Data Exploration, Visualization, and Feature Engineering

Data Science Dojo

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Agenda

- Why data exploration and visualization
- Exploration and visualization using R
  - Core R functionality – iris dataset
  - lattice package – mtcars dataset
  - ggplot2 package – diamonds and G20 datasets
- Story-telling with data
  - Titanic data set

WHY DATA EXPLORATION AND VISUALIZATION

Data beats algorithm but…

- More data usually yields good generalization performance, even with a simple algorithm
- But there are caveats
  - Amount of data may have diminishing returns
  - Data quality and variety matters
  - A decent performing learning algorithm is still needed
  - Most importantly, extracting useful features out of data is important

Why feature engineering matters

- 23:05:33 –5 UTC, April 3, 2014

Hour of date	Day of week	AM/PM

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Dispelling common myths

- There is *NO* single ML algorithm that will take raw data and give you the best model

- You do *NOT* need to know a lot of machine learning algorithms to build robust predictive models

Janitorial work is important

- Not spending time on understanding your data is a source of many problems!
- Remember the 80/20 rule
  - 80% : Data cleaning, data exploration, feature engineering, pre-processing etc…
  - 20% : Model building

EXPLORATION AND VISUALIZATION USING R

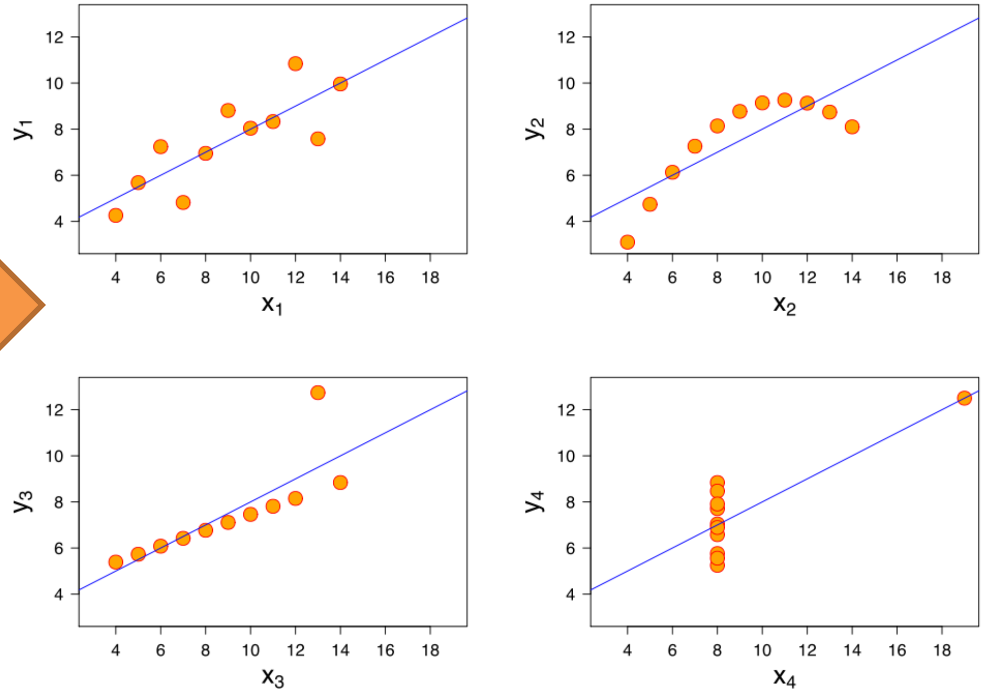
Objectives

- Develop an understanding of the high-level thinking process of data exploration
- Make sense of data using visualization techniques
- Learn to perform feature engineering
- Become a good storyteller

Anscombe's quartet

I		II		III		IV	
x	y	x	y	x	y	x	y
10.0	8.04	10.0	9.14	10.0	7.46	8.0	6.58
8.0	6.95	8.0	8.14	8.0	6.77	8.0	5.76
13.0	7.58	13.0	8.74	13.0	12.74	8.0	7.71
9.0	8.81	9.0	8.77	9.0	7.11	8.0	8.84
11.0	8.33	11.0	9.26	11.0	7.81	8.0	8.47
14.0	9.96	14.0	8.10	14.0	8.84	8.0	7.04
6.0	7.24	6.0	6.13	6.0	6.08	8.0	5.25
4.0	4.26	4.0	3.10	4.0	5.39	19.0	12.50
12.0	10.84	12.0	9.13	12.0	8.15	8.0	5.56
7.0	4.82	7.0	7.26	7.0	6.42	8.0	7.91
5.0	5.68	5.0	4.74	5.0	5.73	8.0	6.89

Plot



Anscombe's quartet

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10.0	8.04	10.0	9.14	10.0	7.46	8.0	6.58
8.0	6.95	8.0	8.14	8.0	6.77	8.0	5.76
13.0	7.58	13.0	8.74	13.0	12.74	8.0	7.71
9.0	8.81	9.0	8.77	9.0	7.11	8.0	8.84
11.0	8.33	11.0	9.26	11.0	7.81	8.0	8.47
14.0	9.96	14.0	8.10	14.0	8.84	8.0	7.04
6.0	7.24	6.0	6.13	6.0	6.08	8.0	5.25
4.0	4.26	4.0	3.10	4.0	5.39	19.0	12.50
12.0	10.84	12.0	9.13	12.0	8.15	8.0	5.56
7.0	4.82	7.0	7.26	7.0	6.42	8.0	7.91
5.0	5.68	5.0	4.74	5.0	5.73	8.0	6.89

Consider the 4 following different datasets

Mean of X	9
Variance of X	11
Mean of Y	7.5
Variance of Y	4.125
Correlation between X & Y	0.816

Awareness

New to R?

- Focus on ideas/concepts rather than exact syntax. R help is your friend. ☺

  ?mean, ?sd
  ??melt *(use two question marks for packages not loaded)*
  help( )
  example( )

- All slides have code samples
- Sample code + slides: 'Data Exploration and Visualization' folder

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Common graphical parameters

- Title of graph using the **main** function, main = “title”
- Label x- axis by using the **xlab** function, xlab = “label x axis”
- Label x- axis by using the **ylab** function, ylab = “label y axis”
- Colors controlled by **col**
- Get legends of layered plots with **auto.key**=TRUE

Exploring data commands

Commands	Description
read.csv() , read.table()	Load data/file into a dataframe
data()	Loads or resets a dataset
names()	List names of variables in a dataframe
head()	First 6 rows of data
tail()	Last 6 rows of data
str()	Display internal structure if R object
View()	View dataset in spreadsheet format in RStudio
dim()	Dimensions(rows and columns) of dataframe
summary()	Display 5-number summary and mean
colnames()	Provide column names

CORE R GRAPHICS

The iris dataset

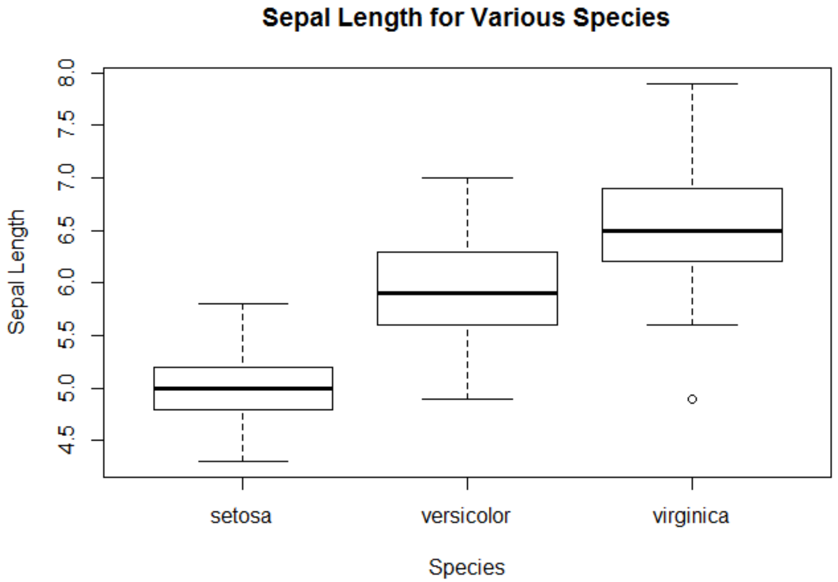
```
data(iris)
head(iris)
```

```
> head(iris)
  Sepal.Length Sepal.Width Petal.Length Petal.Width Species
1          5.1         3.5          1.4         0.2  setosa
2          4.9         3.0          1.4         0.2  setosa
3          4.7         3.2          1.3         0.2  setosa
4          4.6         3.1          1.5         0.2  setosa
5          5.0         3.6          1.4         0.2  setosa
6          5.4         3.9          1.7         0.4  setosa
```

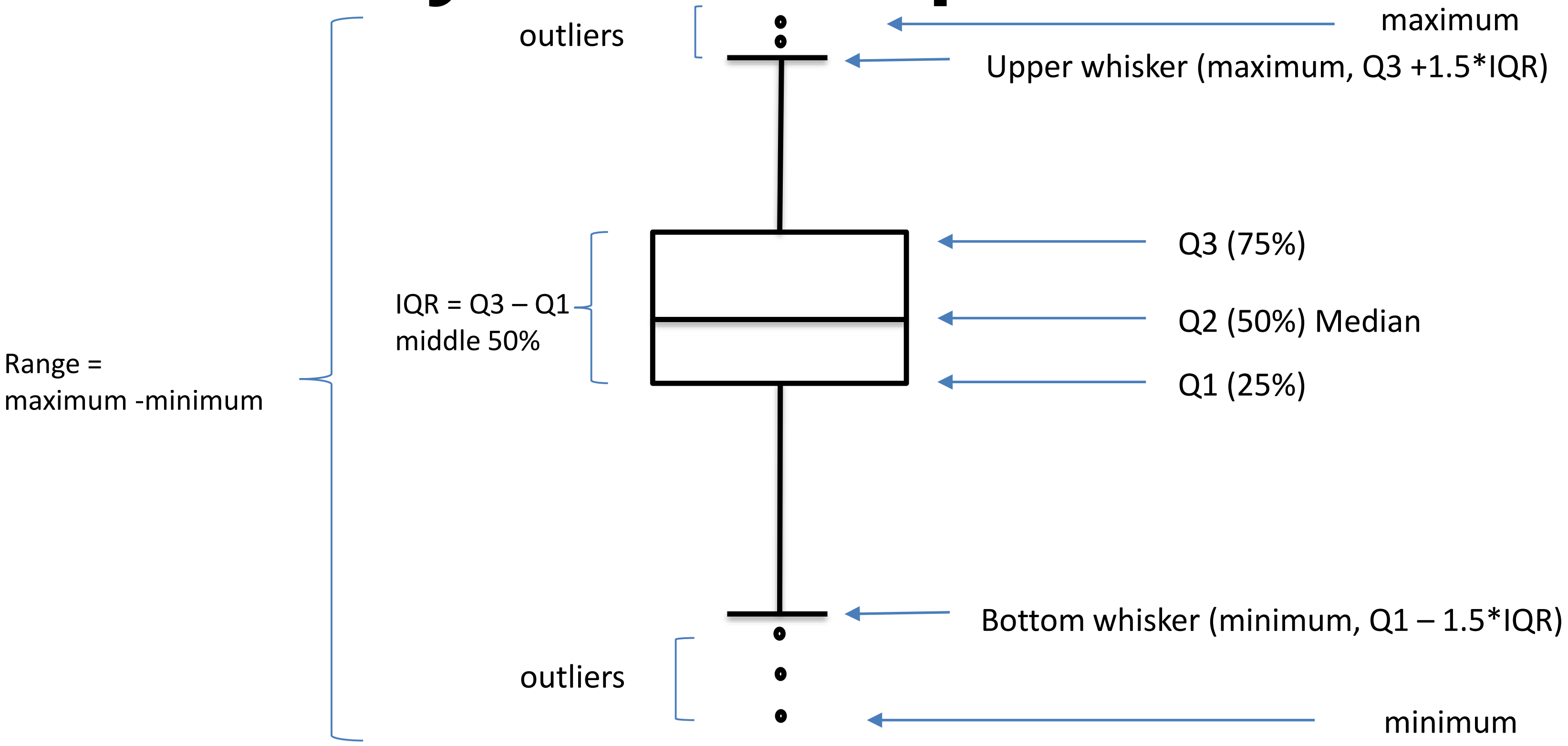
Boxplots

- Summarizes *quantitative/numeric* data

```
# Core Graphics
boxplot(
Sepal.Length ~ Species,
data=iris,
main="Sepal Length for
Various Species",
xlab="Species",
ylab="Sepal Length"
)
```



Anatomy of a boxplot



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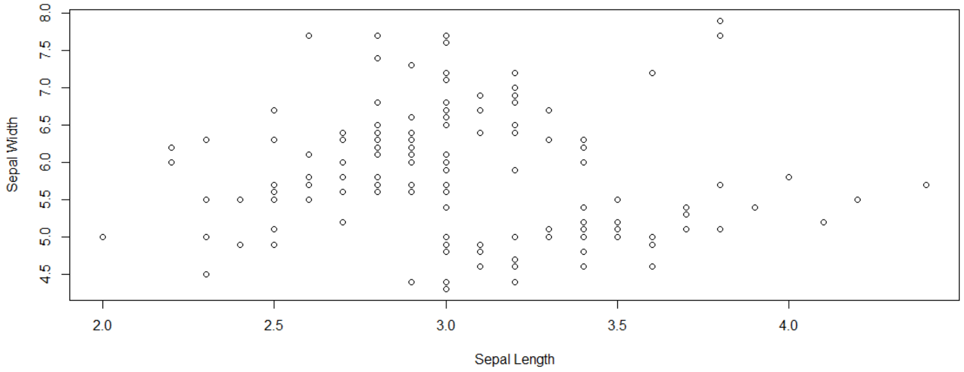
Plot

- Visual depiction of correlation between numeric variables

```
# Core Graphics
plot(Sepal.Length ~ Sepal.Width,
data=iris, xlab= "Sepal Length",
ylab= "Sepal Width")
```

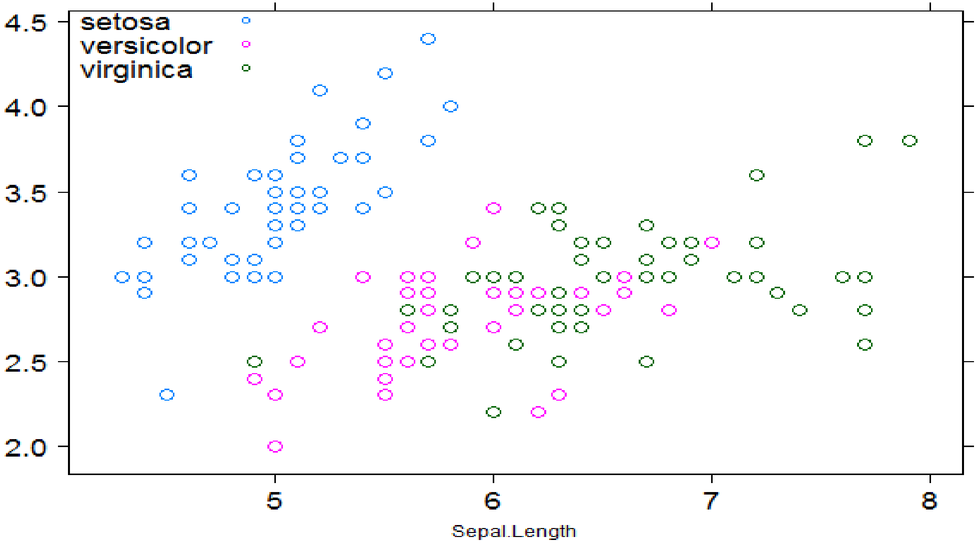


LATTICE GRAPHICS

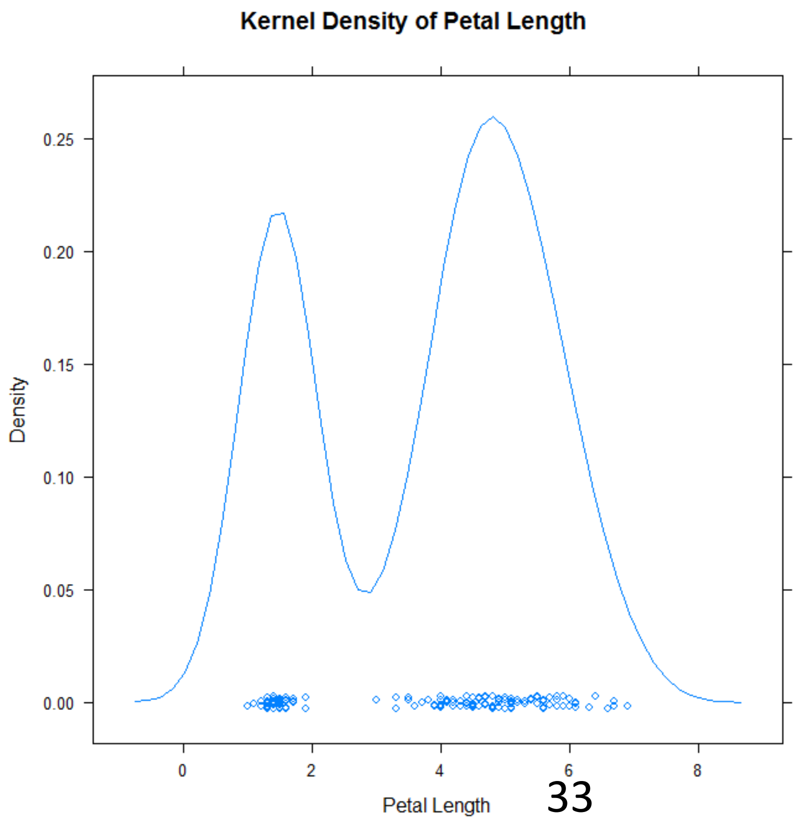
xyplot

- Plot counterpart in lattice package.
- Similar output as core graphics, but easier to color and segment points

```
# Lattice Graphics
library(lattice)
xyplot(Sepal.Width ~
Sepal.Length, data=iris,
groups=Species,
auto.key=TRUE
)
```



Density plots



- Estimates density function from counts
- Area under the curve is always one
- Does not work with missing values

```
densityplot(iris$Petal.Length, main="Kernel Density of Petal Length", xlab="Petal Length")
```

Try adding plot.points=F

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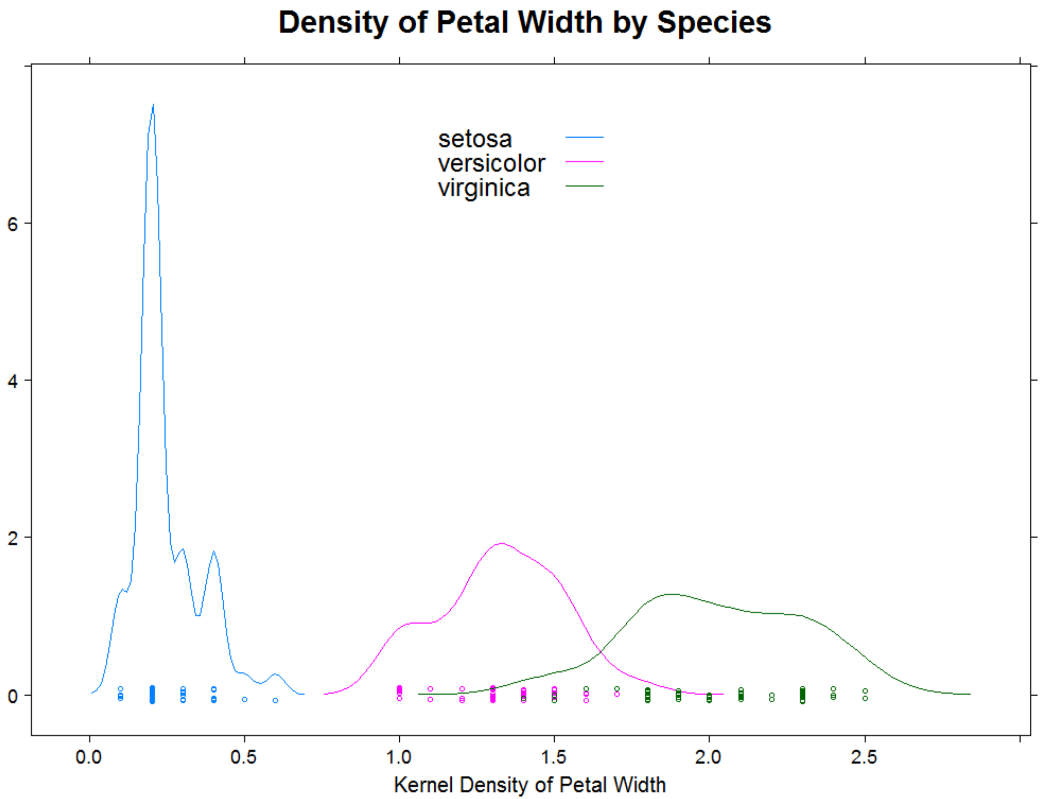
Multiple density plots



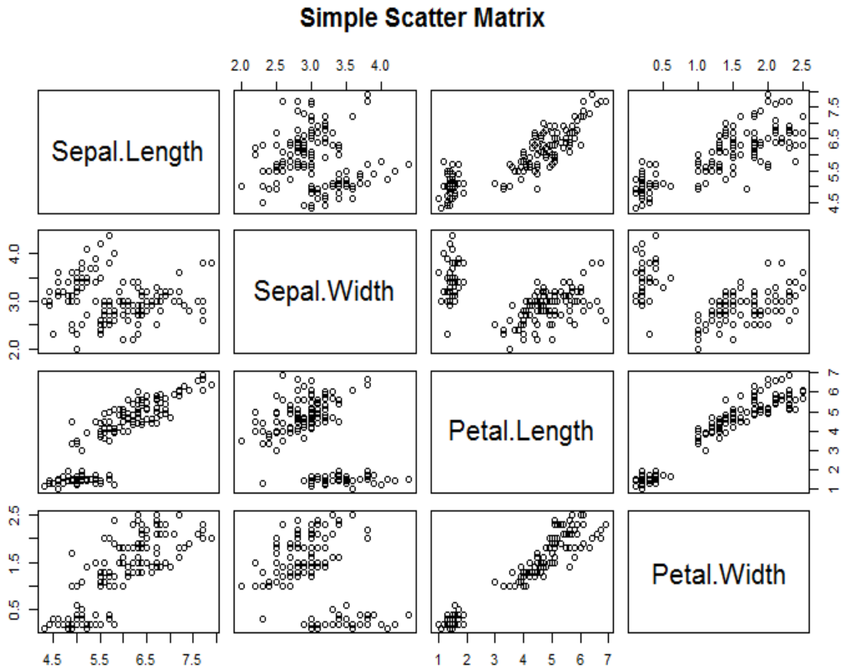

```
densityplot(~Petal.Width,
data=iris,
groups=Species,
auto.key=TRUE,
xlab="Kernel Density of
Petal Width",
ylab="Frequency",
main=list(label="Density
of Petal Width by
Species"))
```

Scatterplot matrix

- Multiple relationships in one graph
- Good for initial explorations

```
# Core Graphics

pairs(
iris[,1:4],
main="Scatterplot Matrix"
)
```



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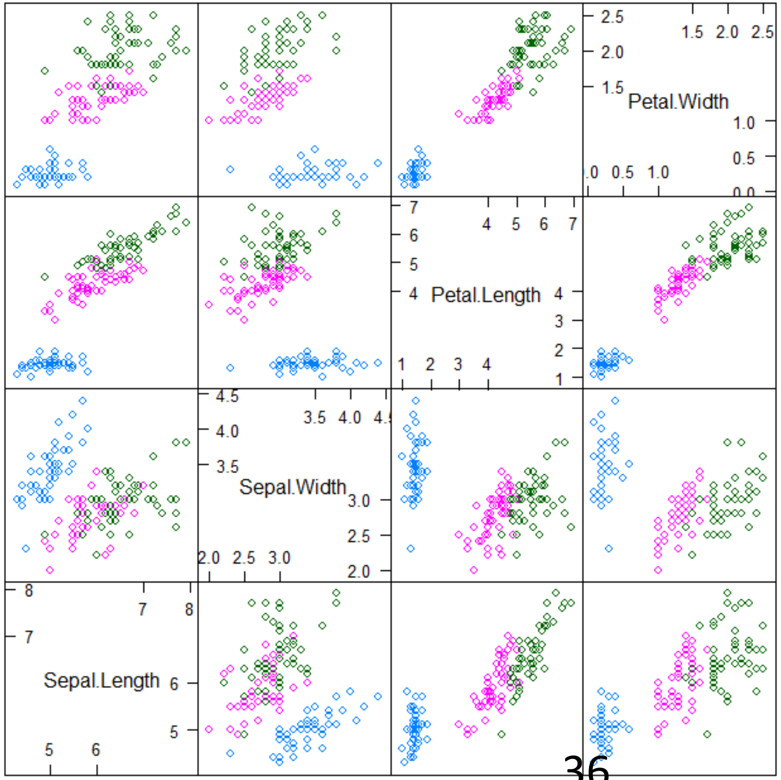
Scatterplot matrix



Scatter Plot Matrix

```
# Lattice Graphics

splom(iris[1:4],
groups=iris$Species)
```

In-class Exercise

- Using the "mtcars" dataset, predict mpg based on other columns.
- Create at least 2 different plots illustrating useful relationships in data and summarize your findings.

The “mtcars” dataset

```
data(mtcars)
head(mtcars)
```

```
> head(mtcars)
                   mpg cyl disp  hp drat    wt  qsec vs am gear carb
Mazda RX4         21.0   6  160 110 3.90 2.620 16.46  0  1    4    4
Mazda RX4 Wag     21.0   6  160 110 3.90 2.875 17.02  0  1    4    4
Datsun 710        22.8   4  108  93 3.85 2.320 18.61  1  1    4    1
Hornet 4 Drive    21.4   6  258 110 3.08 3.215 19.44  1  0    3    1
Hornet Sportabout 18.7   8  360 175 3.15 3.440 17.02  0  0    3    2
Valiant           18.1   6  225 105 2.76 3.460 20.22  1  0    3    1
```

GGPLOT2 GRAPHICS

The “diamonds” dataset

```
library(ggplot2)
data(diamonds)
head(diamonds)
```

```
> head(diamonds)
  carat       cut color clarity depth table price    x    y    z
1  0.23     Ideal     E     SI2  61.5    55   326 3.95 3.98 2.43
2  0.21   Premium     E     SI1  59.8    61   326 3.89 3.84 2.31
3  0.23      Good     E     VS1  56.9    65   327 4.05 4.07 2.31
4  0.29   Premium     I     VS2  62.4    58   334 4.20 4.23 2.63
5  0.31      Good     J     SI2  63.3    58   335 4.34 4.35 2.75
6  0.24 Very Good     J    VVS2  62.8    57   336 3.94 3.96 2.48
```

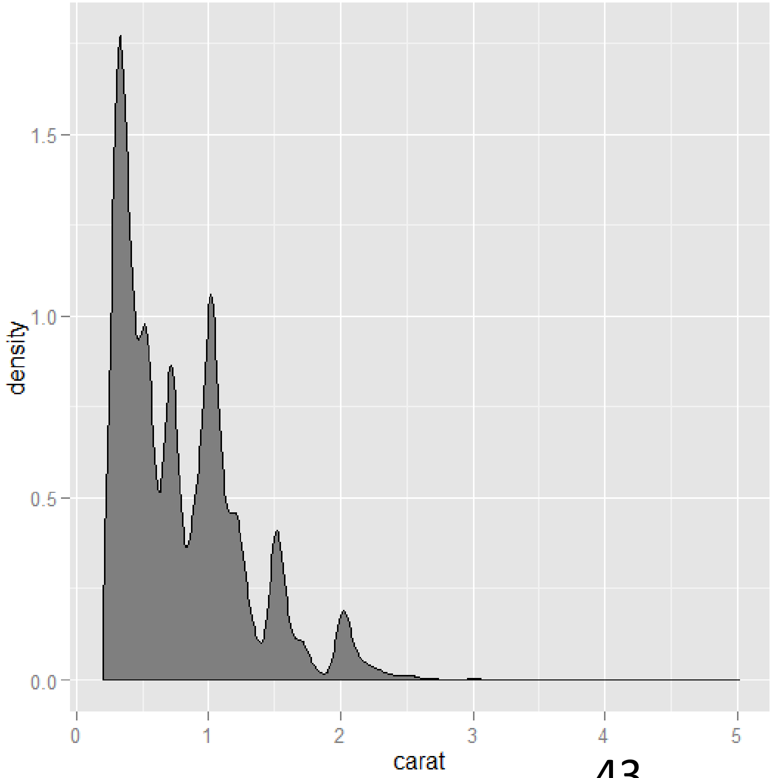
ggplot fundamentals

- ggplot() provides a blank canvas for plotting
- geom_*() creates actual graphical layers
  - geom_point()
  - geom_boxplot()
- aes() defines an "aesthetic" either globally or by layer

Layering

$$\underbrace{\texttt{ggplot(diamonds, aes())}}_{\text{Layer 1}} \quad + \quad \underbrace{\texttt{geom\_point()}}_{\text{Layer 2}}$$

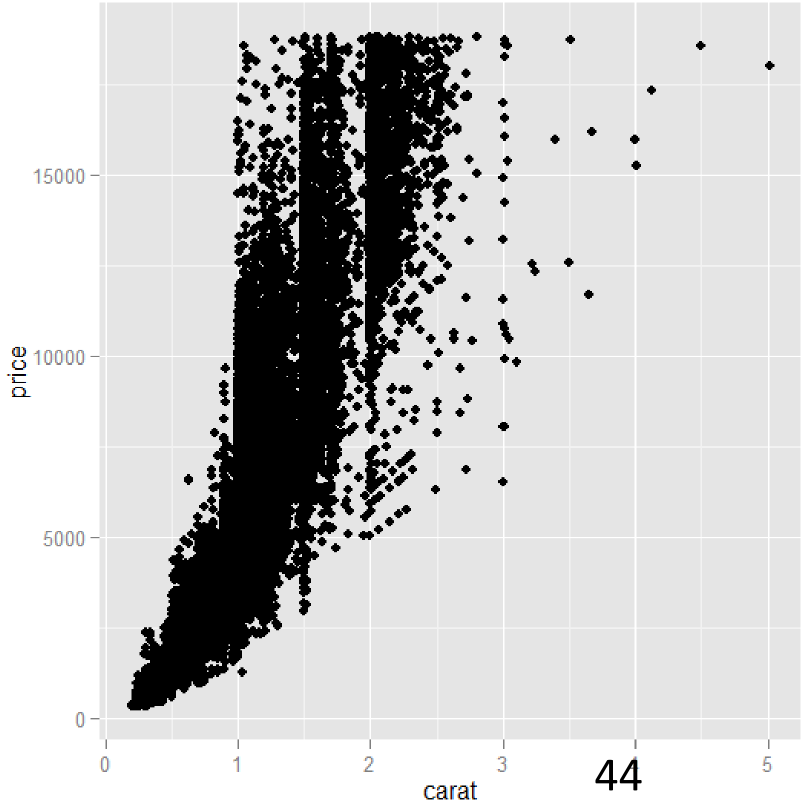
Density plot




```
ggplot(diamonds) +
geom_density(aes(x=carat),
fill="gray50")
```

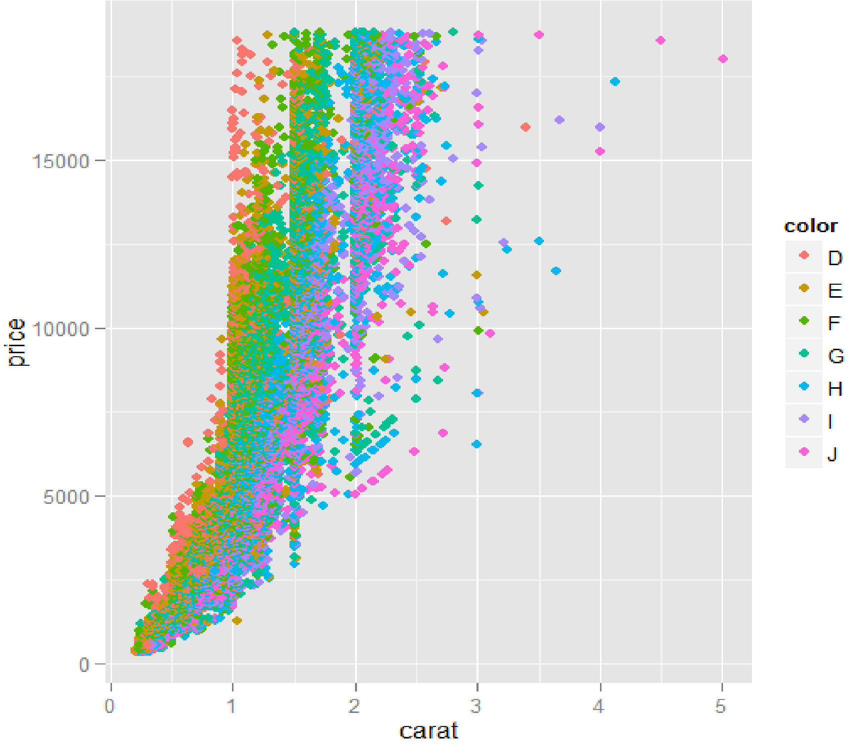
- Note the location of aes()

Scatterplot




```
ggplot(diamonds,
aes(x=carat,y=price)) +
geom_point()
```

ggplot object




```
# Store the plot for future
modification
g <- ggplot(diamonds,
aes(x=carat, y=price))

# add settings specific to
geom_point layer
g + geom_point(aes(color=color))
```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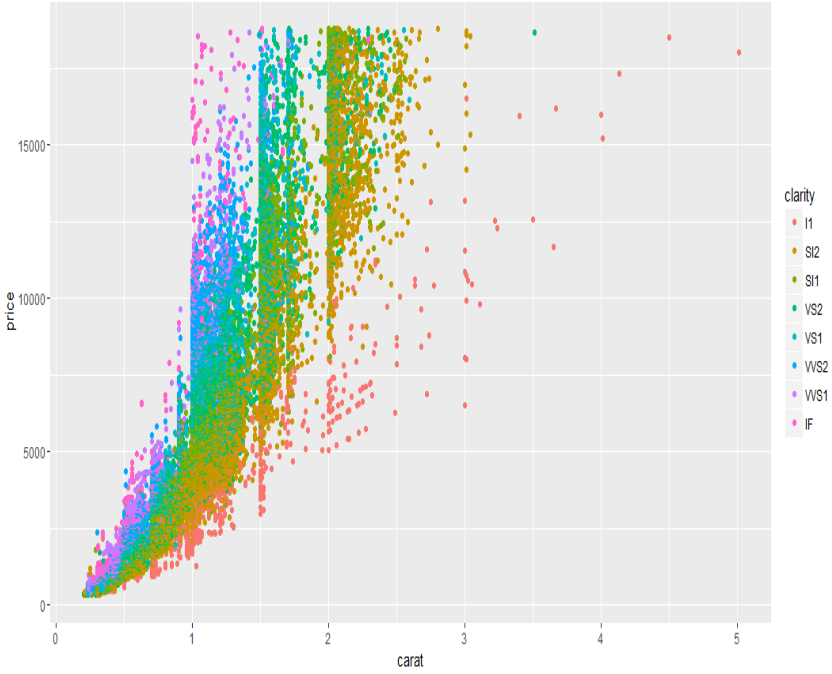
ggplot object




```
# Store the plot for future
modification
g <- ggplot(diamonds,
aes(x=carat, y=price))

# add settings specific to
geom_point layer
g +
geom_point(aes(color=clarity))
```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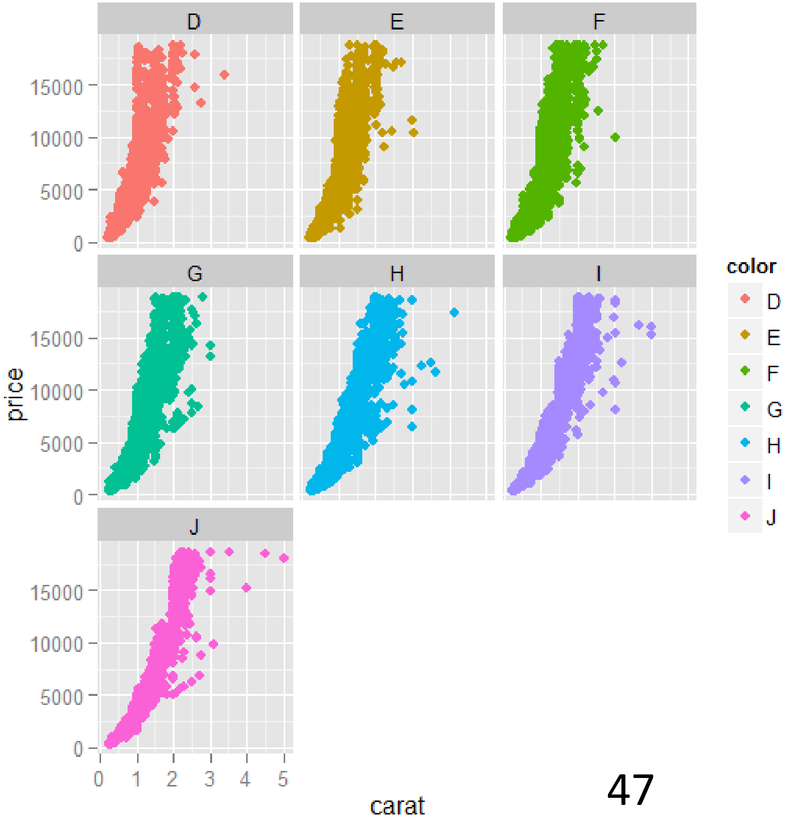
Separating segments




```
g +
geom_point(aes(color=color)) +
facet_wrap(~ color)
```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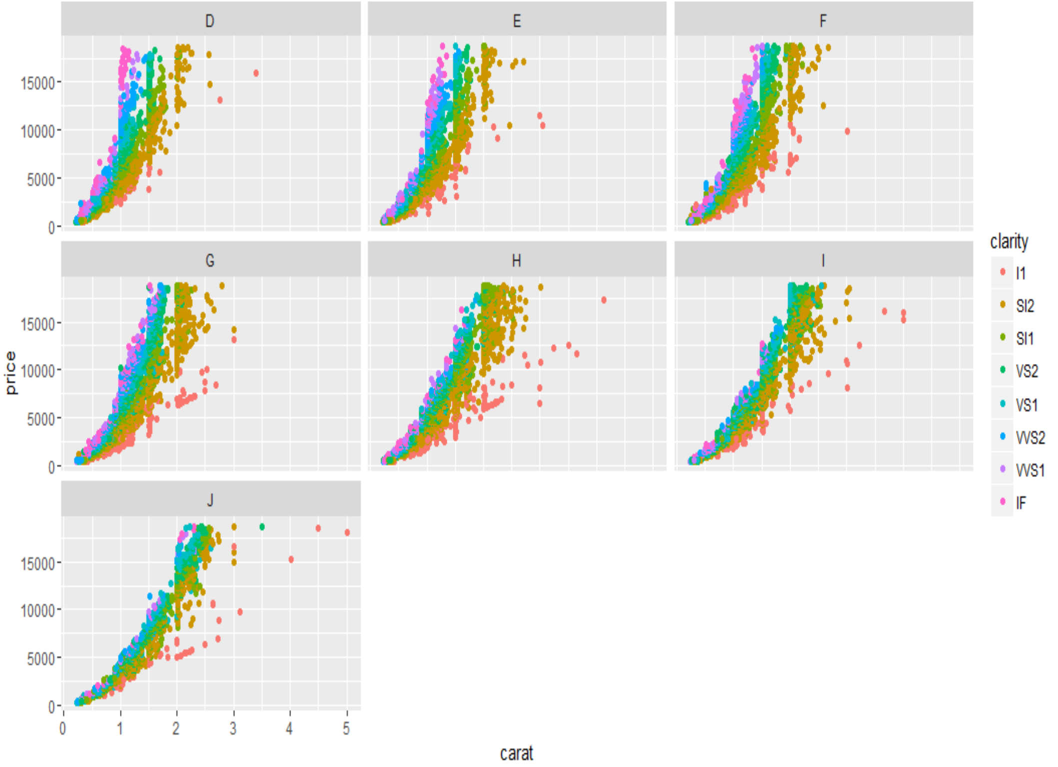
Separating segments




```
g +
geom_point(aes(color
=clarity)) +
facet_wrap(~ color)
```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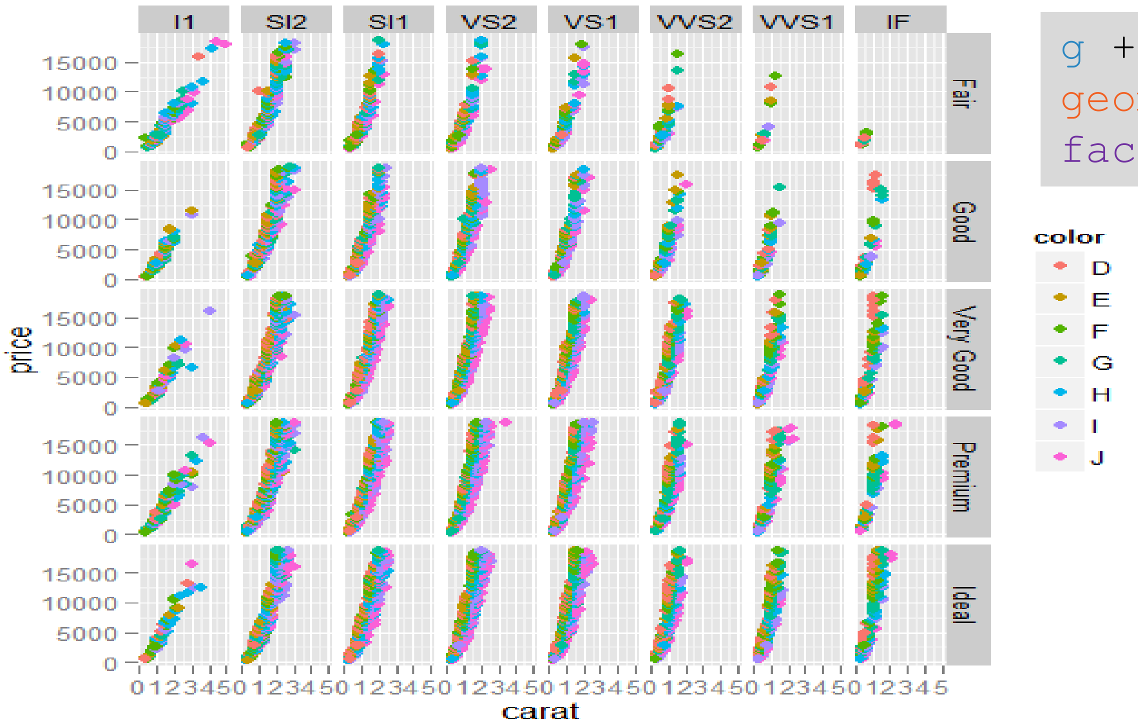
More segments!




```
g +
geom_point(aes(color=color))+
facet_grid(cut ~ clarity)
```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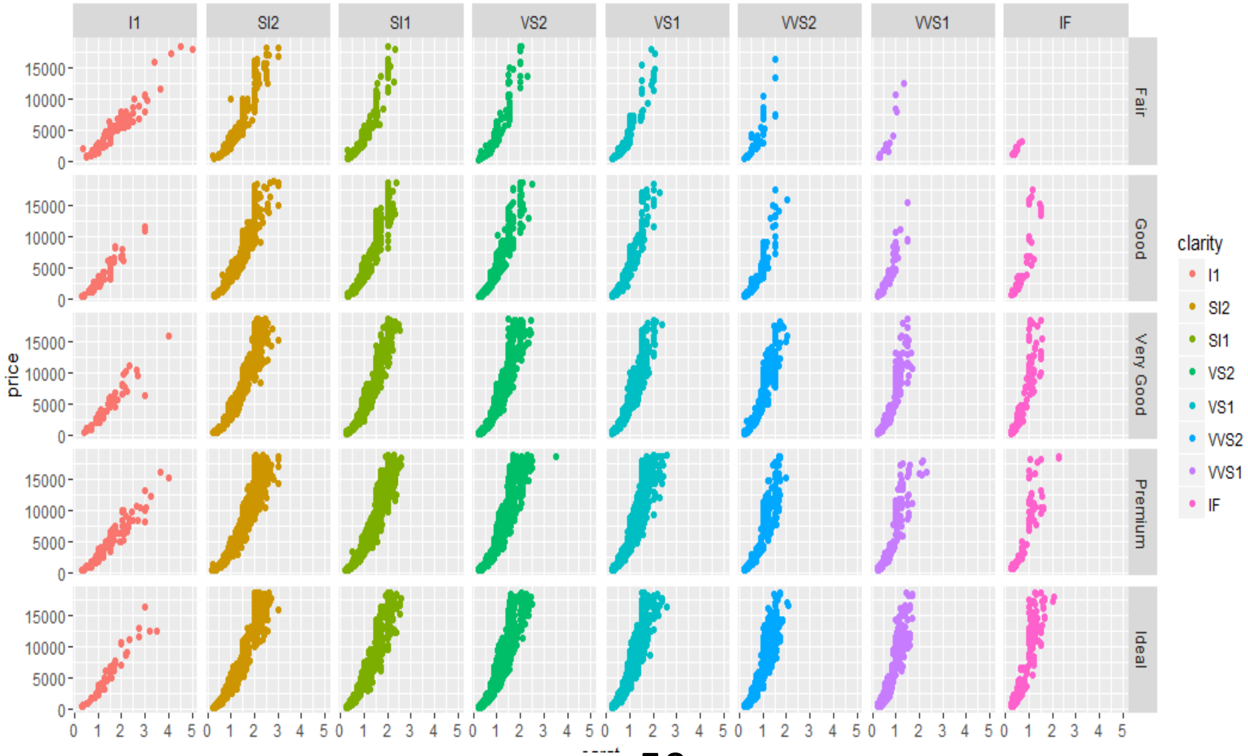
More segments!

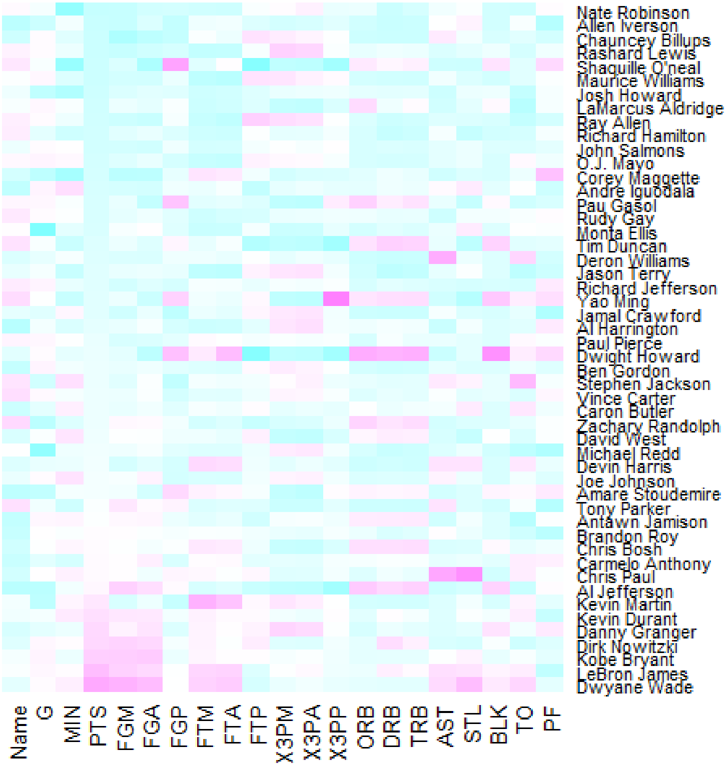



```
g +
geom_point(aes
(color=clarity
)) +
facet_grid(cut
~ clarity)
```

Heatmaps

- Helpful in understanding strength of relationship in a correlation or a distance matrix.
- Visualizes numbers by changing into colored cells.

Heatmaps



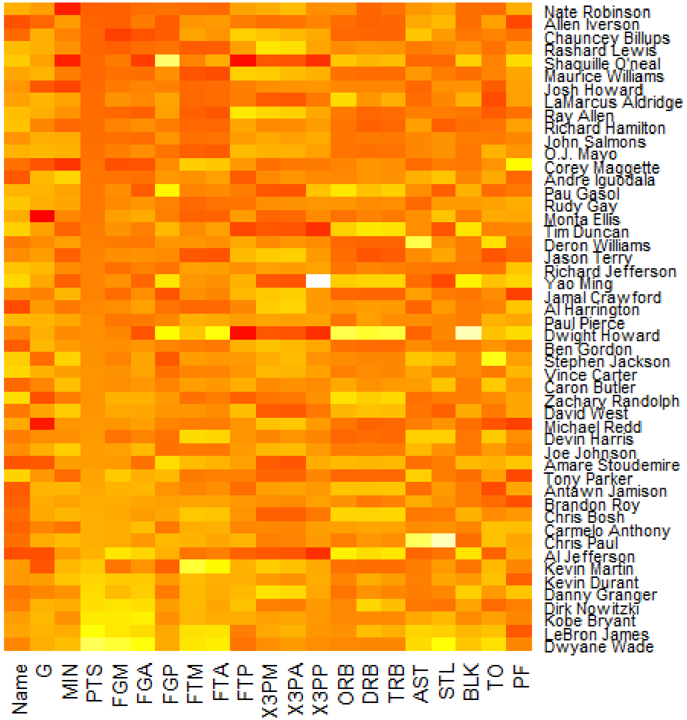

```
nba <-
read.csv("http://datasets.flowingdata.com/
ppg2008.csv", sep=",")
```

```
row.names(nba) <- nba$Name
```

```
nba_matrix <- data.matrix(nba)
nba_heatmap <- heatmap(nba_matrix,
Rowv=NA, Colv=NA, col = cm.colors(256),
scale="column", margins=c(5,10))
```

Source: http://flowingdata.com

Heatmaps




```
nba <-
read.csv("http://datasets.flowingdata.com/
ppg2008.csv", sep=",")
```

```
row.names(nba) <- nba$Name
```

```
nba_heatmap <- heatmap(nba_matrix,
Rowv=NA, Colv=NA, col =
heat.colors(256), scale="column",
margins=c(5,10))
```

Source: http://flowingdata.com

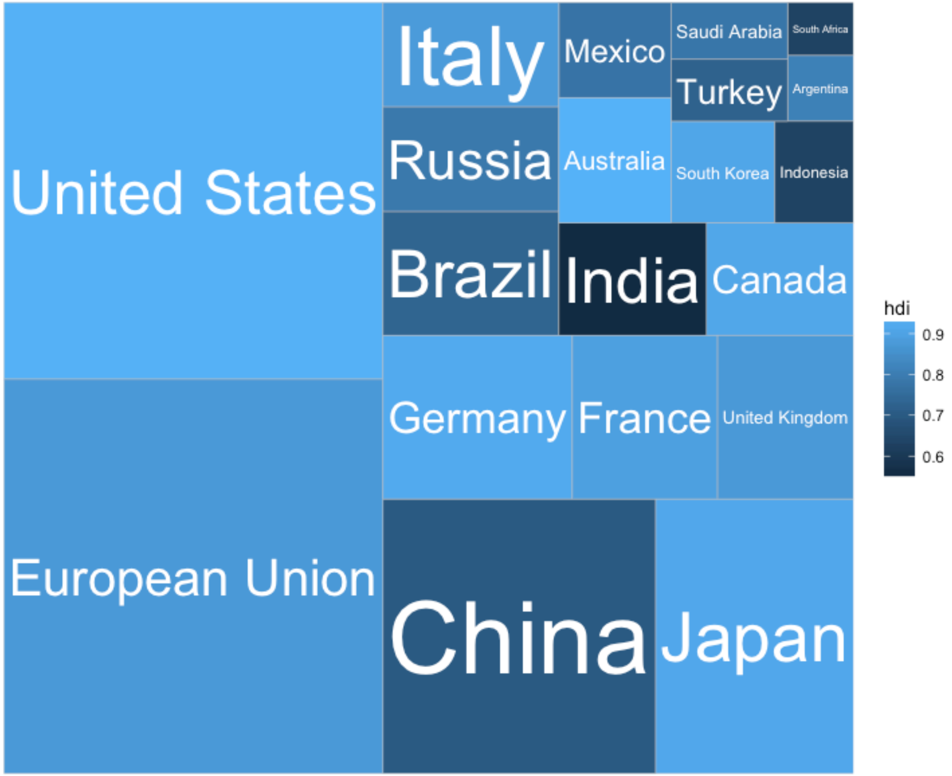
Treemap

- Powerful visualization of multi-dimensional values grouped by nested segments
- Provides an easy at-first-glance overview of a dataset's hierarchical structure
- Must specify variables for: area, fill, and label

Treemap

```
library(treemapify)
```

```
head(G20)
ggplot(G20, aes(area =
gdp_mil_usd, fill = hdi,
label = country)) +
geom_treemap() +
geom_treemap_text(color='whi
te', place='center',grow=T)
```



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Summary

- ✓ Basics of R
- ✓ Graphing in R – core, lattice, ggplot2, and treemapify
- ✓ Look at multiple types of graphs
- ✓ Visualize and segment data to gain more insights
- ✓ Identify key features
- ✓ Summarize findings

STORYTELLING WITH TITANIC

Finding the data set

- Set your working directory to the bootcamp root
- Load data in from "Datasets/titanic.csv"

Looking at the first few rows

```
titanic <- read.csv("titanic.csv")`
head(titanic)
```

```
> head(titanic)
  PassengerId Survived Pclass                                                Name    Sex Age SibSp Parch           Ticket    Fare Cabin Embarked
1           1        0      3                             Braund, Mr. Owen Harris   male  22     1     0        A/5 21171  7.2500                S
2           2        1      1 Cumings, Mrs. John Bradley (Florence Briggs Thayer) female  38     1     0         PC 17599 71.2833   C85        C
3           3        1      3                              Heikkinen, Miss. Laina female  26     0     0 STON/O2. 3101282  7.9250                S
4           4        1      1        Futrelle, Mrs. Jacques Heath (Lily May Peel) female  35     1     0           113803 53.1000  C123        S
5           5        0      3                            Allen, Mr. William Henry   male  35     0     0           373450  8.0500                S
6           6        0      3                                    Moran, Mr. James   male  NA     0     0           330877  8.4583                Q
> 
```

What features should we consider?

What is the data type of each column?

```
str(titanic)
```

```
'data.frame':		891 obs. of  12 variables:
 $ PassengerId: int  1 2 3 4 5 6 7 8 9 10 ...
 $ Survived   : int  0 1 1 1 0 0 0 0 1 1 ...
 $ Pclass     : int  3 1 3 1 3 3 1 3 3 2 ...
 $ Name       : Factor w/ 891 levels "Abbing, Mr. Anthony",..: 109 191 358 277 16 559 520 629 417 581 ...
 $ Sex        : Factor w/ 2 levels "female","male": 2 1 1 1 2 2 2 2 1 1 ...
 $ Age        : num  22 38 26 35 35 NA 54 2 27 14 ...
 $ SibSp      : int  1 1 0 1 0 0 0 3 0 1 ...
 $ Parch      : int  0 0 0 0 0 0 0 1 2 0 ...
 $ Ticket     : Factor w/ 681 levels "110152","110413",..: 524 597 670 50 473 276 86 396 345 133 ...
 $ Fare       : num  7.25 71.28 7.92 53.1 8.05 ...
 $ Cabin      : Factor w/ 148 levels "","A10","A14",..: 1 83 1 57 1 1 131 1 1 1 ...
 $ Embarked   : Factor w/ 4 levels "","C","Q","S": 4 2 4 4 4 3 4 4 4 2 ...
```

Casting

Set target column as a factor

```
titanic$Survived <- as.factor(titanic$Survived)
```

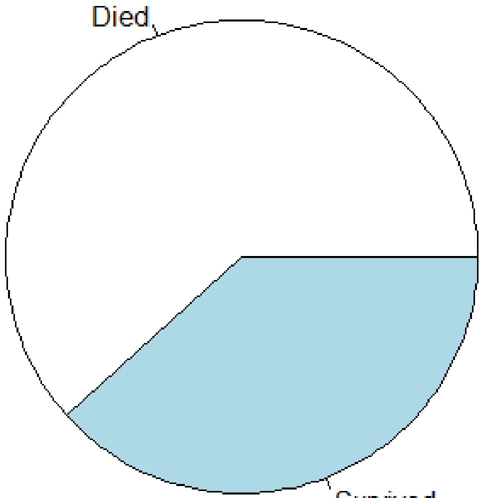
Rename factors and columns

```
levels(titanic$Survived) <- c("Dead", "Survived")
levels(titanic$Embarked) <- c("Unknown", "Cherbourg",
                              "Queenstown", "Southampton")
str(titanic[,c("Embarked","Survived")])
```

```
'data.frame':  891 obs. of  2 variables:
 $ Embarked: Factor w/ 4 levels
"Unknown","Cherbourg",..: 4 2 4 4 4 3 4 ...
 $ Survived: Factor w/ 2 levels "0","1": 1 2 2 2
1 1 1 1 2 2 ...
```

Class distribution: Pie Chart

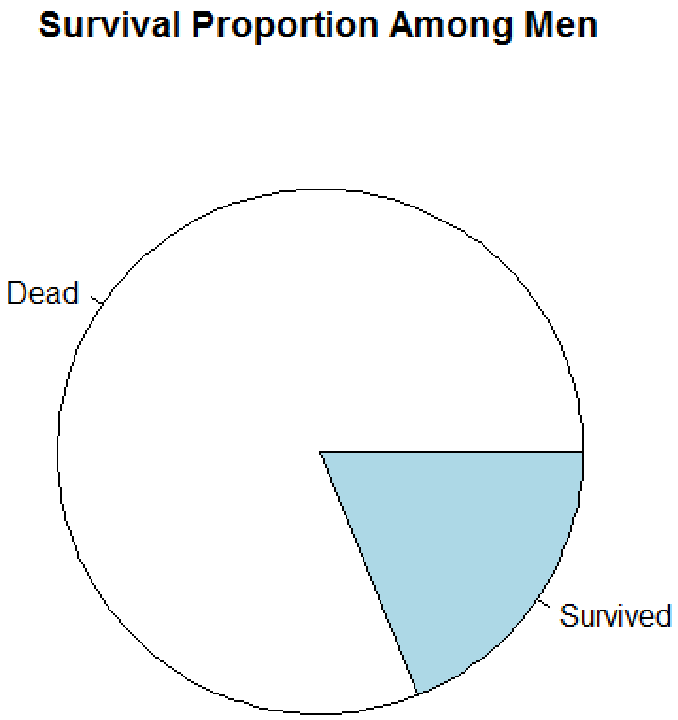
```
survivedTable <- table(titanic$Survived)
pie(survivedTable, labels=c("Died", "Survived"))
```

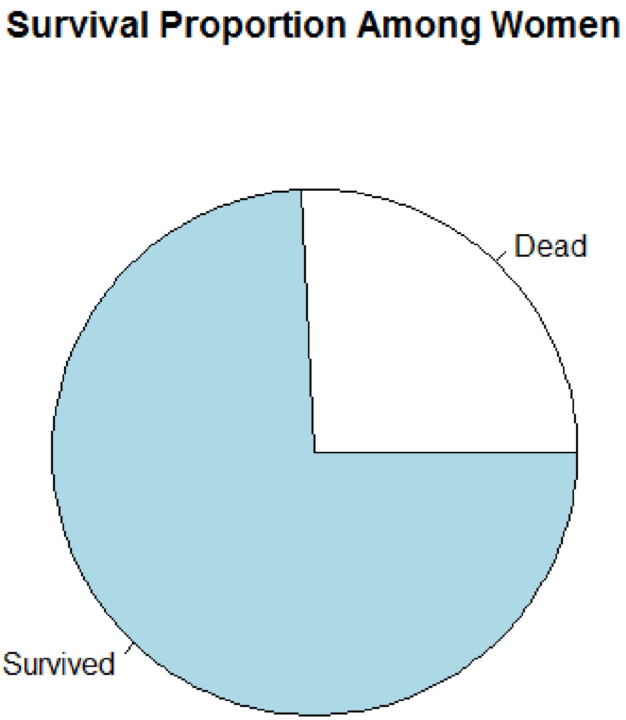


Is Sex a good predictor?

```
#Identify where sex = male for all columns
male <- titanic[titanic$Sex == "male",]

#Identify where sex = female for all columns
female <- titanic[titanic$Sex == "female",]

par(mfrow=c(1,2)) #two figures arranged in 1 row and 2 columns
pie(table(male$Survived), labels=c("Dead","Survived"))

pie(table(female$Survived), labels=c("Dead","Survived"))
```

Is Sex a good predictor?





Is Age a good predictor?

```
summary(titanic$Age)
```

Min.	1st Qu.	Median	Mean	3rd Qu.	Max.	NA's
0.42	20.12	28.00	29.70	38.00	80.00	177

- How about by Survival?

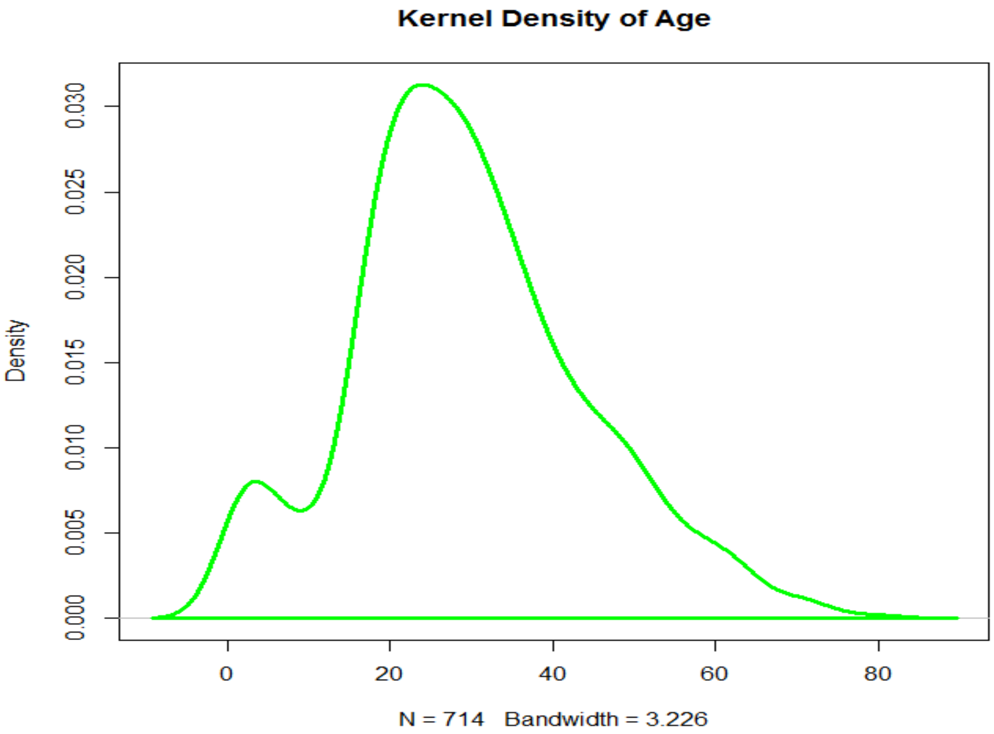
```
summary(titanic[titanic$Survived
=="Dead",]$Age)
```

Min.	1st Qu.	Median	Mean	3rd Qu.	Max.	NA's
1.00	21.00	28.00	30.63	39.00	74.00	125

```
summary(titanic[titanic$Survived
=="Survived",]$Age)
```

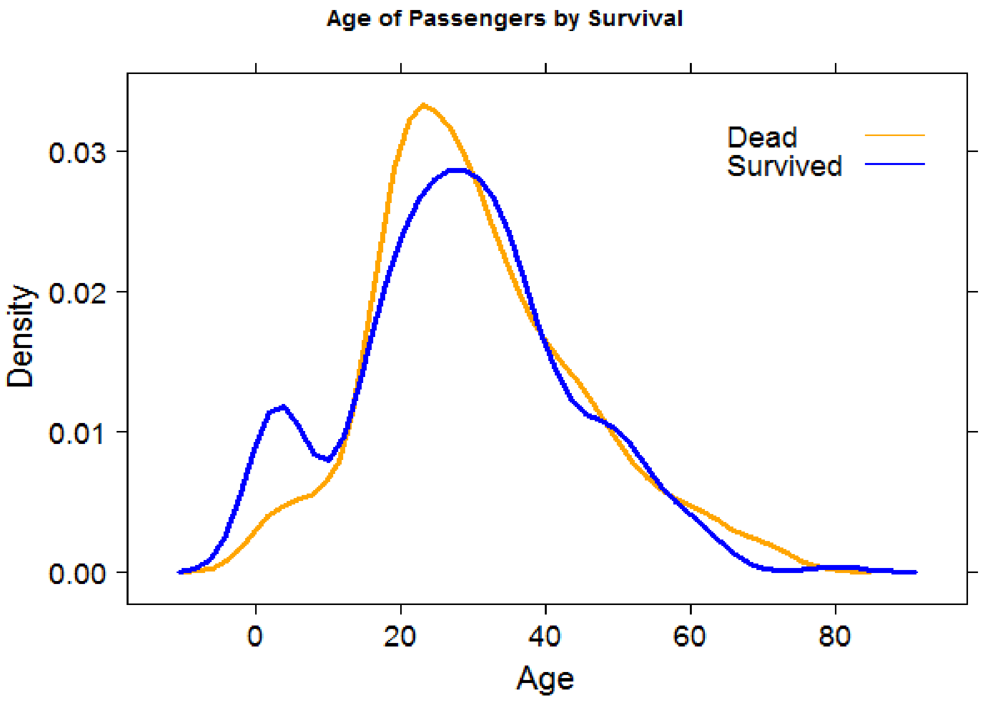
Min.	1st Qu.	Median	Mean	3rd Qu.	Max.	NA's
0.42	19.00	28.00	28.34	36.00	80.00	52

Sample solution




```
density(titanic$Age)
#NAs prevent this
> d <-
density(na.omit(titanic$Ag
e))
> plot(d, main="Kernel
Density of Age")
>
polygon(d,border="green",l
wd=3)
```

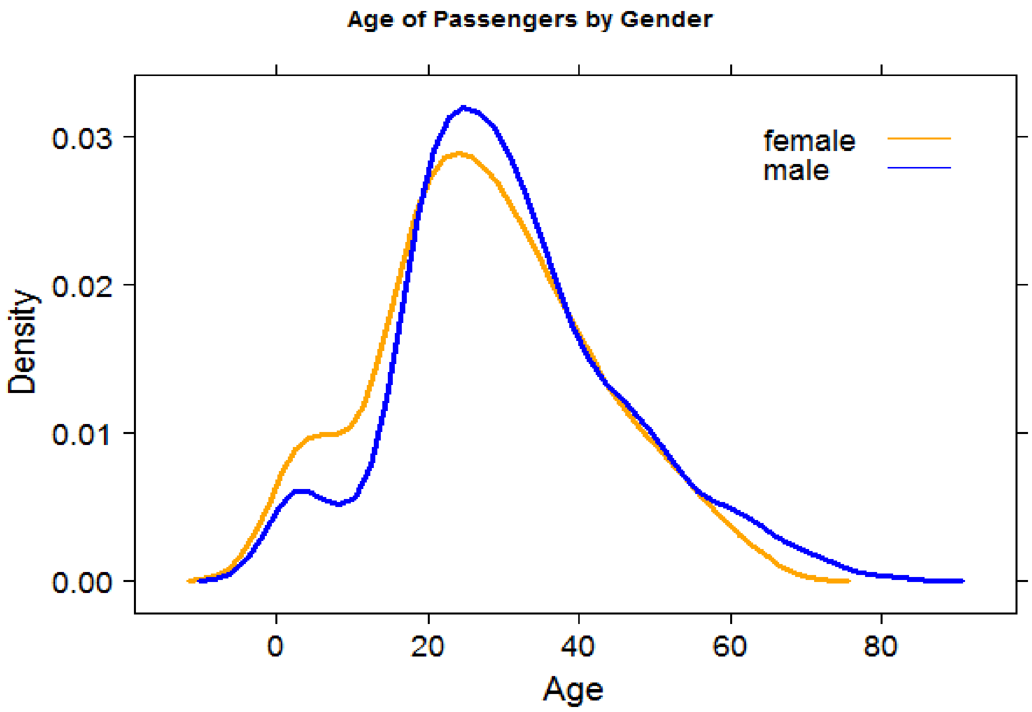
Is Age a good predictor for Survival?




```
densityplot(~Age,data=titanic,groups=Survived,plot.points=F, lwd=3)
```

Note: won't work with missing values

Is Age a good predictor for Gender?




```
densityplot(~ Age,
data=titanic, groups=Sex,
plot.points=F, lwd=3)
```

QUESTIONS