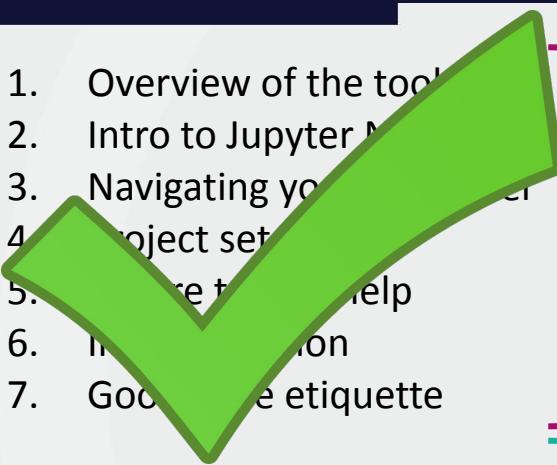


The background features a complex, glowing purple network of interconnected lines and dots, resembling a digital or biological structure. A large, semi-transparent white circle is positioned on the right side. In the top left corner, there's a white curved shape containing a small cluster of purple dots and a few short line segments.

Day 2



Outline

- 
- 1. Overview of the tools
 - 2. Intro to Jupyter Notebook
 - 3. Navigating your environment
 - 4. Project setup
 - 5. Where to find help
 - 6. Introduction to Python
 - 7. Good practice etiquette
 - 8. Data processing with pandas
 - 9. Automation with python
 - 10. Reproducibility
 - 11. Version control
- 
- Day 1
- Day 2

Intro to Pandas

Open the notebook called:
06_Intro_to_pandas.ipynb

Intro to Automation

Open the notebook called:
07_Intro_to_automation.ipynb

Day 2 – Reproducibility



Reproducibility

What is it?

The ability for you, or others, to redo your work in order to:

1. Check for correctness (confirmation)
2. Use the same method on new data (adaptation)
3. Use a different method on the same data (modification)



Is reproducibility important?

The reproducibility crisis in psychology

RESEARCH ARTICLE

Estimating the reproducibility of psychological science

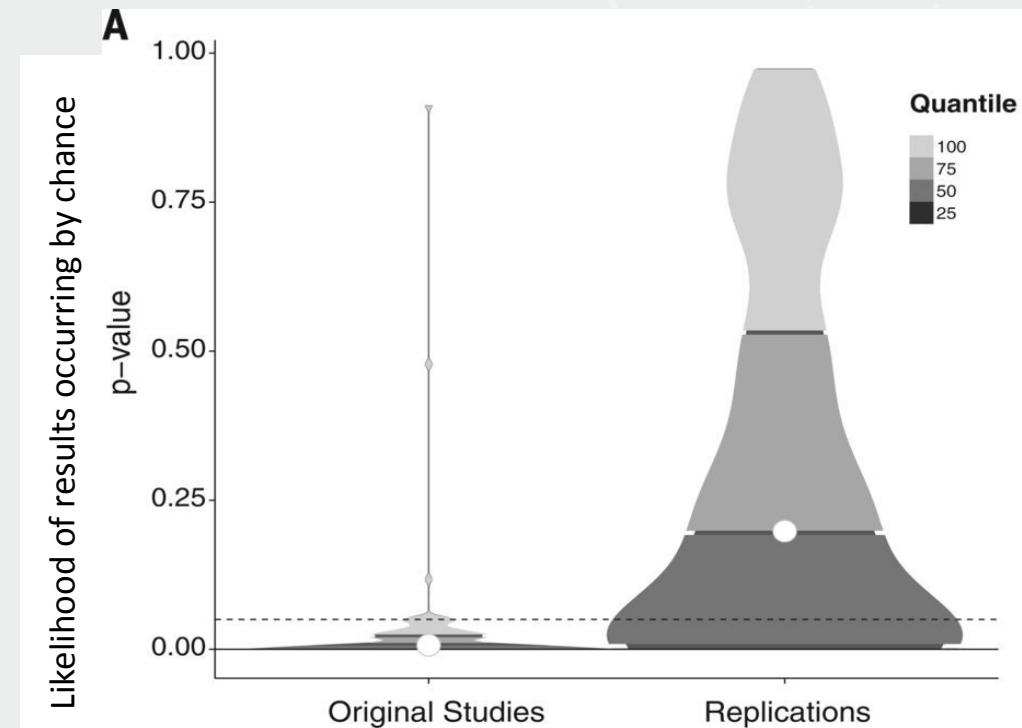
Open Science Collaboration^{*,†}

+ Author Affiliations

[†]Corresponding author. E-mail: nosek@virginia.edu

Science 28 Aug 2015:
Vol. 349, Issue 6251,
DOI: 10.1126/science.aac4716

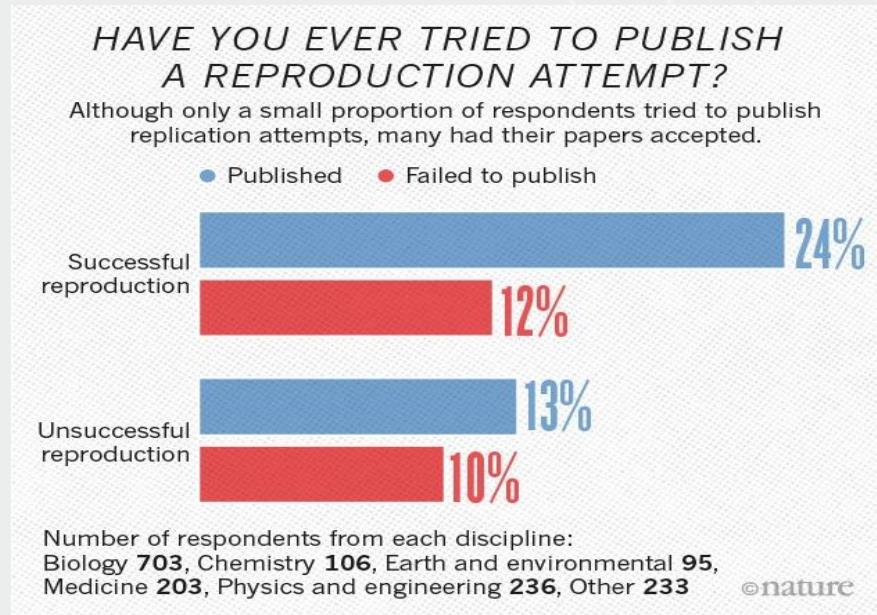
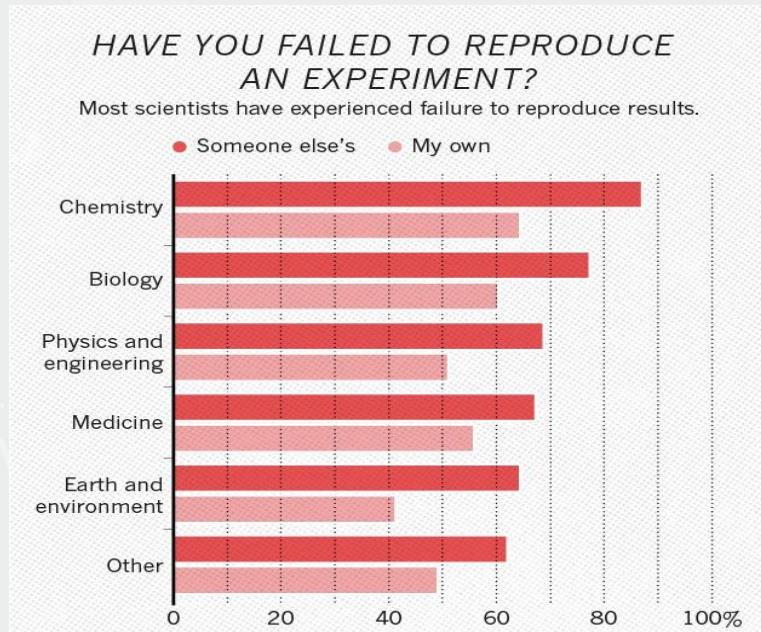
<http://dx.doi.org/10.1126/science.aac4716>





We should all be worried about reproducibility

Science is based on reproducible results, but many results cannot be reproduced.





Reproducibility

Why is it important?

If you cannot recreate your results with the same data and same method then you should not trust your results.

Confirmation is important for building confidence and trust.

Adaptation allows you to reuse methods that you are confident with.

Modification is a natural part of the development process and ultimately saves time.



Scenario - You predict a big loss/gain for your area

5:00
Stop

You want to be sure of your results before you present them

Rerunning your analysis from a fresh start and getting the same result builds confidence

How do you achieve a “fresh run” of your analysis?

1. Full automation
2. Human readable recipe
3. Scripted process
4. Start again and hope that you can find all the relevant post-it-notes

Be brave and share your experience



Scenario - You predict a big loss/gain for your area

You want to be sure of your results before you present them

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How do you achieve a “fresh run” of your analysis?

1. Full automation
2. Human readable recipe
3. Scripted process
4. Start again and hope that you can find all the relevant post-it-notes

A year later your predictions don't pan out.

Can you still defend your analysis, and identify where your model diverged from reality?



Scenario - Loss of personnel

5:00
Stop

A key person in your org has moved to another group/company.

This person was responsible for generating reports that are critical to your business. They use a semi-automated system, and it normally takes them 30 mins to complete a report.

- Could you continue their work?
 - Where is the data, code, info on what they are doing?
- How long would it take you to run their reporting?
- How can you improve this scenario?

Does your process pass the lotto factor?



Documentation

Documentation is key to reproducibility

- Documentation is the idea of documenting your procedures for your experiment/process so that an outsider could understand the workings of your team
 - The audience is a user
- Remember the Lotto factor
- Documentation is a love letter to your future self - Damian Conway



Nomenclature and Naming Conventions

Language is important, precise language even more so

- Clear and concise language aids transparency
- Standard names and terminology makes language accessible
- Avoid jargon/TLAs where possible
 - Describe/introduce when necessary
 - A reference document for terminology is useful
- Apply to all communication channels



Nomenclature and Naming Conventions

Language is important, precise language even more so

- File [+ Program/Process/Function/Variable] naming should be consistent
 - Team members won't have to over think the naming process
- Easier to facilitate access, retrieval and storage of files
- Easier to browse through files saving time and effort
- Harder to lose!



Case study

Former PhD student and subsequent founder of the Figshare platform, Mark Hahnel, typified a common challenge:

“During my PhD I was never good at managing my research data. I had so many different file names for my data that I always struggled to find the correct file quickly and easily when it was requested. My former PI was so horrified upon seeing the state of my data organisation that she held an emergency lab book meeting with the rest of my group when I was leaving.”

Research Information, April/May 2014



Example

As previously suggested, consistent and meaningful naming of files and folders can make everyone's life easier. See this example below:

YYYYMMDD_SiteA_SensorB.CSV to encode Date Location Sensor

Which when applied, would look like this below:

20150621_Yaouk_Humidity.CSV

Some characters may have special meaning to the operating system so avoid using these characters when you are naming files. These characters include the following: / \ " ' * ; - ? [] () ~ ! \$ { } < # @ & | **space** tab newline



Discussion

5:00
Stop

Good or Bad names

1. Myfile.docx
2. 20200130_core_instructions_draft.docx
3. Fig 2.png
4. 20200130_core_instructions_v1.docx
5. My figure (3) v2.png
6. fig02_core_instructions_presentation.png



Ten Simple Rules for Reproducible *Computational* Research

1. For every result, keep track of how it was produced
2. Avoid manual data manipulation steps
3. Archive the exact versions of all external programs used
4. Version control all custom scripts
5. Record all intermediate results, when possible in standardized formats
6. For analyses that include randomness, note underlying random seeds
7. Always store raw data behind plots
8. Generate hierarchical analysis output, allowing layers of increasing detail to be inspected
9. Connect textual statements to underlying results
10. Provide public access to scripts, runs, and results



Ten Simple Rules for Reproducible *Computational* Research

1. For every result, keep track of how it was produced
2. Avoid manual data manipulation steps

”

As a **minimum**, you should at least record sufficient details on programs, parameters, and manual procedures to allow yourself, in a year or so, to approximately reproduce the results.

”

If working at the UNIX command line, manual modification of files can usually be replaced by the use of standard UNIX commands or small custom scripts.

”

Do by hand to learn/test/explore, then encode this in a script or instruction set that future you, and future others, could understand and run



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Learn from the mistakes of others

Don't be "An Astronomer"

OPEN ACCESS Freely available online

PLOS ONE

CrossMark click for updates

How Do Astronomers Share Data? Reliability and Persistence of Datasets Linked in AAS Publications and a Qualitative Study of Data Practices among US Astronomers

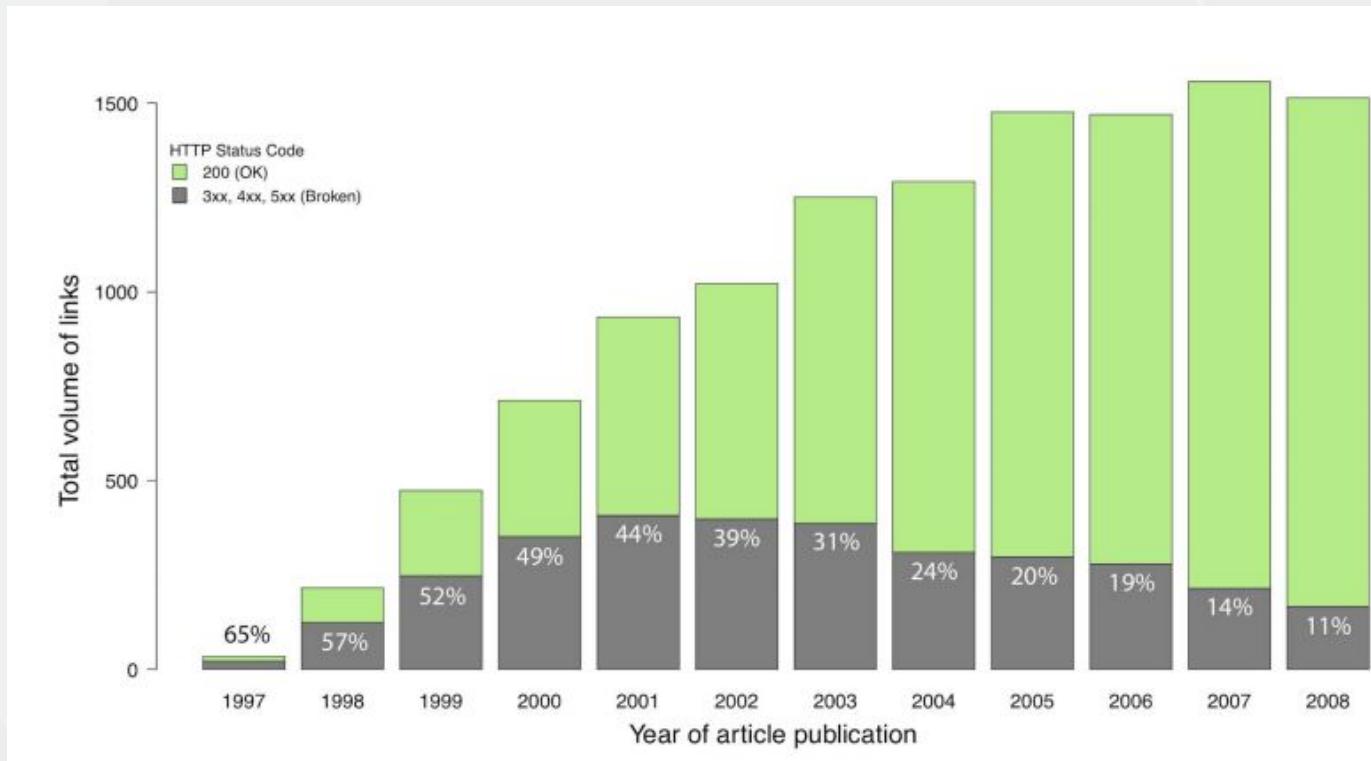
Alberto Pepe^{1,2*}, Alyssa Goodman^{1,2}, August Muench¹, Merce Crosas², Christopher Erdmann¹

¹ Harvard-Smithsonian Center for Astrophysics, Cambridge, Massachusetts, United States of America, ² Institute for Quantitative Social Science, Harvard University, Cambridge, Massachusetts, United States of America

No, I don't have a website where I store these data. Most of it is in various stages of mess. —An Astronomer



Learn from the mistakes of others





Learn from the mistakes of others

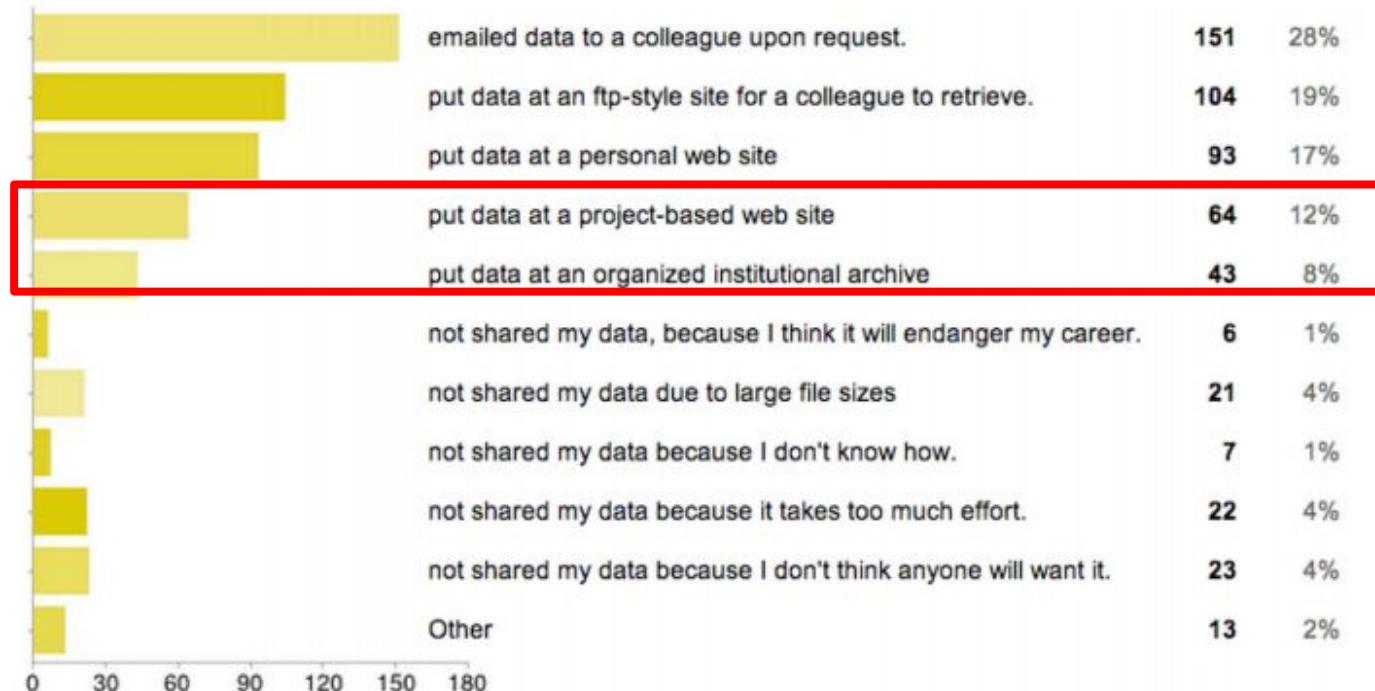


Figure 5. Survey results to question 2: Data sharing practices. Survey results to question: When it comes to sharing DATA you've created, collected or curated, you have?
doi:10.1371/journal.pone.0104798.g005



Best Practice

Make reproducibility part of your daily routine

Retrofitting is difficult

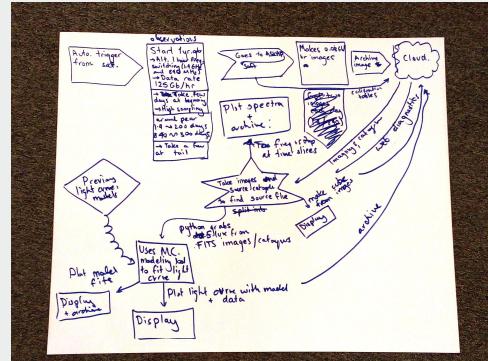
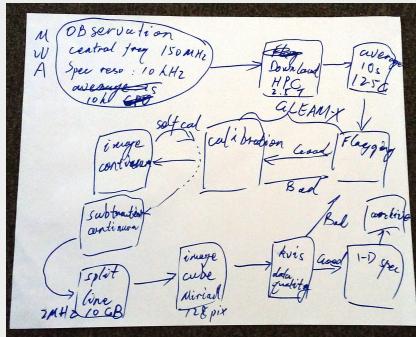
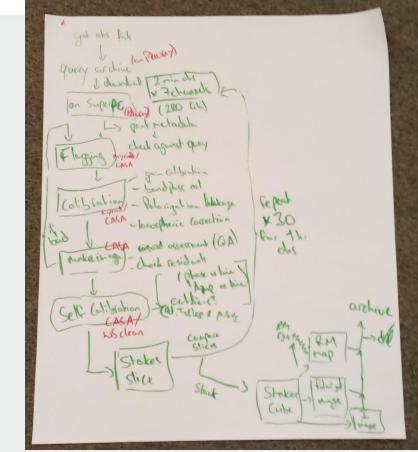
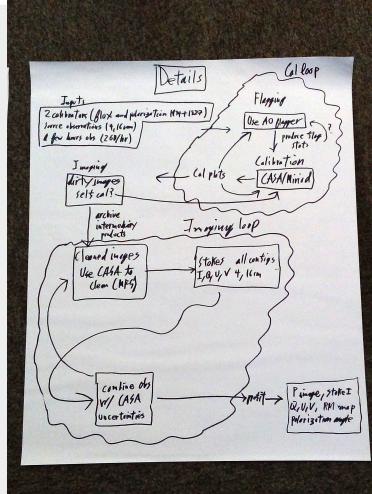
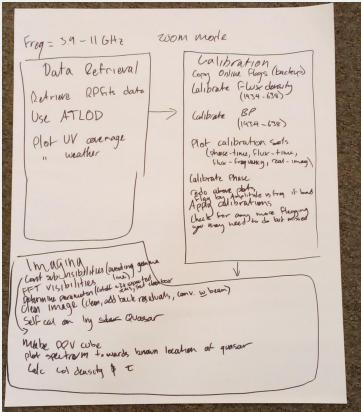
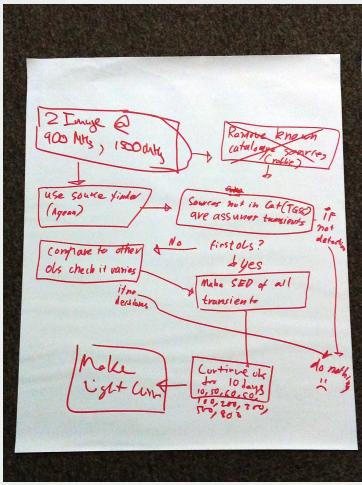
Commit to making your **next** project reproducible from day 1

Good automation, documentation, and organisation make reproducibility almost free

Visualise each task/project as a workflow which needs to be orchestrated with inputs, outputs, and processes.

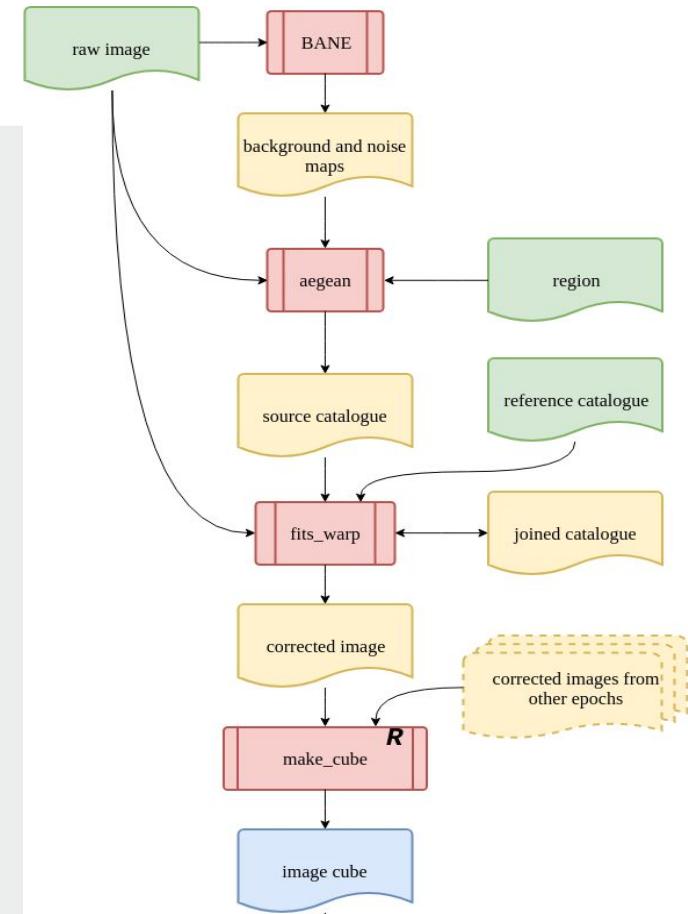
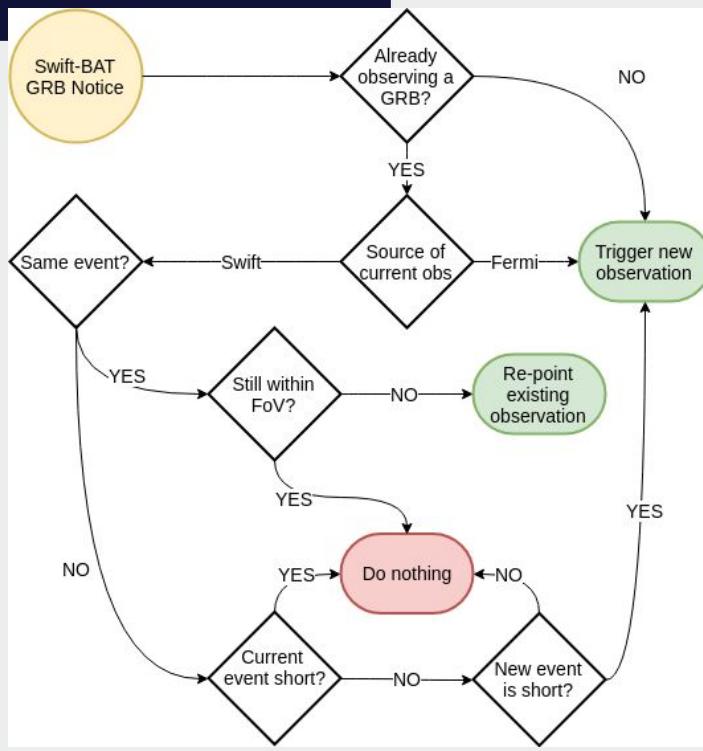


Day 0 Plans





Final implemented workflows



Reproducibility Recap

Making your work/research/reporting/decision making reproducible:

- Builds confidence in results
- Allows trusted methods to be applied to new information
- Saves time and effort by allowing extension/modification

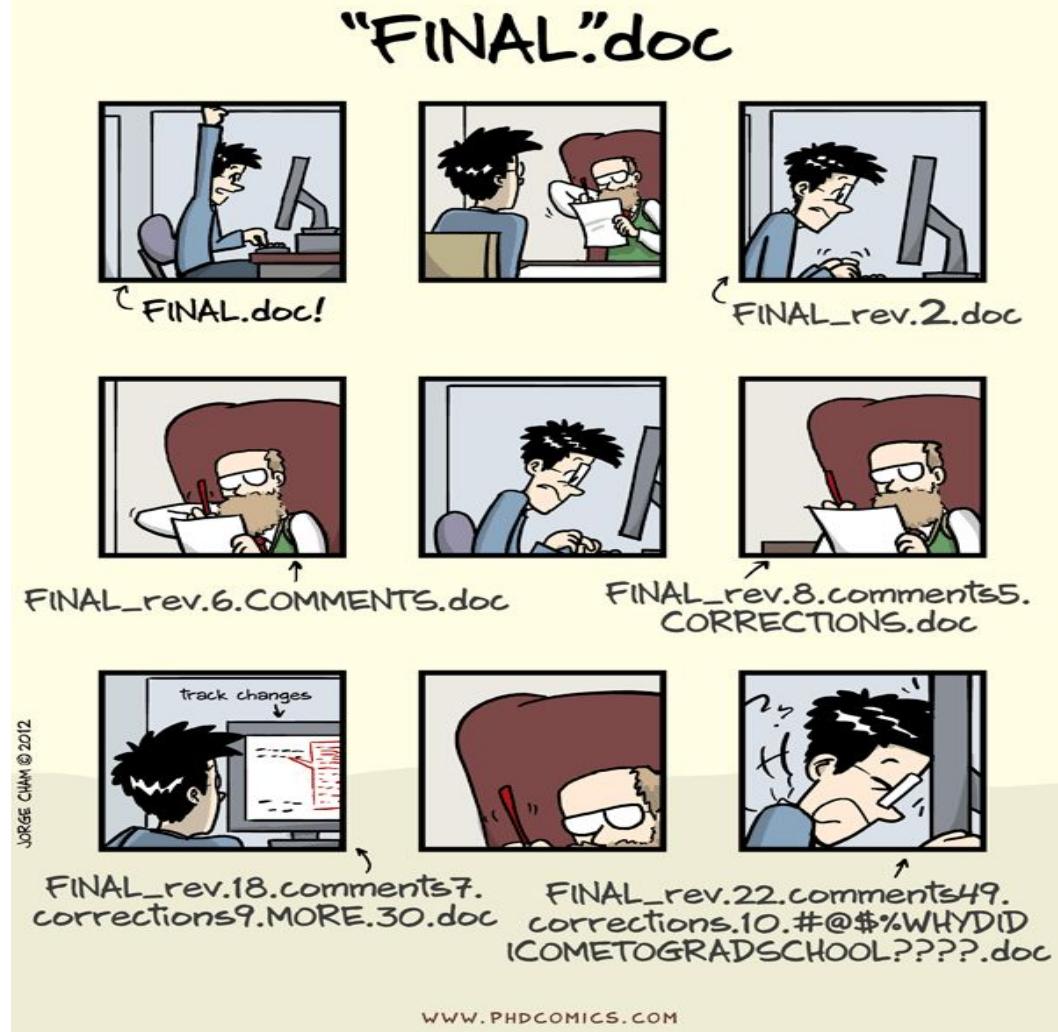
Reproducibility is easy when:

- Workflows are automated
- Metadata is stored
- You embrace it on day 1, not day N-1.

Day 2 – Version Control



The Problem





Why Version Control?

As data scientist, we spend much of our time writing code, whether it be for data cleaning, machine learning, or visualisation. As such, our codes are often constantly evolving. By putting all of our code under version control we can:

- **tag code** versions for later reference (*via tags*).
- record a **unique identifier** for the exact code version used to produce a particular plot or result (*via commit identifiers*).
- **roll back** our code to previous states (*via checkout*).
- **identify** when/how **bugs** were introduced (*via diff/blame*).
- **keep multiple versions** of the same code in sync with each other (*via branches/merging*).
- efficiently **share and collaborate** on our codes with others (*via remotes/online hosting*).



Why Version Control?

It's important to also realise that many of the advantages of version control are not limited to just managing code. For example, it can also be useful when writing papers/reports. Here we can use version control to:

- **bring back** that paragraph we accidentally deleted last week.
- **try out a different structure** and simply disregard it if we don't like it.
- **concurrently work on a report** with a collaborator and then **automatically merge** all of our **changes** together.

The upshot is ***you should use version control for almost everything***. The benefits are well worth it...



Introducing Git

Many version control systems exist, we love Git

*Git is a free and open source **distributed** version control system designed to handle everything from small to very large projects with speed and efficiency.*

Git is easy to learn and has a tiny footprint with lightning fast performance. It outclasses SCM tools like Subversion, CVS, Perforce, and ClearCase with features like cheap local branching, convenient staging areas, and multiple workflows. [Git website](#)

- *Distributed* -> everyone has their own complete copy of the entire repository and can make changes as they like
- Committing to a ‘central’ repository can be done once happy with the changes
- As opposed to *Subversion*, access to the central repo is not required to make changes
- Git is fast (primarily written in C & shell script) and lightweight (as you only track changes)
- Written by Linus Torvalds (creator of Linux)



Introducing Git

Git is a free and open source distributed version control system designed to handle everything from small to very large projects with speed and efficiency.

Git tracks changes made to files. You can initialise a Git Repository (the .git/ folder inside a project folder) for a project to build a history of your project over time:

- Added files/folders
- Changed files/folders
- Renamed files/folders
- Removed files/folders

Including by who, and when

Why GitHub or GitLab

- ❑ Version Control remotely
- ❑ Visible code and reproducibility
- ❑ Open code and reuse
- ❑ Collaborative code development
- ❑ Open code development



GitHub or GitLab advantages

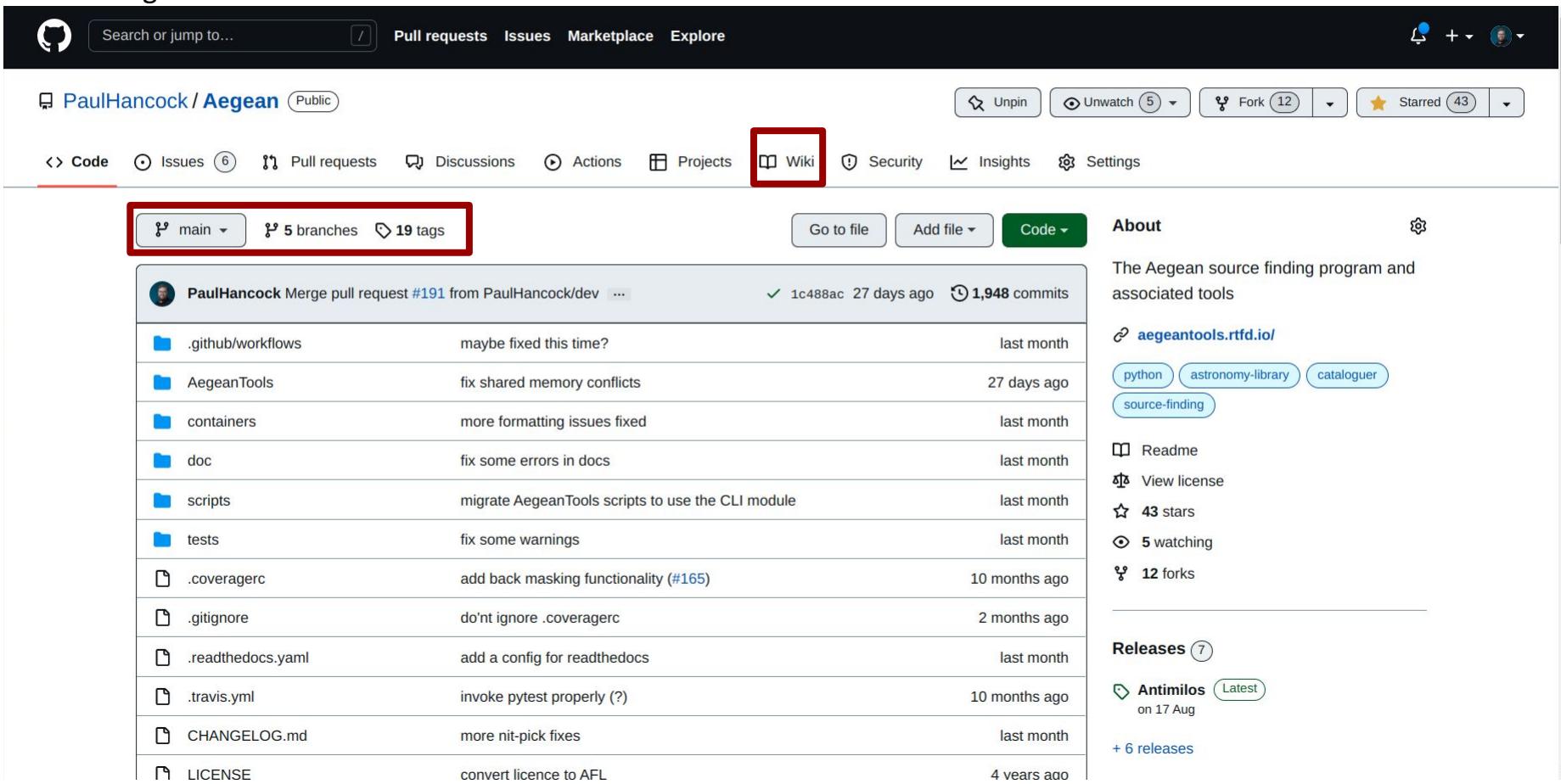
Why we like GitHub/GitLab

- Collaboration tools
- Developer tools
- 3rd party integrations
- Online backup of your work
- Easy/free publish/share work (or keep private)

Example: <https://github.com/PaulHancock/Aegean>

Easy Branch/Tag Management

“Onsite” Documentation



The screenshot shows a GitHub repository page for **PaulHancock / Aegean**. The top navigation bar includes links for **Pull requests**, **Issues**, **Marketplace**, and **Explore**. On the right side, there are buttons for **Unpin**, **Unwatch** (5), **Fork** (12), **Starred** (43), and user profile icons.

The main repository page has tabs for **Code**, **Issues** (6), **Pull requests**, **Discussions**, **Actions**, **Projects**, and **Wiki**. The **Wiki** tab is highlighted with a red box. Below the tabs, there are buttons for **Go to file**, **Add file**, and **Code**.

The repository summary section shows **main** branch, **5 branches**, and **19 tags**. It also displays a merge pull request from **PaulHancock / dev** with commit details: **1c488ac** 27 days ago, **1,948 commits**. A list of recent commits follows:

File / Commit Message	Date
.github/workflows maybe fixed this time?	last month
AegeanTools fix shared memory conflicts	27 days ago
containers more formatting issues fixed	last month
doc fix some errors in docs	last month
scripts migrate AegeanTools scripts to use the CLI module	last month
tests fix some warnings	last month
.coveragerc add back masking functionality (#165)	10 months ago
.gitignore do't ignore .coveragerc	2 months ago
.readthedocs.yaml add a config for readthedocs	last month
.travis.yml invoke pytest properly (?)	10 months ago
CHANGELOG.md more nit-pick fixes	last month
LICENSE convert licence to AFL	4 years ago

The **About** section provides information about the project: "The Aegean source finding program and associated tools". It includes a link to **aegeantools.rtd.io/** and tags for **python**, **astronomy-library**, **cataloguer**, and **source-finding**. It also lists **Readme**, **View license**, **43 stars**, **5 watching**, and **12 forks**.

The **Releases** section shows 7 releases, with the latest being **Antimilos** (Latest on 17 Aug) and 6 other releases.

Issue tracking and collaborative tools

Automated tools for testing and deployment

The screenshot shows a GitHub repository page for 'PaulHancock / Aegean'. The top navigation bar includes 'Pull requests', 'Issues', 'Marketplace', and 'Explore' tabs. Below the navigation is a search bar and several action buttons: 'Unpin', 'Unwatch (5)', 'Fork (12)', and 'Starred (43)'. The main navigation bar features 'Code', 'Issues (6)' (highlighted with a red box), 'Pull requests', 'Discussions', 'Actions' (highlighted with a red box), 'Projects', 'Wiki', 'Security', 'Insights', and 'Settings'.

Below the navigation is a search bar with filters: 'Filters ▾', 'Q is:issue is:open', 'Labels (16)', 'Milestones (1)', and a 'New issue' button. The main content area displays a list of issues:

Open/Closed	Title	Author	Label	Milestone	Assignee	Comments
6 Open ✓ 120 Closed						
replace dependency on lmfit with scipy fitting	Aegean	on hold				1
Support for different initial guess modes	enhancement					
Insight or guidance to improve bad results	question					3
Update API documentation for AegeanTools	documentation					5
user option to force source shapes during blind source finding	Aegean enhancement					
simultaneous fitting across multiple images	Aegean enhancement					

Each issue row includes a checkbox, the issue number, the title, the author, the label, the milestone, the assignee, and the number of comments. A 'ProTip!' at the bottom suggests mixing and matching filters to narrow down the search.

💡 ProTip! Mix and match filters to narrow down what you're looking for.



Version Control Recap

Version control helps:

- Identify when/where issues are introduced
- Facilitate safe testing of new ideas through branches
- Track milestones/versions through tagging
- Collaboratively work with *fewer* conflicts

Online Git repositories such as GitHub/GitLab add:

- A centralised repo
- Collaborative tools
- Development tools
- An automated deploy/test/document loop via 3rd party plugins
- A natural platform to share/advertise your excellent work

Saving our progress

Compress and download our notebooks

Select “terminal” from the launcher tab.

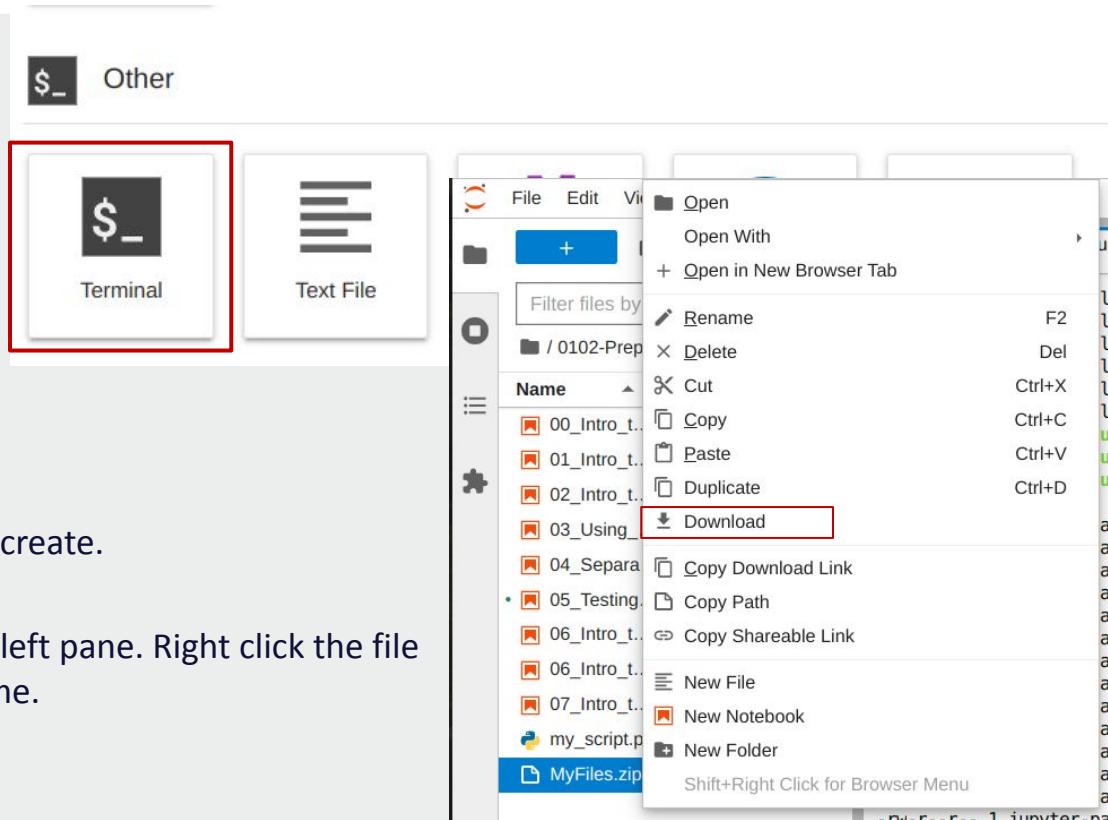
Move to the notebooks directory.

Zip the notebooks using:

```
> tar -caf MyFiles.zip *
```

Where MyFiles.zip is the file that we want to create.

You should see the file in the browser on the left pane. Right click the file and select “download” to take your work home.



Day 2 – Reflections



COREHUB.COM.AU/SKILLS

