Lecture 2
Reproducibility

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Data Science for Economics

Note: Materials for this lecture are drawn from Ted Miguel's Development Economics course at UC Berkeley

Agenda

1. Overview of reproducibility and transparency
2. Organizing files
3. Coding transparency and portability
4. Writing code in Python
5. More Python basics

Key references on research reproducibility

- Casey, Glennerster, and Miguel. (2012). “Reshaping Institutions: Evidence on Aid Impacts Using a Pre-analysis Plan”, Quarterly Journal of Economics, 127(4), 1755-1812.
- Miguel et al. (2014). “Promoting Transparency in Social Science Research”, Science, 10.1126/science.1245317.
- Christensen and Miguel. (2018). “Transparency, Reproducibility, and the Credibility of Economics Research”, Journal of Economic Literature, 56(3), 920-980.
- Ferguson et al. (2023). “Survey of open science practices and attitudes in the social sciences”, Nature Communications, 14.
- Christensen, Freese, and Miguel. (2019). Transparent and Reproducible Social Science Research: How to Do Open Science, University of California Press.

Threats to validity of research

- **Fraud**: undermines public trust in science
  - Open data and code can help uncover
- **Publication bias**: missing studies/ "file-drawer" problem
  - Wasted research effort, misguided policy decisions
  - Author manipulation/"p-hacking"
  - Pre-registration as a counter
- **Failure to replicate**: within study (reproduction) and across settings (replication)
  - Increasing journal data posting requirements
  - Difficulty of getting funding or publishing replications of studies

What do transparency and reproducibility mean in economics research?

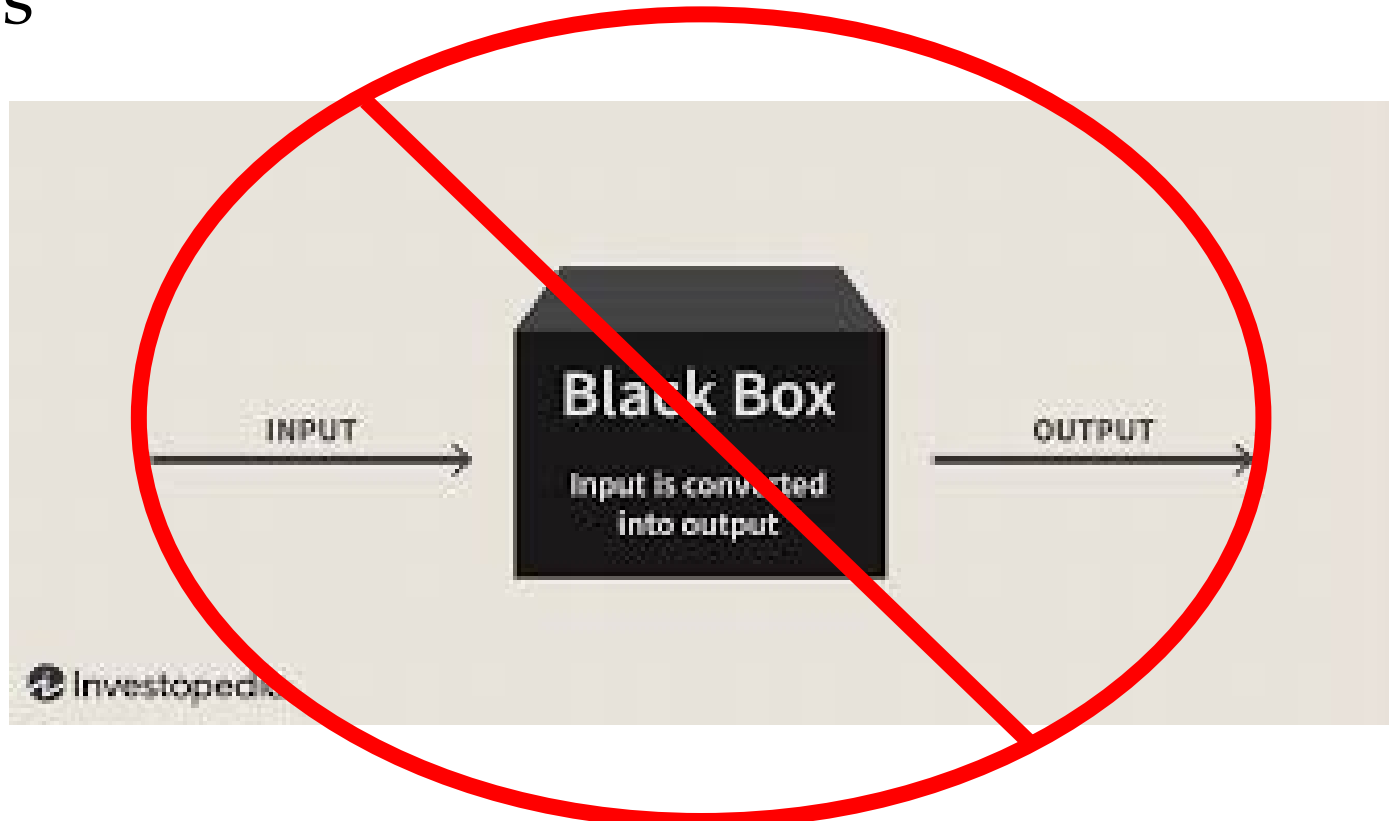
- **Transparency**:
  - Ensuring that all data, methods, and analyses are openly shared and clearly documented.
  - Allows others to understand and evaluate the research process, and to test for investigator bias or sensitivity to alternative methods.
- **Reproducibility**:
  - The ability of others to reproduce the results of a study using the original data and code provided by the researchers.
  - Critical for validating findings, building trust, and advancing knowledge.

Key principles of reproducibility

- **Accessible Data**: Provide well-documented, cleaned, and publicly available datasets (where ethically and legally possible).
- **Open Code**: Share analysis scripts and code in a version-controlled repository (e.g., GitHub) with clear instructions for execution.
- **Clear Methodology**: Document all steps in data collection, preprocessing, and analysis to ensure clarity and consistency.
- **Replication Workflow**: Design workflows that allow others to re-run analyses seamlessly, ensuring compatibility across systems and tools.

Takeaways for this course

- The quality and clarity of the research process matters
- Focus of this lecture: organizing and documenting research methods



Organizing files

The first step to a clear research process is file organization. There should be a clear structure to your folder hierarchy and file names.

```
README.pdf
data/
   raw/
      cps0001.dat
   analysis/
      combined_data.dta
      combined_data.csv
      combined_data_codebook.pdf
code/
  01_create/
      01_readcps.R
      02_readfred.R
  02_analysis/
      01_table1-5.R
      02_figures1-4.R
results/
  table1.tex
  table2.tex
  ...
  figure1.pdf
  figure2.pdf
```

Source

Example: Jigawa Floods Project

Basic structure:

1. Code
    1. High-frequency checks for data collection
    2. Analysis: separate scripts for different tasks
2. Data
    1. Raw
    2. Clean
3. Output
    1. Figures
    2. Maps
    3. Tables

Example: Locusts and Conflict Project

- Work in progress
- Mix of archived and active files
- Clear overarching structure
- Clean replication package

Example: Folder of data sources

1. Sub-folders for main data sources/data types
2. Clear organization within folders
3. Example: LSMS-ISA
    1. Country sub-folders
    2. Year/round sub-folders
    3. Data, Questionnaires, Resources sub-folders
    4. Zipped raw data and unzipped folders
        1. Always keep a copy of the original data as a backup!
4. Readme documents
    1. Notes on where/when data were accessed
    2. Example: ARES data

https://databank.illinois.edu/datasets/IDB-1107366

Retrieved 2/14/23

1. ARES, crop-specific exposure to temperature and moisture shocks
    a. 0.25 degree cells by year by crop
    b. 1961-2014
    c. /Users/pierrebiscaye/Dropbox/Data/Spatial/ARES
    d. .nc files with crop layers

File organization in this class

- Folders for each section
- Subfolders for data, images
- Separate Jupyter Notebooks for different topics
- Clear file naming conventions

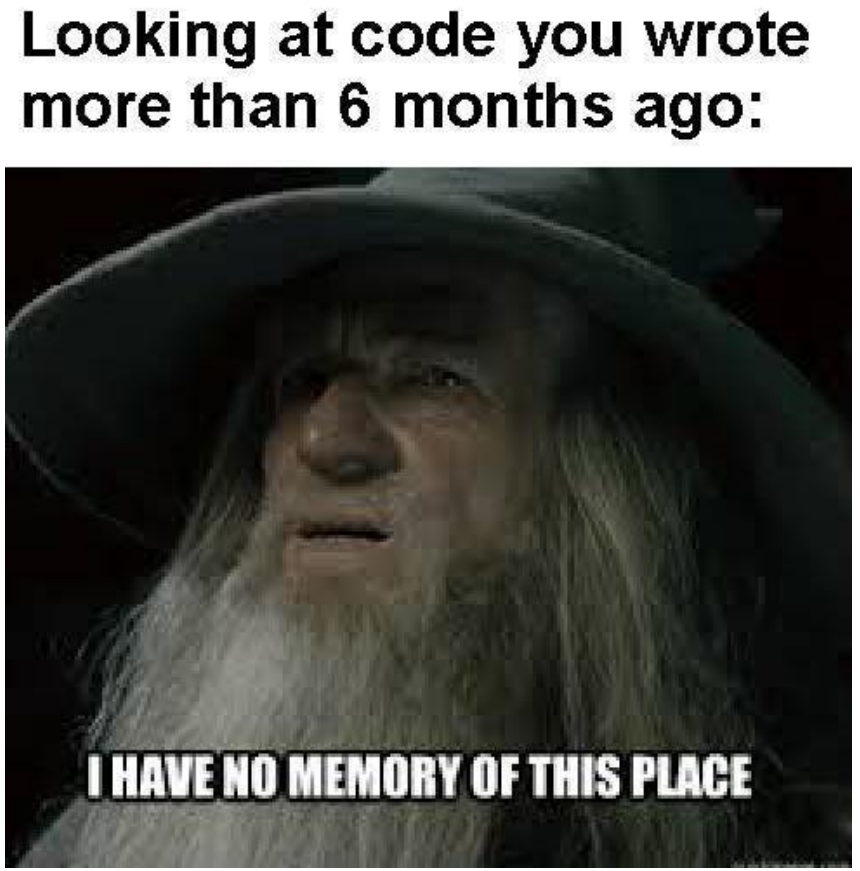
Transparent and reproducible code

1. Comment and document thoroughly
2. Use modular design
3. Adopt consistent naming conventions
4. Version control
5. Ensure code is portable

This is important for both your future self and for potential collaborators and reviewers!

1. Comment and document thoroughly

- Include clear comments in your code to explain the purpose of each section and the logic behind key steps.
- Best practice: Use a README file to provide an overview of the project, dependencies, and how to run the code.



Source

Commented code

```
//
// Prep spatial datasets
//

* Locust Swarms
{
* Data from FAO Locust Hub, coordinates and dates of locust swarm observations globally from 1985-2023
* Some additional detail included there but lots of missing data so focus just on swarm location and timing
import delimited "$data/Locust Hub/Retrieved 2.13.23/Swarms.csv", encoding(UTF-8) clear
drop if objectid==.
drop if missing(x) | missing(y)
destring x, replace
destring y, replace
gen double lat=round((y+0.125)*4)/4 - 0.125
gen double lon=round((x+0.125)*4)/4 - 0.125
gen date = date(substr(startdate,1,10), "YMD")
format date %td
gen year=year(date)
gen month=month(date)

* Output for mapping
preserve
keep x y year
export delimited "$clean/mapping_swarms.csv", replace
restore

* Match to countries, identify countries in Africa and Arabian Peninsula with at least 10 swarms in analysis period
1996-2018
geoinpoly y x using "$data/Country boundaries/Country raw/UIA_World_Countries_Boundaries/WORLD_coor.dta"
merge m:1 _ID using "$data/Country boundaries/Country raw/UIA_World_Countries_Boundaries/WORLD_data.dta", keep(1 3)
gen swarm=1 if year>1995 & year<2019
egen swarms_1996_2018=sum(swarm),by(COUNTRY)
bys COUNTRY: gen first=_n==1
br COUNTRY swarms_1996_2018 if swarms_1996_2018>10 & first==1
* Well over 10: Algeria, Burkina Faso, Chad, Egypt, Eritrea, Ethiopia, Gambia, Guinea, Guinea-Bissau, Libya, Mali,
Mauritania, Morocco/Western Sahara, Niger, Oman, Saudi Arabia, Senegal, Somalia, Sudan, Tunia, Yemen
* Cabo Verde 42, India 127, Iran 19, Israel 11, Kenya 22, Pakistan 108 also meet the swarm count criteria but outside
area of interest or are borderline cases (Israel, Kenya)

* Set target geographic area to trim other datasets, based on countries to target
drop if lat<-2.5
drop if lat>37.5
drop if lon<-17.5
drop if lon>60.25

```

Data Readme

This document outlines the data sources used in this study.

****
A. DATA INCLUDED IN REPLICATION PACKAGE
****

1. Locust swarms data

* Retrieved from FAO Locust Hub 2/13/2023 at https://locust-hub-hqfao.hub.arcgis.com/search?collection=dataset
* These data are no longer publicly accessible, so I include the raw downloaded data in the package
* The data includes coordinates and dates of locust swarm observations globally from 1985-2023
* Some additional detail included there but lots of missing data so focus just on swarm location and timing

Raw data file: Swarms.csv

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2. ACLED Conflict data

* raw dataset is full conflict data downloaded from ACLED on 11/26/20, https://acleddata.com/
* second datafile downloaded covers 02/11/20-02/03/23 and is merged with the first
* the data records all conflicts and actors from all events going as far back as the database - 1997
* selected geographies: East Africa, Middle Africa, Middle East, North Africa, West Africa, South Asia

Raw data files:
* 1997-01-01-2020-11-26-Eastern_Africa-Middle_Africa-Middle_East-Northern_Africa-South_Asia-Western_Africa.csv
* 2020-02-11-2023-02-03-Eastern_Africa-Middle_Africa-Middle_East-Northern_Africa-South_Asia-Western_Africa.csv

_____

3. Aggregated analysis data

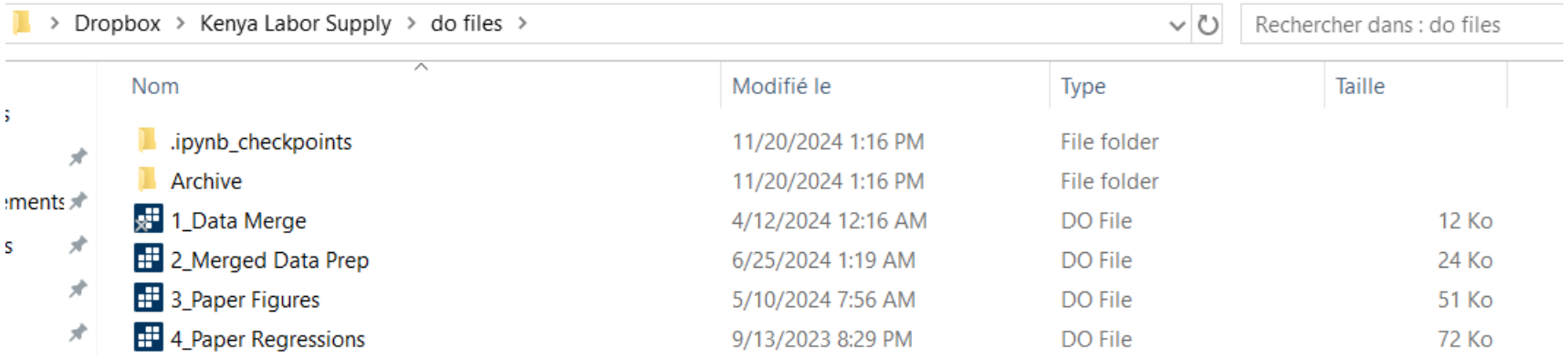
* Dataset at 0.25 degree grid cell by year level produced by aggregating swarm, conflict, and other data
* Description of inputs is below

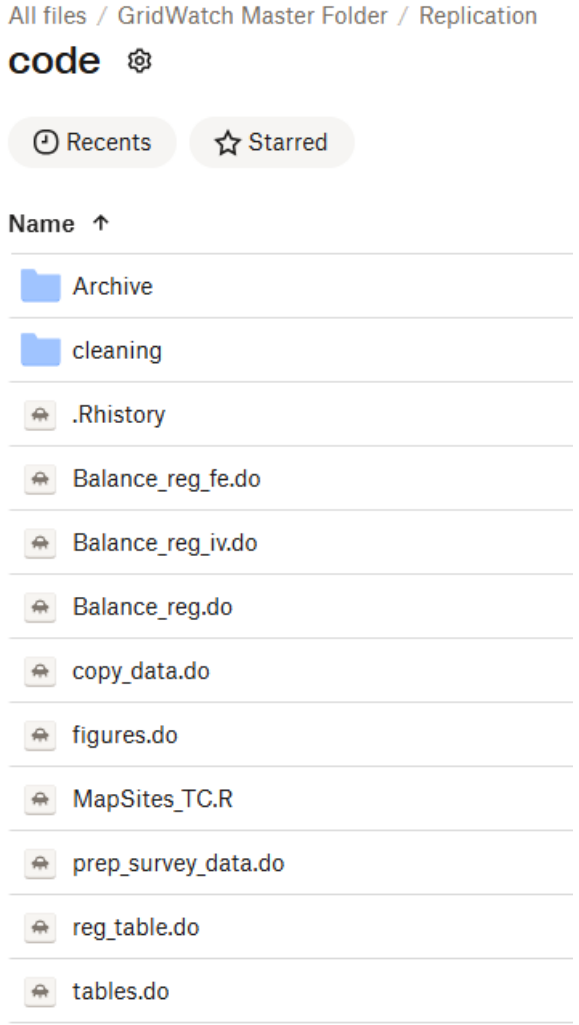
Processed data file: temp_2025.dta

****
B. INPUTS TO TEMP_2025.DTA
****

2. Use modular design

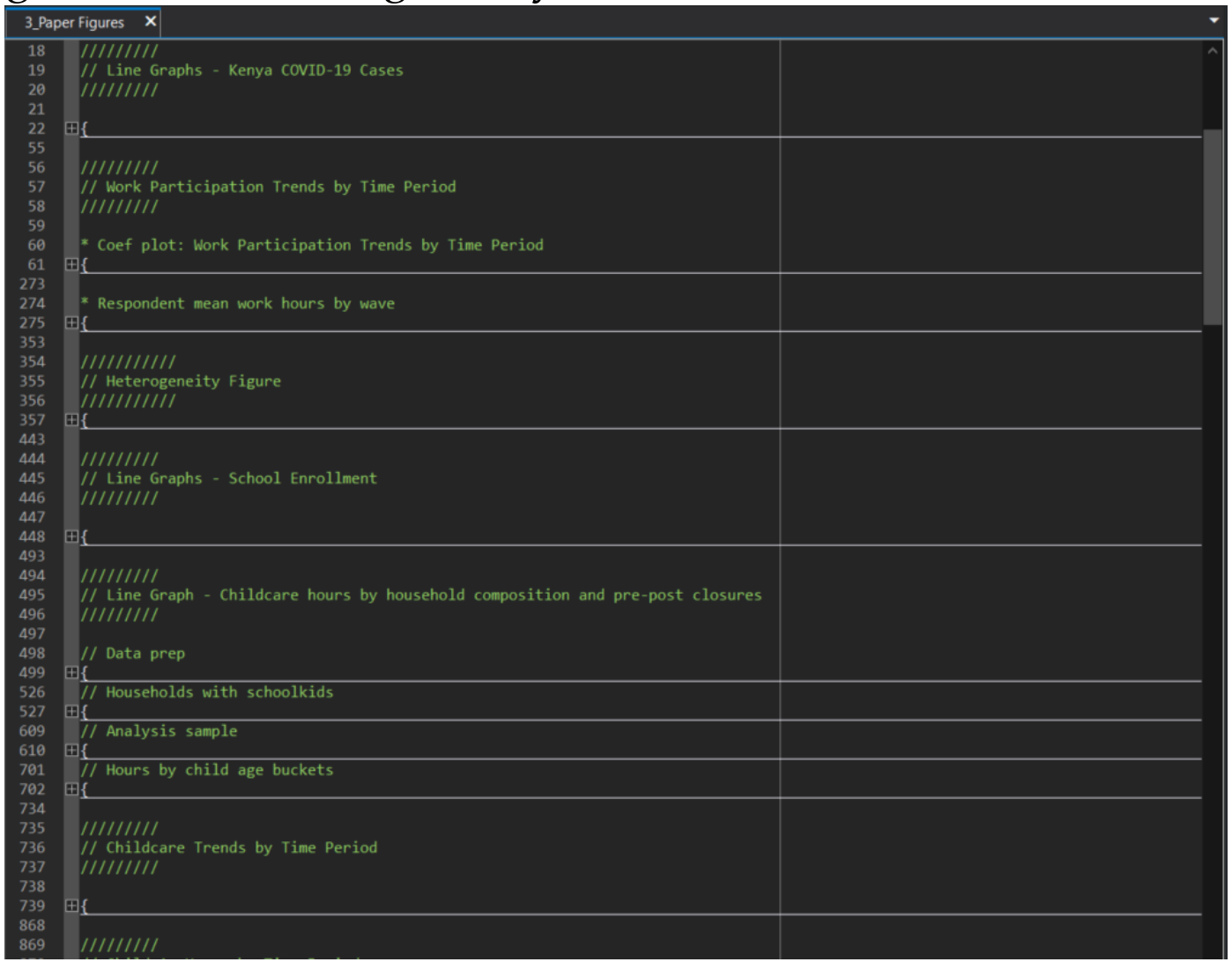
- Break the code into smaller, reusable functions or scripts to make it easier to debug, update, and understand. Separate data cleaning, analysis, and visualization steps logically.



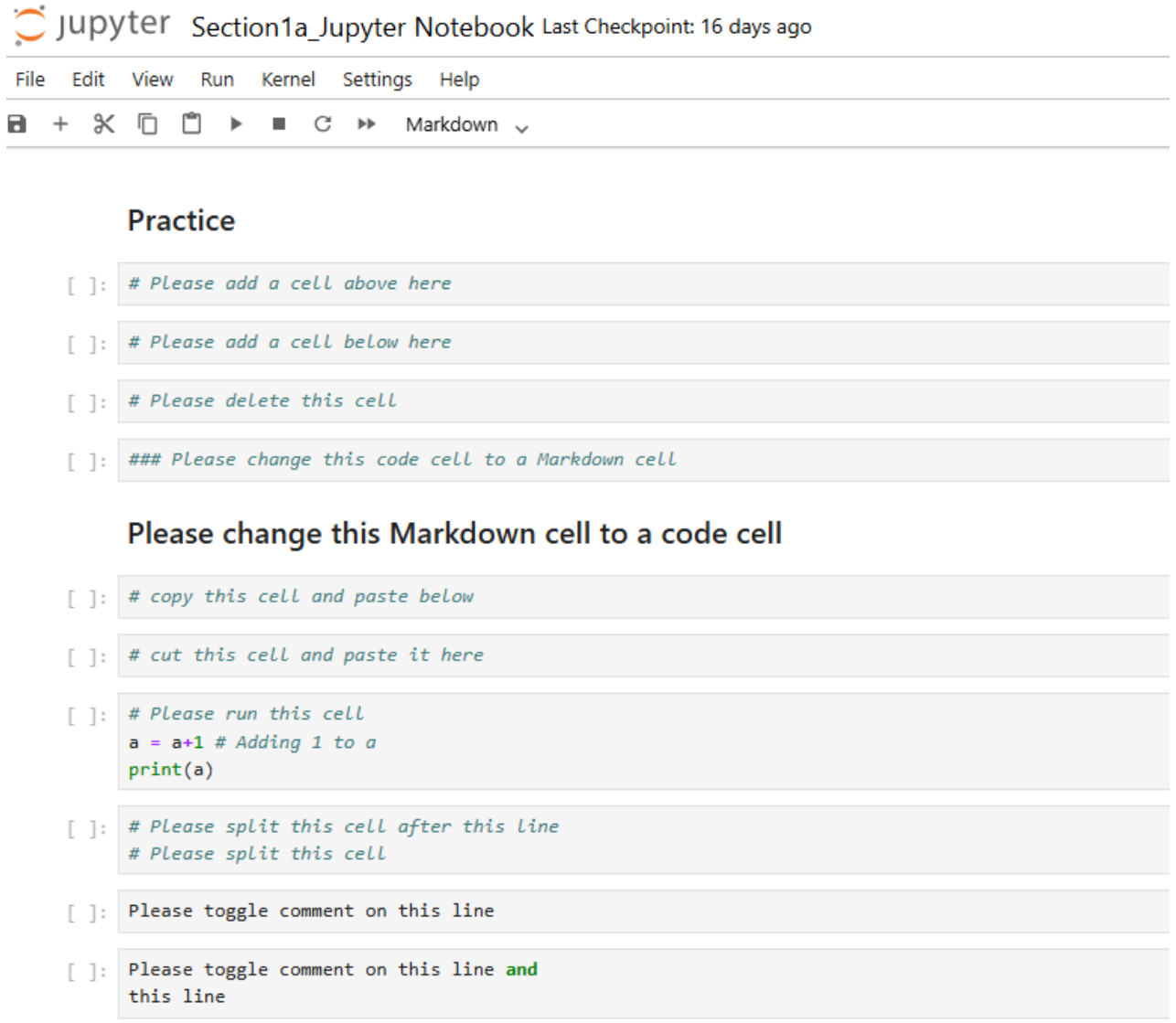


Modular design within scripts

Use clear headings and labels to organize your code



Jupyter Notebook: modular by default

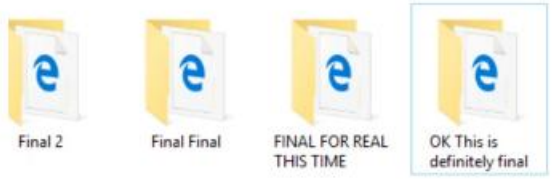


3. Adopt consistent naming conventions

- Use meaningful, consistent names for variables, functions, and files to enhance readability and reduce confusion.
- For example, use clean_data() instead of cd() for function names.

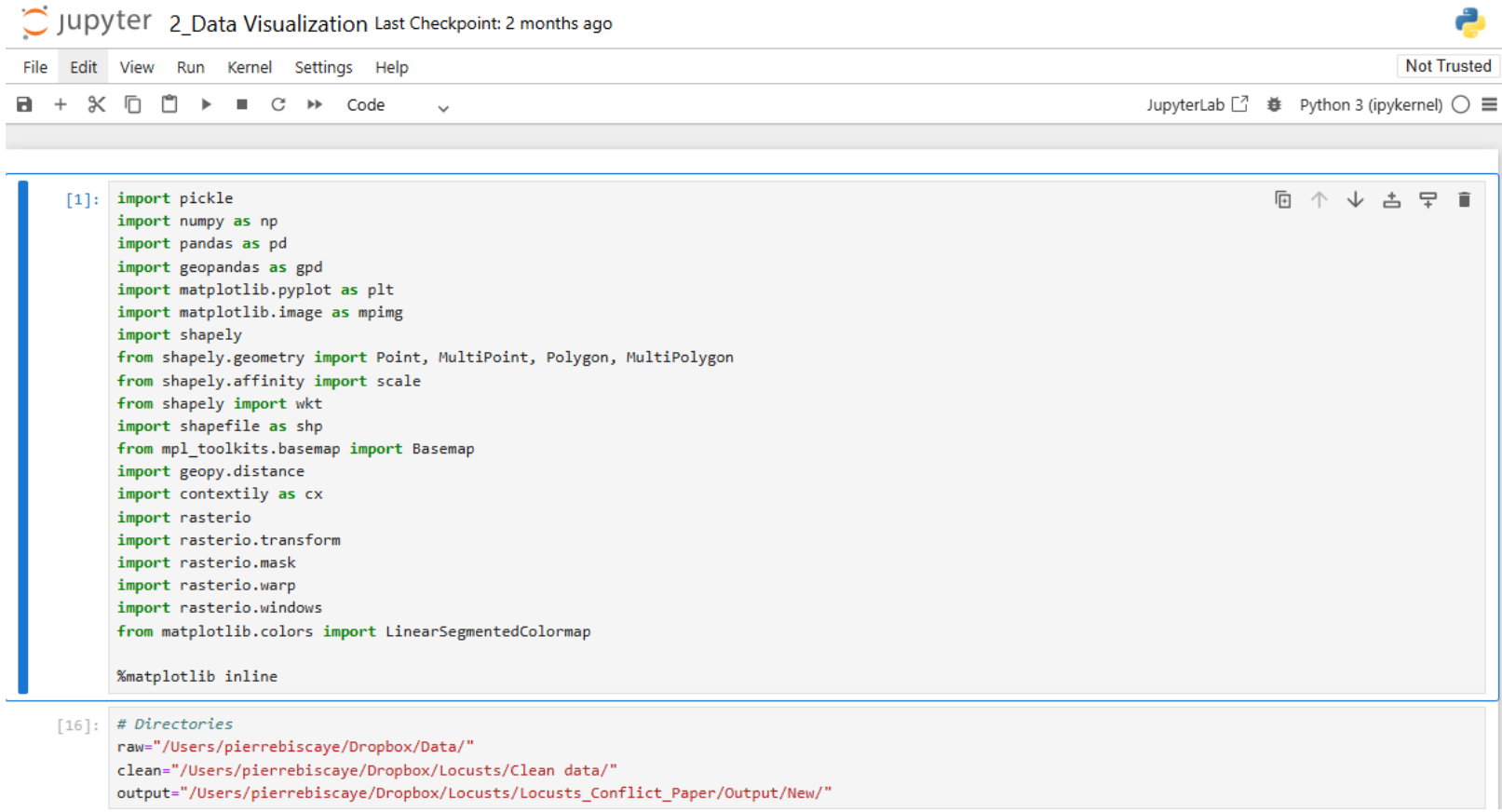
4. Version control

- Use version control systems to track changes in your code and collaborate efficiently.
- Commit changes with descriptive messages and maintain a well-organized repository structure.
- Gold standard: GitHub repository
- Minimum: clear files names indicating version history, well-structured archive folder
  - Back up your code, don't just always overwrite



5. Ensure code is portable

- Avoid hardcoding file paths or machine-specific dependencies.
- Use relative paths and specify software environments to ensure others can run the code seamlessly.
- Often useful to have a "0_setup script" with paths and packages to run first.

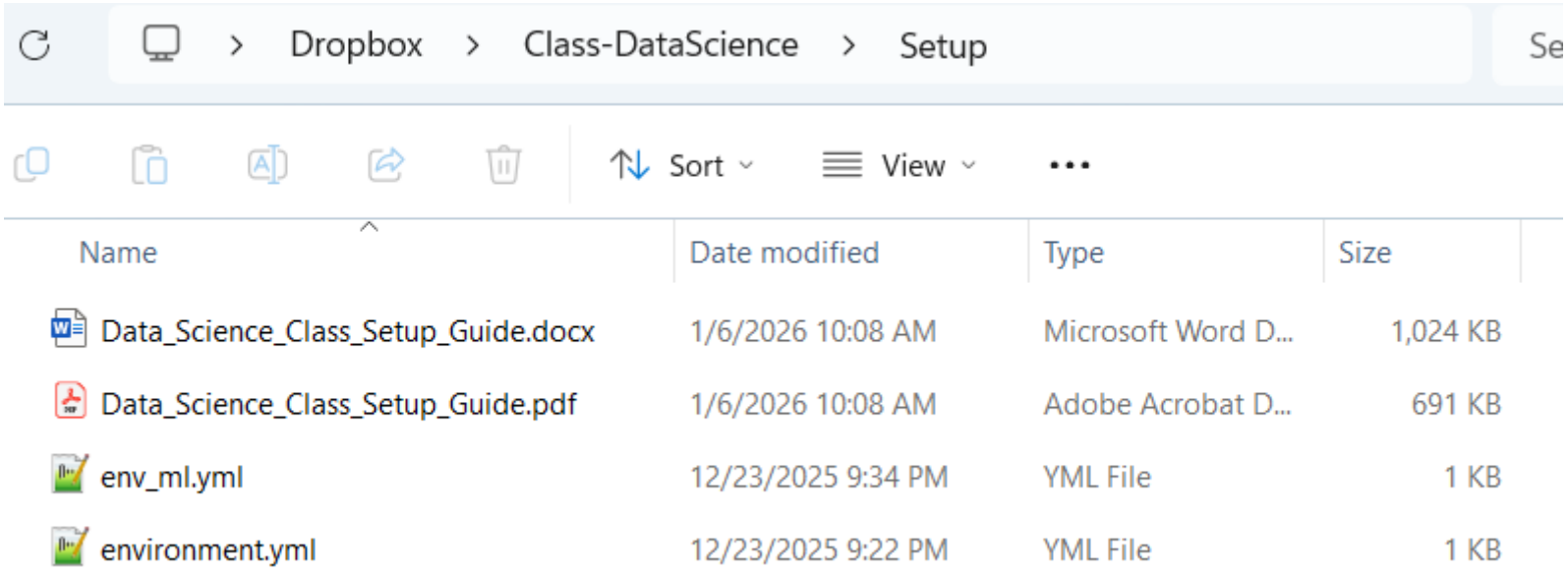


Example: directory setting code file

Setup.do

```
clear all
set more off
set trace off
cap log close
set maxvar 10000
pause off

* Set paths
if "`c(username)'"=="pierrebiscaye"{
    global home "/Users/pierrebiscaye/Dropbox"
}
if "`c(username)'"=="pibiscay"{
    global home "C:\Users\pibiscay\Dropbox"
}

    global rawghsp "$home/Data/LSMS-ISA/Nigeria"
    global w1 "$rawghsp/Wave 1 (2010-11)/Data/NGA_2010_GHSP-W1_v03_M_STATA/All One Folder"
    global w2 "$rawghsp/Wave 2 (2012-13)/Data/NGA_2012_GHSP-W2_v02_M_STATA/All One Folder"
    global w3 "$rawghsp/Wave 3 (2015-16)/Data/NGA_2015_GHSP-W3_v02_M_Stata"
    global w4 "$rawghsp/Wave 4 (2018-19)/Data/NGA_2018_GHSP-W4_v03_M_Stata12"
    global epar "$home/Data/EPAR Analysis/335 Indicator Curation/Nigeria GHS/Data"
    global harmo "$home/Data/LSMS-ISA/Harmonized/Data"
    global clean "$home/Nigeria Floods GHSP/Clean Data/GHSP"
    global code "$home/Nigeria Floods GHSP/Code"
    global tab "$home/Nigeria Floods GHSP/Output/Tables/New"
    global fig "$home/Nigeria Floods GHSP/Output/Figures/New"

include "$code/Balance_reg_fe.do"
include "$code/Balance_reg.do"

set scheme s1color // s1color, s2color, tab1
```

Example: reproducible package environment



Writing Python code

- Into Jupyter!