



R-Ladies Lausanne

ThundeR Talks - July 2, 2019

Ninon Mounier

WHAT'S THAT ALL ABOUT?

WORKFLOW





workflowr: *organized + reproducible + shareable data science in R*

- **Organized**

Provides a project template with organized subdirectories

Mixes code and results with R Markdown

Uses Git to version both source code and results

- **Reproducible**

Displays the code version used to create each result

Runs each analysis in an isolated R session

Records the session information of each analysis

Sets the same seed for random number generation for each analysis

- **Shareable**

Creates a website to present your research results

Documents how to host your website for free via [GitHub Pages](#) or [GitLab Pages](#)

Creates links to past versions of results

◦

source: <https://github.com/jdblischak/workflowr>





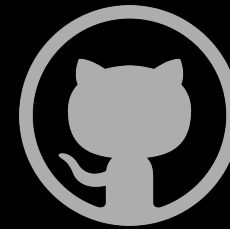
Let's take a leap in the dark together...

Getting started...

- **Define git parameters**
- **Create a new project**
- **Understand project template/structure/management**
 - `wflow_build()`
 - `wflow_status()`
 - `wflow_view()`
 - `wflow_git_commit()`
 - `wflow_publish()`

Getting started...

- **Deploy the website (github)**



- `wflow_use_github()`
- `wflow_push()`

- **Customise the website**

- → use Bootstrap themes or custom CSS files

Use option `dry_run = TRUE` to preview what a function will do

My opinion

Worth a try!

- ❑ How to integrate workflow and HPC?

 - ➔ use the data/ code/ output/ folders

 - check if output files already created, if not use slurm?

- ❑ Github structure:

 - Projects as subdirectories?

 - Private repositories?

See: [Migrating an existing project to use workflowr](#)

