# Statistics and Data Analysis Using R

#### Fall, 2013

Statistics and Data Analysis Using R was developed to be a brief but systematic introduction to the R statistical software suite for biomedical scientists. We assume that you have at least a passing familiarity the plots and statistical analyses that are most commonly used in biomedical papers, but no formal background in statistics or programming is necessary. The primary objective is learning to use R, but the course also emphasizes the standards of practice that programmers and data analysts have implemented to ensure transparency, accuracy and accountability. The 5 week course is very hands on, we will review and expand on weekly programming assignments in dedicated lab sessions held every friday. To get the most out of the course, you should bring a laptop with R already installed on it (see installation instructions below) and spend some time out of class on each weeks programming assignment, and come to the Wednesday lab meetings ready to discuss your coding problems as well as solutions.

### Why R?

R (http://cran.r-project.org/) is free open-source statistical software available for PC, OSX and Linux platforms. R is a *command-line* program, meaning that in order to perform analyses, the user must type commands into a terminal rather than make selections from a menu. If all you will ever need to do is a t-test there are simpler tools available, but no menu driven programs can offer the analytic flexibility that cutting-edge science will demand. There are a number of other excellent statistical programs available, including SAS, Stata, and S-Plus, but we like R best for several reasons:

- R is free
- it is available for any operating system
- R is supported by a huge library of user-contributed packages
- Its graphical capabilities are unparalleled.

These features have made R the first choice of professional statisticians working in public health and medicine, you will probably never need another program, and this is a great opportunity to learn the basics.

## Installing R

You can download the latest versions of R for Windows, Mac OS X, and Linux as well as manuals and detailed installation instructions from the U.S homepage for R. http://cran.us.r-project.org R-3.0.2 is the current version but it is updated frequently and version numbers will change every few months.

R includes an editor and facilities for workspace management, but you may want to install Rstudio as well, http://rstudio.org/. Rstudio is a free and open source integrated development environment (IDE) for R that makes it much easier to manage your work sessions.

#### Instructors

Luigi Marchionni, MD, PhD (marchion@jhu.edu) Leslie Cope, PhD (cope@jhu.edu) Rob Scharpf, PhD (rscharpf@jhu.edu) Elana Fertig, PhD (ejfertig@jhmi.edu)

#### Time and Place

The course meets on wednesdays and fridays between Nov 6 and Dec 13 from 10:30am-12:30pm in PCTB 113/114. Please note that the Nov. 20 session will meet in PCTB G18/19 instead.

#### Course Website

Course materials can be found online at http://tinyurl.com/Intro2R-Fall-2013

### Course Textbook

There is no required course text book, and in fact we would rather have you become comfortable with the extensive built in documentation than depend on external sources for the basics. However, references to external sources will be provided for the interested reader as the course progresses.

#### Course Schedule

- Week 1- Nov 6-8 Getting started in R, data input and output (Rob Scharpf)
- Week 2 Nov 13-15 Plotting and Statistical Analysis (Leslie Cope)
- Week 3 Nov 20-22 Introduction to programming, loops and control structures (Elana Fertig)
- Week 4- Dec. 4-6 Reproducibility, crash class on regular expressions and string manipulations. (Luigi Marchionni)
- Week 5 Dec 11-13 Writing your own functions, good programming practice (Staff)

#### Course Work

Weekly programming assignments will be reviewed and extended in Friday lab sessions.