# CS 270 Final Exam Review

Jake Cahoon

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# Introduction

## 1.1 Note From Jake

I hope the last review document was helpful, and I hope this one is as well. To make these, I've gone through the Exam Study Guide topics posted on Learning Suite. With the hope that I don't miss topics that will be on the final, I went through every slide and included all that I deem noteworthy. Despite this endeavor, I will miss topics, so I suggest you use this as a supplement to your study rather than relying solely on it.

I've included a table of contents this time around, so you can jump to the topics you need to study. That being said, let's friggin' do this.

# Data Mining Process Model

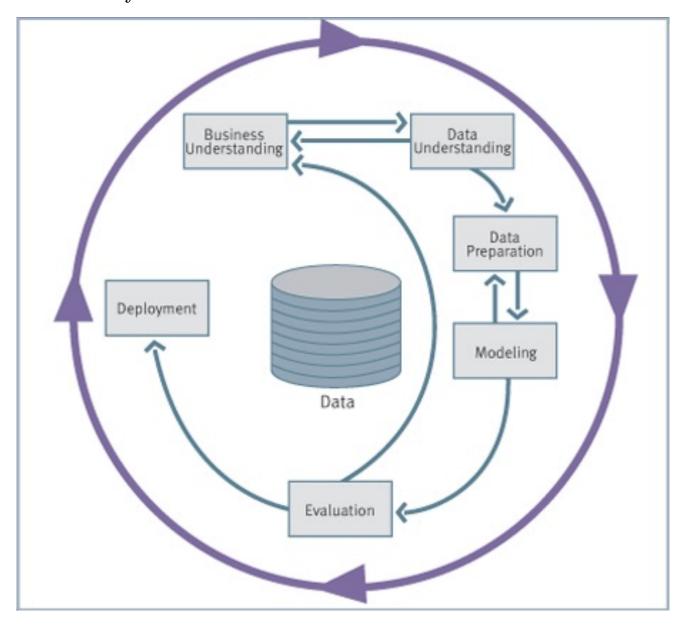
## 2.1 The Process

#### Note:-

I'm pulling straight from the slides here. Know this process at a high level.

- 1. Identify and define the task (business understanding)
  - Understand the context, audience, and problem
  - Tell the story
- 2. Gather and prepare the data
  - Build a dataset for the task
  - Select/transform/derive features
  - Conduct exploratory data analysis
  - Clean the data
- 3. Build and evaluate model
- 4. Deploy the model
  - Evaluate business related results
- 5. Iterate and improve the model

## 2.2 The Cycle Picture



# Bayesian Learning

There are two approaches to statistical learning: frequentist and Bayesian. Frequentist statistics is based on the idea of repeated sampling, while Bayesian statistics is based on the idea of starting with prior beliefs and then updating beliefs based on new information.

## 3.1 Bayes Theorem

#### Definition 3.1.1

Let C and A be events. Then

$$P(C|A) = \frac{P(A|C)P(C)}{P(A)}$$

Note:-

The right side of the equation is based on our current data, while the left side is what we want to find.

#### 3.1.1 Notes

- Prior probabilities are based on prior knowledge. They are the initial beliefs.
- Posterior probabilities are the updated beliefs based on new information.
- P(C) is the prior probability of the Class.
- P(A) is the prior probability of the Attribute.
- P(A|C) is the likelihood of the Attribute given the Class.
- P(C|A) is the posterior probability of the Class given the Attribute.

## 3.2 Bayesian Classifiers

Given a set of attributes  $\{A_1, A_2, \dots, A_n\}$  and a class C, we can use Bayes Theorem to find the output class C that maximizes  $P(C|A_1, A_2, \dots, A_n)$ . For each output class C, do

$$P(C|A_1, A_2, \dots, A_n) = \frac{P(A_1, A_2, \dots, A_n | C)P(C)}{P(A_1, A_2, \dots, A_n)}$$

## 3.3 Maximum A Posteriori (MAP) Estimation

#### Definition 3.3.1

Let D be a dataset and let H be the set of all hypotheses. Then

$$\hat{h}_{MAP} = \operatorname*{argmax}_{h \in H} P(h|D)$$

where  $\hat{h}_{MAP}$  is the maximum a posteriori hypothesis.

This is guaranteed to be "best", but it is computationally expensive and impractical for large hypothesis spaces.

## 3.4 Bayes Optimal Classifier

TODO: Figure out what to add here.

## 3.5 Naive Bayes Classifiers

A simple classifier that assumes that the attributes are conditionally independent given the class. This is a naive assumption, but it works well in practice. We assume that:

$$P(A_1, A_2, \dots A_n | C) = P(A_1 | C) \cdot P(A_2 | C) \cdot \dots \cdot P(A_n | C)$$

In other words, the probability of all the attributes given the class is the product of the probabilities of each attribute given the class. Then for each  $A_i \in A$  and for each  $C_j \in C$  we can estimate  $P(A_i|C_j)$ , which you did when you calculated the probabilities in the HW.

Once we have the probabilities, we can classify a new instance X by

$$\hat{C} = \underset{C_j \in C}{\operatorname{argmax}} \left( P(C_j) \cdot \prod_{i=1}^n P(A_i | C_j) \right)$$

#### 3.5.1 Notes

- Various  $P(C_j)$  and  $P(A_i|C_j)$  are estimated from the training data.
- Stores the probabilities in a table.
- For a new instance X, the classifier calculates the probability of each class given the attributes.
- $\hat{C}$  is the class with the highest probability.
- The true probability is the normalized probability.
- Independence assumption may not hold for some attributes.

## **Ensembles**

Two heads are better than one, not because either is infallible, but because they are unlikely to go wrong in the same direction. - C.S. Lewis

## 4.1 Bias and Variance

- Bias is the error due to overly simplistic assumptions in the learning algorithm.
- Variance is the error due to the algorithm's sensitivity to fluctuations in the training data.

Bias and variance are inversely related. As one goes up, the other goes down. The goal is to minimize both.

## 4.2 Why are Ensembles Helpful?

By using ensembles, we can reduce the bias and variance of our models. Ensembles combine multiple models to create a stronger model. The idea is that the models will make different errors, and by combining them, we can reduce the overall error. See Dr. Snell's slides for examples.

#### 4.2.1 Four Important Criteria

- 1. Independence: The models should be independent.
- 2. Diversity: The models should be different enough to make different errors.
- 3. Decentralization: The models should be trained on different subsets of the data.
- 4. Aggregation: The models should be combined in a way that reduces error.

## 4.3 Voting Ensemble

Let T be the training set,  $A = \{A_1, A_2, \dots, A_n\}$  be the set of models, and C be the set of classes. Define a function  $\delta$  as follows:

$$\delta(a,b) = \begin{cases} 1 & \text{if } a = b \\ 0 & \text{otherwise} \end{cases}$$

Then the voting ensemble is defined as

- 1. For k = 1 to N,  $h_k$  is a model trained on T using learning algorithm  $A_k$ .
- 2. For a new instance X, the ensemble predicts

$$\hat{C} = \underset{c \in C}{\operatorname{argmax}} \sum_{k=1}^{N} \delta(c, h_k(X))$$

#### 4.3.1 Notes

- Key issues: diversity and independence (too small of a set A and too similar models will not help).
- The models should be trained on different subsets of the data.

## 4.4 Bagging

Let T be the training set, A be the learning algorithm, N be the number of samples (or bags) of size d drawn from T, C be the set of classes, and  $\delta$  be defined as in the Voting Ensembles section. Then the bagging ensemble is defined as

- 1. For k = 1 to N,  $S_k \subseteq T$  is a sample of size d drawn from T, with replacement and  $h_k$  is a model trained on  $S_k$  using learning algorithm A.
- 2. For a new instance X, the ensemble predicts

$$\hat{C} = \underset{c \in C}{\operatorname{argmax}} \sum_{k=1}^{N} \delta(c, h_k(X))$$

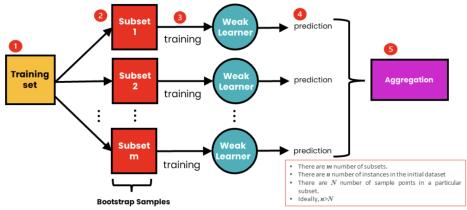
#### 4.4.1 Random Forest

- A bagging ensemble extension of decision trees
- Each tree is trained on a different bootstrap sample
- Two random variables: the bootstrap sample and the feature subset

#### 4.4.2 Notes

- Train N models on N different bootstrap samples
- Combines the outputs by voting ( $\delta$  function)
- Decreases error by reducing variance due to unstable learning algorithms
- · Homogeneous models are used

## The Process of Bagging (Bootstrap Aggregation)



## 4.5 Boosting

<sup>&</sup>quot;Boost weak learners into strong learners."

#### 4.5.1 AdaBoost

AdaBoost learns from mistakes by increasing the weights of the misclassified instances. His slides go quite in depth on AdaBoost (adaptive boosting), so I would recommend reviewing them.

- 1. Start with uniform weights
- 2. Train a weak learner
- 3. Update the weights based on the performance of the weak learner
- 4. Repeat

## 4.5.2 Gradient Boosting

Unlike AdaBoost, Gradient Boosting learns from residual errors, rather than directly updating the weights.

- 1. Train a weak learner
- 2. Compute the residuals (errors) on the training set
- 3. Train a new weak learner to predict the residuals
- 4. Repeat from step 2 until the residuals are small

#### 4.5.3 Notes

- Great for reducing bias
- Combines the outputs by weighted voting/averaging
- Homogeneous models are used
- Weak learners need to be better than random guessing
- Construct a strong learner by weighted voting of the weak learners

# Training set Subset Subset Weak False prediction Weak False prediction Weak Folse prediction Overall Prediction

## 4.6 Stacking

Let T be the base-level training set, N be the number of base-level learning algorithms,  $A = \{A_1, A_2, \dots, A_n, A_{\text{meta}}\}$  be the set of base-level learning algorithms, and  $A_{\text{meta}}$  be the chosen meta-level learner. Then the stacking ensemble is defined as

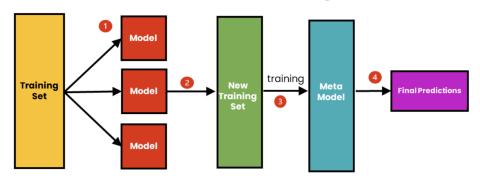
- 1. For i = 1 to N,  $h_i$  is a model trained on T using learning algorithm  $A_i$ .
- 2. Let  $T_m$  be the meta-level training set.  $T_m = \emptyset$ .
- 3. For k = 1 to |T|,  $E_k = \{h_1(X_k), h_2(X_k), \dots, h_N(X_k), y_k\}.$
- 4.  $T_m = T_m \cup E_k$ .
- 5.  $h_{\text{meta}}$  is a model trained on  $T_m$  using learning algorithm  $A_{\text{meta}}$ .
- 6. For a new instance X, the ensemble predicts

$$\hat{C} = h_{\text{meta}}(h_1(X), h_2(X), \dots, h_N(X))$$

#### 4.6.1 Notes

- Improves accuracy by combining the outputs of multiple models
- Hetereogeneous models are used at the base level
- The meta-level model is trained on the outputs of the base-level models

### The Process of Stacking



# Clustering

Clustering is an important type of unsupervised learning (PCA is unsupervised). The goal of clustering is to group similar data points together in a cluster.

## 5.1 K-Means

K-Means is a simple and popular clustering algorithm. The goal is to partition the data into k clusters. The algorithm works as follows:

## 5.2 Hierarchical Clustering

- 1. Start with an  $n \times n$  adjacency matrix
- 2. Initialize each point as its own cluster
- 3. Merge the two "closest" clusters (closeness is either single, complete, or average linkage)
- 4. Repeat until there is only one cluster

## 5.2.1 Walkthrough

#### Example 5.2.1

Compute the clusters for the following adjacency matrix using single linkage.

Note:-

I recommend x-ing out the values in the clusters you've already merged.

#### Solution:

1. Merge: b and d since the distance between them is 0.1.

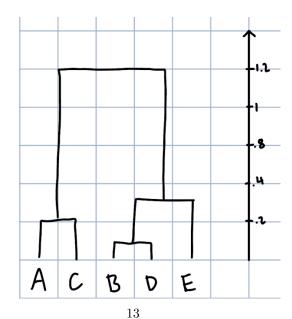
2. Merge: a and c since the distance between them is 0.2.

3. Merge:  $\{b, d\}$  and e since the distance between them is 0.3.

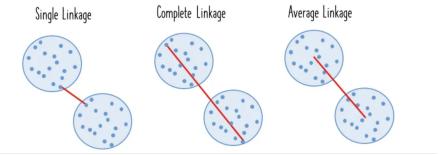
$$\begin{bmatrix} a & b & c & d & e \\ a & 0 & 1.4 & \times & 1.3 & 1.2 \\ b & 1.4 & 0 & 1.6 & \times & 0.4 \\ c & 1.6 & 0 & 1.5 & 1.4 \\ d & 1.3 & \times & 1.5 & 0 & 0.3 \\ e & 1.2 & 0.4 & 1.4 & 0.3 & 0 \end{bmatrix} \implies \begin{bmatrix} a & b & c & d & e \\ 0 & 1.4 & \times & 1.3 & 1.2 \\ 1.4 & 0 & 1.6 & \times & 0.4 \\ \times & 1.6 & 0 & 1.5 & 1.4 \\ 1.3 & \times & 1.5 & 0 & 0.3 \\ e & 1.2 & 0.4 & 1.4 & 0.3 & 0 \end{bmatrix}$$

4. Merge:  $\{a,c\}$  and  $\{b,d,e\}$  since the distance between them is 1.2.

5. Draw the dendrogram dude



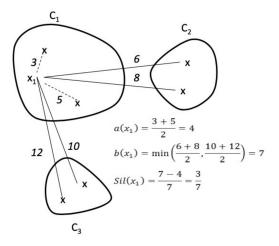
#### 5.2.2 Closeness



- Single linkage can lead to long chained clusters
- Complete linkage finds more compact clusters
- Average linkage is used less because it is computationally expensive

## 5.3 Silhouette Score

- A measure of how similar an object is to its own cluster compared to other clusters
- Ranges from -1 to 1 with 1 being the best
- A score of 1 means the object is well matched to its own cluster and poorly matched to neighboring clusters
- A score of 0 means the object is on the boundary of two clusters (ambiguous))
- A score of -1 means the object is probably in the wrong cluster



#### 5.3.1 Bois and Gurls to Write Down

Let i be a point in a cluster. Then

- a(i): The average distance (dissimilarity) between i and all other points in the same cluster (minimize a)
- b(i): The minimum average distance between i and all points in any other cluster (maximize b)
- Once we have a and b, we can calculate the silhouette score by

$$s(i) = \frac{b(i) - a(i)}{\max(a(i), b(i))}$$

## 5.3.2 Walkthrough

#### Example 5.3.1

Compute the silhouette scores for each point, each cluster, and the overall score for the following:

Cluster	Point	$\boldsymbol{x}$	y	
1	a	0.8	0.7	
1	b	0.9	0.8	
2	c	0.6	0.6	
2	d	0	0.2	
2	e	0.2	0.1	

Note:-

I would highly recommend making an adjacency matrix for easy calculation.

**Solution:** I recommend drawing a table and filling it out. Here's what it should look like:

Cluster	Point	x	y	a(i)	b(i)	s(i)
1	a	0.8	0.7	0.2	0.93	0.785
1	b	0.9	0.8	0.2	1.13	0.832
2	С	0.6	0.6	0.95	0.4	-0.579
2	d	0	0.2	0.65	1.4	0.536
2	e	0.2	0.1	0.6	1.3	0.538

I'll walk through two points, a and c.

$$a(a) = \frac{0.1 + 0.1}{1}$$

$$= 0.2$$

$$b(a) = \frac{0.3 + 1.3 + 1.2}{3}$$

$$= 0.93$$

$$s(a) = \frac{0.93 - 0.2}{\max(0.2, 0.93)}$$

$$= \frac{0.73}{0.93}$$

$$\approx 0.785$$

$$a(c) = \frac{1 + 0.9}{2}$$

$$= 0.95$$

$$b(c) = \frac{0.5 + 0.3}{2}$$

$$= 0.4$$

$$s(c) = \frac{0.4 - 0.95}{\max(0.95, 0.4)}$$

$$\approx -0.579$$

Now take the average of each cluster for the cluster scores:

$$s(1) = \frac{0.785 + 0.832}{2} \approx 0.804$$
  
$$s(2) = \frac{-0.579 + 0.536 + 0.538}{3} \approx 0.165$$

Finally, take the average of the cluster scores for the overall score:

$$s = \frac{0.785 + 0.832 - 0.579 + 0.536 + 0.538}{5} \approx 0.421$$

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# Reinforcement Learning

6.1 Q-Learning

Basic Precision/Recall

# **CNNs**

- 8.1 Structure
- 8.1.1 Convolutional Layers
- 8.1.2 Pooling Layers
- 8.1.3 Fully Connected Layers

# Other Deep Learning Topics

- 9.1 GANs
- 9.2 RNNs
- 9.3 LSTMs
- 9.4 Transformers