Lecture 2: Classification & Clustering

STATS 202: Data Mining and Analysis

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Announcements



- ► HW1 is (still) live
- ► Review session this Friday
 - ▶ Probability, Statistics, Calculus, & Linear algebra
 - ▶ Will be held live over Zoom
- ► TAs will be available to answer questions over chat

Outline



- Classification
 - K-nearest neighbors
 - ► Naive Bayes
- Clustering
 - K-means
 - ► Hierarchical clustering

Bayes classifier



- f₀ gives us a probability of the observation belonging to each class.
- ▶ To select a class, we can just pick the element in $f_0 = [p_1, p_2, ..., p_K]$ that's the largest
 - ► Called the Bayes Classifier
- ► As a classifier, produces the lowest error rate

Bayes error rate

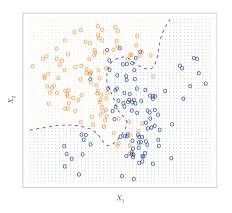
$$1 - \mathbb{E}_0 \left[\max_{y} \mathbb{P}_0[Y = y | X_1, X_2, ..., X_{\rho}] \right]$$
 (1)

Analogous to the irreducible error described previously

Bayes classifier



Example: Classifying in 2 classes with 2 features.



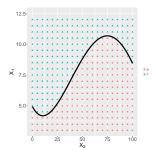
The Bayes error rate is 0.1304.

Bayes classifier

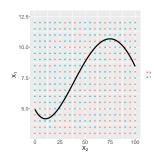


Note: $C(\mathbf{x}) = \underset{y}{\operatorname{arg max}} f_0(y)$ may seem easier to estimate

 \triangleright Can still be hard, depending on the distribution f_0 , e.g.



Bayes error = 0.0

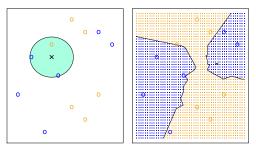


Bayes error = 0.3



How do we estimate Bayes classifier C(x)?

► Could just vote based on the *K* nearest neighbors (where *K* is some positive integer)

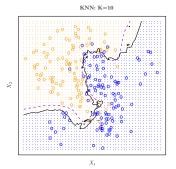


The KNN approach, using K = 3.



Using KNN (i.e. \hat{f}_n^{knn}) as a classifier $\mathcal{C}(\mathbf{x})$, we can estimate Bayes boundary f_0^* .

▶ Despite simplicity, \hat{f}_n^{knn} can be surprisingly close



The KNN (K=10) and Bayes decision boundaries.



Mathematically, we can represent KNN as

K-nearest neighbors

$$\mathbb{P}(Y=j|X=x_0)=\frac{1}{K}\sum_{i\in\mathcal{N}_0}\mathbb{I}(y_i=j)$$
 (2)

We can apply Bayes rule to the resulting probabilities to get our classifier.



Some details to consider in our KNN implementation:

- \blacktriangleright Are all our X_i 's on the same scale?
 - ► Typically, will standardize all features to be mean 0 and variance 1.



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$$d_{(i)} = ||x_{(i)} - x_0|| \tag{3}$$



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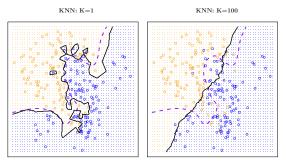
$$d_{(i)} = ||x_{(i)} - x_0|| \tag{3}$$

- Ties are typically broken randomly
- ▶ What size *K* do we use?
 - Estimated with e.g. test set



Higher values of K will result in smoother decision boundaries

► You're trading off higher variance for higher bias

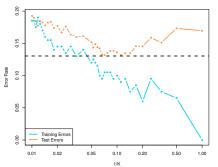


Two KNN boundary estimates (K = 1 and K = 100).



More flexibility (i.e. lower K) will result in over-fitting

► Similar to regression setting



KNN training/test errors as a function of K. Black line is Bayes error.



Another simple estimator is Naive Bayes.

By Bayes Theorem, we have

$$\mathbb{P}_{0}(Y|X_{1},X_{2}) = \frac{\mathbb{P}_{0}(Y)\mathbb{P}_{0}(X_{1},X_{2}|Y)}{\mathbb{P}_{0}(X_{1},X_{2})}$$
(4)

$$= \frac{\mathbb{P}_0(X_1, X_2, Y)}{\mathbb{P}_0(X_1, X_2)}$$
 (5)

We only care about the numerator

▶ It's a function of Y



Typically, we have

$$\mathbb{P}_{0}(X_{1}, X_{2}, Y) = \mathbb{P}_{0}(Y) \cdot \mathbb{P}_{0}(X_{1}|Y) \cdot \mathbb{P}_{0}(X_{2}|X_{1}, Y)$$
 (6)

However, we "naively" assume independence such that

$$\mathbb{P}_0(X_2|X_1,Y) \approx \mathbb{P}_0(X_2|Y) \tag{7}$$

Consequently, we have

$$\mathbb{P}_0(Y|X_1,X_2) \propto \mathbb{P}_0(Y)\mathbb{P}_0(X_1,X_2|Y)$$
 (8)

$$\approx \mathbb{P}_0(Y) \prod_{i=1}^2 \mathbb{P}_0(X_i | Y)$$
 (9)



$$\mathbb{P}_0(Y|X_1,X_2) \approx \frac{1}{Z}\mathbb{P}_0(Y)\prod_{i=1}^2\mathbb{P}_0(X_i|Y)$$

We can estimate \mathbb{P}_0 empirically

- e.g. kernel density estimation
- Could also use parametric models (e.g. Gaussian distribution)
- Question: What if the feature is categorical?



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Remark: we don't need Z if we're just classifying

Just take the class with the max value, e.g.

Example naive bayes classifier

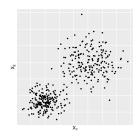
$$\hat{y}_n = \mathcal{C}(X_1, X_2) = \underset{y \in \{\text{Orange}, \text{Blue}\}}{\operatorname{arg max}} \mathbb{P}_0(y) \prod_{i=1}^2 \mathbb{P}_0(X_i | y) (10)$$

Clustering



Sometimes, we do not have the classes as our output Y. But we still want to assign each observation to a group.

- ► This is referred to as *Clustering*
- Falls into unsupervised learning (i.e. no clearly defined outcome of interest)
- Our goal is to find homogeneous subgroups among the observations



Clustering



There are many types of clustering algorithms.

We will cover three:

- K-means clustering
- Hierarchical clustering
- Expectation maximization algorithm
 - ► Beyond scope of our class



Clusters all observations into K clusters

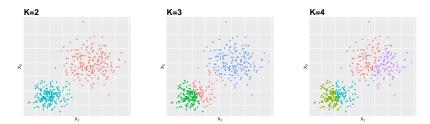
- ► K must be specified a-priori
- ▶ Algorithm then assigns every point to one of the *K* clusters
- ▶ Object is to minimize the *within-cluster variation*, i.e.

K-means clustering

$$\min_{C_1, C_2, ..., C_k} \sum_{\ell=1}^K W(C_\ell) : W(C_\ell) = \frac{1}{|C_\ell|} \sum_{i, j \in C_\ell} \mathcal{D}^2(\mathbf{x}_i, \mathbf{x}_j)$$

 $\mathcal{D}(\mathbf{x},\mathbf{y})$ measures the distance between \mathbf{x} and \mathbf{y} (typically the Euclidean distance, i.e. $\sqrt{\sum_{j=1}^p (x_j-y_j)^2}$).





Results from applying K-means clustering with different K's.



Algorithm steps

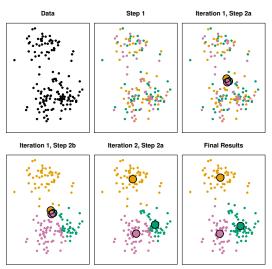
K-means clustering

- 1. Assign each observation (randomly) to one of the *K* clusters.
- 2. Iterate the 2 following steps until cluster assignments stop changing:
 - a Find the centroid of each of the K clusters

$$\bar{\mathbf{x}}_{\ell} = \frac{1}{|C_{\ell}|} \sum_{i \in C_{\ell}} \mathbf{x}_{i} \tag{11}$$

b Reassign each sample to the nearest centroid (using $\mathcal{D}^2(\mathbf{x}, \mathbf{y})$)





Visualization of k-means at different steps.



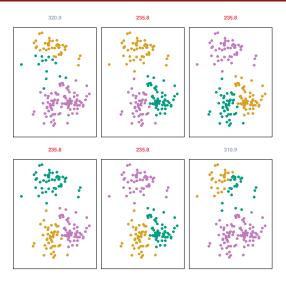
Some properties of K-means

▶ The algorithm always converges to a local minimum of

$$\min_{C_1, C_2, \dots, C_k} \left\{ \sum_{\ell=1}^K \frac{1}{|C_\ell|} \sum_{i, j \in C_\ell} \mathcal{D}^2(\mathbf{x}_i, \mathbf{x}_j) \right\}$$
(12)

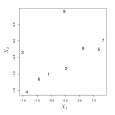
- The algorithm is random
 - ▶ Each initialization can result in a different minimum
 - Can run with with multiple initializations and select lowest minimum



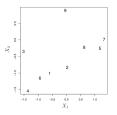


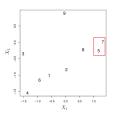
Example of running K-means 6 different times (K = 3).



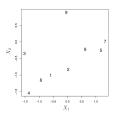


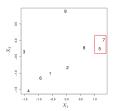


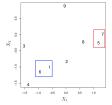




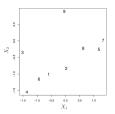


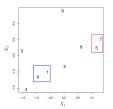


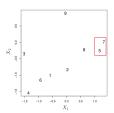


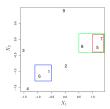






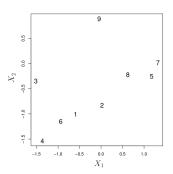


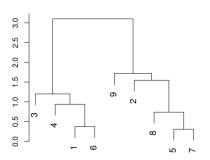






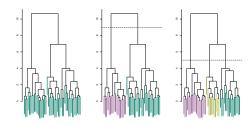
- ▶ The algorithm results in a dendogram
- ► *Hierarchical* in the sense that lower clusters are nested within higher clusters







- ► The number of clusters does not need to be specified a-priori
- Clusters created by cutting dendogram at a vertical point
- ▶ **Note**: Not all segmentation problems are nested clusters.
 - e.g. Market segmentation for consumers of 2 genders from 3 different nationalities.
 - Wierd to divide into 2 groups, and then to further divide 1 in half





In each iteration, we fuse the 2 clusters *closest* to each other.

► While we can use the Euclidean distance, what if a cluster has multiple observations?



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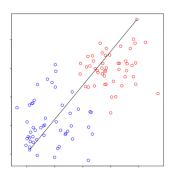
► While we can use the Euclidean distance, what if a cluster has multiple observations?

Linkage defines the dissimilarity between two clusters

Four primary types:

- 1. Complete
- 2. Average
- 3. Single
- 4. Centroid

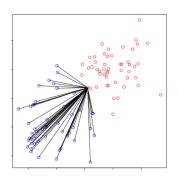




Complete linkage:

► The distance between 2 clusters is the maximum distance between any pair of samples, one in each cluster.

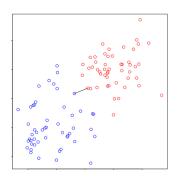




Average linkage:

► The distance between 2 clusters is the average of all pairwise distances.

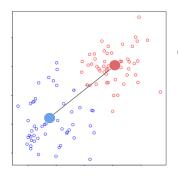




Single linkage:

- The distance between 2 clusters is the minimum distance between any pair of samples, one in each cluster.
- ► Suffers from chaining phenomenon

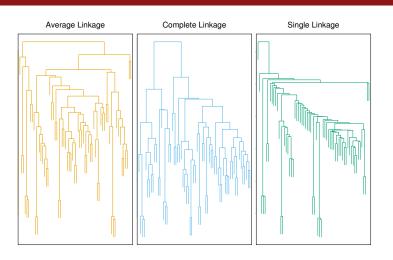




Centroid linkage:

- ► The distance between 2 clusters is the distance between each centroid.
- ► Suffers from inversions





 ${\sf Examples} \ of \ hierarchical \ clustering \ using \ different \ linkages.$



Clustering is riddled with questions and choices

- ► Is clustering appropriate? i.e. Could a sample belong to more than one cluster?
 - ▶ Mixture models, soft clustering, topic models.



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- How many clusters are appropriate?
 - Choose subjectively depends on the inference sought.
 - Some formal methods based on gap statistics, mixture models, etc.



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- How many clusters are appropriate?
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 - Some formal methods based on gap statistics, mixture models, etc.
- Are the clusters robust?
 - ► Run the clustering on different random subsets of the data. Is the structure preserved?
 - Try different clustering algorithms. Are the conclusions consistent?
 - Most important: temper your conclusions.

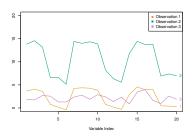


Questions on distance

- ▶ Should we scale the variables before doing the clustering.
 - Variables with larger variance have a larger effect on the Euclidean distance between two samples.
- Does Euclidean distance capture dissimilarity between samples?

Correlation distance



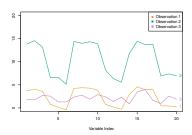


Example: Suppose that we want to cluster customers at a store for market segmentation.

- ► Samples are customers
- ► Each variable corresponds to a specific product and measures the number of items bought by the customer during a year.

Correlation distance



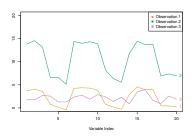


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- ▶ We **could** use Euclidean distance
 - Would cluster all customers who purchase few things (orange and purple)

Correlation distance





Example: Suppose that we want to cluster customers at a store for market segmentation.

- ▶ We **could** use Euclidean distance
 - Would cluster all customers who purchase few things (orange and purple)
- ▶ What if: we want to cluster customers who purchase similar things?
 - ▶ Correlation distance may be a more appropriate measure

References



- [1] ISL. Chapters 2.2.3, 10.3
- [2] ESL. Chapter 6.6.3