

# APMA E4990.02: Introduction to Regression

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Lecture 2

# Checklist

- Finish the basic UNIX example I made an announcement for?
- Install Github, Anaconda, Cygwin (if using Windows).
- Clone Github repo AND repo for the assignment - see Homework 1 in the homework folder. Due next week.

# Waitlist?

- We only got approval for additional TA resources very recently SO:
- If you want any of the students in your waitlist to enroll AFTER THE CAP IS CHANGED, they will have to fill out a Registration Adjustment Form  
<http://registrar.columbia.edu/sites/default/files/content/reg-adjustment.pdf>
- OR PICK UP A FORM AT THE FRONT ON YOUR WAY OUT.
- **Tiffany Simon:** tms26@columbia.edu

# Unix Importance

- All production level data science code is run on Linux/Unix servers. The scripts used to pull in data, merge it and run models is all run via scripts on these machines.
- Commonly need to move, edit files on remote servers, scp various pieces of code/data, start AWS clusters, etc. All of this requires basic knowledge of the command line.
- My opinion is that learning without doing is pointless, so we will learn Linux by example when you set up your own web server.

# Unix Knowledge

## File manipulation first

- mkdir, ls, cd, touch, mv (-r), cp (-r)

## Making life on the command line easier

- emacs shortcuts on the command line
- up arrow, CTRL-R history search, CTRL-E, CTRL-U

## A terminal based editor

- pico, nano, vim or emacs.

## Text manipulation and display

- cat head tail grep (-r)

## Network

- curl, wget

## STDIN/OUT

- redirection (>1 >2),

<http://ryanstutorials.net/linuxtutorial/cheatsheet.php>

## Process chaining and standard in/out

- pipe (|), redirecting

## Process manipulation

- bg, CTRL-Z, CTRL-C
- environmental variables
- ps (list of processes)
- what is TOP
- lsof (what files are open by what processes)

## Common tasks:

- operating on a bunch of files using xargs or a loop
- shell scripting
- .bashrc file
- working on remote machines with sshd
- awk/sed for file processing.



```
pi@raspberrypi:~ $ ls -l /usr/lib
total 11
drwxr-xr-x 3 bin 128 Sep 22 05:45 dict
drwxr-xr-x 2 dmr 32 Sep 22 05:48 dmr
drwxr-xr-x 5 bin 416 Sep 22 05:46 games
drwxr-xr-x 3 sys 496 Sep 22 05:42 include
drwxr-xr-x 10 bin 520 Sep 22 05:43 lib
drwxr-xr-x 11 bin 176 Sep 22 05:45 libn
drwxr-xr-x 3 bin 208 Sep 22 05:46 mdec
drwxr-xr-x 2 bin 80 Sep 22 05:46 pub
drwxr-xr-x 6 root 96 Sep 22 05:45 spool
drwxr-xr-x 13 root 208 Sep 22 05:42 src
# ls -l /usr/libm
total 0
#
```

## Basic Navigation

### pwd

Where am I in the system.

### ls [path]

Perform a listing of the given path or your current directory.

Common options: -l, -h, -a

### cd [path]

Change into the given path or into your home directory.

### Path

A description of where a file or directory is on the filesystem.

### Absolute Path

One beginning from the root of the file system (eg. ./etc/sysconfig ).

### Relative Path

One relative to where you currently are in the system (eg. Documents/music ).

### ~ (tilde)

Used in paths as a reference to your home directory (eg. ~/Documents ).

### . (dot)

Used in paths as a reference to your current directory (eg. ./bin ).

### .. (dot dot)

Used in paths as a reference to your current directories parent directory (eg. ../bin ).

### TAB completion

Start typing and press TAB. The system will auto complete the path. Press TAB twice and it will show you your alternatives.

## More About Files

### file [path]

Find out what type of item a file or directory is.

### Spaces in names

Put whole path in quotes ( " ) or a backslash ( \ ) in front of spaces.

### Hidden files and directories

A name beginning with a . (dot) is considered hidden.

## Permissions

### r (read) w (write) x (execute)

### Owner or User, Group and Others

### ls -l [path]

View the permissions of a file or all items in a directory.

### chmod <permissions> <path>

Change permissions. Permissions can be either shorthand (eg. 754) or longhand (eg. g+x).

## Manual Pages

### man <command>

View the man page for a command.

### man -k <search term>

Search for man pages containing the search term.

### Press q to exit man pages

## Vi / Vim

### View our Vim Cheat sheet

## Filters

### head

Show the first n lines.

### tail

Show the last n lines.

### sort

Sort lines in a given way.

### wc

How many words, characters and lines.

### grep

Search for a given pattern.

### See more on our Grep Cheat sheet

More filters can be found [here](#).

## File Manipulation

### mkdir <directory name>

Create a directory

### rmdir <directory name>

Remove a directory (only if empty).

### touch <file name>

Create a blank file.

### cp <source> <destination>

Copy the source file to the destination.

### mv <source> <destination>

Move the source file to the destination.

### rm <path>

Remove a file or directory.

### Common options: -r -f

## Wildcards

### May be used anywhere in any path.

### \*

Zero or more characters (eg. b\*).

### ?

Single character (eg. file.???).

### []

Range (eg. b[aio]).

## Wildcards

### May be used anywhere in any path.

### \*

Zero or more characters (eg. b\*).

### ?

Single character (eg. file.???).

### []

Range (eg. b[aio]).

## Process Management

### CTRL + C

Cancel the currently running process.

### kill <process id>

Cancel the given process.

### Include the option -9 to kill a stubborn process.

### ps

Obtain a listing of processes and their id's.

Including the option aux will show all processes.

### CTRL + Z

Pause the currently running process and put it in the background.

### jobs

See a list of current processes in the background.

### fg <job number>

Move the given process from the background to the foreground.

# Review from Last Time

- Introduced examples of machine learning applications and methods.
  - Used examples from Amazon, Netflix, The New York Times, Booking.com and Tinder.
  - Discussed **Predictive**, **Descriptive** and **Prescriptive** methods of machine learning.
  - Linear Regression, Decision Trees, Graph Diffusion models.
- How do we learn from data?
  - Overall pipeline: Data pulling -> Merging/Cleaning -> Model Training -> Evaluation.
  - MNIST digit recognition via KNN.
  - Logistic Regression for churn at The New York Times.
- Model Evaluation and Complexity
  - Cross Validation - having a hold-out set for testing.
  - Hyperparameter optimization and overfitting. How complex is “just right”?

# Outline

- Introduction to Linear Regression
  - Cross Validation reminder - the test/train split.
  - Definition, and comparison of  $L_p$  norms.
  - How do we find the solution? The special case of  $p=2$ .
  - When do unique solutions exist?
  - The problem of dependent features.
- General Solution Finding Procedure - Gradient Descent
  - Gradient Descent, and Introduction to Convex Optimization.
  - Concrete Examples of Linear Regression and Gradient Descent in Python in an iPython Notebook.
- Introduction to Decision Trees
  - How decision trees are constructed - variance reduction.
  - Concrete Examples of Decision Trees in Python with an iPython Notebook.

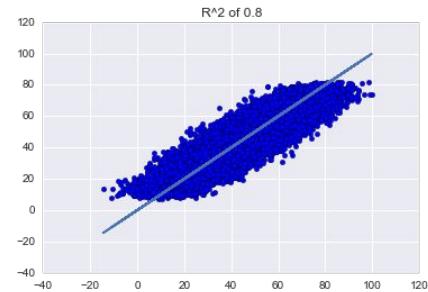
# What is a model?

- We generally have some input data  $X$  and wish you predict some outcome  $y$ .
- A simple and common assumption is that the data has some linear response:

$$y = \beta^T \cdot x$$

- But we can't predict the future - there is a natural uncertainty present in any model. Thus  $y$  must follow some probability distribution.

$$p(y, x)$$

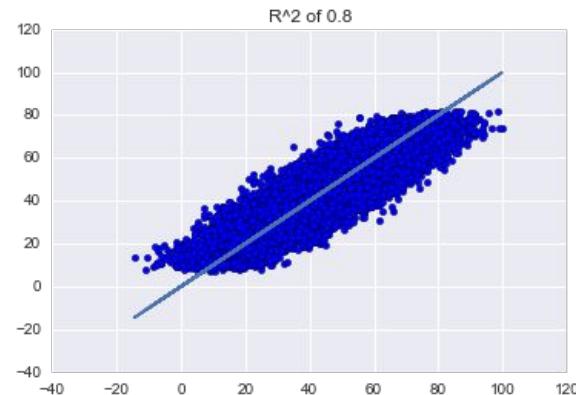


# Introduction to Regression

Given a collection of points to  $(x_i, y_i)$   
learn from:

Can we find a function  $f : X \rightarrow Y$   
minimizing the distance to the  
data.

$$\mathcal{L}(y, f) := \sum_{i=1}^N d(y_i, f(x_i))$$



$d(x, y)$  - Is a metric (distance between two points)

# Testing/Training Split

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We always learn from training data

# Splitting data into Testing and Training

In [72]: `x_train.head()`

Out[72]:

	account	const	hotel_rating	location	price_per_night_avg	purchase_velocity_lastweek	rooms_left	sellouts_total
0	34961	1	8.9	8.5	409	232	226	3
1	25510	1	4.5	7.1	189	325	265	4
2	95668	1	7.7	6.0	377	287	475	4
3	38365	1	1.7	8.9	350	205	289	8
4	51131	1	0.6	5.2	542	301	66	2

In [73]: `x_test.head()`

Out[73]:

	account	const	hotel_rating	location	price_per_night_avg	purchase_velocity_lastweek	rooms_left	sellouts_total
40	33846	1	7.8	3.1	294	85	371	1
41	17771	1	7.8	6.8	572	377	384	6
42	10967	1	0.9	6.9	191	95	79	8
43	37812	1	2.9	3.2	418	65	4	5
44	61968	1	3.5	4.2	222	201	356	3

The image shows two side-by-side screenshots of hotel booking platforms. The left screenshot displays a listing for 'Dorsett Shepherds Bush' in London, showing a double room with a 34% discount. It includes a photo of the room, guest reviews (4.5 stars), and booking details like 'Free cancellation - PAY LATER'. The right screenshot shows a listing for 'City Marque Albert Serviced Apartments' in Central London, featuring a studio apartment. Both screenshots include a 'Book now' button.

Let  $y$  be the number of rooms booked at a hotel tomorrow.

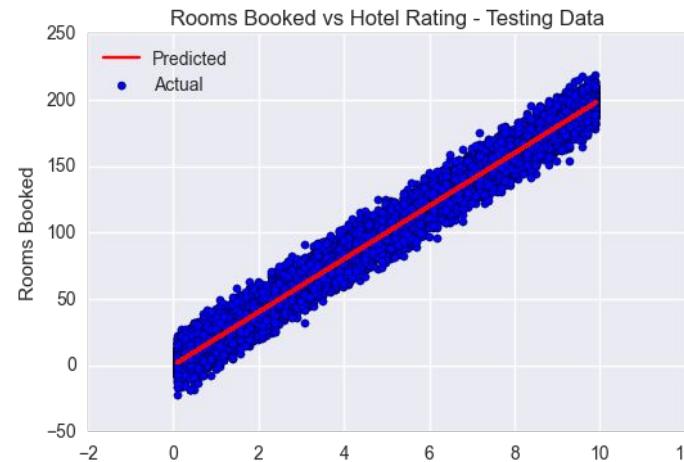
- First, we randomly split our data into **testing and training subsets**.
- The most important thing in machine learning is **generalizability** - we must be able to **make predictions on unseen data** that works.
- If you make a model **complex enough**, you can **often get very good accuracy** on the **training data**, but it **won't generalize** (next lecture).

# Splitting data into Testing and Training



```
In [72]: x_train.head()  
Out[72]:
```

	account	const	hotel
0	34961	1	8.9
1	25510	1	4.5
2	95668	1	7.7
3	38365	1	1.7
4	51131	1	0.6



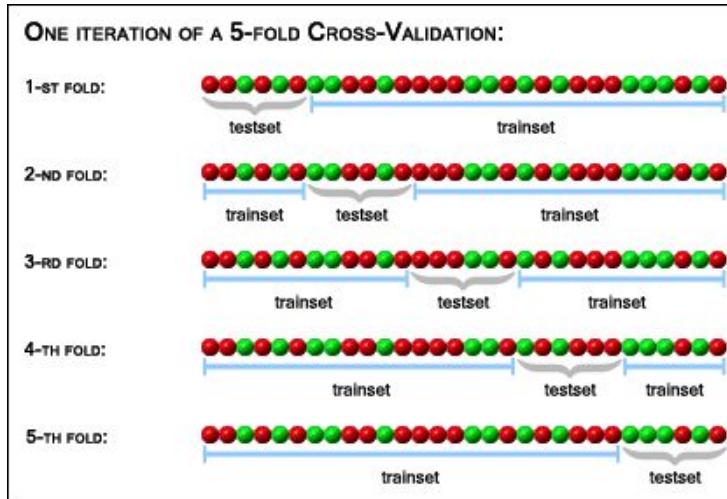
```
In [73]: x_test.head()  
Out[73]:
```

	account	const	hotel
40	33846	1	7.8
41	17771	1	7.8
42	10967	1	0.9
43	37812	1	2.9
44	61968	1	3.5

- We must use the **training data to learn the model**.
- Then from the model we've learned, we **evaluate it on the testing data**.

$$y = 20 * \text{hotel-rating} + \text{noise}$$

# Cross Validation



- Generally instead of just taking say **80% of your data for training, 20% for testing (for example)**, we randomly split the data into several ‘folds’.
- In **k-fold cross-validation**, the original sample is randomly partitioned into **k equal sized subsamples**. Of the **k** subsamples, **a single subsample is retained as the validation data** for testing the model, and **the remaining  $k - 1$  subsamples are used as training data**.

# Linear Regression

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# Why $p = 2$ ?

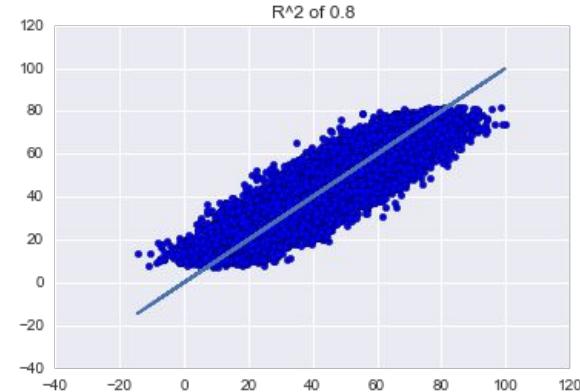
$$\mathcal{L}(y, \beta \cdot x) = \frac{1}{N} \sum_{i=1}^N |y_i - \beta \cdot x_i|^p$$

## Most common case:

- $p=2$ 
  - Penalizes outliers much more.
  - Less sparse coefficients.
  - Has an **analytical solution**.
  - Unique solution when features are linearly independent.
  - Compatible with CLT (will see later!)
- $p=1$ 
  - More sparse coefficients (zero valued).
  - Helps eliminate collinear features.
  - Degenerate solutions.
  - Deals with outliers better (why?)

## Linear assumption:

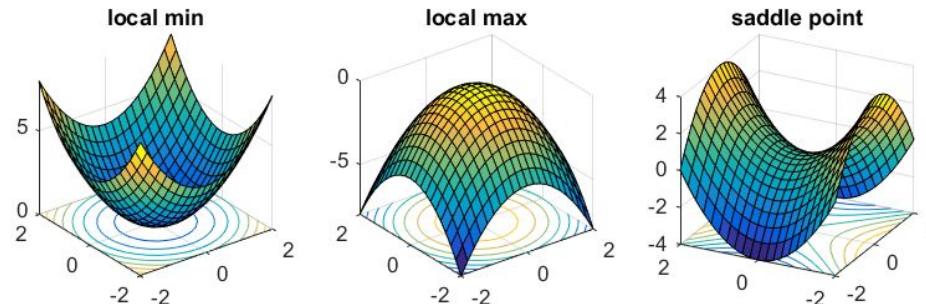
$$f(x_i) = \beta \cdot x_i$$



**Question:** Why would  $p = \infty$  be a bad choice for machine learning?

# Convex Analysis (Calculus)

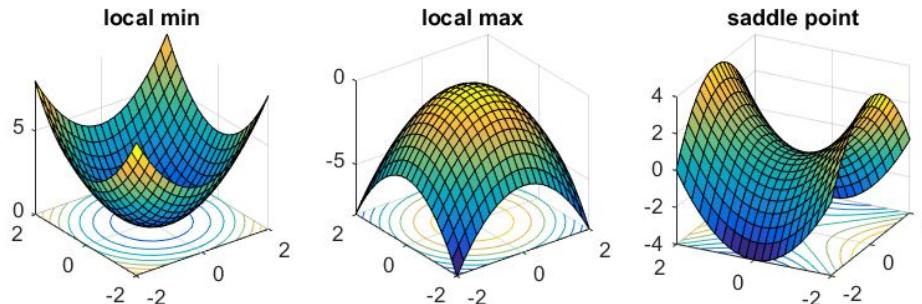
- Recall from last time that our goal is often find some convex function  $\mathcal{L} : \mathbf{S} \rightarrow \mathbb{R}$  such that we learn the rules of our model by its minimization (ie. coefficients). These are referred to as **parametric methods**.
- We can also “fit” data to a probability by **maximizing** the **likelihood** - concave problem.
- This is **not always the case**, with algorithms such as **decision trees** or **neural nets**. However, it covers many cases in supervised learning. These are called **non-parametric methods** (to be covered in this class or next).



# Convex Analysis (Calculus)

- Assume  $\mathcal{L} : \mathbf{S} \rightarrow \mathbb{R}$  is continuously differentiable and convex, ie.  
$$\mathcal{L}''(\beta) \geq 0 \text{ for all } \beta \in \mathbf{S}$$
  
$$\exists \beta_0 \text{ such that } \mathcal{L}(\beta_0) \leq \mathcal{L}(\beta) \text{ for all } \beta \in S$$
- There f has a minimum value. Moreover, the minimum is unique when f is strictly convex, ie.

$$\mathcal{L}''(\beta) \geq c > 0 \text{ for all } \beta \in \mathbf{S}$$



# Ordinary Least Squares ( $p=2$ )

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Analytical Solutions and Stability Analysis

# Analytical Solution to OLS

$$\mathcal{L}(\beta) = \frac{1}{N} \sum_{i=1}^N |y_i - \beta \cdot x_i|^2$$

**Claim:** When  $p = 2$  and the matrix  $X^T X$  is invertible, the above is minimized uniquely by

$$\beta = (X^T X)^{-1} X^T y$$

The solution can make sense when  $X^T X$  isn't invertible (in practice it often can be "almost not invertible", but solutions will be unstable and degenerate (more later))

# Analytical Solution to OLS when p=2

**Proof:**

$$\mathcal{L}(\beta) = \frac{1}{N} \sum_{i=1}^N |y_i - \beta \cdot x_i|^2$$

- Need linearly independent features for a unique inversion.

Differentiating, we have

$$\frac{\partial \mathcal{L}}{\partial \beta_j} = -\frac{2}{N} \sum_{i=1}^N (y_i - \beta \cdot x_i)x_j$$

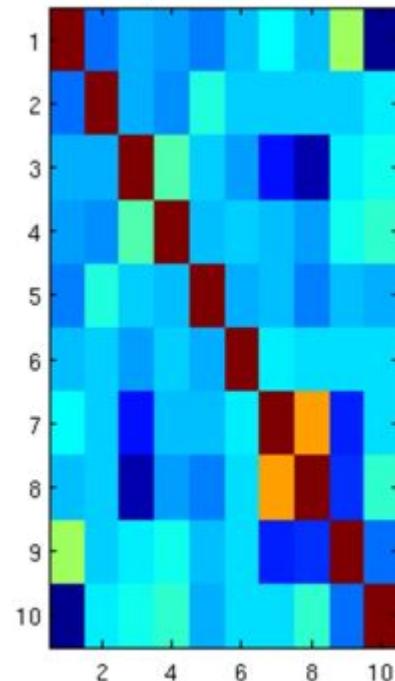
$$\frac{\partial \mathcal{L}}{\partial \beta_j} = 0 \Rightarrow \boxed{\frac{1}{N} \sum_{i=1}^N (\beta \cdot x_i)x_{ij} = \frac{1}{N} \sum_{i=1}^N y_i x_{ij}.} \Rightarrow X^T X \beta = X^T y.$$

$$\Rightarrow \beta = (X^T X)^{-1} X^T y$$

# Analytical Solution to OLS

It easily follows that  $\frac{d^2\mathcal{L}}{d^2\beta} = \frac{2}{N}X^T X$

- From this, it follows there is a **unique solution** iff the **eigenvalues** of the above matrix are **strictly positive**.
- This occurs precisely when **X has linearly independent features**.
- If X is **mean centered and normalized**, the above matrix is equivalent to the **correlation matrix**.



# Advertisement Example

What are the features?

**TV:** advertising dollars spent on TV for a single product in a given market (in thousands of dollars)

**Radio:** advertising dollars spent on Radio

**Newspaper:** advertising dollars spent on Newspaper

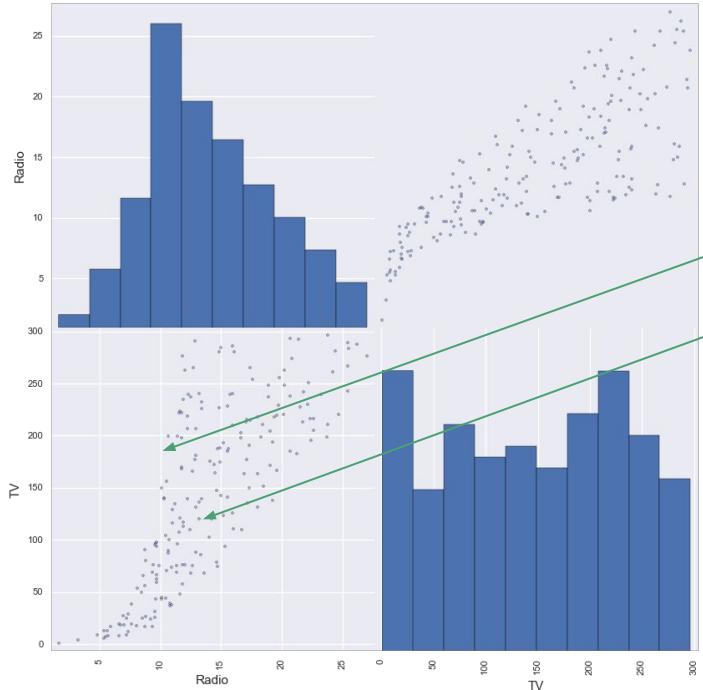
**Goal:** Predict the number of sales in a given market based on the advertising in TV, Radio and Newspaper.

```
In [263]: # read data into a DataFrame
import pandas as pd
import pylab as plt
import seaborn
from sklearn.linear_model import LinearRegression
import numpy as np
import random
import json
pd.set_option('display.max_columns', 500)
%matplotlib inline

df = pd.read_csv('http://www-bcf.usc.edu/~gareth/ISL/Advertising.csv', index_col=0)
df.head()
```

	TV	Radio	Newspaper	Sales
1	230.1	37.8	69.2	22.1
2	44.5	39.3	45.1	10.4
3	17.2	45.9	69.3	9.3
4	151.5	41.3	58.5	18.5
5	180.8	10.8	58.4	12.9

# Understanding correlation



Vectors are in n dimensions  
(number of data points)

Note: This is not the same as the data you will do for the homework!

$$\mathbb{R}^{2 \times n} \times \mathbb{R}^{n \times 2}$$

$$X^T X = \begin{bmatrix} 150 & 160 & \dots \\ 17 & 12 & \dots \\ \dots & \dots & \dots \end{bmatrix} \begin{bmatrix} 150 & 17 \\ 160 & 12 \\ \dots & \dots \end{bmatrix}$$

$$[X^T X]_{ij} = \mathbf{x}_i \cdot \mathbf{x}_j$$

$i =$  Radio



$j =$  TV



$$[\text{Corr}]_{ij} = \frac{\mathbf{x}_i \cdot \mathbf{x}_j}{\|\mathbf{x}_i\| \|\mathbf{x}_j\|} = \cos(\delta_{ij})$$

# Eigenvalues of Covariance Matrix

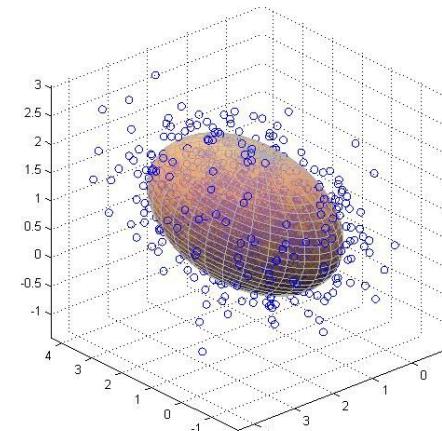
$$\lambda_1 = \max_{\|y\|=1} y^T A y$$

$$y^T A y = \langle y, A y \rangle$$

$$y = v + \epsilon w$$

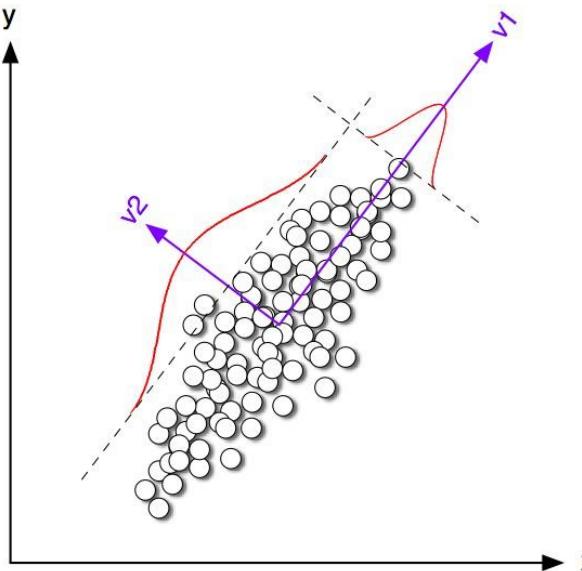
$$\nabla_\epsilon (y^T A y) \Big|_{\epsilon=0} = \langle w, A v \rangle + \langle v, A w \rangle = \lambda \langle v, w \rangle$$

$$2 \langle A v, w \rangle = \lambda \langle v, w \rangle \longrightarrow A v = \tilde{\lambda} v$$



The others are obtained by maximizing in the orthogonal complement to the vector  $v$ .

# Eigenvalues and Instability



- Strongly correlated variables result in an orthogonal direction with a small eigenvalue.
- This creates the instability we have been discussing.

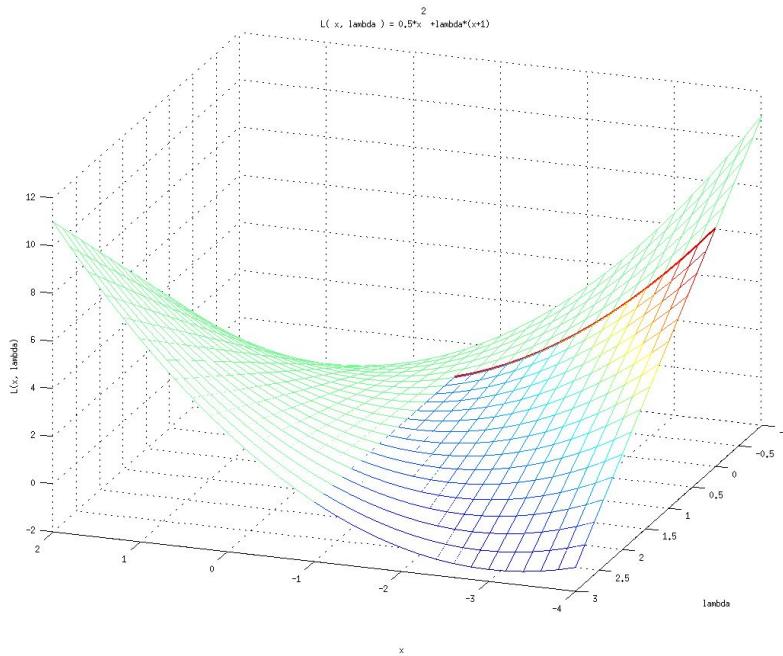
$$Av = \tilde{\lambda}v$$

# Dependent features - what goes wrong

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Why we want to eliminate collinear/dependent features

# Correlation results in instability



- Flat directions in the feature space mean less stability and more uncertainty about what the true coefficients are!
- When we see gradient descent, the parameters involved can significantly affect solutions when there is no stability.

$$\frac{d^2 \mathcal{L}}{d^2 \beta} = \frac{2}{N} X^T X$$

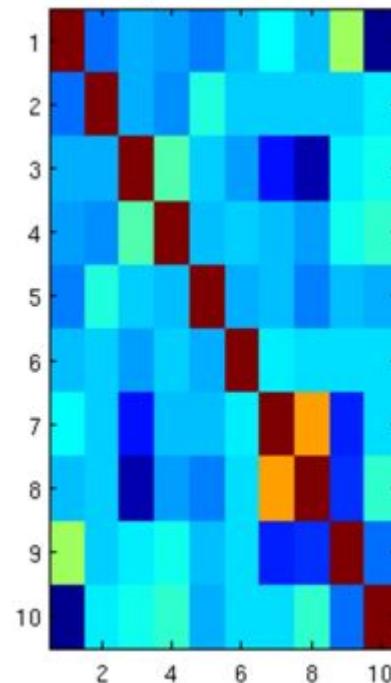
# Correlation results in instability

- $X^T X$  is **symmetric** and therefore has **nonnegative eigenvalues**.
- They are **positive** precisely when the features of  $X$  are **linearly independent**.
- Let's assume for simplicity that  **$X$  is mean centered** (**fine but sometimes reasons why you might not**).

Imagine that  $X$  has two columns which are factors of one another. What can go wrong?

$$y = \alpha x_1 + \beta x_2$$

But our real rule is:  $y = 5x_1$



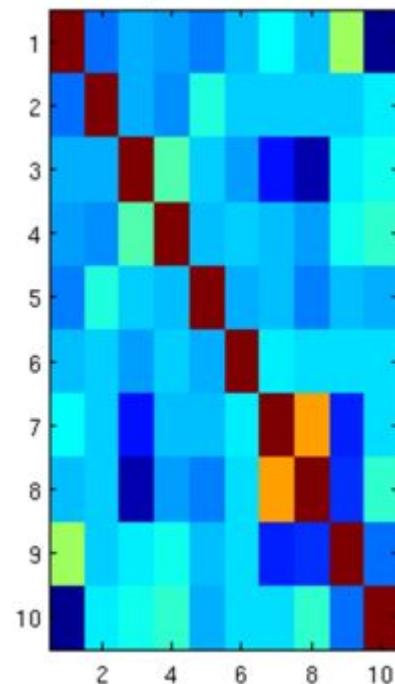
# Correlation results in instability

$$y = \alpha x_1 + \beta x_2$$

But our real rule is:  $y = 5x_1$

Then  $y = -1000x_1 + 10005x_2$  is also a solution

Why is this a problem?



# Correlation results in instability

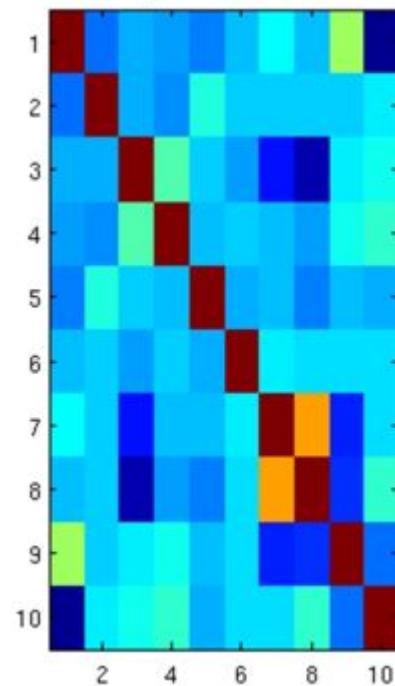
$$y = \alpha x_1 + \beta x_2$$

But our real rule is:  $y = 5x_1$

Then  $y = -1000x_1 + 10005x_2$  is also a solution

## Why is this a problem?

- Creates problems when finding the minimum (next).
- Causes more uncertainty in coefficient estimates (after gradient descent).



# A simple illustration - correlated

$$y = x_1 + \epsilon$$

$$x_2 = 100 * x_1 + \epsilon$$

$$\hat{y} = \beta_1 x_1 + \beta_2 x_2$$

```
n=10000
x1 = np.linspace(0,0.01,n)

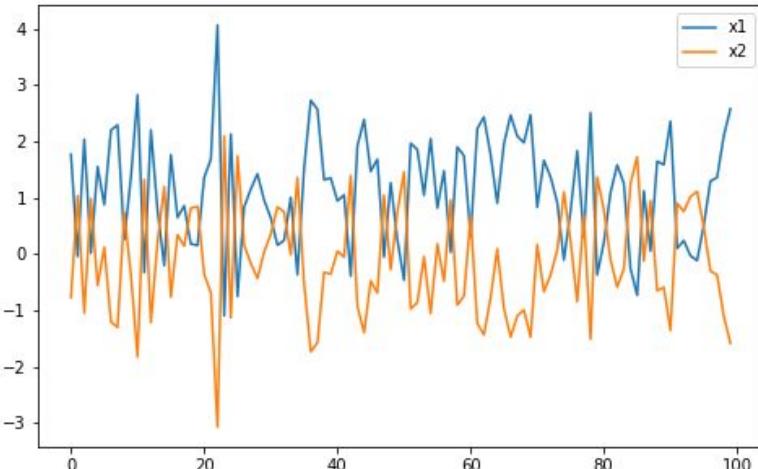
s = np.random.normal(0, 0.001, n)
x2 = 100*x1 + s

df=pd.DataFrame({'x1':x1,'x2':x2})

y = x1 + np.random.normal(0, 0.01, n)
coefs1=[]
coefs2=[]
scores_perp=[]

for i in range(0,100):
    y = x1 + np.random.normal(0, 0.001, n)
    regr = linear_model.LinearRegression()
    xdf
    # Train the model
    regr.fit(x,y)
    coefs1.append(regr.coef_[0])
    coefs2.append(regr.coef_[1]*100)
    scores_perp.append(regr.score(x,y))

plt.figure(figsize=(8,5))
plt.plot(coefs1,label='x1')
plt.plot(coefs2,label='x2')
plt.legend()
```



Number of iterations solving the same problem.

# A simple illustration - orthogonal features

$$y = x_1 + \epsilon$$

$$x_1 \perp x_2$$

$$\hat{y} = \beta_1 x_1 + \beta_2 x_2$$

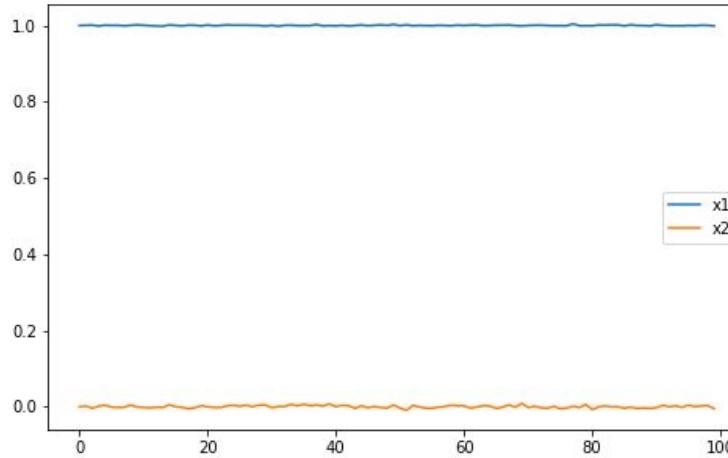
```
k = np.random.normal(0,0.01,n)/np.linalg.norm(k)**2
s = np.random.normal(0, 0.001, n)

x2 = 100*np.linspace(0,0.01,n)
x2 -= x1.dot(k) * k / np.linalg.norm(k)**2
x1=k
df=pd.DataFrame({'x1':x1,'x2':x2})

y = x1 + np.random.normal(0, 0.01, n)
coefs1=[]
coefs2=[]
scores_perp=[]

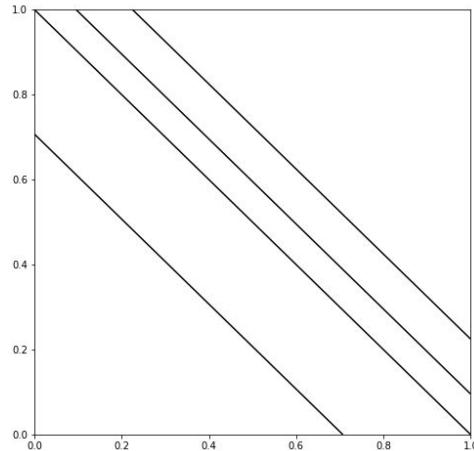
for i in range(0,100):
    y = x1 + np.random.normal(0, 0.001, n)
    regr = linear_model.LinearRegression()
    x=df
    # Train the model
    regr.fit(x,y)
    coefs1.append(regr.coef_[0])
    coefs2.append(regr.coef_[1]*100)
    scores_perp.append(regr.score(x,y))

plt.figure(figsize=(8,5))
plt.plot(coefs1,label='x1')
plt.plot(coefs2, label='x2')
plt.legend()
plt.show()
```

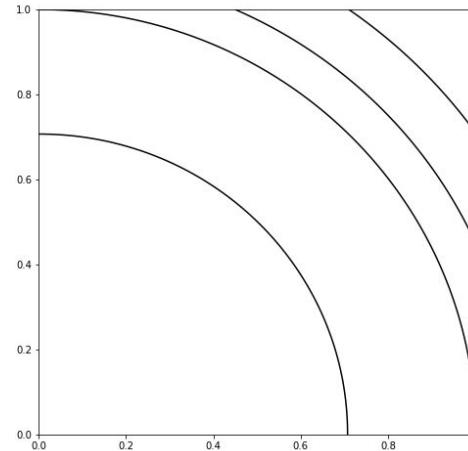


Number of iterations solving the same problem.

# Comparison of the second derivatives



Correlated



Uncorrelated

Notice how in the correlated case, there is clear degeneracy - the solution is not unique. Any value along that line is constant, so there is no unique solution. This is what accounts for the instability in the solution.

# Gradient Descent

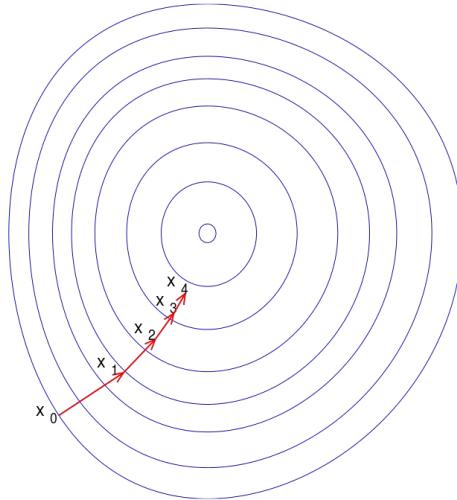
---

How do we minimize a function when there is no analytical solution?

# We can't always solve analytically!

$$\mathcal{L}(\beta) = \frac{1}{N} \sum_{i=1}^N |y_i - \beta \cdot x_i|^2$$

- In these cases, we use the method of gradient descent.
- Almost all other models don't have any explicit, analytical solution, so we have to use gradient descent.



# Continuous Gradient Descent

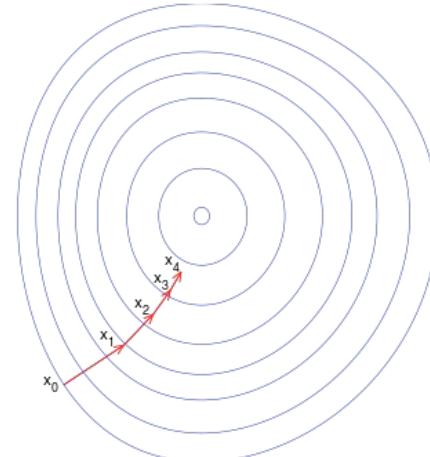
Assume that  $\beta(t)$  solves the equation:

$$\dot{\beta}(t) = -\nabla \mathcal{L}(\beta(t))$$

And that:  $\beta \mapsto \mathcal{L}(\beta)$

- Is strictly convex.
- Twice differentiable.

Then:  $\beta(t)$  converges to the minimum of  $f$  exponentially fast.



# Proof of Convergence of Gradient Descent

Differentiating and using the gradient flow, we have

$$\frac{d}{dt}(\beta(t) - \beta_0)^2 = -2\nabla\mathcal{L}(\beta(t))(\beta(t) - \beta_0)$$

**Next:** How do we use strict convexity to obtain an estimate?

**\*Note\***: In reality we always have a discrete version of the above, and must choose a ‘learning rate’ (ie. time step size) - we will gloss over this for now.

# Proof of Convergence of Gradient Descent

Using Taylor's Law we have

$$\mathcal{L}(\beta) - \mathcal{L}(\beta_0) = \nabla \mathcal{L}(\beta) \cdot (\beta - \beta_0) + \frac{1}{2}(\beta - \beta_0)^T D^2 \mathcal{L}(\xi)(\beta - \beta_0)$$

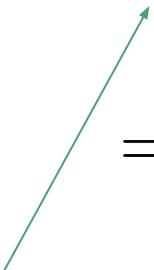
$$\mathcal{L}(\beta_0) - \mathcal{L}(\beta) = \nabla \mathcal{L}(\beta_0) \cdot (\beta_0 - \beta) + \frac{1}{2}(\beta - \beta_0)^T D^2 \mathcal{L}(\tilde{\xi})(\beta - \beta_0)$$

$$\frac{d}{dt}(\beta(t) - \beta_0)^2 = -2\nabla \mathcal{L}(\beta(t))(\beta(t) - \beta_0)$$

$$\frac{d}{dt}(\beta - \beta_0)^2 \leq -m|\beta - \beta_0|^2$$

Therefore:

$$\Rightarrow |\beta(t) - \beta_0| \leq C e^{-mt}$$



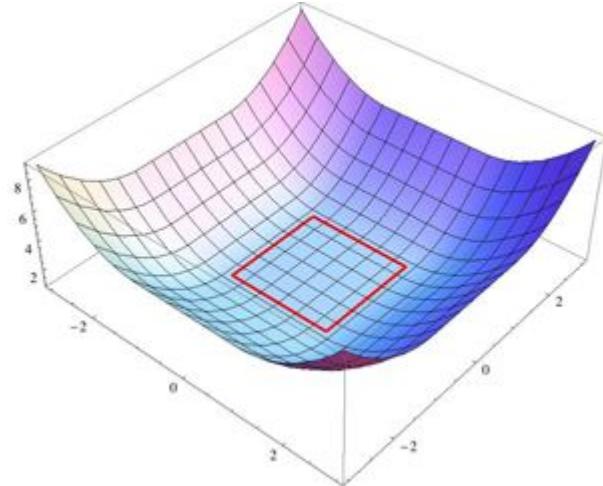
$$-\nabla \mathcal{L}(\beta - \beta_0) = -(\beta - \beta_0)^T (D^2 \mathcal{L}(\xi) + D^2 \mathcal{L}(\tilde{\xi}))(\beta - \beta_0) \leq -M|\beta - \beta_0|^2$$

# Proof of Convergence of Gradient Descent

$$-\nabla \mathcal{L}(\beta - \beta_0) = -(\beta - \beta_0)^T (D^2 \mathcal{L}(\xi) + D^2 \mathcal{L}(\tilde{\xi})) (\beta - \beta_0) \leq -M |\beta - \beta_0|^2$$

$$\Rightarrow |\beta(t) - \beta_0| \leq C e^{-mt}$$

Stability is directly related to how large the eigenvalues are of the correlation matrix!



# How is it computed in python?

$$\frac{\partial \mathcal{L}}{\partial \beta_j} = -\frac{2}{N} \sum_{i=1}^N (y_i - \beta \cdot x_i) x_j$$

$$\beta_n = \beta_{n-1} - \kappa \nabla_{\beta} \mathcal{L}(\beta_{n-1})$$

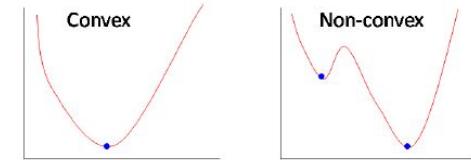
```
def step_gradient(b_current, m_current, points, learningRate):
    b_gradient = 0
    m_gradient = 0
    N = float(len(points))
    for i in range(0, len(points)):
        x = points[i, 0]
        y = points[i, 1]
        b_gradient += -(2/N) * (y - ((m_current * x) + b_current))
        m_gradient += -(2/N) * x * (y - ((m_current * x) + b_current))
    new_b = b_current - (learningRate * b_gradient)
    new_m = m_current - (learningRate * m_gradient)
    return [new_b, new_m]
```

$\kappa$  is known as the learning rate - when computing we must choose a time step size - Homework 1 covers this.

*But if the learning rate is too big then, gradient descent may overshoot the minimum and oscillate back and forth. Why?*

# The importance of gradient descent.

- All **parametric problems** in predictive machine learning seek to **minimize the distance of some a priori function of the data to the observed values.**
- It only makes sense to try to **minimize functions** which are **convex** (at least locally). Otherwise it's common to get stuck in local minima.
- When there is no analytical solution, the solution must be obtained by following the **steepest descent from a starting point to the minimum of the function.**
- This will be true when we work with more advanced probabilistic methods as well, and more important to understand.
- Stability is directly related to the eigenvalues of the covariance matrix, and also depends on how we choose our learning rate.



# Lecture Outline

- Review Homework Solutions and relate it back to the theory we covered. Github 101 from Chris!
- Coefficient Estimate Confidence (Lecture 2 - Intro to Regression Continued)
- Lecture 3 Notes - Complexity, Regularization and Over Fitting.
  - Homework 2 - Applying these tools to the TV/Radio/Newspaper Sales data.

# Linear Regression II: Estimators and Confidence

---

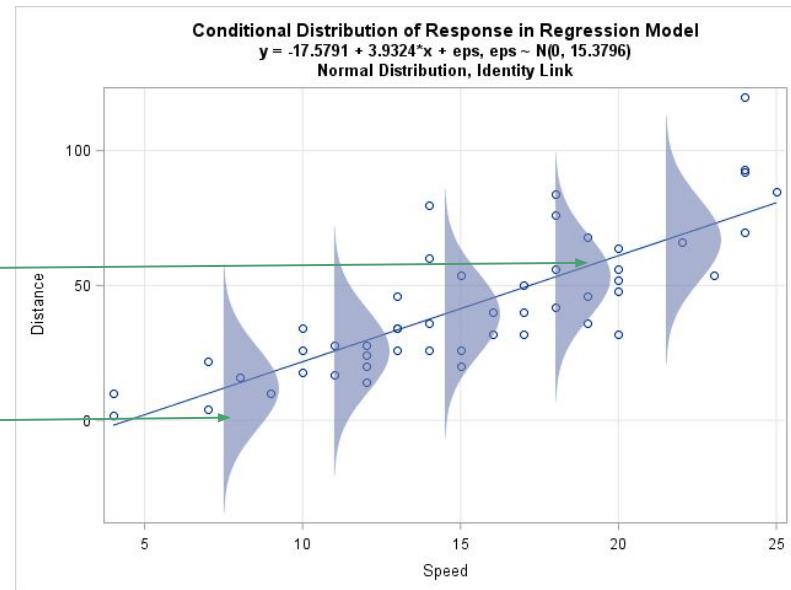
# Understanding uncertainty

$$y = \hat{y} + \epsilon$$

$\hat{y}$  Estimator

$\epsilon$  Error

In machine learning, our goal is to find the best **estimator** in the presence of natural uncertainty (**error**).



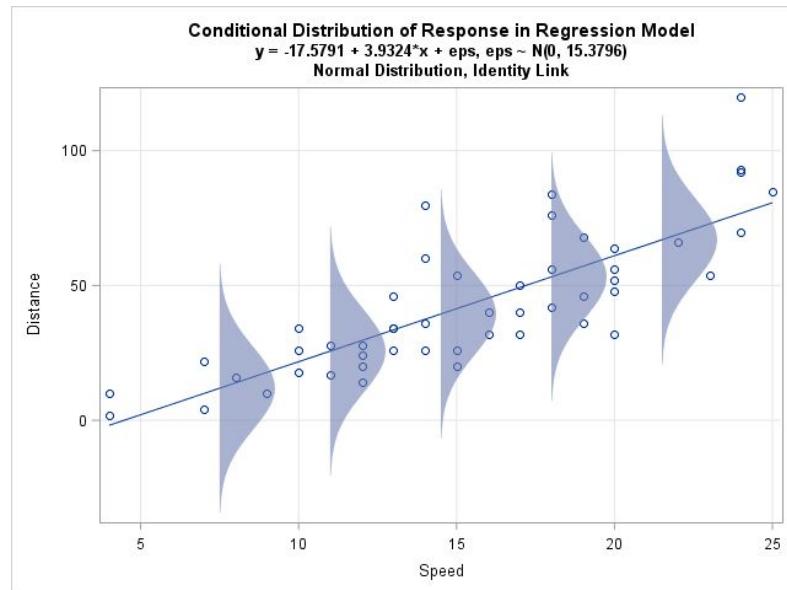
# What is the error?

$$y = \hat{y} + \epsilon$$

$\epsilon$  Is assumed to be from some distribution, ie.

$$\epsilon_i \sim \mathcal{N}(0, \sigma^2)$$

**For example:** you could never predict the height of someone from their age, gender, ethnicity or any other collection of variables alone - there is a natural variance in the heights of individuals.



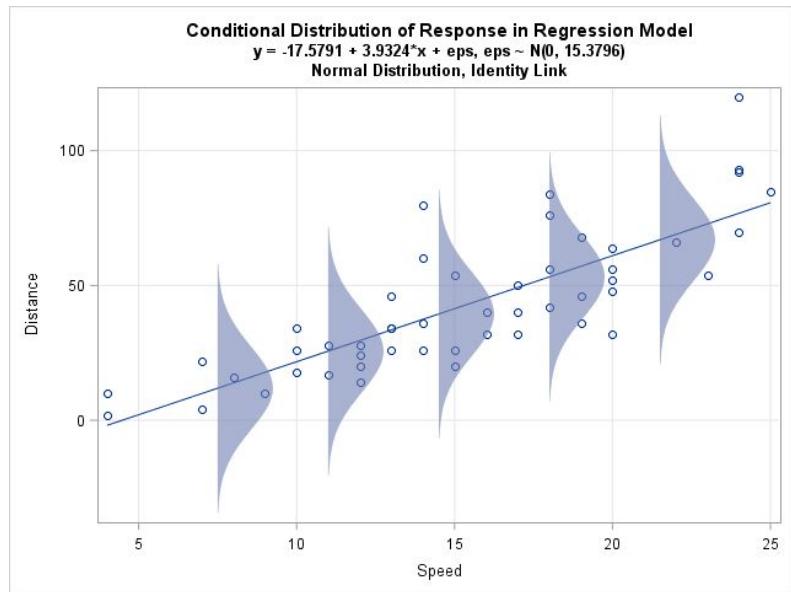
# What is the error?

$$y = (\hat{\beta} + \epsilon_\beta) \cdot x + \epsilon$$

$\epsilon$  Is assumed to be from some distribution, ie.

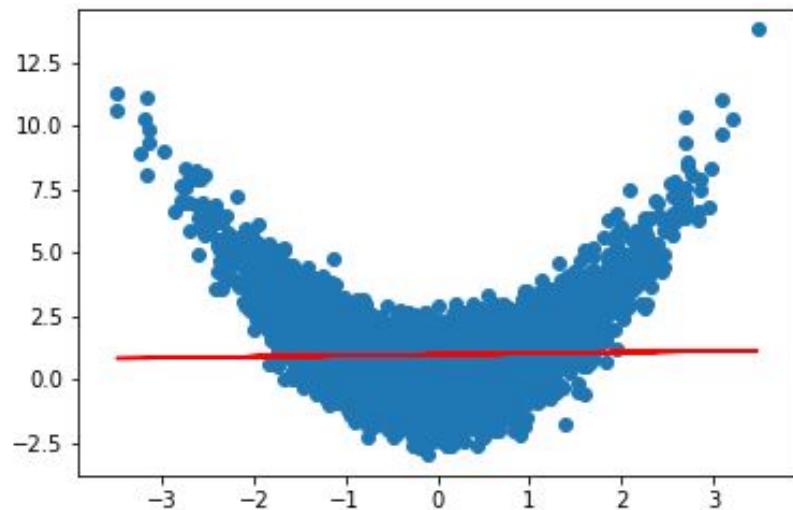
$$\epsilon_i \sim \mathcal{N}(0, \sigma^2)$$

$$\mathbb{E}(y|x) = \beta \cdot x$$



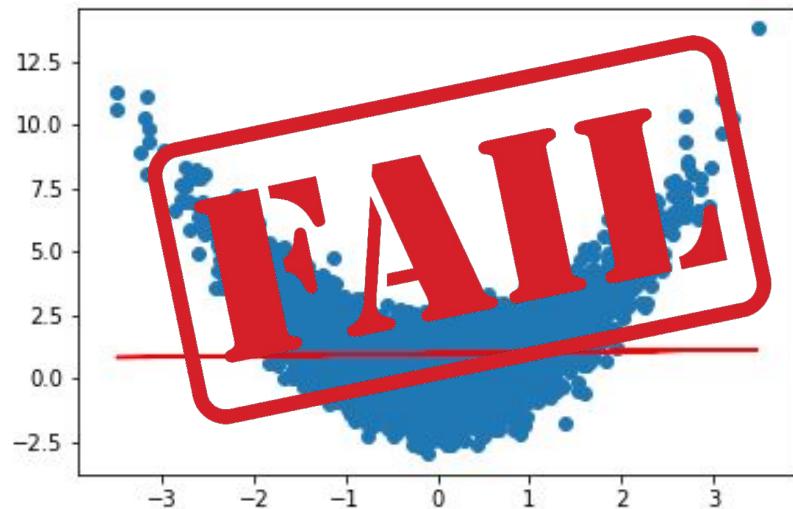
# What assumptions does this imply?

Assumption 1 - Linear relationship between dependent and independent variables.



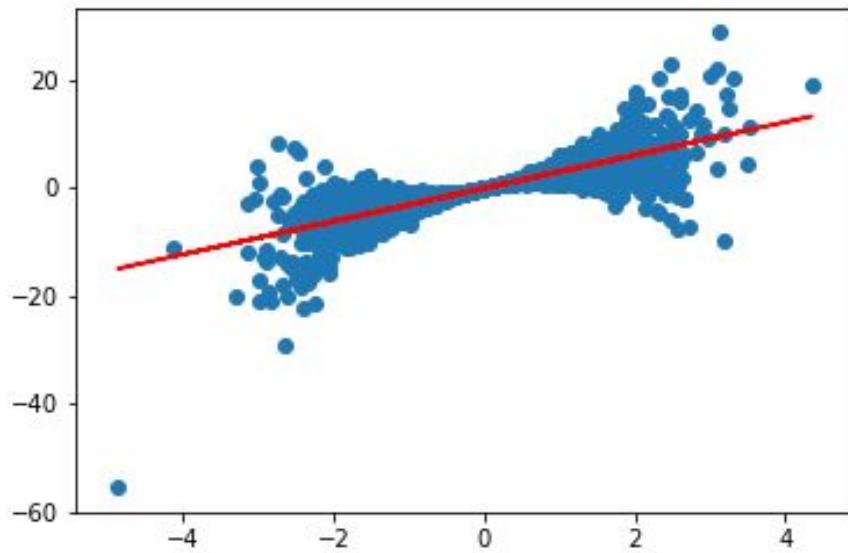
# What assumptions does this imply?

Assumption 1 - Linear relationship between dependent and independent variables.



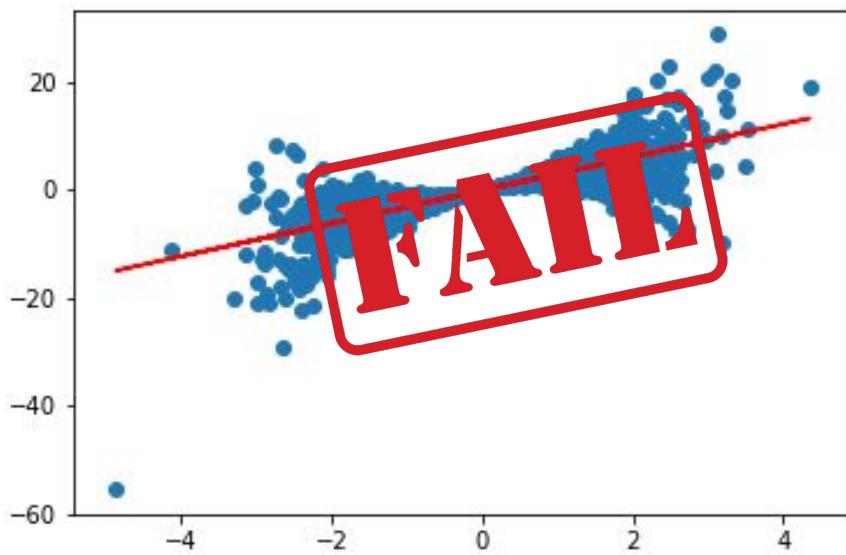
# What assumptions does this imply?

Assumption 2 - Heteroscedasticity: constant variance for any value of  $\mathbb{E}(y|x) = \beta \cdot x$



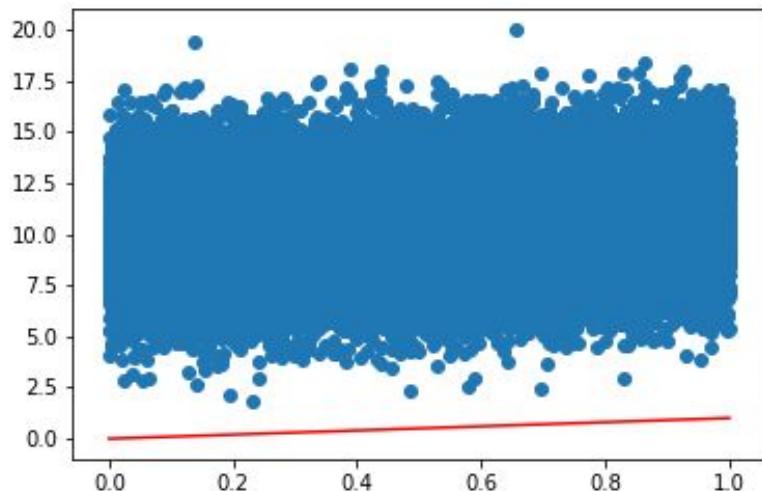
# What assumptions does this imply?

Assumption 2 - Heteroscedasticity: constant variance for any value of  $\mathbb{E}(y|x) = \beta \cdot x$



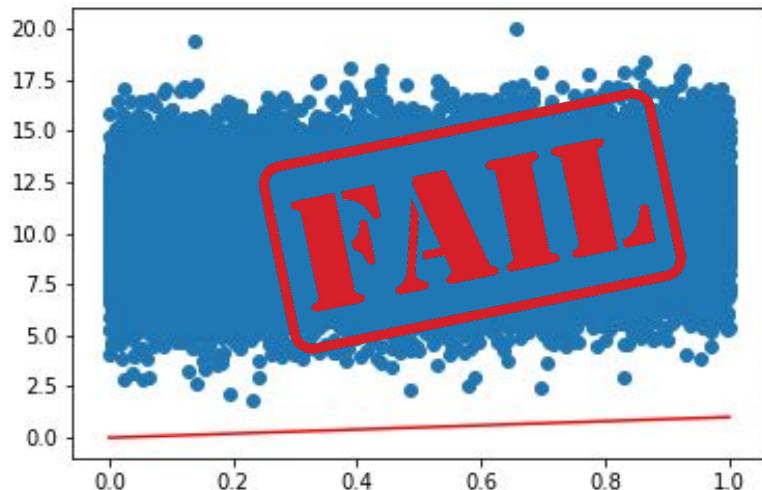
# What assumptions does this imply?

Assumption 3 - The residuals are all normally distributed with zero mean.



# What assumptions does this imply?

Assumption 3 - The residuals are all normally distributed with zero mean.



# From Last Time

## Things we covered

- Linear Regression: Exact solution, stability and gradient descent methods.
- Normal Residuals - how the error around the predicted mean is normally distributed.
- Feature Importance - Why normalizing is essential.

## Today

- Review above material and some linear algebra.
- Go through a concrete example - predicting Booking.com sellouts.
- Do an in class session working through some problems and applying what we've covered so far.
- Please collaborate! But don't simply copy.

# Variable Importance and Confidence

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# Advertisement Example

What are the features?

**TV:** advertising dollars spent on TV for a single product in a given market (in thousands of dollars)

**Radio:** advertising dollars spent on Radio

**Newspaper:** advertising dollars spent on Newspaper

**Goal:** Predict the number of sales in a given market based on the advertising in TV, Radio and Newspaper.

```
In [263]: # read data into a DataFrame
import pandas as pd
import pylab as plt
import seaborn
from sklearn.linear_model import LinearRegression
import numpy as np
import random
import json
pd.set_option('display.max_columns', 500)
%matplotlib inline

df = pd.read_csv('http://www-bcf.usc.edu/~gareth/ISL/Advertising.csv', index_col=0)
df.head()
```

	TV	Radio	Newspaper	Sales
1	230.1	37.8	69.2	22.1
2	44.5	39.3	45.1	10.4
3	17.2	45.9	69.3	9.3
4	151.5	41.3	58.5	18.5
5	180.8	10.8	58.4	12.9

# Advertisement Example

```
In [263]: # read data into a DataFrame
import pandas as pd
import pylab as plt
import seaborn
from sklearn.linear_model import LinearRegression
import numpy.random as nprnd
import random
import json
pd.set_option('display.max_columns', 500)
%matplotlib inline

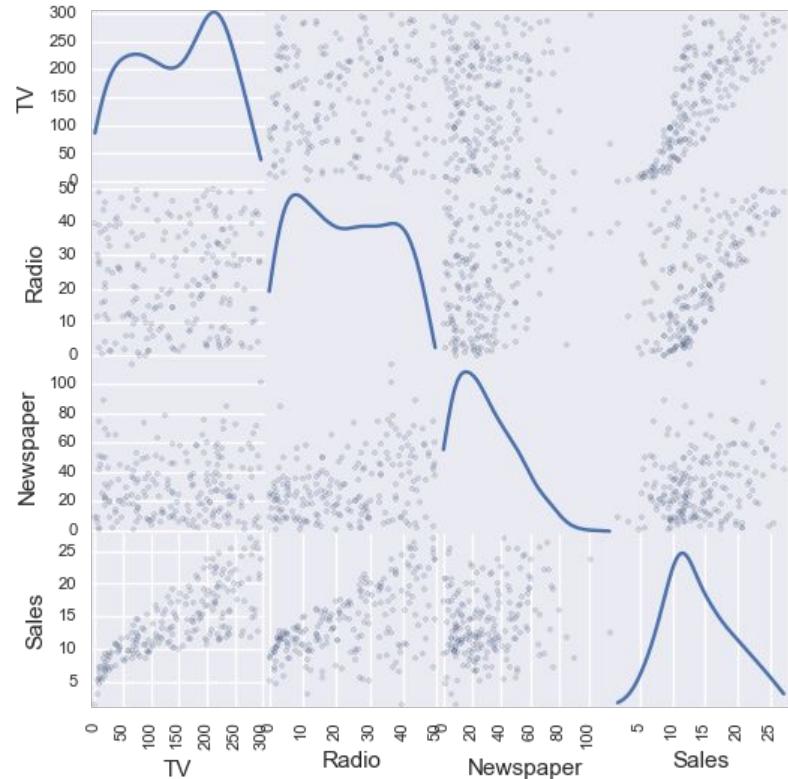
df = pd.read_csv('http://www-bcf.usc.edu/~gareth/ISL/Advertising.csv', index_col=0)
df.head()
```

Out[263]:

	TV	Radio	Newspaper	Sales
1	230.1	37.8	69.2	22.1
2	44.5	39.3	45.1	10.4
3	17.2	45.9	69.3	9.3
4	151.5	41.3	58.5	18.5
5	180.8	10.8	58.4	12.9

# Inference vs Prediction

- Sometimes people care more about inference - which is understanding how variables affect the outcome variable.
- **Example: Does advertising on TV, Radio or Newspaper positively affect sales?**



\*Homework 1\*

# Feature importance & Inference

$$y_i = \sum_{j=1}^N \beta_j X_{ij}$$

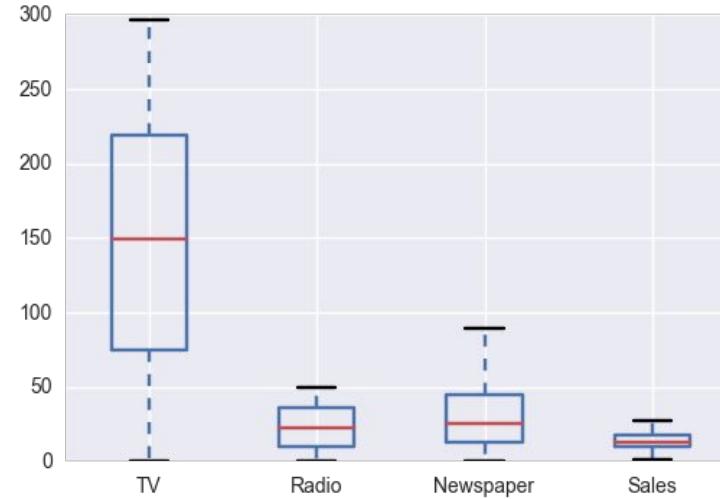
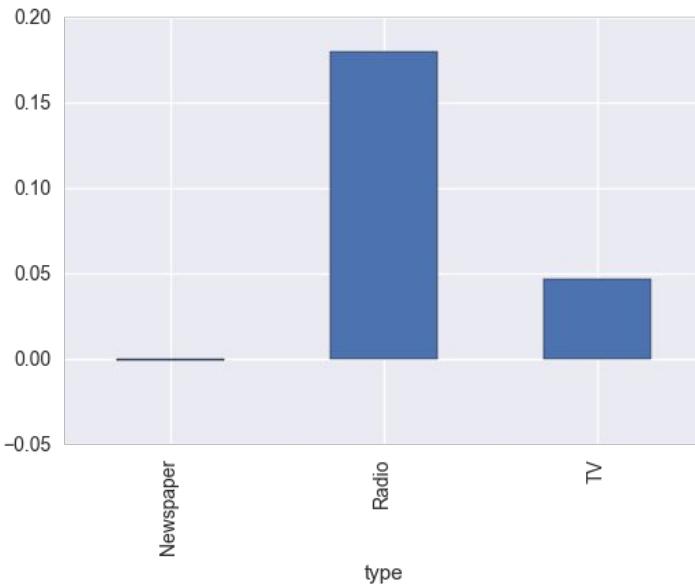
When there is no relationship between the features, we can conclude that

$$\frac{\partial y_i}{\partial X_{ik}} = \beta_k$$

- This allows us to interpret the significance of each variable, in terms of how much it changes the response variable.
- **Does not work if there is any interaction or dependence between the features however.**
- To **compare coefficients**, we need to **normalize the columns to have the same scale**.

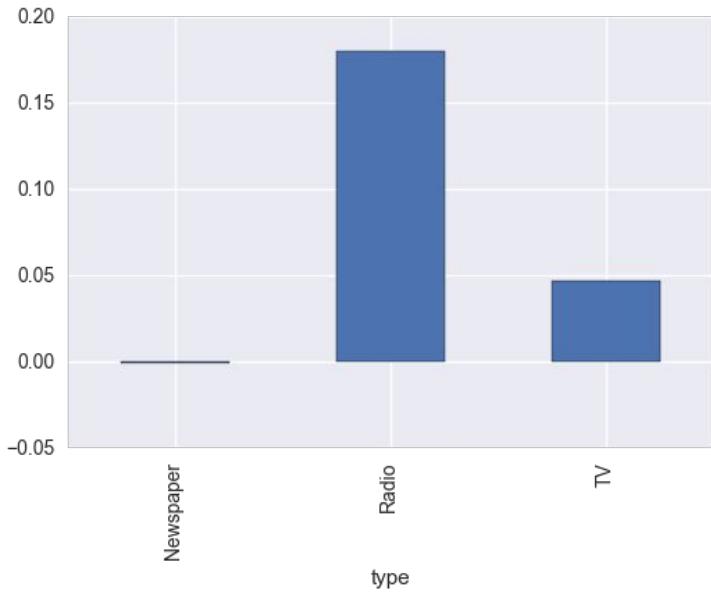
- But how confident are we?

# What's wrong with this picture?

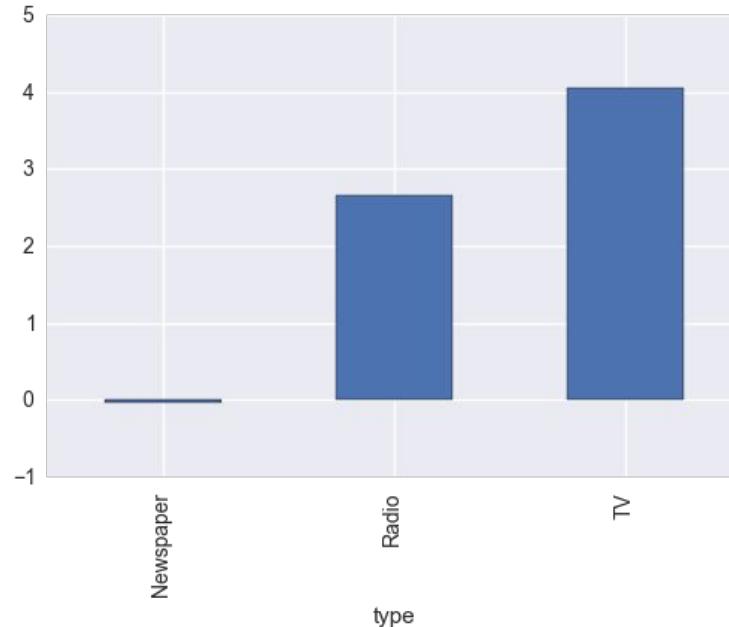


**Conclusion:** Radio is the most significant variable. ?

# After standardizing



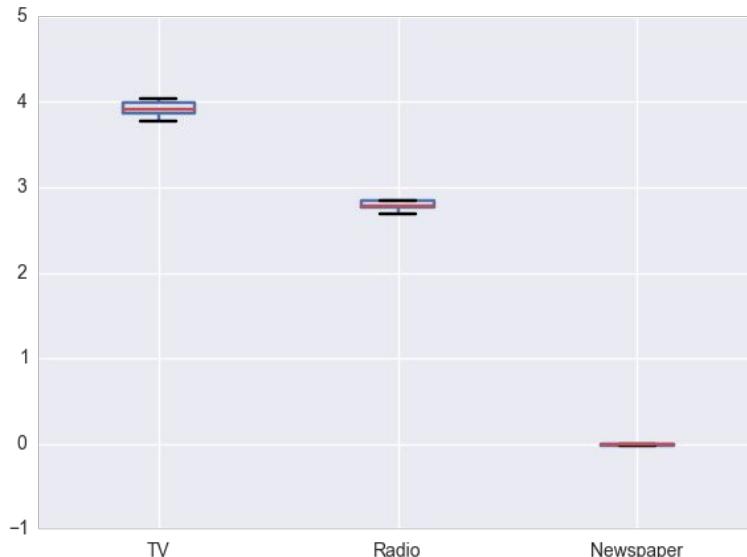
Non-Standardized



Standardized

- Each feature is normalized to have **mean zero** and **unit variance**

# Confidence: via Cross Validation



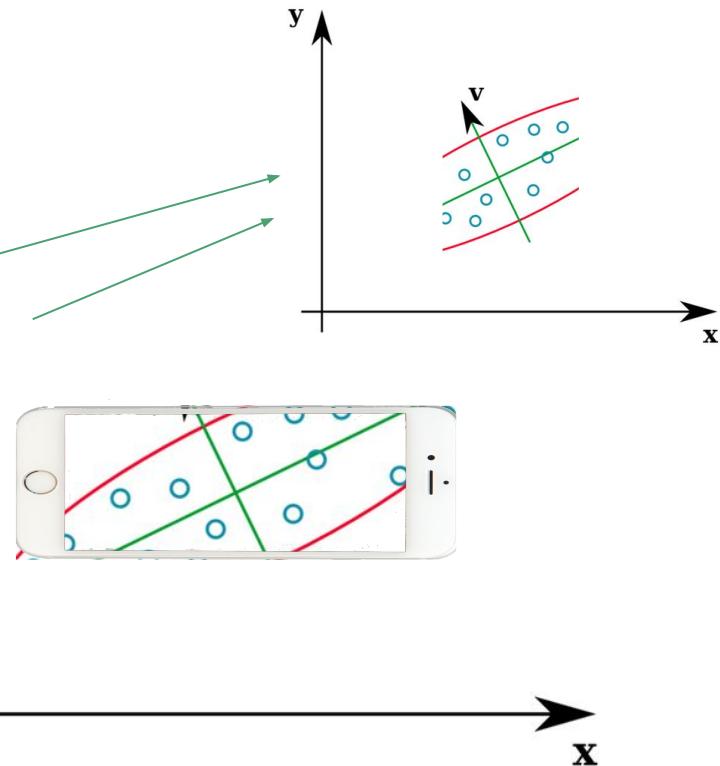
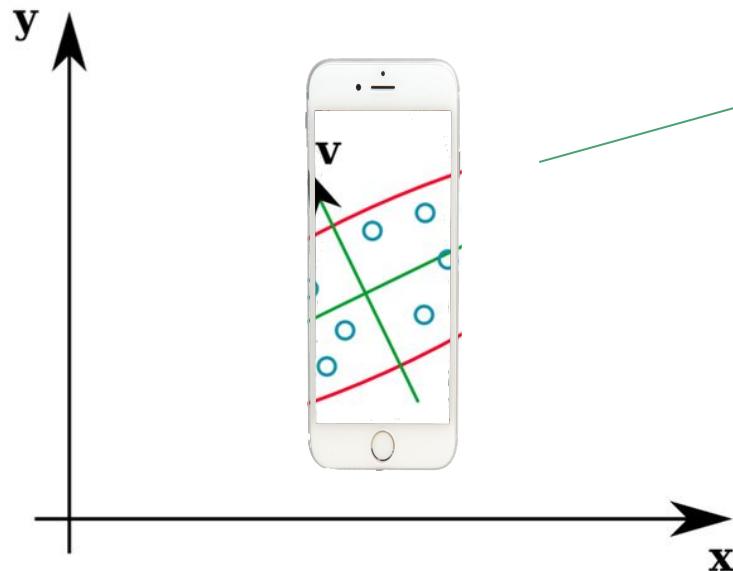
- Average your model **over many random folds, and standardize.**
- Look at the variance in the coefficients you obtain over many folds.

Average over 5-fold cross validation.

# Great questions from last time

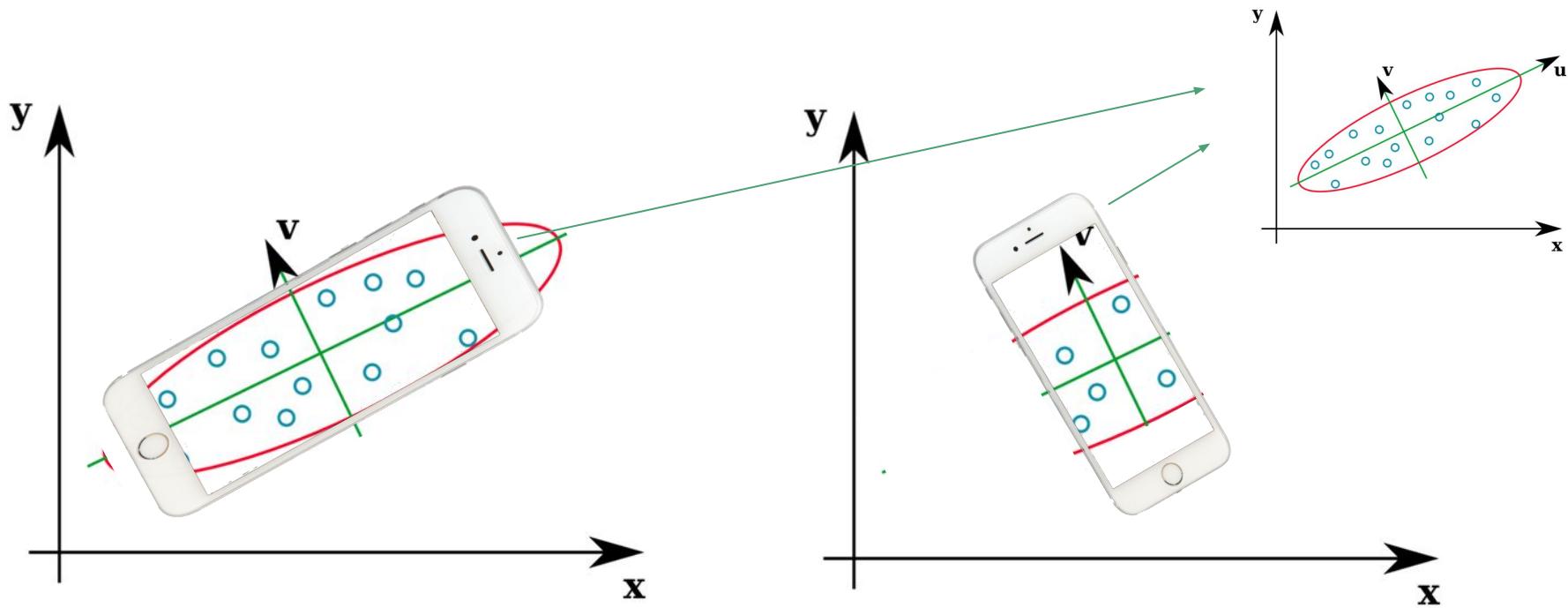
1. PCA - why are eigenvectors better than our original axes? What are they?
  
2. If we do K-fold cross validation, which coefficient do we choose? The mean, max? (**Answer: None of them! That's not the point.**)
  
3. Does standardizing data result in better or worse performance?  
**(No: But not doing it prevents you from doing regularization, so it can in that way)**
  
4. When to use **standardization vs normalization?** (these words are used in different ways by different people in the literature)

# Normal Coordinates



Imagine you can only take a photo of these points in two directions - which ones would you choose?

# Eigenvectors



By rotating our camera to “good” angles, we capture more about the distribution of the data! These are the eigenvectors (ie. PCA).

# How to make predictions with K-Fold?

1. If we do K-fold cross validation, which coefficient do we choose? The mean, max? (Answer: None of them! That's not the point).

The **purpose k-fold cross validation is to measure the confidence of your model “procedure”**- ie. feature selection, model selection (linear regression, random forest, etc), regularization (hyperparameter selection).

It can also give you a sense of how wildly parameters change and thus a sense of '**confidence**' about your variables, with regards to generalizing to unseen data.

But ultimately you **verify your model with k-fold cross validation**, then **train on the entire data set** once you're ensured it generalizes to new examples!

More than one model? Do your best on the K folds with each model, and compare the total error over each fold:

**More explanations:**

<https://stats.stackexchange.com/questions/52274/how-to-choose-a-predictive-model-after-k-fold-cross-validation>

# Should you always normalize your data?

- It depends a lot on your data and your model, but generally for linear models, **normalizing is not only safer, it is usually needed**
- When we penalize the size of error terms (regularization), we will need to normalize for this to make any sense.
- For tree-based models, it doesn't matter (we will see why later).
- **Long story short:** if your model is linear, you probably want to normalize unless you have a reason not to. Ie. is a 1cm change in height the same as a 1kg change in weight?

# Types of standardization/normalization

## Normalizing to mean zero and unit variance

- **Regression/Classification**
- **Clustering**, standardization may be especially crucial in order to compare similarities between features based on certain distance measures.
- **Principal Component Analysis**, where we usually prefer standardization over Min-Max scaling.
- Coefficient comparison when variables represent different categories.
- Helps with gradient descent convergence - why?

## Min-Max Scaling

- Image processing, where pixel intensities have to be normalized to fit within a certain range (i.e., 0 to 255 for the RGB color range).
- I've honestly never seen real world examples other than this where this is done.

**Conclusion: Always use standardization (mean zero and unit variance) unless you believe there is a strong reason not to.**

# Standardization or Normalization?

There are some definitions (which I may have mixed up in lecture potentially):

**Normalization** transforms your data into a range between 0 and 1.

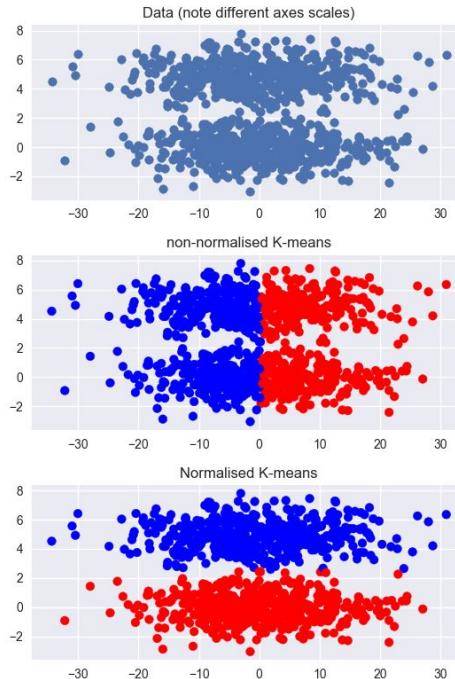
**Standardization** transforms your data such that the resulting distribution has a mean of 0 and a standard deviation of 1.

**It's quite rare for people to use min/max normalization in regression problems in practice. In general people want to standardize so that the mean is 0 and the variance is 1. So you would have:**

In general you care less about having your data all be within the same range, **but instead want measurements of change to be the same across all variables**. This is important when comparing coefficients, and when we will cover regularization, which puts bounds on the coefficients. There are indeed some exceptions though, such as when your data doesn't vary much but you want it to have the same range.

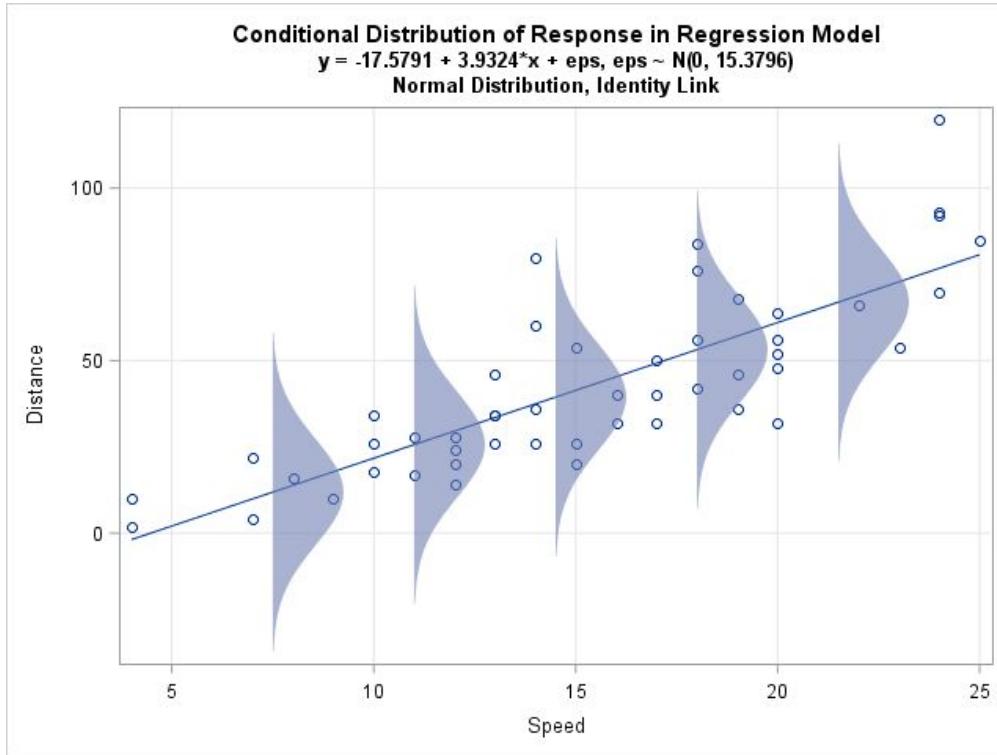
**Taking max or min also isn't really stable for outliers ( can you see why? Just one large point will mess things up).**

# Clustering Example



- Here we see that the plot above has a different range than the data below.
- When we normalize our data, we don't get the natural split we want.
- Although this is classification, the idea generalizes as well.

# Errors are normally distributed



- $$y_i = \beta \cdot x_i + \epsilon_i$$
- $$\epsilon_i \sim \mathcal{N}(0, \sigma^2)$$
- We assume the errors for ordinary least squares are assumed to be normally distributed (we will see why later).
  - This assumption will be justified by the central limit theorem.

# Properties of summing random variables

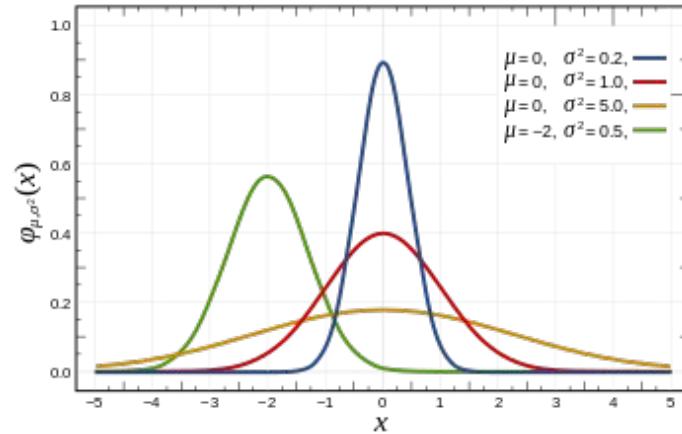
$$X \sim \mathcal{N}(\mu_X, \sigma_X^2)$$

$$Y \sim \mathcal{N}(\mu_Y, \sigma_Y^2)$$

$$Z = X + Y$$

If X and Y are **independent**,  
then:

$$Z \sim \mathcal{N}(\mu_X + \mu_Y, \sigma_X^2 + \sigma_Y^2)$$



## Intuition of Reason

- The number of heads when flipping a coin many times is (approximately) normally distributed.
- If you flip **two coins** N times, it's the same as flipping one coin with the average mean of the two coins 2N times (ie. another coin!), which is also normally distributed.

# Distribution of slope estimator

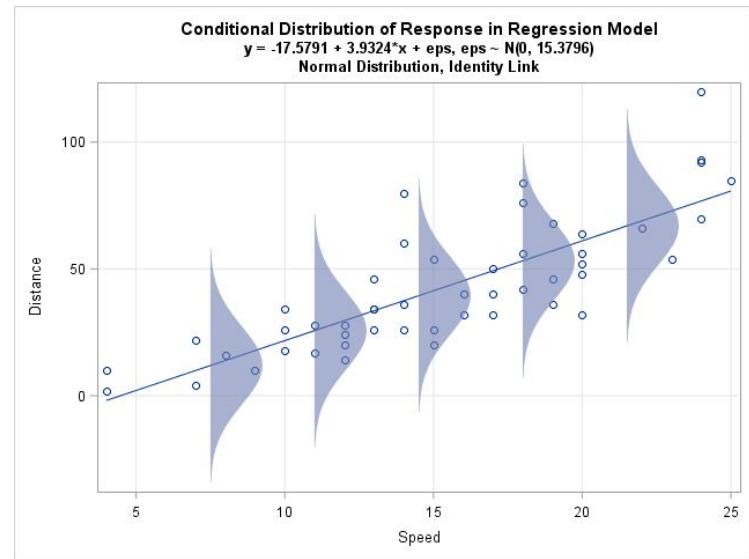
$$y_i = \beta \cdot x_i + \epsilon_i$$

$$\sum_i \beta \cdot x_i^2 \sim \mathcal{N} \left( \sum_i y_i \cdot x_i, \sum_i x_i^2 \right)$$

$$\beta \sim \mathcal{N} \left( \frac{\sum_i y_i \cdot x_i}{\sum_i x_i^2}, (\sum_i x_i^2)^{-1} \right)$$

$$\beta \sim \mathcal{N} \left( \hat{\beta}, (\sum_i x_i^2)^{-1} \right)$$

This variance gives us a confidence bound on our estimate of the coefficient.



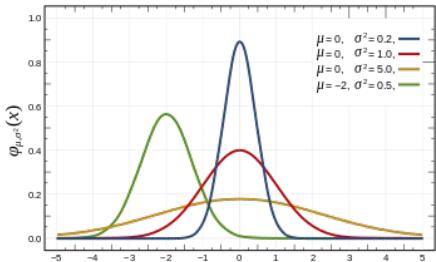
# Confidence of coefficients

$$y_i = \beta \cdot x_i + \epsilon_i$$

$$\hat{\beta}_1 = \frac{\sum_i x_i y_i}{\sum x_i^2}$$

$$= \frac{\sum x_i (\beta_1 x_i + \epsilon)}{\sum x_i^2}$$

$$= \beta_1 + \frac{\sum \epsilon_i x_i}{\sum x_i^2}$$



$$f(x | \mu, \sigma^2) = \frac{1}{\sqrt{2\sigma^2\pi}} e^{-\frac{(x-\mu)^2}{2\sigma^2}}$$

$\epsilon_i$  is normally distributed and independent.

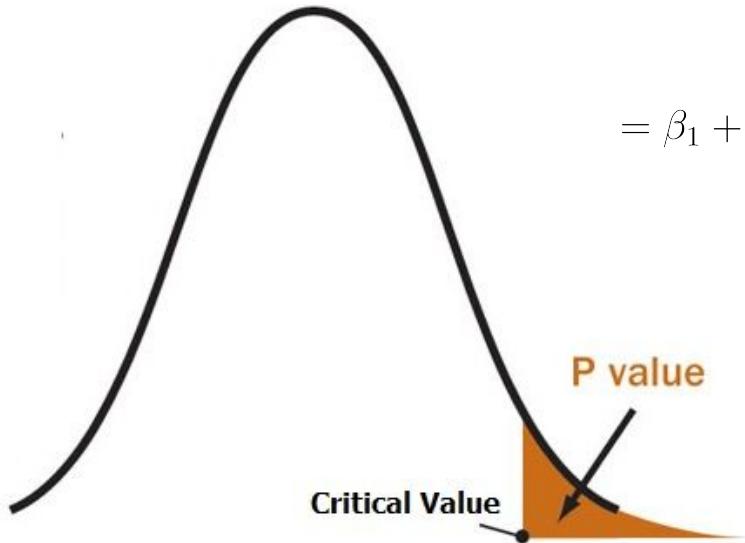
$$\epsilon_i \sim \mathcal{N}(0, \sigma^2)$$

$$\frac{\sum_i \epsilon_i x_i}{\sum x_i^2} \sim \mathcal{N}\left(0, \frac{\sigma^2}{\sum x_i^2}\right)$$

$$\rightarrow \mathcal{N}(0, (X^T X)^{-1} \sigma^2)$$

This is how you measure p values of regression coefficients.

# Confidence via p values



$$= \beta_1 + \frac{\sum \epsilon_i x_i}{\sum_i x_i^2}$$

$$\frac{\sum_i \epsilon_i x_i}{\sum x_i^2} \sim \mathcal{N}(0, (X^T X)^{-1} \sigma^2)$$

$$t = \frac{\beta_1 - 0}{\text{SE}(\beta_1)}$$

Number of standard deviations away from 0 - how rare is it to observe this value under the hypothesis H0?

$$\text{SE}(\beta_1)^2 \sim \frac{1}{N} \frac{\sum_i (y_i - \hat{y}_i)^2}{\sum_i (x_i - \bar{x})^2}$$

**H0:** The coefficient is zero and has no effect on y.

**H1:** The coefficient is non-zero and has an impact on y.

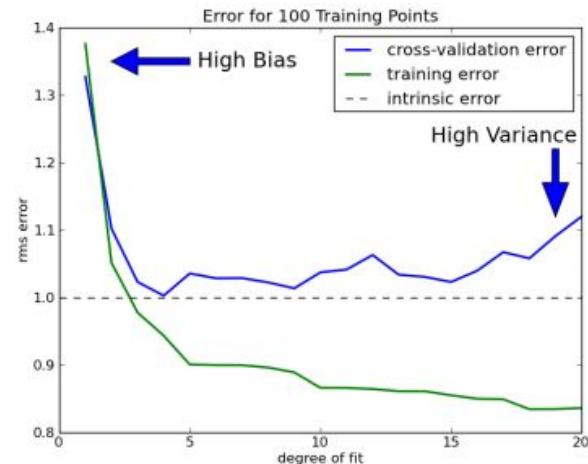
This is an example of **Hypothesis Testing**

# Eliminating bad variables?

$$\beta = (X^T X)^{-1} X^T y$$

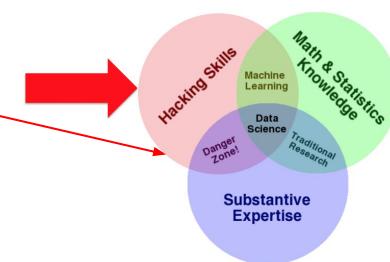
- 1) For building predictive models, the best strategy is to **optimize for performance by restricting the size of the coefficients, or dimensionality reduction.**
- 2) **If you don't have a lot of data, use many fold cross validation** to avoid the chance of having a spurious model.

\*Both will be covered in detail **next lecture.** \*



# Quick Summary

- Derived the analytical solution for L2 linear regression.
- Related the stability of the solution and confidence of the estimate of the coefficients to the dependence of features on one another (covariance/correlation). **P values or cross validation help to understand the confidence in our coefficients.**
- Under all of these assumptions, we can make sense of ‘feature importance’ - but we have ~~to be~~ very careful!



# Evaluating on testing data

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We have a model - does it generalize?

# How to evaluate on testing data?

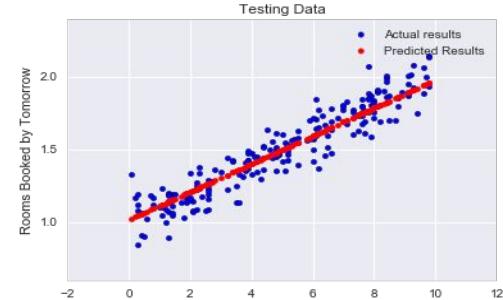
For regression problems, we generally use two possible metrics:

- **Root mean squared error:**

$$\sqrt{\frac{1}{N} \sum_{i=1}^N (y_i - f(x_i))^2}$$

- **R-squared:**

$$1 - \frac{\sum_{i=1}^N (f(x_i) - y_i)^2}{\sum_{i=1}^N (y_i - \bar{y})^2} \quad \longleftarrow$$

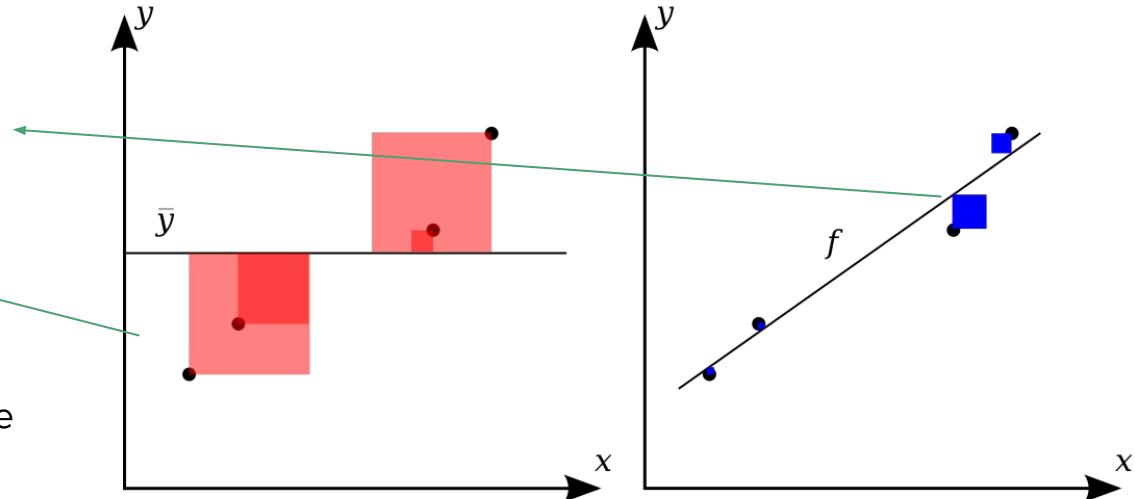


- This is the fraction of variance that our model is able to explain.

# Coefficient of Determination ( $R^2$ )

$$1 - \frac{\sum_{i=1}^N (f(x_i) - y_i)^2}{\sum_{i=1}^N (y_i - \bar{y})^2}$$

- This is the percentage of variance that our model is able to explain
- A perfect model would have the second term be 0, resulting in 1.
- If the model was simply the mean, the result would be 0.



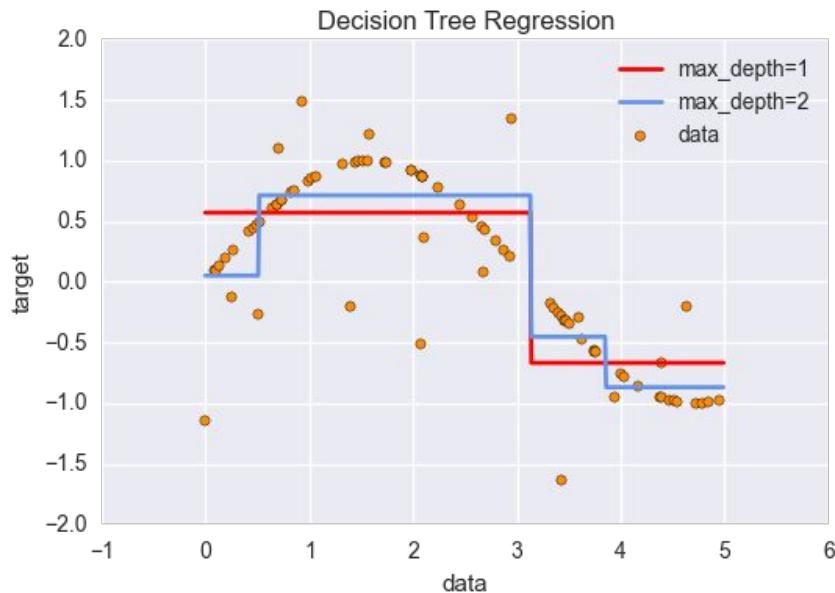
# What have we shown?

- Given any norm, by iterating under its gradient flow, we can obtain a solution to the minimization problem (it's unique iff the norm is strictly convex).
- Most minimization problems don't have analytical solutions like the L2 norm - **this is how scikit-learn codes these algorithms (with GD).**
- We will now see a live demo of this in Python.
- [https://github.com/Columbia-Intro-Data-Science/APMAE4990/blob/master/not\\_ebooks/Lecture1%20-%20Introduction-to-Regression.ipynb](https://github.com/Columbia-Intro-Data-Science/APMAE4990/blob/master/not_ebooks/Lecture1%20-%20Introduction-to-Regression.ipynb)
- You will be asked to apply these methods that we work through now in the homework assignment:  
<https://github.com/Columbia-Intro-Data-Science/APMAE4990/blob/master/homework/Homework%201%20-%20Github%20and%20Pandas.ipynb>

# Decision Tree Regression

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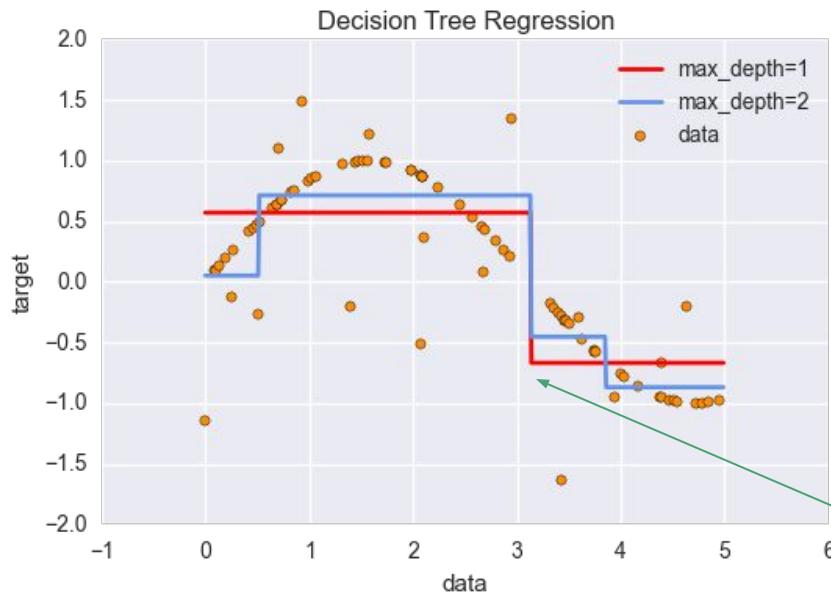
# First: A 1d example



$$y = \sin(x) + \text{noise}$$

- Decision trees make a recursive series of decisions which lead to an outcome, real valued or labeled (for regression, real valued)
- The goal is to make splitting decisions on the data to minimize a norm, in particular, the L2 distance to the mean on that subset of the data, also known as the **variance**.

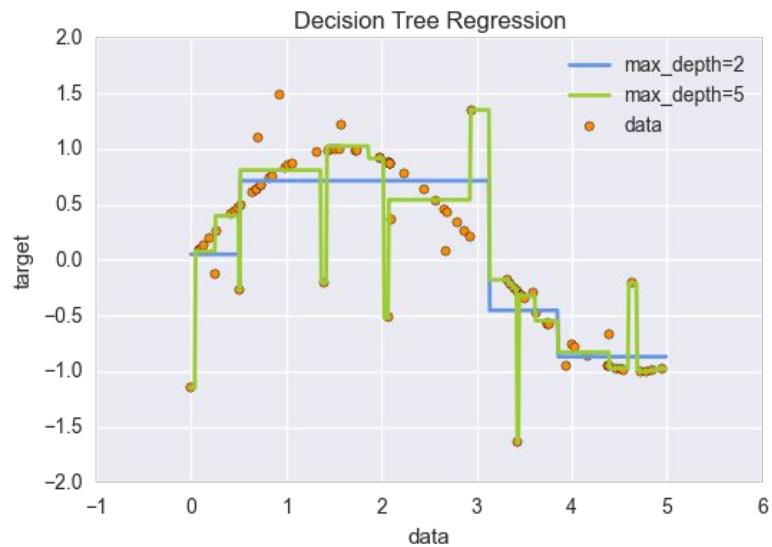
# First: A 1d example



$$y = \sin(x) + \text{noise}$$

- For a depth of **zero**, one chooses the **mean**.
- How does one choose the splitting point when the depth is larger than zero?
- For a depth of one, the algorithm searches through all values between (-1,6) (in this example), and chooses the split which **minimizes the variance on the two segments which it creates**.

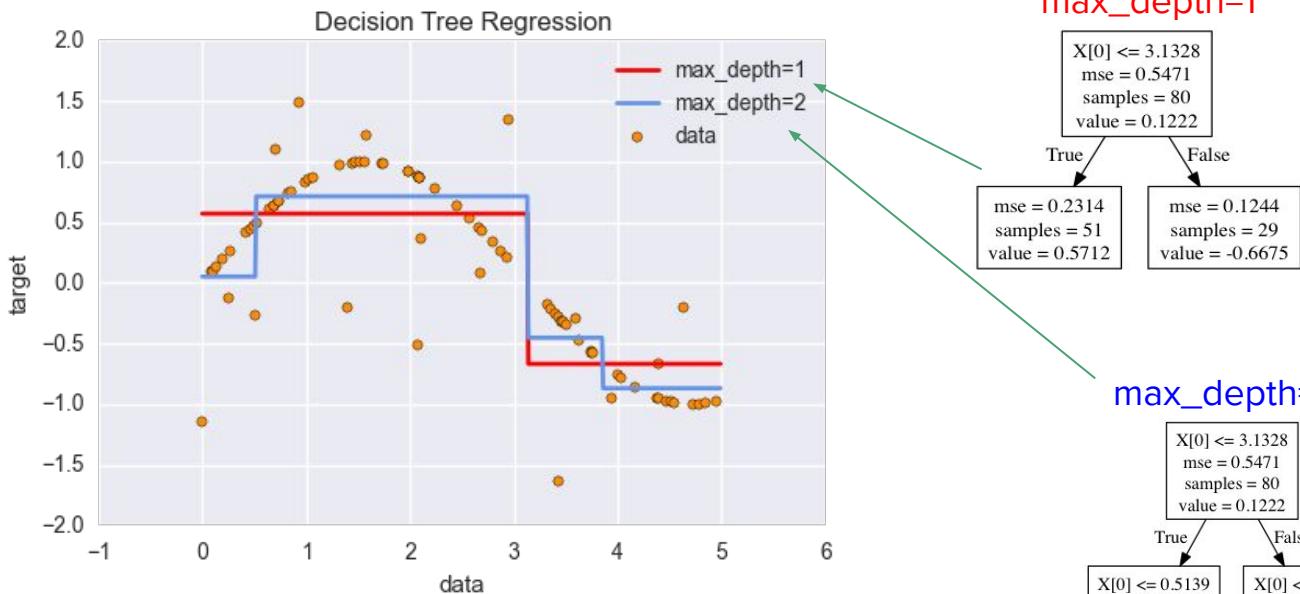
# Increased depth can be bad



$$y = \sin(x) + \text{noise}$$

- In general, if one adds more depth to the tree, there is a risk of overfitting. Look at the unwanted variance we have picked up in this example.
- We will learn next lecture how to choose the perfect depth number!

# How do the decisions work?



max\_depth=1

X[0] <= 3.1328  
mse = 0.5471  
samples = 80  
value = 0.1222

True  
mse = 0.2314  
samples = 51  
value = 0.5712

False  
mse = 0.1244  
samples = 29  
value = -0.6675

max\_depth=2

X[0] <= 3.1328  
mse = 0.5471  
samples = 80  
value = 0.1222

True  
X[0] <= 0.5139  
mse = 0.2314  
samples = 51  
value = 0.5712

False  
X[0] <= 3.8502  
mse = 0.1244  
samples = 29  
value = -0.6675

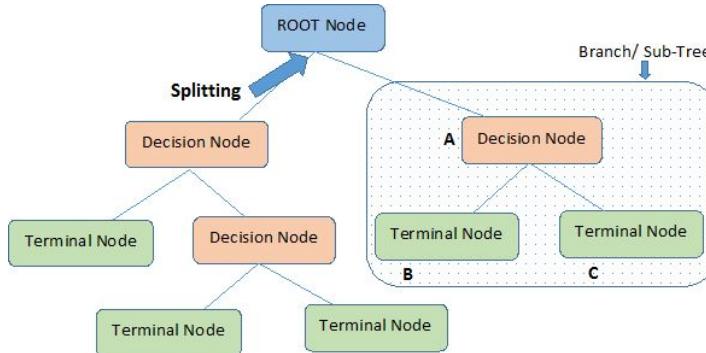
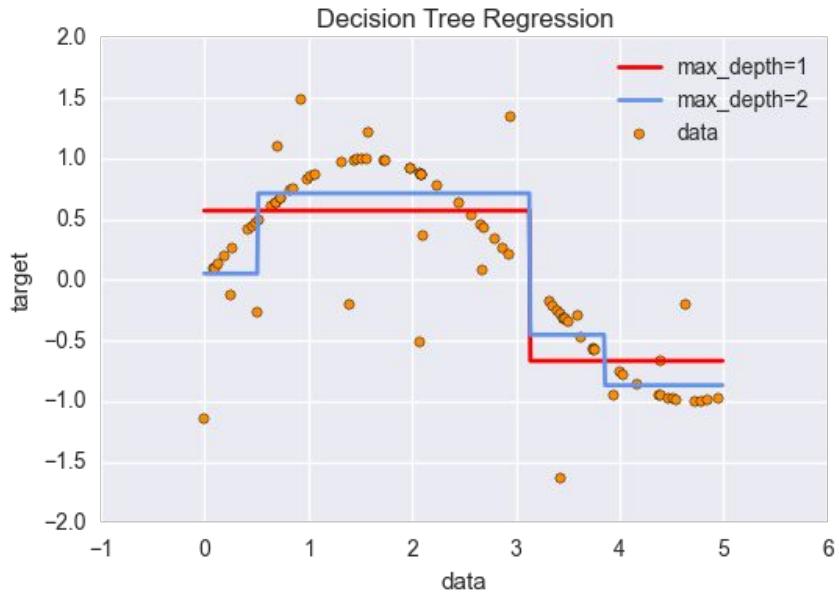
mse = 0.1919  
samples = 11  
value = 0.0524

mse = 0.1479  
samples = 40  
value = 0.7138

mse = 0.1241  
samples = 14  
value = -0.4519

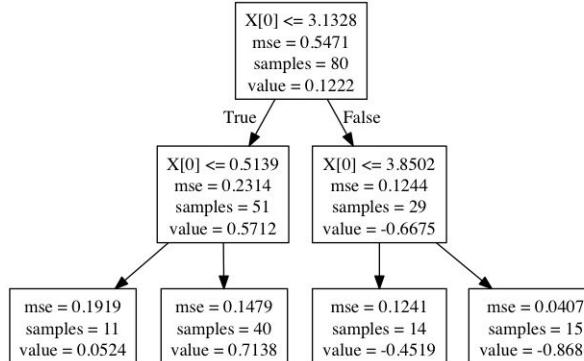
mse = 0.0407  
samples = 15  
value = -0.8686

# Some terminology

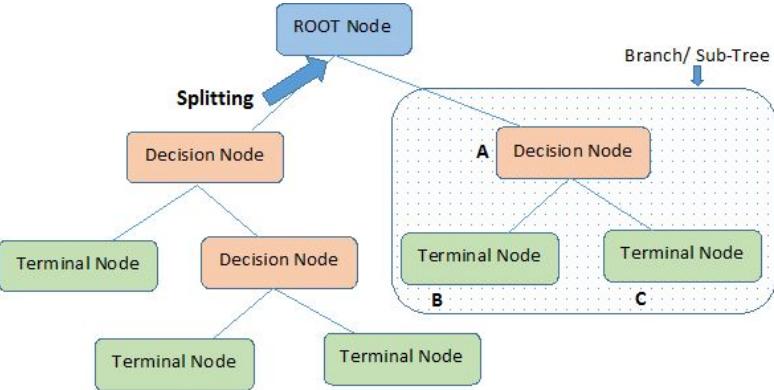


Note:- A is parent node of B and C.

**max\_depth=2**



# Some terminology

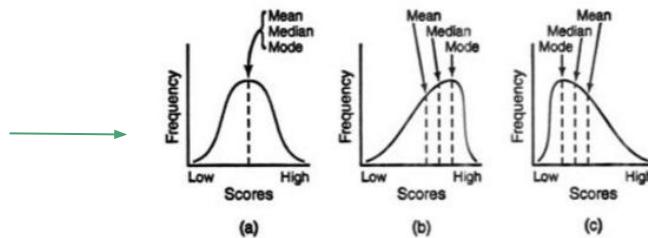


Note:- A is parent node of B and C.

1. **Root Node:** It represents entire population or sample and this further gets divided into two or more homogeneous sets.
2. **Splitting:** It is a process of dividing a node into two or more sub-nodes.
3. **Decision Node:** When a sub-node splits into further sub-nodes, then it is called decision node.
4. **Leaf/ Terminal Node:** Nodes do not split is called Leaf or Terminal node.
5. **Pruning:** When we remove sub-nodes of a decision node, this process is called pruning. You can say opposite process of splitting.
6. **Branch / Sub-Tree:** A sub-section of entire tree is called branch or sub-tree.
7. **Parent and Child Node:** A node, which is divided into sub-nodes is called parent node of sub-nodes where as sub-nodes are the child of parent node.

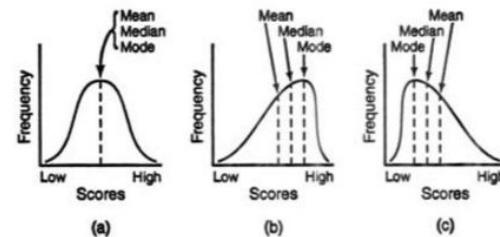
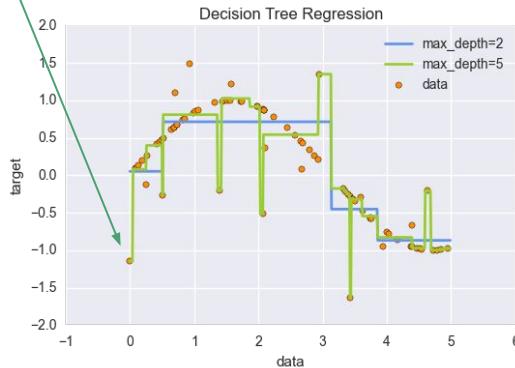
# What are the advantages of decision trees?

- Very easy to interpret.
- Makes **no a-priori assumptions about the structural form of the data** (as linear models do). For instance, works well with non-linear data.
- More “robust” than linear models, meaning **less sensitive to outliers**. This is for the same reason that the median is less sensitive to outliers than the mean.
- Well defined notion of ‘**most significant variables**’, so good for data exploration (will explain).
- Handles categorical and real-valued variables (doesn’t require one-hot encoding as linear models always do - Exercise: Why?)



# What are the disadvantages?

- Prone to **overfitting** (more on this next lecture).
- Can lose information when dealing with continuous variables, since it needs to make a finite number of splits.



# More than one variable?

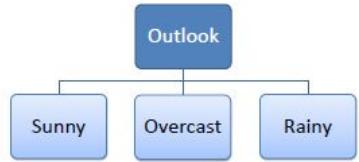


- A decision tree is simply a recursive set of decisions which leads to a result.
- It can be a real value or a class.
- Decision trees are very simple to understand, but are somewhat complex to explain when multiple features are involved.

**Machine Learning Objective:** Let's try to predict the number of hours played on a golf course in a single day by all of the golfers.

Example taken from: [http://chem-eng.utoronto.ca/~datamining/dmc/decision\\_tree\\_reg.htm](http://chem-eng.utoronto.ca/~datamining/dmc/decision_tree_reg.htm)

# How do we choose the root node?



Outlook	Temp	Humidity	Windy	Hours Played
Sunny	Mild	High	FALSE	45
Sunny	Cool	Normal	FALSE	52
Sunny	Cool	Normal	TRUE	23
Sunny	Mild	Normal	FALSE	46
Sunny	Mild	High	TRUE	30
Rainy	Hot	High	FALSE	25
Rainy	Hot	High	TRUE	30
Rainy	Mild	High	FALSE	35
Rainy	Cool	Normal	FALSE	38
Rainy	Mild	Normal	TRUE	48
Overcast	Hot	High	FALSE	46
Overcast	Cool	Normal	TRUE	43
Overcast	Mild	High	TRUE	52
Overcast	Hot	Normal	FALSE	44

		Hours Played (StDev)
Outlook	Overcast	3.49
	Rainy	7.78
	Sunny	10.87
SDR=1.66		

		Hours Played (StDev)
Temp.	Cool	10.51
	Hot	8.95
	Mild	7.65
SDR=0.17		

		Hours Played (StDev)
Humidity	High	9.36
	Normal	8.37
SDR=0.28		

		Hours Played (StDev)
Windy	False	7.87
	True	10.59
SDR=0.29		

- Here we've tried splitting first by the variable "Outlook".
- From here, we can compute the conditional standard deviations in the three subsets created.
- Our goal is to find the variable that, when split, reduces the stdev the most.
- Original stdev of "Hours Played" is 9.33.
- We can try this for every variable, and then choose the one which reduces the stdev the most.
- Subsequent decisions are made by recursively iterating this procedure.

# More precise description

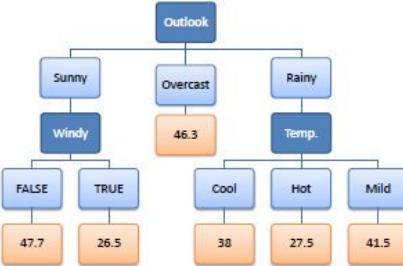
Predictors				Target
Outlook	Temp.	Humidity	Windy	Hours Played
Rainy	Hot	High	False	26
Rainy	Hot	High	True	30
Overcast	Hot	High	False	48
Sunny	Mild	High	False	46
Sunny	Cool	Normal	False	62
Sunny	Cool	Normal	True	23
Overcast	Cool	Normal	True	43
Rainy	Mild	High	False	36
Rainy	Cool	Normal	False	38
Sunny	Mild	Normal	False	48
Rainy	Mild	Normal	True	48
Overcast	Mild	High	True	62
Overcast	Hot	Normal	False	44
Sunny	Mild	High	True	30



For zero depth we minimize:

$$\text{Var}(y) := \frac{1}{N} \sum_{i=1}^N (y_i - \bar{y})^2$$

Solution is:  $\hat{y} = \bar{y}$



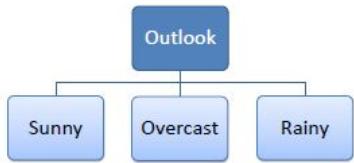
$$\begin{aligned} \text{Var}(Y|X = x_j) &= \sum_{i=1}^N (y_i - \bar{y}_j)^2 p(y_i|x_j) \\ &= \frac{1}{N} \sum_{i,x_i=x_j} (y_i - \bar{y}_j)^2 |X = x_j| \end{aligned}$$

$$\text{VarRed}(Y, X) = \text{Var}(Y) - \sum_j \text{Var}(Y|X = x_j)$$

For depth one we minimize the conditional variance for each variable:

Variance Reduction

# Let's try Outlook first



Outlook	Temp	Humidity	Windy	Hours Played
Sunny	Mild	High	FALSE	45
Sunny	Cool	Normal	FALSE	52
Sunny	Cool	Normal	TRUE	23
Sunny	Mild	Normal	FALSE	46
Sunny	Mild	High	TRUE	30
Rainy	Hot	High	FALSE	25
Rainy	Hot	High	TRUE	30
Rainy	Mild	High	FALSE	35
Rainy	Cool	Normal	FALSE	38
Rainy	Mild	Normal	TRUE	48
Overcast	Hot	High	FALSE	46
Overcast	Cool	Normal	TRUE	43
Overcast	Mild	High	TRUE	52
Overcast	Hot	Normal	FALSE	44

$$\begin{aligned} \text{Var}(Y|X = x_j) &= \sum_{i=1}^N (y_i - \bar{y}_j)^2 p(y_i|x_j) \\ &= \frac{1}{N} \sum_{i,x_i=x_j} (y_i - \bar{y}_j)^2 |X = x_j| \end{aligned}$$

$$\text{VarRed}(Y, X) = \text{Var}(Y) - \sum_j \text{Var}(Y|X = x_j)$$

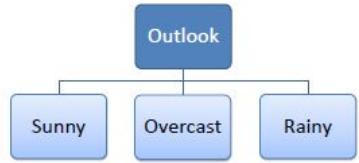
	Hours Played (StDev)	Count
Outlook	Overcast	3.49
	Rainy	7.78
	Sunny	10.87
		14

$$\text{Var}(y) := \frac{1}{N} \sum_{i=1}^N (y_i - \bar{y})^2 = 9.32^2$$

$$\sum_j \text{Var}(Y|\text{Outlook}) = \frac{4}{14}(3.49)^2 + \frac{5}{14}(7.78)^2 + \frac{5}{14}(10.87)^2$$

- Therefore there is a variance reduction of **19.56 for the outlook variable.**

# Now we've split based on Outlook



Outlook	Temp	Humidity	Windy	Hours Played
Sunny	Mild	High	FALSE	45
Sunny	Cool	Normal	FALSE	52
Sunny	Cool	Normal	TRUE	23
Sunny	Mild	Normal	FALSE	46
Sunny	Mild	High	TRUE	30
Rainy	Hot	High	FALSE	25
Rainy	Hot	High	TRUE	30
Rainy	Mild	High	FALSE	35
Rainy	Cool	Normal	FALSE	38
Rainy	Mild	Normal	TRUE	48
Overcast	Hot	High	FALSE	46
Overcast	Cool	Normal	TRUE	43
Overcast	Mild	High	TRUE	52
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		Hours Played (StDev)
Humidity	High	9.36
	Normal	8.37
SDR=0.28		

		Hours Played (StDev)
Windy	False	7.87
	True	10.59
SDR=0.29		

- These groups now have less variance in the data. And Outlook reduced variance the most. Now we continue recursively.
- A branch set with standard deviation more than 0 needs further splitting - we need pure classes on the final leaf nodes.

# Common Questions

- **What about real valued inputs?** Create a histogram of the variable, then do binary splits.
- **Are splits always binary?** In Python, yes, but not for a general algorithm. Any 3 way split can be seen as a one versus all split (ie. A&B or C versus A, B or C). It depends on the algorithm.

Follow the notebook here:

[https://github.com/doriang102/Columbia\\_Data\\_Science/blob/master/notebooks/Lecture1%20-%20Introduction-to-Regression.ipynb](https://github.com/doriang102/Columbia_Data_Science/blob/master/notebooks/Lecture1%20-%20Introduction-to-Regression.ipynb)