Frictionless Data Intro

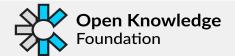


Reproducible RDM workflows TU Delft April 2022

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Hi! I'm Lilly

- Product Manager, Frictionless Data @ Open Knowledge Foundation
- Neuroscience PhD
- Love open data + open science
- **Based in Austin, TX, USA**



Frictionless Data for Reproducible Research

Removing the "friction" in research data to move from data to insight faster

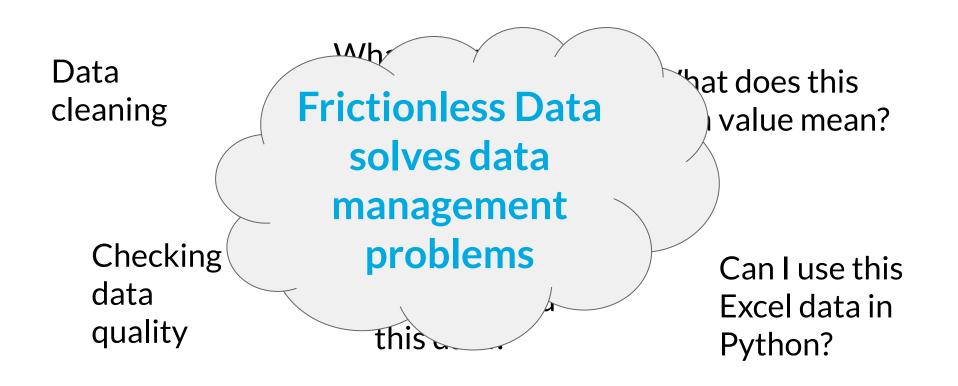


Open source & community focused:

https://github.com/frictionlessdata



What are some "frictions" in data?





Frictionless Data is open code, tools +



https://github.com/frictionlessdata

https://frictionlessdata.io/software

https://frictionlessdata.io/standards

standards

Reproducible Data Workflows lead to better science

- What is a Reproducible Data Workflow?
- Why does it matter?
- We'll see how you can use Frictionless to create a Reproducible Data Workflow!



Frictionless Python Framework has 4 main functions









Describe Data

Extract Data

Validate Data

Transform Data





Example of a data dictionary

- Data collected: the date the data were collected in YYYY-MM-DD format
- **Species**: a code for the species of the animal detected See below for a table of what the codes stand for
- Sex: the sex of the animal detected M for male and F for female
- Weight: the weight of the animal detected measured in grams

Missing data is coded as **NA**.

species code	scientific name	common name
PF	Perognathus flavus	Silky pocket mouse
ОТ	Onychomys torridus	Southern grasshopper mouse
NA	Neotoma albigula	White-throated woodrat

Is this Machine Readable?

Slide text from https://eirini-zormpa.github.io/frict ionless-data-workshop/data-orga nisation

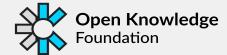
Keep your data + metadata together in a datapackage

Data e.g., experiment.csv



Metadata
+ schema
(this is like the data dictionary)

https://frictionlessdata.io/standards/



Create machine readable metadata + schemas with Describe



Data importing and exporting problems

We've known since **2016** that many genetics papers (**1 in 5**) have mistakes because Microsoft Excel interprets gene names as dates.

MICROSOFT REPORT SCIENCE

Scientists rename human genes to stop Microsoft Excel from misreading them as dates

Sometimes it's easier to rewrite genetics than update Excel

By James Vincent | Aug 6, 2020, 8:44am EDT

Also, for any **language scientists** out there working with a language that's not English, always use **UTF-8 text encoding**.

How could data validation help with this issue?



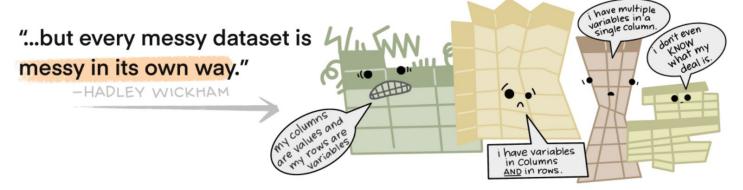
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Transform your data by cleaning it

The standard structure of tidy data means that "tidy datasets are all alike..."









Illustrations from the <u>Openscapes</u> blog <u>Tidy Data for reproducibility</u>, <u>efficiency</u>, <u>and collaboration</u> by Julia Lowndes and Allison Horst.

@lilscientista

Let's start coding!

In the terminal (Mac) or command prompt (Windows):

- Make sure you are in the same folder (directory) as the data you downloaded
- Type `jupyter notebook`
- This will open a Jupyter notebook in your internet browser
- If this does not work, raise your hand & we will help you
- As a backup, here is the code notebook:

 https://github.com/frictionlessdata/frictionless-py/blob/main/docs/tutorials/
 s/notebooks/frictionless-RDM-workflows.ipynb
- Helpers say hi!





Welcome to Day 2!









Describe Data

Extract Data

Validate Data









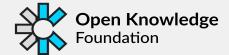
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Data e.g., experiment.csv

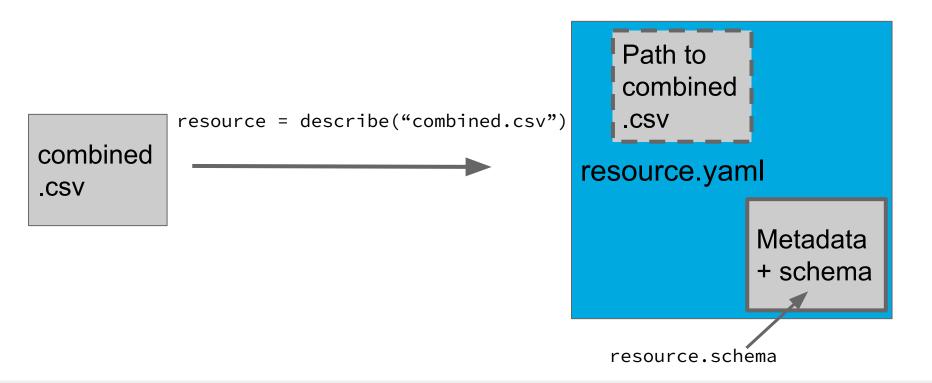


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Resource descriptor contains data + metadata/schema





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