# Data Mining & Informatics Lecture #3

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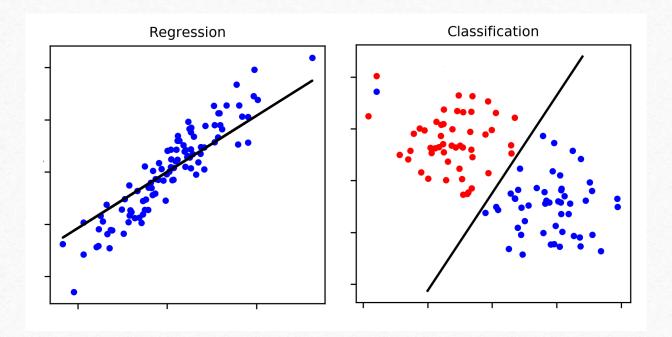
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#### NOTE

- Extra lessons
- Extra lessons will be on Tuesday 5-7 pm
- The first class will be next week and will be dedicated to Python basics

#### Classification

- Introduction
- There are two main supervised learning methods: Classification and Regression



#### Introduction

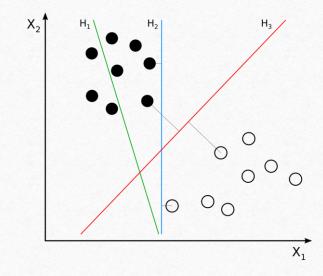
- In all supervised learning methods there are 3 steps:
- Defining the model: In this step, the model is defined and hyperparameters are set.
- Fitting the model (training): In this step, the training data are used to find the optimum model parameters using a learning algorithm (for example gradient descent in neural network)
- Testing the model (prediction): after setting the model hyperparameter and finding the model's parameter in the training state, in this step, test data are used to check how accurate the model's predictions are.

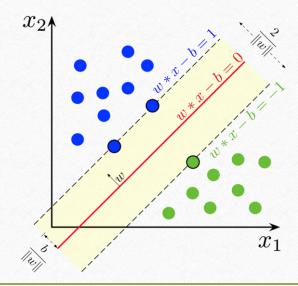
#### Classification

- Classification is a supervised learning method in which the target variables (the variables that need to be predicted) are categorical.
- If there are only two categories, it is called binary classification. If there are more than two categories, it is called multiclass classification.
- There are different classification methods. Here we discuss support vector machine (SVM), decision tree (DT), random forest (RF), and K-nearest neighbor (KNN) classifiers.

#### Classification - SVM

- The objective is to find a hyperplane in an n-dimensional space that separates the data points to their potential classes. The hyperplane should be positioned with the maximum distance to the data points
- The data points with the minimum distance to the hyperplane are called Support Vectors

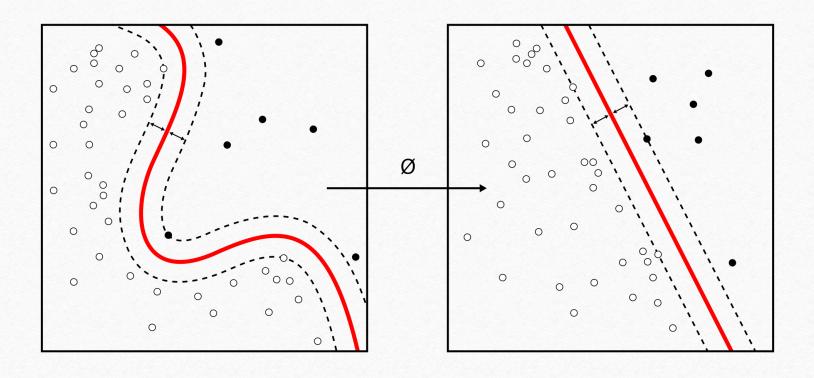




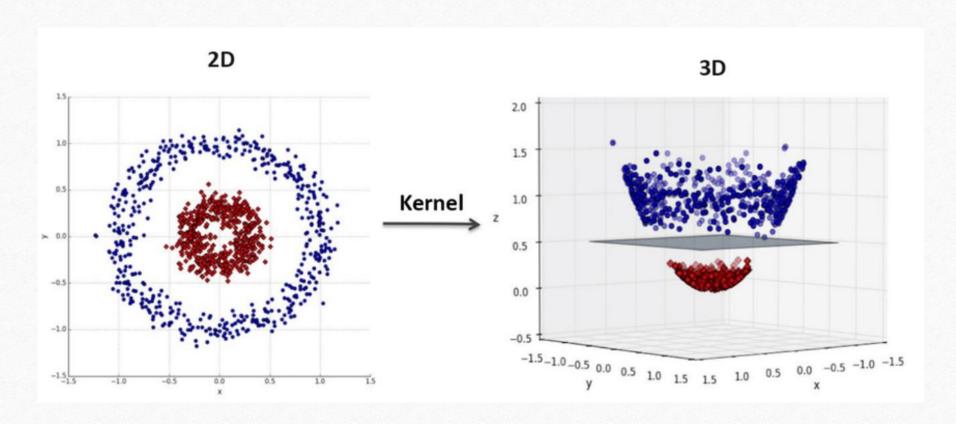
#### Classification - SVM

- The original maximum-margin hyperplane algorithm proposed by Vapnik in 1963 constructed a linear classifier that works when data are linearly separable (using a line or hyperplane)
- For data that are not linearly separable, we need to use a kernel function and transform data to another space where they are linearly separable. This is called the kernel trick.
- There are different kernel functions including linear function, Polynomial function, Radial basis function (RBF), and Sigmoid function.

• Classification - SVM



#### • Classification - SVM



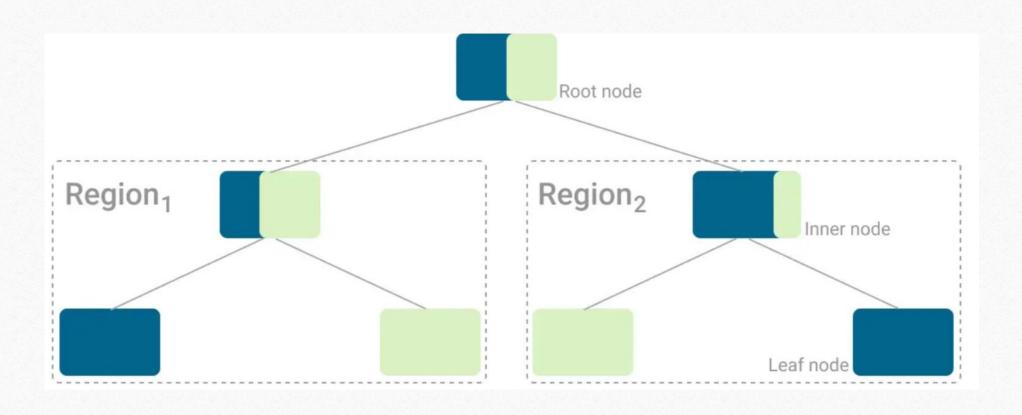
#### Classification - SVM

- There are different kernel functions, including linear function, Polynomial function, Radial basis function (RBF), and Sigmoid function.
- The kernel type is a hyperparameter and should be selected using a trial and error procedure.
- The SVM classification is used for binary classification by definition.

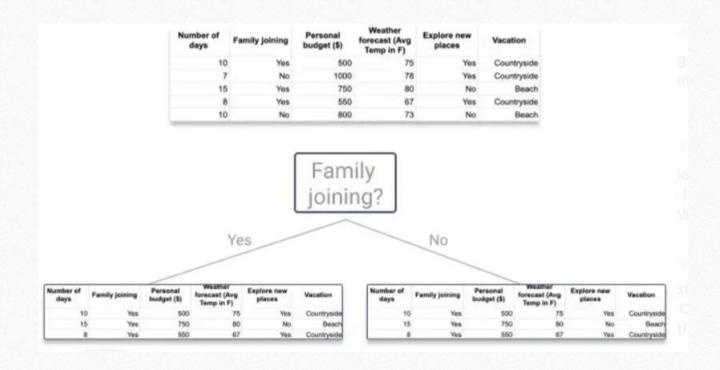
#### Classification – Decision tree

- Decision Tree is a Supervised Machine Learning Algorithm that uses a set of rules to make decisions, similar to how humans make decisions.
- The intuition behind Decision Trees is that you use the dataset features to create yes/no questions and continually split the dataset until you isolate all data points belonging to each class.
- Every time you ask a question you're adding a **node** to the tree. And the first node is called the **root node**. Each node is connected to other nodes using **branches**.
- If you decide to stop the process after a split, the last nodes created are called **leaf** nodes.
- Every time you answer a question, you're also creating branches and segmenting the feature space into disjoint **regions**.

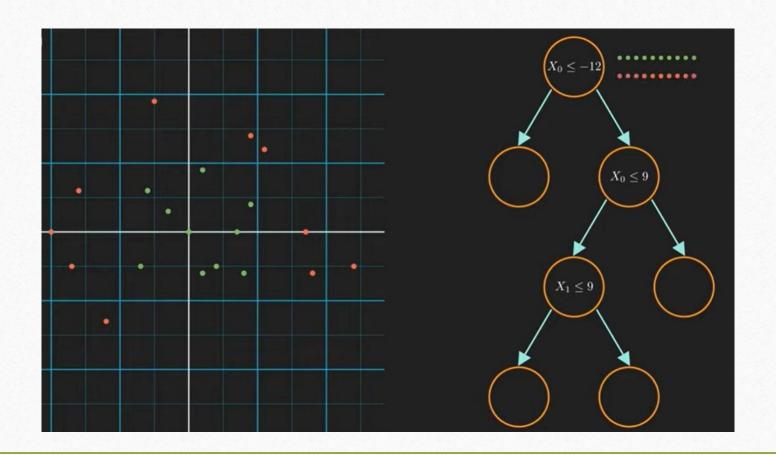
• Classification – Decision tree



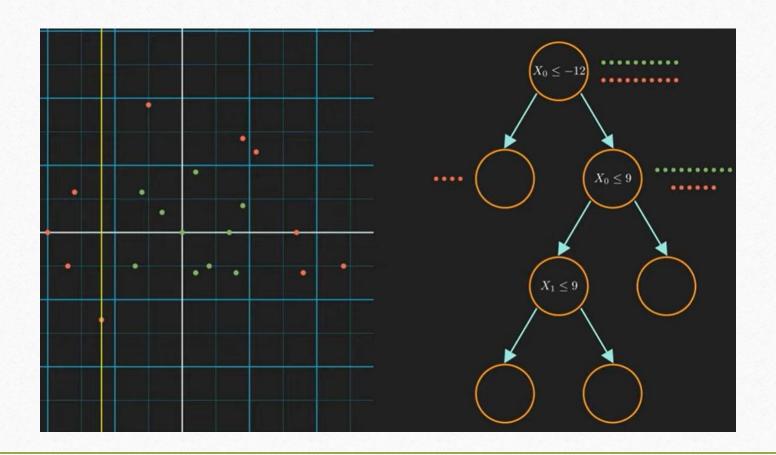
Classification – Decision tree



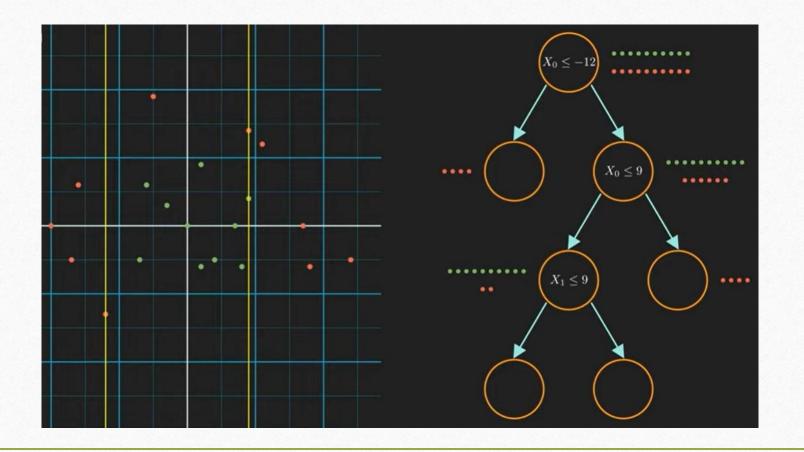
• Classification – Decision tree



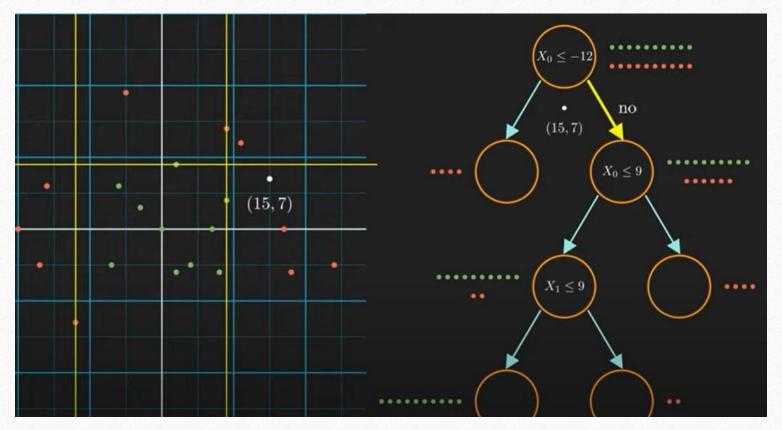
• Classification – Decision tree



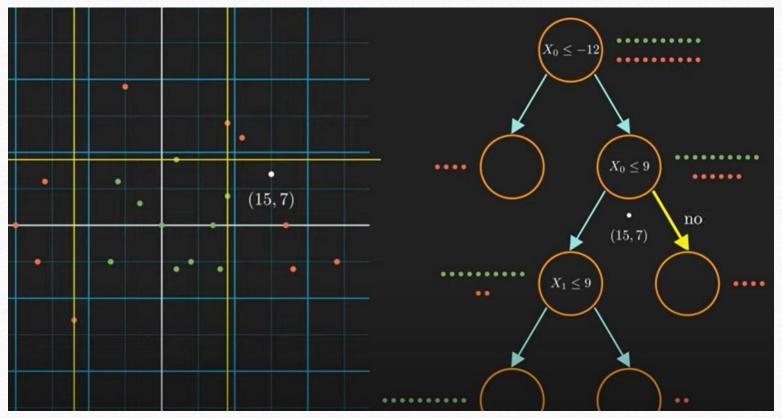
• Classification – Decision tree



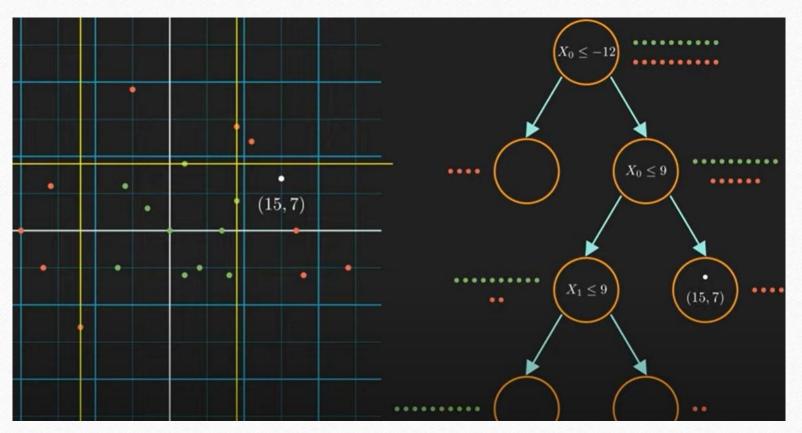
Classification – Decision tree



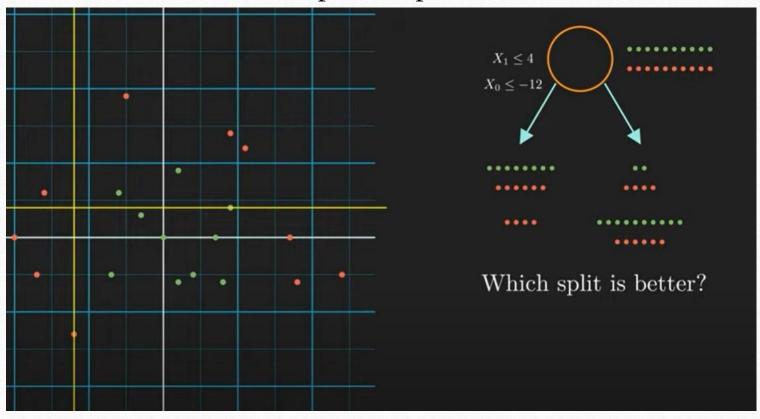
Classification – Decision tree



• Classification – Decision tree



- Classification Decision tree
- But how can we determine the optimal split and condition?



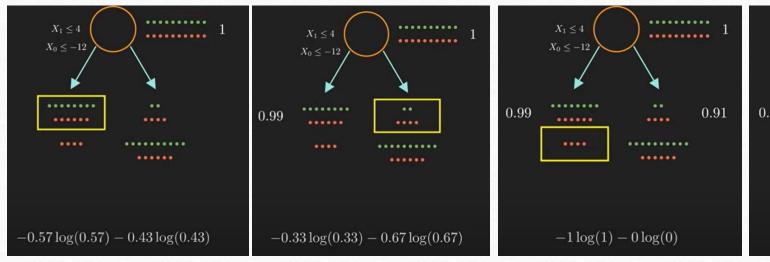
- Classification Decision tree
- We can decide the best split by calculating the information gain resulting from each split using the following formula (it is  $log_2$ ):

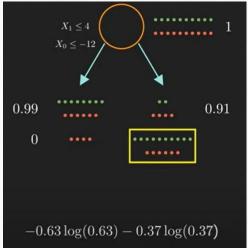
$$Entropy = \sum_{i} - p_i \log(p_i)$$

$$p_i = \text{probability of class i}$$

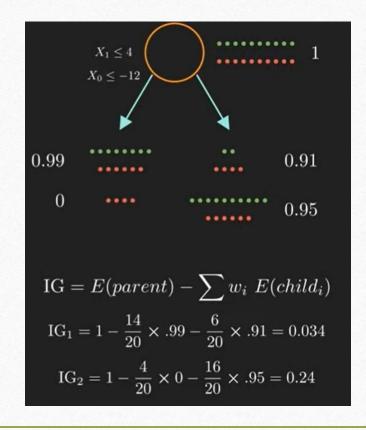
$$IG = E(parent) - \sum w_i \ E(child_i)$$

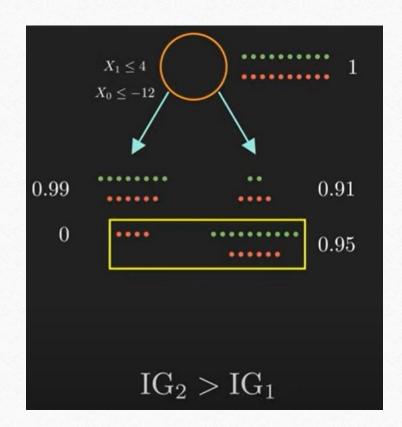
Classification – Decision tree





#### Classification – Decision tree

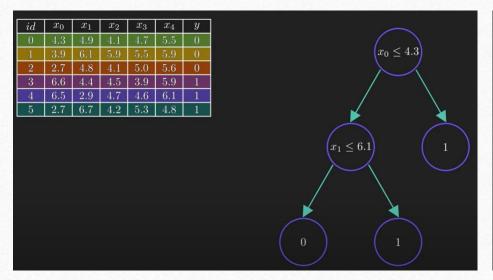


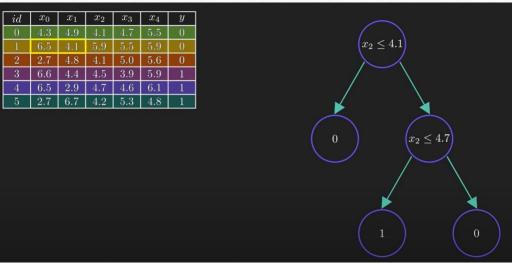


#### Classification – Decision tree

- In reality, the model traverses through every possible feature and feature value to find the best split.
- A pure leaf node is a leaf node that contains data from only one class.
- Although it's desirable to have data from only one class in each leaf node, this will not happen in most cases. If we have data from different classes in leaf nodes, we can use the **majority voting** technique to decide which class the leaf node belongs to.
- According to the majority voting, the class of the leaf node is the same class as the class of the majority of data inside the leaf.

- Classification Random Forest
- Decision trees are highly sensitive to the training data resulting in a high variance
- This problem will limit the generalization of the trained model





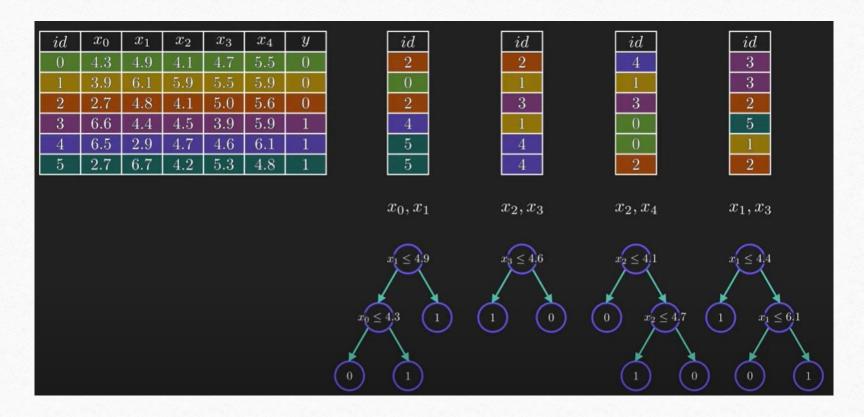
- Classification Random Forest
- Random forest is a collection of multiple random decision trees
- Random forest algorithm can be summarized as follows:
- **Step1**: Choose random subsets of the original dataset with replacement so that the total number of rows in the selected subset is the same as the original dataset. This process is called bootstrapping.

						15-1	5					
id	$x_0$	$x_1$	$x_2$	$x_3$	$x_4$	y	id	id	id	id		
0	4.3	4.9	4.1	4.7	5.5	0	2	2	4	3		
1	3.9	6.1	5.9	5.5	5.9	0	0	1	1	3		
2	2.7	4.8	4.1	5.0	5.6	0	2	3	3	2		
3	6.6	4.4	4.5	3.9	5.9	1	4	1	0	5		
4	6.5	2.9	4.7	4.6	6.1	1	5	4	0	1		
5	2.7	6.7	4.2	5.3	4.8	1	5	4	2	2		
								Bootstrapped Datasets				

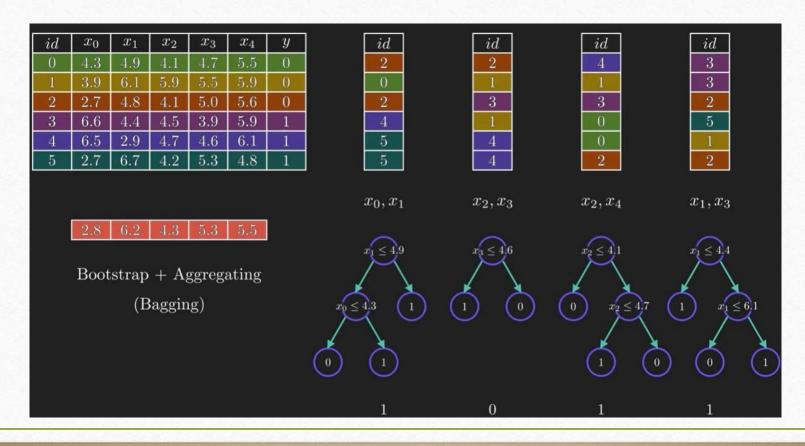
- Classification Random Forest
- NOTE: The number of subsets is a hyperparameter. Here we chose four subsets. However, it is usually set to 100.
- **Step2**: Randomly select a subset of features for each dataset. The number of selected features is a hyperparameter usually set to the square root or log of the total number of features.

id	$x_0$	$x_1$	$x_2$	$x_3$	$x_4$	y	$\lceil id \rceil$	id	$\lceil id \rceil$	id
0	4.3	4.9	4.1	4.7	5.5	0	2	2	4	3
1	3.9	6.1	5.9	5.5	5.9	0	0	1	1	3
2	2.7	4.8	4.1	5.0	5.6	0	2	3	3	2
3	6.6	4.4	4.5	3.9	5.9	1	4	1	0	5
4	6.5	2.9	4.7	4.6	6.1	1	5	4	0	1
5	2.7	6.7	4.2	5.3	4.8	1	5	4	2	2
							$x_0, x_1$	$x_2, x_3$	$x_2, x_4$	$x_1, x_3$

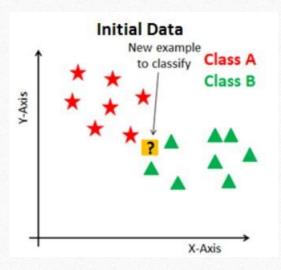
- Classification Random Forest
- Step3: Train one tree for each selected dataset and set of features.



- Classification Random Forest
- Step4: For new unseen data, make predictions using the majority voting

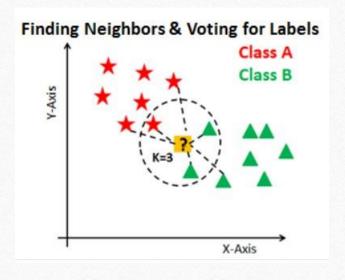


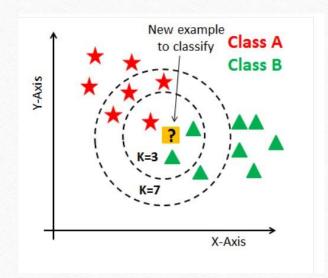
- Classification K-nearest neighbors
- Suppose P1 is the point, for which label needs to predict. First, you find the k closest point to P1 and then classify points by majority vote of its k neighbors. Each object votes for their class and the class with the most votes is taken as the prediction



- Classification K-nearest neighbors (KNN)
- For finding closest similar points, you find the distance between points using distance measures such as Euclidean distance, Hamming distance, Manhattan distance and Minkowski distance.
- KNN algorithm has the following three steps:
- **Step1**: Calculate the distance between the new data point and K neighboring data
- Step2: Find closest neighbors
- Step3: Use majority voting to predict the class of the new data

• Classification – K-nearest neighbors (KNN)





**NOTE**: K is a hyperparameter and is usually set to the square root of the total number of samples. You may get different results using different K.

Classification – K-nearest neighbors (KNN)

NOTE: KNN and RF can be used for multiclass classification

**NOTE**: There are also KNN and RF regression algorithms that use similar procedures to the classifiers but are used for regression.

- Classification Evaluation metrics
- Evaluation metrics are used to measure how good the predictions of the system are
- The most important evaluation metrics for classification include Accuracy, Precision, Recall, F1 Score, Specificity, and Sensitivity

#### • Classification – Evaluation metrics

#### **Evaluation Matrix**

#### **Confusion Matrix**

	<b>Actual Positive</b>	Actual Negative
	(Apple)	(Non-apple)
<b>Predicted Positive</b>	TP	FP
(Apple)		
<b>Predicted Negative</b>	FN	TN
(Non-apple)		

TP: true positive FP: false positive FN: false negative TN: true negative

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

**NOTE:** Accuracy is NOT always a good measure!

### • Classification – Evaluation metrics

#### **Confusion Matrix**

**Total fruits: 20** 

	<b>Actual Positive</b>	<b>Actual Negative</b>
	(Apple)	(Non-apple)
<b>Predicted Positive</b>	10	5
(Apple)	TP	FP
<b>Predicted Negative</b>	2	3
(Non-apple)	FN	TN

TP: true positive

FN: false negative

FP: false positive

TN: true negative

#### • Classification – Evaluation metrics

**TP**: The number of positive samples that are predicted correctly as positive (**T**ruly classified as **P**ositive)

**TN**: The number of negative samples that are predicted correctly as negative (**T**ruly classified as **N**egative)

**FP**: The number of negative samples that are predicted incorrectly as positive (**F**alsely classified as **P**ositive)

**FN**: The number of positive samples that are predicted incorrectly as negative (Falsely classified as Negative)

### • Classification – Evaluation metrics

#### Limitation

#### Suppose

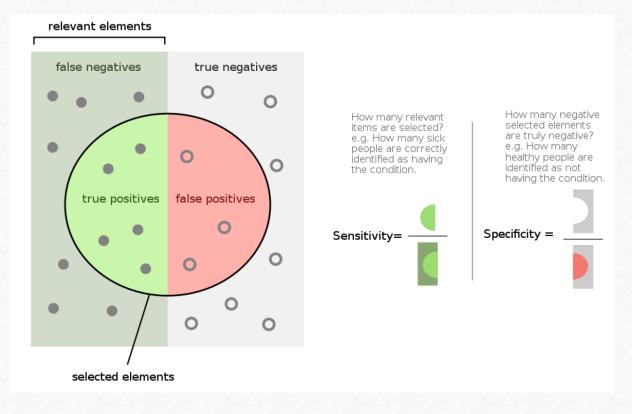
- Total number of fruits in the testing examples = 10,000
- Number of Non-apple = 9990
- Number of Apple = 10

#### Can you classify Apple?

If model predicts everything to be of class non-apple, the accuracy is 9990/10000 = 99.9 %!!!

Here, accuracy is misleading because model cannot detect any Apple at all, still achieving high accuracy.

Classification – Evaluation metrics



Sensitivity and specificity. (2024, January 24). In Wikipedia. https://en.wikipedia.org/wiki/Sensitivity\_and\_specificity

#### Classification – Evaluation metrics

sensitivity, recall, hit rate, or true positive rate (TPR)

$$TPR = \frac{TP}{P} = \frac{TP}{TP + FN}$$

specificity, selectivity or true negative rate (TNR)

$$ext{TNR} = rac{ ext{TN}}{ ext{N}} = rac{ ext{TN}}{ ext{TN} + ext{FP}}$$

precision or positive predictive value (PPV)

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

Sensitivity and specificity. (2024, January 24). In Wikipedia. https://en.wikipedia.org/wiki/Sensitivity\_and\_specificity

### • Classification – Evaluation metrics

#### **Evaluation Matrix**

	Actual Positive (Apple)	Actual Negative (Non-apple)
Predicted Positive (Apple)	TP	FP
Predicted Negative (Non-apple)	FN	TN

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

$$Recall = \frac{TP}{TP+FN}$$

$$F_1 = 2 * \frac{Precision * Recall}{Precision + Recall}$$

### • Classification – Evaluation metrics

**NOTE 1**: These metrics are defined for binary classification, but can be extended to multi-class classification

**NOTE 2**: It is optional to consider either of the classes as negative or positive. However, once you have defined a class as negative or positive, you should stick to your definition.

### • Classification – Evaluation metrics

**NOTE 1**: These metrics are defined for binary classification, but can be extended to multi-class classification

**NOTE 2**: It is optional to consider either of the classes as negative or positive. However, once you have defined a class as negative or positive, you should stick to your definition.

## • Evaluation metrics - example

**Question**: Assume we have 1000 fruits, and we want to classify them into two classes as follows: Apple and Non-apple. Assume we have 10 apples in the dataset, and the rest of the fruits are not apples. If the classifier classifies all fruits as Non-apple, calculate the evaluation metrics for this classifier.

Solution: We define the Non-apple class as positive and the Apple class as negative.

TP= 990 (How many Non-apple fruits are classified as Non-apple)

TN= 0 (How many Apple fruits are classified as Apple)

FP= 10 (How many Apple fruits are classified as Non-apple)

FN= 0 (How many Non-apple fruits are classified as Apple)

### • Evaluation metrics - example

Accuracy = (TP+TN)/(TP+TN+FP+FN)=(990+0)/(990+0+10+0)=0.99 (99%) Interpretation: The model is prediction 99% of cases correctly.

Sensitivity= TP/(TP+FN)=990/(990+0)=1 (100%)

Interpretation: The model is predicting 100% of Non-apple fruits correctly

Specificity= TN/(TN+FP)=0/(0+10)=0 (0%)

Interpretation: The model is predicting 0% of Apple fruits correctly

Precision= TP/(TP+FP)= 990/(990+10)=0.99 (99%)

Interpretation: 99% of predicted Non-apple fruits are correct

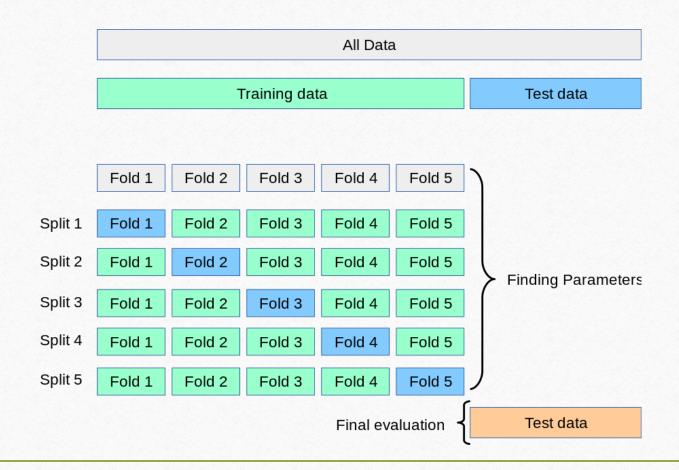
#### Evaluation – Cross validation

- In classification, we use a portion of data for training the model. This portion is called **training data**
- We use the rest of the data for resting the model. This portion is called **testing** data.
- There is a challenge. Which part of data should be used for training the model and which part should be used for testing?

#### Evaluation – Cross validation

- In the cross-validation (CV) method, we divide data into K folds. Then in each iteration, one of the folds is reserved for training the model and one portion is reserved for testing the model.
- We repeat this process until all folds have been used for testing at least once.
- This procedure is also called K-fold cross-validation.
- The accuracy of the model will be the average accuracy of all iterations.
- The best practice is to test the model on separate test data after the cross-validation

• Evaluation – Cross validation



### • Examples

Please see Lecture #3-examples notebook for examples of classification models.