

# Introduction to R

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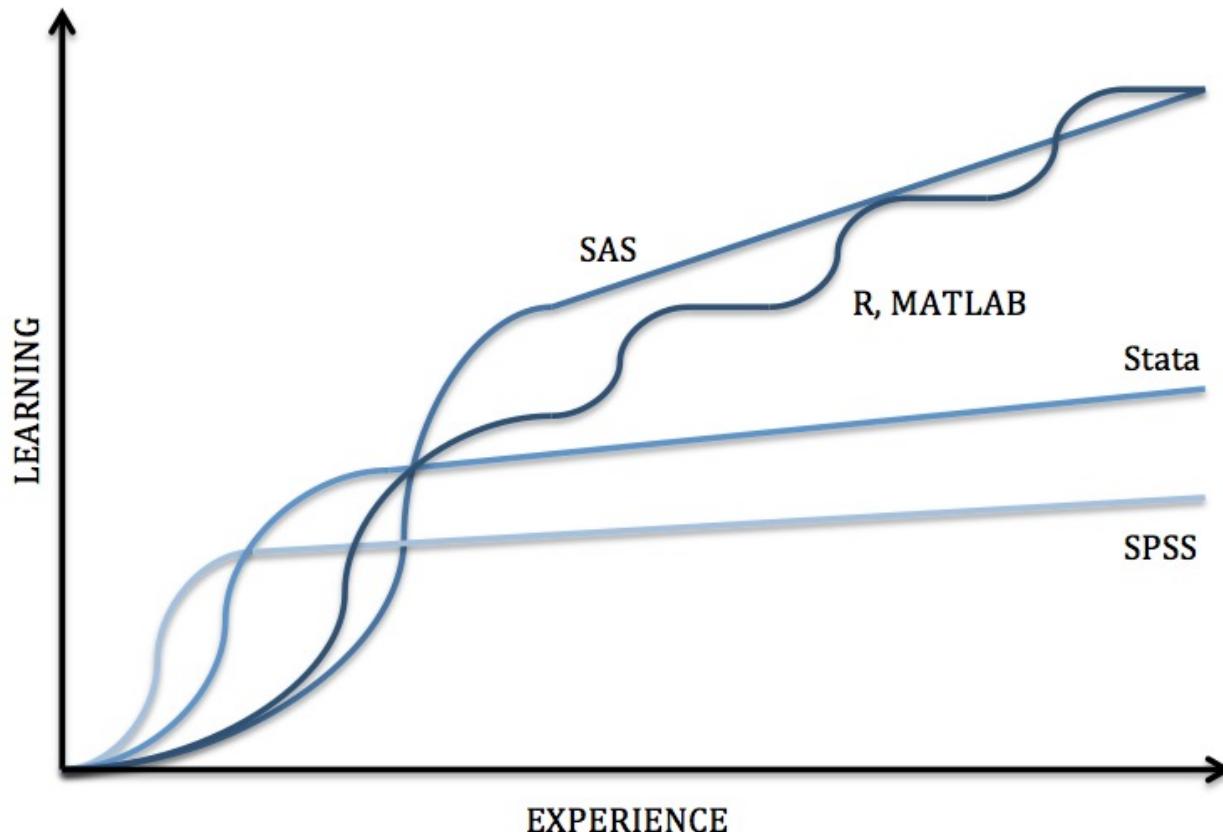
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# Outline

- Why use R?
- R Basics
- R for Database Management
  - Reading-in data, merging datasets, reshaping, recoding variables, sub-setting data, etc.
- R for Statistical Analysis
  - Descriptive and Regression Analysis
- Other topics in R
  - Tidyverse
  - Parallel Processing
  - R Studio
    - R Markdown
- Applied Example
- R Resources

# Learning Curves of Various Software Packages



Source: <https://sites.google.com/a/nyu.edu/statistical-software-guide/summary>

# Summary of Various Statistical Software Packages

Software	Interface*	Learning Curve	Data Manipulation	Statistical Analysis	Graphics	Specialties
SPSS	<b>Menus &amp; Syntax</b>	Gradual	Moderate	Moderate Scope Low Versatility	Good	Custom Tables, ANOVA & Multivariate Analysis
Stata	<b>Menus &amp; Syntax</b>	Moderate	Strong	Broad Scope Medium Versatility	Good	Panel Data, Survey Data Analysis & Multiple Imputation
SAS	Syntax	Steep	Very Strong	Very Broad Scope High Versatility	Very Good	Large Datasets, Reporting, Password Encryption & Components for Specific Fields
R	Syntax	Steep	Very Strong	Very Broad Scope High Versatility	Excellent	Packages for Graphics, Web Scraping, Machine Learning & Predictive Modeling
MATLAB	Syntax	Steep	Very Strong	Limited Scope High Versatility	Excellent	Simulations, Multidimensional Data, Image & Signal Processing

\* The primary interface is bolded in the case of multiple interface types available.

Source: <https://sites.google.com/a/nyu.edu/statistical-software-guide/summary>

# Goals of Today's Talk

- Provide an overview of the use of R for database management
  - By doing so, we can hopefully lower the learning curve of R, thereby allowing us to take advantage of its “very strong” data manipulation capabilities
- Provide an overview of the use of R for statistical analysis
  - This includes descriptive analysis (means, standard deviations, frequencies, etc.) as well as regression analysis
  - R contains a wide number of pre-canned routines that we can use to implement the method we'd like to use

# Part I

## R Basics

R Console

R version 3.2.2 (2015-08-14) -- "Fire Safety"  
Copyright (C) 2015 The R Foundation for Statistical Computing  
Platform: x86\_64-apple-darwin13.4.0 (64-bit)

R is free software and comes with ABSOLUTELY NO WARRANTY.  
You are welcome to redistribute it under certain conditions.  
Type 'license()' or 'licence()' for distribution details.

Natural language support but running in an English locale

R is a collaborative project with many contributors.  
Type 'contributors()' for more information and  
'citation()' on how to cite R or R packages in publications.

Type 'demo()' for some demos, 'help()' for on-line help, or  
'help.start()' for an HTML browser interface to help.  
Type 'q()' to quit R.

[R.app GUI 1.66 (6996) x86\_64-apple-darwin13.4.0]  
[Workspace restored from /Users/adrianrohitdass/.RData]  
[History restored from /Users/adrianrohitdass/.Rapp.history]

>

Command Window

Untitled

<functions>

Q Help search

Syntax Window

# Programming Language

- Programming language in R is generally *object oriented*
  - Roughly speaking, this means that data, variables, vectors, matrices, characters, arrays, etc. are treated as “objects” of a certain “class” that are created throughout the analysis and stored by name.
  - We then apply “methods” for certain “generic functions” to these objects
- Case sensitive (like most statistical software packages), so be careful

# Classes in R

- In R, every object has a *class*
  - For example, character variables are given the class of **factor** or **character**, whereas numeric variables are **integer**
- Classes determine how objects are handled by generic functions. For example:
  - the `mean(x)` function will work for **integers** but not for **factors** or **characters** - which generally makes sense for these types of variables

# Packages available (and loaded) in R by default

Package	Description
<code>base</code>	Base R functions (and datasets before R 2.0.0).
<code>compiler</code>	R byte code compiler (added in R 2.13.0).
<code>datasets</code>	Base R datasets (added in R 2.0.0).
<code>grDevices</code>	Graphics devices for base and grid graphics (added in R 2.0.0).
<code>graphics</code>	R functions for base graphics.
<code>grid</code>	A rewrite of the graphics layout capabilities, plus some support for interaction.
<code>methods</code>	Formally defined methods and classes for R objects, plus other programming tools, as described in the Green Book.
<code>parallel</code>	Support for parallel computation, including by forking and by sockets, and random-number generation (added in R 2.14.0).
<code>splines</code>	Regression spline functions and classes.
<code>stats</code>	R statistical functions.
<code>stats4</code>	Statistical functions using S4 classes.
<code>tcltk</code>	Interface and language bindings to Tcl/Tk GUI elements.
<code>tools</code>	Tools for package development and administration.
<code>utils</code>	R utility functions.

Source: <https://cran.r-project.org/doc/FAQ/R-FAQ.html>

For database management, we usually won't need to load or install any additional packages, although we might need the "foreign" package (available in R by default, but not initially loaded) if we're working with a dataset from another statistical program (SPSS, SAS, STATA, etc.)

# Packages in R

- Functions in R are stored in *packages*
  - For example, the function for OLS (`lm`) is accessed via the “**stats**” package, which is available in R by default
  - Only when a package is *loaded* will its contents be available. The full list of packages is not loaded by default for computational efficiency
  - Some packages in R are not installed (and thus loaded) by default, meaning that we will have to install packages that we will need beforehand, and then load them later on

# Packages in R (Continued)

- To load a package, type `library(packagename)`
  - Ex: To load the foreign package, I would type `library(foreign)` before running any routines that require this package
- To install a package in R:
  - Type `install.packages("packagename")` in command window
  - For example, the package for panel data econometrics is `plm` in R. So, to install the `plm` package, I would type `install.packages("plm")`.
    - Note that, although installed, a package will not be loaded by default (i.e. when opening R). So, you'll need `library(package)` at the top of your code (or at least sometime before the package is invoked).
  - Some packages will draw upon functions in other packages, so those packages will need to be installed as well. By using `install.packages("")`, it will automatically install dependent packages

# Some Basic Operations in R

- Q: If  $x = 5$ , and  $y = 10$ , and  $z = x + y$ , what is the value of  $z$ ?
- Let's get R to do this for us:

```
> x = 5
> y = 10
> z = x + y
> z
[1] 15
```

- In this example, we really only used the ‘+’ operator, but note that ‘-’, ‘/’, ‘\*’, ‘^’, etc. work the way they usually do for scalar operations

# Some Basic Operations in R

- Now suppose we created the following vectors:

A =	<table border="1"><tr><td>1</td></tr><tr><td>2</td></tr><tr><td>3</td></tr></table>	1	2	3	B =	<table border="1"><tr><td>2</td></tr><tr><td>4</td></tr><tr><td>6</td></tr></table>	2	4	6
1									
2									
3									
2									
4									
6									

- What is A + B?

```
> A = c(1,2,3)
> B = c(2,4,6)
> Z = A + B
> Z
[1] 3 6 9
```

In R, c() is used to combine values into a vector or list. Since we have multiple values, we need to use it here

- Note that with vectors, '+', '-', '/', '\*', '^' perform element-wise calculations when applied to vectors. So, vectors need to be the same length.

# Working with Matrices in R

- A matrix with typical element  $(i,j)$  takes the following form:

(1,1)	(1,2)	(1,3)
(2,1)	(2,2)	(2,3)
(3,1)	(3,2)	(3,3)

- Where  $i = \text{row number}$  and  $j = \text{column number}$
- In R, the general formula for extracting elements (i.e. single entry, rows, columns) is as follows:
  - `matrixname[row #, column #]`
- If we leave the terms in the brackets blank (or leave out the whole bracket term) R will spit out the whole matrix

# Working with Matrices in R (Continued)

- Example: Suppose we had the following matrix:

1	4	7
2	5	8
3	6	9

- To create this matrix in R, type:

```
> matrix = matrix(c(1, 2, 3, 4, 5, 6, 7, 8, 9), nrow=3, ncol=3)
```

- Extract the element in row #2, column #3

```
> matrix[2,3]
```

8

- Extract the second row

```
> matrix[2,]
```

2 5 8

- Extract the last two columns

```
> matrix[,c(2,3)]
```

4 7

5 8

6 9

Since we require  
multiple  
columns, we  
need to use c()  
here

# Working with Matrices in R (Continued)

- Example: Suppose now we had the following vector, with typical element 'i':

1
2
3

- Extract the third element of the vector

```
> vector[3]
```

```
3
```

- Suppose the 2<sup>nd</sup> element should be 5, not 2. How do we correct this value?

```
> vector[2] = 5
```

```
> vector
```

```
1
```

```
5
```

```
3
```

# But wait a minute...

- Q: If this is a tutorial on the use of R for database management/statistical analysis, then why are we learning about vectors/matrices?
- A: The way we work with data in R is very similar/identical to how we work with vectors/matrices
  - This is different from other statistical software packages, which may be a contributing factor to the “high” learning curve in R
- The importance of vector/matrices operations will become more clear as we move

# Part II

# R for Database Management

# Reading Data into R

What format is the data in?

- Data from Comma Separated Values File (.csv)
  - Package: `utils`
  - Formula: `read.csv(file, header = TRUE, sep = ",", quote = "\"\"", dec = ".", fill = TRUE, comment.char = "")`, ...)
- Data from Excel File (.xlsx)
  - Package: `xlsx`
  - Formula: `read.xlsx(file, sheetIndex, sheetName=NULL, rowIndex=NULL, startRow=NULL, endRow=NULL, colIndex=NULL, as.data.frame=TRUE, header=TRUE, colClasses=NA, keepFormulas=FALSE, encoding="unknown", ...)`
- Data from STATA (.dta)
  - Package: `foreign`
  - `read.dta(file, convert.dates = TRUE, convert.factors = TRUE, missing.type = FALSE, convert.underscore = FALSE, warn.missing.labels = TRUE)`

Other Formats: See package “`foreign`”

<https://cran.r-project.org/web/packages/foreign/foreign.pdf>

# Reading Data into R

Examples:

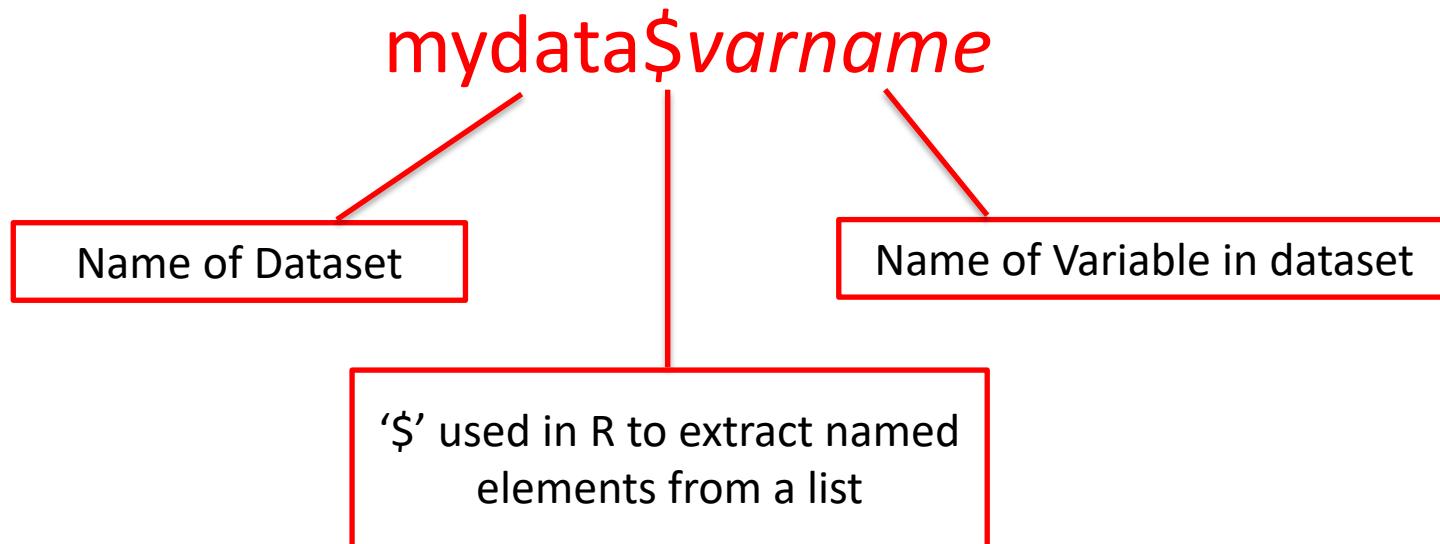
- CSV file with variable names at top
  - `data = read.csv("C:/Users/adrianrohitdass/Documents/R Tutorial/data.csv")`
- CSV file with no variable names at top
  - `data = read.csv("C:/Users/adrianrohitdass/Documents/R Tutorial/data.csv", header=F)`
- STATA data file (12 or older)
  - `library(foreign)`
  - `data = read.dta("C:/Users/adrianrohitdass/Documents/R Tutorial/data.dta")`
- STATA data file (13 or newer)
  - `library(readstata13)`
  - `data = read.dta13("C:/Users/adrianrohitdass/Documents/R Tutorial/data.dta")`

# Comparison and Logical Operators

Operator	Description	Example
=	Assign a value	x = 5
==	Equal to	sex ==1
!=	Not equal to	LHIN != 5
>	Greater than	income >5000
<	Less than	healthcost < 5000
>= or <=	Greater than or equal to Less than or equal to	income >= 5000 healthcost <= 5000
&	And	sex==1 & age>50
	Or	LHIN==1   LHIN ==5

# Referring to Variables in a Dataset

- Suppose I had data stored in “mydata” (i.e an object created to store the data read-in from a .csv by R). To refer to a specific variable in the dataset, I could type



# Creating a new variable/object

- No specific command to generate new variables (in contrast to STATA’s “gen” and “egen” commands)
  - `x = 5` generates a 1x1 scalar called “x” that is equal to 5
  - `data$age = year - data$dob` creates a new variable “age” in the dataset “data” that is equal to the year – the person’s date of birth (let’s say in years)

# Looking at Data

- Display the first or last few entries of a dataset:
  - Package: **utils**
  - View entire dataset in separate window
    - `View(x, title)`
  - First few elements of dataset (default is 5):
    - `head(x, n, ...)`
  - Last few elements of dataset (default is 5):
    - `tail(x, n, ...)`
- List of column names in dataset
  - Package: **base**
  - Formula: `colnames(x)`

# Missing Values

Missing Values are listed as “NA” in R

- Count number of NA's in column  
 $\text{sum(is.na(x))}$
- Recode Certain Values as NA (i.e. non responses coded as -1)  
 $x[x == -1] = \text{NA}$

# Renaming Variables (Columns)

A few different ways to do this:

- To rename the ‘ith’ column in a dataset
  - `colnames(data)[i] = “My Column Name”`
- Can be cumbersome – especially if don’t know column # of the column you want to rename (just it’s original name)
- Alternative:
  - `colnames(data)[which(colnames(data) == “R1482600”)] = “race”`

Grabs column names  
from specified  
dataset

Look-up that returns  
the column #

New column name

# Subsetting Data

- Subsetting can be used to restrict the sample in the dataset, create a smaller data with fewer variables, or both
- Recall: extracting elements from a matrix in R
  - `matrixname[row #, column #]`
- What's the difference between a matrix and a dataset?
  - Both have row elements
    - Typically the individual records in a dataset
  - Both have column elements
    - Typically the different variables in the dataset
- If we think of our dataset as a matrix, then the concept of subsetting in R becomes a lot easier to digest

# Subsetting Data (Continued)

Examples:

- Restrict sample to those with age  $\geq 50$   
`> datas1 = data[data$age >=50,]`
- Create a smaller dataset with just ID, age, and height  
`> datas2 = data[, c("ID", "age", "height")]`
- Create a smaller dataset with just ID, age, and height; with age  $\geq 50$   
`> datas3 = data[data$age>=50, c("ID", "age", "height")]`

# Recoding Variables in R

- Usually done with a few lines of code using comparison and logical operators
- Ex: Suppose we had the following for age:  
`> data$age = [19, 20, 25, 30, 45, 55]`
- If we wanted to create a categorical variable for age (say, <20, 20-39, 40-59), we could do the following:  
`> data$agecat[data$age <20] = 1  
> data$agecat[data$age >=20 & data$age <40] = 2  
> data$agecat[data$age >=40 & data$age <60] = 3  
> data$agecat  
> [1, 2, 2, 2, 3, 3]`

# Merging Datasets

Suppose we had the following 2 datasets:

Data1			Data2	
Id	Age	Income	Id	Health Care Cost
1	55	49841.65	1	188.1965
2	63	46884.78	2	172.2420
3	65	45550.87	3	102.8355
4	69	26254.15	4	150.2247
5	52	22044.73		

Our first dataset contains some data on age and income, but not health care costs to the public system. Dataset 2 contains this data, but was not initially available to us. It also doesn't have age or income.

The common element between the two datasets is "Id", which uniquely identifies the same individuals across the two datasets.

Note that, for some reason, individual 5 does not have a reported health care cost

# Merging Datasets (Continued)

- Command: merge
  - Package: **base**
- For our example:
  - **Datam = merge(Data1, Data2, by="Id", all=T)**
  - Resulting Dataset

Unique identifier  
across datasets

Optional, but  
default is F,  
meaning those  
who can't be  
matched will be  
excluded

Datam			
Id	Age	Income	Health Care Cost
1	55	49841.65	188.1965
2	63	46884.78	172.2420
3	65	45550.87	102.8355
4	69	26254.15	150.2247
5	52	22044.73	NA

## Part II

# R for Statistical Analysis

# Descriptive Statistics in R

- Mean
  - Package: **base**
  - Formula: **mean(x, trim = 0, na.rm = FALSE, ...)**
- Standard Deviation
  - Package: **stats**
  - Formula: **sd(x, na.rm = FALSE)**
- Correlation
  - Package: **stats**
  - Formula: **cor(x, y = NULL, use = "everything", method = c("pearson", "kendall", "spearman"))**

# Descriptive Statistics (Example)

- Suppose we had the following data column in R (transposed to fit on slide):
  - Vector = [5,5,6,4]
- What is the mean of the vector?
- In R, I would type
  - > `mean(Vector)`
  - > 5

# Descriptive Statistics (Example)

- Suppose now we had the following:
  - Vector = [5,5,6,4,NA]
- What is the mean of the vector?
- In R, I would type
  - > `mean(Vector)`
  - > `NA`
- Why did I get a mean of NA?
  - Our vector included a missing value, so R couldn't compute the mean as is.
- To remedy this, I would type
  - > `mean(Vector, na.rm=T)`
  - > `5`

# Tabulations R

- Tabulations of categorical/ordinal variables can be done with R's *table* command:
  - Package: **base**
  - Formula: **table(..., exclude = if (useNA == "no") c(NA, NaN), useNA = c("no", "ifany", "always"), dnn = list.names(...), deparse.level = 1)**

Ex: Table Sex Variable, with extra column for missing values (if any)

```
> mytable = table(pdata$sex, exclude=NULL)
> mytable
```

	Female	Male	<NA>
	17540	18396	0

# Graphing Data in R

- Generic X-Y Plotting
  - Package: `graphics`
  - Formula: `plot(x, y, ...)`

Example:

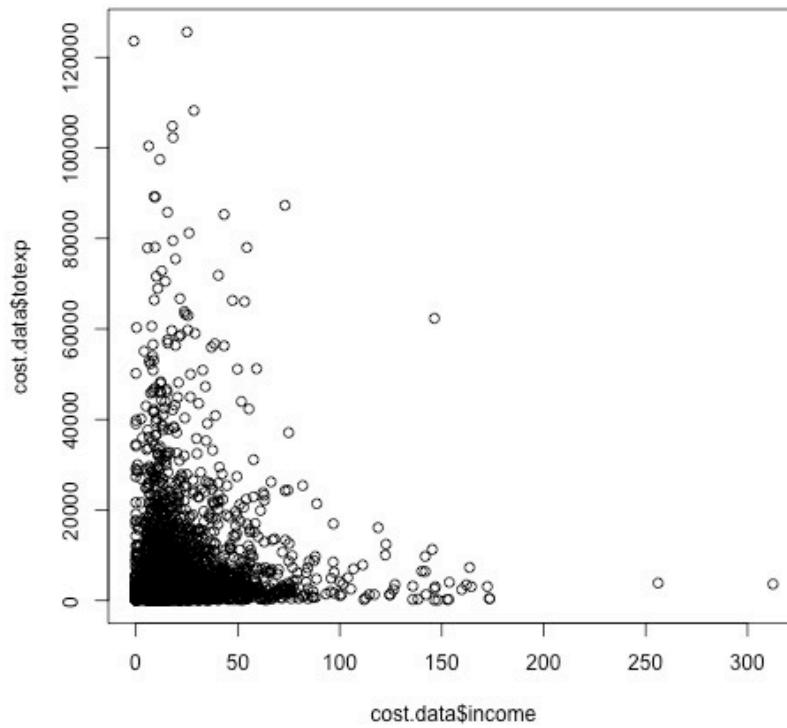
```
plot(cost.data$income,cost.data$totexp)
```

- Plotting with `ggplot()` function
  - Package: `ggplot2`
  - Formula: `ggplot(data = NULL, mapping = aes(), ..., environment = parent.frame())`

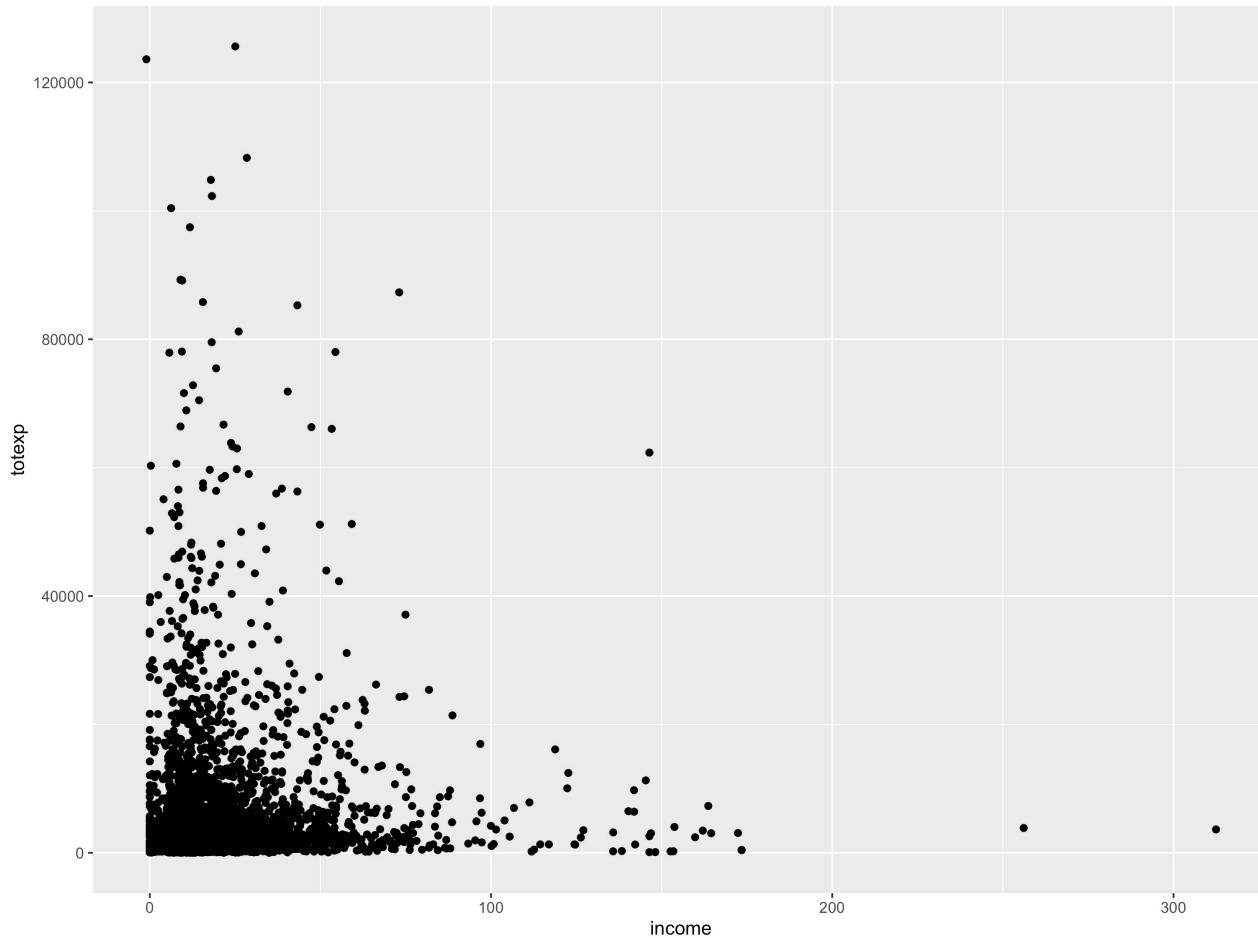
Example:

```
ggplot(cost.data, aes(x=income, y=totexp)) + geom_point()
```

# Resulting Graph (Generic)



# Resulting Graph (ggplot2)



See <https://github.com/rstudio/cheatsheets/raw/master/data-visualization.pdf> for ggplot cheatsheet

# Ordinary Least Squares

- The estimator of the regression intercept and slope(s) that minimizes the sum of squared residuals (Stock and Watson, 2007).
  - Package: `stats`
  - Formula: `lm(formula, data, subset, weights, na.action, method = "qr", model = TRUE, x = FALSE, y = FALSE, qr = TRUE, singular.ok = TRUE, contrasts = NULL, offset, ...)`

Examples:

Regression of “total health care expenditure” on “age, gender, household income, supplementary insurance status (insurance beyond Medicare), physical and activity limitations and the total number of chronic conditions” using dataset “cost.data” from Medical Expenditure Panel Survey (65+)

```
ols.costdata = lm(totexp ~ age + female + income + suppins + phylim + actlim + totchr,  
data = cost.data)
```

Online Help File

<https://stat.ethz.ch/R-manual/R-devel/library/stats/html/lm.html>

# Ordinary Least Squares

```
> ols.costdata = lm(totexp ~ age + female + income + suppins + phylin + actlim + totchr, data = cost.data)
> summary(ols.costdata)
```

Call:

```
lm(formula = totexp ~ age + female + income + suppins + phylin +
    actlim + totchr, data = cost.data)
```

Residuals:

Min	1Q	Median	3Q	Max
-17311	-5000	-2318	716	113095

Coefficients:

	Estimate	Std. Error	t value	Pr(> t )
(Intercept)	8358.954	2597.715	3.218	0.00131 **
age	-85.363	34.317	-2.487	0.01292 *
female	-1383.290	427.485	-3.236	0.00123 **
income	6.469	9.568	0.676	0.49904
suppins	724.863	433.889	1.671	0.09490 .
phylin	2389.019	534.738	4.468	8.21e-06 ***
actlim	3900.491	582.991	6.690	2.65e-11 ***
totchr	1844.377	172.919	10.666	< 2e-16 ***

---

Signif. codes: 0 ‘\*\*\*’ 0.001 ‘\*\*’ 0.01 ‘\*’ 0.05 ‘.’ 0.1 ‘ ’ 1

Residual standard error: 11290 on 2947 degrees of freedom

Multiple R-squared: 0.1163, Adjusted R-squared: 0.1142

F-statistic: 55.42 on 7 and 2947 DF, p-value: < 2.2e-16

Example adapted from Jones (2013) *Applied Health Economics*

# Post-Estimation

Package: `lmtest`

- Breusch-Pagan test for heteroskedasticity.

`bptest(formula, varformula = NULL, studentize = TRUE, data = list())`

- Ramsey's RESET test for functional form.

`resettest(formula, power = 2:3, type = c("fitted", "regressor", "princomp"), data = list())`

Package: `car`

- Variance Inflation Factor (VIF)

`vif(model)`

Package: `sandwich`

- Heteroskedasticity-Consistent Covariance Matrix Estimation

`coeftest(ols.costdata, vcovHC(ols costdata, type = "HC1"))`

Notes: need to combine with `lmtest coeftest()` command, and use `type = "HC1"` to get the same results as STATA's "robust" command

# Extracting Beta coefficients, standard errors, etc. from model

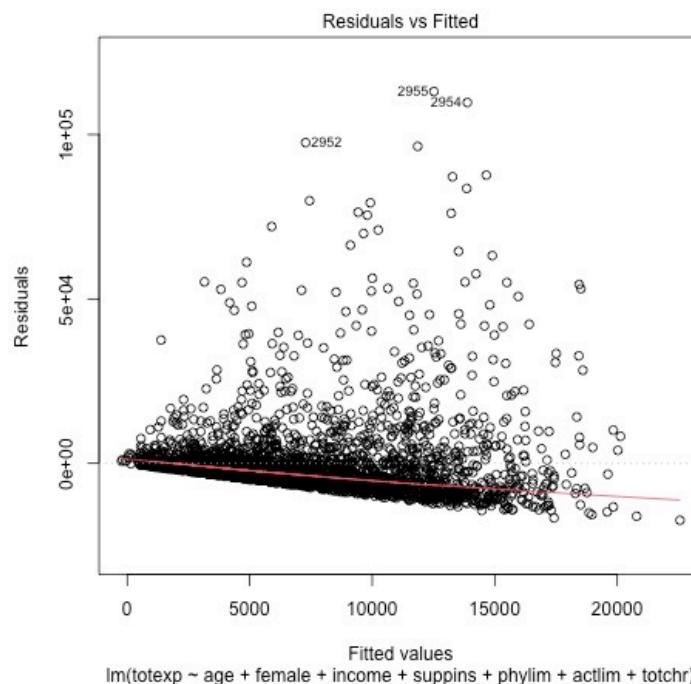
- A couple of ways to do this, but most of the information we're after is stored in the coefficients object returned from summary:

```
> summary(ols.costdata)$coefficients
            Estimate Std. Error    t value   Pr(>|t|)
(Intercept) 8358.95394 2597.71486  3.2178104 1.305733e-03
age          -85.36264   34.31701 -2.4874733 1.292031e-02
female       -1383.28982  427.48537 -3.2358764 1.226119e-03
income        6.46894    9.56821  0.6760867 4.990386e-01
suppins      724.86321  433.88874  1.6706200 9.490295e-02
phylim       2389.01859  534.73836  4.4676402 8.206489e-06
actlim       3900.49083  582.99135  6.6904781 2.651802e-11
totchr       1844.37687  172.91874 10.6661482 4.356843e-26
```

- The above is a matrix, so we can get the information we need through column extractions:
  - Beta coefficients: `summary(ols.costdata)$coefficients[,1]`
  - Standard errors: `summary(ols.costdata)$coefficients[,2]`
  - T-value: `summary(ols.costdata)$coefficients[,3]`
  - P-value: `summary(ols.costdata)$coefficients[,4]`

# Residuals vs Fitted Values

- For Residuals vs Fitted Values (RVFV) Plot, use generic `plot()` function on regression object. First plot is RVFV
- Formula: `plot(ols.costdata, 1)`



\*The other 5 plots are: Normal Q-Q, Scale-Location, Cook's distance, Residuals vs Leverage, and Cook's distance vs Leverage

# Models for Binary Outcomes

- R does not come with different programs for binary outcomes. Instead, it utilizes a unifying framework of generalized linear models (GLMs) and a single fitting function, `glm()` (Kleiber & Zeileis (2008))

Package: `stats`

Formula: `glm(formula, family = gaussian, data, weights, subset, na.action, start = NULL, etastart, mustart, offset, control = list(...), model = TRUE, method = "glm.fit", x = FALSE, y = TRUE, contrasts = NULL, ...)`

- For binary outcomes, we specify `family="binomial"` and `link= "logit"` or `"probit"`
- Can be extended to count data as well (`family="poisson"`)

Online help: <https://stat.ethz.ch/R-manual/R-devel/library/stats/html/glm.html>

# Models for Binary Outcomes

Example: Probit Analysis: factors associated with being arrested

```
> probit = glm(arrestbin~age + male, data = subdata, family="binomial"(link="probit"))
> summary(probit)

Call:
glm(formula = arrestbin ~ age + male, family = binomial(link = "probit"),
     data = subdata)

Deviance Residuals:
    Min      1Q  Median      3Q      Max 
-0.5115 -0.4497 -0.3339 -0.2652  2.6550 

Coefficients:
            Estimate Std. Error z value Pr(>|z|)    
(Intercept) -2.88106   0.24974 -11.536 < 2e-16 ***
age          0.07088   0.01527   4.641 3.46e-06 ***
male         0.44323   0.04422  10.023 < 2e-16 ***
---
Signif. codes:  0 ‘***’ 0.001 ‘**’ 0.01 ‘*’ 0.05 ‘.’ 0.1 ‘ ’ 1

(Dispersion parameter for binomial family taken to be 1)

Null deviance: 4214.0 on 8360 degrees of freedom
Residual deviance: 4087.5 on 8358 degrees of freedom
(623 observations deleted due to missingness)
AIC: 4093.5

Number of Fisher Scoring iterations: 5
```

# Instrumental Variables

A way to obtain a consistent estimator of the unknown coefficients of the population regression function when the regressor,  $X$ , is correlated with the error term,  $u$ . (Stock and Watson, 2007).

Package: [AER](#)

Formula: `ivreg(formula, instruments, data, subset, na.action, weights, offset, contrasts = NULL, model = TRUE, y = TRUE, x = FALSE, ...)`

Online documentation: <https://cran.r-project.org/web/packages/AER/AER.pdf>

# IV Example

Example: Determinants of Income (As a function of Health)

```
> require(AER)
> iv = ivreg(Income ~ Health + Age | ParentHealth + Age)
> summary(iv, diagnostics = TRUE)

Call:
ivreg(formula = Income ~ Health + Age | ParentHealth + Age)

Residuals:
    Min      1Q  Median      3Q     Max 
-3.1557 -0.6261  0.0130  0.6495  2.8700 

Coefficients:
            Estimate Std. Error t value Pr(>|t|)    
(Intercept) 2.03965   0.06817  29.92   <2e-16 ***
Health       0.99773   0.01186  84.16   <2e-16 ***
Age          2.00177   0.07256  27.59   <2e-16 ***

Diagnostic tests:
                    df1 df2 statistic p-value    
Weak instruments  1  997      1427   <2e-16 ***
Wu-Hausman       1  996      2271   <2e-16 *** 
Sargan           0   NA        NA      NA      
---
Signif. codes:  0 '****' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

Residual standard error: 0.9742 on 997 degrees of freedom
Multiple R-Squared: 0.9573, Adjusted R-squared: 0.9572
Wald test: 1.067e+04 on 2 and 997 DF,  p-value: < 2.2e-16
```

Prints out F-test for Weak Instruments, Hausman Test Statistic (vs ols) and Sargan's Test for Over-identifying Restrictions (if more than one instrument use)

# Other Regression Models

- Panel Data Econometrics
  - Package: **plm**
  - <https://cran.r-project.org/web/packages/plm/vignettes/plm.pdf>
- Linear and Generalized Linear Mixed Effects Models
  - Package: **lme4**
  - <https://cran.r-project.org/web/packages/lme4/lme4.pdf>
- Quantile Regression
  - Package: **quantreg**
  - <https://cran.r-project.org/web/packages/quantreg/quantreg.pdf>

# Part III

## Other topics in R

# Tidyverse

# Tidyverse

From Tidyverse website:

“The tidyverse is an opinionated collection of R packages designed for data science. All packages share an underlying design philosophy, grammar, and data structures...tidyverse makes data science faster, easier and more fun”

Source: <https://www.tidyverse.org>

- Packages within tidyverse: **ggplot2, dplyr, tidyr, readr, purrr, tibble, stringr, andforcats**
- To get, type: **install.packages("tidyverse")** in R console

# Tidyverse (Continued)

## Package: `dplyr`

- Description: provides a flexible grammar of data manipulation.
- Example Commands:
  - Restrict sample to those with age  $\geq 50$ 
    - `subdata = filter(data, age >= 50)`
  - Create a smaller dataset with just ID, age, and height
    - `subdata = select(data, ID, age, height)`
  - Create a smaller dataset with just ID, age, and height;  
with age  $\geq 50$ 
    - `subdata = data %>%  
filter(age >= 50) %>%  
select(ID, age, height)`

# Tidyverse (Continued)

## Package: **dplyr**

- Example Commands (continued):
  - Create new variable (age) in existing dataset
    - `data = mutate(data, age = year - dob)`
  - Rename a variable in a dataset (new name = old name)
    - `data = rename(data, race = R1482600)`
- <https://cran.r-project.org/web/packages/dplyr/dplyr.pdf>

# Tidyverse (Continued)

Other (selected) packages in Tidyverse:

- Package: **readr**
  - Description: The goal of 'readr' is to provide a fast and friendly way to read rectangular data (like 'csv', 'tsv', and 'fwf')
  - <https://cran.r-project.org/web/packages/readr/readr.pdf>
- Package: **tidyr**
  - Description: Tools for reshaping data, extracting values out of string columns, and working with missing values
  - <https://cran.r-project.org/web/packages/tidyr/tidyr.pdf>

# Parallel Processing

# Parallel Processing in R

- Parallel computing: From Wikipedia: “Parallel computing is a type of computation in which many calculations or the execution of processes are carried out simultaneously. Large problems can often be divided into smaller ones, which can then be solved at the same time.”
  - See here for more:  
[https://en.wikipedia.org/wiki/Parallel\\_computing](https://en.wikipedia.org/wiki/Parallel_computing)
- Modern day computers typically contain:
  - Single-core
  - Multicore (Dual, Quad, Hexa, Octo, etc.)
- May also contain hyperthreading

# Parallel Processing in R (Continued)

- Parallel processing can be used in many situations, including:
  - Bootstrapping
  - Microsimulation models
  - Monte Carlo experiments
  - Probabilistic Sensitivity Analysis
- By utilizing parallel processing, we can significantly speed up the processing time of our calculations

# Parallel Processing in R (Continued)

- There are many packages to perform parallel processing in R, including
- **parallel**
  - Available in R by default
  - Handles large chunks of computations in parallel
  - <https://stat.ethz.ch/R-manual/R-devel/library/parallel/doc/parallel.pdf>
- **doParallel**
  - “parallel backend” for the “foreach” package
  - provides a mechanism needed to execute foreach loops in parallel
  - <https://cran.r-project.org/web/packages/doParallel/vignettes/gettingstartedParallel.pdf>

# Example: Monte Carlo Experiment

```
> RNGkind("L'Ecuyer-CMRG") #Special type of seed for parallel processing
> set.seed(12345) #Set the seed
> require(doParallel) #Load "doParallel" package
> b0 = 1 #True value on constant
> b1 = 2 #True value on X1
> b2 = 3 #True value on X2
> n = 10000 #Sample size
> S = 50000 #Number of simulations
>
> ncores = 2 #Set as appropriate depending on your hardware
> registerDoParallel(cores=ncores) # Shows the number of Parallel Workers to be used
>
> ###Parallel Processing###
> p = Sys.time()
> olsmparresults = foreach(i=1:S, .combine='cbind', .multicombine=TRUE) %dopar%
+ {
+   x1 = rnorm(n, mean = 1, sd = 1)
+   x2 = rnorm(n, mean = 1, sd = 1)
+   e = rnorm(n, mean = 0, sd = 1)
+   y = b0 + b1*x1 + b2*x2 + e
+   data = data.frame(cbind(y, x1, x2))
+
+   ols = lm(y~x1 + x2, data = data)
+   betahatols = coefficients(ols)
+ }
> comp.time = Sys.time() - p
> comp.time
Time difference of 2.668144 mins
```

# Example: Monte Carlo Experiment (Continued)

```
> ###Run everything through 1 core (for comparison purposes)###
> p = Sys.time()
> olsmcresults = foreach(i=1:S, .combine='cbind', .multicombine=TRUE) %do%
+ {
+   x1 = rnorm(n, mean = 1, sd = 1)
+   x2 = rnorm(n, mean = 1, sd = 1)
+   e = rnorm(n, mean = 0, sd = 1)
+   y = b0 + b1*x1 + b2*x2 + e
+   data = data.frame(cbind(y, x1, x2))
+
+   ols = lm(y~x1 + x2, data = data)
+   betahatols = coefficients(ols)
+ }
> comp.time = Sys.time() - p
> comp.time
Time difference of 4.718112 mins
```

Notice we changed %dopar%  
to %do% to run everything  
through a single core

# R Studio

# What is R Studio?

From R Studio Website:

- An integrated development environment (IDE) for R.  
Includes:
  - A console
  - Syntax highlighting editor
  - Tools for plotting, history, debugging, and workspace history
- Can think of it as a more user friendly version of R
- A free version is available as well
- For more information, see <https://www.rstudio.com>

RStudio

**Syntax Window**

```
olsgmmfunction.R *
1 set.seed(198901)
2 require(gmm)
3 b0 = 0.5
4 b1 = 1
5 b2 = -1
6 b3 = 1
7 n = 100000
8 nrs = 10
9 S = 1
10 b0ols = rep(0,S)
11 b1ols = rep(0,S)
12 b2ols = rep(0,S)
13 b3ols = rep(0,S)
14 b0gmmm1 = rep(0,S)
15 b1gmmm1 = rep(0,S)
16 b2gmmm1 = rep(0,S)
17 b3gmmm1 = rep(0,S)
18 b0elm1 = rep(0,S)
19 b1elm1 = rep(0,S)
20 b2elm1 = rep(0,S)
21 b3elm1 = rep(0,S)
22
```

1:17 (Top Level) ▾

**Environment** History

**List of datasets/variables**

dat	num [1:10, 1:5] -2.55 -1.06 -3.29 -1.3 -1.03 ...
data	num [1:10, 1:5] -2.55 -1.06 -3.29 -1.3 -1.03 ...
data2	1000 obs. of 2 variables
datanrsp	Large matrix (204940 elements, 1.6 Mb)
datapop	Large matrix (500000 elements, 3.8 Mb)
results	num [1:11, 1:8] -1.07 0 0 0 0 ...
results1	num [1, 1:8] -1.073 NA 0.998 NA -0.17 ...
results10	num [1, 1:8] 0 NA 0 NA 0 NA 0 NA
results11	num [1, 1:8] 0 NA 0 NA 0 NA 0 NA

Files Plots Packages Help Viewer

R: Generalized method of moment estimation ▾ Find in Topic

**Files, plots, packages, help, and viewer**

**Generalized method of moment estimation**

**Description**

Function to estimate a vector of parameters based on moment conditions using the GMM method of Hansen(82).

**Usage**

```
gmm(g, x, t0=NULL, gradv=NULL, type=c("twoStep", "cue", "iterative"),
wmatrix = c("optimal", "ident"), vcov=c("HAC", "MDS", "iid", "True
kernel=c("Quadratic Spectral", "Truncated", "Bartlett", "Parzen
crit=10e-7, bw = bwAndrews, prewhite = 1, ar.method = "ols", ap
tol = 1e-7, itermax=100, optfct=c("optim", "optimize", "nlminb",
model=TRUE, X=FALSE, Y=FALSE, TypeGmm = "baseGmm", centeredVco
weightsMatrix = NULL, traceIter = FALSE, data, eqConst = NULL,
eqConstFullVcov = FALSE, ...)
evalGmm(g, x, t0, tetw=NULL, gradv=NULL, wmatrix = c("optimal", "id
vcov=c("HAC", "iid", "TrueFixed"), kernel=c("Quadratic Spectral"
"Bartlett", "Parzen", "Tukey-Hanning"), crit=10e-7, bw = bwAndre
prewhite = FALSE, ar.method = "ols", approx="AR(1)".tol = 1e-7
```

**Command/Results Window**

```
Console ~ / 
> results3 = rbind(mean(b0gmmm2),sd(b0gmmm2),mean(b1gmmm2),sd(b1gmmm2),mean(b2gmmm2),sd(b2gmmm2),mean(b3gmmm2),sd(b3gmmm2))
> results4 = cbind(mean(b0gmmm3),sd(b0gmmm3),mean(b1gmmm3),sd(b1gmmm3),mean(b2gmmm3),sd(b2gmmm3),mean(b3gmmm3),sd(b3gmmm3))
> results5 = cbind(mean(b0gmmm4),sd(b0gmmm4),mean(b1gmmm4),sd(b1gmmm4),mean(b2gmmm4),sd(b2gmmm4),mean(b3gmmm4),sd(b3gmmm4))
> results6 = cbind(mean(b0gmmm5),sd(b0gmmm5),mean(b1gmmm5),sd(b1gmmm5),mean(b2gmmm5),sd(b2gmmm5),mean(b3gmmm5),sd(b3gmmm5))
> results7 = cbind(mean(b0elm1),sd(b0elm1),mean(b1elm1),sd(b1elm1),mean(b2elm1),sd(b2elm1),mean(b3elm1),sd(b3elm1))
> results8 = cbind(mean(b0elm2),sd(b0elm2),mean(b1elm2),sd(b1elm2),mean(b2elm2),sd(b2elm2),mean(b3elm2),sd(b3elm2))
> results9 = cbind(mean(b0elm3),sd(b0elm3),mean(b1elm3),sd(b1elm3),mean(b2elm3),sd(b2elm3),mean(b3elm3),sd(b3elm3))
> results10 = cbind(mean(b0elm4),sd(b0elm4),mean(b1elm4),sd(b1elm4),mean(b2elm4),sd(b2elm4),mean(b3elm4),sd(b3elm4))
> results11 = cbind(mean(b0elm5),sd(b0elm5),mean(b1elm5),sd(b1elm5),mean(b2elm5),sd(b2elm5),mean(b3elm5),sd(b3elm5))
>
> results = rbind(results1, results2, results3, results4, results5, results6, results7, results8, results9, results10,
results11)
> |
```



# R Markdown (In R Studio)

# What is R Markdown?

From R Markdown website:

“R Markdown provides an authoring framework for data science. You can use a single R Markdown file to both

- save and execute code
- generate high quality reports that can be shared with an audience”

Source: <https://rmarkdown.rstudio.com/lesson-1.html>

With R Markdown, you can render to a variety of formats, which includes PDF (uses [LaTeX](#)) and Microsoft Word

To create a R Markdown file, go to File → New File → R Markdown



```
1 ---  
2 title: "Untitled"  
3 output: word_document  
4 ---  
5  
6 ```{r setup, include=FALSE}  
7 knitr::opts_chunk$set(echo = TRUE)  
8 ````  
9  
10 ## R Markdown  
11  
12 This is an R Markdown document. Markdown is a simple formatting syntax for authoring HTML, PDF, and MS Word documents. For more details on using R Markdown see  
http://rmarkdown.rstudio.com.  
13  
14 When you click the **Knit** button a document will be generated that includes both content as well as the output of any embedded R code chunks within the document. You can  
embed an R code chunk like this:  
15  
16 ```{r cars}  
17 summary(cars)  
18 ````  
19  
20 ## Including Plots  
21  
22 You can also embed plots, for example:  
23  
24 ```{r pressure, echo=FALSE}  
25 plot(pressure)  
26 ````  
27  
28 Note that the `echo = FALSE` parameter was added to the code chunk to prevent printing of the R code that generated the plot.  
29
```

“Knit”, or generate document

Global options for document here (echoing of R code, loading packages, etc.)

# for Document Sections

R code chunk for output (summary of “cars” data)

R code chunk for output (to insert a plot available in R memory)

# Page 1 (of 2)

The screenshot shows a Microsoft Word document titled "Untitled". The document is an R Markdown file. It contains the following content:

This is an R Markdown document. Markdown is a simple formatting syntax for authoring HTML, PDF, and MS Word documents. For more details on using R Markdown see <http://rmarkdown.rstudio.com>.

When you click the Knit button a document will be generated that includes both content as well as the output of any embedded R code chunks within the document. You can embed an R code chunk like this:

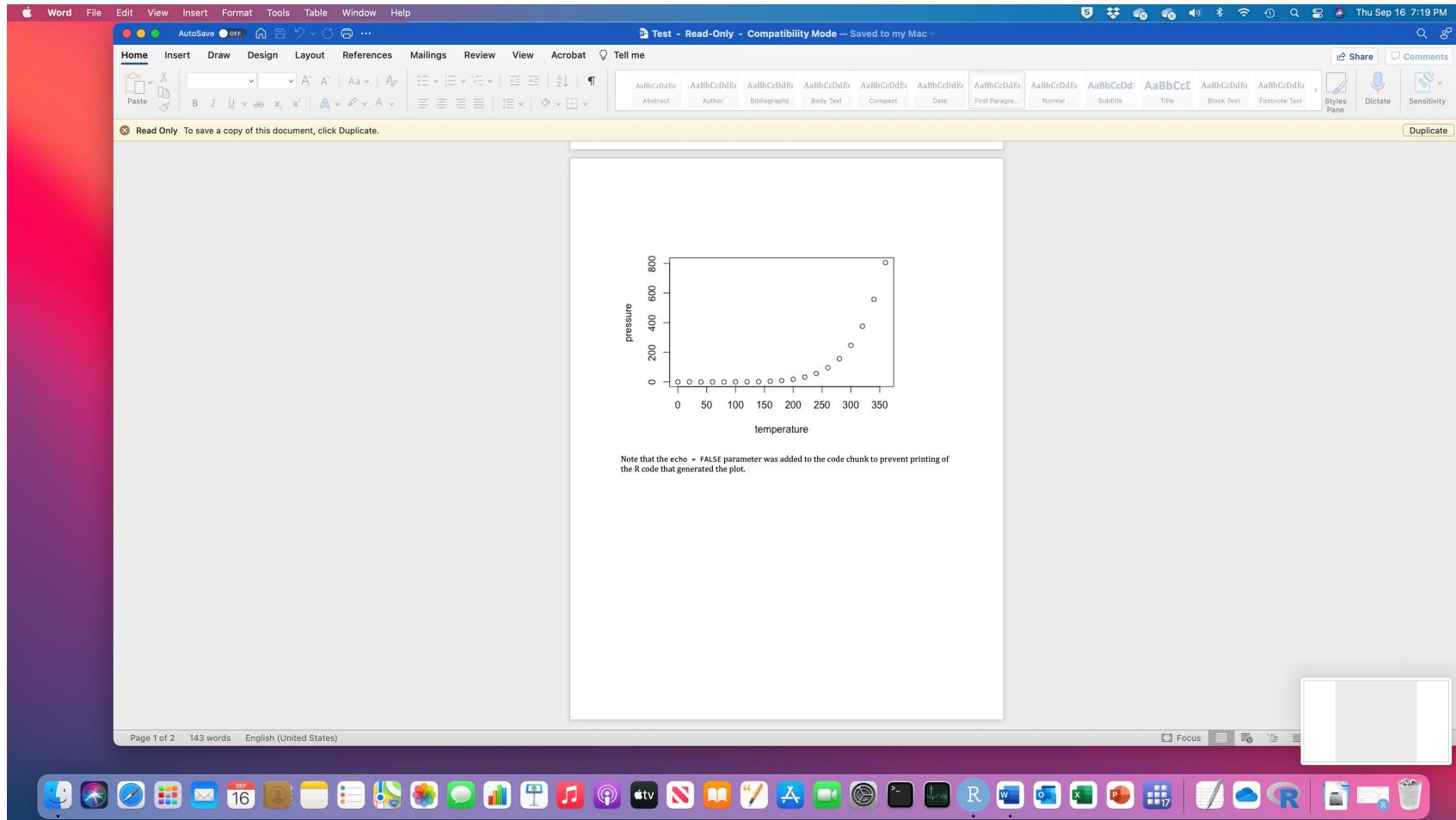
```
summary(cars)
```

## speed dist  
## Min. :4.0 Min. : 2.00  
## 1st Qu.:12.0 1st Qu.: 26.00  
## Median :15.0 Median : 36.00  
## Mean :15.4 Mean : 42.98  
## 3rd Qu.:19.0 3rd Qu.: 56.00  
## Max. :25.0 Max. :120.00

**Including Plots**

You can also embed plots, for example:

Page 2 (of 2)



# Tips for Outputting In MS Word

<b>Output Option</b>	<ul style="list-style-type: none"><li>The <code>word_document2</code> (<a href="#">Bookdown</a>) and <code>rdocx_document</code> (<a href="#">Officedown</a>) formats are generally superior to <code>word_document</code> (default in R Markdown), particularly for automatic numbering of figures/tables, and cross-referencing of figures/tables.</li><li>The <code>rdocx_document</code> lets you easily switch between landscape and portrait</li></ul>
<b>Tables</b>	Default <code>knitr::kable()</code> function works, but <code>flextable()</code> function <a href="#">flextable</a> creates “pretty” tables with a large amount of flexibility (customize cell padding and column widths, table footnotes, long tables, etc.)
<b>Figures</b>	Use <code>knitr::include_graphics(filepath)</code> for previously saved figures to include in the document
<b>References</b>	<ul style="list-style-type: none"><li>Default reference style is Chicago. Visit <a href="#">Zotero Style Repository</a> to search for additional Citation Style Language (CSL) files (Vancouver, APA, journal specific styles, etc.). Can modify existing reference style, which may be necessary for certain journals (<a href="https://editor.citationstyles.org/about/">https://editor.citationstyles.org/about/</a>)</li><li>Add citations with markdown syntax by typing <code>[@cite]</code> or <code>@cite</code>.</li><li>Store references in plain text BibTeX database (*.bib)</li><li>Can also look up and Insert Citations dialog in the Visual Editor by clicking the <code>@</code> symbol in the toolbar or by clicking Insert &gt; Citation</li></ul>
<b>Document formatting</b>	To modify font sizes, text alignment, etc., need to create a reference style document following these instructions: <a href="https://rmarkdown.rstudio.com/articles_docx.html">https://rmarkdown.rstudio.com/articles_docx.html</a>

Please also see the R Markdown cheat sheet:

<https://github.com/rstudio/cheatsheets/raw/master/rmarkdown-2.0.pdf>

# Applied Example

- Analysis of Health Expenditure Data in Jones et al. (2013) *Chapter Three*
- The data covers the medical expenditures of US citizens aged 65 years and older who qualify for health care under Medicare.
  - Outcome of interest is total annual health care expenditures (measured in US dollars).
  - Other key variables are age, gender, household income, supplementary insurance status (insurance beyond Medicare), physical and activity limitations and the total number of chronic conditions.
- Data can be downloaded from here (mus03data.dta):  
<https://www.stata-press.com/data/musr.html>

# R Markdown Code From Example

```
---
```

```
title: "Untitled"
output: word_document
---
```

```
```{r setup, include=FALSE}
knitr::opts_chunk$set(echo = FALSE)
```

# Regression Results

```{r regresults}
load("cost.data.results.RData")
knitr::kable(cost.data.results)
```

# Plot

```{r plot}
knitr::include_graphics("RVFV.jpg")
```
```

# Conclusions

- R has extremely powerful database management capabilities
  - Is fully capable of performing the same sort of tasks as commercial software programs
  - Can be enhanced through Tidyverse package for a more user friendly experience
- R is very capable of statistical analysis
  - Is fully capable of calculating summary statistics and performing regression analysis right out of the box
  - Can install additional packages to perform other sorts of analysis, depending on the research question of the user
  - Performance can be improved by the use of parallel processing
- R, and the additional packages available to enhance the use of R, are available free of charge

# R Resources

# R Online Resources

- A list of R packages is contained here:

[https://cran.r-project.org/web/packages/available\\_packages\\_by\\_date.html](https://cran.r-project.org/web/packages/available_packages_by_date.html)

- By clicking on a particular package, you'll be taken to a page with more details, as well as a link to download the documentation
- Typing `help(topic)` in R pulls up a brief help file with syntax and examples, but the online manuals contain more detail

# R Online Resources

- UCLA Institute for Digital Research and Education
  - List of topics and R resources (getting started, data examples, etc.) can be found here:  
<http://www.ats.ucla.edu/stat/r/>
- RStudio Cheatsheets
  - <https://www.rstudio.com/resources/cheatsheets/>

# Other R Resources

1. Kleiber, C., & Zeileis, A. (2008). *Applied econometrics with R*. Springer Science & Business Media.
  - Great reference for the applied researcher wanting to use R for econometric analysis. Includes R basics, linear regression model, panel data models, binary outcomes, etc.
2. Jones, A. M., Rice, N., d'Uva, T. B., & Balia, S. (2013). *Applied health economics*. Routledge.
  - Excellent reference for applied health economics. Examples are all performed using STATA, but foreign package should help here.
3. CRAN Task View: Econometrics
  - A listing of the statistical models used in econometrics, as well as the R package(s) needed to perform them. Available at: <https://cran.r-project.org/view=Econometrics>

Thanks for Listening  
Good luck with R!