

### About Us

#### John Muschelli

Assistant Scientist, Department of Biostatistics

PhD in Biostatistics, ScM in Biostatistics

Email: jmusche1@jhu.edu

### About Us

#### Kristin Linn

Assistant Professor, Department of Biostatistics and Epidemiology Perelman School of Medicine, University of Pennsylvania

PhD in Biostatistics

Email: klinn@mail.med.upenn.edu

# Learning Objectives

- Introduction to the Statistical Software R
- Reading and Writing Images
- Visualization of Images and Image Data
- ► Inhomogeneity Correction
- Brain Extraction
- Image Segmentation
- Coregistration Within and Between MRI Studies
- Intensity Normalization
- Segmentation of MS lesions

## Course Website/Materials

The Course overview is located at (with slides): http://johnmuschelli.com/ISBI\_2017.html

All materials for this course (including source for the slides) is located at:

https://github.com/muschellij2/imaging\_in\_r

#### RStudio Server

For this course, we will use an RStudio Server because installing all the packages can be a lengthy process. Mostly all the code we show requires a Linux/Mac OSX platform for FSL and other systems:

http://johnmuschelli.com/rstudio\_server.html

The code to make the server is located at: https://github.com/muschellij2/rneuro/blob/master/ms\_rstudio\_droplet.sh

### Installing R: Local

If you want to install things locally:

- You can install the latest R from http://cran.r-project.org/
- Install RStudio
- Directions for a local setup is http://johnmuschelli.com/ imaging\_in\_r/installing\_everything\_locally/



### What is R?

- ▶ R is a language and environment for statistical computing and graphics
- ▶ R is the open source implementation of the S language, which was developed by Bell laboratories
- ▶ R is both open source and open development

```
(source: http://www.r-project.org/)
```

## Why R?

- Powerful and flexible
- Free (open source)
- Extensive add-on software (packages)
- Designed for statistical computing
- High level language

# Why not R?

- ► Fairly steep learning curve
  - "Programming" oriented
  - Minimal interface
- Little centralized support, relies on online community and package developers
- Annoying to update
- ► Slower, and more memory intensive, than the more traditional programming languages (C, Java, Perl, Python)

## Working with R

- ► The R Console "interprets" whatever you type
  - Calculator
  - Creating variables
  - Applying functions
- "Analysis" Script + Interactive Exploration
  - Static copy of what you did (reproducability)
  - ▶ Try things out interactively, then add to your script

R essentially is a command line with a set of

functions loaded

### R Uses Functions, in Packages

- R revolves around functions
  - Commands that take input, performs computations, and returns results
  - When you download R, it has a "base" set of functions/packages (base R)
- Functions are enclosed in packages
  - ► These written by R users/developers (like us) some are bad
  - ▶ Think of them as "R Extensions"

## RStudio (the software)

RStudio is an Integrated Development Environment (IDE) for R

- It helps the user effectively use R.
- Makes things easier
- ▶ Is NOT dropdown statistical tools (such as Stata)
  - See Rcmdr or Radiant.
- ➤ All snapshots in these slides are taken from http://ayeimanol-r.net/2013/04/21/289/

### **RStudio**

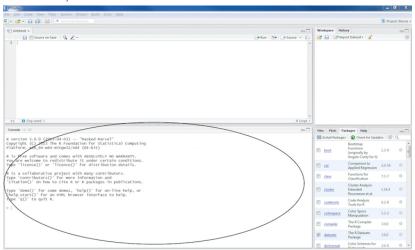
#### Easier working with R

- Syntax highlighting, code completion, and smart indentation
- Easily manage multiple working directories and projects

#### More information

- Workspace browser and data viewer
- Plot history, zooming, and flexible image and PDF export
- ▶ Integrated R help and documentation
- Searchable command history

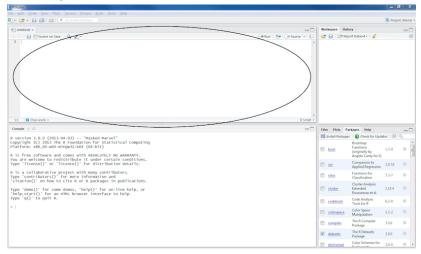
# RStudio/R Console



## RStudio/R Console

- Where code is executed (where things happen)
- ▶ You can type here for things interactively
- Code is **not saved** on your disk

### Source/Editor

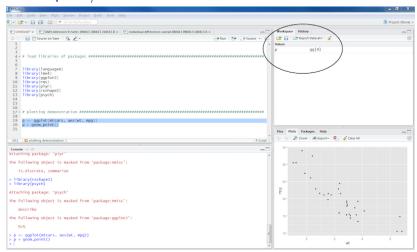


### Source

- Where files open to
- Have R code and comments in them
- Can highlight and press (CMD+Enter (Mac) or Ctrl+Enter (Windows)) to run the code

In a .R file (we call a script), code is saved on your disk

# Workspace/Environment



# Workspace/Environment

- ► Tells you what **objects** are in R
- ▶ What exists in memory/what is loaded?/what did I read in?

#### History

- Shows previous commands. Good to look at for debugging, but don't rely on it as a script. Make a script!
- Also type the "up" key in the Console to scroll through previous commands

#### Other Panes

- Files shows the files on your computer of the directory you are working in
- ▶ Viewer can view data or R objects
- ▶ **Help** shows help of R commands
- Plots pretty pictures
- Packages list of R packages that are loaded in memory