Statistical Software Camp: Introduction to R

Day 1

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1 Introduction

1.1 Why Use R?

- Widely-used (ever-increasingly so in political science)
- Free
- Power and flexibility
- Graphical capabilities
- Pedagogical usefulness
- Learning curve is initially steep, but it gets easier later

1.2 Installing R

- 1. Go to http://www.r-project.org.
- 2. Follow the link under the Download section on the left hand side of the page.
- 3. Make sure to select an appropriate mirror (the closer the faster) and operating system.
- 4. Once the download is finished, you can install R in the same way you install other software.

1.3 Using R

1.3.1 Typing Directly

- R as a calculator
- <- or = to assign a name to an object (<- is recommended)
- # for comments
 - > apple <- 7
 > apple
 - [1] 7

```
> ## Can use = but not recommended
> orange = 12
> orange
[1] 12
```

• Hello \neq hello \neq HELLO

```
> obama <- "president" # character string
> Obama # generates error
```

- Variable names cannot begin with numbers
- One command per line or use;

```
> mathcamp09 <- "Fun"; enrollment <- 15
> mathcamp09
[1] "Fun"
```

> enrollment

[1] 15

• Space between syntaxes will be ignored, but not in ""

```
> pol.571 <- "first in quant sequence"
> pol.571
[1] "first in quant sequence"
```

- Can use <up> and <down> to navigate through previously entered commands
- An example: calculate the amount of Princeton's endowment the university could spend for each student without harming the principle.

```
> 15787000000 / (4918+2416)  # Endowment divided by the total number of students

[1] 2152577

> endow.ps <- 15.787*(10^9) / (4918+2416)  # Stores the result as `endow.ps'

> yield.ps <- endow.ps*0.08  # Creates new object `yield.ps'

> yield.ps
```

1.3.2 Using a Text Editor

- Almost always want to use a text editor
- Efficiency, Replicability
- Built-in R editor; what we'll use in the camp
- Notepad, TextPad, TextEdit, etc.
- More fancy editors: WinEdt (http://www.winedt.com) for Windows and Aquamacs (http://aquamacs.org) for Mac OS X
- Send code from R editor to **console** by:
 - Copy & Paste
 - Highlight and press <Ctrl + R> on Windows
 - Highlight and press < Apple + Return > on Mac
- An example: Repeat the endowment calculation above using the R editor.

1.4 Numeric Vectors

A **vector** is simply string of numeric values connected in a specific order.

1.4.1 Creating, Indexing and Combining

- Use c() to enter a data vector
- Use square brackets ([]) to access elements of a vector
- c() can also combine more than one vectors

```
> endow <- c(9928, 11207, 13045, 15000) #Princeton endowment 2004-2007 > endow
```

- [1] 9928 11207 13045 15000
- > endow[2] # Use square brackets to access elements of a vector
- [1] 11207
- > endow[c(2, 4)]
- [1] 11207 15000
- > endow[4] <- 15787
- > endow
- [1] 9928 11207 13045 15787
- > endow[-1]

```
[1] 11207 13045 15787

> endow <- endow / 1000
> endow

[1] 9.928 11.207 13.045 15.787

> endow.old <- c(8.398, 8.359, 8.320, 8.730) #Years 2000-2003
> endow <- c(endow.old, endow) #combine vectors
> endow

[1] 8.398 8.359 8.320 8.730 9.928 11.207 13.045 15.787
```

1.4.2 Manipulating

• Named vectors can be manipulated like numbers

```
> endow.penn <- c(3.201,3.382,3.393,3.547,4.019,4.370,5.313,6.635) #UPenn endowment
> endow.penn

[1] 3.201 3.382 3.393 3.547 4.019 4.370 5.313 6.635

> ratio <- endow / endow.penn
> ratio

[1] 2.623555 2.471614 2.452107 2.461235 2.470266 2.564531 2.455298 2.379352
```

1.4.3 Basic Functions

- General format of a function: output <- functionname(input)
- More than one input: output <- functionname(input1, input2, ...)
- Inputs are often called **arguments**
- The colon operator (:) creates a simple sequence
- Use seq() for more complex sequences

```
> 1:10
[1] 1 2 3 4 5
```

• names() gives named entries to a vector

```
> names(ratio) <- 2000:2007
> ratio
```

```
2000
               2001
                         2002
                                  2003
                                           2004
                                                     2005
                                                              2006
  2.623555 2.471614 2.452107 2.461235 2.470266 2.564531 2.455298 2.379352
• Simple calculations: min(), max(), range(), length(), sum() and mean()
  > min(ratio) # Shows the minimum value of our vector
  [1] 2.379352
  > max(ratio) # Same, but for the maximum value
  [1] 2.623555
  > range(ratio)
  [1] 2.379352 2.623555
  > length(ratio) # Tells us the number of values in our vector
  [1] 8
  > ratio[length(ratio)]
      2007
  2.379352
  > sum(ratio) # Sum of all values in the vector
  [1] 19.87796
  > sum(ratio)/length(ratio) # Long way to calculate the mean
  [1] 2.484745
  > mean(ratio)
                # Easier
  [1] 2.484745
   Object Class
• An object belongs to a certain class (e.g. numeric, character, function)
• The function class() returns the class of an object
• The function summary() to get a summary of an object
```

2007

1.5

- - > endowment <- c(9928, 11207, 13045, 15787)> class(endowment)
 - [1] "numeric"
 - > summary(endowment)

```
Min. 1st Qu. Median Mean 3rd Qu. Max.
9928 10890 12130 12490 13730 15790
```

- > schools <- c("Harvard", "Princeton", "Yale", "MIT", "Stanford", "CalTech")
 > class(schools)
- [1] "character"
- > summary(schools) #Not so useful when dealing with "character" vectors

Length Class Mode 6 character character

- factor is more useful for categorical variables than character
- The function as.factor() coerces an object into a factor (as.numeric(), as.character())
- The function levels() returns the categories of a factor
 - > regions <- c("Africa", "Africa", "Asia", "Asia", "Africa", "Middle East")
 > regions
 - [1] "Africa" "Africa" "Asia" "Africa"
 - [6] "Middle East"
 - > class(regions)
 - [1] "character"
 - > regions <- as.factor(regions)</pre>
 - > regions
 - [1] Africa Africa Asia Asia Africa Middle East Levels: Africa Asia Middle East
 - > class(regions)
 - [1] "factor"
 - > levels(regions) # Displays all possible categories for our factor
 - [1] "Africa" "Asia" "Middle East"
 - > summary(regions) # Useful output for factors

Africa Asia Middle East
3 2 1

1.6 Workspace

- The workspace is the sandbox where R temporarily saves everything you've created or loaded into R
- The functions objects() and ls() list all objects in the workspace
- To remove an object from the workspace, use remove() or rm()
- You can remove multiple objects via rm(object1, object2, ...)

```
> objects()
 [1] "apple"
                   "endow"
                                "endow.old"
                                              "endow.penn" "endow.ps"
                   "enrollment" "mathcamp09"
                                              "orange"
                                                            "pol.571"
 [6] "endowment"
[11] "ratio"
                   "regions"
                                "schools"
                                              "vield.ps"
> rm(schools)
> 1s()
 [1] "apple"
                   "endow"
                                "endow.old"
                                              "endow.penn" "endow.ps"
 [6] "endowment"
                   "enrollment" "mathcamp09" "orange"
                                                            "pol.571"
                                "yield.ps"
[11] "ratio"
                   "regions"
```

1.7 Saving Objects and Workspace

- Use save() to save an object or multiple objects (or go to the pulldown menu under File: "Save Workspace...")
- The file name should be xxx.RData

```
> save(endowment, file = "H:/endowment.RData")
> save(endowment, regions, file = "H:/endowment.RData") # save multiple objects
```

• If you want to save all objects (i.e., the entire workspace), then use save.image()

```
> save.image("H:/Handout1.RData")
```

- The working directory is where R loads and saves your files.
- For example, save(endowment, file = "endowment.RData") will save endowment.RData in the working directory
- When you close down R, you will be asked if you'd like to save your current workspace. You normally want to choose NO.
- If you choose YES, R will save your files in the working directory as ".RData" (empty file name + extension).
- getwd() displays the working directory

```
> getwd()
```

- [1] "/home/teppei/Documents/teaching/software09/handouts"
- setwd() changes the working directory (or go to the pulldown menu)
 - > setwd("H:/") # Start saving to H drive
- Change the working directory to an appropriate directory at the beginning of your code

1.8 Getting R Help

- Use help() or ? to look up help on a function
- help.search("xxx") to look for functions that mention the word xxx

```
> help("save")
> ?save  # Same thing
> ?read.table
> help.search("save")
```

• Many resources on the web (e.g., search, mailing list at http://cran.r-project.org)

2 Reading Data and Doing Simple Analysis in R

2.1 Loading Workspace and Data

- Data analysis almost always begins by reading some data.
- If you have the file named ".RData" in the working directory, it will get automatically loaded when you start R in that directory
- load() for a previously saved R object/workspace (xxx.RData)
- read.table() for a simple text (ASCII) file (xxx.txt, etc.)
- read.csv() for a comma-delimited file (xxx.csv)
- read.delim() for a tab-delimited file (xxx.tab, etc.)
- Use header = T (for "TRUE") for a file with variable name headers in the first row
- read.table() defaults to header = T; others to F (= "FALSE")
- The object class for datasets is data.frame

```
> load("Africa.RData") # Looks for the file in our working directory
> Africa <- read.table("Africa.txt", header = TRUE)
> Africa <- read.csv("Africa.csv", header = TRUE)
> class(Africa)
[1] "data.frame"
```

• We can also pull data straight from the web

```
> primates <- "http://www.hopkinsmedicine.org/FAE/primate_structural_properties.txt"
> hop <- read.delim(primates, header=T)</pre>
```

2.2 Summary Statistics

- After reading new data, we usually compute summary statistics to get an overview.
- Various functions for summary statistics are available in R.
- mean() returns the (arithmetic) mean of an imput vector.
- max() and min() return the largest and smallest element of a numeric vector, respectively. range() gives you the distance between these two numbers.
- The function summary() will provide the mean, median, minimum, maximum, and quartiles of a numeric object and a table for a factor object (you can also use table() for this)

```
> mean(Africa$GDP.pc) # Simple mean
[1] 4616.115
> Africa$pop <- Africa$GDP / Africa$GDP.pc # Population of each country
> mean(Africa$GDP.pc)
[1] 4616.115
> median(Africa$GDP.pc)
[1] 2162.5
> range(Africa$GDP.pc)
[1]
      500 23294
> summary(Africa$GDP.pc)
   Min. 1st Qu.
                 Median
                           Mean 3rd Qu.
                                            Max.
    500
           1366
                   2162
                           4616
                                           23290
                                   5569
```

2.3 Graphs

- Graphs are an excellent device to summarize and present quantitative information. They are often way superior to numerical summaries, such as tables.
- Great graphics: Easy to understand the "story" without much explanation or eye-balling
- Bad graphics:
 - Inefficient (leaving out easy-to-add information)
 - Potentially misleading (leaving the reader with the wrong impression)
 - Too complicated (taking too much time to understand)
- The function barplot() will produce a barplot figure
- The function pie() will produce a pie chart figure

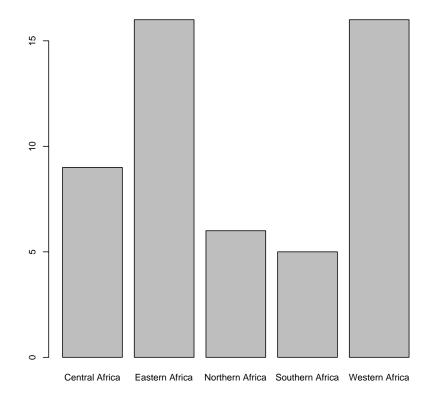
- main = Title to put on the graphic.
- xlab = Label for the x-axis. Similarly for ylab.
- xlim = Specify the x-limits, as in <math>xlim = c(0,10), for the interval [0,10]. Similar argument for the y-axis is ylim.
- type = Type of plot to make. Use "p" for points (the default), "1" for lines, and "h" for vertical lines.
- pch = The style of point that is plotted. This can be a number or a single character. Numbers between 0 and 25 give different symbols. The command plot(0:25,pch = 0:25) will show those possible.
- lty = When lines are plotted, specifies the type of line to be drawn. e.g., "solid", "dashed",
 "dotted" (See ?par for full details.)
- 1wd = The thickness of lines. Numbers bigger than 1 increase the default.
- col = Specifies the color to use for the points or lines, e.g., "blue", "red", "yellow".

Table 1: Useful arguments for plot() and other graphic functions. From Verzani (2005), Table 3.7 (p. 86).

- Specify arguments within each function to tweak the graphs to our exact specifications
- To learn the arguments for a particular command, use the help function (e.g., ?barplot or help("pie"))
 - > load("Africa.RData")
 - > x <- table(Africa\$Region) # make a table for a factor variable
 - > x

Central Africa Eastern Africa Northern Africa Southern Africa Western Africa 9 16 6 5 16

- > barplot(x) # Bar graph; takes in a numeric vector
- > barplot(x, xlab="Regions", ylab="Frequency", ylim=c(0,20),
- + main="Geographical Distribution of African Countries") # Adds labels
- > pie(x) # Pie chart; takes in a numeric vector



Geographical Distribution of African Countries

