# Lecture 8: Linear Classifiers and More Model Validation

**INFO 1998: Introduction to Machine Learning** 



## **Agenda**

- 1. Perceptron + SVM
- 2. More Cross-Validation techniques



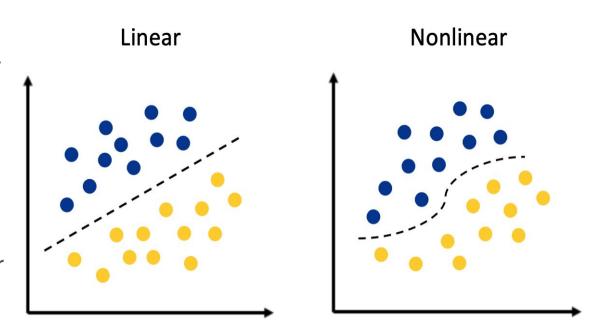
# **Linear Classifiers**



### **Linear Classifiers**

A linear classifier is a hyper plane that is used to classify our data points

A hyperplane is our decision boundary and our goal is to find the hyper plane that best classifies our data



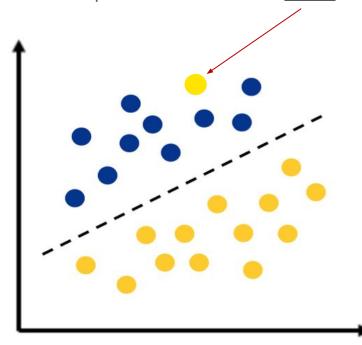


## **Linearly Separable**

In this example, we cannot partition our dataset into yellow and purple with a linear decision boundary. This means that our data is not linearly separable.

Outliers are frequently the reason a data set is not linearly separable.

This data set is not linearly separable because of an <u>outlier</u>





### **Perceptron Learning Algorithm**

Goal: find a normal vector w that perfectly classifies all the points in our data set Algorithm:

Initialize classifier as some random hyperplane
While there exists a misclassified point x:

Tilt classifier slightly so that it classifies x correctly

(or, is a little closer to classifying x correctly)

End While

"Use your mistakes as your stepping stones"





### **Limitations of Perceptron**

The training algorithm will never terminate if your training dataset is not linearly separable  $\cong$ 

Is a great model to understand the intuition behind the training of a linear classifier: iteratively improve classifier by using misclassified points ©

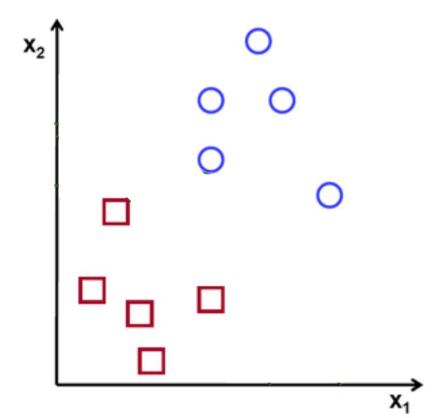




# **SVM**



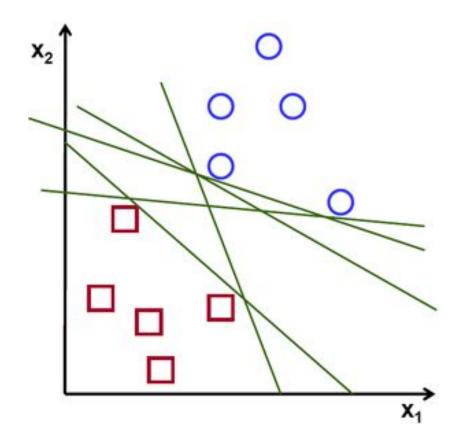
# Classify (+) and (-)







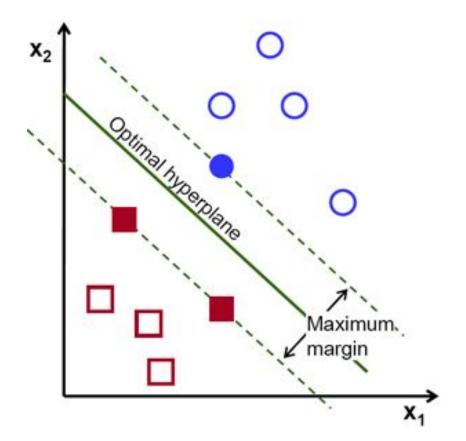
# Which Hyperplane?







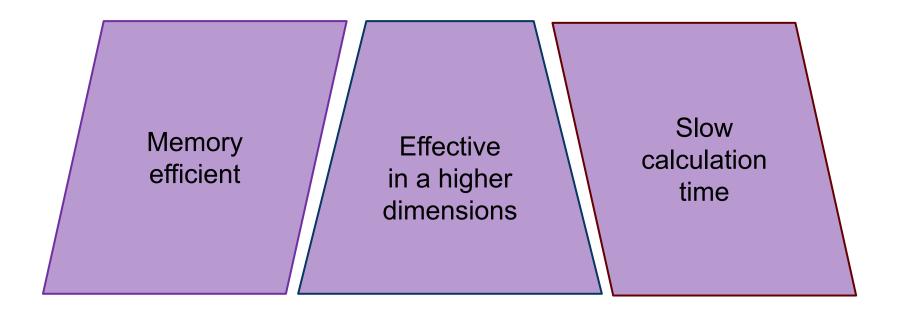
# **Optimal Hyperplane**







## **Support Vector Machine**

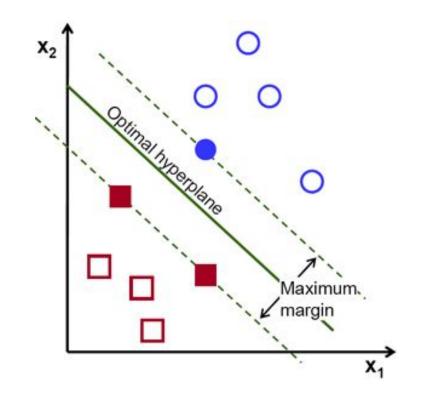






## **Maximal Margin Classifier**

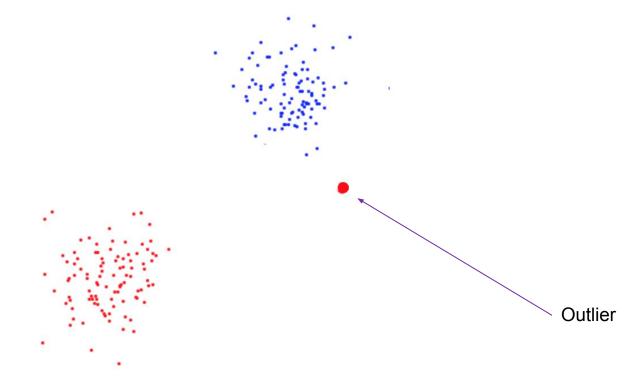
- We want to find a separating hyperplane
- Once we find candidates for the hyperplane, we try to maximize the margin, the normal distance from borderline points
  - Only Support Vectors matter







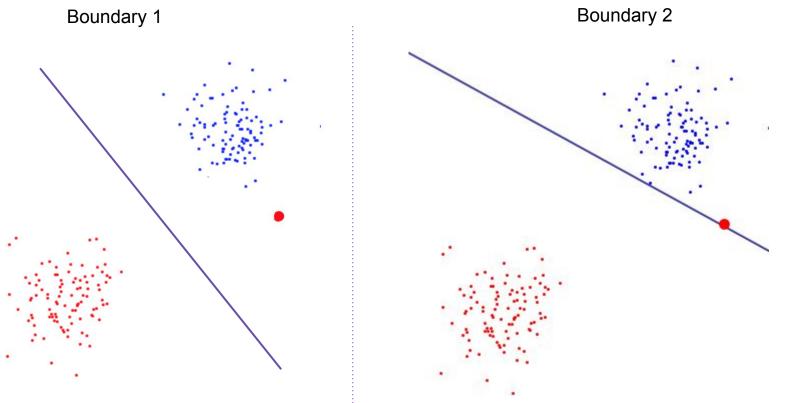
# What if...







# Which Decision Boundary is better?







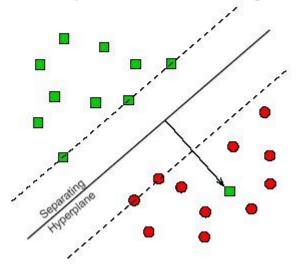
### **Margins**

Use cost function to penalize misclassified points

Choice of cost function makes margin "hard" vs. "soft"

#### Non-separable training sets

Use linear separation, but admit training errors.



Penalty of error: distance to hyperplane multiplied by error cost C.

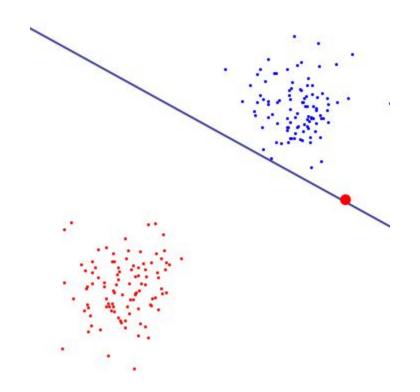




### **Hard Margins**

High penalty value

 The hyperplane can be dictated by a single outlier





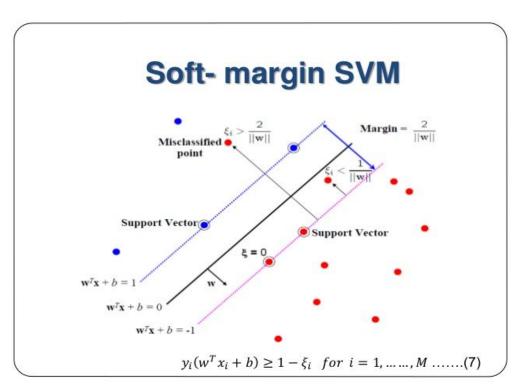


### **Soft Margins**

Used in non-linearly separable datasets

Allow for misclassification

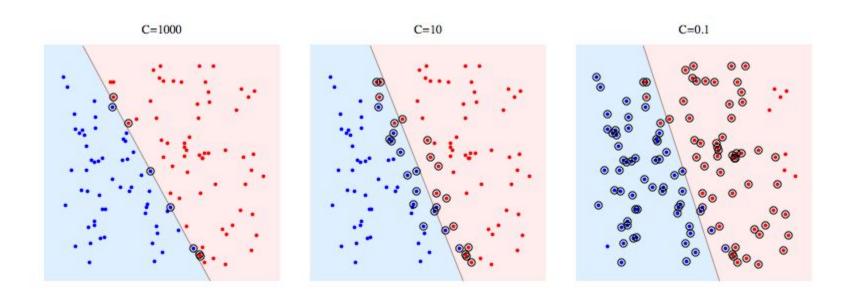
 Can account for "dirty" boundaries







# **Misclassification Penalty C**

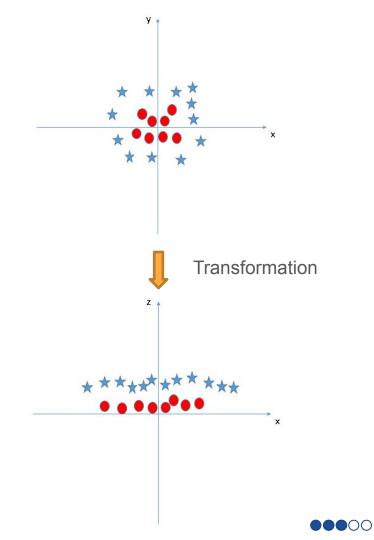






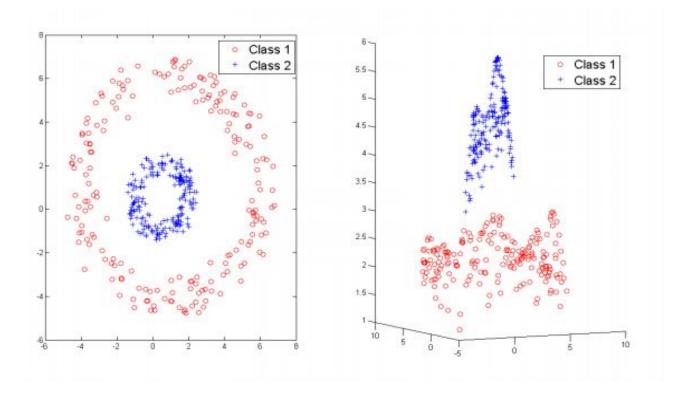
### **Kernels**

- You cannot linearly divide the 2 classes on the xy plane at right
- Introduce new feature,  $z = x^2 + y^2$  (radial kernel)
- Map 2 dimensional data onto 3 dimensional data. Now a hyperplane is easy to find. (Imagine slicing a cone!)





## **Kernels**







### **SVM** has **MANY** Hyperparameters

### SVM

#### C

The "penalty cost" for misclassifications (soft margins)

#### Gamma

How far the influence of a single training example reaches

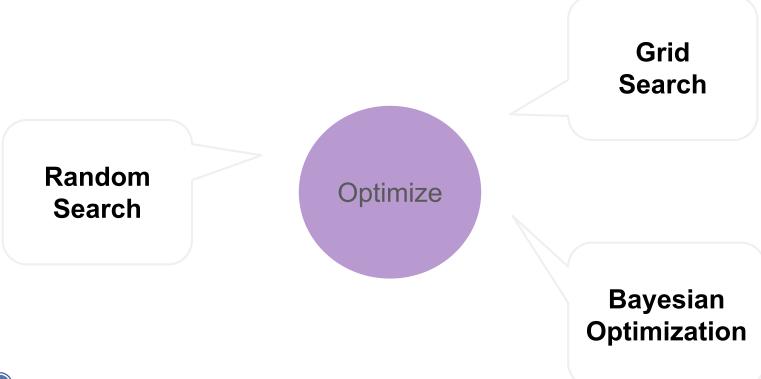
### Kernels

Method of transforming our data set





### **Finding the Best Hyper Parameters**

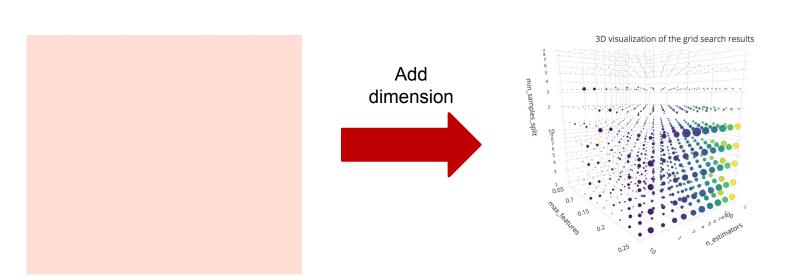


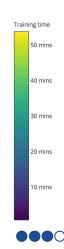




### **Curse of Dimensionality**

Our search space for the optimal hyper-parameters increases **exponentially** as the number of hyper parameters we are considering increases







### **Overview**

Perceptron	SVM
<ul> <li>A very simple model</li> <li>Will perform poorly if data is not linearly separable</li> </ul>	<ul> <li>More complex model because we have to choose the "penalty cost" associated with misclassifications</li> <li>Can transform feature space by choosing a Kernel</li> </ul>





# Demo



### **Bias-Variance Tradeoff**

#### In regression:

Want to predict y for a fixed test data point x

*y* is generated from the true model  $y = f(x) + \epsilon$  ( $\epsilon$  has zero mean and independent of everything else)

Estimate using some model  $\hat{f}$ :  $\hat{y} = \hat{f}(x)$ 

Then:

test error = 
$$\mathbb{E}(\hat{f}(x) - y)^2 = \underbrace{\left(\mathbb{E}\hat{f}(x) - f(x)\right)^2}_{\text{bias}^2} + \underbrace{\text{Var}(\hat{f}(x))}_{\text{variance}} + \underbrace{\text{Var}(\epsilon)}_{\text{irreducible error}}$$





### **Proof (not required, only for reference)**

Suppose we have test set data:  $y = f(x) + \mathcal{E}$ . The test data y is random, since  $\mathcal{E}$  is random. The estimator f hat is also random, because it is fitted on random training data.

Then from the previous slide "MSE = bias^2 + variance" for unknown quantity  $\theta$  and its random estimator  $\theta$  hat, we have:

$$\mathbb{E}(\hat{f}(x) - y)^{2}$$

$$= \mathbb{E}\left[\mathbb{E}\left[(\hat{f}(x) - y)^{2} | y\right]\right]$$

$$= \mathbb{E}\left[\left(\mathbb{E}\hat{f}(x) - y\right)^{2} + \text{Var}(\hat{f}(x))\right]$$

$$= \mathbb{E}\left(\mathbb{E}\hat{f}(x) - f(x) - \epsilon\right)^{2} + \text{Var}(\hat{f}(x))$$

$$= \mathbb{E}\left(\mathbb{E}\hat{f}(x) - f(x)\right)^{2} + 2\mathbb{E}\left[\left(\mathbb{E}\hat{f}(x) - f(x)\right)\epsilon\right] + \mathbb{E}\epsilon^{2} + \text{Var}(\hat{f}(x))$$

(For some part of this lecture material, credit goes to Professor Damek Davis, Cornell ORIE department.)





### **Bias-Variance Tradeoff**

$$\underbrace{\mathbb{E}(\hat{f}(x) - y)^{2}}_{\text{test error}} = \underbrace{\left(\mathbb{E}\hat{f}(x) - f(x)\right)^{2}}_{\text{bias}^{2}} + \underbrace{\text{Var}(\hat{f}(x))}_{\text{variance}} + \underbrace{\text{Var}(\epsilon)}_{\text{irreducible error}}$$

Less flexible model:

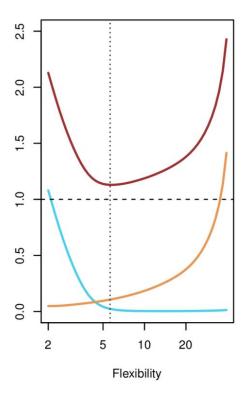
$$\hat{f}(x) = \hat{\beta}_0 + \hat{\beta}_1 x$$

bias high, variance low

More flexible model:

$$\hat{f}(x) = \hat{\beta}_0 + \hat{\beta}_1 x + \hat{\beta}_2 x^2 + \hat{\beta}_3 x^3$$

bias low, variance high



# More Validation Techniques



### Review: Regression vs. Classification

## Regression

- Predict Continuous Data
- "On average, how wrong are we?"

### Classification

- Predict Discrete or Categorical data
- "How many points do we get wrong?"

Numbers



Continuous

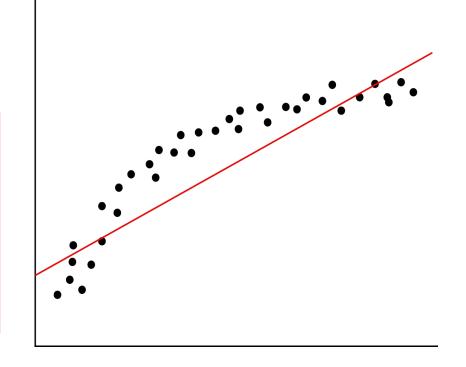




### **Underfitting**

Underfitting means we have <u>high bias</u> and <u>low variance</u>.

- Lack of relevant variables/factor
- Imposing limiting assumptions
  - Linearity
  - Assumptions on distribution
  - Wrong values for parameters



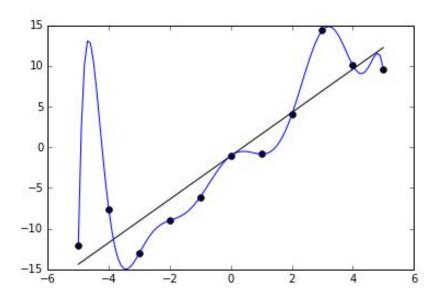




### **Overfitting**

Overfitting means we have <u>low bias</u> and <u>high variance</u>.

- Model fits too well to specific cases
- Model is over-sensitive to sample-specific noise
- Model introduces too many variables/complexities than needed







### **Bias-Variance Tradeoff**

When evaluating the models we are using, one property that we would often refer to is the bias-variance tradeoff.

Bias-variance tradeoff is the property of a set of predictive models where:

- Models with higher flexibility would typically have lower bias and higher variance;
- Models with lower flexibility would typically have higher bias and lower variance.

The graph gives a visualization on the outcomes of different bias and variance if we are doing experiment of throwing darts.

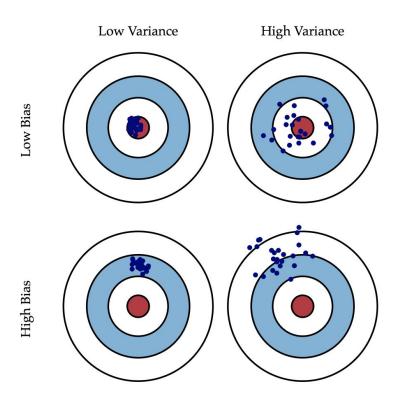


Fig. 1 Graphical illustration of bias and variance.

Graph source: <a href="http://scott.fortmann-roe.com/docs/BiasVariance.html">http://scott.fortmann-roe.com/docs/BiasVariance.html</a>





### **Bias-Variance Tradeoff**

#### Say:

- $\triangleright$   $\theta$  is an unknown quantity we want to estimate
- $\triangleright$   $\hat{\theta}$  is an estimator of  $\theta$  (computed from data, random)

The mean squared error (MSE) decomposes as:

$$\underbrace{\mathbb{E}(\hat{\theta} - \theta)^{2}}_{\text{MSE}} = \underbrace{\left(\mathbb{E}\hat{\theta} - \theta\right)^{2}}_{\text{bias}^{2}} + \underbrace{\text{Var}(\hat{\theta})}_{\text{variance}}$$

$$MSE = bias^2 + variance$$





### Leave-P-out

Let **D** be our whole dataset

Choose a P

For every combination of **P** points in **D**:

Use a train/test split with those **P** points as test, the rest as train





## Leave-P-out: different from K-fold!

Let's say **D** has a size of 4. There are four data points: *a, b, c,* and *d*. K-fold:

- K = 2.
- Each fold has a size of 2: {*a*,*b*} and {*c*,*d*}
- So, we only have 2 possible test sets: {a,b} and {c,d}

#### Leave-P-out:

- P = 2.
- We have 6 possible test sets: {a,b}, {a,c}, {a,d}, {b,c}, {b,d}, and {c,d}





#### Leave-P-out

#### Pros:

- Dependable (not random)
- Representative checks all combinations

#### Cons:

- Slow!
  - Runtime <u>increases</u> with larger datasets
  - Runtime <u>explodes</u> with larger P





#### **Monte Carlo CV**

- Getting accuracy 1 time doesn't tell us much
- Getting accuracy 2 times tells us a bit
- Getting accuracy 3 times tells us a bit more
- ...
- Getting accuracy N times might be good enough!

Take the average of those **N** times





### **Monte Carlo CV**

- Need to use new, random train/test split each time
  - If you use the same train/test split each time, you're not getting any new information!
- Pros:
  - easy to implement
  - easy to make faster/slower by changing number of iterations
- Cons:
  - random -> train/test splits not guaranteed to be representative of dataset
  - harder to calculate how many iterations you need





## The Bootstrap

#### What if we don't have enough data?

- Use **bootstrap datasets** to approximate the test error
- Sample with replacement from the original training dataset (with n samples) to generate bootstrap datasets of size n
  - Some data points may appear more than once in the generated data
  - Some data points may not appear
- Estimate of test error = average error among bootstrap datasets





# Pipeline of the Bootstrap

#### **Algorithm 1** Estimating prediction error via the bootstrap

- Input: dataset {(x<sub>i</sub>, y<sub>i</sub>)}<sup>n</sup><sub>i=1</sub>, loss function ℓ.
   for b = 1, 2, ..., B do
   generate set S<sub>b</sub> by sampling n items with replacement from {(x<sub>i</sub>, y<sub>i</sub>)}<sup>n</sup><sub>i=1</sub>.
   form prediction f̂<sup>b</sup> by fitting the logistic regression model with training data S<sub>b</sub>.
   compute err<sub>b</sub> := 1/n ∑<sup>n</sup><sub>i=1</sub> ℓ(y<sub>i</sub>, f̂<sup>b</sup>(x<sub>i</sub>))
   end for
   return err<sub>boot</sub> := 1/B ∑<sup>B</sup><sub>b=1</sub> err<sub>b</sub>
- Does this give very good estimate of the prediction error?
- If no, why not? Does it underestimate or overestimate the prediction error?





# Pipeline of the Bootstrap

No.

Since the bootstrap is sampling with replacement for B validation folds, each fold would have significant overlap with the original data used for training. Approximately, **2/3** of the training data would appear in each validation fold.

This leads to significant **underestimation** of the prediction error.

**❖** Why **2/3**?





#### **Proof**

Suppose there are n samples (yi, xi) in the training set data, then for each of the bootstrap validation fold b, every sample has probability 1/n of being selected into b. The probability of each data sample (yi, xi) not being in fold b is (1 - 1/n), respectively.

Probability of avoid selecting all n data points in fold b: (1 - 1/n)^n.

When n gets large, we have this probability converges to 1/e.

(Why this converges? Probably should review Calc I, or see:

https://math.stackexchange.com/questions/882741/limit-of-1-x-nn-when-n-tends-to-infinity.)

Therefore, the fraction of overlapping data points in each fold is (1-1/e), which is about 2/3.





# How we fix the problem

- Requires some thoughts when generating validation set, especially for real-world complex data.
- Can partly fix this problem by only using predictions for those observations that did not (by chance) occur in the current bootstrap sample.
- But the method gets complicated.





# Bootstrap vs. k-fold

In K-fold validation, each of the K folds is distinct from the other (K - 1) folds used for training: there is **no overlap**.

This is crucial for its success in estimating prediction error.





# Why we still use bootstrap

- Bootstrap allows us to use a computer to mimic the process of obtaining new data sets.
- Can be used to quantify the uncertainty associated with a given estimator or statistical learning method.
- Provides an estimate of the standard error of a coefficient, or a confidence interval for that coefficient.
  - i.e., the variability of the model!





# **Bagging (Bootstrap Aggregating)**

#### What if we don't have enough data?

- Bagging is a common technique that builds on Bootstrapping
- Main Idea: Do Bootstrapping a bunch and make a classifier for each bootstrap, then majority prediction wins.
- Many weak learners aggregated typically outperform a single learner over the entire set, and overfits less.
  - Principle behind Random Forests





# Demo



# **Measures of overfitting**

- Compare *train* accuracy with *test* accuracy!
- Method 1: Generalization error
  - (Generalization error) = (test error) (train error)
  - Error introduced from trying to generalize to new data
- Method 2: Prediction variation
  - VAR(predictions for a specific point x)
  - How much do predictions change for point x, based on the training data?

Reminder: don't mix up error and accuracy! They're nearly equivalent measures, but confusing them in your code can cause problems





# **Coming Up**

- Assignment 8: Due before next class
- Next Lecture: Applications of Unsupervised Learning

