

# Welcome to coreR

# NCEAS Learning Hub for Delta Stewardship Council

June 2024

#### Week's Schedule

	RM 2-310		RM 2-309		RM 2-310		RM 2-310		
Session Time	<u>Monday</u>		<u>Tuesday</u>		<u>Wednesday</u>		<u>Thursday</u>		<u>Friday</u>
8:30-10:00	Introduction	Camila	Cleaning & Wrangling Data	Angel	Publishing to the Web	Angel	Shiny cont'	Camila	
	Set up				Intro to Data Viz				
10:00-10:30	BREAK		BREAK		BREAK		BREAK		
10:30-12:00	Literate Analysis with Quarto	Camila	Practice Session I		Working with Spatial Data	Angel	Wrap-up: Reproducibility & Provenance	Camila	
							Survey + Q&A		
12:00-1:00	LUNCH		LUNCH		LUNCH		ADJOURN		
1:00-2:30	Introduction to Git & GitHub	Angel	Collaborationg with Git & GitHub	Angel	Practice Session II		Technical		
							Non-Technical		
2:30-3:00	BREAK		BREAK		BREAK		Practice		
3:00-4:30	Tidy Data	Camila	Data Management	Camila	Intro to Shiny	Camila			

# An immersion course in **R programming for environmental data** science.

You will gain experience on how to leverage the use of data science tools to increase your capacity to **collaborate** with your team, create **reproducible** <u>workflows</u>, and learn **best practices for open science**.

#### About this course





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# **Environmental data science:**



Artwork: @allison horst

And building robust workflows.

U.S National Science Foundation (NSF) subcommittee on replicability in science: "reproducibility refers to the **ability of a researcher to duplicate the results of a prior study using the same materials as were used by the original investigator**"

Goodman et al 2016

#### Types of reproducibility

- "Computational reproducibility: When detailed information is provided about code, software, hardware and implementation details."
- "Empirical reproducibility: when detailed information is provided about non-computational empirical scientific experiments and observations. In practice, this enabled by making data freely available as well as details of how data was collected."
- **"Statistical reproducibility:** when detailed information is provided about the choice of statistical tests, model parameters, threshold, values etc. This mostly related to pre-registration of study design to prevent p-values hacking and manipulations."

Victoria Stodden, 2014, <u>ROpenSci Reproducibility Guide</u>

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#### Does this look familiar?

#### Is this $\downarrow$ how you've been working with data? Cool! If it has been working for you, feel good about it.



Allison Horst, EDS 221, Scientific Programming Essentials

#### A common workflow



Paste COPY Re-copy / Re-paste Paste LOON Paste CODV Copy / Paste Re-copy / Re-paste Copy / Paste Paste

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Allison Horst, EDS 221, Scientific Programming Essentials

#### A common workflow



Allison Horst, EDS 221, Scientific Programming Essentials

#### **Room for improvement**

- No history of what has been done to the data. From raw data to final figures/results.
- Lack of documentation on the step by step process.
- What happens if data is updated? We need to repeat every process?
- How do we collaborate with colleagues? Back and forth emails and versions of files with inevitably long file names (final\_report\_v1\_CVP\_AC\_review\_new\_this\_one.docx)
- How do we transfer analysis to final reports? Is this reproducible?

#### Work with your data like it's going to need an alibi



Allison Horst

Do everything in **well-annotated and organized scripts** that contain streamlined and easy-to-follow records of your entire analysis from **raw data** through **final reports** with **unbreakable file paths** and **complete history** of changes made.

Adapted from Allison Horst, EDS 221, Scientific Programming Essentials



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- Hadley Wickham (from <u>fivebooks.com interview</u>)



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#### Reproducibility starting point: Set up a robust structure

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Artwork: @allison horst

#### Reproducibility starting point: Set up a robust structure

- The fundamental idea behind a reproducible analysis is a **clean**, **repeatable script-based workflow**.
- This will allow you to **re-run your analysis as many times** as needed before (and after) the completion of your project.
- The smoother and more **automated the workflow**, the easier, faster and more robust the process of repeating it will be.



1. Use a scripted (programming) language

- 2. Use one folder per project
- 3. Organize the content of your project with sub-folders
- 4. Set up robust file paths

# Talk to your neighbor

- How do you generally organize your files for a project?
- What do you like about your system?
- Do you see any limitations to your system?

# 1. Use a scripted (programming) language



#### How do you tell your code where to find files?

Artwork by Allison Horst

How do you tell your code where to find files?

some\_data <- read.csv("/home/vargaspoulsen/Documentes/Workshops/RLadies-SB/reproducibleworkflows/some\_data.csv") How do you tell your code where to find files?

some\_data <- read.csv("/home/vargaspoulsen/Documentes/Workshops/RLadies-SB/reproducibleworkflows/some\_data.csv")

If I share my script with this file path to my colleagues, would they be able to open the file?

Probably not.





• Provides a **self contained working directory** (folder) that does not depend on the absolute location of your computer.



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• Bundles all your work within a working directory, pointing to relative locations within the project.

• Within this centralize location we can organize all the files involved in our project (inputs data, scripts, outputs, etc.)





When you create an *R Project*, it **creates and Rproj file and a folder in your computer that will be the working** *directory* when your are working in your Rproj.



• An absolute path always starts with the root of your file system and locates files from there.

/home/vargas-poulsen/Documents/Workshops/RLadies-SB/reproducible-workflows/data/data.csv

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\*\*R projects set the file path relative to the project's directory (folder)\*\*



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#### Let's take a look at one example









# Analysis related scripts

















## Example of project organization



3. Organize the content of your project with sub-folders

#### **General organization recommendations**

- Keep your **Raw data Raw** Never edit your raw data.
- Clearly **separate raw** data **from "clean"** processed data.
- **Review external inputs** to make sure they align with the established organization structure.
- Define informative **file naming conventions**.



- Different operating systems use different characters to define file paths.
  - Mac and Linux uses <u>slashes</u> (e.g. plots/diamonds.pdf)
  - Windows uses <u>backslashes</u> (e.g. \plots.pdf).
    - ~ is a convenient shortcut to your home directory on mac
  - Windows doesn't really have the notion of a home directory, so it instead points to your documents directory.

R for Data Science (Grolemund & Wickham)

#### How to make your file paths within your project robust?



Artwork by <u>Allison Horst</u>

"The goal of the here package is to enable easy file referencing in project-oriented workflows. In contrast to using setwd(), which is fragile and dependent on the way you organize your files, here uses the top-level directory of a project to easily build paths to files."

here.r Documentation

It allows us to **navigate through the files in our project** without having to worry about operating system issues. here() starts from the working directory, aka your Rproj folder.



If I'm working within my R project, to read
some\_data.csv, inside the raw folder in this case, I
can use the here::here() function.

#### some\_data ← read\_csv(here::here("data", "raw", "some\_data.csv"))

#### R Projects + here() = robust file paths

#### BIG first step towards reproducible workflows!



Artwork by Allison Horst

# **Organization Wrap up**

# One of the first steps to achieve reproducibility is to **set up a <u>robust structure</u>** for our work.







Scripted analysis

One folder with organized content

Robust file paths

Artwork by Allison Horst

# **Organization Wrap up**



reproducible\_project

reproducible\_project

Self contained project with all file paths relative to folders within the project, analysis can be reproduced elsewhere



- Best Practices for Writing Reproducible Code, University of Utrecht
- <u>A Guide to Reproducible Code in Ecology and Evolution, British Ecological</u> <u>Society</u>
- <u>Reproducibility and Provenance, NCEAS Learning Hub</u>
- Workflows, LTER Scientific Computing Workshops
- <u>Reproducibility Lesson, LTER Synthesis Skills for Early Career Researchers</u>
- EDS 221, Lesson 1 and Lesson 2, UCSB MEDS, By Allison Horst
- <u>GitHub Clinic, Openscapes</u>
- Building reproducible analytical pipelines with R, Bruno Rodrigues