an introduction to R for epidemiologists the basics

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Outline

🚺 about

• Some things people are doing with R

2 installing

3 using

- calculating, assigning, combining
- from calculations to programming
- packages
- 4 getting data into data

under the hood

- how is R different from other programs?
- workspaces and files
- workspaces
- help!

about

But first...



Credit where credit is due...

• Tomas Aragon, MD, DrPH

- Applied Epidemiology Using R
- http://medepi.com/
- John Fox, PhD
 - An Introduction to Statistical Computing in R
 - http://socserv.mcmaster.ca/jfox/Courses/UCLA/index.html
- Bill Venebles, PhD
 - An Introduction to R
 - o cran.r-project.org/doc/manuals/R-intro.pdf
- Phil Spector, PhD
 - Data Manipulation with R

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about





what is R?

A flexible, scalable, **free** tool for the description, analysis, visual display, exploration and interpretation of data.

- a calculator
- a suite of statistical tools
- a graphics creator
- a programming language
- a simulation lab
- a means of scientific documentation and discourse

It is uniquely suited to epidemiological analysis.

about Some things people are doing with R

Making stunning graphics Paul Butler



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Predicting Elections Andrew Gelman, "Red State, Blue State, Rich State, Poor State"



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Crawling and Scraping the Web

John Muschelli, Andrew Jaffe, Jeffrey Leek. Simply Statistics Blog





Making Money



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Publishing Newspapers

New York Times is an R Shop



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R intro

Doing Science Developed by Scientists for Scientists



so, why learn R?

Many options for epidemiological computing: SAS, STATA, SPSS, Mathematica, Excel....

But, if you want to ...

- accomplish many tasks with a single tool
- better understand the methods you use
- use methods not available in any other program
- develop and share your own methods
- collaborate with wide community of scientific colleagues

...R might be for you.

and did I mention it's free?

what R is not

- a GUI experience
- initially easy and intuitive
- warrantied in any way (if it runs, it can be on CRAN)
- a DBMS
- (*traditionally*) well suited to enormous data sets (*but that is changing*...)
 - historical 32-bit limit $(2^{31}-1)$ on size of a vector
 - in R, objects like matrices are actually vectors
 - R stores everything in RAM
 - old rule of thumb: 100,000 rows, 20 variables (*very* conservative), single object 10% of RAM, etc...
 - but, things have changed
 - 64 bit computing, increased RAM
 - interest in 'big data' in R http://www.cybaea.net/Blogs/Data/Big-data-for-R.html
 - packages like optimize read.csv, ff, bigmemory, data.table, rsqlite, python take advantage of "paralellism" (Hadoop, MapReduce)

how to install R

- go to http://www.r-project.org/
- elect CRAN (Comprehensive R Archive Network) from left menu
- Iink to nearby geographic site (e.g., http://software.rc.fas.harvard.edu/mirrors/R/)
- select your operating system
- Chose "Base" installation
- save R-X.X.X-win32.exe (windows) or R-X.X.X-mini.dmg (Mac OS X)
- vun the installation program accepting defaults

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R is a calculator

math operators and functions

arithmetic + , - , * , / power ^

convert 68 degrees Fahrenheit to Celsius ($C^0 = \frac{5}{9}(F^0 - 32)$)

5/9*(68-32)

assignment operator 'memory' key



using calculating, assigning, combining

concatenation function

combine or "vectorize"



c()

7



functions R "apps"

fx()

math operators and functions

mathematical functions - sqrt, log, exp, sin, cos, tan simple functions - max, min, length, sum, mean, var, sort

```
abs(-23) #absolute value
exp(8) # exponentiation
log(exp(8)) # natural logarithm
sqrt(64) # square root
```



- create a vector object called "my.numbers" that consists of the numbers 2,4,6 and 8.
- what is the square root of the sum of "my.numbers"?

about R functions

and their arguments

- function name without parentheses returns source code
 - useful if want to write own code or functions
- args(function) returns brief argument syntax
- some arguments have default values
 - if entered in correct order need not be named

```
args(sample)
data<-1:30
sample(s = 18, x = data, r = T)
# with replacement (spell enough to identify)
sample(s = sample(1:100, 1), x = sample(1:10, 5), r=T)
# arg any valid R expression</pre>
```

write your own function R is a programming language

```
my.function<-function(x){
5/9*(x-32)
}
my.function(68)
[1] 20
a<-c(134,156,222)
my.function(a)
[1] 56.66667 68.88889 105.55556</pre>
```

base R comes with many handy statistical functions

summary statistics

- summary(), fivenum(), stem() examine the distribution of a data set
- qqnorm(), qqline() normal plots
- boxplots() (a, b)

test statistics

- t.test() 2-sample t test, (a, b),
 - R does not by default assume equality of variances, (can use an F test to examine this assumption)
 - var.test() returns an F test, (a,b)

 wilcox.test() returns a two-sample non-parametric Wilcoxon (aka Mann-Whitney) or one-sample Wilcoxon (specify "paired=TRUE") test

Some statistics with R

```
myDat<-data.frame(cbind(outcome1=rnorm(1000,20,5),
outcome2=rpois(1000,5),
grp=factor(sample(c("a","b","c"), 1000, replace=T))))
```

```
summary(myDat$outcome1)
fivenum(myDat$outcome1)
stem(myDat$outcome1)
```

```
boxplot(myDat)
boxplot(outcome1~grp, data=myDat)
```

```
myDat2<-cbind(rnorm(1000,20,5), rpois(1000,5))
boxplot(mvDat2)</pre>
```

```
qqnorm(myDat$outcome1)
qqline(myDat$outcome1)
```

```
t.test(myDat$outcome1, myDat$outcome2)
```

```
wilcox.test(myDat$outcome1, myDat$outcome2)
wilcox.test(myDat$outcome1, myDat$outcome2, paired=T)
```



- run the following code to create a small data set: crashDat<-data.frame(age=rnorm(n=100,mean=22, sd=2), crash=sample(x=c(0,1),size=100,replace=T, prob=c(.2,.8)))
- what is the mean age
- create a box plot comparing age by crash status

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R packages a quick example

- collections of user-written functions
- install.package("my.package") copies the package from CRAN to your installation of R
- *library(my.package)* brings the package into RAM so you can use it

```
# is jello associated with diarrhea?
library(epitools) #load 'epitools' package
data(oswego) \#load Oswego dataset
names(oswego) #get some info
epitab(oswego$jello,oswego$ill) #use epitab for RR
# "pretty" up the results (everything is an object)
round(epitab(oswego$jello, oswego$ill,
method = "riskratio")$tab, 3)
# same function, different method for OR
with(oswego,round(epitab(jello, ill, method = "oddsratio")$tab, 3))
```

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getting data into data

the scan() and cbind() functions R as a spreadsheet

```
weight <- scan()
1: 134 156 222
4:
Read 3 items
height <- scan()
 1: 60 63 72
4: Read 3 items
bmi <- (weight*703)/height^2</pre>
cbind(weight, height, bmi)
    weight height
                       bmi
[1,]
    134 60 26.16722
[2,] 156 63 27.63114
[3,] 222 72 30.10532
```

NB: to scan in character variables use scan(, what = "")

getting "real" data into R "there's a function for that"

- read.table() is how you get data into R
- optimized version read.csv() even better

```
cars<-read.table("http://www.columbia.edu/~cjd11/
charles_dimaggio/DIRE/resources/R/cars.txt",
header=T, stringsAsFactors=F)
```

using read.csv

```
dig<-read.csv("http://www.columbia.edu/~cjd11/charles_dimaggio
DIRE/resources/R/dig.csv", header=T,
  stringsAsFactors=F) #digitalis data
  str(dig)
  table(dig$TRTMT,dig$DEATH)
```



- go to http://www.columbia.edu/~cjd11/charles_dimaggio/ DIRE/styled-4/styled-6/
 - DIRE \rightarrow epidemiology \rightarrow R
- click to download the data file called "sparcs" to your machine
- read the file into R using read.csv()
 - remember to save the dataframe to a named object
- what is the mean age?
- how many males are in county "59" (Brooklyn)?

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about data and procedures

- folks come to R from programs like SAS, SPSS and (gasp) Excel
- data manipulation steps or procs are followed by analytic steps or procs
- these two activities are fairly-well demarcated and differentiated
- data are mutable, procs are immutable

R is different functional programming, and R objects

- functional programming data are immutable, functions return new "objects"
 - could be data, could be something else, e.g. a regression object, a table object
 - you only see minimal information about the new object on your screen
 - you can save the results of a function as a new object
- everything in R (including functions) is an object
 - some objects you will learn about: vectors, matrices, arrays, lists, dataframes
 - objects have "characteristics" that determine how they "behave"

practical implications

- same function will return different results depending on the argument object type
- e.g. plot(numeric.data) returns scatterplot, plot(sp.class.data) returns a map
- you can (and often must) supply a function as an argument to another function
- e.g. plot(table(a,b))
- you will write a lot of parentheses
- "magrittr" allows pipeline of operations

```
plot %>%
    table(a,b)
```

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workspaces and files

workspaces

help!

your workspace

and how to work in it

- R session stores objects, data etc ... in a .RData file
 - at end of session will be prompted to "save workspace"
 - you may or may not want to
 - do save your source document (we'll discuss)
- getwd() to find location current workspace
- setwd() to set it
- *ls()* or *objects ()* to list current objects in workspace
- rm() to remove objects
- *save.image()* to save *current* space
- save() more control
- load() place .RData file back into workspace
- *q()* quit

objects in your workspace

identifying, assessing, removing

- ls() / objects() view the objects in the current workspace, pattern = search for object names that contain phrase eg ls(pattern = "dat")
- rm() / remove() remove workspace objects eg rm(dat, dat2, dat3, dat4) CAUTION - rm(list = ls()) will remove everything
- str() / mode() / class() "go to" functions to assess objects
- data() displays available data sets data(Titanic) Titanic str(Titanic)

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getting help

online community (http://r.789695.n4.nabble.com/), tutorials
(http://www.ats.ucla.edu/stat/r/), search sites
(http://www.r-project.org/search.html), books by folks like
Venebles and Aragon, and built-in help:

- help() opens help page
- apropos()displays all objects matching topic
- library(help=packageName) help on a specific package
- example() ; demo()
- vignette(package="packageName"); vignette(package="topic")
- RSiteSearch("packageName")

```
help(sample) ; ?sample ; ??sample
apropos("sam")
example(sample)
demo(graphics)
```