Automating your R-based research project

Introduction to GitHub Actions

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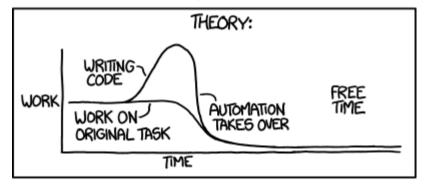
2022-03-13

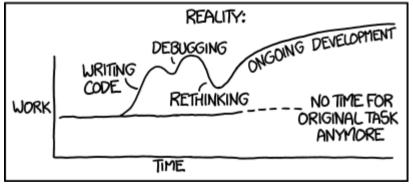
Outline

- Overview of automation
- Introduction to GitHub Actions
- Demonstration of automation

Automation

"I SPEND A LOT OF TIME ON THIS TASK.
I SHOULD WRITE A PROGRAM AUTOMATING IT!"





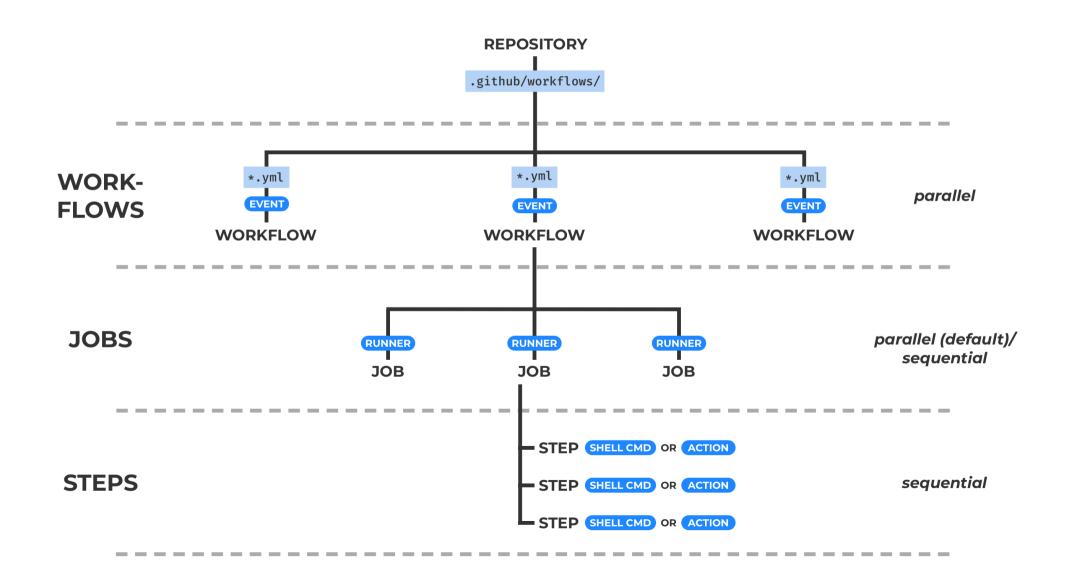
'Automating' comes from the roots 'auto-' meaning 'self-', and 'mating', meaning 'screwing'

What can be automated?

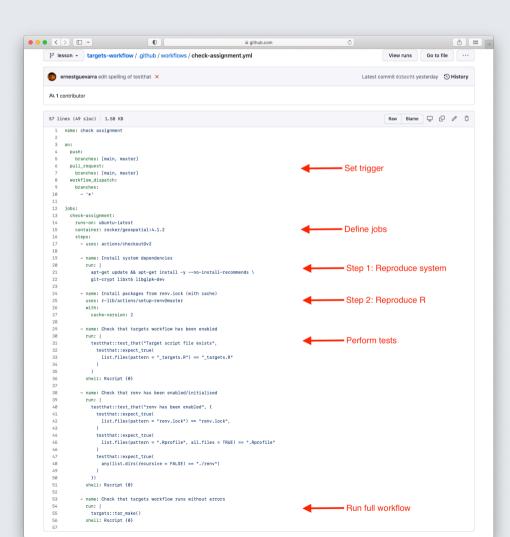
- works best with a project workflow (sometimes called a project pipeline)
 - our recommended solution for this is the {targets} workflow/pipeline
- works best if project workflow is portable
 - our recommended solution for this is {renv}

Why automate?

- to reproduce previous results
- to recreate results deleted by fat fingers
- to rerun the workflow/pipeline with updated software
- to run the same workflow/pipeline on a new dataset



Example GitHub Actions automation



- Example automation built-in to the latest assignment on creating targets workflow
- Can be found at .github/workflows/checkassignment.yml

Setting the trigger for the automation

```
name: check assignment

on:
    push:
        branches: [main, master]
    pull_request:
        branches: [main, master]
    workflow_dispatch:
        branches:
        - '*'
```

Defining jobs to be implemented in the workflow

```
jobs:
   check-assignment:
    runs-on: ubuntu-latest
   container: rocker/geospatial:4.1.2
   steps:
    - uses: actions/checkout@v2
```

Specifying steps in the workflow

Workflow step 1: reproduce the system environment

```
    name: Install system dependencies
        run: |
            apt-get update && apt-get install -y --no-install-recommends \
            git-crypt libxt6 libglpk-dev
```

Workflow step 2: Reproduce the R project-local environment

```
- name: Install packages from renv.lock (with cache)
uses: r-lib/actions/setup-renv@master
with:
   cache-version: 2
```

Specifying steps in the workflow

Workflow step 3: Perform tests/checks

```
- name: Check that targets workflow has been enabled
 run:
   testthat::test that("Target script file exists",
      testthat::expect true(
        list.files(pattern = "_targets.R") == "_targets.R"
 shell: Rscript {0}
- name: Check that renv has been enabled/initialised
 run:
   testthat::test_that("renv has been enabled", {
      testthat::expect true(
        list.files(pattern = "renv.lock") == "renv.lock",
      testthat::expect true(
        list.files(pattern = ".Rprofile", all.files = TRUE) == ".Rprofile"
      testthat::expect true(
        any(list.dirs(recursive = FALSE) == "./renv")
```

Specifying steps in the workflow

Workflow step 4: Run the targets workflow

```
- name: Check that targets workflow runs without errors
run: |
   targets::tar_make()
shell: Rscript {0}
```

Questions?

Thank you!

Slides can be viewed at https://oxford-ihtm.io/open-reproducible-science/session11.html

PDF version of slides can be downloaded at https://oxford-ihtm.io/open-reproducible-science/pdf/session11-automating-r-projects.pdf

R scripts for slides available here