinderteam/status/1082556317842264064>

@turingway, CC-BY 4.0, The Turing Way, DOI: 10.5281/zenodo.6346555

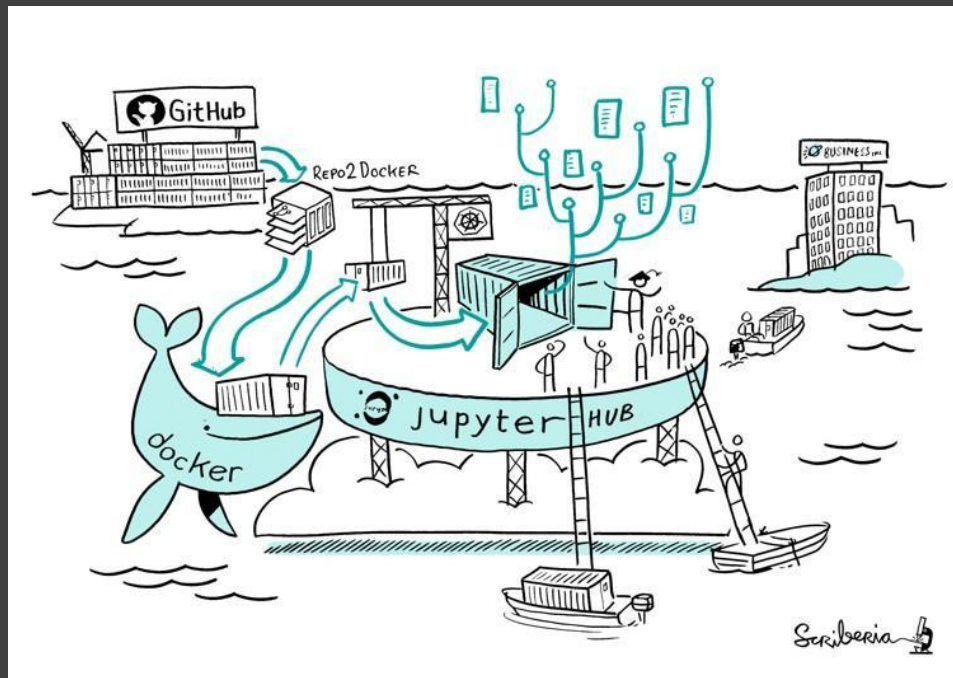


Courtesy of Juliette Taka: <https://twitter.com/mybinderteam/status/1082556317842264064>

@turingway, CC-BY 4.0, The Turing Way, DOI: 10.5281/zenodo.6346555

BinderHub

- cloud-based technology
- can launch a repository of code in a browser
- allows the user to execute and interact with the code

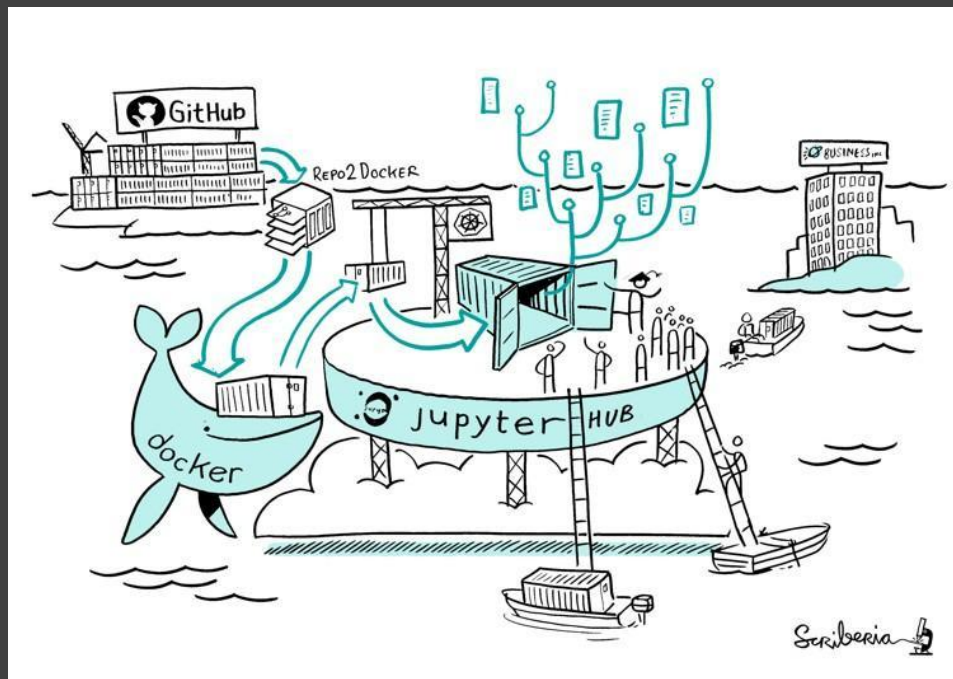


<https://the-turing-way.netlify.app/reproducible-research/binderhub/binderhub-compute.html>

@turingway, CC-BY 4.0, The Turing Way, DOI: 10.5281/zenodo.6346555

BinderHub

- repo2docker
- Kubernetes
- Helm
- JupyterHub
- a cloud service platform




<https://the-turing-way.netlify.app/reproducible-research/binderhub/binderhub-compute.html>

@turingway, CC-BY 4.0, The Turing Way, DOI: 10.5281/zenodo.6346555

mybinder.org

- online service for sharing computational environments from online repositories
- a federation of BinderHub deployments

Thanks to Google Cloud, OVH, GESIS Notebooks and the Turing Institute for supporting us! 🍷



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](#).

Build and launch a repository


GitHub repository name or URL

GitHub

Git ref (branch, tag, or commit) Path to a notebook file (optional)

HEAD File

Copy the URL below and share your Binder with others:

Copy the text below, then paste into your README to show a binder badge:  [launch binder](#)

<https://mybinder.org/>

gke.mybinder.org



Run by [The Binder Team](#)

Funded by [Google Cloud Platform](#)

ovh.mybinder.org



Run by [The OVH Team](#)

Funded by [OVH](#)

gesis.mybinder.org



Run by [The GESIS Notebooks Team](#)

Funded by [GESIS](#)

turing.mybinder.org



Run by [The Turing Way team led by Sarah Gibson](#)


Funded by [The Alan Turing Institute](#)




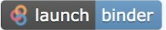

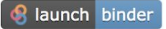




<https://mybinder.readthedocs.io/en/latest/about/about.html>

@turingway, CC-BY 4.0, The Turing Way, DOI: 10.5281/zenodo.6346555

Example 1

This example demonstrates a reproducible 4-step workflow for predicting a protein fold classification using a Machine Learning approach.

Rule 9: Design Your Notebooks to Be Read, Run, and Explored. The nbviewer links below provide a non-interactive preview of notebooks and  buttons launch Jupyter Notebook or Jupyter Lab in your web browser using the Binder (mybinder.org) server (may be slow!). (See the Binder website how to setup links to a Git repository.) The HTML links provide a permanent static record of the notebooks. All notebooks can also be launched directly from the links in the 0-Workflow.ipynb top-level notebook.

Nbviewer	Jupyter Notebook	Jupyter Lab	HTML
0-Workflow.ipynb			HTML
1-CreateDataset.ipynb			HTML
2-CalculateFeatures.ipynb			HTML
3-FitModel.ipynb			HTML
4-Predict.ipynb			HTML

Thanks to Google Cloud, OVH, GESIS Notebooks and the Turing Institute for supporting us! 🦊



Starting repository: `jupyter-guide/ten-rules-jupyter/master`


Take a look at our [gallery of example repositories](#).

Build logs

show

Example 1

This example demonstrates a reproducible 4-step workflow for predicting a protein fold classification using a Machine Learning approach.

Rule 9: Design Your Notebooks to Be Read, Run, and Explored. The nbviewer links below provide a non-interactive preview of notebooks and  buttons launch Jupyter Notebook or Jupyter Lab in your web browser using the Binder (mybinder.org) server (may be slow!). (See the Binder website how to setup links to a Git repository.) The HTML links provide a permanent static record of the notebooks. All notebooks can also be launched directly from the links in the 0-Workflow.ipynb top-level notebook.

Nbviewer	Jupyter Notebook	Jupyter Lab	HTML
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1-CreateDataset.ipynb			HTML
2-CalculateFeatures.ipynb			HTML
3-FitModel.ipynb			HTML
4-Predict.ipynb			HTML

The image shows a Jupyter Notebook interface. On the left is a file browser for the directory `/example1/`. It contains a table of files and folders, all last modified 5 months ago:

Name	Last Modified
data	5 months ago
intermediate_data	5 months ago
0-Workflow.html	5 months ago
0-Workflow.ipynb	5 months ago
1-CreateDataset....	5 months ago
1-CreateDataset....	5 months ago
2-CalculateFeatu...	5 months ago
2-CalculateFeatu...	5 months ago
3-FitModel.html	5 months ago
3-FitModel.ipynb	5 months ago
4-Predict.html	5 months ago
4-Predict.ipynb	5 months ago
mlutils.py	5 months ago
pdbutils.py	5 months ago
protectors.py	5 months ago

The main notebook area is titled "1-CreateDataset.ipynb" and shows the following content:

Create Dataset

This notebook extracts from the Protein Data Bank information about the secondary structure of proteins. The ultimate goal is to assign a fold classification from a protein sequence.

Rule 2: Document the Process, Not Just the Results. Here we describe the steps how to produce the dataset.

Rule 7: Build a Pipeline. Besides documenting all steps, the entire process of dataset creation from the original data files in the `/data` directory is automated. There are no manual steps.

Rule 8: Share and Explain Your Data. To enable reproducibility we provide a `/data` directory with data files and a file that describes the datasets with download locations and dates.

```
[1]: # column names
value_col = "foldClass" # fold class to be predicted

[2]: import pandas as pd
import numpy as np
import pdbutils
```

Kirstie Whitaker

“Binder is great because it also encourages reproducible practices in the communication.”



@kirstie_j

@turingway, CC-BY 4.0, The Turing Way, DOI: 10.5281/zenodo.6346555

Hands-on session

Zero to Binder, a guided tour of building a Binder resource:

* Link to the full Tutorial: <http://bit.ly/zero-to-binder-python>

Preparing to Launch your First Repo

Zero to Binder, a guided tour of building a Binder resource:

* Link to the full Tutorial: <http://bit.ly/zero-to-binder-python>

Preparing a repository for Binder

1. The repository is in a public location online
2. The repository does not require any personal or sensitive information (such as passwords)

Preparing a repository for Binder

1. The repository is in a public location online
2. The repository does not require any personal or sensitive information (such as passwords)
3. The repository contains content designed for people to read

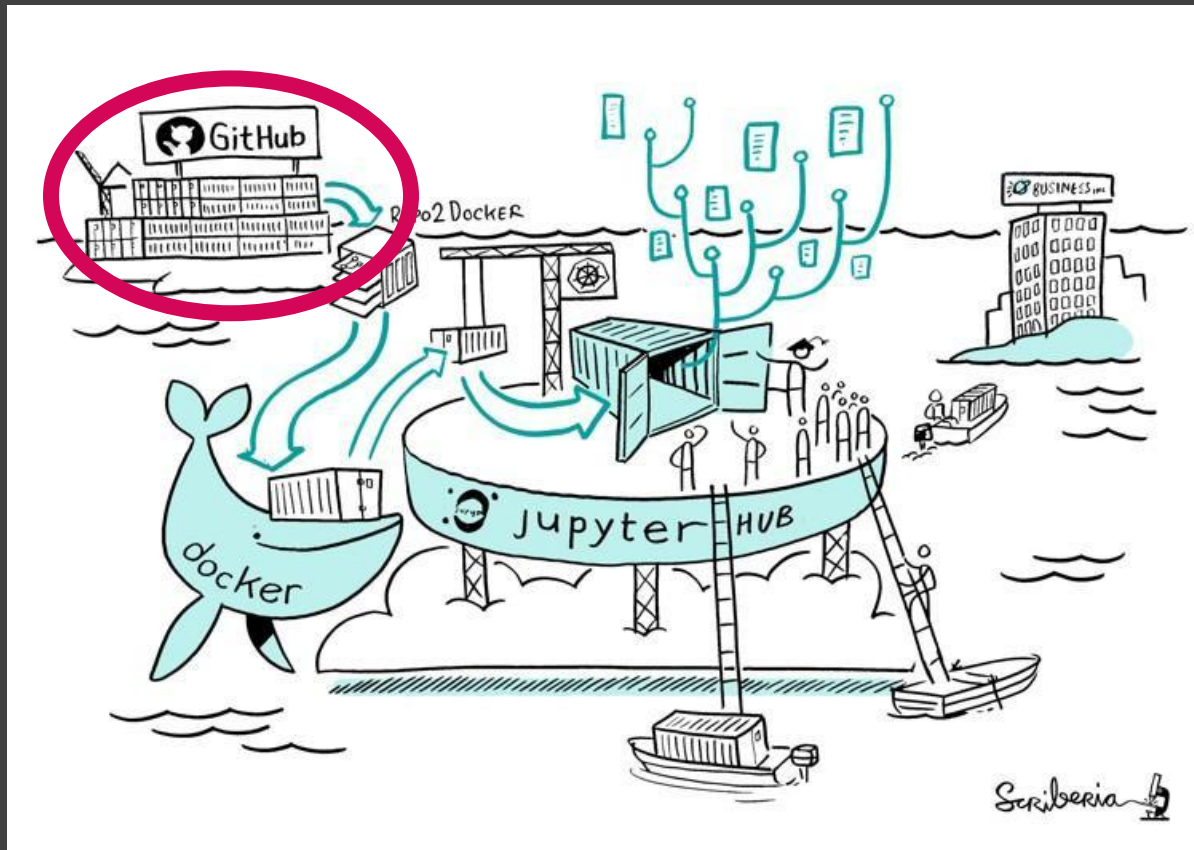
Launch your First Repo

1. Go to <https://mybinder.org>
2. Type the URL of your repo into the "GitHub repo or URL" box. It should look like this: <https://github.com/your-username/my-first-binder>
3. As you type, the webpage generates a link in the "Copy the URL below..." box It should look like this:
<https://mybinder.org/v2/gh/your-username/my-first-binder/HEAD>
4. Copy it, open a new browser tab and visit that URL.
 - You will see a "spinner" as Binder launches the repo.

Preparing a repository for Binder

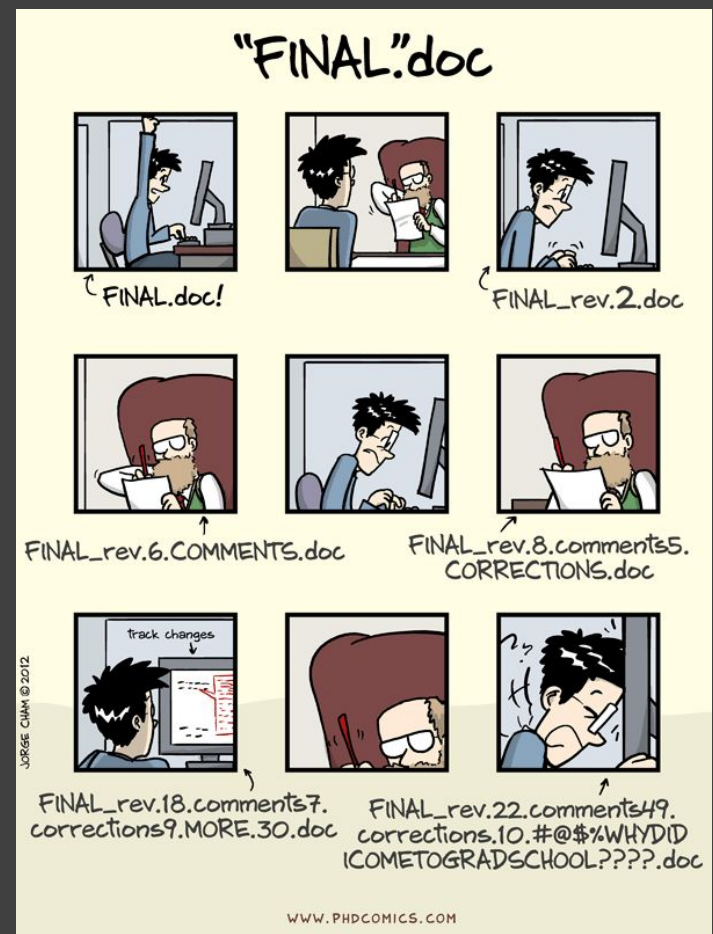
1. The repository is in a public location online
2. The repository does not require any personal or sensitive information (such as passwords)
3. The repository contains content designed for people to read
4. The repository has configuration files that specify its computational environment

How does *Binder* work?



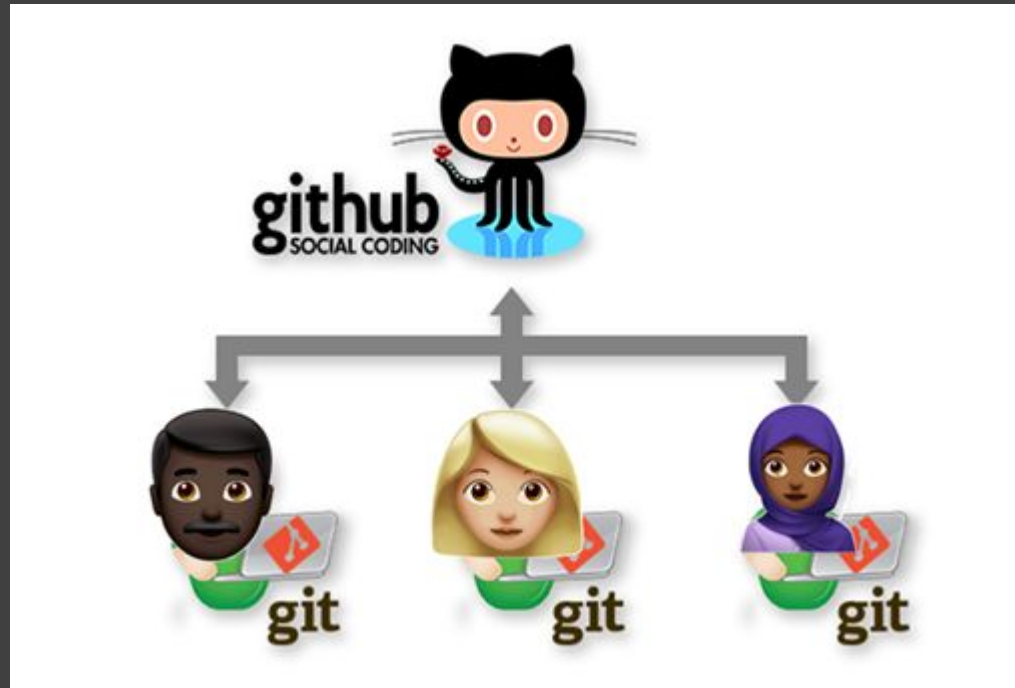


- version control system
- records changes to a file or set of files over time
- provides access to any specific version



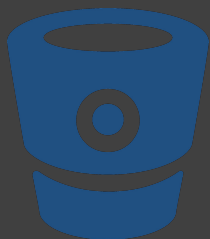


- online hosting platform for git repositories
- enables collaborative work



Preparing a repository for Binder

1. The repository is in a public location online



Preparing a repository for Binder

1. The repository is in a public location online
2. The repository does not require any personal or sensitive information (such as passwords)
3. The repository contains content designed for people to read.
4. The repository has **configuration files** that specify its computational environment

Configuration file

- defines your computational environment

```
requirements.txt ×
1 # Requirements for the demo notebooks
2 # Useful for MyBinder configuration
3 pandas==1.0.5
4 numpy==1.19.0
5 matplotlib==3.2.2
6 datascience
7 folium
8 jupyter-book==0.8.2
9 sphinxcontrib-bibtex==1.0.0
10
```

pip freeze

- Python specific
- captures the versions of all packages that you're currently using
- can print to screen or save in a file named `requirements.txt`

Examples

1. Generate output suitable for a requirements file.

[Unix/macOS](#) [Windows](#)

```
$ python -m pip freeze
docutils==0.11
Jinja2==2.7.2
MarkupSafe==0.19
Pygments==1.6
Sphinx==1.2.2
```

2. Generate a requirements file and then install from it in another environment.

[Unix/macOS](#) [Windows](#)

```
env1/bin/python -m pip freeze > requirements.txt
env2/bin/python -m pip install -r requirements.txt
```

CONDA

- environment, package and version management system
- for multiple languages
- Information about installed software saved in file called `environment.yml`



```
! environment.yml x
1  name: example-environment
2  channels:
3  | - conda-forge
4  dependencies:
5  | - python
6  | - numpy
7  | - pip
8  | - pip:
9  | | - nbgitpuller
10 | | - sphinx-gallery
11 | | - pandas
12 | | - matplotlib
13 | |
```



R environment




- support for R and RStudio with libraries pinned to a specific snapshot on MRAN, defined in `runtime.txt`
- `install.R` specifies one library to install per line

File	Action
<code>.github/workflows</code>	Add workflow to create a binder badge comment on
<code>bus-dashboard</code>	Remove the DESCRIPTION file
<code>LICENSE</code>	Create LICENSE
<code>README.md</code>	Add r-conda to the README as an alternative
<code>index.ipynb</code>	adding example
<code>install.R</code>	Add example Shiny app
<code>runtime.txt</code>	Update runtime.txt

r / runtime.txt

 **betatim** Update runtime.txt 



🔗 2 contributors  





Raw Blame   




1 lines (1 sloc) | 17 Bytes

```
1 r-3.6-2019-04-12
```

r / install.R

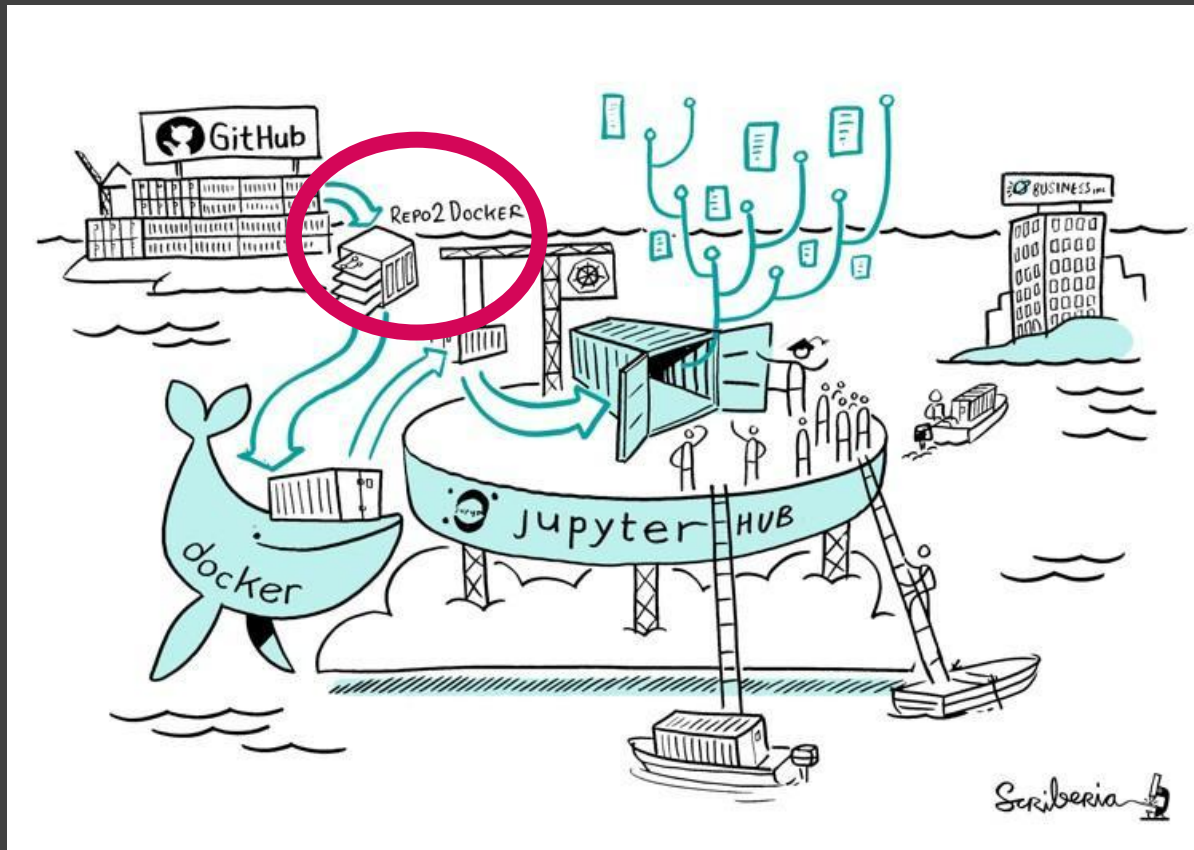
 **betatim** Add example Shiny app 

🔗 4 contributors    

Raw Blame   

5 lines (5 sloc) | 148 Bytes

```
1 install.packages("tidyverse")
2 install.packages("rmarkdown")
3 install.packages("httr")
4 install.packages("shinydashboard")
5 install.packages('leaflet')
```





repo2docker

- automatically builds a Docker image from a code repository given a configuration file

Calling repo2docker

repo2docker is called with this command:

```
jupyter-repo2docker <source-repository>
```

where `<source-repository>` is:

- a URL of a Git repository (<https://github.com/binder-examples/requirements>),
- a Zenodo DOI ([10.5281/zenodo.1211089](https://doi.org/10.5281/zenodo.1211089)),
- a SWHID ([swh:1:rev:999dd06c7f679a2714dfe5199bdca09522a29649](https://zenodo.org/record/1211089/files/swh:1:rev:999dd06c7f679a2714dfe5199bdca09522a29649)), or
- a path to a local directory (`a/local/directory`)

of the source repository you want to build.

For example, the following command will build an image of Peter Norvig's `Pytudes` repository:

```
jupyter-repo2docker https://github.com/norvig/pytudes
```

Building the image may take a few minutes.

Containers



- package a project with all of the parts it needs - such as libraries, dependencies, and system settings
- anyone can then open up a container and work within it
- the computational environment of the container is identical across instances

Containers



- behaves like a virtual machine
- more lightweight -> only contains the individual components needed to operate the project

Containers



master minimal-dockerfile / Dockerfile Go to file ...

betatim Switch USER at the end to not run as ro... Latest commit dbcf36b on 28 Feb 2019 History

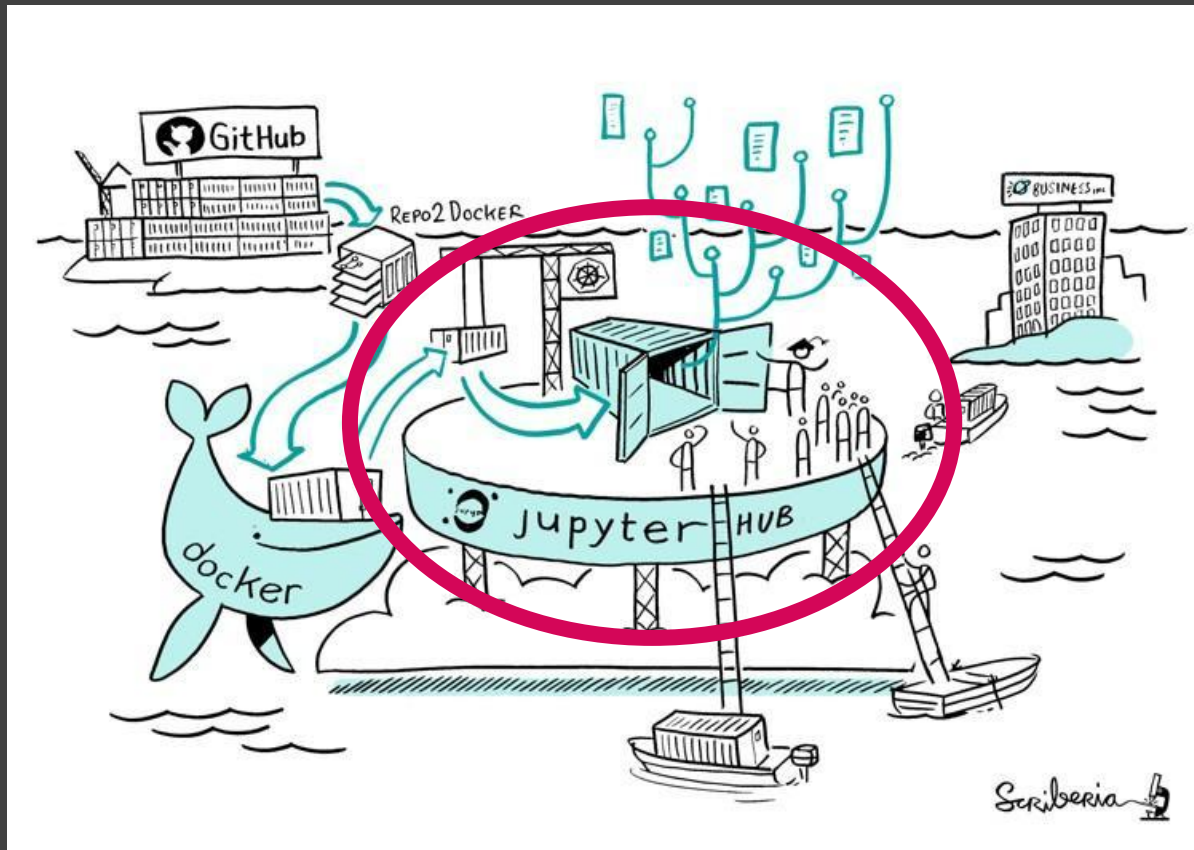
2 contributors

17 lines (15 sloc) 370 Bytes Raw Blame 📄 ✎ 🗑️

```
1 FROM python:3.7-slim
2 # install the notebook package
3 RUN pip install --no-cache --upgrade pip && \
4     pip install --no-cache notebook
5
6 # create user with a home directory
7 ARG NB_USER
8 ARG NB_UID
9 ENV USER ${NB_USER}
10 ENV HOME /home/${NB_USER}
11
12 RUN adduser --disabled-password \
13     --gecos "Default user" \
14     --uid ${NB_UID} \
15     ${NB_USER}
16 WORKDIR ${HOME}
17 USER ${USER}
```

<https://github.com/binder-examples/minimal-dockerfile/blob/master/Dockerfile>

@turingway, CC-BY 4.0, The Turing Way, DOI: 10.5281/zenodo.6346555



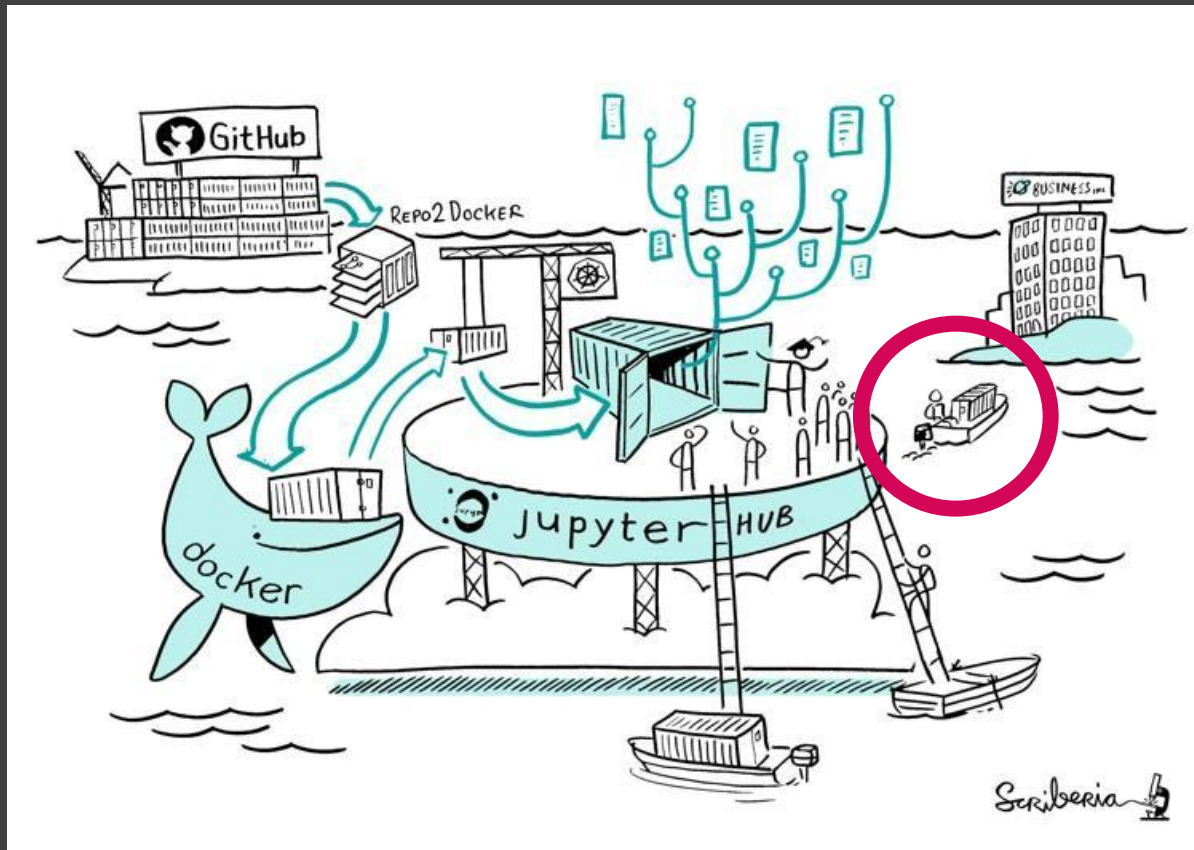
JupyterHub

- allows users to interact with a computing environment through a webpage
- “the cloud is just someone else’s computer” @kirstie_j



JupyterHub

1. JupyterHub creates a Kubernetes pod for the user that serves the built Docker image for the repository.
2. JupyterHub monitors the user's pod for activity, and destroys it after a short period of inactivity.



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<https://mybinder.org/v2/gh/your-username/my-first-binder/HEAD>
4. Copy it, open a new browser tab and visit that URL.
 - You will see a "spinner" as Binder launches the repo.

What happens in the background?

While you wait for the launch, BinderHub (the backend of Binder) is:

- Fetching your repo from GitHub
- Reading the contents
- Creating a Docker file based on your repo requirement
- Launching that Docker image in the Cloud
- Connecting you to it via your browser


Let's run a Python file

- Create a file called `hello.py` via the web interface with `print("Hello from Binder!")` on the first line and commit to the main branch
- In the top right corner, click "New"  "Terminal"
- In the new tab with the terminal, type `python hello.py` and press return

Let's add dependencies

- In your repo, create a file called requirements.txt
- Add a line that says: `numpy==1.14.5`
- Check for typos! Then commit to the main branch.
- Visit <https://mybinder.org/v2/gh/your-username/repo/HEAD> again in a new tab
- *BinderHub will read requirements.txt and install version 1.14.5 of the numpy package.*

Check the Environment

- In the top right corner, click "New"  "Python 3" to open a new notebook
- Type the following into a new cell

```
import numpy  
print(numpy.__version__)  
numpy.random.randn()
```
- Run the cell to see the version number and a random number printed
- Continue with this Binder Lesson:

Sarah Gibson

“It took me a while to feel like I knew enough to contribute to Binder. But the team are always so excited to have my input. It’s really motivating to be part of such a welcoming community.”



Thank you!

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- These slides reuse previously developed materials by Kirstie Whitaker, Sarah Gibson, Martina Vilas, Malvika Sharan, Emma Karoune, + many in *The Turing Way* community
- README and LICENSE section reuses Open Life Science Materials

Useful Links

- **Book:** the-turing-way.netlify.com
- **Twitter:** twitter.com/turingway
- **Newsletter:** tinyletter.com/TuringWay
- **GitHub:** github.com/alan-turing-institute/the-turing-way
- **Original artwork by Scriberia:** <https://doi.org/10.5281/zenodo.3332807>

