

# Reliability and Reproducibility in Neurorehabilitation Research

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# Outline

## Reproducible Science

- What's the problem?
- What can be done?
  - Data Management
  - Data Analysis
  - Data Visualization
- (Bonus topic: Open science!)

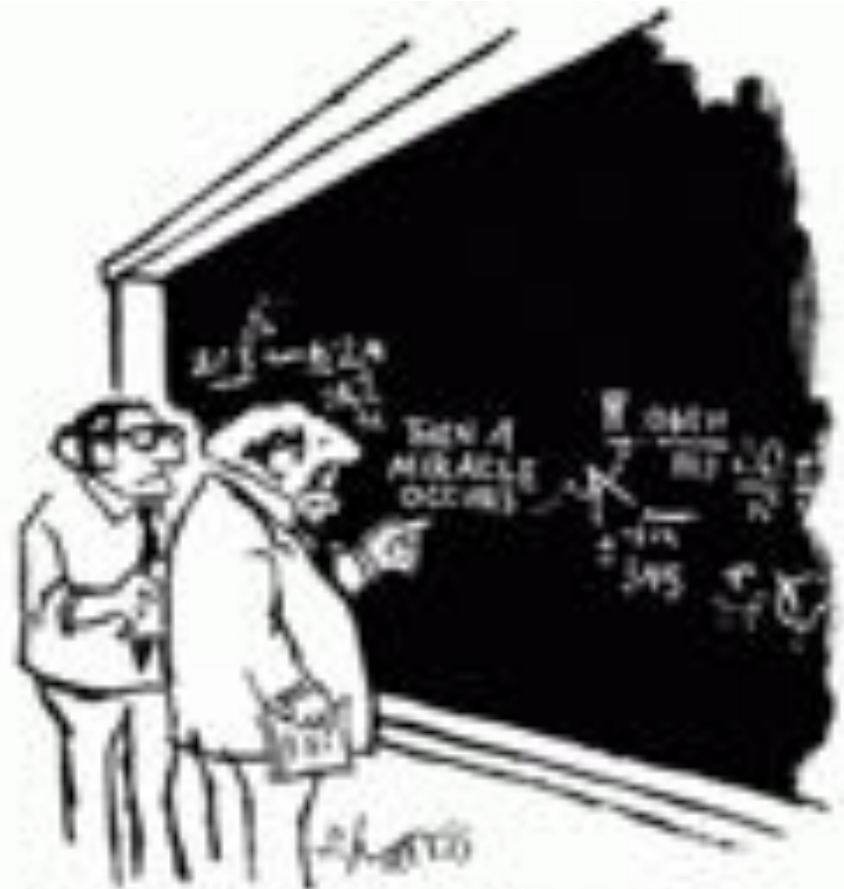
# Reproducible Science

## What is reproducibility?

The ability for someone else (or yourself) to reproduce an entire experiment and results

## What is reliability?

The degree to which the result of a measurement, *calculation*, or specification can be depended on to be accurate.



"I think you should be more explicit here in step two."

# The Reproducibility Crisis

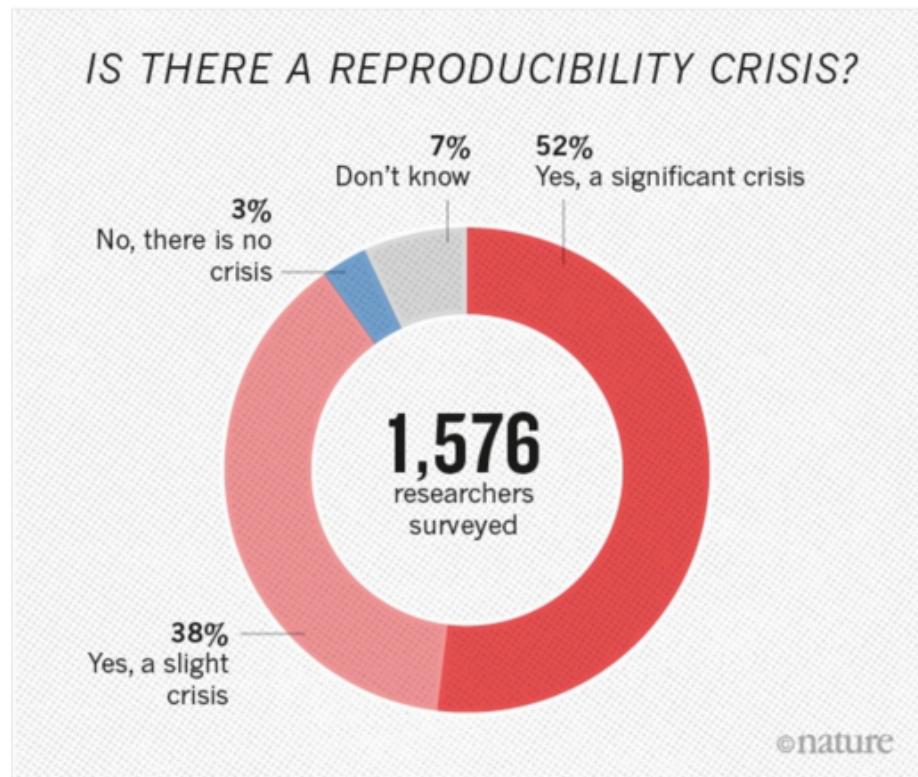
- What is the reproducibility crisis?
- Psychology – only 39 of 100 replication attempts were successful
  - <https://www.nature.com/news/over-half-of-psychology-studies-fail-reproducibility-test-1.18248>
- More than 70% of scientists have tried and failed to reproduce another scientist's experiments:
  - <https://www.nature.com/news/1-500-scientists-lift-the-lid-on-reproducibility-1.19970>

# Contributing Factors

1. Positive publication bias
2. Logistical limitations – limited money, time leading to underpowered samples
3. Heterogeneity and random variance in the sample that provides a positive effect (e.g., a few very special participants)
4. Normal human errors

# Clinical Implications of Irreproducibility

- Small studies with promising results → Large (expensive, time-consuming) studies with null results
- Wasted time/hope for patients
- Setting back science...



# What Can Be Done?

- Collecting really large heterogeneous datasets, either prospectively at one site:
  - UK Biobank
  - All of Us
  - Human Connectome Project
- Or retrospectively pooling data across sites:
  - Consortia or multi-site experiments where experiments can be replicated across sites
  - ADNI (Alzheimer's Disease Neuroimaging Initiative)
  - ABIDE
  - 1000 Functional Connectomes
  - ENIGMA

# What Can Be Done?

An experiment can only be reproduced if all the details of the experiment are properly recorded and shared. It's everyone's job!

## 1. **Data Management**

- Managing your own data
- Managing and compiling data from others

## 2. **Data Analysis**

- Simulating, checking, processing, and analyzing data

## 3. **Data Visualization**

- From quality assurance to full papers, showing individual and group results

## 4. **Reproducible Papers** (Jupyter Notebook, R Markdown, Matlab Markup)



## 4. Reproducible Papers

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- At the very least, report:
  - Actual values (means  $\pm$  SD)
  - Effect sizes (cohen's d, f, hedge's g for smaller samples)
- But better yet is sharing the raw data so people can analyze it for themselves.
- Doing so allows others to replicate your results and power their studies based on your results

# 4. Reproducible Papers

- Using a tool such as Jupyter Notebook or R Markdown, you can provide the code for analyses, as well as interactive results
- Reproducible Neuroimaging:  
<http://www.reproducibleimaging.org/>



ReproNim: A Center for Reproducible Neuroimaging Computation

Welcome to ReproNim!

The ReproNim vision is to help neuroimaging researchers to:

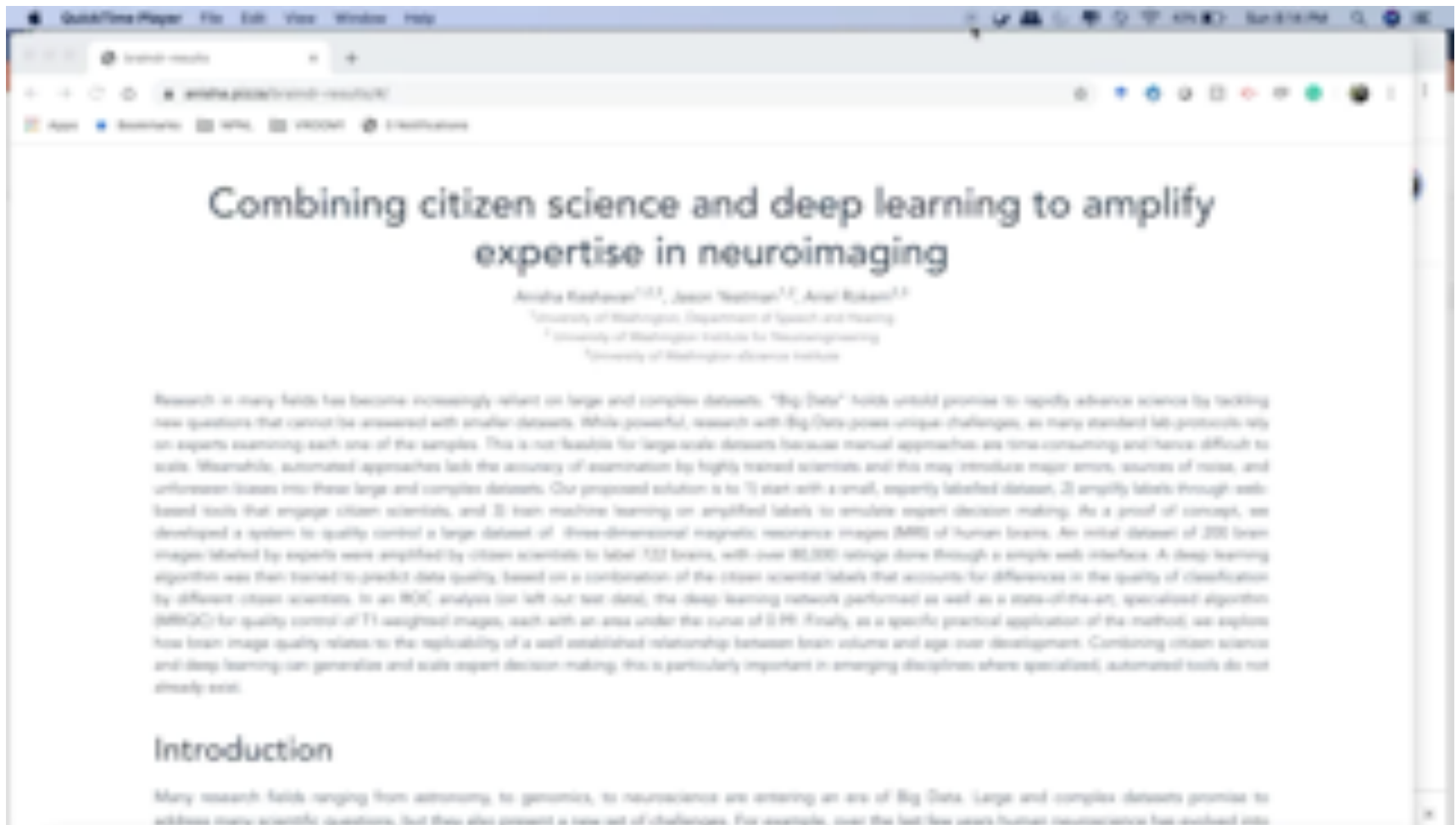
- Find and Share data in a **FAIR** fashion (discover resources with NeuroBLAST)
- Comprehensively describe their data and analysis workflows in precisely replicable fashion (describe research processes with ReproIN and BrainVerse)
- Manage their computational resource options (do analysis with NICEMAN)

so that the outcomes of neuroimaging research are more reproducible.



# 4. Reproducible Papers

Example: Keshavan et al., 2019: [braindr.results.us](https://braindr.results.us)



# A few notes on reproducibility...

- Emphasis on reproducibility is growing!
- Reproducibility awards and competitions
- Emphasis on sharing data for reproducible papers (will discuss more in open science next)
- Efforts for pre-registered reports:  
<https://www.journals.elsevier.com/cortex/news/registered-reports-a-new-article-format-from-cortex>
- And growing acceptance of null results as just as valid as positive results!

# Data Management

# Data Management

1. You should plan to keep your data in a format that can be (easily) shared with anyone at any time.
2. This means thinking about your data and analyses BEFORE you collect it. (And evaluating DURING).



# Data Management

Today we'll cover two types of data management:

1) Data that is yours:

1. Recording your experimental protocol
2. File and naming conventions
3. Recording meta-data

2) Combining data from others:

1. Database tools
2. Data processing pipelines
3. Tools for (reproducibly) checking and manipulating data

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I, too, was young once.



And I wish I had done better with data management!

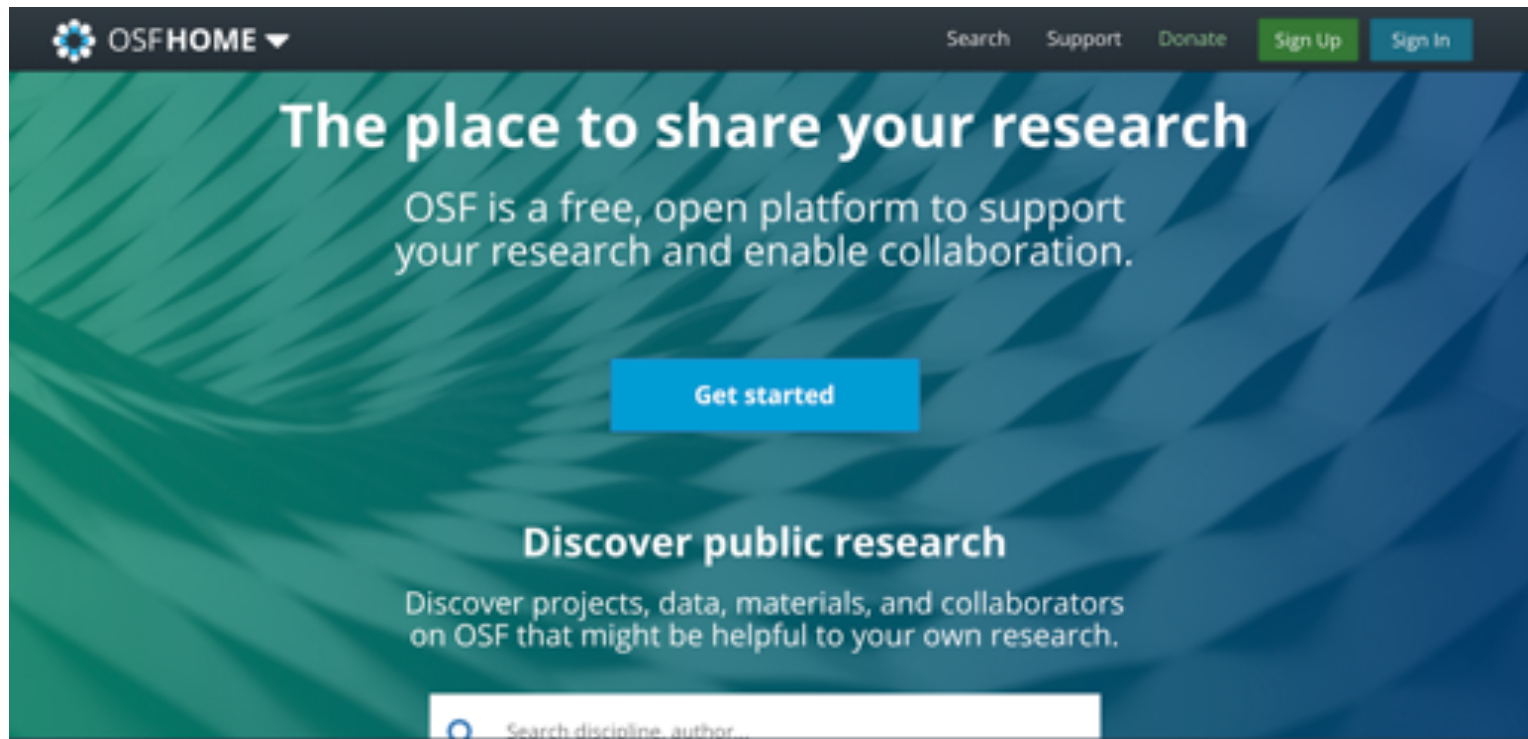
# 1. Experimental Protocols

How do you currently keep track of your experiments?

- Record a “brief” protocol with an overview of all steps, including IRB used, consent, general experimental steps, subject payment amount, etc.
- Record a “detailed” protocol with every step, including instructions to participants
- Have a second person review the detailed protocol with you to ensure they can replicate it as detailed
- Keep this somewhere accessible (e.g., lab google drive)

# 1. Experimental Protocols

Consider an organizational framework, such as a lab google drive with all the necessary components for each project and a similar format per project, or Open Science Framework (<https://osf.io/dashboard>)



# 2. Naming and File Structures

An experiment can only be reproduced if all the details of the experiment are properly recorded and shared

## Basics

- Keep all file names machine readable!
  - Alphanumeric (no spaces, slashes or symbols) – no initials
    - Consistent capitalization
    - Subj01, subj01, subj\_01, Subj\_01, SUBJ01.... – Pick one and stick with it
  - Consistent number of digits - that means anticipating how many subjects, groups, etc. (e.g., padded 0s; subj01, subj02..)
- How to write dates so it becomes logically organized: 20191016
- If you have multiple timepoints on the same day, add time in HHMMSS if needed (military format): 20191016\_133402

# 2. Naming and File Structures

An experiment can only be reproduced if all the details of the experiment are properly recorded and shared

## Basics

- Create templates of subject folders to copy and populate
- After the first subject's data is collected, analyze all data and refine file structures
- Use a consistent planned naming convention with no spaces and few symbols so it can be easily machine-sorted if needed:
  - sub-c01g01s001-t01
    - This way you can easily extract just: \*c01\*, \*g01\*, \*t01\*

# 2. Naming and File Structures

## Basics

- Consistent file structure for each subject:
  - studyFolder
    - subjFolder
      - subjData
      - subjAnalyses
      - subjResults
      - old
    - groupFolder
      - groupData
      - groupAnalyses
      - groupResults
      - old
    - writeups
      - 20180925\_draft
  - strokeStudy
    - subj01
      - data
      - analyses
      - results
      - old
    - group
      - data
      - analyses
      - results
      - old
    - writeups
      - 20180925\_draft
- For neuroimaging - see BIDS format (<https://bids.neuroimaging.io/>)

# 3. Meta-Data

- Useful to also have a “meta-data” file that describes what the various analyses are, if subgroup analyses were performed, what they were and why, etc.
  - Sex: 1=female, 2=male
  - FMUE is out of 60, not 66 points (left out reflexes)
  - 9 hole peg is normalized to other hand
- Think of this as your detailed methods section so you can revisit it 5 years later and make sense of what you did and why
- Keep track of this with the experimental protocol and in the data file – will also allow you to easily combine data with other projects or other collaborators

# Data Management

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3. *Recording meta-data*

2) Combining data from others:

1. Database tools
2. Data processing pipelines
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# ENIGMA Consortium

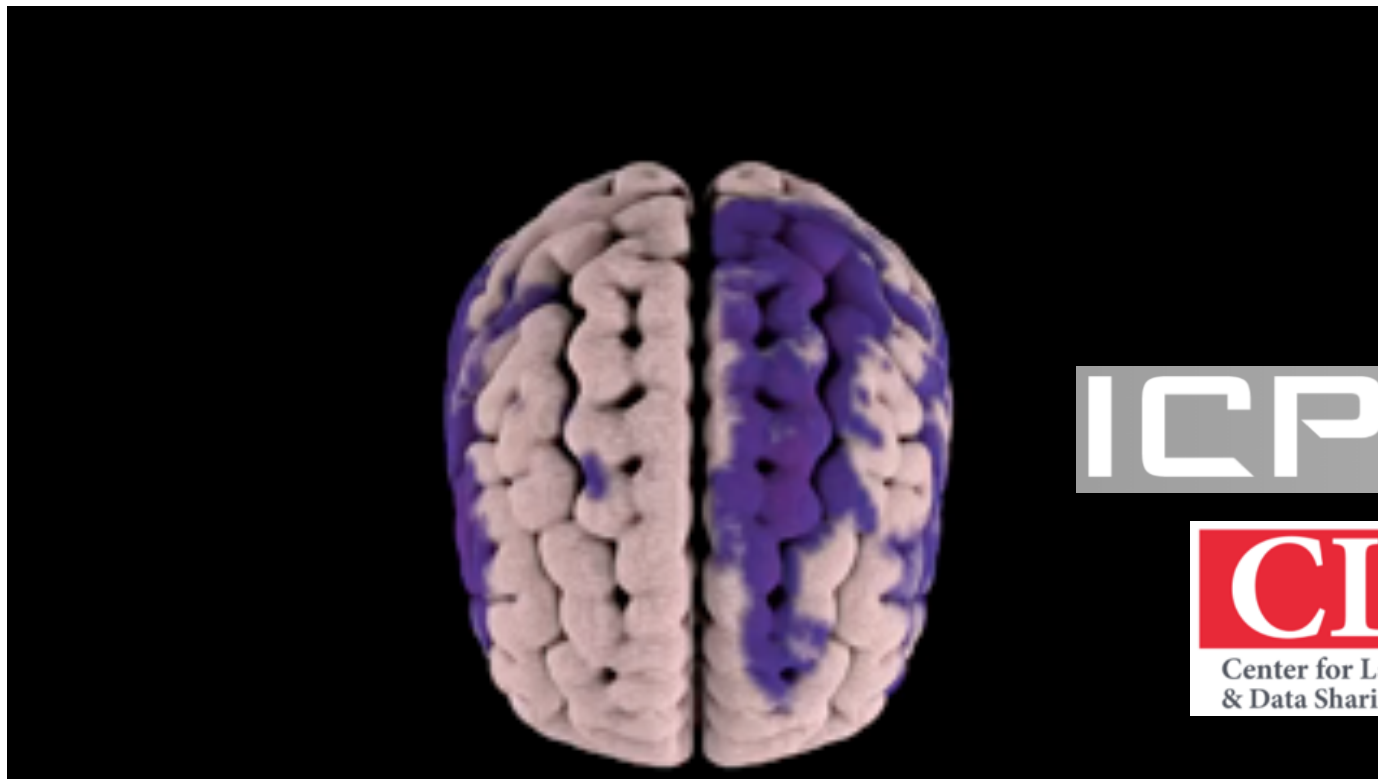
Largest-Ever Worldwide Analysis of Brain Scans and Genetic Data  
53,000+ people from over 35 countries studying 18 brain diseases



# Shared Dataset of Stroke MRIs and Lesion Masks

## **ATLAS: Anatomical Tracings of Lesions After Stroke**

Open-source dataset of 304 stroke T1-weighted MRIs with manually hand-traced lesion masks, along with neuroradiology evaluations and meta-data  
(Liew et al., 2018, *Nature Scientific Data*,  
<https://www.nature.com/articles/sdata201811>)



<https://www.youtube.com/watch?v=Ag5CUsRNY9Q>

# Data Management

## Data versus a database



da·ta·base

/ˈdɑdəˌbās,ˈdādəˌbās/

*noun*

a structured set of data held in a computer, especially one that is accessible in various ways.  
"a database covering nine million workers"

Most data is (hopefully) stored in a database

# Database Management

## 1. CSV file

- Can be read into Matlab, R, Python, etc.
- Matlab: Data structure
- R/Python: Data frame

## 2. If higher dimensional data (e.g., you have multiple CSV files (or excel sheet tabs)), consider a relational database, such as SQL (or SQLite)

- Tables for subject demographics, behavioral measures, brain imaging measure, etc.
- Fast queries, non-redundant fields
- For many columns, it's a little more human readable

# Database Management

Common data elements: What are we going to record, and how are we going to record it?

- What are we going to record? Requires consensus
  - Stroke Recovery and Rehabilitation Roundtable (SRRR)
- How are we going to record it? Requires forethought
  - Time since stroke (days, months, years...)
  - Units (metric, not-metric)
  - Default to the highest resolution where possible

# Database Steps

Key aspects of database management (when receiving data from others)

1. Data schema (organization)
2. Data scrubbing
3. Data scrubbing
4. Data re-organization and data scrubbing
- ...
99. Data analysis

# Database Management

## 1. New Site Data

- MRIs (nifti preferred, dicoms ok)
- CSV with demographics, behavior, scanner parameters

## 2. We Scrub Data

- Ensure subIDs of MRIs/csv match
- BIDSify MRI data and add to `../enigma/new/BIDS/site` folder
- Add BIDS SUBJECT\_ID and SESSION\_ID to csv (see `../scripts/makeBIDS`)
- Rename csv columns to match sqlite database columns (or add new)
- Place original csv in `../BIDS/site` e.g., `R033/sourcedata` as `site_behavior.csv`
- Place formatted csv (with BIDS Ids, renamed columns) in same sourcedata folder as `site_behavior_renamed.csv`

## 3. Run Freesurfer 5.3 and 6.0

- Make new folder in freesurfer `_v*_interim` folder
- In there, make swarm commands:
  - `cd ../enigma/new/scripts/freesurfer/`
  - `MakefreesurferCommands.py` (replace with the version of freesurfer and location for data to go)
  - See `readme.txt` and run resulting `freesurferCommands.txt` file with `Swarm.py` script
- Send outputs to folder in interim folder

## 4. Extract Brain CSV Values and QC

- Cd appropriate interim folder
- Cp `../enigma/new/scripts/ENIGMA_Wrapper_scripts-master/modified`
- Run `2_extractsubcortical_volumes.sh`
- Run `3_extractcortical_volumes.sh`
- Copy master csvs and rename `'_QC'`
- Qc Data – `Landoltvolumes_QC.csv`

## 5. Add New Brain Data To Existing Data

- Copy freesurfer folders (if not directed to `../new/freesurfer_v53` or `../new/freesurfer_v6` already)
- Cd `../scripts/freesurfer/` and run `CopyFolders.sh` (change source)
- Add new brain csvs and QC csvs to masters  
Cd `../scripts/outputcsvs_append` and run `python2 outputcsvs_append.py`  
Make sure to change the output directory to directory with newly generated output csvs)
- Perform lesion analysis (see sheet 2)

## 6. Rerun scripts to generate enigma.db

- Rerun scripts to create updated `enigma.db` sqlite database – include date in name or put in dated folder to keep track of all versions

# Hands-On Database Tools

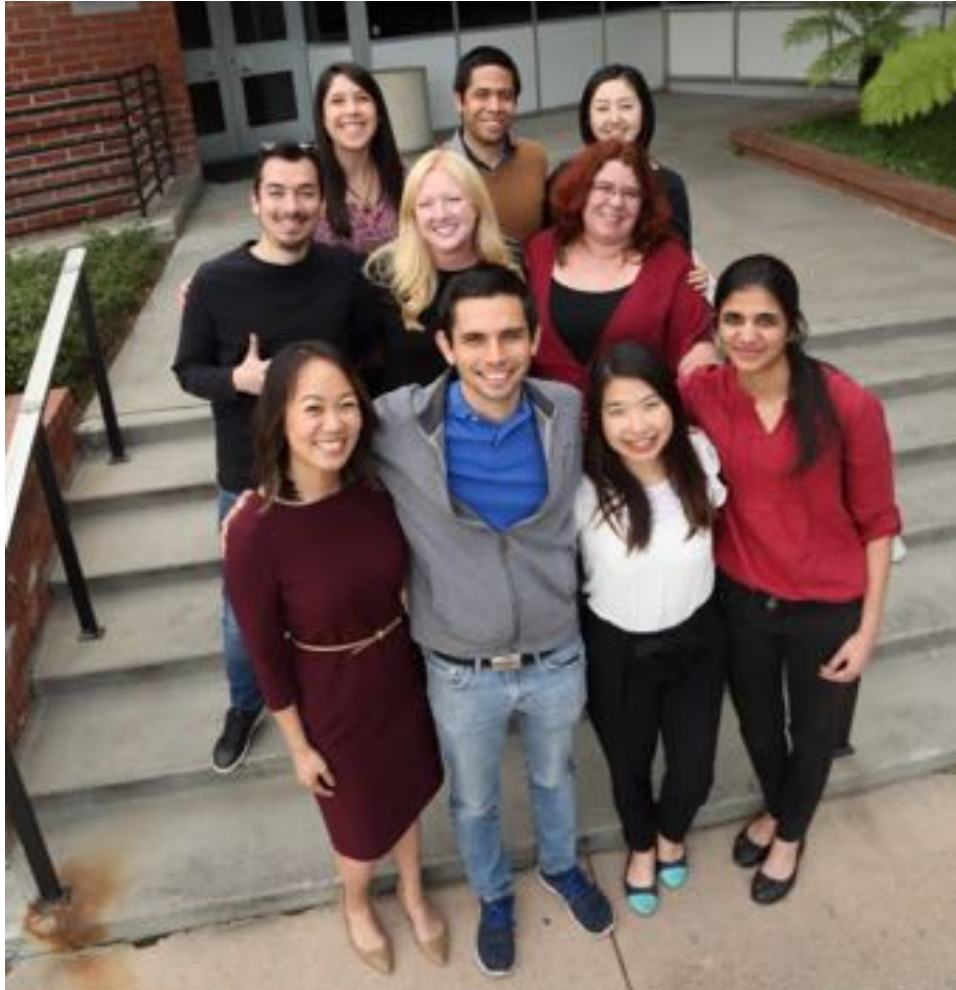
- SQLite (walk-through demo)
- Python with PANDAS to manipulate csv files in jupyter notebook (hands on)



# Thank you!

The Neural Plasticity and Neurorehabilitation Laboratory

<http://npnl.usc.edu>



Email me:

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Twitter us:

@NPNLatUSC





# Bonus Topic!

- Open Science
  - What is it?
  - Why share?
  - Resources

# What is Open Science?

- **Open science** is the movement to make scientific research, data and dissemination accessible to all levels of an inquiring society, amateur or professional.
- Encompasses:
  - Open data – shared data
  - Open source code – shared code and analyses
  - Open methods – shared experimental paradigms
  - Open peer review – comments and responses posted
  - Open access – access to journal articles
  - Open educational resources – online tutorials, etc.

# Why Open Science?

- **Why not share:**
  - Someone else could steal my ideas and publish them before me!
  - I worked hard to collect this data!
  - I worked hard to create this code!

But... open science doesn't mean you give everything up and get nothing in return!

# Why Open Science?

- **Why share:**
  - You can publish a paper on your data or code and get cited for creating these valuable resources
  - Someone else could find new and exciting findings from the data you collected, or build on your ideas, or make use of the code you worked so hard on
  - You can also find new collaborations to keep building on your current work
  - You are contributing not only to your own scientific career but to science in general!

# What Can I Share?

## **Open data – shared data**

- Nature's Scientific Data, GigaScience Data Reports, and others are highly cited
- Many journals now request data shared with your paper submission (hence the repro steps before)
- Many grants now mandate that you share your data on a site, such as github, figshare, etc.
- There are many repositories for archiving data – e.g., ICPSR, FCP/INDI
- See: <https://www.nature.com/sdata/policies/repositories>

# What Can I Share?

## **Open source code – shared code and analyses**

- Github!
  - This allows others to report bugs and recommend enhancements
  - Also allows you to track users of your data
  - Also allows others to contribute to your code and analyses – to make it better
    - The open science community is usually friendly and might recommend a pull to help you out!
    - You can even hire someone via bitcoin bids to help you with your code :D
- <https://guides.github.com/activities/hello-world/>
- <https://product.hubspot.com/blog/git-and-github-tutorial-for-beginners>



# What Can I Share?

Open methods – shared experimental paradigms

- If code, also can be done via Github!
- Provide your exact paradigm so others can try to replicate it in different contexts
- Provide your tools, substrates, etc. so others can try it at their sites (→ usually involves some data or material transfers via your local tech transfer office)

# What Can I Share?

Open peer review – comments and responses posted

- Preprint archives such as bioRxiv:  
<https://www.biorxiv.org>
- Allows for comments from the community, twitter and views metrics (for cover letters), and uploads of revised versions; will linked to published version later
- Possible to get open peer review and point editors to this (but still beta)
  - <https://www.biorxiv.org/content/10.1101/441451v1?versioned=true>

# What Can I Share?

Open access and open educational resources

- Consider publishing in open access journals
- Consider making your tutorials videotaped and shared on youtube
- I just learned that Quicktime lets you record your screen plus audio easily :D  
<https://support.apple.com/guide/quicktime-player/record-your-screen-qtp97b08e666/mac>
- Watch out for this as it will grow in the future!

# A few other notes...

- Some people may still try to misuse your data or be entitled, so it's important to have adequate protections over it (e.g., an education license for any code shared on github)
- You don't actually own your work (your university does) so you do need to disclose any work you create
- Open science is growing and always changing
  - Exciting to get into it now because it's likely to become mandated soon :D
  - It's a great community full of supportive people 😊

# Some Resources

- <http://www.reproducibleimaging.org/module-reproducible-basics/>
- <https://software-carpentry.org/>
- <https://datacarpentry.org/>