# Decision trees Bagging

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## Recap

Lecture 4: SVM, PCA kNN indexes



- 1. Support Vector Machine (SVM)
  - Hinge loss
  - Kernel trick
- 2. Dimensionality reduction and PCA
  - Problem statement
  - Singular Value Decomposition
  - Eckart–Young theorem
  - Equivalent definitions
  - Data normalization
- 3. k Nearest Neighbors
  - o kNN indexes
  - > HNSW

## Outline

- 1. Intuition
- 2. Construction procedure
- 3. Information criteria
- 4. Special highlights
  - Dealing with missing data
  - Binarization
  - Decision tree as linear model
  - Standards
  - Hyperparameters
- 5. Bootstrap and Bagging
- 6. Random Forest



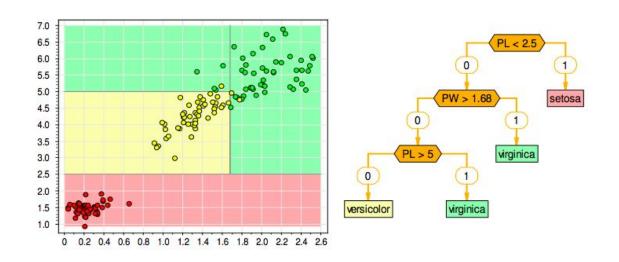
## Intuition

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#### **Decision tree for Iris data set**

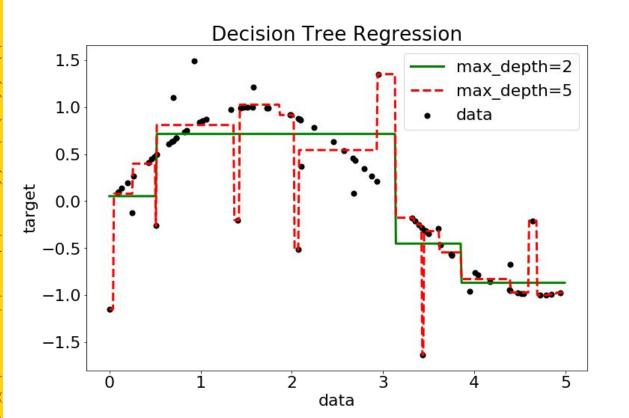




setosa
$$r_1(x) = [PL \leqslant 2.5]$$
virginica $r_2(x) = [PL > 2.5] \land [PW > 1.68]$ virginica $r_3(x) = [PL > 5] \land [PW \leqslant 1.68]$ versicolor $r_4(x) = [PL > 2.5] \land [PL \leqslant 5] \land [PW < 1.68]$ 

## **Decision tree in regression**





Green - decision tree of depth 2

Red - decision tree of depth 5

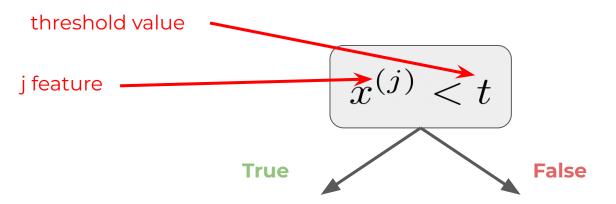
Every leaf corresponds to some constant.

# **Construction procedure**

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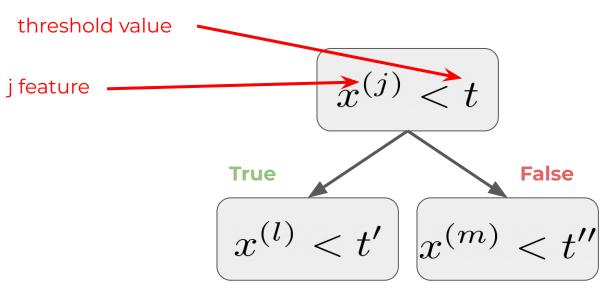






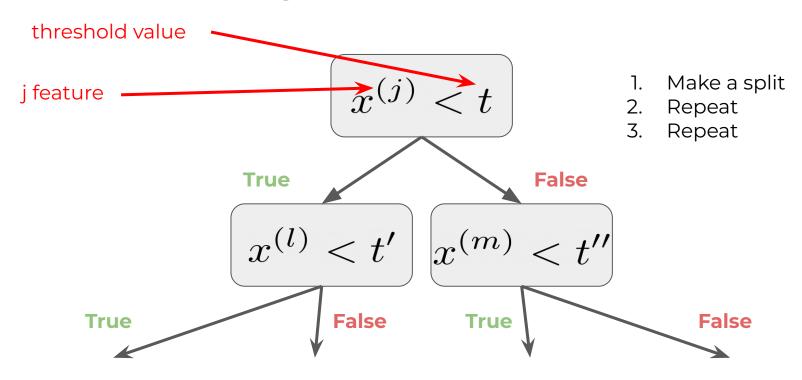
1. Make a split



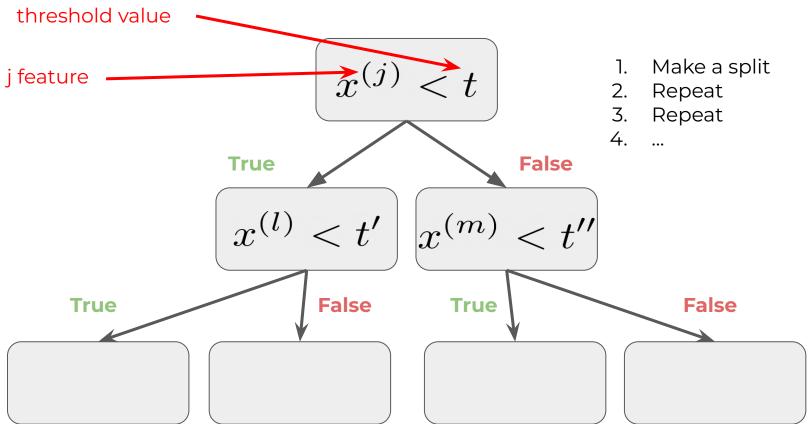


- 1. Make a split
- 2. Repeat











threshold value

i feature

# WHAT IF I TOLD YOU TO TELL ME THAT I SHOULD TELL YOU WHAT IF I TOLD YOU

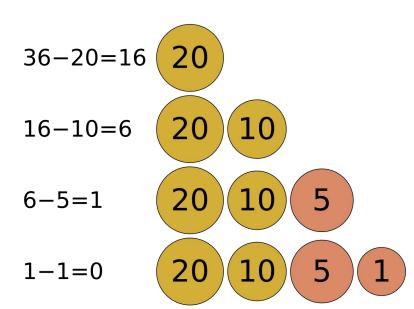
True

#### **Greedy algorithm**



A greedy algorithm is any algorithm that follows the problem-solving heuristic of making the locally optimal choice at each stage.

In many problems, a greedy strategy does not produce an optimal solution, but a greedy heuristic can yield locally optimal solutions that approximate a globally optimal solution in a reasonable amount of time.



#### How to answer in leaf?



#### Classification:

- most popular
- sample with frequencies of classes

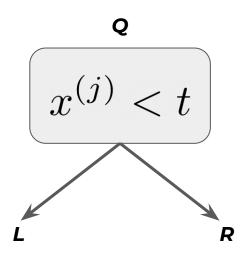
#### Regression:

Depends on loss function!

- for MSE
  - o average in node
- for MAE
  - o median in node

#### How to split data properly?



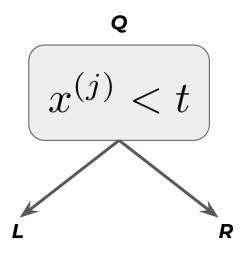


We can not use gradient this time because solution set is discrete.

So let's apply discrete optimization!

#### How to split data properly?





$$\frac{|L|}{|Q|}H(L) + \frac{|R|}{|Q|}H(R) \longrightarrow \min_{j,t}$$

#### How to choose concrete split?

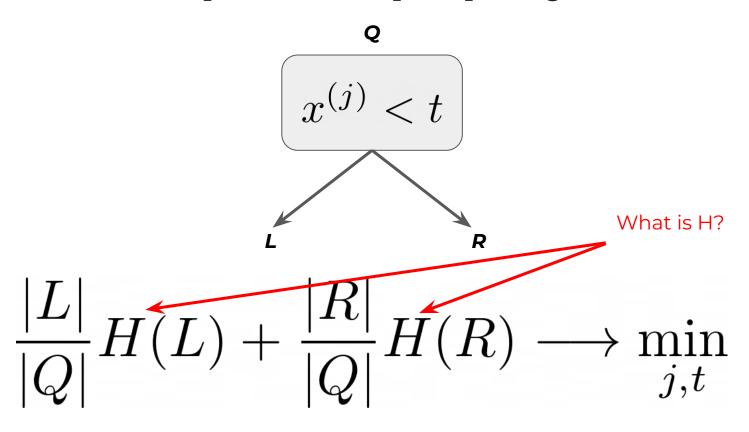


Brute force algorithm will take too much time.

Random splits are chosen and compared.

#### How to split data properly?





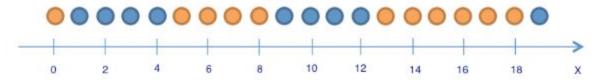
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H(R) is measure of "heterogeneity" of our data.

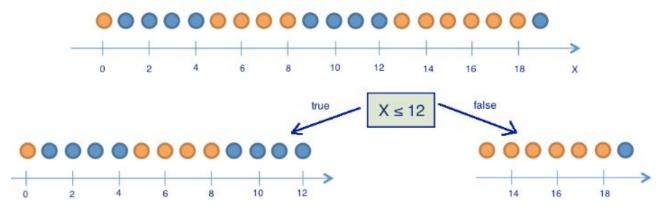
Consider binary classification problem:





H(R) is measure of "heterogeneity" of our data.

Consider binary classification problem:





H(R) is measure of "heterogeneity" of our data.

Consider binary classification problem:

Obvious way:

Misclassification criteria:

$$H(R) = 1 - \max\{p_0, p_1\}$$

1. Entropy criteria: 
$$H(R) = -p_0 \log p_0 - p_1 \log p_1$$

2. Gini impurity: 
$$H(R) = 1 - p_0^2 - p_1^2 = 2p_0p_1$$



H(R) is measure of "heterogeneity" of our data.

Consider multiclass classification problem:

Obvious way:

Misclassification criteria:

$$H(R) = 1 - \max_{k} \{p_k\}$$

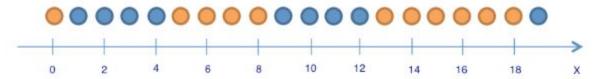
$$H(R) = -\sum_{k=0}^{\infty} p_k \log p_k$$

$$H(R) = 1 - \sum_{k} (p_k)^2$$



H(R) is measure of "heterogeneity" of our data.

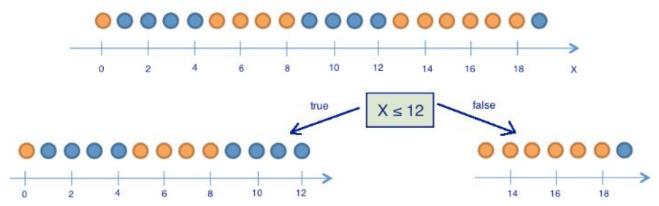
Consider binary classification problem:





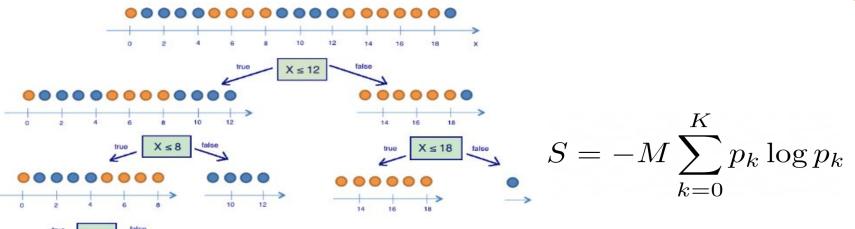
H(R) is measure of "heterogeneity" of our data.

Consider binary classification problem:



#### Information criteria: Entropy





In binary case N = 2

$$S = -p_{+} \log_{2} p_{+} - p_{-} \log_{2} p_{-} = -p_{+} \log_{2} p_{+} - (1 - p_{+}) \log_{2} (1 - p_{+})$$

## **Information criteria: Gini impurity**



$$G = 1 - \sum_{k} (p_k)^2$$

In binary case N = 2

$$G = 1 - p_+^2 - p_-^2 = 1 - p_+^2 - (1 - p_+)^2 = 2p_+(1 - p_+)$$



H(R) is measure of "heterogeneity" of our data.

Consider multiclass classification problem:

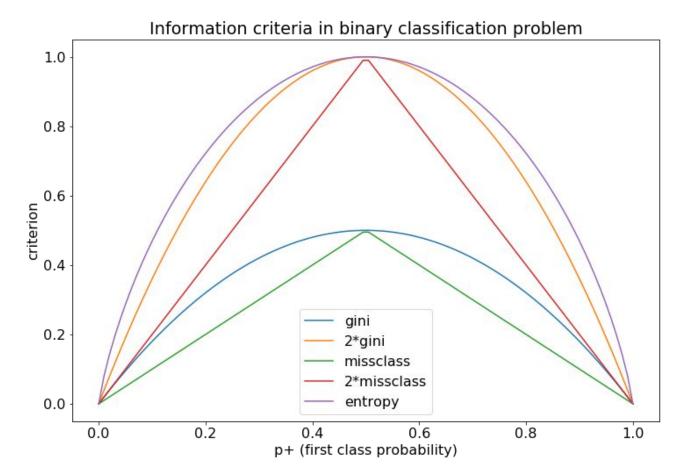
Obvious way: Misclassification criteria:

$$H(R) = 1 - \max_{k} \{p_k\}$$

1. Entropy criteria: 
$$H(R) = -\sum_k p_k \log_2 p_k$$

2. Gini impurity: 
$$H(R) = 1 - \sum_k (p_k)^2$$







H(R) is measure of "heterogeneity" of our data.

Consider regression problem:

1. Mean squared error

$$H(R) = \min_{c} \frac{1}{|R|} \sum_{(x_i, y_i) \in R} (y_i - c)^2$$

What is the constant?

$$c^* = \frac{1}{|R|} \sum_{y_i \in R} y_i$$

#### **Hyperparameters**

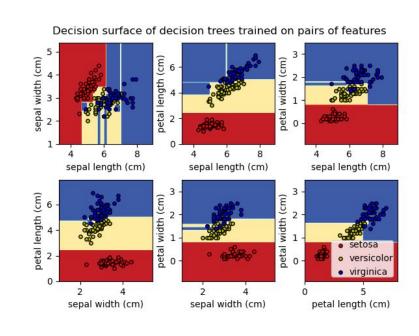


- max\_depth: min 1
- min\_samples\_split: min 2
- min\_samples\_leaf: min 1
- min\_impurity\_decrease

#### Minor

- criterion:
  - o gini, entropy, log\_loss for classification
  - MSE or MAE for regression
- splitter: best, random
- max\_features: sqrt, log2

As of sklearn implementation



#### **Standards**



- <u>ID-3</u>
  - Entropy criteria; Stops when no more gain available
- C4.5
  - Normalised entropy criteria; Stops depending on leaf size; Incorporates pruning
- C5.0
  - Some updates on C4.5
- CART
  - Gini criteria; Cost-complexity Pruning; Surrogate predicates for missing data;
- etc.

Read more

# Special highlights

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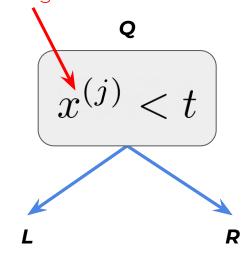
#### Missing values in Decision Trees



If the value is missing, one might use both sub-trees and average their predictions.

But this will negatively affect model computational performance.

#### Missing value



$$\hat{y} = \frac{|L|}{|Q|} \hat{y}_L + \frac{|R|}{|Q|} \hat{y}_R$$

#### Missing values in Catboost



Forbidden: Missing values are not supported, their presence is interpreted as an error

**Min**: Missing values are processed as the minimum value (less than all other values) for the feature. It is guaranteed that a split that separates missing values from all other values is considered when selecting trees.

**Max**: Missing values are processed as the maximum value (greater than all other values) for the feature. It is guaranteed that a split that separates missing values from all other values is considered when selecting trees.

The **default** processing mode **is Min** 

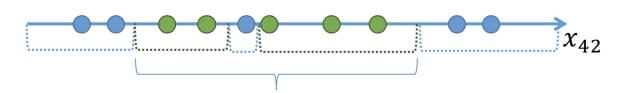
**Documentation** 

#### **Binarization**



Idea: instead selecting one threshold define several for one feature.





e.g. <u>Border count hyperparameter</u> in Catboost (defaults to 254)

#### **Decision Trees as Linear models**



Let J be the subspace of the original feature space, corresponding to the leaf of the tree.

Prediction takes form

$$\hat{y} = \sum_{j} w_j [x \in J_j]$$

# **Bootstrap and Bagging**

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### **Bootstrap**



Consider dataset X containing m objects.

Pick m objects with return from X and repeat in N times to get N datasets.

Error of model trained on Xj: 
$$\varepsilon_j(x) = b_j(x) - y(x), \qquad j = 1, \ldots, N,$$

Then 
$$\mathbb{E}_x(b_j(x)-y(x))^2=\mathbb{E}_x\varepsilon_j^2(x)$$
.

The mean error of N models: 
$$E_1=rac{1}{N}\sum_{j=1}^N \mathbb{E}_x arepsilon_j^2(x).$$

## **Bootstrap**



Consider the errors unbiased and uncorrelated:

$$\mathbb{E}_x \varepsilon_j(x) = 0;$$

$$\mathbb{E}_x \varepsilon_i(x) \varepsilon_j(x) = 0, \quad i \neq j.$$

$$E_N = \mathbb{E}_x \left( \frac{1}{N} \sum_{j=1}^n b_j(x) - y(x) \right)^2 =$$

The final model averages all predictions:

$$a(x) = \frac{1}{N} \sum_{i=1}^{N} b_j(x).$$

$$= \mathbb{E}_{x} \left( \frac{1}{N} \sum_{j=1}^{N} \varepsilon_{j}(x) \right)^{2} =$$

$$= \frac{1}{N^{2}} \mathbb{E}_{x} \left( \sum_{j=1}^{N} \varepsilon_{j}^{2}(x) + \sum_{i \neq j} \varepsilon_{i}(x) \varepsilon_{j}(x) \right) =$$

$$= 0$$

## **Bootstrap**



Consider the errors unbiased and unco

$$\mathbb{E}_x \varepsilon_j(x) = 0;$$

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The final model averages all predictions:

$$a(x) = \frac{1}{N} \sum_{j=1}^{N} b_j(x).$$

Error decreased by N times!

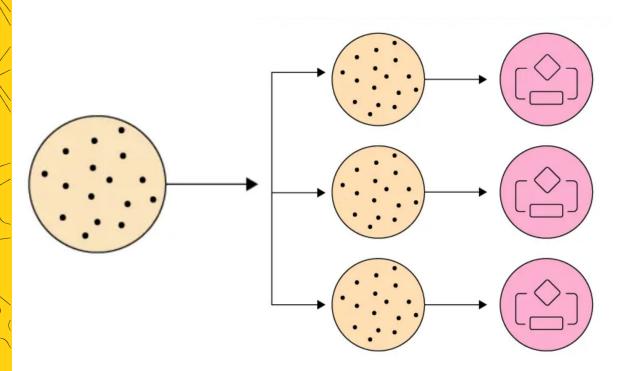
$$= \mathbb{E}_{x} \left( \frac{1}{N} \sum_{j=1}^{N} \varepsilon_{j}(x) \right)^{2} =$$

$$= \frac{1}{N^{2}} \mathbb{E}_{x} \left( \sum_{j=1}^{N} \varepsilon_{j}^{2}(x) + \underbrace{\sum_{i \neq j} \varepsilon_{i}(x) \varepsilon_{j}(x)}_{=0} \right) =$$

## Bagging = Bootstrap aggregating

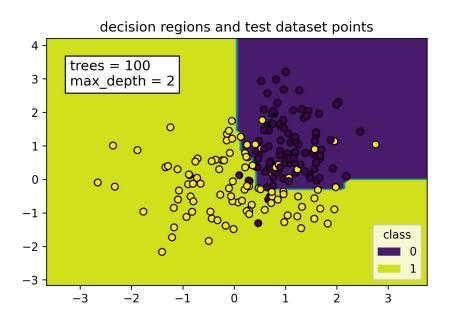


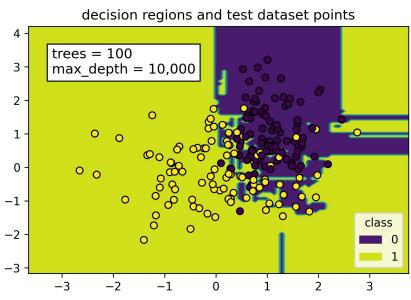
Decreases the variance if the basic algorithms are not correlated



### **Bagging overfitting**







# Random Forest

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### **RSM - Random Subspace Method**



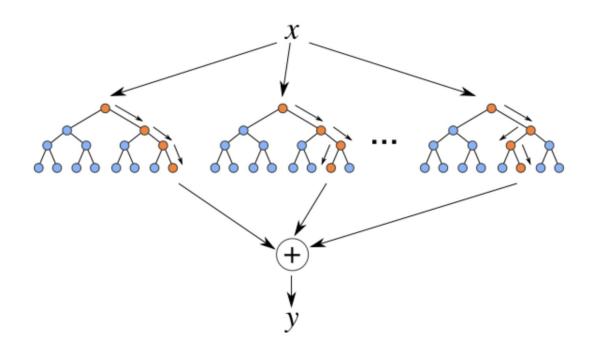
Same approach, but with features.

Just subsample of features for each bootstraped dataset

### **Random Forest**



Bagging + RSM = Random Forest



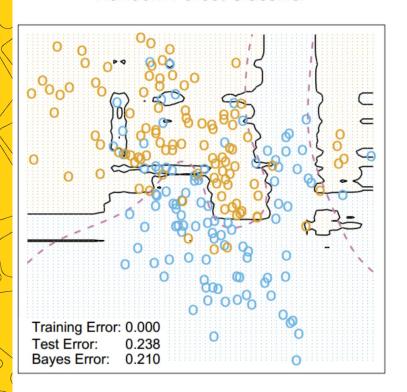
#### **Random Forest**



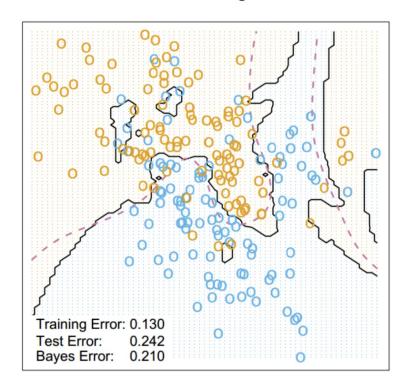
- One of the greatest "universal" models
- There are some modifications: Extremely Randomized Trees, Isolation Forest, etc



#### Random Forest Classifier



#### 3-Nearest Neighbors



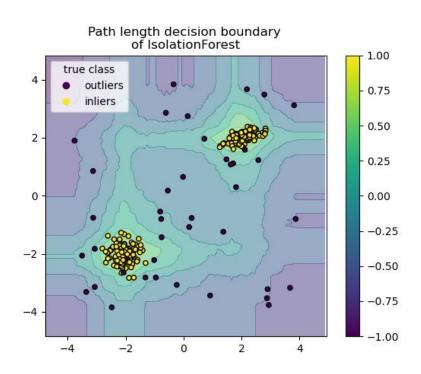
# Isolation forest

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#### Method to search for anomalies





#### Method to search for anomalies



Isolation Forest 'isolates' observations by randomly selecting a feature and then randomly selecting a split value between the maximum and minimum values of the selected feature.

This path length, averaged over a forest of such random trees, is a measure of normality and our decision function.

Random partitioning produces noticeably shorter paths for anomalies. Hence, when a forest of random trees collectively produce shorter path lengths for particular samples, they are highly likely to be anomalies.

https://scikit-learn.org/stable/modules/outlier\_detection.html#isolation-forest

https://alexanderdyakonov.wordpress.com/2017/04/19/%D0%BF%D0%BE%D0%B8%D0%BA-%D0%B0%D0%BD%D0%BE%D0%BC%D0%B0%D0%BB%D0%B8%D0%B9-anomaly-detection/

# Revise

- 1. Intuition
- 2. Construction procedure
- 3. Information criteria
- 4. Decision trees special highlights
  - o Decision tree as linear model
  - Dealing with missing data
  - Categorical features
- 5. Bootstrap and Bagging
- 6. Random Forest



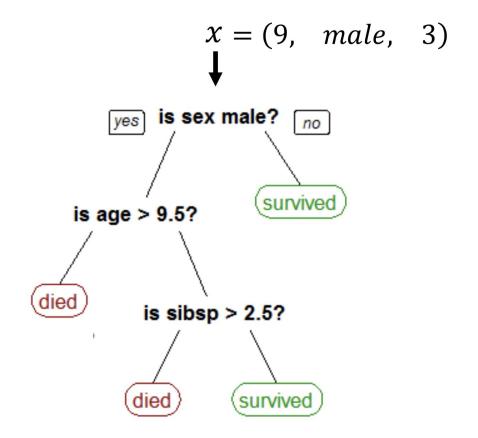
# Thanks for attention!

Questions?

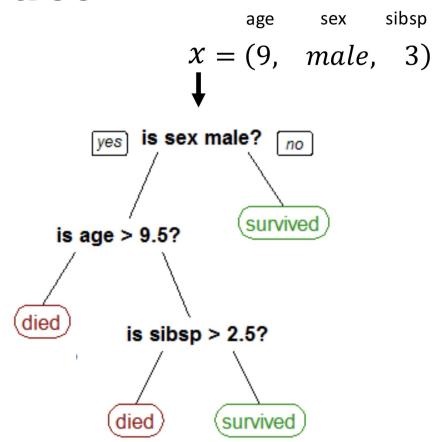




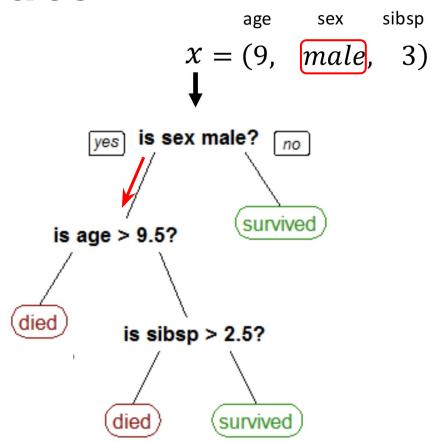




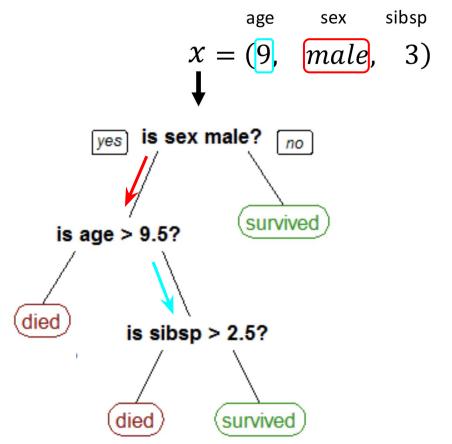




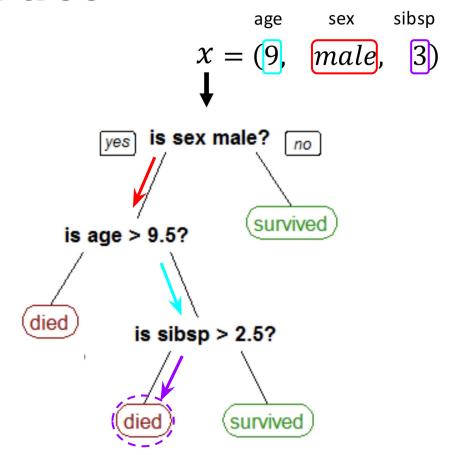




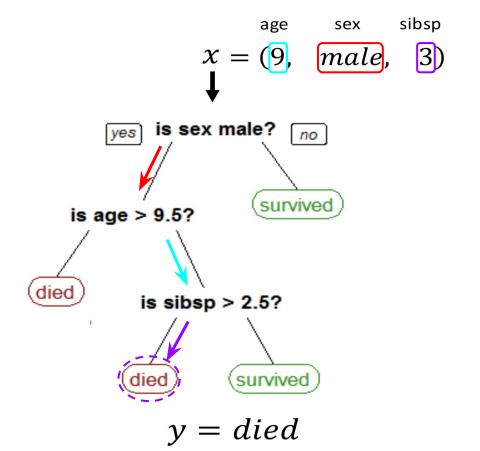




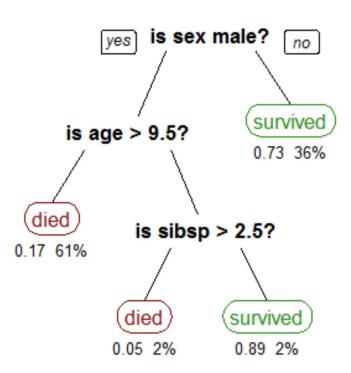




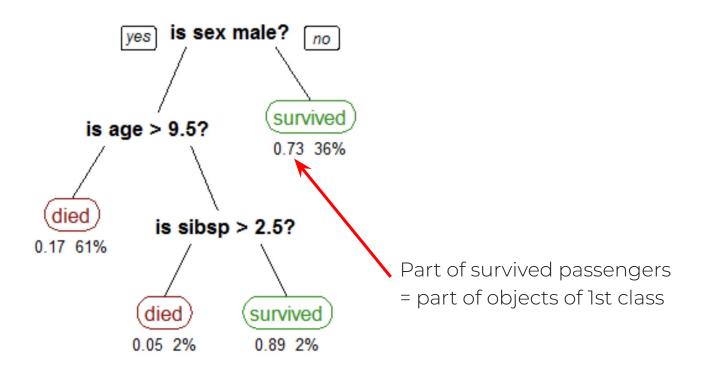




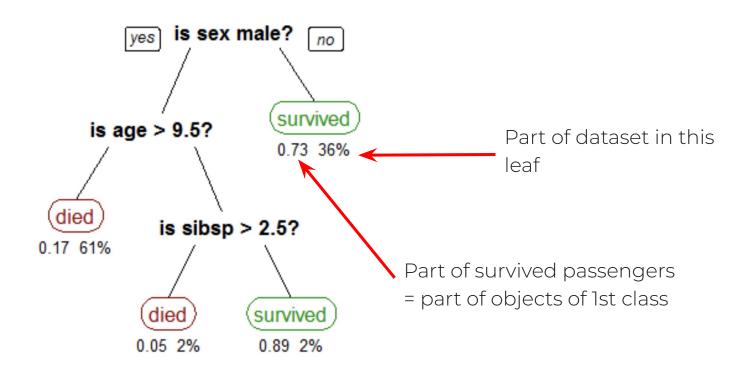




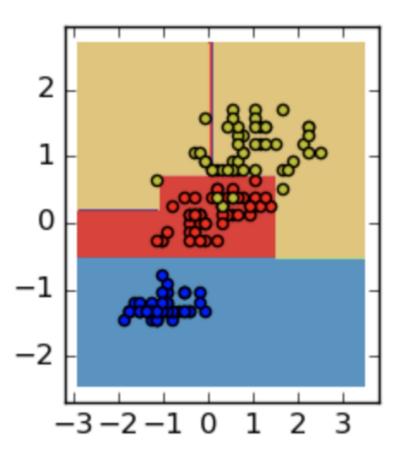












Classification problem with 3 classes and 2 features.