Reproducible Tools and Workflows

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Tools we'll see this week

R, RStudio

- https://cran.r-project.org/
- https://www.rstudio.com/
- make (and other command line tools)
 - For Mac/Linus: pre-installed
 - For Windows:

https://cran.r-project.org/bin/windows/Rtools/

🛛 git

- git (https://git-scm.com/)
- github (https://github.com/)
- gitkraken (https://www.gitkraken.com/)
- any text editor
 - any command line terminal

Introductions

Me:

Thomas Political Scientist, Methodology Department R

Vou:

- Name
- Field/Department
- Tools/Software

Learning Objectives

- Understand how to organize a reproducible research project
- 2 Recognize different approaches to reproducibility and tools for implementing various reproducible workflows
- 3 Th: Apply various workflows to your own work
- Th: Understand how to collaborate reproducibly

1 Organizing Things

2 Building Things

3 Keeping and Changing Things

4 Thursday: Hands-On

1 Organizing Things

2 Building Things

3 Keeping and Changing Things

4 Thursday: Hands-On

Activity!

How do you organize your files for a project?

Wait, but why do we care?

If we're going to be transparent *in the end* (e.g., at verification or data archiving stage), what do we need to provide?

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A well-organized, reproducible analysis!

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A well-organized, reproducible analysis!

So rather than make that an annoying, post-hoc exercise related to publication, try to get organized and stay organized throughout your project from the very beginning.





"Reproducibility is collaboration with people you don't know, incl. yourself next week." – @philipbstark #openscience

4 t3 ***** …

The single most important part of reproducibility is naming things!







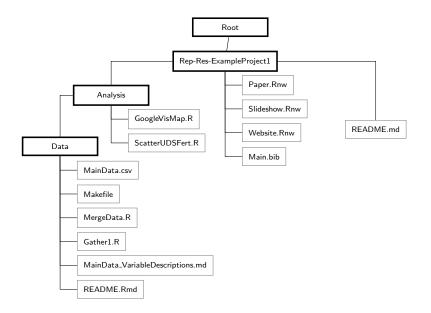




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📝 Dissertation - Prospectus 0.doc	2015-07-15 09:17	Microsoft Word 9	18 KB
Dissertation - Prospectus 1.doc	2015-07-15 09:18	Microsoft Word 9	142 KB
💼 Dissertation - Prospectus 2.doc	2015-07-15 09:18	Microsoft Word 9	246 KB
💼 Dissertation - Prospectus 3.doc	2015-07-15 09:18	Microsoft Word 9	250 KB
Dissertation - Prospectus 4.doc	2015-07-15 09:19	Microsoft Word 9	250 KB
Dissertation - Prospectus 5.doc	2015-07-15 09:19	Microsoft Word 9	263 KB
🖬 Dissertation - Prospectus 6.doc	2015-07-15 09:19	Microsoft Word 9	287 KB
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🚾 Dissertation - Prospectus 9 (For Jamie).pdf	2015-07-15 09:19	PDF File	328 KB
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🖬 Dissertation - Prospectus 10 (Big Question).doc	2015-07-15 09:18	Microsoft Word 9	19 KB
🖬 Dissertation - Prospectus 10 (New) hml.doc	2015-07-15 09:18	Microsoft Word 9	179 KB
🖬 Dissertation - Prospectus 10 (New).doc	2015-07-15 09:18	Microsoft Word 9	280 KB
🖬 Dissertation - Prospectus 10a (Big Question).doc	2015-07-15 09:18	Microsoft Word 9	19 KB
🖬 Dissertation - Prospectus 10b (Big Question).doc	2015-07-15 09:18	Microsoft Word 9	32 KB
📷 Dissertation - Prospectus 10c (Big Question).doc	2015-07-15 09:18	Microsoft Word 9	34 KB
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🖬 Dissertation - Prospectus 13 (Outline).doc	2015-07-15 09:18	Microsoft Word 9	159 KB
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🖬 Dissertation - Prospectus 13b (Outline).doc	2015-07-15 09:18	Microsoft Word 9	266 KB
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What makes up the ideal reproducible research product?

- Gandrud's template
- rOpenSci's "Research Compendium"
- Project TIER
- AJPS Replication/Verification Policy



project

- |- DESCRIPTION # project metadata and dependencies
- README.md
- # top-level description of content
- |- data/ # raw data, not changed once created | +- my_data.csv # data files in open formats
- |- analysis/ # any programmatic code | +- my scripts.R # R code used to analyse data

Original Data Original data files Importable data files (if necessary) Metadata The Metadata Guide Supplementary metadata documents	Documents The final paper The Data Appendix The Read Me file
(if necessary) Analysis Data Analysis data files	Command Files

Don't be this guy:

🕌 Old Material	🔁 APSA2011 Handouts.pdf	APSA2011.aux	APSA2011.log
APSA2011.nav	APSA2011.out	1 APSA2011.pdf	APSA2011.snm
APSA2011.synctex.gz	APSA2011.tex	APSA2011.toc	APSR Reviews.txt
AQMW2011 Handouts.pdf	AQMW2011.aux	AQMW2011.log	AQMW2011.nav
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📭 hovland.png	🔁 imp-noinfo.pdf	🔁 imp-repeat.pdf	🔁 imp-search.pdf
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Tables 2011-12-28.docx	Tables.doc	Tables.xlsx	

mkdir code

mkdir data

mkdir figures

echo # My Project > README.md

Everything you do should be plain text*

Everything you do should be plain text*

* Exceptions to this are images (sometimes)

- Plain text is always compatible Every single operating system has a plain text editor and they are all
 compatible up to the encoding of the text. This means that if you develop your lecture notes on a Mac and I
 develop them on a PC then we can still easily share no worrying if you have the right software.
- Plain text is easy to mix and match If your lecture materials are in a simple plain text format like
 markdown you can copy and paste the materials from one lecture into another and when the document is
 compiled make all the formatting/colors/etc. match. No more looking at hodgepodges of borrowed slides
 some with one ppt format and some with another.
- Plain text is easy to maintain We mostly work on scalable education here at the JHU Data Science Lab.
 We often think of scalable education in terms of the number of students, but here we have also run into the problem of scaling the number of courses/instructor. I am currently the lead instructor on more than half a dozen classes running all the time. Every time I have to re-record a video it takes set up time, recording time, editing time. If I have an error in a markdown file it is a quick edit to a text file.
- Plain text is lightweight Images can be stored online and the lecture notes themselves are small. This
 might not matter where internet access is good, but in places with limited resources or wifi, this can be the
 difference from easily accessible lecture notes and bad ones.
- Plain text is always forward compatible Regardless of the next platform, if we have all of our knowledge/lecture notes stored in plain text it will be easy to extact them. When you switch platform, or compiling software, or style, there isn't a worry about the files not working appropriately.

```
https://simplystatistics.org/2017/06/13/
the-future-of-education-is-plain-text/
```

Additionally...

- Easy to use in version control
- Easy to dynamically update as part of an analysis "pipeline"

File	Good format(s)
Document	.md, .tex, .Rmd, .Rnw
Presentation	.tex, .Rmd, .Rnw
Code	.R, .Rmd, .py, .do, .ado
Data	.tsv, .csv
Codebook	.txt
Citations	.bib
Images	.svg, .pdf, .png
References	.bib

Is it possible to take the plain text ideology too far?

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Questions?

File names

Which of these do we like best?

- PhD Comics style
- Sequential version numbers
- Datestamps

File names

Which of these do we like best?

- PhD Comics style
- Sequential version numbers
- Datestamps
- None of the above (Git!)

1 Organizing Things

2 Building Things

3 Keeping and Changing Things

4 Thursday: Hands-On

Activity!

What's your analytic workflow? How do you get results into a paper, poster, or presentation?

Make figure/table/analysis in R

- Make figure/table/analysis in R
- Copy/paste into Word document

- Make figure/table/analysis in R
- 2 Copy/paste into Word document
- Adjust figure/table numbering

- Make figure/table/analysis in R
- 2 Copy/paste into Word document
- Adjust figure/table numbering
- 4 Double check references

- Make figure/table/analysis in R
- 2 Copy/paste into Word document
- Adjust figure/table numbering
- 4 Double check references
- 5 Save as PDF
- 6 Change something in 1, repeat 2-5

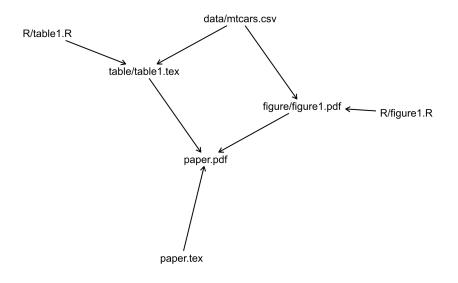
- Make figure/table/analysis in R
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- 5 Save as PDF
- 6 Change something in 1, repeat 2-5
- Get feedback (f*ck!!), repeat 1-5

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- 2 Copy/paste into Word document
- Adjust figure/table numbering
- 4 Double check references
- 5 Save as PDF
- 6 Change something in 1, repeat 2-5
- Get feedback (f*ck!!), repeat 1-5
- B Get reviews (f*ck!!!!!), repeat 1-5

- Make figure/table/analysis in R
- 2 Copy/paste into Word document
- Adjust figure/table numbering
- 4 Double check references
- 5 Save as PDF
- 6 Change something in 1, repeat 2-5
- Get feedback (f*ck!!), repeat 1-5
- B Get reviews (f*ck!!!!!), repeat 1-5
- Repeat 7 (f*ck!!!!!!!!!!!), repeat 1-5

Workflows as DAGs

- Reproducibility means executing a DAG
- DAG
 - Directed
 - Acyclic
 - Graph
- Files are *nodes*; workflows are *arrows*
- Example: https: //github.com/leeper/make-example



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	Endogenous covariates						Pop, 18 and older	
	Sample-selection models						Pop, 65 and older	
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	Multivariate time series		Panel data			marriage	Number of marriages	
	Longitudinal/panel data		Mixed-effects liper regression			divorce	Number of divorces	
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(1980 Census data	SEM (structural equation modeling)							
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Lose track of the DAG

- Lose track of the DAG
- Won't comply with DA-RT verification policies

- Lose track of the DAG
- Won't comply with DA-RT verification policies
- You will make mistakes!

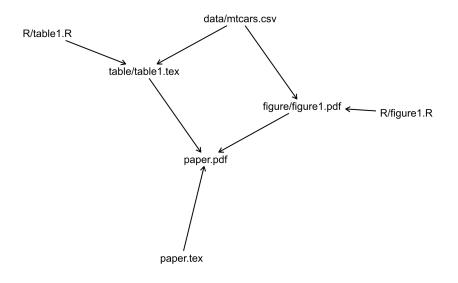
- Lose track of the DAG
- Won't comply with DA-RT verification policies
- You will make mistakes!
- Eventually, you will have wasted your entire life manually fixing references, figure/table cross-references, and making sure that all of your numbers are correctly rounded and p-values have the correct number of stars next to them!

The prevalence of statistical reporting errors in psychology (1985–2013)

Michèle B. Nuijten¹ · Chris H. J. Hartgerink¹ · Marcel A. L. M. van Assen¹ · Sacha Epskamp² · Jelte M. Wicherts¹

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Abstract This study documents reporting errors in a sample of over 250,000 p-values reported in eight major psychology journals from 1985 until 2013, using the new R package "statcheck." statcheck retrieved null-hypothesis significance testing (NHST) results from over half of the articles from this period. In line with earlier research, we found that half of all published psychology papers that use NHST contained at least one p-value that was inconsistent with its test statistic and degrees of freedom. One in eight papers contained a grossly inconsistent p-value that may have affected the statistical conclusion. In contrast to earlier findings, we found that the average prevalence of inconsistent p-values has been stable over the years or has declined. The prevalence of gross inconsistencies was higher in p-values reported as significant than in p-values reported as nonsignificant. This could indicate a systematic bias in favor of significant results. Possible solutions for the high prevalence of reporting inconsistencies could be to encourage charing data to let co-authors check results in a co-called Most conclusions in psychology are based on the results of null hypothesis significance testing (NHST; Cumming et al., 2007; Hubbard & Rvan, 2000; Sterling, 1959; Sterling, Rosenbaum, & Weinkam, 1995). Therefore, it is important that NHST is performed correctly and that NHST results are reported accurately. However, there is evidence that many reported p-values do not match their accompanying test statistic and degrees of freedom (Bakker & Wicherts, 2011; Bakker & Wicherts, 2014; Berle & Starcevic, 2007; Caperos & Pardo, 2013; Garcia-Berthou & Alcaraz, 2004; Veldkamp, Nuijten, Dominguez-Alvarez, Van Assen, & Wicherts, 2014; Wicherts, Bakker, & Molenaar, 2011). These studies highlighted that roughly half of all published empirical psychology articles using NHST contained at least one inconsistent p-value and that around one in seven articles contained a gross inconsistency, in which the reported p-value was significant and the computed n-value was not or vice versa



Do everything in one file

1 Do everything in one file

Master file calls code for one-file-per-output

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make ("code within workflow")

Do everything in one file

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4 knitr/rmarkdown ("workflow within code")

Everything in One File

```
# Brexit Deservingnes Experiment Analysis
# setwd("c:/users/thomas/dropbox/brexitdeservingness/")
```

```
# load data
dat <- rio::import("data/LSE_Hobolt_May18_Client.sav")</pre>
stopifnot(identical(dim(dat), c(3273L, 62L)))
# Regression analysis: perceived deservingness
stargazer::stargazer(
  # reduced model (only leavers and remainers) with interaction
  lm(opinion ~ identity * condition, data = subset(dat, identity %in% c("A Leav
  type = "tex",
  out = "figures/results-deservingness.tex",
  star.char = c("*").
  star.cutoffs = c(0.05),
  notes = c("* $p<0.05$"),
  notes.append = FALSE,
  model.numbers = FALSE,
  float = FALSE.
  digits = 2,
  align = TRUE
```

One-File-Per-Output

Preference Trial Experiment Analysis
Thomas J. Leeper
2018-06-25
#setwd("C:/Users/Thomas/Dropbox/KnowledgeGaps")

```
# code
library("car")
library("xtable")
library("GK2011")
source("Analysis/functions.R")
```

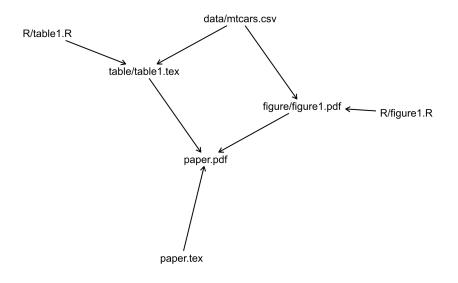
```
# recoding
source("Analysis/experiment_cleaning.R")
```

```
# demographics
source("Analysis/experiment_demographics.R", echo = TRUE)
```

```
## Main analysis
source("Analysis/experiment_knowledge.R")
```

```
## Appendix
source("Analysis/experiment_appendix.R")
```

What's missing from these workflows?



make with a makefile

```
all: paper.pdf
figure/figure1.pdf: R/figure1.R data/mtcars.csv
    Rscript R/figure1.R
table/table1.tex: R/table1.R data/mtcars.csv
    Rscript R/table1.R
paper.pdf: paper.tex figure/figure1.pdf table/table1.tex
    pdflatex $<
    p
```

Dynamic documents: rmarkdown

1 YAML metadata header

```
title: My Manuscript
author: Thomas J. Leeper
---
```

2 Document contents in markdown

A header
A subhead
This is my manuscript, **bold** and *italic*.

3 Code in "code chunks":

```
''`{r chunk1}
# R code
hist(rnorm(1000))
'''
```

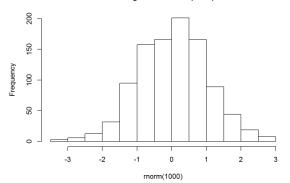
- title: My Manuscript
- author: Thomas J. Leeper
- date: 2017-09-21
- output: pdf_document

```
This is my manuscript.
```

```
'``{r chunk1}
# R code
hist(rnorm(1000))
'''
```

This is my manuscript.

R code hist(rnorm(1000))



Histogram of rnorm(1000)

What about Stata?

1 \checkmark Do everything in one file

 \mathbf{Z} \checkmark Master file calls code for one-file-per-output

3 √ make ("code within workflow")

A ? Nothing as powerful as rmarkdown/knitr

How do you pick a workflow?

- There is no one-size-fits-all workflow!
- Decide what works for you for a given project with particular collaborators
- I use multiple workflows on different projects

Questions?

1 Organizing Things

2 Building Things

3 Keeping and Changing Things

4 Thursday: Hands-On

Activity!

What tools do you use to store, share, and/or archive your research materials?

- Collaborating with yourself or others in the future
 - Going back in time for long-lived projects
 - Verification at publication stage

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- 2 Collaborating with others now
 - Collaborating simultaneously
 - Collaborating asynchronously

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 - Going back in time for long-lived projects
 - Verification at publication stage
- 2 Collaborating with others now
 - Collaborating simultaneously
 - Collaborating asynchronously
- Collaborating with others after you die
 Future reproducibility requests

Live Collaboration

Other Collaboration

Live Collaboration Google Docs Overleaf Dropbox/Box/etc. Email?

Other Collaboration

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Google Docs

Overleaf

Dropbox/Box/etc.

Other Collaboration

- Active project: Version control (git)
- Backup: Dropbox, GDrive, S3, Github

Email?

Live Collaboration

Google Docs

Overleaf

Email?

Dropbox/Box/etc.

Other Collaboration

- Active project: Version control (git)
- Backup: Dropbox, GDrive, S3, Github
- Archiving:
 Dataverse, Zenodo,
 Figshare, OSF

- Git is "an open-source distributed version control system"
- Developed in 2005 by Linus Torvalds
- Widely used in software development world

Helps you keep and annotate snapshots of your project over time

- Better than renaming your files all the time
- Better than using within-file VCS (e.g., Word)
- Better than single-stream sharing (e.g., Dropbox)

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- Helps you keep and annotate snapshots of your project over time
 - Better than renaming your files all the time
 - Better than using within-file VCS (e.g., Word)
 - Better than single-stream sharing (e.g., Dropbox)
- Facilitates collaboration (incl. with future you)
- It's FOSS with lots of clients, tools, and community support
 - Widely used in software development world

Version control helps you stay organized

Version control helps you stay organized
 What's important to keep around?

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- 1 What's important to keep around?
- 2 What's not important to keep around?

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- 3 What is all this crap?

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Think "tracked changes" for all of your files Save history of changes/versions

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- Save history of changes/versions
- Experiment non-destructively

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- Save history of changes/versions
- Experiment non-destructively
- Collaborate
- You're probably already version controlling informally!

Learning Objectives

- Understand how to organize a reproducible research project
- 2 Recognize different approaches to reproducibility and tools for implementing various reproducible workflows
- 3 Th: Apply various workflows to your own work
- Th: Understand how to collaborate reproducibly

 Once you work reproducibly, you'll never want to go back to your old workflow

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- "Advanced" workflows (e.g., make, git) get complicated — StackOverflow is your friend
- Collaborators probably don't know how to (or want to) use these tools
- Reproducibility is selfish first and for science second!

Questions?

1 Organizing Things

2 Building Things

3 Keeping and Changing Things

4 Thursday: Hands-On

- 5 Hands-On
 - Introductory Git
 - Git Branches & History
 - Collaborating with Git
 - Intermediate Git
 - Rmarkdown/knitr
 - make

5 Hands-On ■ Introductory Git

- Git Branches & History
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make

Goal of Hands-On Practice

- **I** Work together on migrating a workflow
- 2 Dig through replication archives
- Work individually or in pairs on making workflow more reproducible

Let's vote: What should we do?

Using Git

 Git create a "local repository" file that you can interact with using a number of tools

- Command-line git
- Git Bash
- Git GUI
- GitHub Desktop
- RStudio (via "Projects")
- GitHub/Bitbucket/GitLab web interfaces
- Gitkraken
- git2r (R package)
- . . .

Initializing a Project Structure

There's no single best way to organize a project

But, some words of wisdom:

- Put like with like
- Avoid excessive hierarchy
- Not everything needs to go into git
- Steal others' structures!

```
git --version
git
git config --global user.name "My Name"
git config --global user.email "me@example.com"
git config --list
```

```
git init
git status
echo Hello world! > README.md
git add README.md
git status
git rm --cached README.md
git status
git add --all
git commit -m "my first commit!"
git status
git log
```

- 1 stage
- 2 commit
- 3 branch
- 4 merge
- 5 push and pull

- 1 stage
 - add/stage: select files to be recorded in a "snapshot" of the project
 - rm/unstage: remove files from the snapshot (but not from your computer)
- 2 commit
- 3 branch
- 4 merge
- 5 push and pull

- 1 stage
- 2 commit
 - commit: record a permanent snapshot of the staged files, labelled with a "commit message"
 commends and if (trainedby the mest message)
 - amend: modify (typically the most recent) commit with new changes or commit message
- 3 branch
- 4 merge
- 5 push and pull

- 1 stage
- 2 commit
- 3 branch
 - produce a complete *local* copy of the project where changes can be made independently of the "master" branch
- 4 merge
- 5 push and pull

- 1 stage
- 2 commit
- 3 branch
- 4 merge

 update a branch with changes from another local branch (or a remote); you can change multiple branches independently.

5 push and pull

- 1 stage
- 2 commit
- 3 branch
- 4 merge
- 5 push and pull
 - **push**: send the project (any new commits) to a remote server (like GitHub)
 - **pull**: grab new commits from a remote server

- 1 stage
- 2 commit
- 3 branch
- 4 merge
- 5 push and pull

90% of What You Need

- git add (stage) or git rm (unstage)
- git commit
- git status
- git log
- git remote
 - git push
 - git pull
- git branch
 - git merge

5 Hands-On

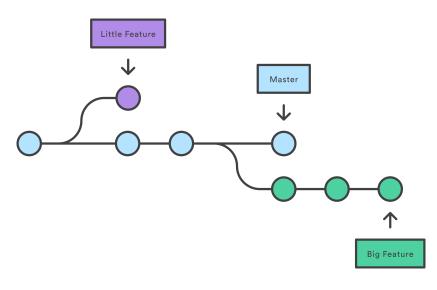
Introductory Git

Git Branches & History

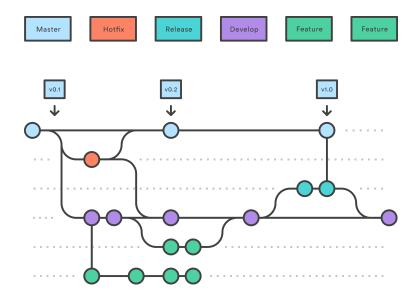
- Collaborating with Git
- Intermediate Git
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Branches

- Branches are local, parallel versions of your entire project
- Useful for multiple things:
 - Experimentation
 - Manuscript submissions
 - Collaboration



Source: https://www.atlassian.com/git/tutorials



Source: https://www.atlassian.com/git/tutorials

Simple branch and merge

git status git checkout -b thomas git status # do something git add --all git commit -m "thomas's commit" git checkout master git branch git log --graph --oneline git merge thomas

GUIs

- You can do everything in Git on the command line
- GUIs can be helpful for:
 - Exploring history
 - Visualizing branches
 - Confirming what you're doing

Merge conflicts

```
git checkout -b thomas
git status
# do something to README.md
git add --all
git commit -m "change on thomas"
git checkout master
# do something to README.md
git add --all
git commit -m "change on master"
git merge thomas
git log
```

Navigating History

- git status
- git log
- git checkout <commit hash>
- git status
- ls
- cat README.md
- git checkout master

```
git status
git log
git checkout <commit hash>
git status
ls
echo aaaaaah!>manuscript.txt
git checkout master
```

Remotes

- A server ("cloud") instance of the Git repository
- Useful for multiple things:
 - Collaboration
 - Transparency
 - Archiving/backups
 - Using web-based Git interfaces

Remotes

Three major players in cloud Git

- GitHub
- Atlassian Bitbucket
- GitLab
- Why choose one or the other?
 - Cost
 - Collaborators
 - Private repositories

```
git status
git remote add github
https://github.com/leeper/rt2
git remote
git remote set-url
git remote rename
git remote remove
```

```
git push github master -u
git fetch github
git fetch github master
git checkout -b new-idea
git push github new-idea
git checkout master
git pull github master
git pull
```

git status

```
git status
git tag -a v0.0.1 -m "v0.0.1"
git push --tags
```

```
git tag -d v0.0.1
```

Tags versus Branches

Branches are for working versions of project

- Collaborator-specific branches
- Submission-specific branches
- Experimental or "bug fix" branches
- Tags are for marking particular snapshots
 - Significant moments in project history
 - Journal submission or conference version
 - Formal "releases"

5 Hands-On

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Collaboration

Technical aspects

- Give collaborators access on GitHub (or wherever)
- Work on separate branches
- Merge agreed changes into master

Human factors aspects

- Requires agreeing on workflow
- Communication about what goes in "master"
- Can feel awkward if moving from a Dropbox- or email-based collaboration style

Try it with a partner!

- 1 Partner A create a GitHub repo; give Partner B access
- 2 Partner B should git fetch/git pull the repo
- 3 Partner B should create a local branch and git push
- 4 Partner A should git fetch the branch
- 5 Partner A should git merge the branch to master and git push
- 6 Partner B should git pull from master
- 7 Both use git log to compare

5 Hands-On

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```
git status
git diff README.md
git diff HEAD README.md
git diff HEAD~1 README.md
git diff HEAD~2 README.md
git diff HEAD~3 README.md
git diff HEAD~20 README.md
git diff <commit hash> README.md
git diff <commit hash>
```

!! DANGER: Amend Commit !!

```
git status
git log --oneline
# maybe add/rm files
git amend
# enter the hell of vim
```

```
git config --global core.editor
"<executable> <options>"
```

Safe reversion

```
git status
git log --oneline
git revert <commit hash>
# enter the hell of vim
# or something else terrible
git revert --abort
```

!! DANGER: Unsafe reversion !!

The StackOverflow Question

```
git status
echo "bad bad bad" > bad.txt
git status
echo bad.txt > .gitignore
git status
echo bad bad bad > bad1.txt
echo bad bad bad > bad2.txt
echo bad* > .gitignore
git status
git add bad1.txt -f
git status
```



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Rmarkdown

YAML metadata header

```
title: My Manuscript
author: Thomas J. Leeper
```

2 Document contents in markdown

A header
A subhead
This is my manuscript, **bold** and *italic*.

3 Code in "code chunks":

''`{r chunk1}
R code
hist(rnorm(1000))
'''

- title: My Manuscript
- author: Thomas J. Leeper
- date: 2017-09-21
- output: pdf_document

```
This is my manuscript.
```

```
'``{r chunk1}
# R code
hist(rnorm(1000))
'''
```

Markdown Basics

Markdown is a very simple markup language for formatting simple texts:

italics
bold
'preformatted'
Heading
Heading
Heading
[link](https://google.com)

italics **bold** preformatted Heading Level 1 Heading Level 2 Heading Level 3 link

Chunk Options

- '``{r chunk1, eval=TRUE, echo=TRUE}
 2 + 2
 '``
- '``{r chunk2, eval=TRUE, echo=FALSE}
 2 + 2
 '``

'``{r chunk3, echo=FALSE, results="hide"}
2 + 2
'``

Global Chunk Options

Basic Tables

```
'``{r table1, results = "asis"}
xtable::xtable(table(mtcars$cyl, mtcars$gear))
```

```
knitr::kable(head(mtcars))
....
```

Regression Results Tables

```
''`{r table2, results = "asis"}
library("stargazer")
stargazer(
  x1 <- lm(mpg ~ disp + wt,
           data = mtcars),
  x2 <- lm(mpg ~ disp + wt + vs,
           data = mtcars),
  header = FALSE
(((
```

Figures

```
''`{r fig1,
    fig.cap = "Fuel Economy by Weight",
    fig.height = 4,
    fig.width = 6}
library("ggplot2")
ggplot(mtcars,
    aes(x = wt)
        y = mpg,
        colour = factor(cyl)) +
  geom point()
...
```

You can work in LaTeX, too!

```
\neg \Box
knitr_basics.Rnw ×
       📄 🗳 🔍 🔂 Format - 🔁 Compile PDF 🛛 🛃
                                                          📑 Run 🔄 🔁 Chunks 🕶
     \documentclass{article}
  1
  2
  3
    \begin{document}
  4
  5
    Here is a code chunk.
  6
  7 - <<>>=
  8
     a < -1+1
  9
     а
 10
     a
 11
 12
    You can also write inline expressions, \Sexpr{a}.
 13
 14
     \end{document}
 15
      C (Top Level) $
                                                                        R Sweave
15:1
```

You can work in LaTeX, too!

```
\begin{document}
Here is a code chunk.
\begin{knitrout}
\definecolor{shadecolor}{rgb}{0.969, 0.969, 0.969}\color{fgcolor}\begin{kframe}
\begin{alltt}
\hlstd{a} \hlkwb{<-} \hlnum{1}\hlopt{+}\hlnum{1}</pre>
\hlstd{a}
\end{alltt}
\begin{verbatim}
## [1] 2
\end{verbatim}
\end{kframe}
\end{knitrout}
You can also write inline expressions, 2.
```

```
\end{document}
```

You can work in LaTeX, too!

Here is a code chunk.

a <- 1+1 a ## [1] 2

You can also write inline expressions, 2.

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make

makefiles

all: <final-target>

<target-1>: <source-file> <source-file> <script to produce target from source-file(s)>

<target-2>: <source-file> <target-1> <script to produce target from source-file(s)>