



github.com/  
JuliaDynamics



# DrWatson

The perfect sidekick to your scientific inquiries

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latest version:  
v0.3.0

## Motivation

- ▶ I am tired of typing `savename = "w=2_f=2.5_x=5.dat"` can't I do it automatically?
- ▶ I moved my folders and my load commands don't work anymore!
- ▶ Do I have to produce a dataframe of my simulations AGAIN?!
- ▶ Yeah you've sent me your project but none of the scripts work...

Sounds familiar...? **DrWatson** tries to eradicate such bad thoughts and bedtime nightmares!

## Functionality

DrWatson is a scientific project assistant software

### Project Setup

- Well-designed project structure
- Robust navigation of project
- Works no matter where it is on disk
- Fully reproducible project

### Naming Simulations

- Deterministic scheme for creating names given *any* parameter collection
- Fully customizable but also with sensible defaults

### Saving Tools

- Automatically add Git commit and source file information to saved data
- Safe-saving over existing files
- Produce or load existing files flexibly

### Running & Listing Simulations

- Preparing multiple simulation runs
- Automatically creating a parameter table (dataframe) of completed runs
- Support for serial & parallel clusters

## Principles

### non-invasive

no strict rules to follow, no command line, no changing the way you work

### simple

easy to understand and use, function based (just call a function)

### reproducible

reproducibility of any result (versioning: code and dependencies)

### consistent

functionality identical across all projects

### modular

flexible design, use only what you need in your project

### incremental

you can add more, completely new, simulations/data

## Navigating a reproducible project

- ▶ Call the function `initialize_project(folder, name)`
- ▶ Always find the e.g. data folder by calling `datadir()`, no matter where the calling script is located
- ▶ Load source: `include(srcdir()*"unitcells.jl")`
- ▶ Automatically add (current) Git commit ID to some data to save: `tagsave(datadir()*"exponents.bson", data)`
- ▶ Send your colleague the `folder` and they have to do:  
`Pkg.activate("path/to/folder")`  
`Pkg.instantiate()`  
It is now **guaranteed** that all code that run for you, runs for them!

## Getting a name out of a container

- ▶ Define a (named) parameter container like so:  
`c = (T = 100, dt = 0.1, N = 25, mode = "bi")`
- ▶ Then calling `savename(c)` gives  
`"N=25_T=100_dt=0.1_mode=bi"`
- ▶ Doing instead `savename(datadir(), c, "dat")` gives  
`"path/to/data/N=25_T=100_dt=0.1_mode=bi.dat"`
- ▶ This works for **any** named container, including **any Julia struct**
- ▶ Sensibly excludes non-fitting fields (e.g. vector-valued fields) but can be customized to full extent

## Producing a dataframe from saved files

- ▶ You have run some simulations with parameters  $a = 1$  or  $2$ ,  $b = "x"$  or  $"y"$  and  $c = 0.3$ . You save them in directory "data"
- ▶ `df1 = collect_results!("results.bson", "data/")`

4x3 DataFrame			
Row	a Int64?	b String?	c Float64?
1	1	x	0.3
2	1	y	0.3
3	2	x	0.3
4	2	y	0.3

- ▶ You now run a new simulation with  $a=4$ ,  $b="z"$ , a **new parameter**  $d=0.5$  or  $0.6$  and the parameter  $c$  **does not exist** anymore!
- ▶ `df2 = collect_results!("results.bson", "data/")`

6x4 DataFrame				
Row	a Any	b Any	c Any	d Float64?
1	1	x	0.3	missing
2	1	y	0.3	missing
3	2	x	0.3	missing
4	2	y	0.3	missing
5	4	z	missing	0.5
6	4	z	missing	0.6

wanted

beta  
testers

more  
contributors

feature  
requests

<https://github.com/JuliaDynamics/DrWatson.jl>

scan me!

