# Lecture 4 Big Data Predictive Analytics

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## Agenda

- Introduction
- K-Nearest Neighbor
- Applying KNN on Big Data
- Naïve Bayes classifier
- Applying Naïve Bayes classifier on Big Data
- Performance Evaluation of classifiers



➤ Predictive Analytics use Predictive Data Mining techniques which use Supervised Machine Learning techniques.

Predictive Analytics Predictive Use Supervised Machine Learning

- Classification is an instance of Supervised Machine Learning and is widely used for prediction purposes.
- In classification, a classifier is given a set of examples that are already classified (i.e. given a class label), and from these examples, the classifier learns to assign a label to unseen examples.
- Examples of classification problems include:
  - Given an email, classify if it is spam or not.
  - Given a handwritten character, classify it as one of the known characters.

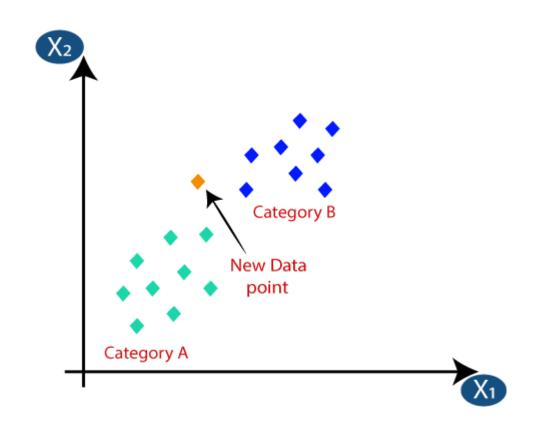
- >Examples of Classification Techniques:
  - K-Nearest Neighbor (KNN)
  - Naïve Bayes
  - Decision Trees (DT)
  - Support Vector Machines (SVM)
  - Neural Networks

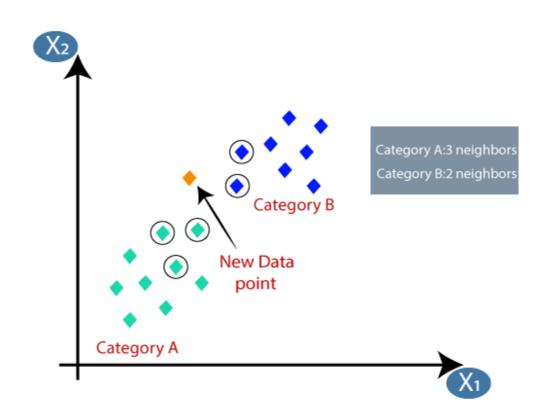


K-nearest neighbors is an algorithm that stores all available cases and classifies new cases based on a **similarity measure** (e.g., distance functions).

KNN algorithm at the training phase just stores the dataset and when it gets new data, then it classifies that data into a category that is much similar to the new data.

- The K-NN working can be explained on the basis of the below algorithm:
- Step 1 For implementing any algorithm, we need dataset. So during the first step of KNN, we must load the training as well as test data.
- Step 2 Next, we need to choose the value of K i.e. the nearest data points. K can be any integer.
- **Step 3** For each point in the test data do the following
  - 3.1 Calculate the distance between test data and each row of training data with the help of any of the method namely: Euclidean, Manhattan or Hamming distance.
  - 3.2 Now, based on the distance value, sort them in ascending order.
  - **3.3** − Next, it will choose the top K rows from the sorted array.
  - 3.4 Now, it will assign a class to the test point based on most frequent class of these rows.





- **▶** Distance functions:
- 1. Euclidean Distance:

$$D = \sqrt{\sum_{i=1}^{n} (x_i - y_i)^2}$$

n is the number of features

2. Manhattan Distance:

$$D = \sum_{i=1}^{n} |x_i - y_i|$$

n is the number of features

#### > Distance functions:

- 3. Hamming Distance:
  - It is a measure of the number of instances in which corresponding symbols are different in two strings of equal length. It is suitable for categorial features.

$$D_{H} = \sum_{i=1}^{n} |x_{i} - y_{i}|$$
$$x = y \Rightarrow D = 0$$

 $x \neq y \Rightarrow D = 1$ 

n is the number of features

#### >Example:

Height (in cms) Weight (in kgs) T Shirt Size					
158	58	М			
158	59	М			
158	63	М			
160	59	М			
160	60	M			
163	60	M			
163	61	M			
160	64	L			
163	64	L			
165	61	L			
165	62	L			
165	65	L			

New customer has height 161cm and weight 61kg.

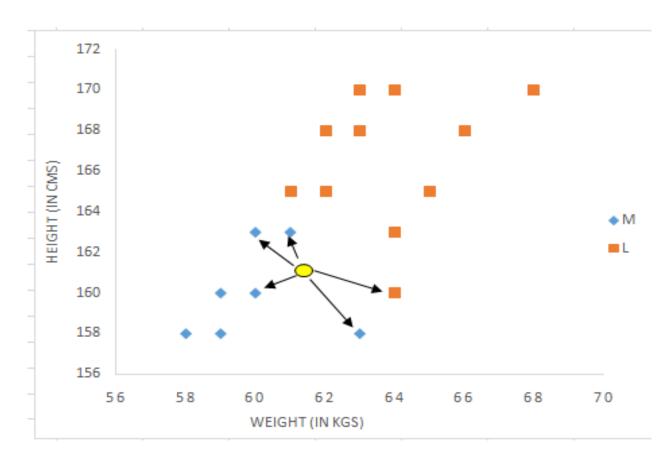
What is his T Shirt Size?

### >Example:

Height (in cms)	Weight (in kgs)	T Shirt Size	Distance	
158	58	M	4.2	
158	59	M	3.6	
158	63	M	3.6	
160	59	М	2.2	3
160	60	М	1.4	1
163	60	М	2.2	3
163	61	М	2.0	2
160	64	L	3.2	5
163	64	L	3.6	
165	61	L	4.0	
165	62	L	4.1	
165	65	L	5.7	

Euclidean Distance is used. For K=5, T shirt Size=M

>Example:



What are the limitations of KNN?

#### Normalization and Standardization:

- When independent variables in training data are measured in different units, it is important to **scale the variables** before calculating distance.
- For example, if one variable is based on height in cms, and the other is based on weight in kgs then **height will influence more** on the distance calculation.
- Scaling the variables can be done by any of the following methods:

$$Xs = \frac{X - min}{\max - min}$$

$$Xs = \frac{X - mean}{s.d.}$$

Normalization

Standardization

- > Handling categorial features:
  - Hamming Distance can be used
  - Can assign a number to each category

#### ➤ How to find best K value?

- **Cross-validation** is a way to find out the **optimal K value**. It estimates the validation error rate by holding out a subset of the training set from the model building process.
- Cross-validation (let's say 10 fold validation) involves randomly dividing the **training set** into 10 groups, or folds, of approximately equal size. 90% data is used to train the model and remaining 10% to validate it. The error rate is then computed on the 10% validation data. This procedure repeats 10 times each time with a different fold. It results to 10 estimates of the validation error which are then averaged out.
- The process is repeated for different values of K. The value of K that yields the smallest average error is selected.



- Despite the promising results shown by the k-NN in a wide variety of problems, it lacks scalability to address Big datasets.
- The main problems found to deal with large-scale data are:
  - Runtime: The complexity of the traditional k-NN algorithm is O((n · D)), where n is the number of instances and D the number of features.
  - **Memory consumption:** For a rapid computation of the distances, the k-NN model may normally require to store the training data in memory. When TR is too big, it could easily exceed the available RAM memory.

These drawbacks motivate the use of Big Data techniques to distribute the processing of KNN over a cluster a nodes.

A MapReduce-based approach for k-Nearest neighbor classification can be applied.

This allows us to simultaneously classify large amounts of unseen cases (test examples) against a big (training) dataset.

- First, the training data will be divided into multiple splits.
- The map phase will determine the k-nearest neighbors in the different splits of the data.
- As a result of each map, the k nearest neighbors together with their computed distance values will be emitted to the reduce phase.

Afterwards, the **reduce phase** will compute the definitive neighbors from the list obtained in the map phase.

The reduce phase will determine which are the final k nearest neighbors from the list provided by the maps.

This parallel implementation provides the exact classification rate as the original k-NN model.



### Naïve Bayes

Naïve Bayes is a probabilistic classification method based on Bayes' theorem (or Bayes' law).

Bayes' theorem gives the relationship between the probabilities of two events and their conditional probabilities.

A naïve Bayes classifier assumes that the presence (or absence) of a particular feature of a class is unrelated to the presence (or absence) of other features.

### Naïve Bayes

The input variables are generally **categorical**, but variations of the algorithm can accept continuous variables.

There are also ways to convert continuous variables into categorical ones. This process is often referred to as the *discretization of continuous* variables.

The output typically includes a class label and its corresponding probability score.

### Naïve Bayes: Bayes' Theorem

The *conditional probability* of event C occurring, given that event A has already occurred, is denoted as P(C|A), which can be found using the following equation:

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

where C is the class label C  $\in$  {c1, c2,...cn} and A is the observed attributes  $A = \{a1, a2, ...am\}$ 

### Naïve Bayes: Bayes' Theorem

A more general form of Bayes' theorem assigns a classified label to an object with multiple attributes  $A = \{a1, a2, ...am\}$  such that the label corresponds to the largest value of  $P(ci \mid A)$ .

$$P(c_{i}|A) = \frac{P(a_{1}, a_{2}, ..., a_{m}|c_{i}) \cdot P(c_{i})}{P(a_{1}, a_{2}, ..., a_{m})}, i = 1, 2, ... n$$

- With two simplifications, Bayes' theorem can be extended to become a naïve Bayes classifier.
- The first simplification is to use the conditional independence assumption. That is, each attribute is conditionally independent of every other attribute given a class label ci.
- The second simplification is to ignore the denominator P (a1, a2,..am) because it appears in the denominator for all values of i, removing the denominator will have no impact on the relative probability scores and will simplify calculations.

Naïve Bayes classification applies the two simplifications mentioned earlier and, as a result,  $P(ci \mid a1, a2,..am)$  is proportional to the product of  $P(aj \mid ci)$  times P(ci).

$$P(c_i|A) \propto P(c_i) \cdot \prod_{j=1}^m P(a_j|c_i)$$
  $i = 1, 2, ...n$ 

- Building a naïve Bayes classifier requires knowing certain statistics, all calculated from the training set.
- The first requirement is to collect the probabilities of all class labels, P(ci).
- The second thing the naïve Bayes classifier needs to know is the conditional probabilities of each attribute aj given each class label ci, namely  $P(aj \mid ci)$ . For each attribute and its possible values, computing the conditional probabilities given each class label is required.

- For a given attribute assume it can have the following values  $\{x,y,z\}$  and assume that we have two class labels  $\{c1 \text{ and } c2\}$ .
- Then the following probabilities need to be computed:
  - P(x | c1)
  - $P(x \mid c2)$
  - P(y | c1)
  - P(y | c2)
  - P(z | c1)
  - P(z | c2)

- After that, the naïve Bayes classifier can be tested over the testing set.
- For each record in the testing set, the naïve Bayes classifier assigns the classifier label ci that maximizes:

$$P(c_i) \cdot \prod_{j=1}^{m} P(a_j | c_i)$$

#### where:

- m is the number of features (dimensions)
- i is the index of class labels
- j is the index of features (dimensions)
- a1 is the value of the first feature in the test record
- a2 is the value of the second feature in the test record......



# Applying Naïve Bayes classifier on Big Data

# Applying Naïve Bayes classifier on Big Data

- Applying MapReduce with Naïve Bayes classifier significantly decreases computation times allowing its application on Big Data problems.
- First, the training data will be divided into multiple splits.
- During the map phase, each map processes a single split, and computes statistics of the input data.
- For each attribute, the map outputs a <Key, Value> pair, where
  - Key is the class label,
  - Value is {AttributeValue, the frequency of attribute value within that class label}

## Applying Naïve Bayes classifier on Big Data

- In the **reduce phase**, the reduce function aggregates the number of each attribute value within each class label.
- For each attribute, the reduce function outputs a <Key, Value> pair, where:
  - Key is the class label,
  - Value is {AttributeValue,  $\Sigma$ i the frequency of attribute value within that class label}
    - where i is the number of mappers.



- A *confusion matrix* is a specific table layout that allows visualization of the performance of a classifier.
- The following figure shows the confusion matrix for a two-class classifier:

		Predicted Class	
		Positive	Negative
	Positive	True Positives (TP)	False Negatives (FN)
Actual Class	Negative	False Positives (FP)	True Negatives (TN)

- > True positives (TP) are the number of positive instances the classifier correctly identified as positive.
- False positives (FP) are the number of instances in which the classifier identified as positive but in reality are negative.
- > True negatives (TN) are the number of negative instances the classifier correctly identified as negative.
- False negatives (FN) are the number of instances classified as negative but in reality are positive.

- The *accuracy* (or the *overall success rate*) is a metric defining the rate at which a model has classified the records correctly.
- It is defined as the sum of TP and TN divided by the total number of instances, as shown in the following equation:

$$Accuracy = \frac{TP + TN}{TP + TN + FP + FN} \times 100\%$$

- The *false positive rate* (FPR) shows what percent of negatives the classifier marked as positive.
- The FPR is also called the *false alarm rate* or the *type I error rate*.
- >FPR is computed as follows:

$$FPR = \frac{FP}{FP + TN}$$

➤ *Precision* is the percentage of instances marked positive that really are positive. It is computed as follows:

$$Precision = \frac{TP}{TP + FP}$$

➤ **Recall** is the percentage of positive instances that were correctly identified. It is also called **true positive rate** (TPR). It is computed as follows:

$$TPR (or Recall) = \frac{IP}{TP + FN}$$

> *F1-score* is the harmonic mean of the precision and recall:

$$F_1 = 2 \cdot rac{ ext{precision} \cdot ext{recall}}{ ext{precision} + ext{recall}}$$

- ➤ Micro-average and Macro-average:
  - The micro-average precision and recall scores are calculated from the sum of classes' true positives (TPs), false positives (FPs), and false negatives (FNs) of the model.
  - The macro-average precision and recall scores are calculated as arithmetic mean (or weighted mean) of individual classes' precision and recall scores.
  - The macro-average F1-score is calculated as arithmetic mean (or weighted mean) of individual classes' F1-score.

>Exercise:

No	Actual	Predicted	Match
1	Airplane	Airplane	<b>√</b>
2	Car	Boat	X
3	Car	Car	✓
4	Car	Car	✓
5	Car	Boat	X
6	Airplane	Boat	X
7	Boat	Boat	✓
8	Car	Airplane	X
9	Airplane	Airplane	✓
10	Car	Car	✓

#### >Exercise:

	Airplane	Boat	Car
Airplane	2	1	0
Boat	0	1	0
Car	1	2	3

	Label	True Positive (TP)	False Positive (FP)	False Negative (FN)
	Airplane	2	1	1
	Boat	1	3	0
1	Car	3	0	3

**Confusion matrix** 

TP, FP, FN calculated from the confusion matrix

#### >Exercise:

• Calculate micro-average, macro-average, and weighted macro-average of precision, recall and F1-score.

