BMI 826-003

Tools for Reproducible Research

Karl Broman

Biostatistics & Medical Informatics, UW-Madison

biostat.wisc.edu/~kbroman
 github.com/kbroman
 @kwbroman
Course web: bit.ly/tools4rr

Reproducible

Reproducible

VS.

Replicable

Reproducible

VS.

Correct

Levels of quality

- Are the tables and figures reproducible from the code and data?
- Does the code actually do what you think it does?
- In addition to what was done, is it clear why it was done?

```
(e.g., how were parameter settings chosen?)
```

- Can the code be used for other data?
- Can you extend the code to do other things?

Basic principles

- Everything via code
- Everything automated
 Workflow and dependencies clearly documented
- Get the data in the most-raw form possible
- Get any/all data and meta-data possible
- Keep track of the provenance of all data files
- Be self-sufficient

Why do we care?

- Avoid embarrassment
- More likely correct
- ► Save time, in the long run
- Greater potential for extensions; higher impact

Your closest collaborator is you six months ago, but you don't reply to emails.

What could go wrong?

- "The attached is similar to the code we used."
- "Where did this data file come from?!"
- "Can you repeat the analysis, omitting subject X?"
- "This part of your script is now giving an error."

Need to avoid

- Open a file to extract as CSV
- Open a data file to do even a slight edit
- Paste results into the text of a manuscript
- Copy-paste-edit tables
- Copy-paste-adjust figures

Basic tools

- Automation with Make
- ► Unix command line
- Latex and Markdown
- ▶ Knitr
- Version control with git
- R packages
- ► Python (or Ruby or Perl)

Other topics

- Organizing projects
- Writing clear code
- ▶ Don't Repeat Yourself (DRY)
- Testing and debugging
- Handling big jobs
- Licenses; human subjects data

Don't Repeat Yourself

- ► In code, in documentation, etc.
- Repeated bits of code are harder to maintain Write a function
- ► Use documentation systems like Roxygen2

 Documentation in just one place
- Make use of others' code

This course

- Brief intro to various tools and concepts
- Try everything out as we go along Ask questions!
- ► I don't know everything Make suggestions!
- Project
 - Write a bit of R code
 - Use version control
 - Make it an R package
 - Write a vignette

Automation with GNU Make

- Make is for more than just compiling software
- ► The essence of what we're trying to do
- Automates a workflow
- Documents the workflow
- Documents the dependencies among data files, code
- Re-runs only the necessary code, based on what has changed

Example Makefile

```
# Example Makefile for a paper
mypaper.pdf: mypaper.bib mypaper.tex Figs/fig1.pdf Figs/fig2.pdf
    pdflatex mypaper
    bibtex mypaper
    pdflatex mypaper
    pdflatex mypaper

    pdflatex mypaper

# cd R has to be on the same line as R CMD BATCH
Figs/fig1.pdf: R/fig1.R
    cd R;R CMD BATCH fig1.R fig1.Rout

Figs/fig2.pdf: R/fig2.R
    cd R;R CMD BATCH fig2.R fig2.Rout
```

Fancier example

```
FIG_DIR = Figs
mypaper.pdf: mypaper.tex ${FIG_DIR}/fig1.pdf ${FIG_DIR}/fig2.pdf
    pdflatex mypaper

# One line for both figures
${FIG_DIR}/%.pdf: R/%.R
    cd R;R CMD BATCH $(<F)

# Use "make clean" to remove the PDFs
clean:
    rm *.pdf Figs/*.pdf</pre>
```

How do you use make?

- ► If you name your make file Makefile, then just go into the directory containing that file and type make
- ► If you name your make file something.else, then type make -f something.else
- Actually, the commands above will build the first target listed in the make file. So I'll often include something like the following.

```
all: target1 target2 target3
```

Then typing make all (or just make, if all is listed first in the file) will build all of those things.

► To be build a specific target, type make target. For example, make Figs/fig1.pdf