

# How I Learned to Stop Worrying and Love the R Console

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# Outline

- 1 Introduction
  - Familiar Examples
- 2 R Console
- 3 Importing Data
- 4 Packages
- 5 Sample Analysis and Visualizations
  - Descriptive Visualizations
  - Modeling
- 6 Reporting
- 7 Where to Go Next?

# Who am I?

Irfan Kanat, PhD Candidate

R user since 2006

Open Source Evangelist

# Before We Begin

Got R & R Studio Installed?

Get your workshop documents:

<https://github.com/iekanat/rworkshop>

# What is this about?

A brief introduction to R.

- R Console
- Importing Data
- Packages
- Sample analyses
- Basic visualization
- Where to get help?



# What is R?

From R project web site:

R is a language *and* an environment for statistical computing and graphics.

- Language
- Environment
- Statistics and Visualization



# What is R?

All this means R is very flexible, which played a huge role in its success.

My take: Low cost, high quality, open source solution for your analysis needs.

# When to Use R?

R is very strong for your classical machine learning and statistical analysis. Thousands of packages address almost all analysis needs. It is a logical first step to start analysis.



# When to Use R?

R is very strong for your classical machine learning and statistical analysis. Thousands of packages address almost all analysis needs. It is a logical first step to start analysis.

Yet it's core design is starting to show its age. There are certain down sides to traditional R:

- Everything is stored in memory<sup>1</sup>
- R is single core<sup>1</sup>

---

<sup>1</sup>Except when it is not. There are packages to overcome these issues.

# Best Part of R

**Packages** CRAN houses over 7K packages. Providing functionality way beyond what is available in commercial packages.

**Community** Millions of users mean, all your questions are either already answered or will be in hours.

**Performance** While memory and core restrictions are real, for the cost of a single user license of a commercial package, you can buy better hardware to run R. Furthermore, with the packages providing multicore and flatfile functionality, R performance is on par or better than commercial packages

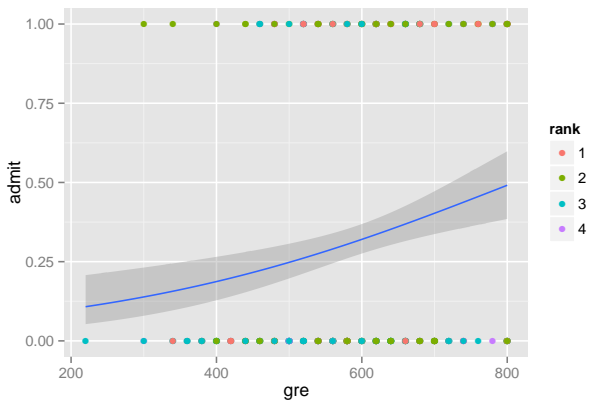
# Logistic Regression

```
# Fit the model
logit_0 <- glm(admit ~ ., admitData, family = "binomial")
# Display fitted model
summary(logit_0)
```

```
##
## Call:
## glm(formula = admit ~ ., family = "binomial", data = admitData)
##
## Deviance Residuals:
##      Min       1Q   Median       3Q      Max
## -1.6268  -0.8662  -0.6388   1.1490   2.0790
##
## Coefficients:
##              Estimate Std. Error z value Pr(>|z|)
## (Intercept) -3.989979   1.139951  -3.500  0.000465 ***
## gre          0.002264   0.001094   2.070  0.038465 *
## gpa          0.804038   0.331819   2.423  0.015388 *
## rank2       -0.675443   0.316490  -2.134  0.032829 *
## rank3       -1.340204   0.345306  -3.881  0.000104 ***
## rank4       -1.551464   0.417832  -3.713  0.000205 ***
## ---
```

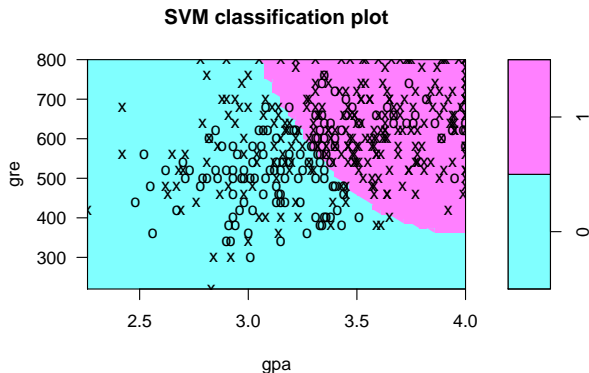
# Logistic Regression

```
ggplot(admitData, aes(x = gre, y = admit)) + geom_point(aes(colour = rank)) +  
  stat_smooth(method = "glm", family = "binomial", se = T)
```

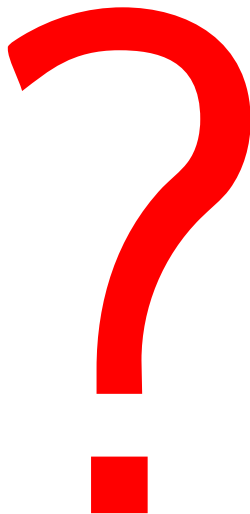


# Support Vector Machine

```
# Fit the model  
svm_0 <- svm(admit ~ ., data = admitData, type = "C-classification")  
# Plot the results  
plot(svm_0, admitData, gre ~ gpa) # Let us plot the results
```



# Questions



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# Command Driven Interface

Command line may be intimidating

Power over Convenience

Consider the number of

- Functions
- Parameters
- Data sources
- Variables
- Replications



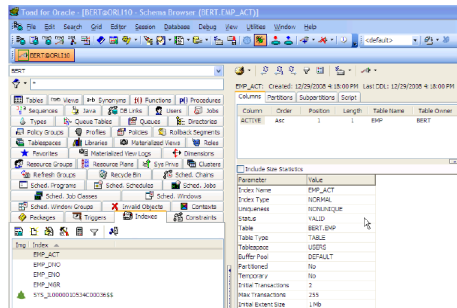
# Command Driven Interface

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# R Studio

The screenshot displays the R Studio interface with four main panels:

- Script Editor (Top Left):** Contains an R script with comments and code. A red box highlights the first 40 lines, and a large red letter 'A' is overlaid on the text. The code includes a comment about saving the project, a function definition for 'get\_wet', and a series of console output lines (39-44).
- Environment (Top Right):** Shows the current environment with variables: 'admtData' (400 obs. of 4 variables), 'csv' (24 obs. of 3 variables), 'd' (10 obs. of 3 variables), 'xLsx' (29 obs. of 3 variables), 'a' (5), 'b' (num [1:30]), 'c' (num [1:30]), 'country' (Large SpatialPolygonsDataFrame (244 elements, 1.5 Mb)), 'fac' (chr [1:10]), 'logit\_0' (List of 30), 'svm\_0' (List of 30), and 'hello' (function(x)). A large blue letter 'C' is overlaid on the 'xLsx' variable.
- Console (Bottom Left):** Shows the output of the script, including the dispersion parameter for binomial family, null deviance, residual deviance, AIC, and the number of Fisher Scoring iterations. A large green letter 'B' is overlaid on the text.
- Plots (Bottom Right):** Displays an 'SVM classification plot' showing 'gpa' on the x-axis (ranging from 2.5 to 4.0) and 'pre' on the y-axis (ranging from 300 to 800). The plot shows two classes of data points (circles and crosses) separated by a decision boundary. A yellow circle highlights a specific data point. A legend on the right indicates two classes: 0 (cyan) and 1 (magenta).

# New Project

File > New Project

Empty Directory > Empty Project > Directory Name: Workshop

# R as a Calculator I

```
# Arithmetics
```

```
2 + 2
```

```
## [1] 4
```

```
2 * 3
```

```
## [1] 6
```

```
2^3
```

```
## [1] 8
```

```
log(100, 10)
```

```
## [1] 2
```

# R as a Calculator II

```
# Logic
1 == 2

## [1] FALSE

1 != 2

## [1] TRUE

2 < 3

## [1] TRUE
```

# Variables I

```
A <- 2
```

```
A
```

```
## [1] 2
```

```
a # Case sensitive
```

```
## Error in eval(expr, envir, enclos): object 'a' not found
```

```
"A" != "a" # Explanation
```

```
## [1] TRUE
```

```
B <- 7
```

```
A + B
```

```
## [1] 9
```

# Variables II

```
C <- c(1, 3, 7, 9)  # A list can be in a variable
```

```
C
```

```
## [1] 1 3 7 9
```

```
C + A
```

```
## [1] 3 5 9 11
```

```
C * A
```

```
## [1] 2 6 14 18
```

```
C < 5
```

```
## [1] TRUE TRUE FALSE FALSE
```

# Indexes and Data Frames I

```
1:30
```

```
## [1] 1 2 3 4 5 6 7 8 9 10 11 12 13 14 15 16 17 18 19 20 21 22 23  
## [24] 24 25 26 27 28 29 30
```

```
C[3]
```

```
## [1] 7
```

```
C[c(2, 3)]
```

```
## [1] 3 7
```

```
C[1:3]
```

```
## [1] 1 3 7
```



# Indexes and Data Frames II

```
Countries <- data.frame(names=c("US", "TR", "DE"), supply=c(10, 8, 7),  
                        those=c(TRUE, FALSE, FALSE))
```

Countries

```
##   names supply those  
## 1    US     10  TRUE  
## 2    TR      8 FALSE  
## 3    DE      7 FALSE
```

# Indexes and Data Frames III

```
Countries[2, ]
```

```
##      names supply those  
## 2      TR          8 FALSE
```

```
Countries[, 3]
```

```
## [1] TRUE FALSE FALSE
```

```
Countries[2, 3]
```

```
## [1] FALSE
```

```
Countries[1:2, ]
```

```
##      names supply those  
## 1      US      10  TRUE  
## 2      TR       8  FALSE
```

# Indexes and Data Frames IV

```
Countries[, "names"]
```

```
## [1] US TR DE  
## Levels: DE TR US
```

```
Countries$names
```

```
## [1] US TR DE  
## Levels: DE TR US
```

```
Countries$those
```

```
## [1] TRUE FALSE FALSE
```

# Loops in R

## CAUTION!

R is notoriously inefficient with your classic loops

- Structure of the Data Frame
- Memory Management

Try to use an apply function instead.

Vectorize your operations.

# For Loop in R

```
for (i in 1:3) print(i)

## [1] 1
## [1] 2
## [1] 3

# Iterating through a data frame
for (i in 1:nrow(Countries)) {
  print(Countries[i, ])
}

##      names supply those
## 1      US      10  TRUE
##      names supply those
## 2      TR       8 FALSE
##      names supply those
## 3      DE       7 FALSE
```

# Functions I

```
mean(C)  # Takes parameters
```

```
## [1] 5
```

```
mean(C, trim = 0.1, na.rm = T)  # Takes multiple parameters
```

```
## [1] 5
```

```
log(sum(C)/length(C))  # Can be combined
```

```
## [1] 1.609438
```

# Functions II

```
HelloWorld <- function(x, y = 1) {  
  for (i in 1:y) {  
    print(paste("Hello", x))  
  }  
}
```

```
HelloWorld("MSBA")
```

```
## [1] "Hello MSBA"
```

```
HelloWorld("MSBA", 2)
```

```
## [1] "Hello MSBA"
```

```
## [1] "Hello MSBA"
```

# Functions III

```
HelloWorld  # Review the source code
```

```
## function(x, y = 1) {  
##   for (i in 1:y) {  
##     print(paste("Hello", x))  
##   }  
## }
```

```
ls
```

```
## function (name, pos = -1L, envir = as.environment(pos), all.names = FALSE,  
##   pattern, sorted = TRUE)  
## {  
##   if (!missing(name)) {  
##     pos <- tryCatch(name, error = function(e) e)  
##     if (inherits(pos, "error")) {  
##       name <- substitute(name)  
##       if (!is.character(name))  
##         name <- deparse(name)  
##       warning(gettextf("%s converted to character string",  
##         sQuote(name)), domain = NA)  
##       pos <- name
```



# Commonly Used Functions I

```
ls() # Get a list of objects in the workspace

## [1] "A"          "admitData"  "B"          "C"          "Countries"
## [6] "HelloWorld" "i"          "logit_0"    "svm_0"
```

```
ls(pattern = "*_0") # partial match on object search

## [1] "logit_0" "svm_0"
```

```
rm("svm_0") # Remove an object from the workspace

# rm(list=ls(pattern=ls())) # This would remove everything if ran
```

# Commonly Used Functions II

```
mean(A)  # Mean

## [1] 2

sd(admitData[, "gre"])  # Standard Deviation

## [1] 115.5165

AIC(logit_0)

## [1] 470.5175
```

# Commonly Used Functions III

```
str(Countries) # Look at the structure of objects
```

```
## 'data.frame': 3 obs. of 3 variables:
## $ names : Factor w/ 3 levels "DE","TR","US": 3 2 1
## $ supply: num 10 8 7
## $ those : logi TRUE FALSE FALSE
```

```
summary(Countries) # Get summary of data
```

```
## names      supply      those
## DE:1   Min.    : 7.000   Mode :logical
## TR:1   1st Qu.: 7.500   FALSE:2
## US:1   Median : 8.000   TRUE :1
##        Mean    : 8.333   NA's :0
##        3rd Qu.: 9.000
##        Max.    :10.000
```

# Commonly Used Functions IV

```
summary(logit_1) # Get summary of model

##
## Call:
## glm(formula = admit ~ gre, family = "binomial", data = admitData)
##
## Deviance Residuals:
##      Min       1Q   Median       3Q      Max
## -1.1623  -0.9052  -0.7547   1.3486   1.9879
##
## Coefficients:
##              Estimate Std. Error z value Pr(>|z|)
## (Intercept) -2.901344   0.606038  -4.787 1.69e-06 ***
## gre          0.003582   0.000986   3.633 0.00028 ***
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## (Dispersion parameter for binomial family taken to be 1)
##
##      Null deviance: 499.98  on 399  degrees of freedom
## Residual deviance: 486.06  on 398  degrees of freedom
## AIC: 490.06
##
## Number of Fisher Scoring iterations: 4
```

# Commonly Used Functions V

```
cor(admitData[, 1:3]) # Get correlation matrix
```

```
##           admit           gre           gpa
## admit 1.0000000 0.1844343 0.1782123
## gre   0.1844343 1.0000000 0.3842659
## gpa   0.1782123 0.3842659 1.0000000
```

# Questions



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# Importing Data

R allows importing data from a wide variety of sources.

- Comma Separated Values (CSV)
- Databases
- Flat files
- Lesser statistical packages
- and more



# Importing CSV Files

CSV has certain advantages that make it popular.

- Compatibility
- Flexibility
- Simplicity

## Sample

```
"iso2", "Supply", "Those"
"AU", 20, 0
"TR", 80, 1
"US", 100, 0
"GB", 50, 0
"DE", 70, 0
```

We use `read.csv()` or `read.csv2()` commands to import the csv files.

```
saveData <- read.csv("PathToCSV", header = TRUE, sep = ",", quote = "\"", )
```

# Working with Excel Files

Much like CSV, except it lacks the simplicity, flexibility, and compatibility of CSV.

```
# Load the necessary library  
library(xlsx)  
# Read in the data from excel file  
xlsx <- read.xlsx("country.xlsx", sheetIndex = 1)
```

# Working with Databases

No speed advantage.

Data larger than memory.

Working with databases:

- Work in the database.
- Import data from database.



# Working with Databases

```
# Load the necessary library
library(RMySQL)
# Establish connection to the database.
channel <- dbConnect(MySQL(), user = "uname", password = "pwd", host = "127.0.0.1",
  dbname = "exampledata")
# Send query and save results in R workspace
sql <- dbGetQuery(channel, "SELECT * FROM table;")
```

# Lesser Statistical Packages :P

Foreign Package

Newer file formats

- sas7bdat
- readstata13



# Questions



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# Packages: Source of R's Power

Encountered already

Make R extendible

Like libraries

Collection of:

- functions
- documentation
- data files





# Gifts from the Community

Currently over 7000 packages

for

- Statistical Modeling
- Machine Learning
- Data Manipulation
- Visualization
- ...

from

- Economics
- Computer Science
- Statistics
- Medicine
- ...



## Great but Where are My Gifts?

# Comprehensive R Archive Network (CRAN)

A Group of FTP and HTML servers hosting R packages.

R has built in package management facilities.

Most of these can be achieved through the R Studio GUI. (Area D, packages pane)

# Package Management

```
# Installing a package
install.packages("e1071") # Notice the quotes around package name
# Loading package into memory
library(e1071) # Notice the lack of quotes
# Unload package
detach("package:e1071", unload = TRUE) # Notice the package: prepended

# Get the list of packages loaded
(.packages())
# Get list of all installed packages (output omitted)
.packages(all.available = T)
```

# How to Find Packages

If you want to search a certain word in installed packages' documentation, you can always use `??` or `help.search()`

```
??mixed  
help.search("mixed model")
```

Internet searches are a bit problematic as R can be a bit ambiguous until Google learns you are interested in the statistical computing environment.

Comprehensive R Archive Network (CRAN)

R Forge

R site search also available with command `RSiteSearch()`

R seek

# Commonly Used Packages: Data Manipulation

**data.tables** Replaces traditional `data.frame`.

- Faster access/write
- Improved selection
- Improved subsetting
- Improved aggregation

Not a drop-in replacement as it breaks compatibility in some cases.

## **ddplyr**

Additional functionality for:

- selection
- filtering
- aggregation

Provides efficient back-end data structures to speed things up.

Works with databases as well.

## Commonly Used Packages: Statistics

Multiple Regression: Stats package, `lm()` (loaded by default)

Generalized Linear Models: Stats package, `glm()`

Traditional Econometric Models: `plm` package

Mixed Modeling: `nlme` and `lme4` packages

# Commonly Used Packages: Machine Learning

Most probably all you need is caret package.

Caret package is a wrapper for a host of classification and regression model training functions. It eases visualizations, data manipulation, and analytics among others. **It currently supports over 150 types of models.**

If you insist on using individual packages:

Classifiers: class package

Support Vector Machines: kernlab, e1071 packages

Clustering: Base package (kmeans(), hclust()), mclust package

Neural Networks: neuralnet package.

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# Motor Trends Dataset

We will use 1974 Motor Trend dataset. It has 32 observations and 11 variables.

- mpg: Miles per gallon
- cyl: Number of cylinders
- disp: Displacement
- hp: Horse Power
- drat: Rear axle ratio
- wt: Weight
- qsec: quarter mile time
- vs: V - S
- am: 0 automatic, 1 manual
- gear: Gears
- carb: Number of carburetors

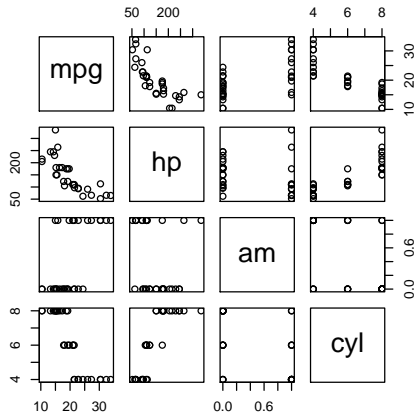
# Motor Trends Dataset I

```
summary(mtcars)
```

```
##           mpg           cyl           disp           hp
##  Min.    :10.40   Min.    :4.000   Min.    : 71.1   Min.    : 52.0
##  1st Qu.:15.43   1st Qu.:4.000   1st Qu.:120.8   1st Qu.: 96.5
##  Median :19.20   Median :6.000   Median :196.3   Median :123.0
##  Mean   :20.09   Mean   :6.188   Mean   :230.7   Mean   :146.7
##  3rd Qu.:22.80   3rd Qu.:8.000   3rd Qu.:326.0   3rd Qu.:180.0
##  Max.   :33.90   Max.   :8.000   Max.   :472.0   Max.   :335.0
##
##      drat           wt           qsec           vs
##  Min.    :2.760   Min.    :1.513   Min.    :14.50   Min.    :0.0000
##  1st Qu.:3.080   1st Qu.:2.581   1st Qu.:16.89   1st Qu.:0.0000
##  Median :3.695   Median :3.325   Median :17.71   Median :0.0000
##  Mean   :3.597   Mean   :3.217   Mean   :17.85   Mean   :0.4375
##  3rd Qu.:3.920   3rd Qu.:3.610   3rd Qu.:18.90   3rd Qu.:1.0000
##  Max.   :4.930   Max.   :5.424   Max.   :22.90   Max.   :1.0000
##
##      am           gear           carb
##  Min.    :0.0000   Min.    :3.000   Min.    :1.000
##  1st Qu.:0.0000   1st Qu.:3.000   1st Qu.:2.000
##  Median :0.0000   Median :4.000   Median :2.000
##  Mean   :0.4062   Mean   :3.688   Mean   :2.812
##  3rd Qu.:1.0000   3rd Qu.:4.000   3rd Qu.:4.000
##  Max.   :1.0000   Max.   :5.000   Max.   :8.000
```

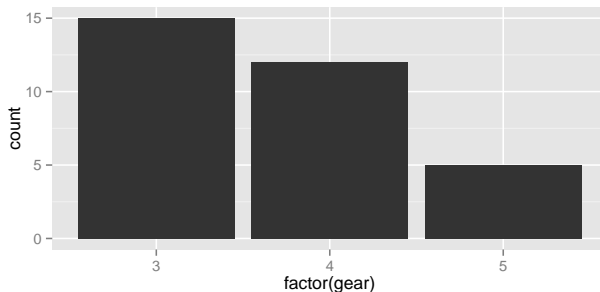
# Motor Trends Dataset II

```
pairs(mtcars[, c("mpg", "hp", "am", "cyl")]) # Visualize Correlations
```



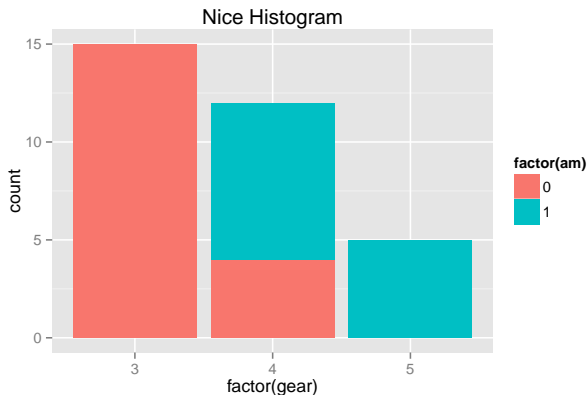
# Histograms I

```
data(mtcars) # Load the Dataset
library(ggplot2) # Load the ggplot package
# Nr of cars by number of gears
qplot(factor(gear), data = mtcars, geom = "bar")
```



# Histograms II

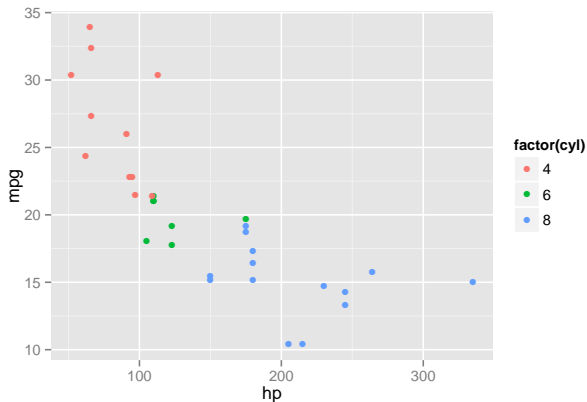
```
# If we are interested in a third categorical variable vs:  
qplot(factor(gear), data=mtcars, geom="bar", fill=factor(am)) +  
ggtitle('Nice Histogram') # This is how you add a title
```



# Scatter Plots I

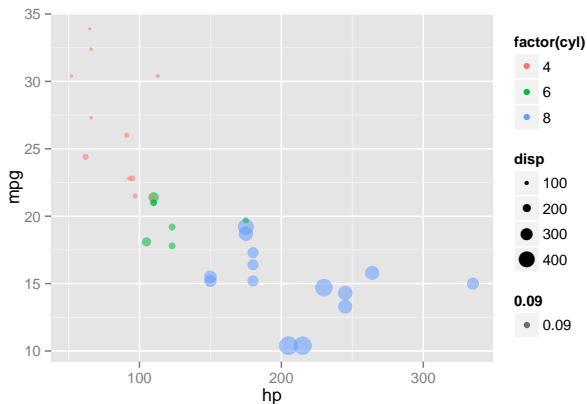
```
# Two continuous variables
```

```
qplot(hp, mpg, data = mtcars, color = factor(cyl))
```



# Scatter Plots II

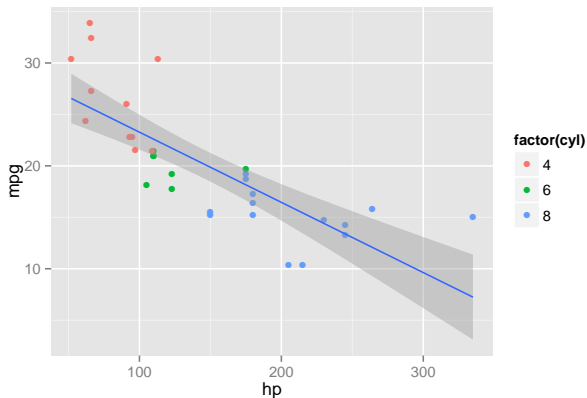
```
# Add two more variables represented by color and size of points  
qplot(hp, mpg, data = mtcars, color = factor(cyl), size = disp, alpha = 0.09)
```





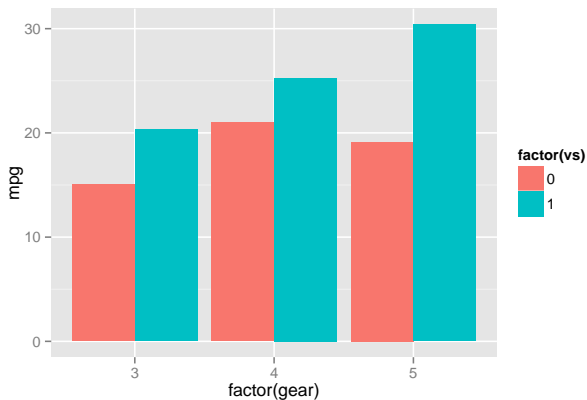
# Scatter Plots III

```
ggplot(mtcars, aes(x = hp, y = mpg)) + geom_point(aes(color = factor(cyl))) +  
  # Add a regression line  
  geom_smooth(method = lm)
```



# Bar Charts I

```
ggplot(mtcars, aes(x = factor(gear), y = mpg, fill = factor(vs)), color = factor(vs))  
  stat_summary(fun.y = mean, position = position_dodge(), geom = "bar")
```



# Multiple Regression

We will keep using motor trends data set.

Pay close attention to how we specify the model.

## Formula

$$Y = \beta_0 + \beta_1 x_1 + \beta_2 x_2 + \epsilon$$

## R Model

$$Y \sim x_1 + x_2$$

This basic structure will remain constant across many R packages.

# Regression I

```
# Let us estimate gas milage
reg_0 <- lm(mpg ~ hp + cyl + am, mtcars)
summary(reg_0)

##
## Call:
## lm(formula = mpg ~ hp + cyl + am, data = mtcars)
##
## Residuals:
##      Min       1Q   Median       3Q      Max
## -4.864 -1.811 -0.158  1.492  6.013
##
## Coefficients:
##              Estimate Std. Error t value Pr(>|t|)
## (Intercept)  30.88834    2.78422   11.094 9.27e-12 ***
## hp          -0.03688    0.01452   -2.540  0.01693 *
## cyl         -1.12721    0.63417   -1.777  0.08636 .
## am           3.90428    1.29659    3.011  0.00546 **
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Residual standard error: 2.807 on 28 degrees of freedom
## Multiple R-squared:  0.8041, Adjusted R-squared:  0.7831
## F-statistic: 38.32 on 3 and 28 DF,  p-value: 4.791e-10
```

# Regression II

```
# Access Fitted Values View first 3 predictions
```

```
reg_0$fitted.values[1:3]
```

```
##      Mazda RX4 Mazda RX4 Wag      Datsun 710  
##      23.97302      23.97302      26.85433
```

```
# Bonus: Are the residuals normally distributed
```

```
shapiro.test(reg_0$residuals)
```

```
##
```

```
## Shapiro-Wilk normality test
```

```
##
```

```
## data:  reg_0$residuals
```

```
## W = 0.98366, p-value = 0.8961
```

# Regression III

```
# PREDICTING NEW DATA BASED ON MODEL
newCar <- mtcars[3, ] # 3rd observation is Datsun 710
newCar$am <- 0 # What if it was automatic?
reg_0$fitted.values[3] # Previous estimate

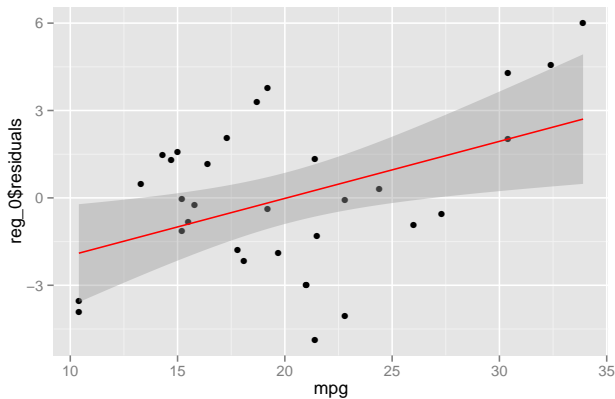
## Datsun 710
## 26.85433

predict(reg_0, newdata = newCar) # Datsun with automatic transmission

## Datsun 710
## 22.95005

## Plot the residuals against observation
qplot(data=mtcars, x = mpg, y = reg_0$residuals) + #
  stat_smooth(method = "lm", col = "red")
```

# Regression IV



# Regression V

```
## COMPARE MODELS
reg_1 <- lm(mpg ~ hp + cyl + am + wt, mtcars) # add weight
anova(reg_0, reg_1)

## Analysis of Variance Table
##
## Model 1: mpg ~ hp + cyl + am
## Model 2: mpg ~ hp + cyl + am + wt
##   Res.Df    RSS Df Sum of Sq    F    Pr(>F)
## 1       28 220.55
## 2       27 170.00   1    50.555 8.0295 0.008603 **
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

# AIC of the model
AIC(reg_0)

## [1] 162.5849

AIC(reg_1)

## [1] 156.2536
```



# Logistic Regression

Dependent variable will be type (binary).

It is basically a regression with a binomial link function.

## Formula

$$\log \left( \frac{p}{1-p} \right) = \beta_0 + \beta_1 x_1 + \epsilon$$

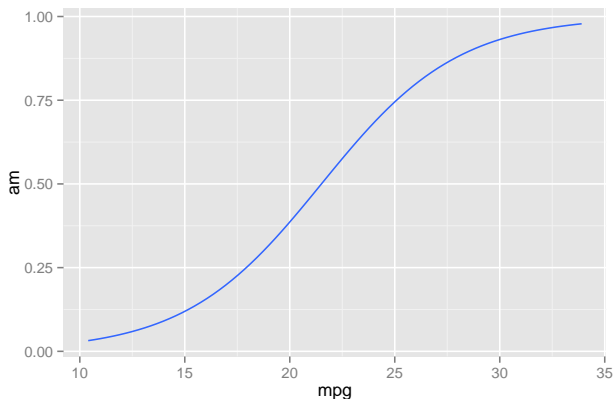
# Logit I

```
logit_2 <- glm(am ~ mpg + drat + cyl, data = mtcars, family = "binomial")
summary(logit_2)
```

```
##
## Call:
## glm(formula = am ~ mpg + drat + cyl, family = "binomial", data = mtcars)
##
## Deviance Residuals:
##      Min       1Q   Median       3Q      Max
## -1.58367  -0.31020  -0.03757   0.17972   1.75395
##
## Coefficients:
##              Estimate Std. Error z value Pr(>|z|)
## (Intercept)  -49.4548    24.1280  -2.050   0.0404 *
## mpg           0.6378     0.4266   1.495   0.1349
## drat          7.2595     3.2702   2.220   0.0264 *
## cyl           1.6115     1.0801   1.492   0.1357
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## (Dispersion parameter for binomial family taken to be 1)
##
##      Null deviance: 43.23  on 31  degrees of freedom
## Residual deviance: 17.03  on 28  degrees of freedom
## AIC: 25.03
##
## Number of Fisher Scoring iterations: 7
```

# Logit II

```
# VISUALIZE mpg - Transmission RELATION  
ggplot(mtcars, aes(x = mpg, y = am)) +  
  stat_smooth(method="glm", family="binomial", se=FALSE) #
```



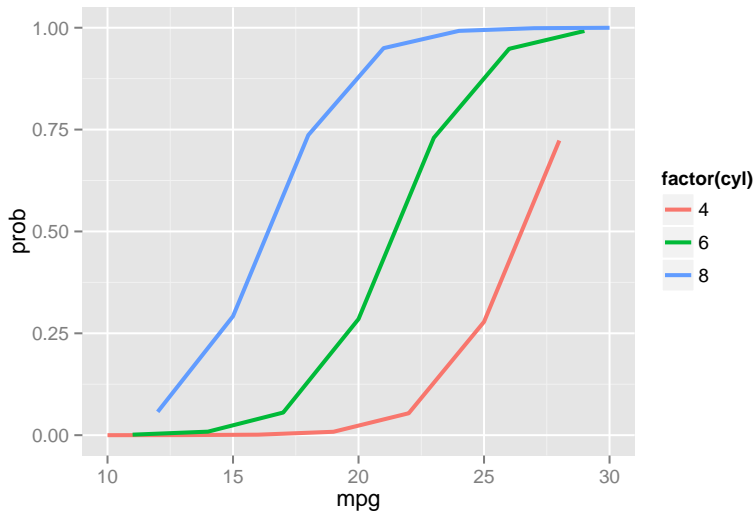
# Logit III

```
## VISUALIZE Mpg - Transmission FOR DIFFERENT NUMBERS OF CYLINDERS

# Create a new dataset with varying number of cylinders and other variables
# fixed at mean levels.
mtcars2 <- data.frame(mpg = rep(10:30, 3), drat = mean(mtcars$drat), disp = mean(mtcars$disp),
  cyl = rep(c(4, 6, 8), 21))
# Predict probability of new data
mtcars2$prob <- predict(logit_2, newdata = mtcars2, type = "response")

# Plot the results
ggplot(mtcars2, aes(x = mpg, y = prob)) + geom_line(aes(colour = factor(cyl)),
  size = 1) # a different color for each category$
```

# Logit IV



# Logit V

```
## DIAGNOSTICS

# Let us compare predicted values to real values
mtcars$prob <- predict(logit_2, type = "response")
# Prevalence of Manual Transmission
mean(mtcars$am)

## [1] 0.40625

# Create predict variable
mtcars$pred <- 0
# If probability is greater than .6 (1-prevalence), set prediction to 1
mtcars[mtcars$prob > 0.6, "pred"] <- 1
```

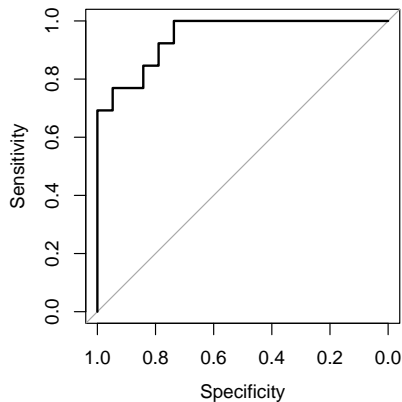
# Logit VI

```
## ROC CURVE

# Load the necessary library
library(pROC)
# Calculate the ROC curve using the predicted probability vs actual values
logit_2_roc <- roc(am ~ prob, mtcars)
# Plot ROC curve
plot(logit_2_roc)

##
## Call:
## roc.formula(formula = am ~ prob, data = mtcars)
##
## Data: prob in 19 controls (am 0) < 13 cases (am 1).
## Area under the curve: 0.9474
```

# Logit VII





# Logit VIII

```
library(caret) # Needed for Confusion Matrix
confusionMatrix(table(mtcars[, c("am", "pred")]))
```

```
## Confusion Matrix and Statistics
##
##      pred
## am  0  1
##    0 18  1
##    1  3 10
##
##               Accuracy : 0.875
##               95% CI : (0.7101, 0.9649)
##      No Information Rate : 0.6562
##      P-Value [Acc > NIR] : 0.005004
##
##               Kappa : 0.7344
##  Mcnemar's Test P-Value : 0.617075
##
##      Sensitivity : 0.8571
##      Specificity : 0.9091
##      Pos Pred Value : 0.9474
##      Neg Pred Value : 0.7692
##      Prevalence : 0.6562
##      Detection Rate : 0.5625
##      Detection Prevalence : 0.5938
##      Balanced Accuracy : 0.8831
##
##      'Positive' Class : 0
##
```

# Caret Package

The Caret package is a wrapper that combines functionality from 27 R packages.

Functions Provided:

- Visualization
- Data Manipulation
- Model Training & Selection
- Parallel Processing

Since so many packages involved, the installation takes a while.

```
install.packages("caret", dependencies = c("Depends", "Suggests"))
```

For this part of the exercise I will focus on Caret Package, following its [vignette](#).

# Data Splitting I

```
## OBTAIN DATASET Dataset comes with mlbench package
library(mlbench)
# Load dataset into the current workspace
data(Sonar)
# 208 observations and 61 variables
```

```
## SPLIT THE DATA

# caret provides functionality
library(caret)
# Set random number seed for reproducibility
set.seed(107)
# Create an index of observations to be included in Training
indexTrain <- createDataPartition(y = Sonar$Class, p = 0.75, list = FALSE)

# $plit the data using the index
Train <- Sonar[indexTrain, ]
Test <- Sonar[-indexTrain, ]
```

# Train a PLS Discriminant Model I

```
## Declare Tuning Control Parameters
ctrl <- trainControl(method = "repeatedcv", # K-fold cross validation
                     repeats = 3, # Repeat resampling 3 times
                     classProbs = TRUE, # Calculate predicted prob for ROC
                     summaryFunction = twoClassSummary) # Set performance metrics for

## Train the Classifier
plsFit <- train(Class ~ ., data = Train, method = "pls",
               tuneLength = 10, # Number of component sets to be evaluated (more
               trControl = ctrl, # Use control parameters from above
               metric = "ROC", # Criteria ROC
               preProc = c("center", "scale")) # Center and scale the predictors
```

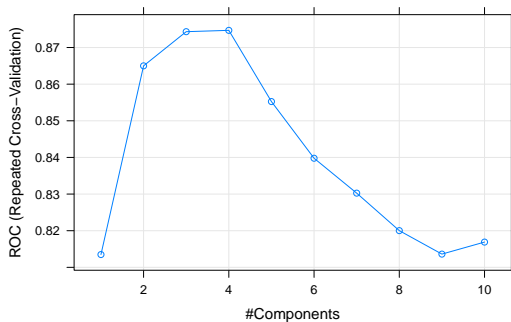
# Train a PLS Discriminant Model II

```
plsFit
```

```
## Partial Least Squares
##
## 157 samples
## 60 predictors
## 2 classes: 'M', 'R'
##
## Pre-processing: centered (60), scaled (60)
## Resampling: Cross-Validated (10 fold, repeated 3 times)
## Summary of sample sizes: 142, 141, 142, 142, 142, 142, ...
## Resampling results across tuning parameters:
##
##      ncomp  ROC      Sens      Spec      ROC SD      Sens SD      Spec SD
##      1      0.8134921 0.7291667 0.7291667 0.11844879 0.1387811 0.1935289
##      2      0.8649967 0.7694444 0.8041667 0.08381907 0.1416676 0.1521373
##      3      0.8743221 0.7476852 0.8363095 0.08548836 0.1726683 0.1375303
##      4      0.8746858 0.7578704 0.7642857 0.08443793 0.1512983 0.1539276
##      5      0.8552497 0.7152778 0.7767857 0.09587112 0.1771056 0.1584577
##      6      0.8397817 0.7337963 0.7732143 0.09814150 0.1726297 0.1749730
##      7      0.8302579 0.7101852 0.7916667 0.10762747 0.1992674 0.1655561
##      8      0.8200231 0.7157407 0.7607143 0.12724476 0.1805642 0.1621515
##      9      0.8136161 0.7245370 0.7696429 0.12961432 0.1847286 0.1593987
##     10      0.8168981 0.7203704 0.7559524 0.11678705 0.1912620 0.1436503
##
## ROC was used to select the optimal model using the largest value.
## The final value used for the model was ncomp = 4.
```

# Train a PLS Discriminant Model III

```
# evaluate the performance of different number of components extracted  
plot(plsFit)
```



# Validate with Test Data I

```
plsPredict <- predict(plsFit, newdata = Test) # Predict results  
head(plsPredict) # View predictions
```

```
## [1] R M M R M R  
## Levels: M R
```

```
head(Test$Class) # View actual$
```

```
## [1] R R R R R R  
## Levels: M R
```

# Validate with Test Data II

```
# Get confusion matrix (predicted vs actual)
confusionMatrix(data = plsPredict, Test$Class) ##

## Confusion Matrix and Statistics
##
##           Reference
## Prediction  M  R
##           M 17  6
##           R 10 18
##
##           Accuracy : 0.6863
##           95% CI : (0.5411, 0.8089)
##           No Information Rate : 0.5294
##           P-Value [Acc > NIR] : 0.01674
##
##           Kappa : 0.3761
##           Mcnemar's Test P-Value : 0.45325
##
##           Sensitivity : 0.6296
##           Specificity : 0.7500
##           Pos Pred Value : 0.7391
##           Neg Pred Value : 0.6429
##           Prevalence : 0.5294
##           Detection Rate : 0.3333
##           Detection Prevalence : 0.4510
##           Balanced Accuracy : 0.6898
##
##           'Positive' Class : M
##
```



# Questions



# Outline

- 1 Introduction
  - Familiar Examples
- 2 R Console
- 3 Importing Data
- 4 Packages
- 5 Sample Analysis and Visualizations
  - Descriptive Visualizations
  - Modeling
- 6 Reporting**
- 7 Where to Go Next?

# Professional Looking Documents

All the documentation of this workshop has been prepared in R Studio.

Look into the Rmd documents to see how this was done.

`summary()` function does not provide the best looking model summaries.  
Try the `texreg` package (do as I say, not as I do).

# Questions



# Outline

- 1 Introduction
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# Quo Vadis?

8 hours is not enough!

RTFM<sup>2</sup>: Documentation is your friend, read the vignettes and manuals.

For tutorials: [Institute For Digital Research and Education, UCLA](#)

For questions: [StackExchange](#)<sup>3</sup>

- [Cross Validated](#) specializes on Statistics
- [Stack Overflow](#) for R programming

R Programming is quirky, learn about efficient R programming.

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<sup>2</sup>Read the Friendly Manual

<sup>3</sup>Word of caution, [pay attention to how you ask your questions](#). It is more important than you realize.