#### Week Goals

Decision Trees: training and application

Overfitting in Decision Trees

Working with categorical features

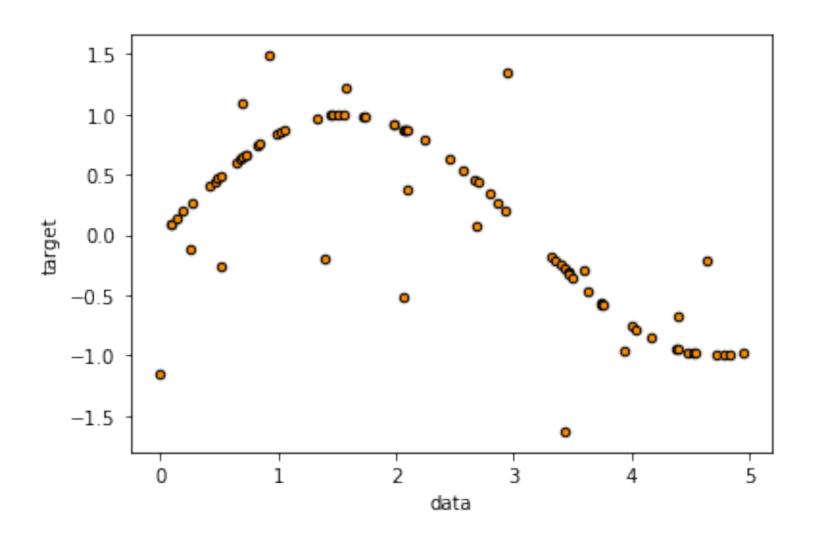
• Target: price of an apartment

• Features: Area, Floor, Distance to nearest metro station, etc.

• Linear Model:

$$a(x) = w_0 + w_1 * (area) + w_2 * (floor) + w_3 * (dist.) + ...$$

• It is likely that the dependency is not linear



Linear Model:

$$a(x) = w_0 + w_1 * (area) + w_2 * (floor) + w_3 * (dist.) + ...$$

• It is likely that the features are interconnected

• Linear model with polynomial features:

$$a(x) = w_0 + w_1 * (area) + w_2 * (floow) + w_3 * (dist.)$$
  
+ $w_4 * (area)^2 + w_5 * (floor)^2 + w_6 * (dist.)^2$   
+ $w_7 * (area) * (floor) + ...$ 

• Linear model with polynomial features:

$$a(x) = w_0 + w_1 * (area) + w_2 * (floow) + w_3 * (dist.)$$
  
+ $w_4 * (area)^2 + w_5 * (floor)^2 + w_6 * (dist.)^2$   
+ $w_7 * (area) * (floor) + ...$ 

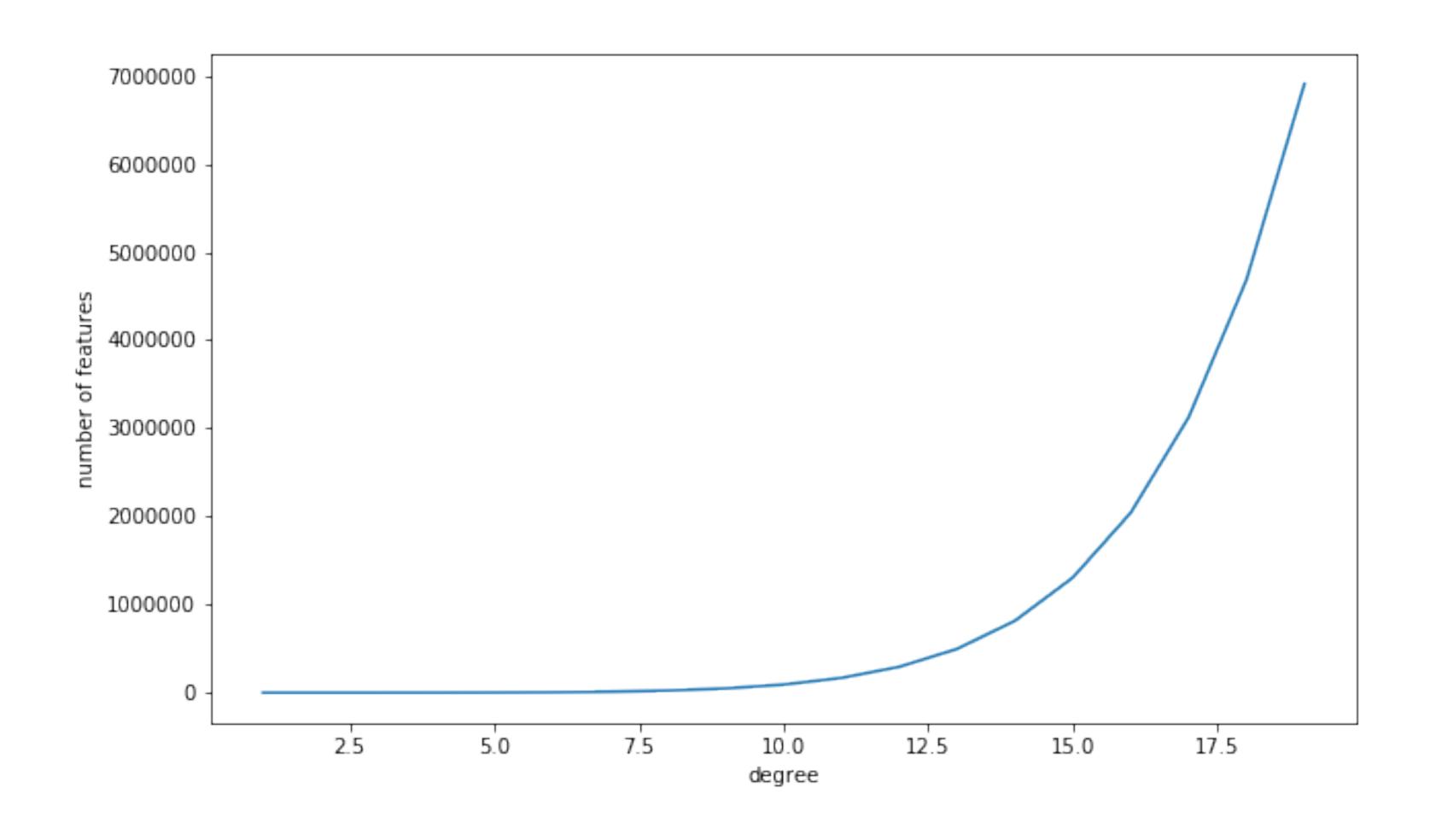
- It maybe hard to interpret the result
- What does  $(dist.) * (floor)^2$  mean?

• Linear model with polynomial features:

$$a(x) = w_0 + w_1 * (area) + w_2 * (floow) + w_3 * (dist.)$$
  
+ $w_4 * (area)^2 + w_5 * (floor)^2 + w_6 * (dist.)^2$   
+ $w_7 * (area) * (floor) + ...$ 

- Assume we had 10 features
- Polynomial of power 2 → 55 extra features
- Polynomial of power 3 → 220 extra features

Linear model with polynomial features:



Add intervals of the features:

$$a(x) = w_0 + w_1 * [30 < area < 50] + w_2 * [50 < area < 80]$$
  
 $+w_{20} * [2 < floor < 5] + ...$   
 $+w_{100} * [30 < area < 50] [2 < floor < 5] + ...$ 

• It is easier to interpret features:

• But we will have even more features!

# Summary

 We need an efficient algorithm to find non-linear dependency between features and target variables

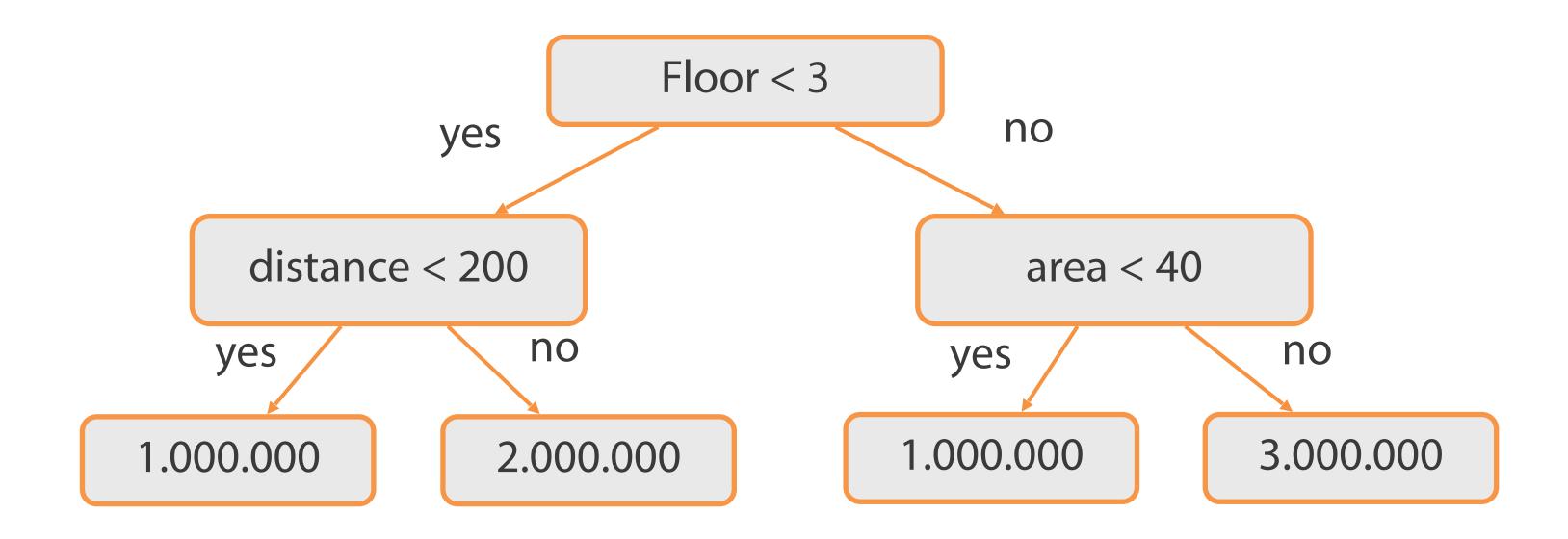
Decision trees will allow us to do that

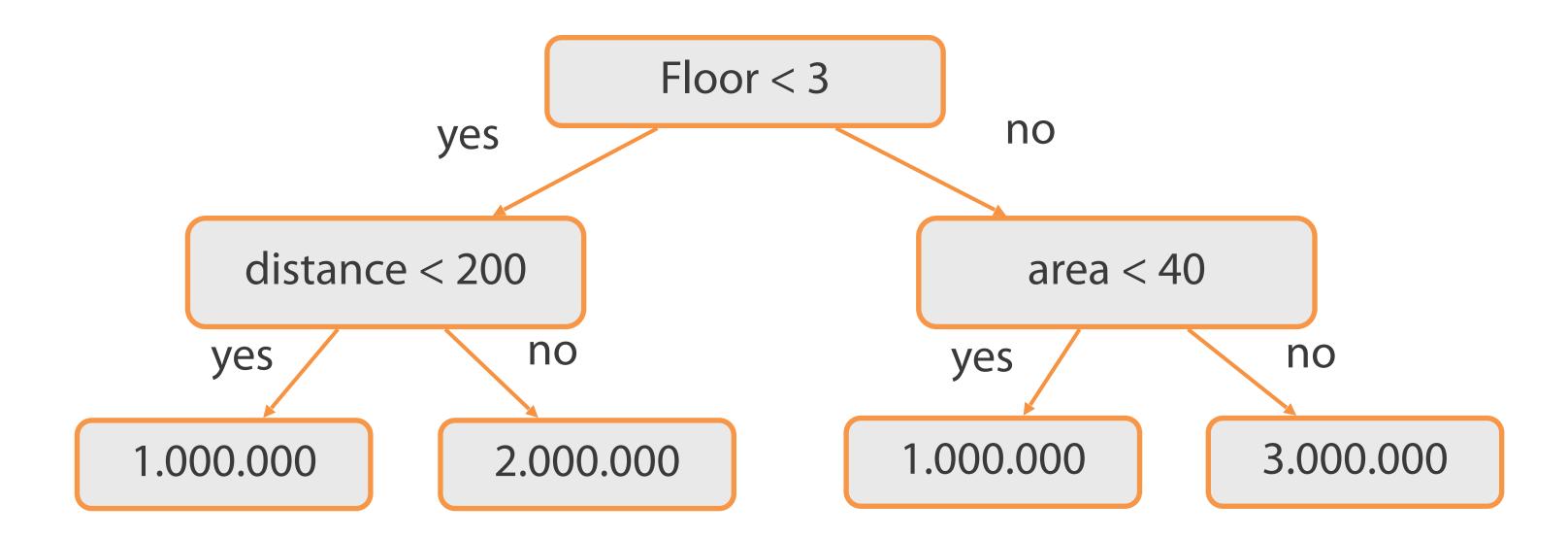
## Logical Rule

$$[30 < area < 50][2 < floor < 5][500 < dist. < 1000]$$

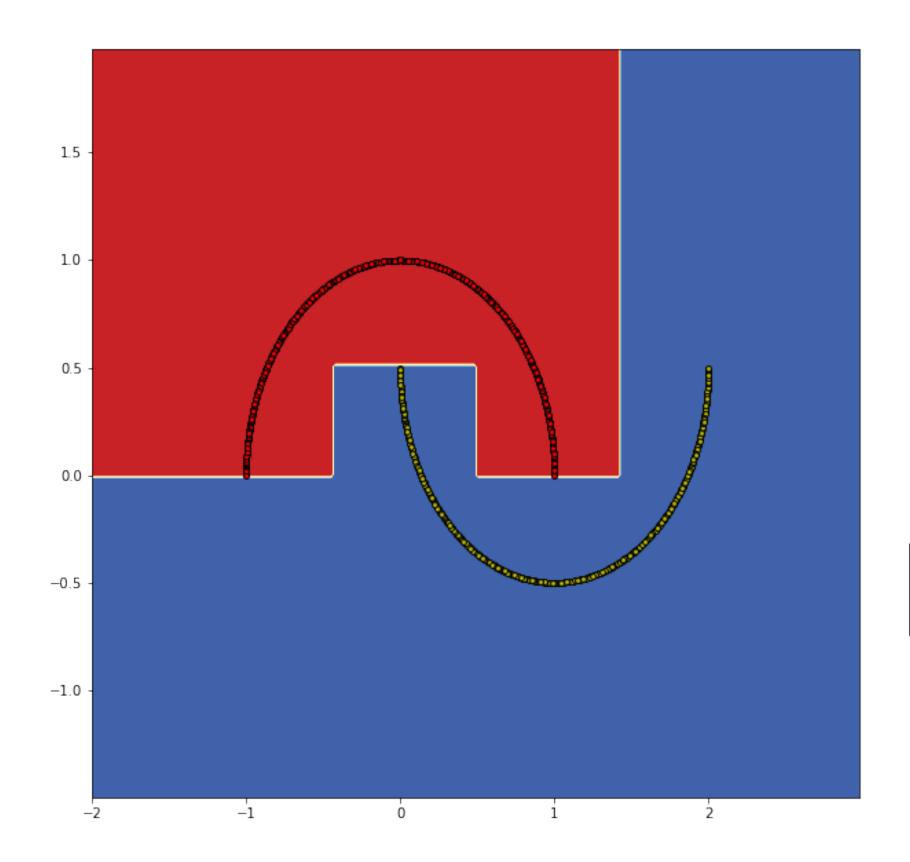
- Easy to explain
- Find non-linear dependencies

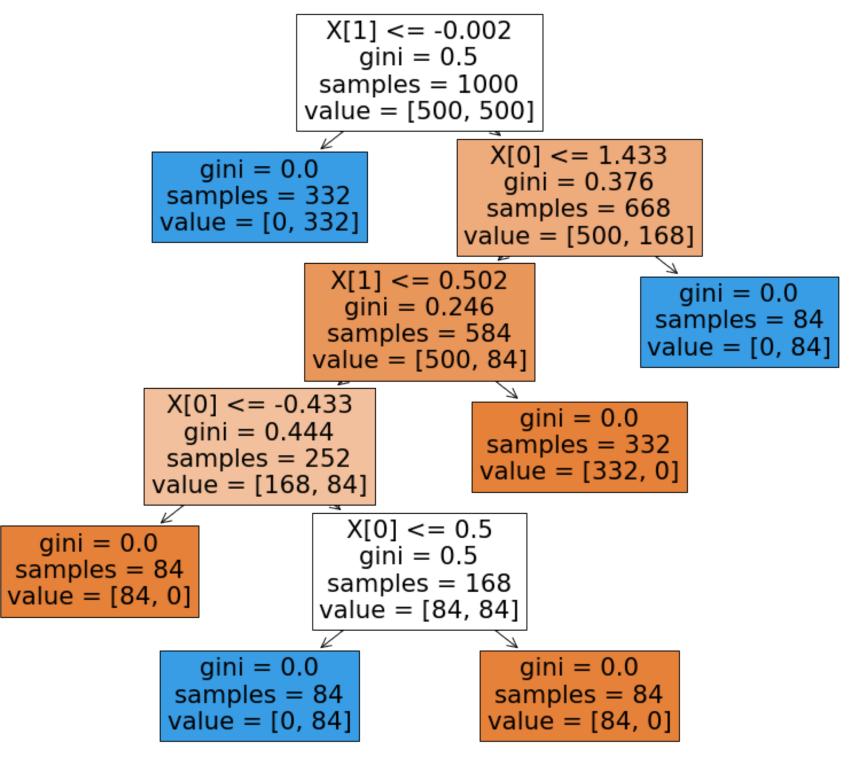
- We need to find good logical rules
- We need to build models out of them

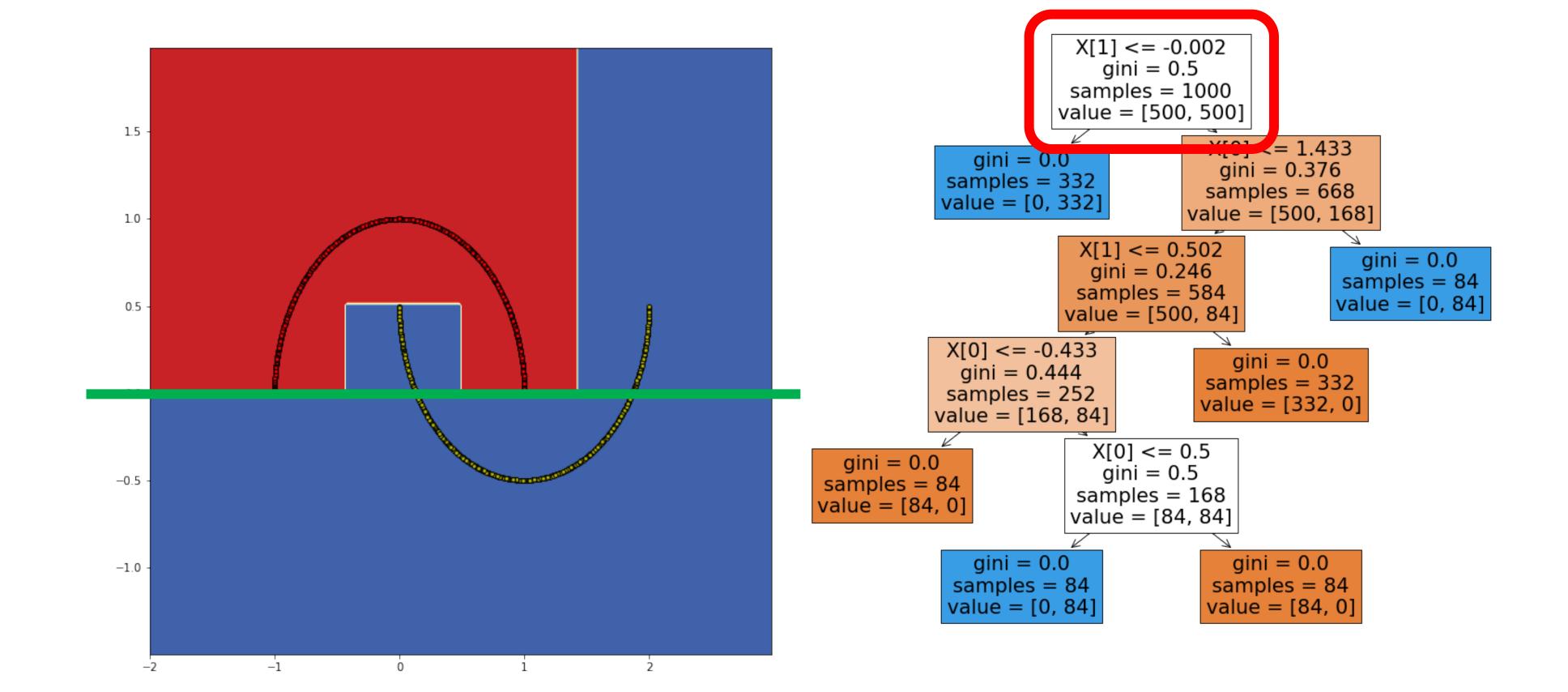


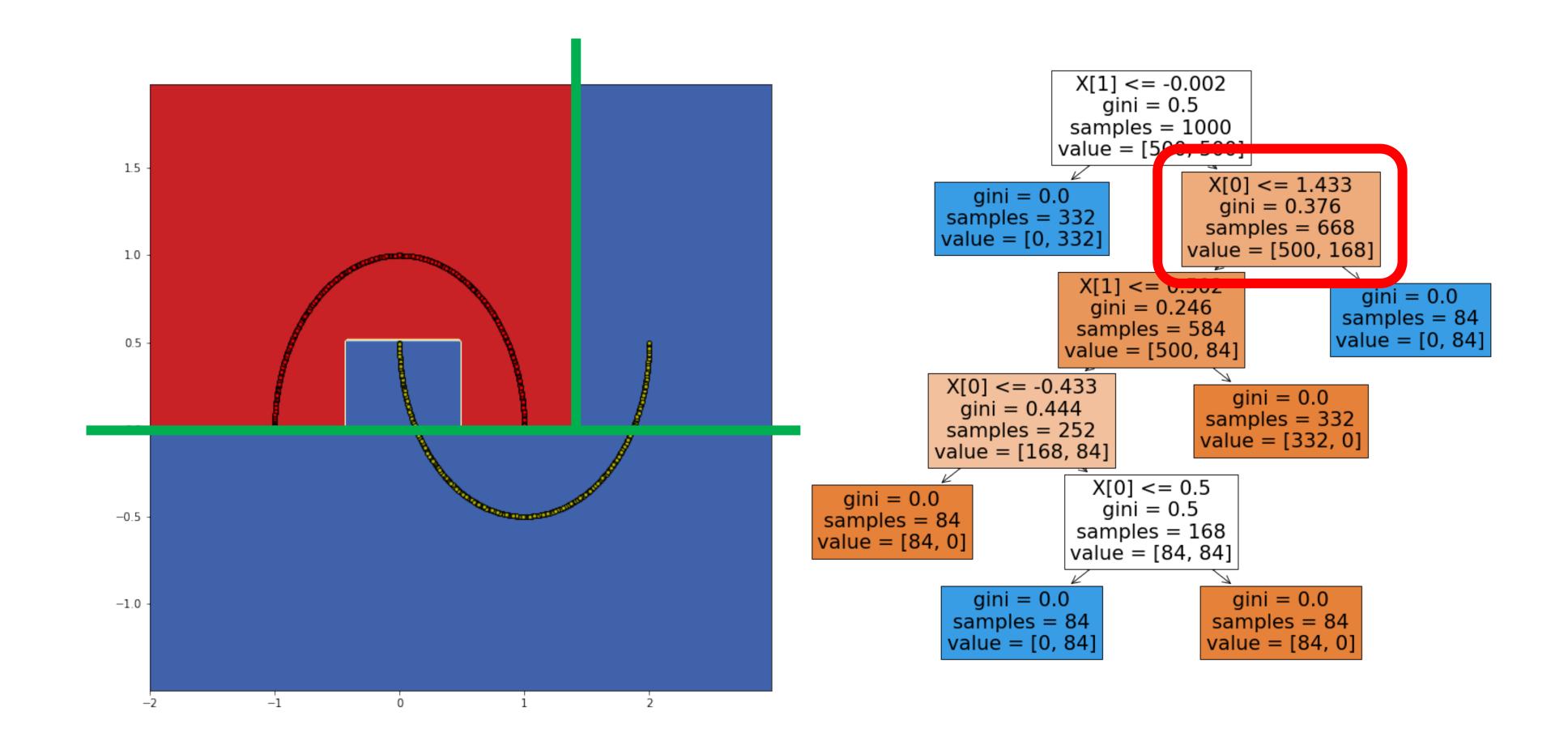


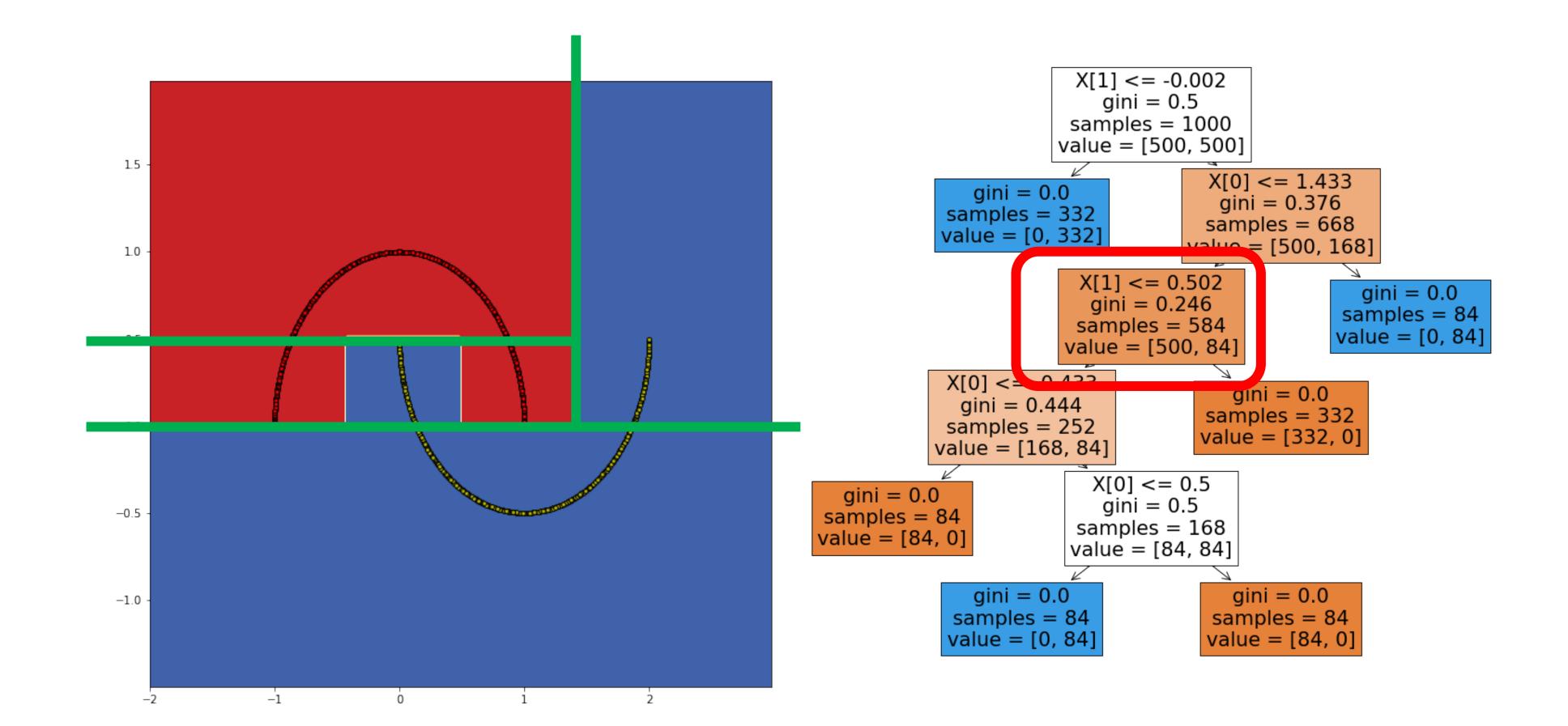
- Internal Nodes: splitting criterion  $\left[x_j < t\right]$
- Leaves: predictions  $c \in \mathbb{Y}$

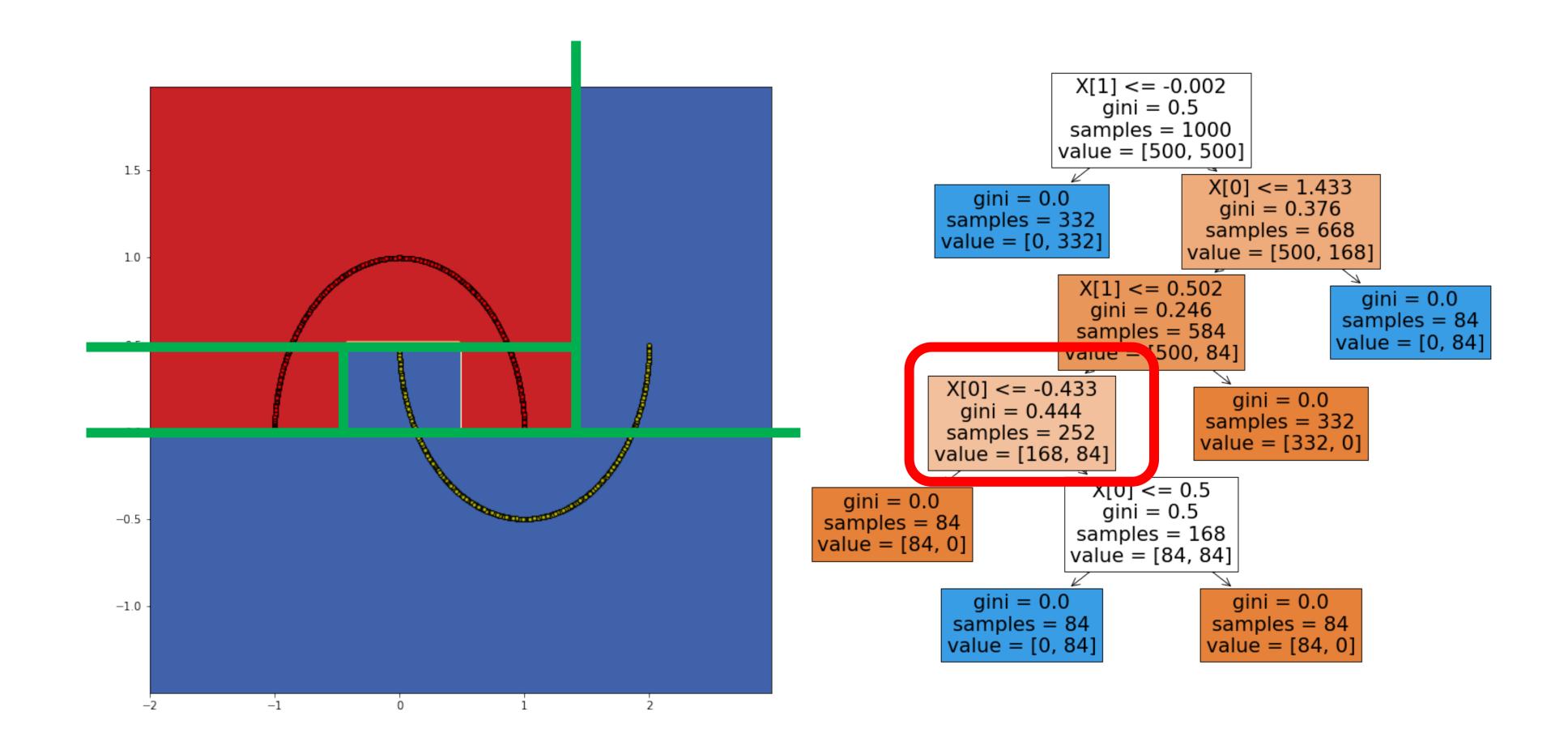


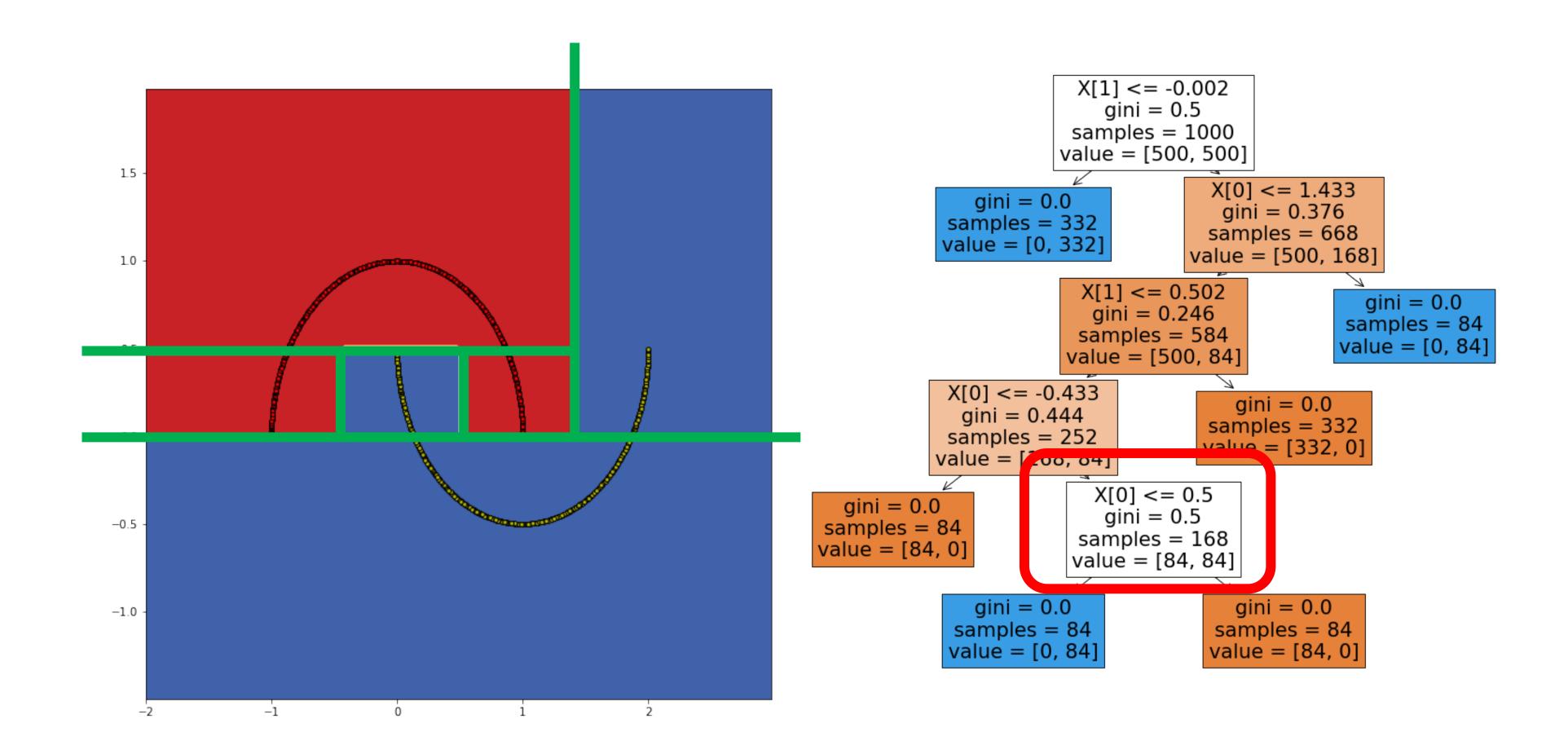




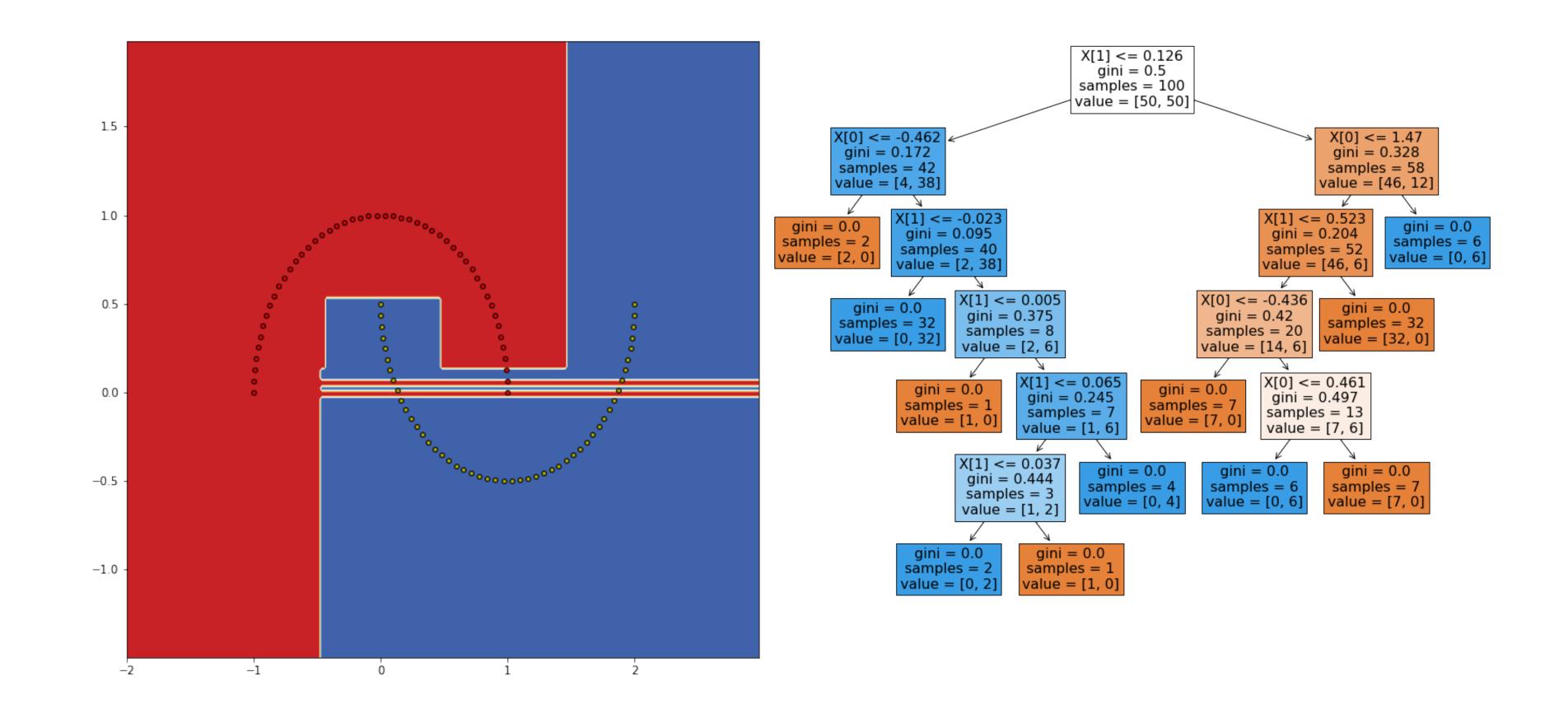








# **Decision Trees and Overfitting**

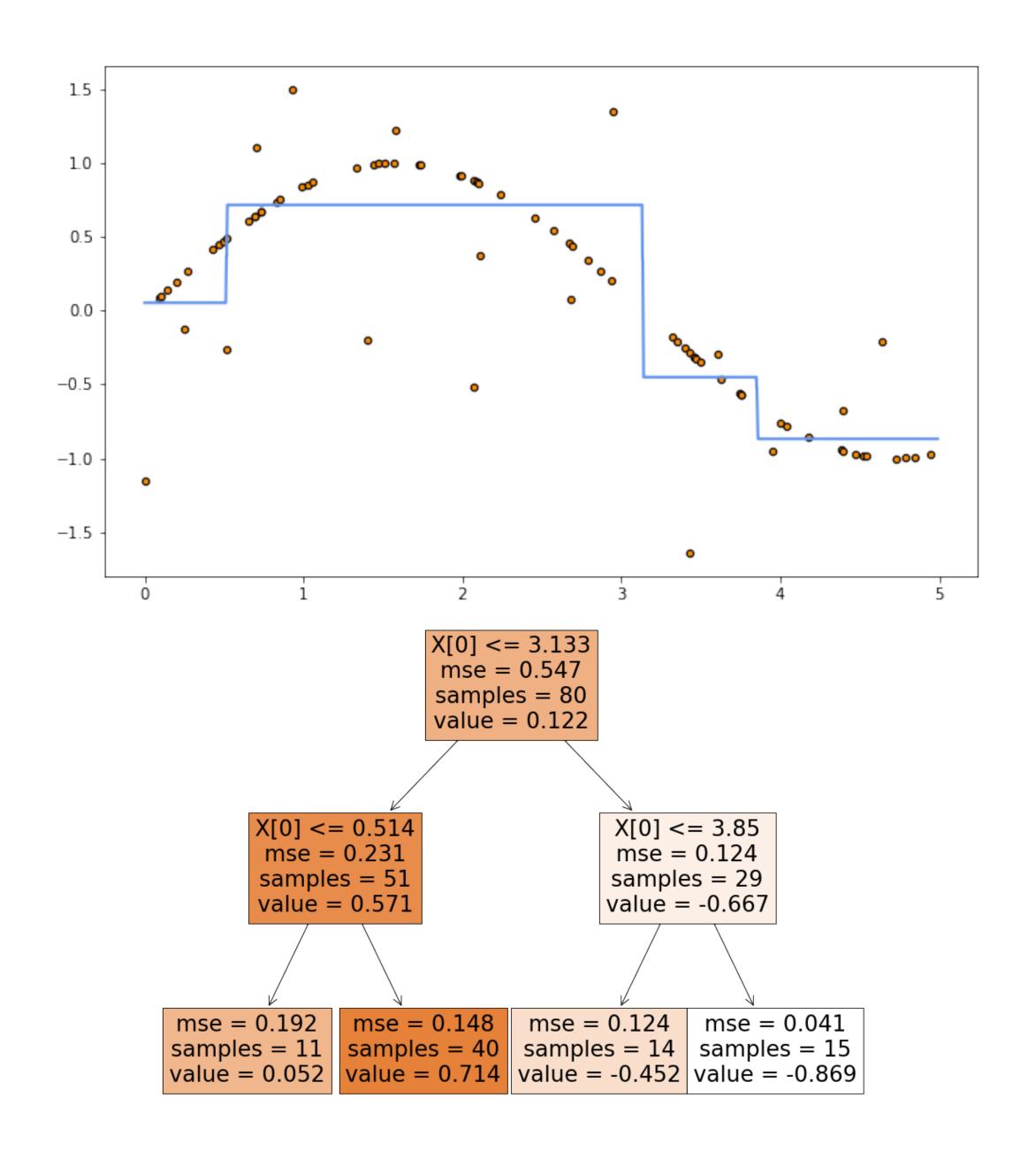


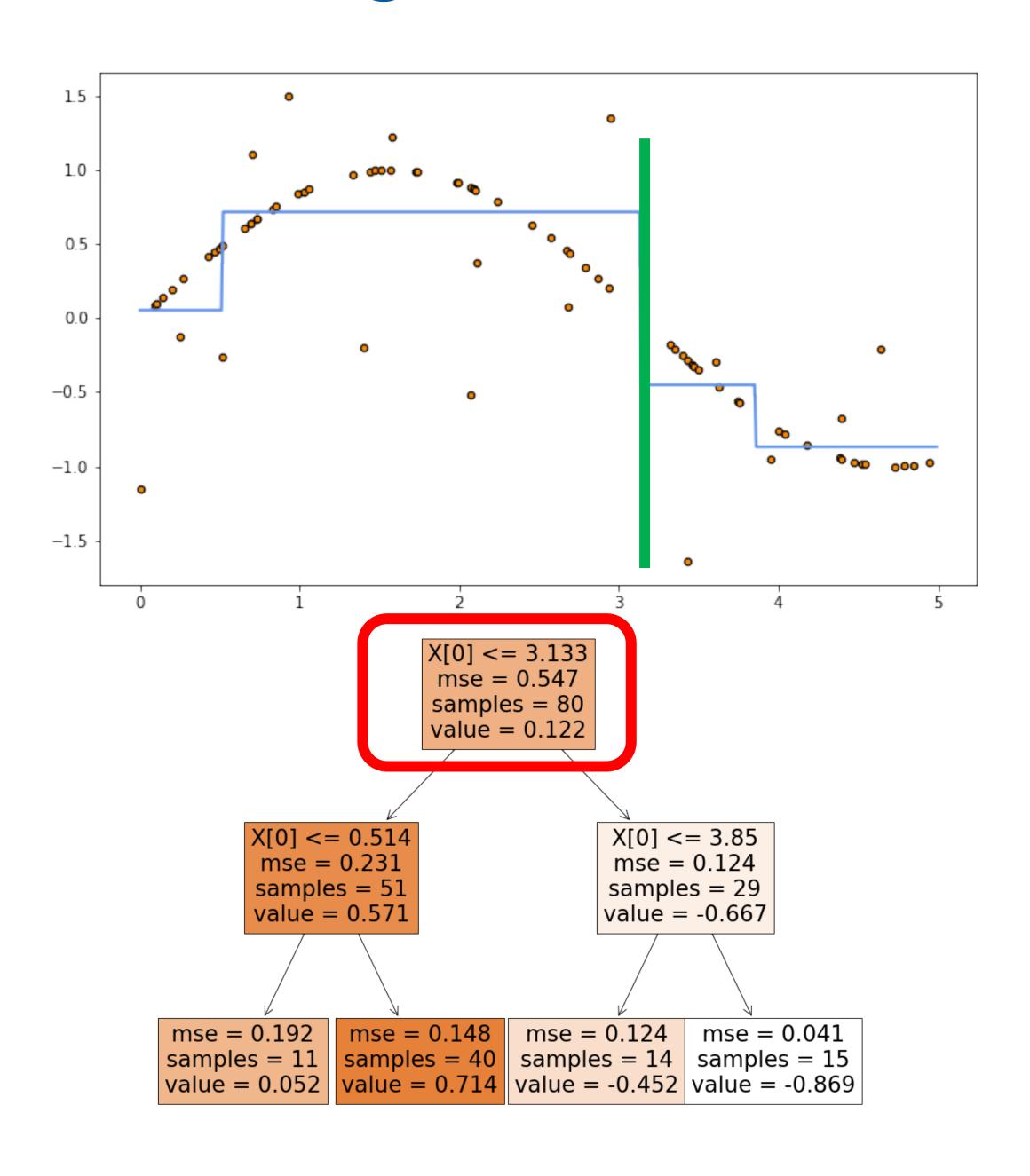
# **Complexity of Decision Trees**

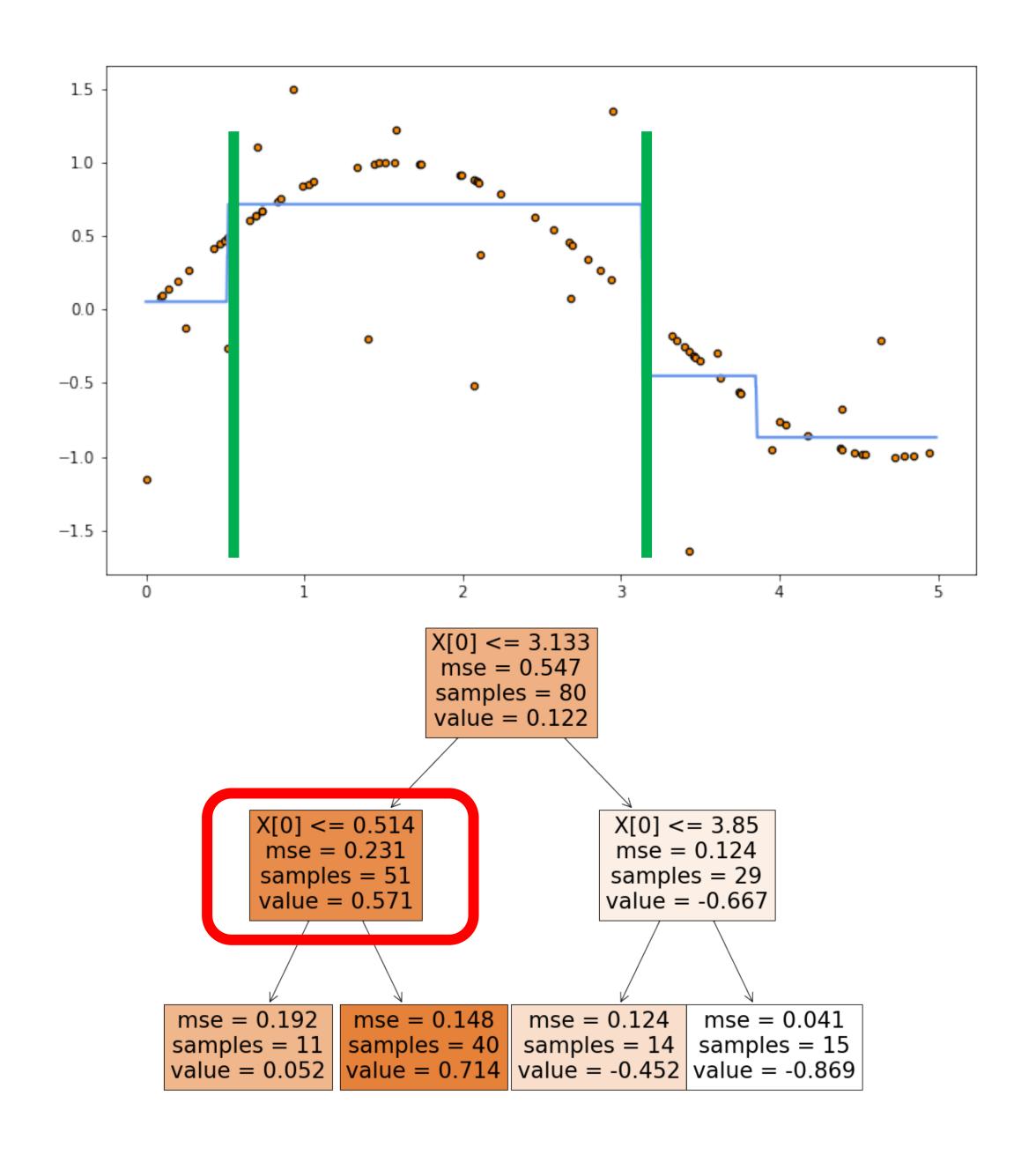
• We can keep splitting until we have only one object in each leaf

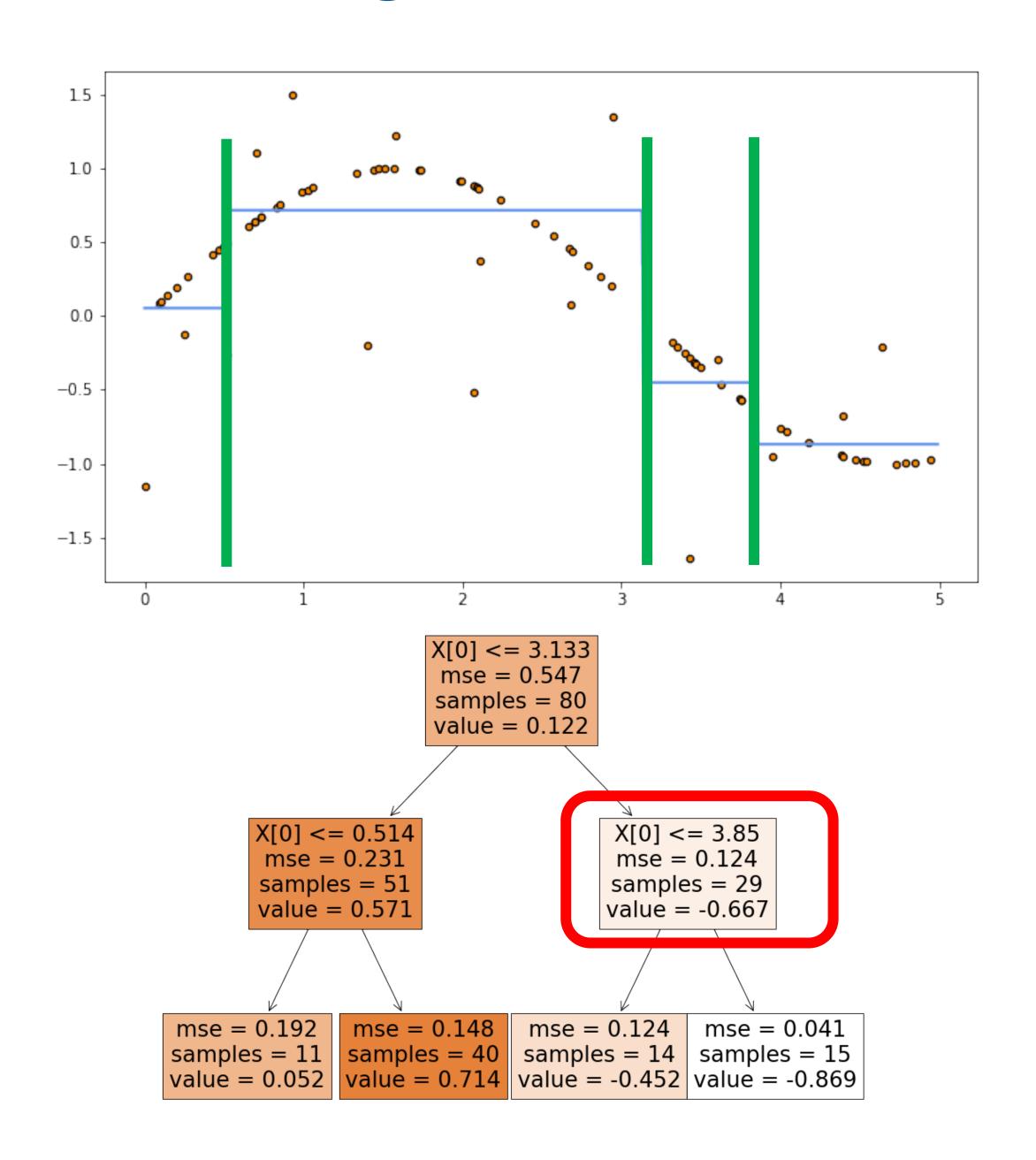
• We can ideally fit any training data

 Unless there are several objects with the same features and different target values

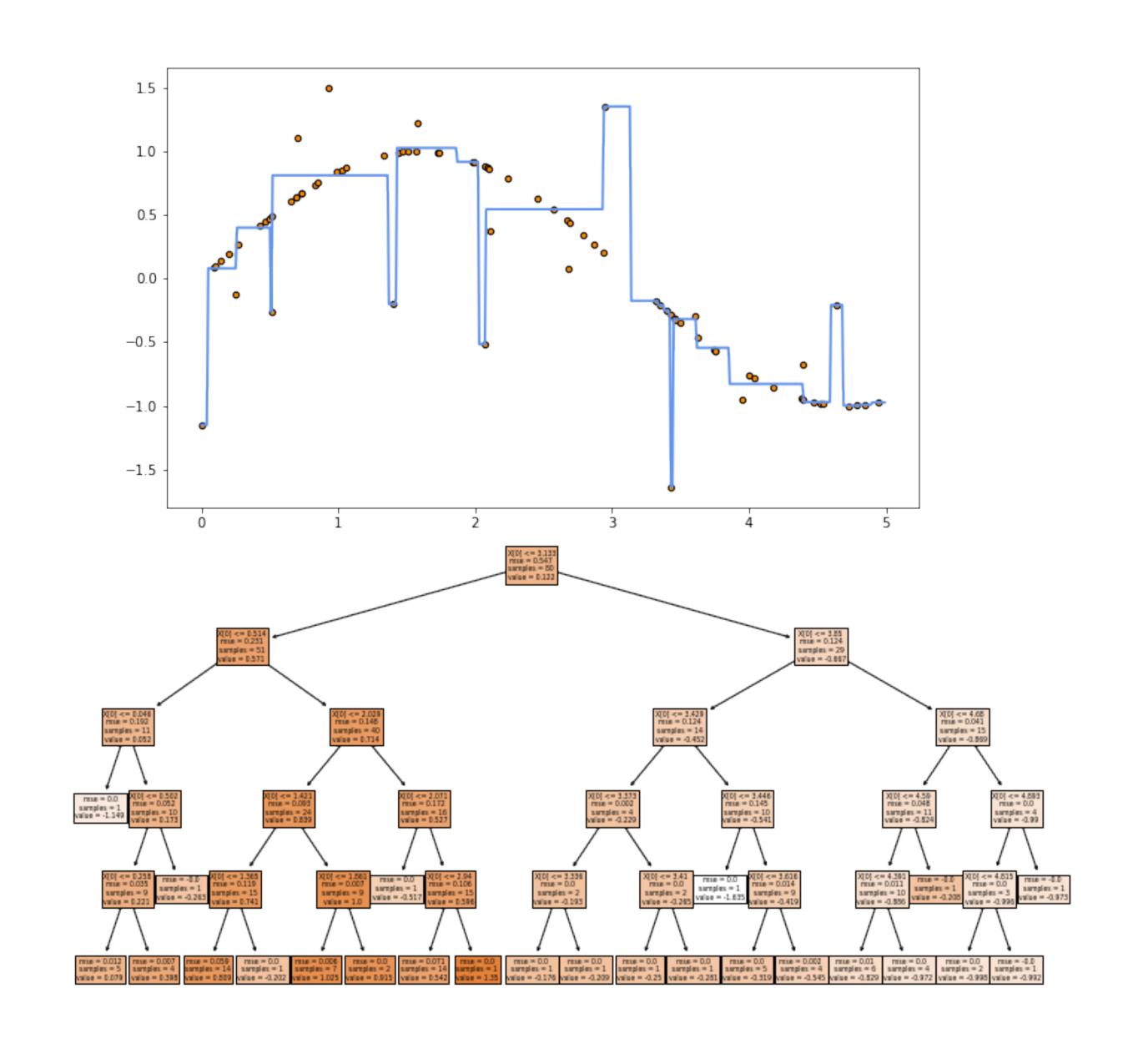








# Decision Trees for Regression and Overfitting



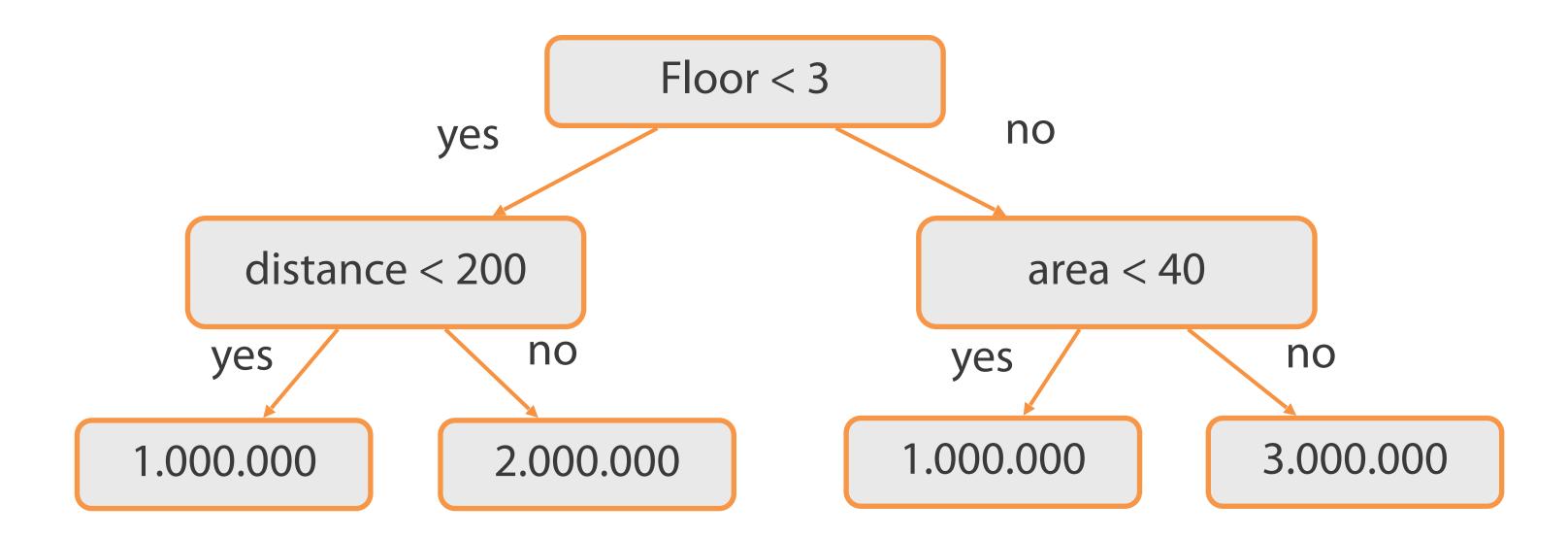
# Summary

• Decision trees — combination of simple logic rules

 Decision Trees split feature space into several areas with constant prediction in each of them

• It is quite easy to overfit, when you use decision trees

## **Structure of Decision Trees**



- Internal Nodes: splitting criterion
- Leaves: predictions  $c \in \mathbb{Y}$

# **Splitting Criterion**

- Step function:  $x_j < t$  not the only option
- Use a linear model:  $[\langle w, x \rangle < t]$
- A specific metric:  $\left[ \rho(x, x_0) < t \right]$

•

• But we can build arbitrary complex models even with the most simple predicates

## Predictions in Leaves: Regression

- We will use constant predictions  $c_v \in \mathbb{Y}$
- Average value:

$$c_v = \frac{1}{|R_v|} \sum_{(x_i, y_i) \in R_v} y_i$$

### **Predictions in Leaves: Classification**

- We will use constant predictions  $c_v \in \mathbb{Y}$
- The most common class:

$$c_v = \underset{k \in \mathbb{Y}}{\operatorname{argmax}} \sum_{k \in \mathbb{Y}} [y_i = k]$$
$$(x_i, y_i) \in R_v$$

#### **Predictions in Leaves: Classification**

- We will use constant predictions  $c_v \in \mathbb{Y}$
- The most common class:

$$c_v = \underset{k \in \mathbb{Y}}{\operatorname{argmax}} \sum_{k \in \mathbb{Y}} [y_i = k]$$
$$(x_i, y_i) \in R_v$$

Class probabilities

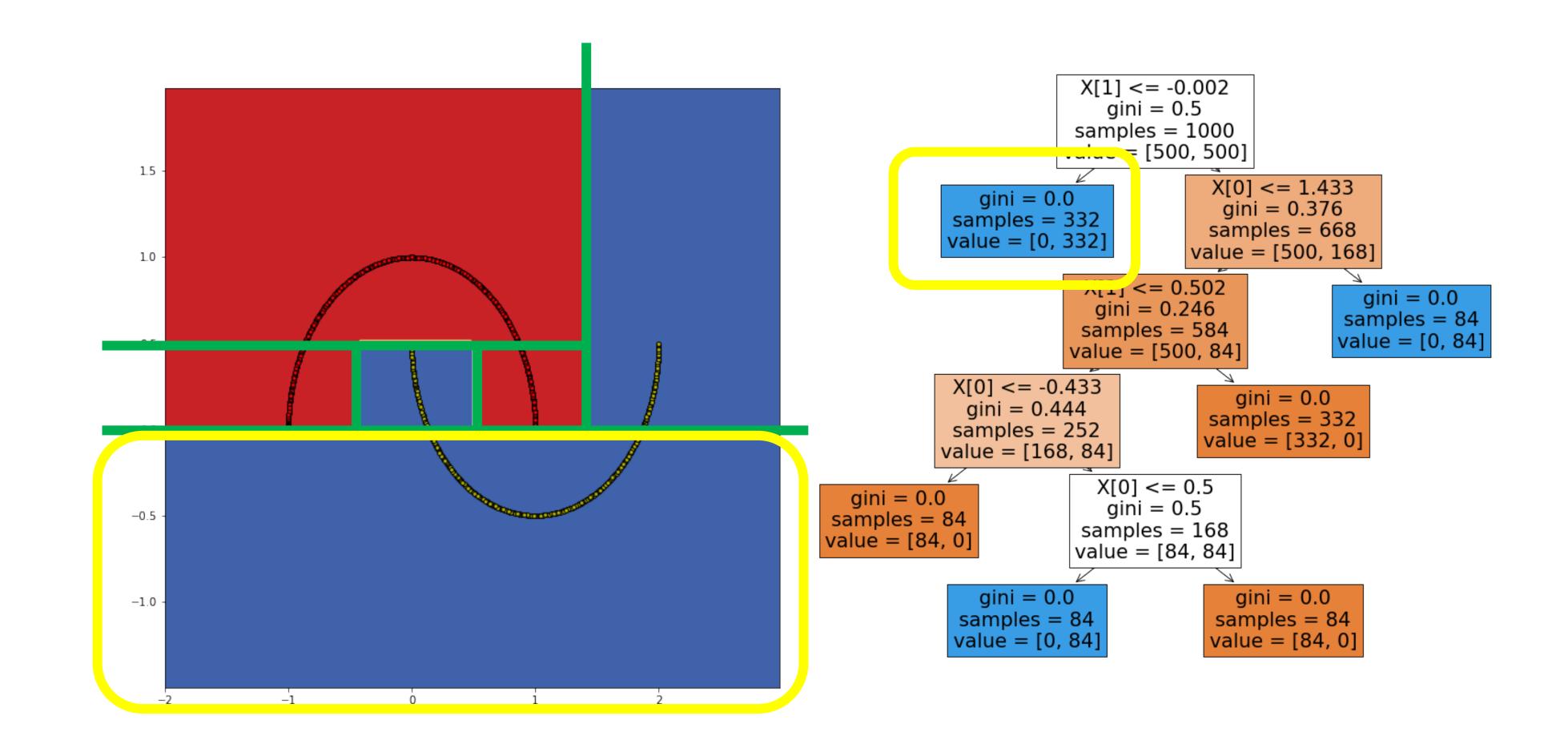
$$c_{vk} = \frac{1}{|R_v|} \sum_{(x_i, y_i) \in R_v} [y_i = k]$$

#### **Predictions in Leaves**

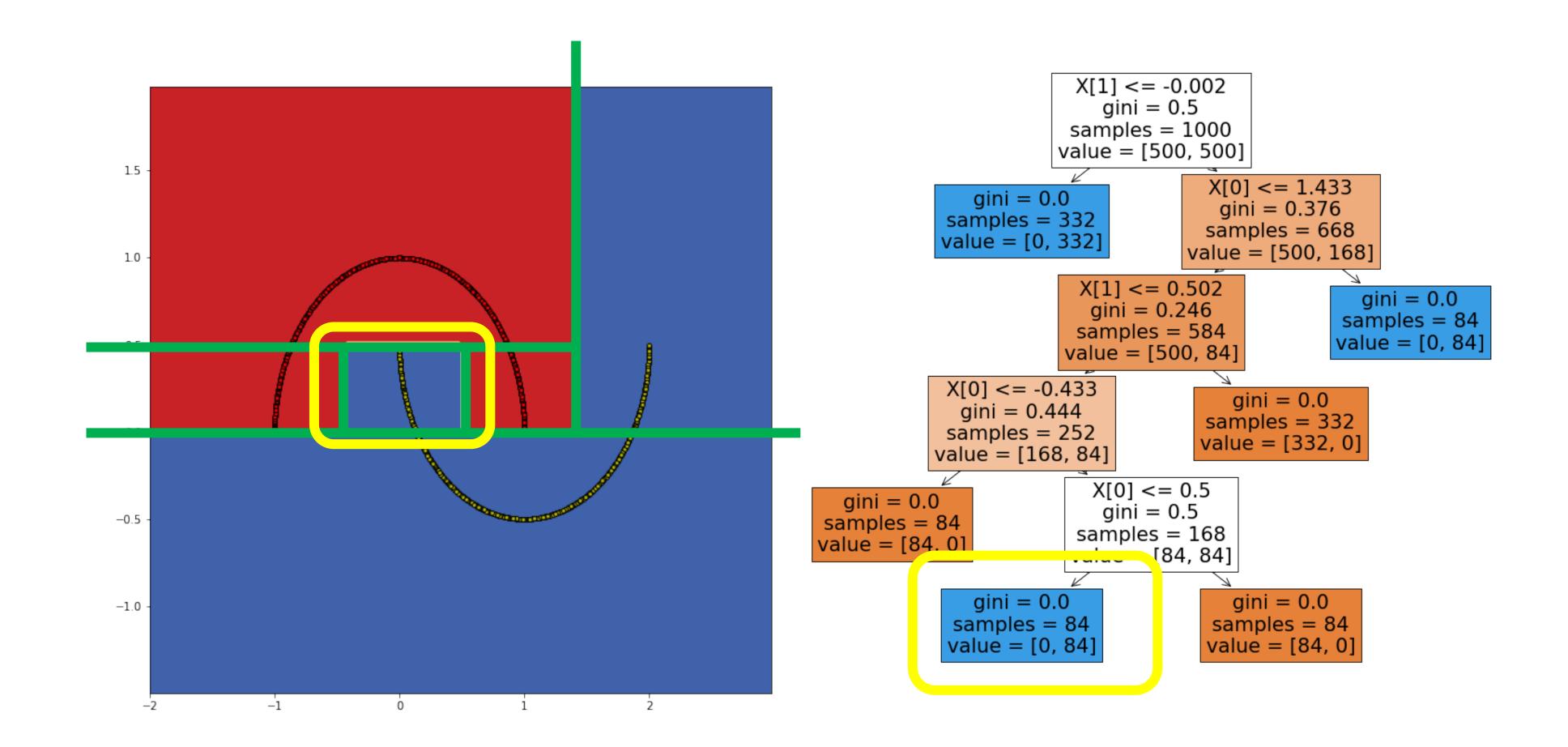
- We could use more complex prediction functions in leaves
- E.g. linear regression:

$$c_v(x) = \langle w_v, x \rangle$$

#### **Decision Tree**



#### **Decision Tree**



## **Decision Tree: Interpretation**

- Tree splits feature space on disjoint sub-spaces  $R_1, \, \dots, \, R_J$
- Each sub-space  $R_j$  corresponds to the leaf
- At each sub-space  $R_j$  prediction  $c_j$  is constant

$$a(x) = \sum_{j=1}^{J} c_j \left[ x \in R_j \right]$$

## **Decision Tree: Interpretation**

$$a(x) = \sum_{j=1}^{J} c_j \left[ x \in R_j \right]$$

- Decision tree constructs new powerful features
- The predictions then are linear combination of new features

# Summary

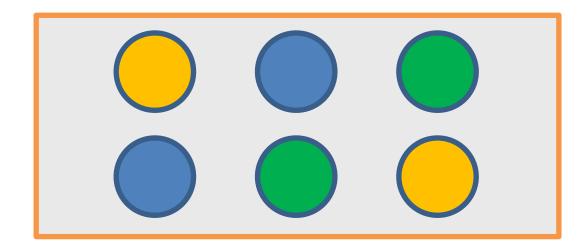
• One could use different approaches in splitting and making predictions in leaves. Usually the simplest one is good enough.

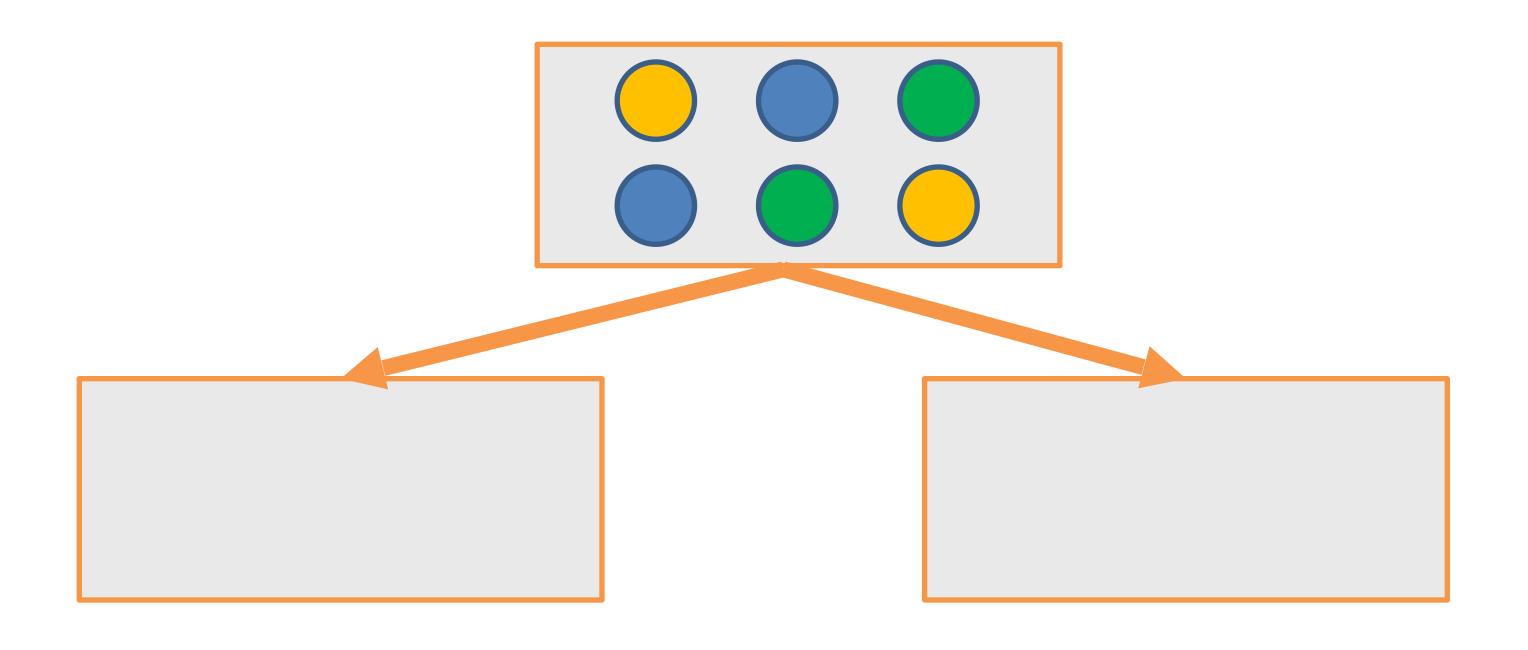
 One could think about decision tree as linear model over new features

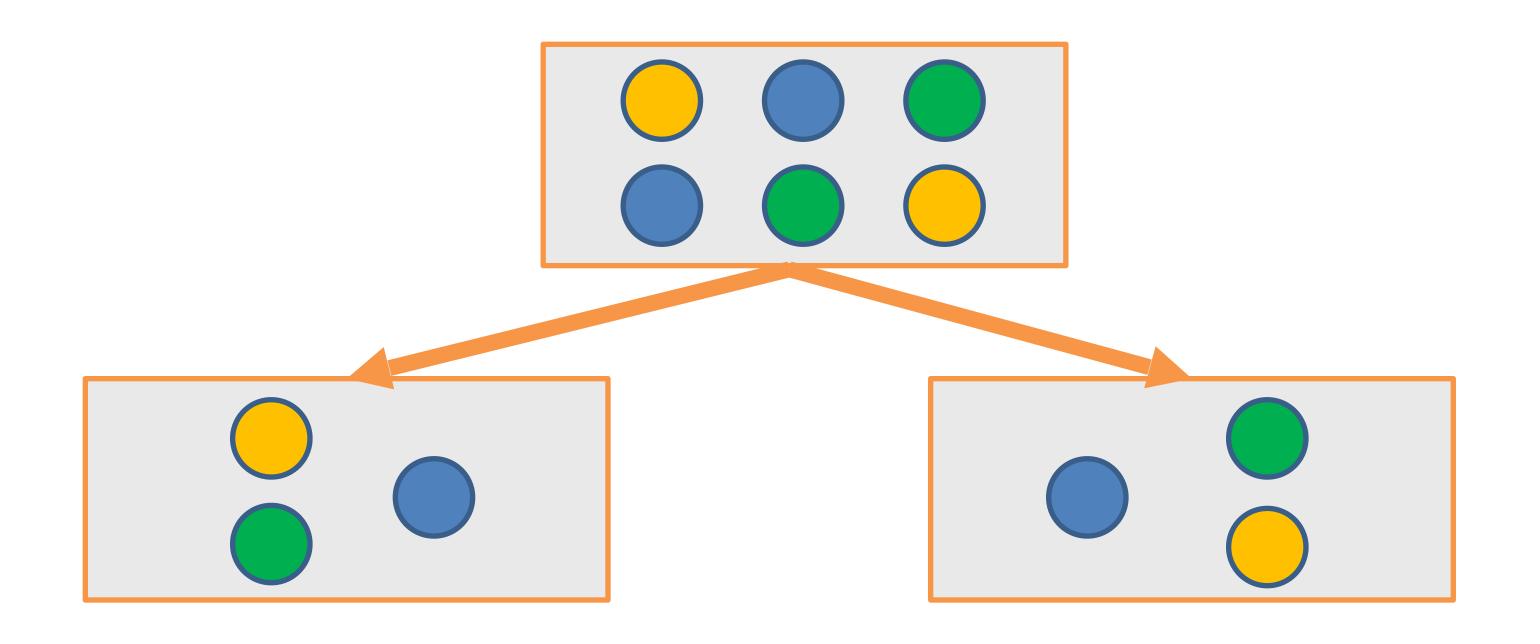
### **How to Train Decision Trees?**

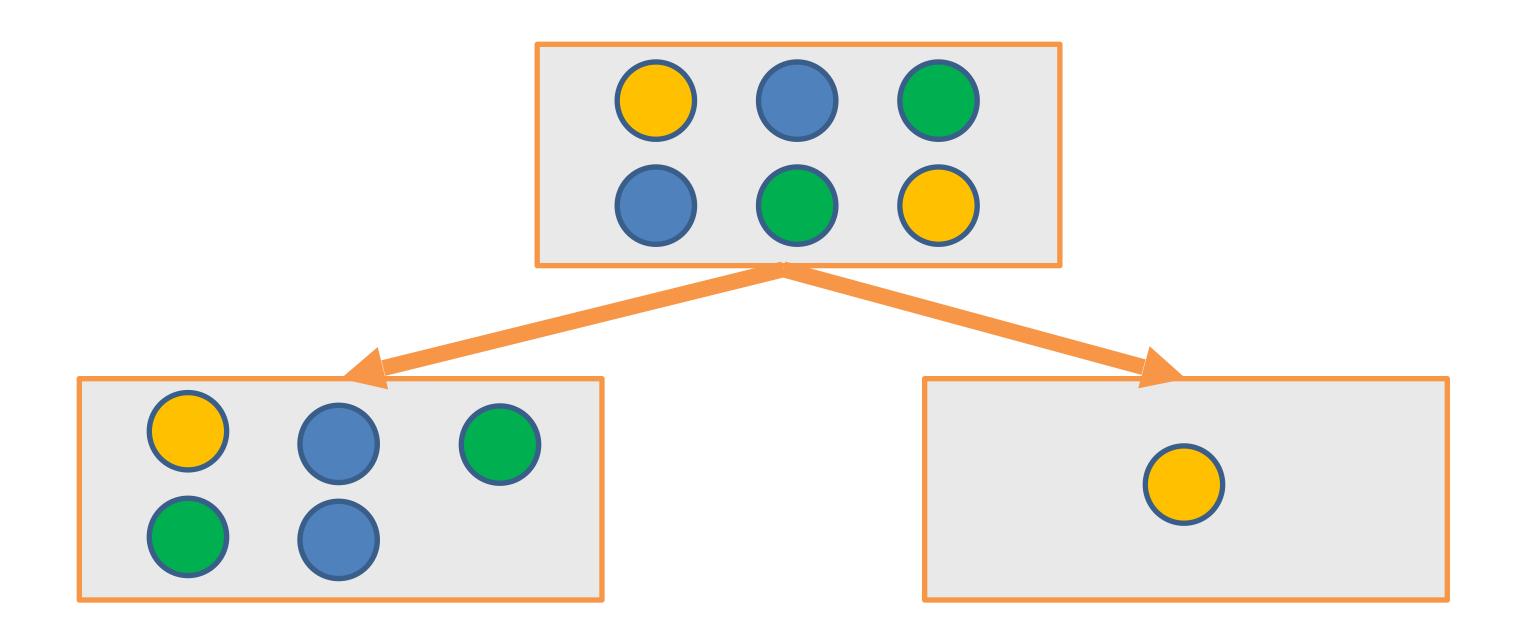
• Let's develop the algorithm on the example

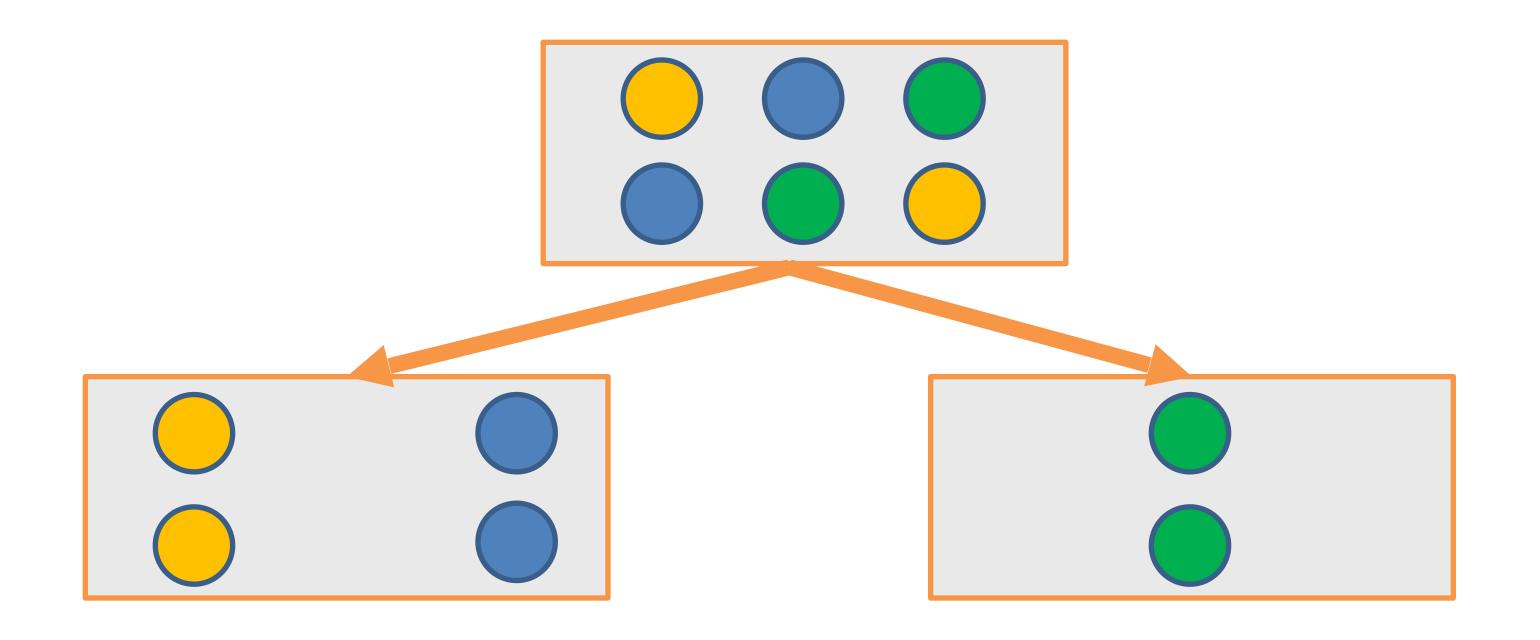
We start from the classification task

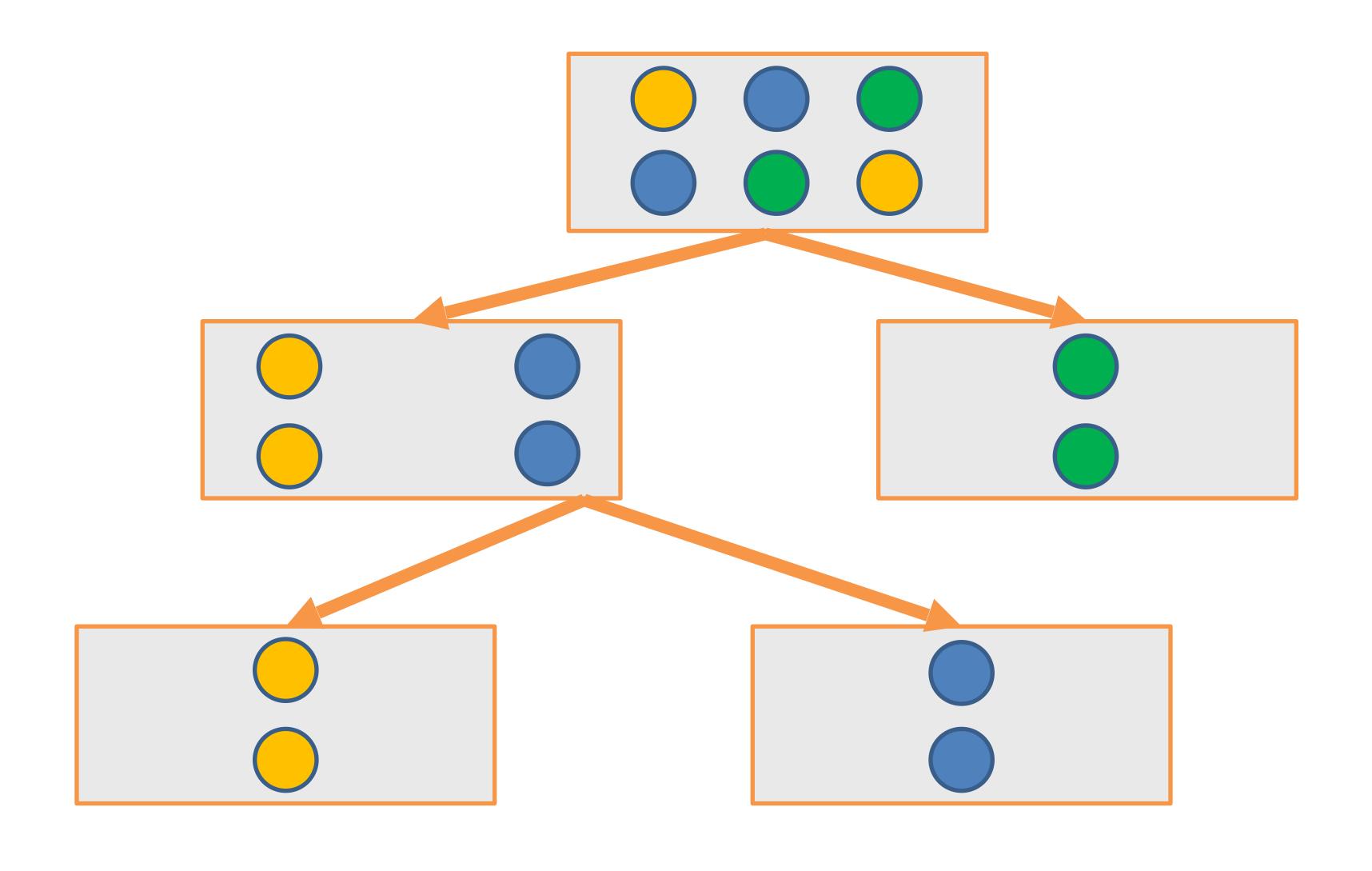




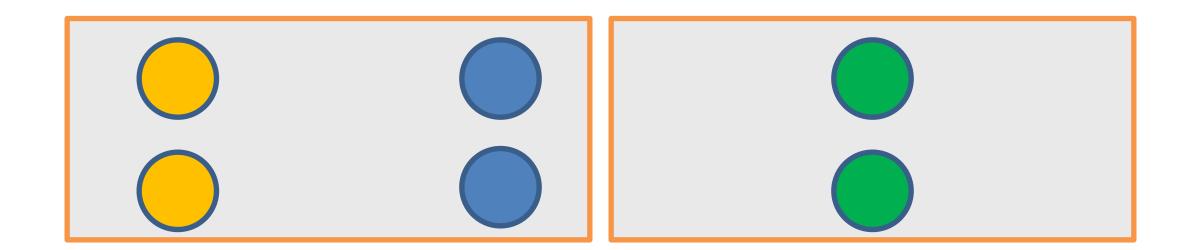




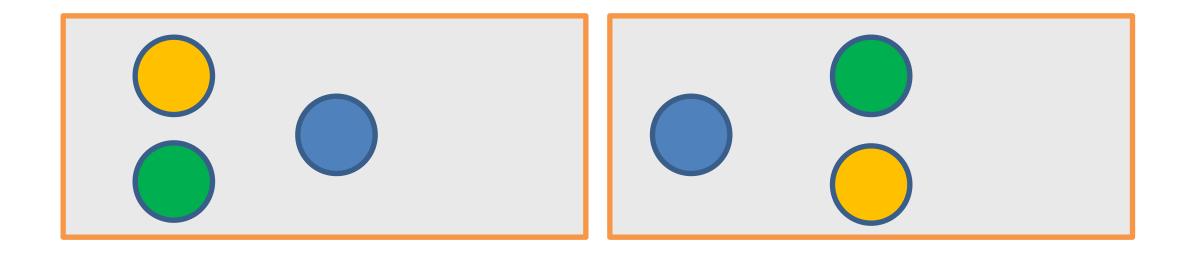




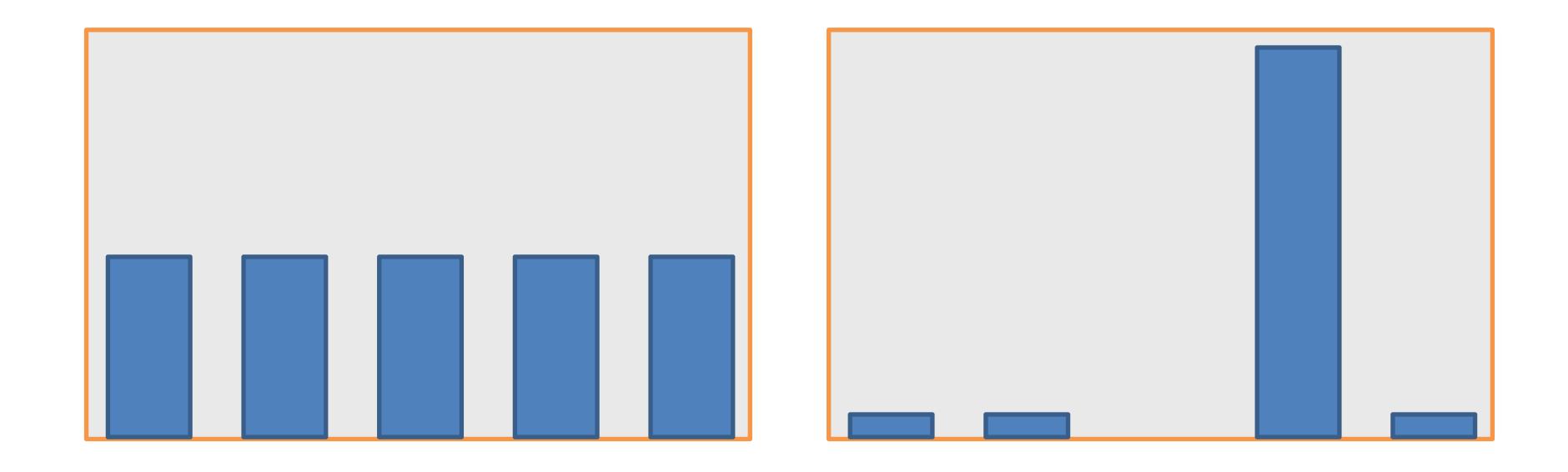
# How to Choose Between Two Splits?



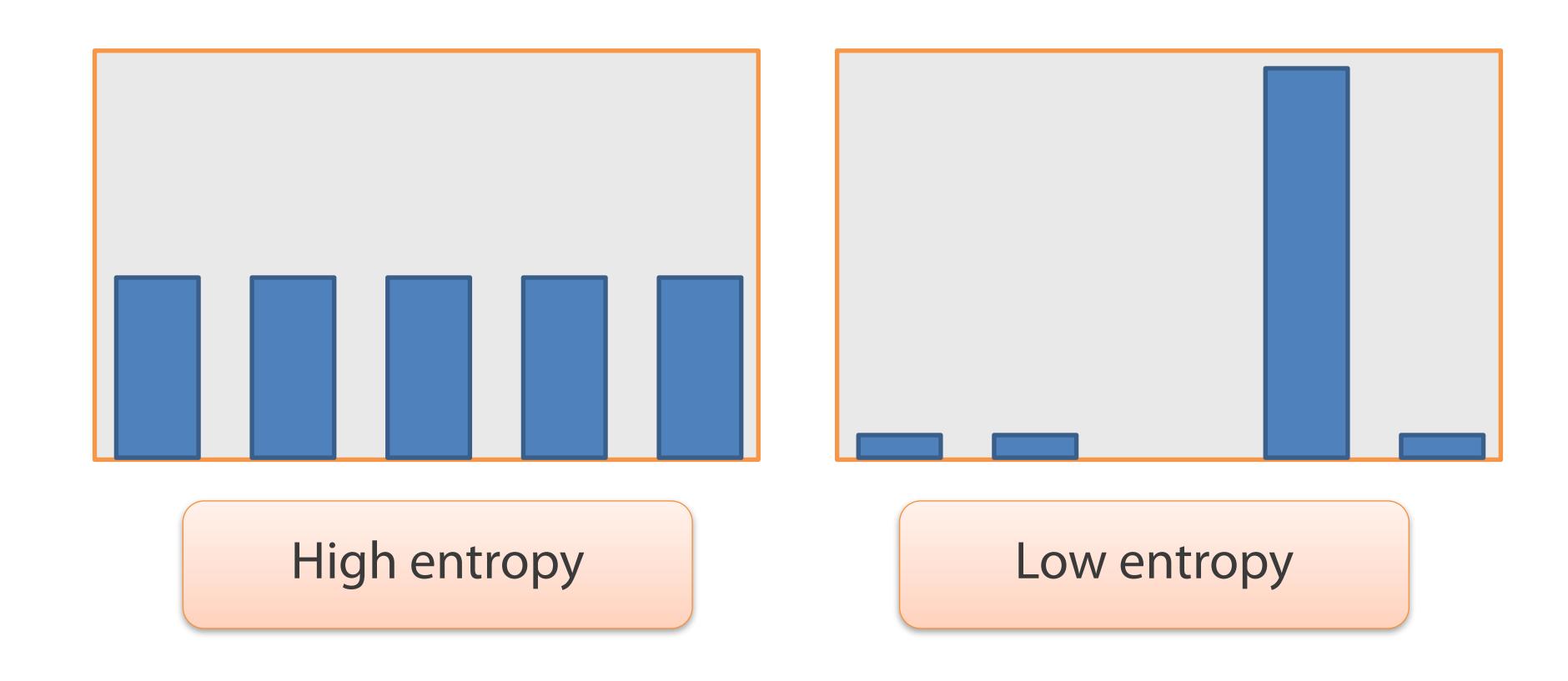
or



We consider entropy as a way to measure the uncertainty of an experiment's outcome



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- Assume we are given discrete distribution with n possible outcomes
- Probability of outcomes:  $p_1, p_2, ..., p_n$
- Entropy of distribution:

$$H(p_1, ..., p_n) = -\sum_{i=1}^{n} p_i \log_2 p_i$$

$$H(p_1, ..., p_n) = -\sum_{i=1}^{n} p_i \log_2 p_i$$

- p = (0.2, 0.2, 0.2, 0.2, 0.2)
  - H = 2.3219

$$H(p_1, ..., p_n) = -\sum_{i=1}^{m} p_i \log_2 p_i$$

- p = (0.2, 0.2, 0.2, 0.2, 0.2)
  - H = 2.3219
- p = (0.9, 0.05, 0.05, 0, 0)
  - H = 0.5689

$$H(p_1, ..., p_n) = -\sum_{i=1}^{n} p_i \log_2 p_i$$

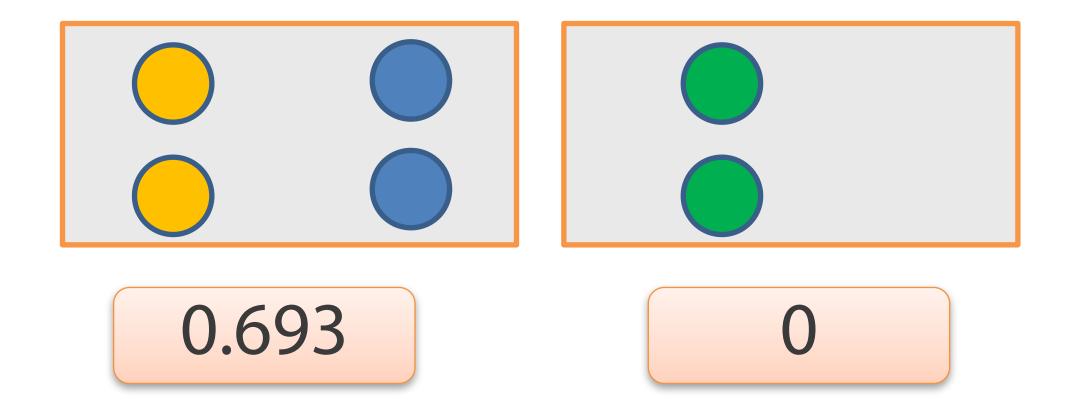
- p = (0.2, 0.2, 0.2, 0.2, 0.2)
  - H = 2.3219
- p = (0.9, 0.05, 0.05, 0, 0)
  - H = 0.5689
- p = (0, 0, 0, 1, 0)
  - H = 0

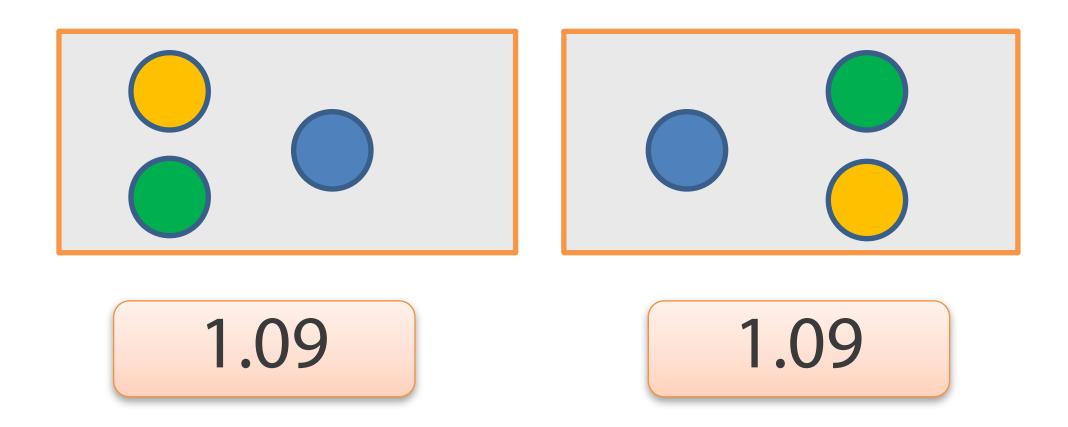
- In classification tasks the number of possible outcomes is the number of classes  $\boldsymbol{K}$
- Probability to be at the class k fraction of objects of class k

$$p_k = \frac{1}{|R|} \sum_{(x_i, y_i) \in R} [y_i = k]$$

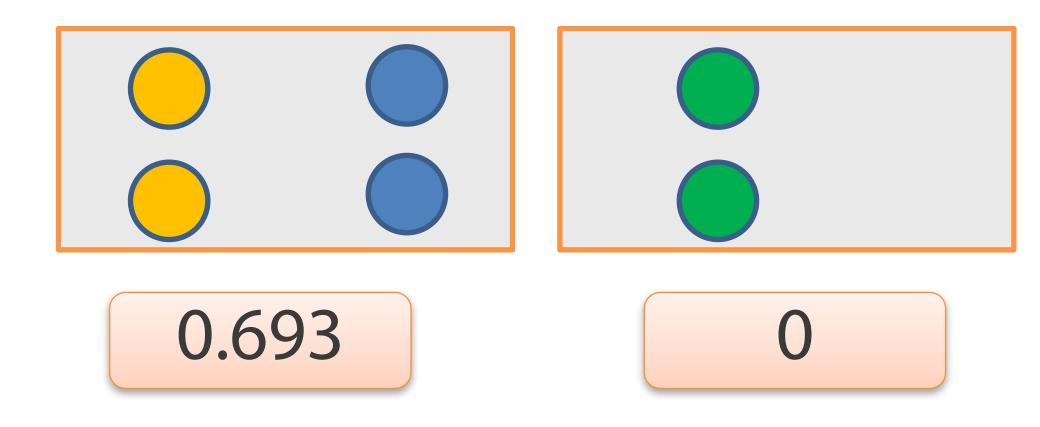
- Zero Entropy there are only objects form one class at the leaf
- Max. Entropy there are equal proportion of objects from each class

# How to Select Between Two Splits?

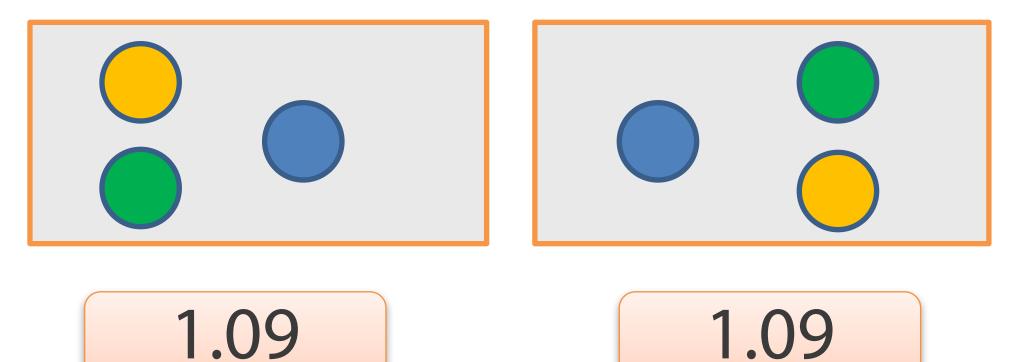




## How to Select Between Two Splits?



- (0.5, 0.5, 0) and (0, 0, 1)
- 0.693 + 0 = 0.693



- (0.33, 0.33, 0.33) and (0.33, 0.33, 0.33)
- 1.09 + 1.09 = 2.18

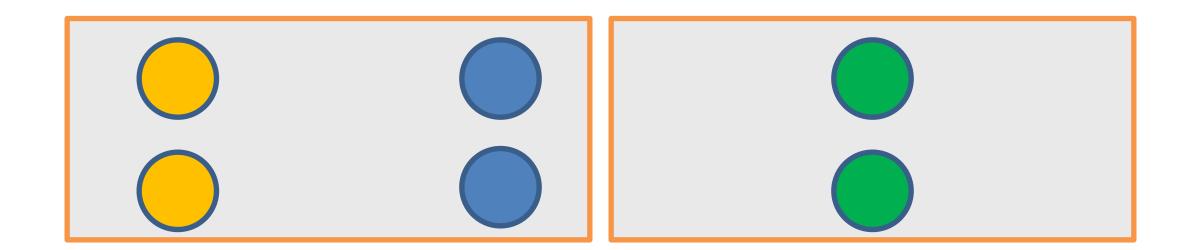
# Summary

 Decision tree could be constructed in a greedy manner from the root node to the leaves

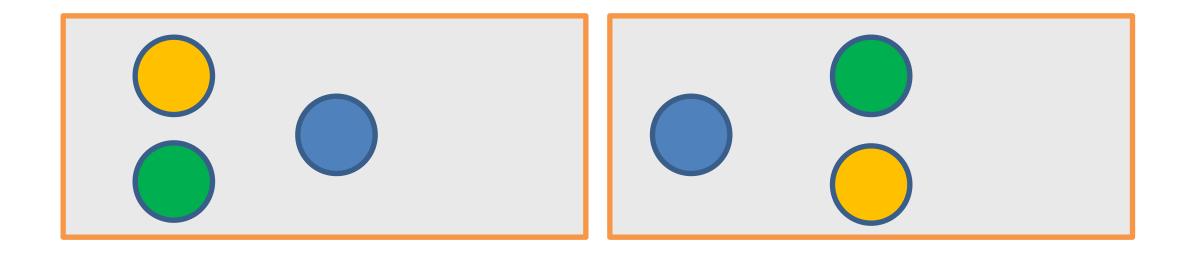
 For classification task we could choose a split, so that it minimizes class diversity at resulting groups

# Measures of Impurity

# How to Choose Between Two Splits?



or



$$H(p_1, ..., p_K) = -\sum_{i=1}^{K} p_i \log_2 p_i$$

Measure of diversity at the node

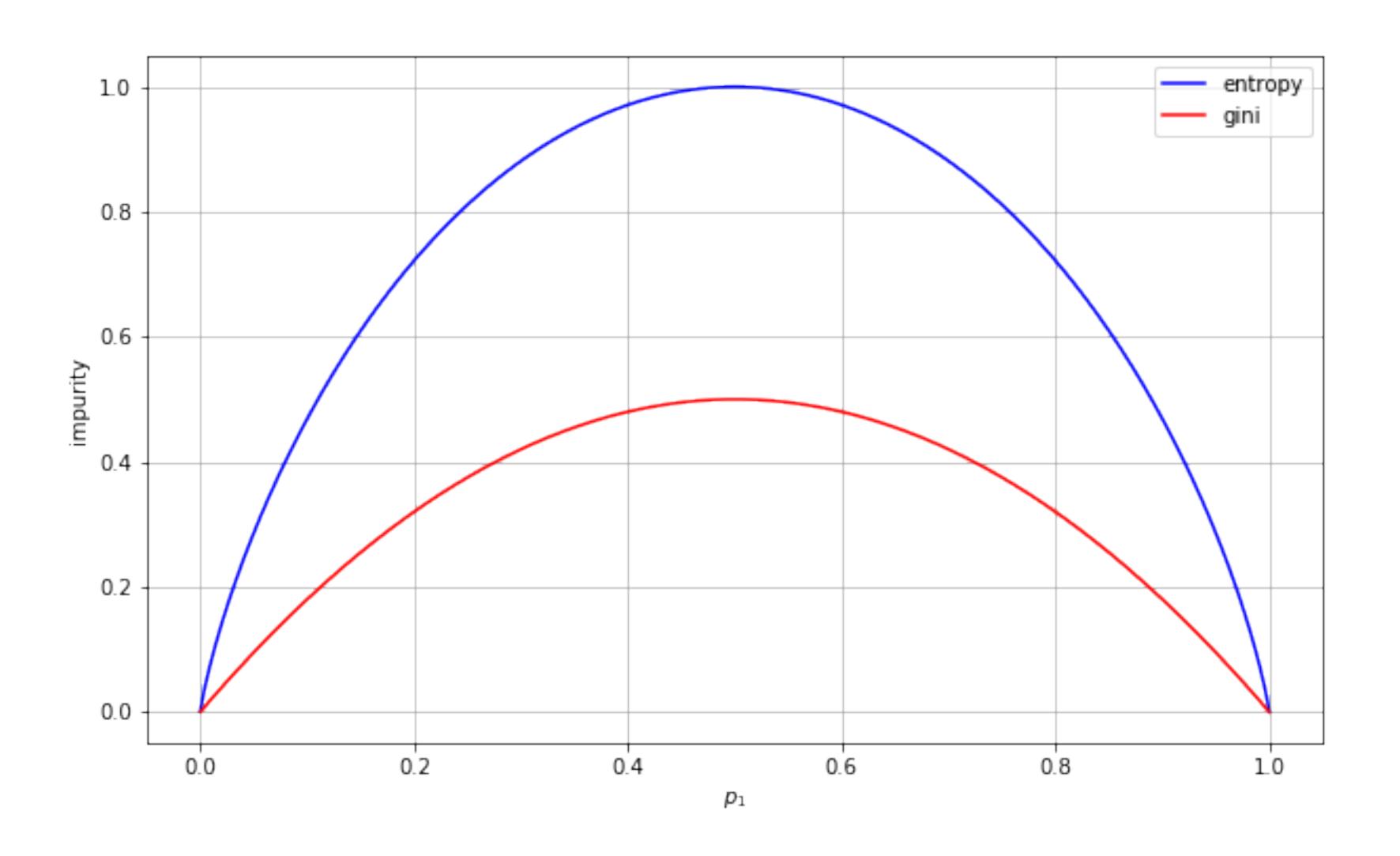
#### Gini Index

$$H(p_1, ..., p_K) = \sum_{i=1}^K p_i (1 - p_i)$$

• Consider a classifier, which outputs class k with probability  $p_k$ 

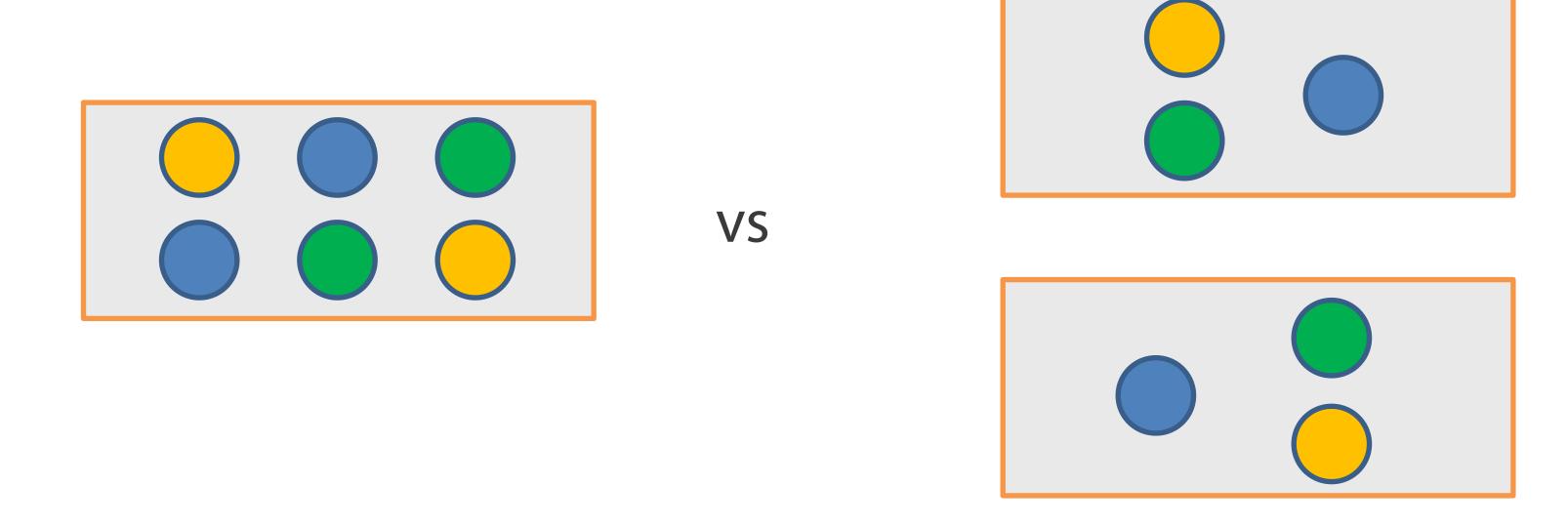
• Gini index is a probability that the object will be classified incorrectly if the class is assigned with probabilities  $p_1, \ldots, p_k$ 

# Gini Index vs Entropy



# **Impurity Criterions**

- How to decide which split is better?
- Compare the impurity before the split (in the initial node R) and in the two nodes after the split ( $R_\ell$  and  $R_r$ )



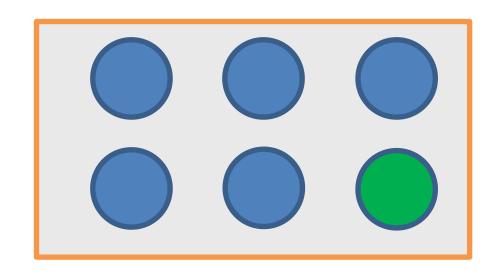
# **Impurity Criterions**

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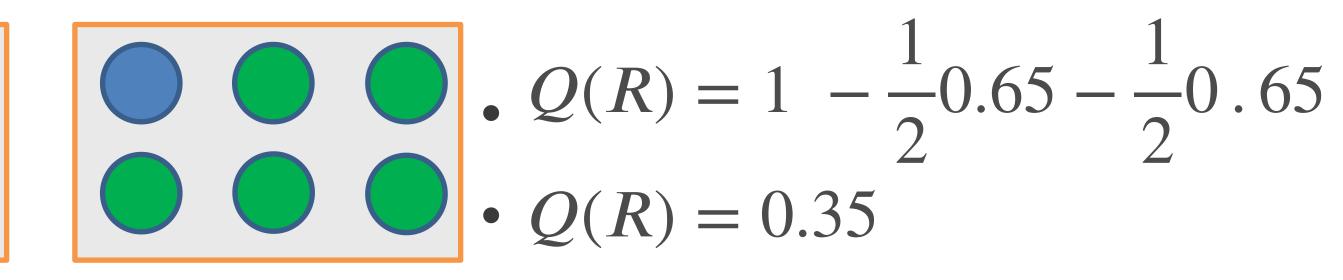
$$Q(R, j, t) = H(R) - \frac{|R_{\ell}|}{|R|} H(R_{\ell}) - \frac{|R_r|}{|R|} H(R_r) \to \max_{j, t}$$

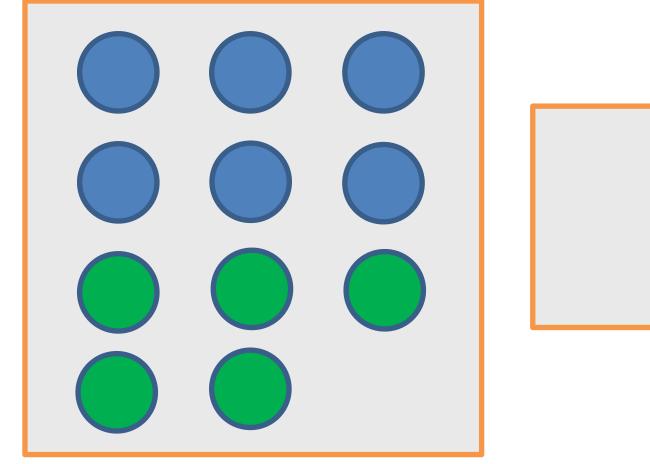
## **How to Compare Two Splits?**

$$H(R) = 1$$



0.65



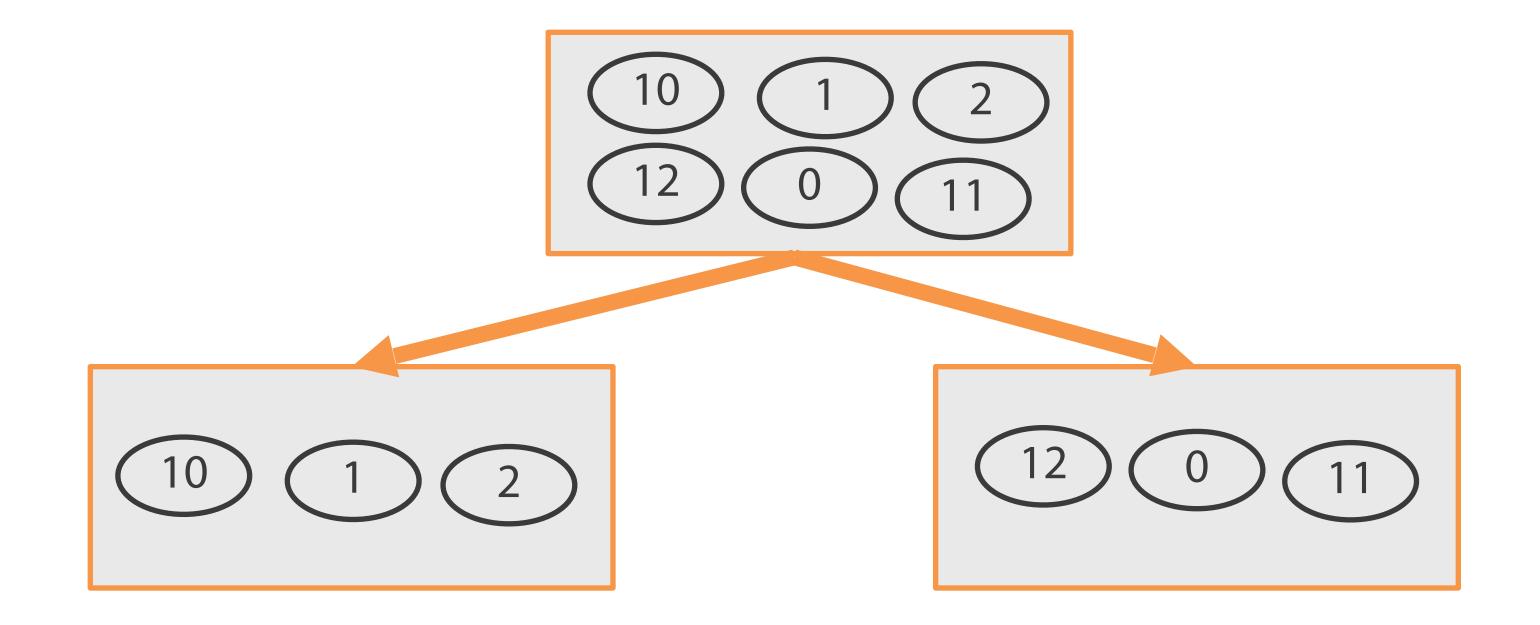


0.994

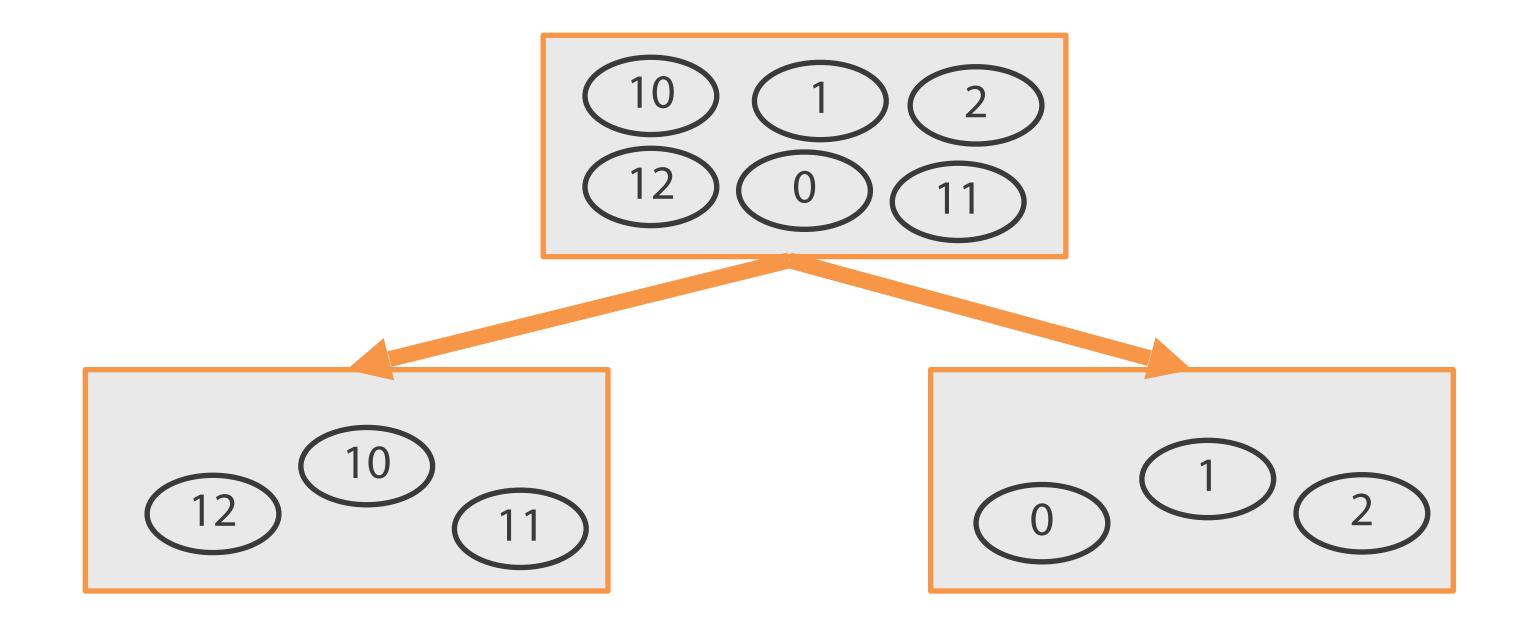
• 
$$Q(R) = 1 - \frac{11}{12}0.994 - \frac{1}{12}0$$
  
•  $Q(R) = 0.088$ 

$$Q(R) = 0.088$$

# **Greedy Construction: Regression**



# **Greedy Construction: Regression**



# Regression Task

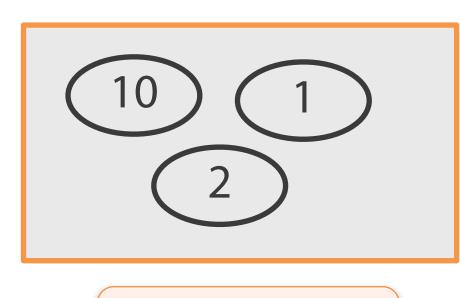
$$H(R) = \frac{1}{|R|} \sum_{(x_i, y_i) \in R} (y_i - y_R)^2$$

$$y_R = \frac{1}{|R|} \sum_{(x_i, y_i) \in R} y_i$$

• So we can measure the variance of answers in the node

# **How to Compare Two Splits?**

$$H(R) = 25.6$$

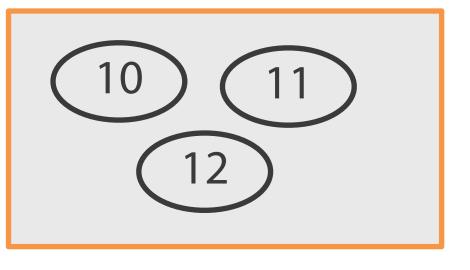


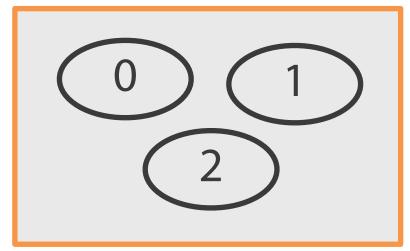
• 
$$Q(R) = 25.6 - \frac{1}{2}16.2 - \frac{1}{2}29.6$$
  
•  $Q(R) = 2.7$ 

• 
$$Q(R) = 2.7$$

16.2

29.6





• 
$$Q(R) = 25.6 - \frac{1}{2}0.7 - \frac{1}{2}0.7$$
  
•  $Q(R) = 24.9$ 

$$Q(R) = 24.9$$

0.7

0.7

### Summary

 We can choose the split, so that it reduces the diversity of answers in the resulting nodes

We use impurity criterion to measure the quality of the split

- There are different criterions that might be used.
   The most popular are:
  - Entropy and Gini for classification
  - Variance for regression

# **Greedy Tree Construction**

#### **How to Construct a Tree?**

- Optimal option:
  - Try all possible trees and select the smallest one

That is too computationally expensive

#### How to Construct a Tree?

• We know how to choose the best split for a given node

• Let's use greedy algorithm

 We start from the root node and will be splitting until some stopping criterion is satisfied

### **Stopping Criterions**

- Restrict the maximal depth
- Restrict the number of leaves
- Fix the minimal number of objects in the node
- Set the minimal decrease in the diversity when splitting
- etc.

- 1. Put the whole dataset into the root:  $R_1 = X$
- 2. Start the tree construction: SplitNode  $(1, R_1)$

SplitNode 
$$(m, R_m)$$

- 1. If stopping criterion is satisfied, then quit
- 2. Find the best split (feature and threshold):

$$j, t = \underset{j,t}{\operatorname{argmax}} Q(R_m, j, t)$$

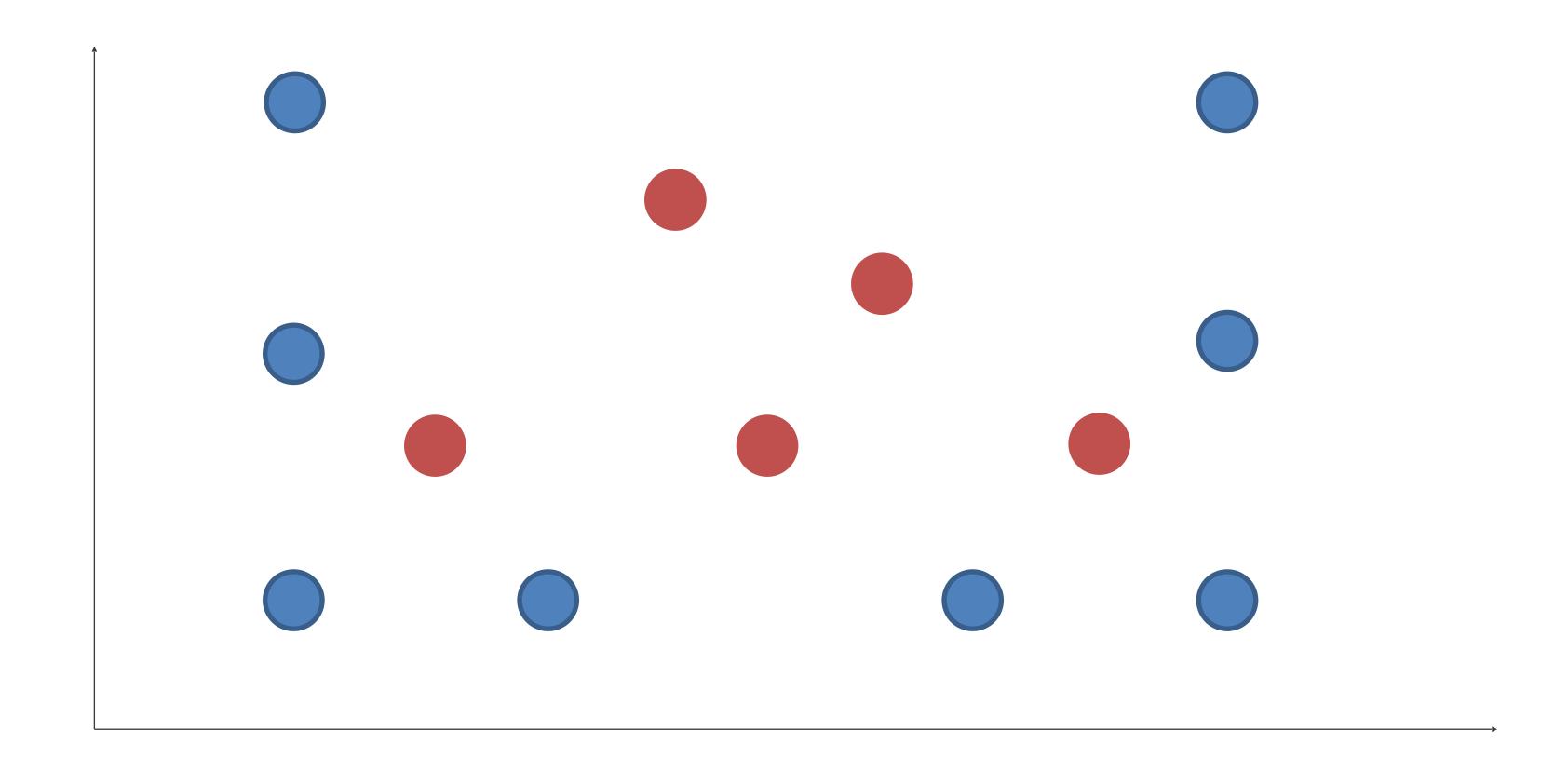
3. Split the objects:

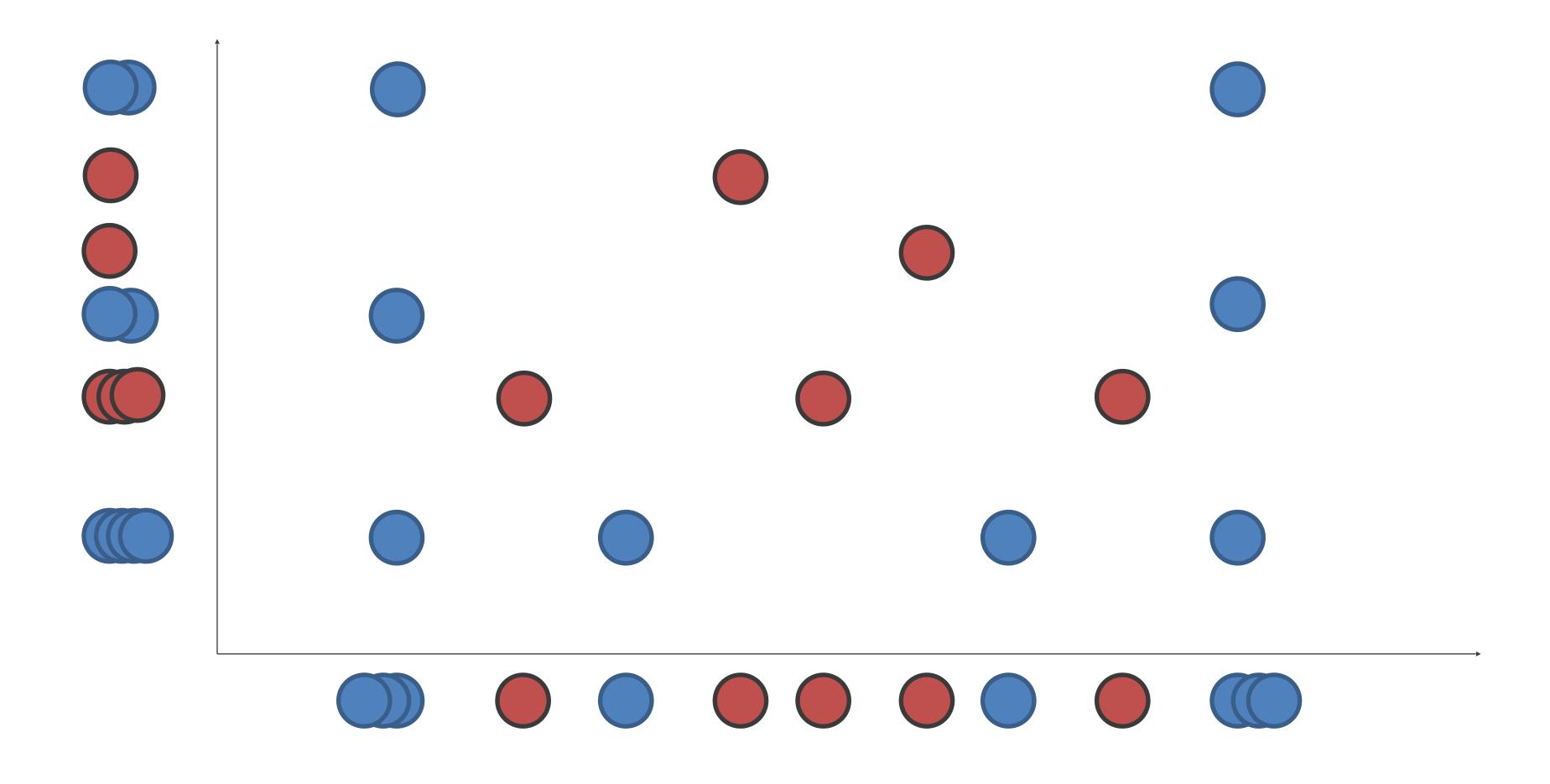
$$R_{\ell} = \left\{ \left\{ \left( x, y \right) \in R_{m} \middle| \left[ x_{j} < t \right] \right\}, \right.$$

$$R_{r} = \left\{ \left\{ \left( x, y \right) \in R_{m} \middle| \left[ x_{j} \ge t \right] \right\} \right\}$$

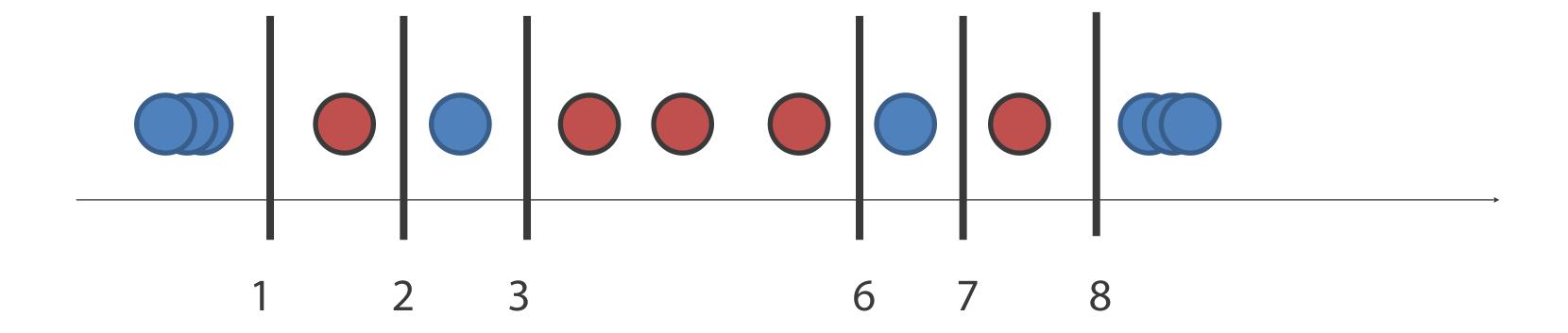
4. Repeat for the child nodes:

SplitNode 
$$(\ell, R_{\ell})$$
 and SplitNode  $(r, R_{r})$ 

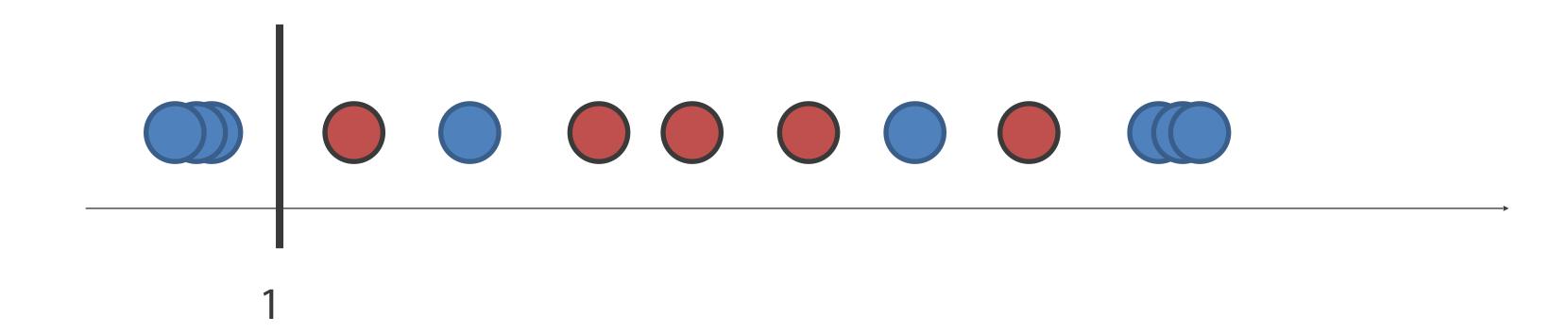




#### Splitting on the first feature



### Splitting on the first feature

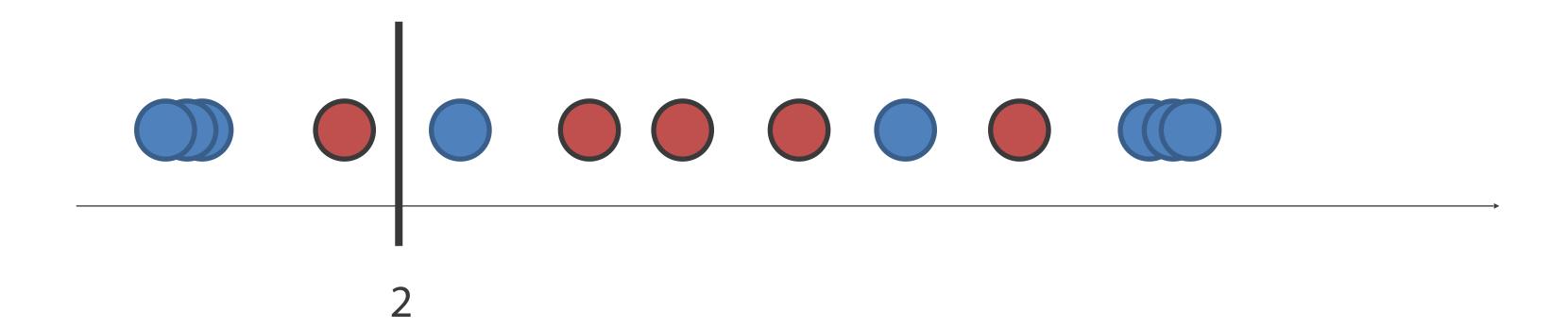


$$(1, 0)$$
  
 $H(p) = 0$ 

$$(1/2, 1/2)$$
  
 $H(p) = 1$ 

$$\frac{3}{13}H(p_l) + \frac{10}{13}H(p_r) = 0.76$$

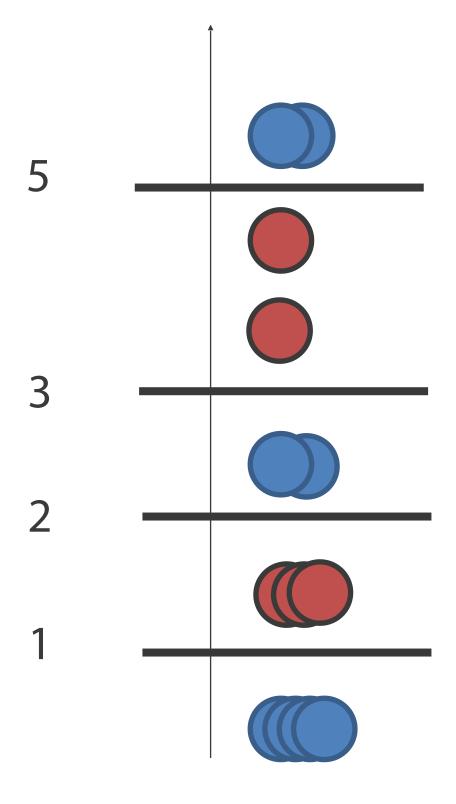
### Splitting on the first feature

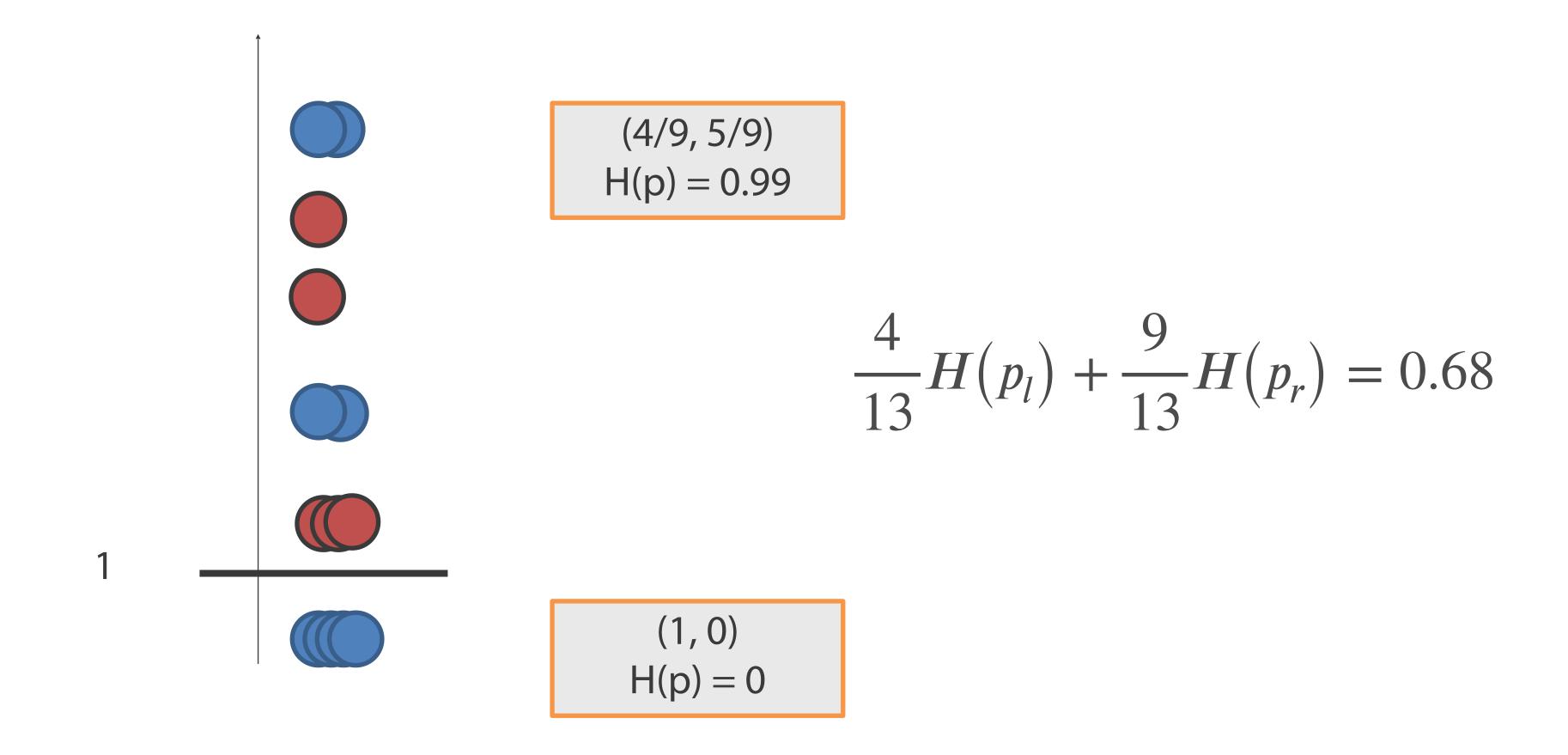


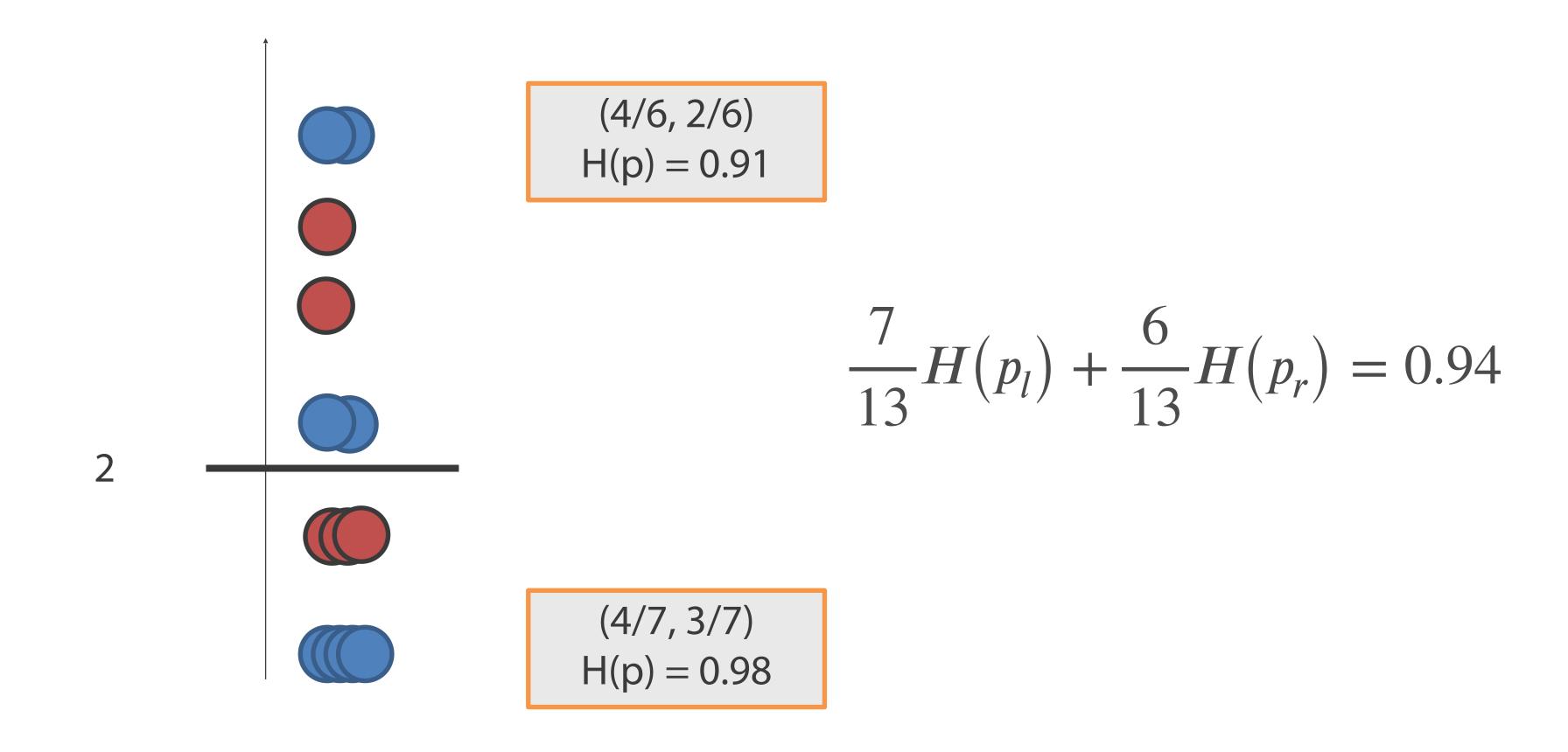
(3/4, 1/4)H(p) = 0.81

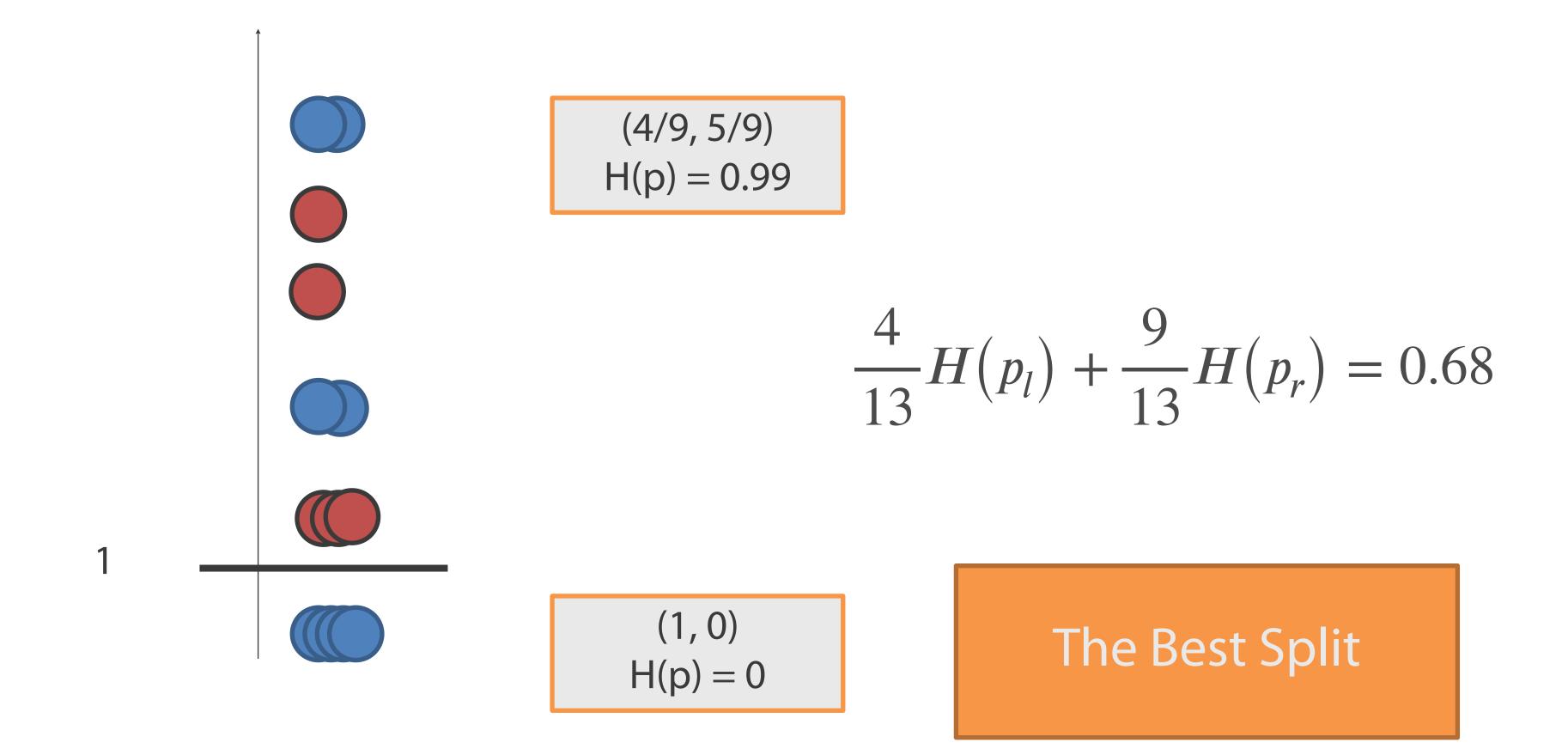
(5/9, 4/9)H(p) = 0.99

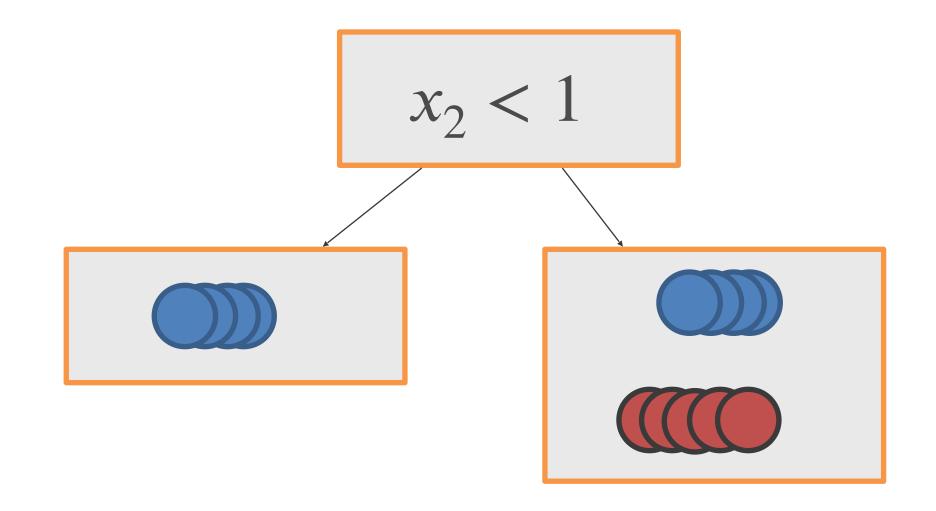
$$\frac{4}{13}H(p_l) + \frac{9}{13}H(p_r) = 0.93$$

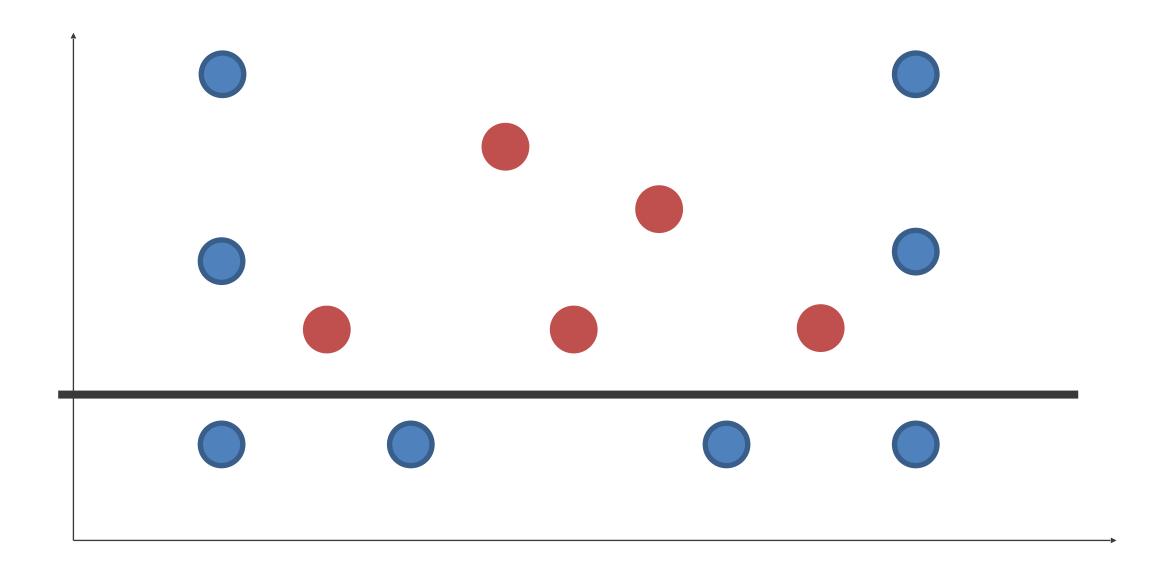


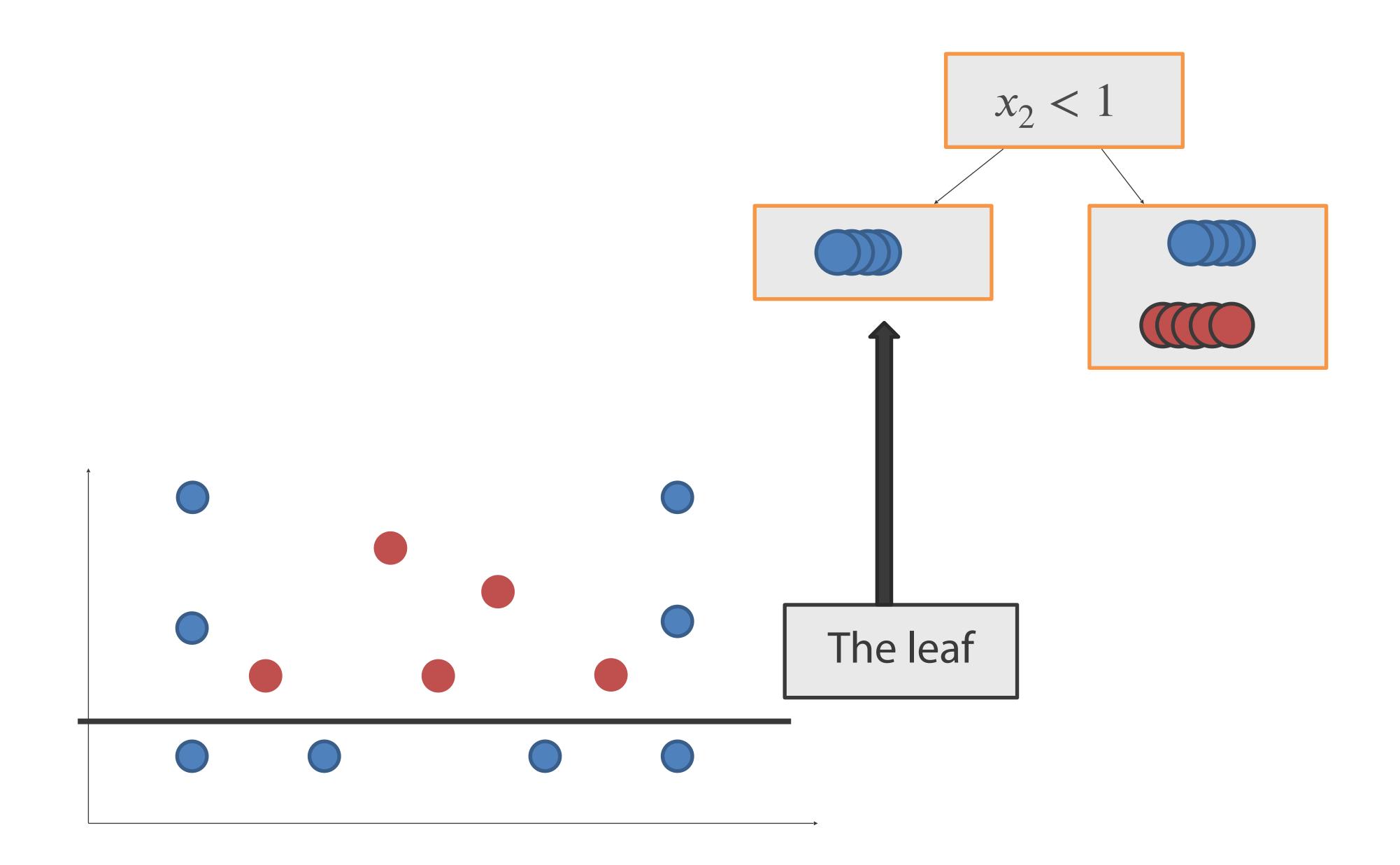


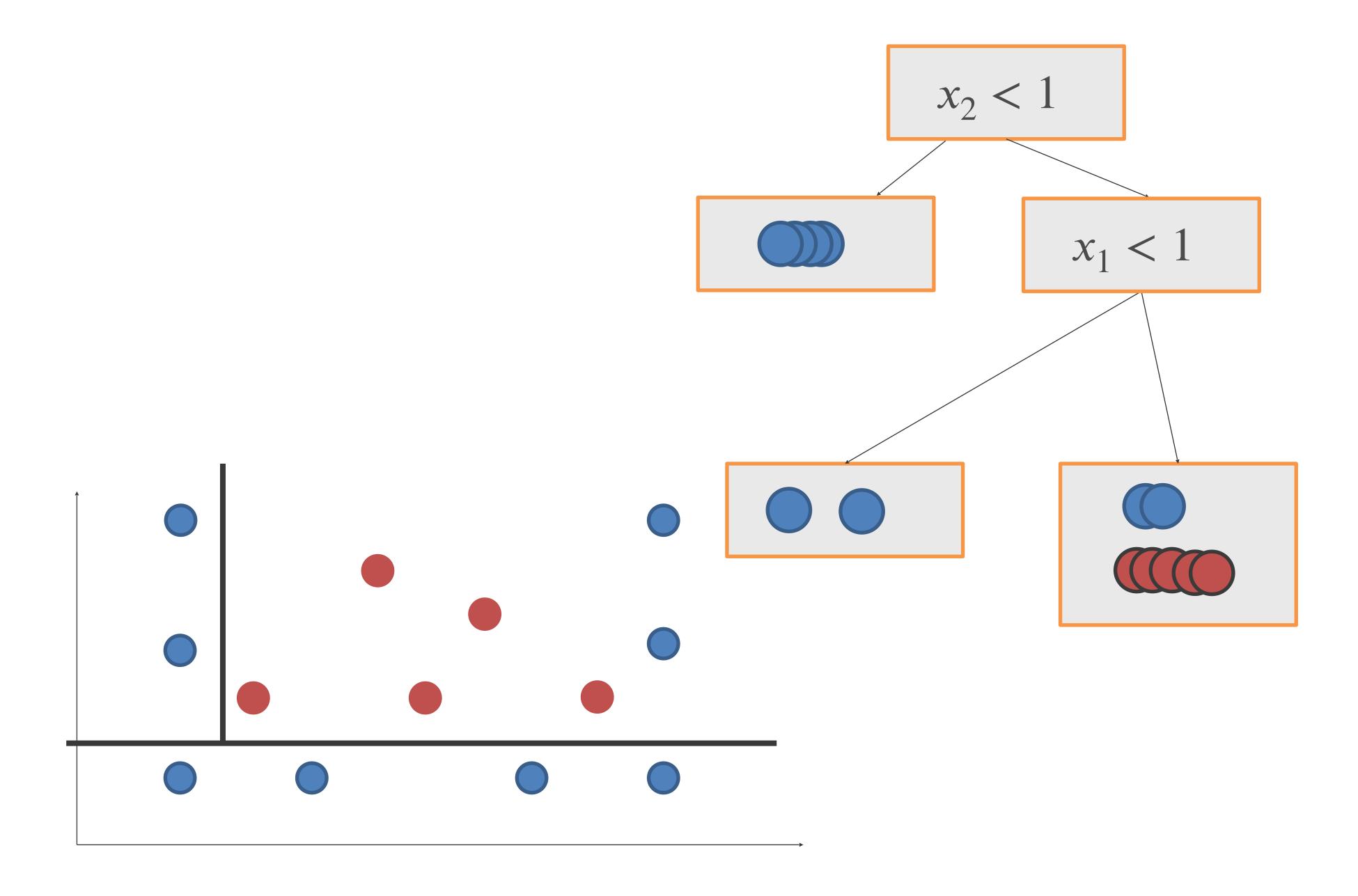


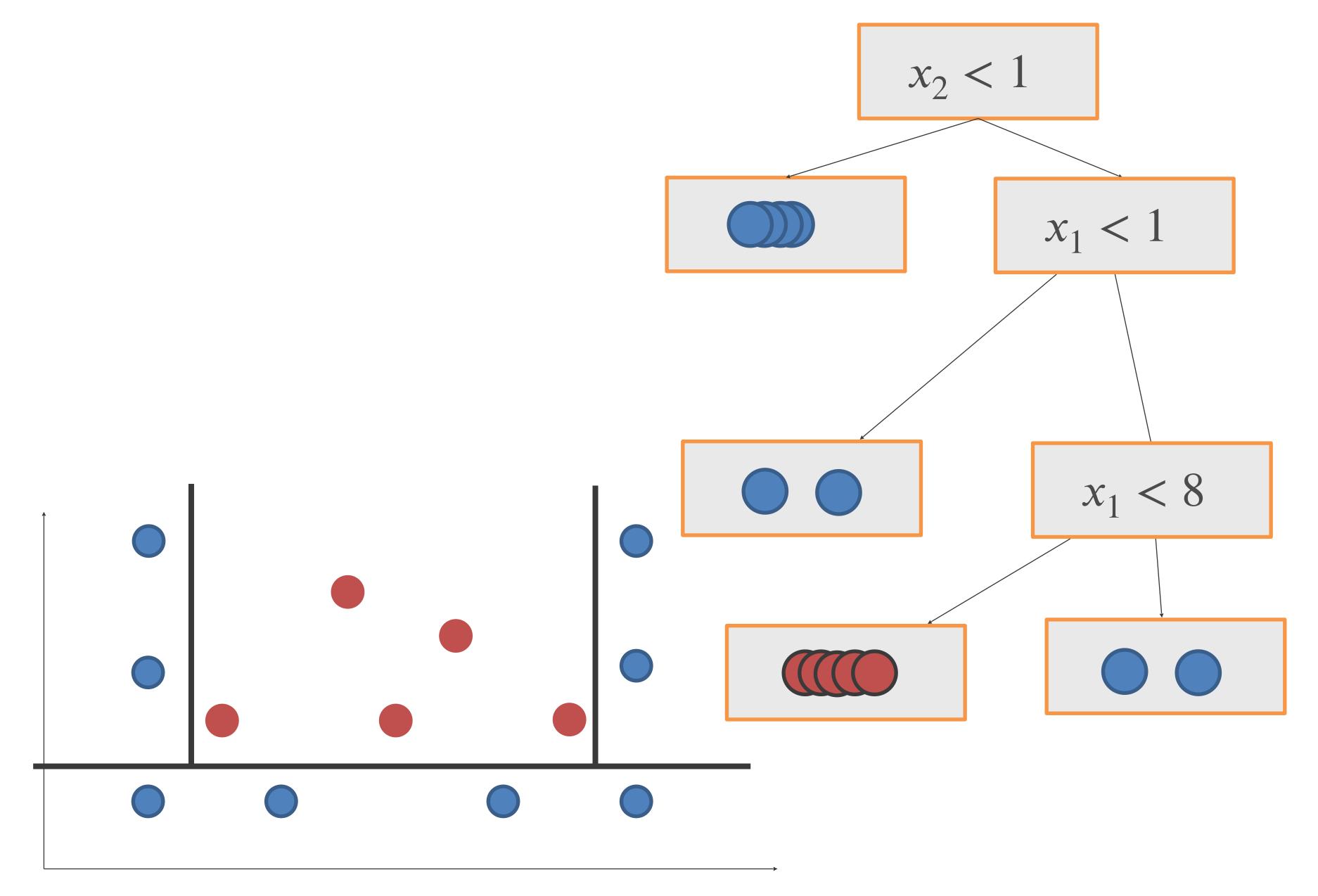












### Summary

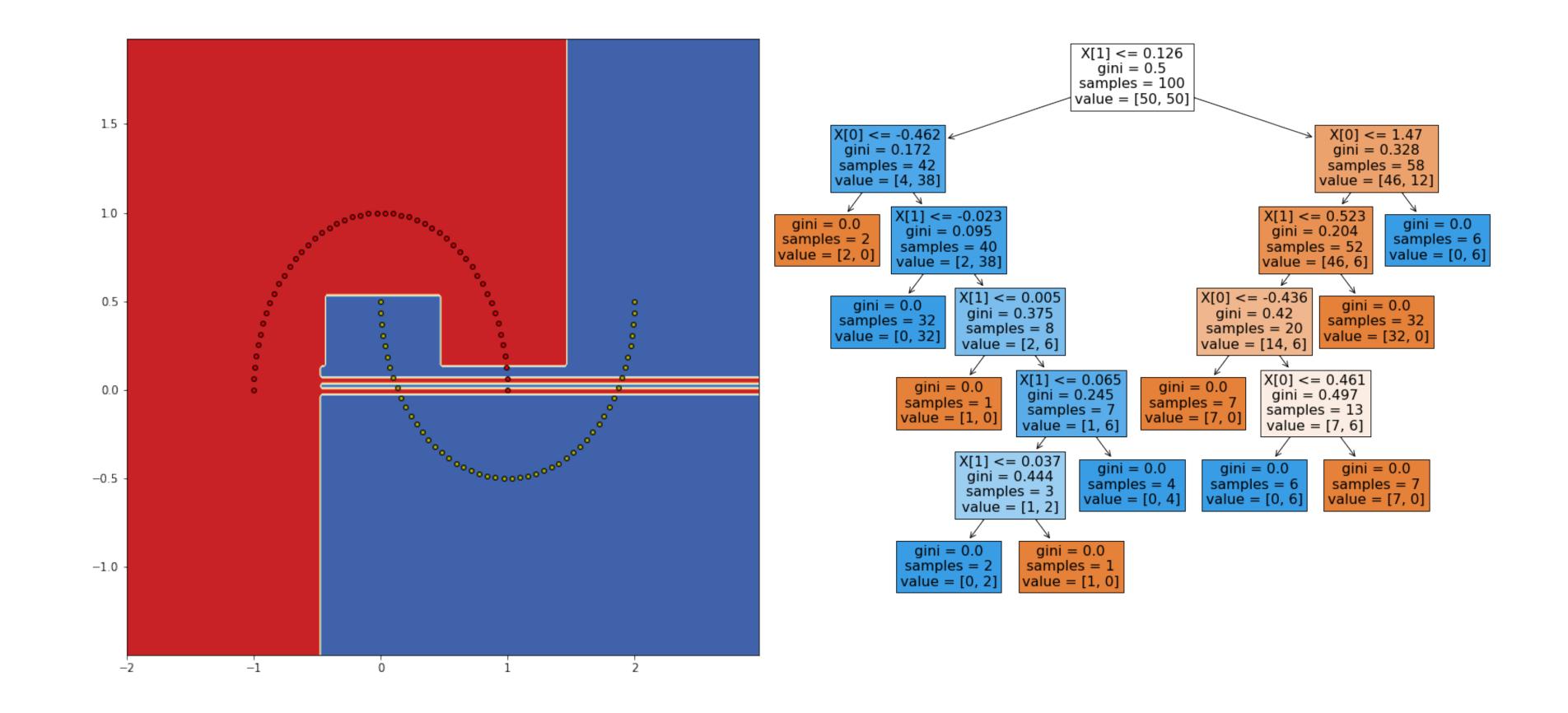
• We use greedy algorithm to construct a tree

At each step we select the best split

• The algorithm is quite complex and requires brute force of all the splits at each step

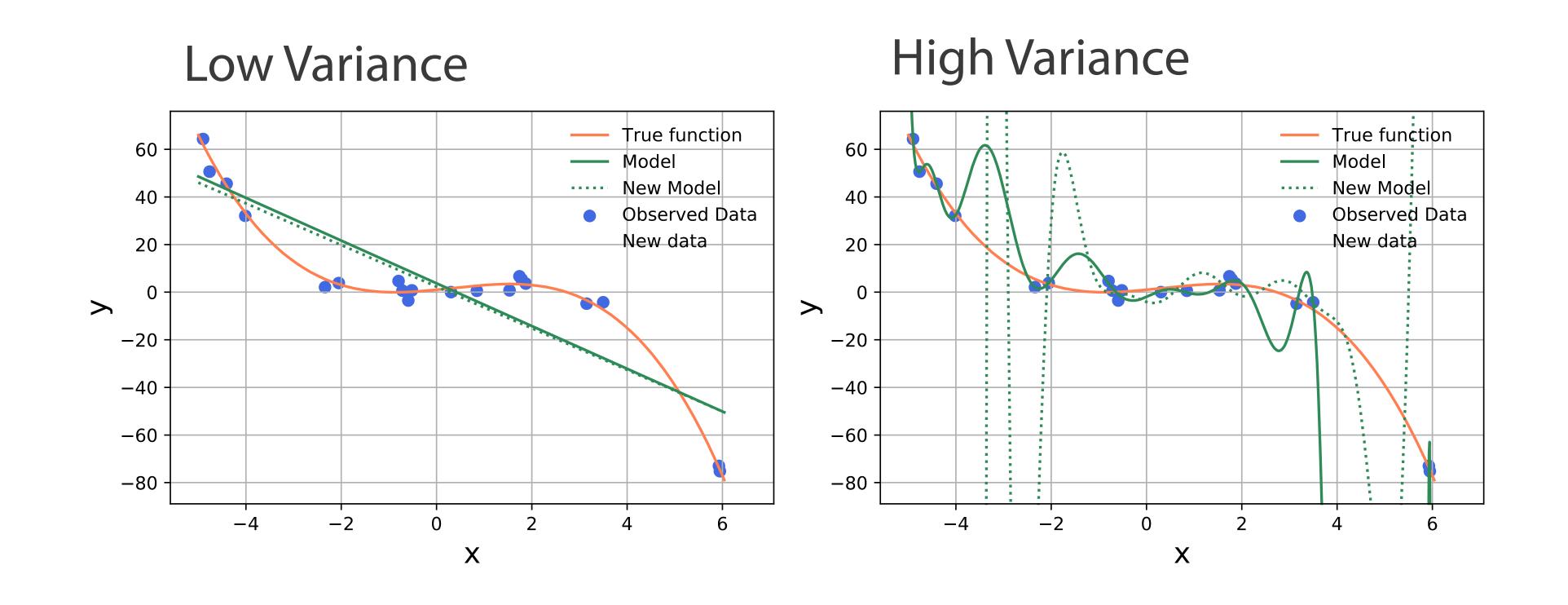
# **High Variance of Decision Trees**

### **Decision Trees and Overfitting**



#### Variance

Variance – sensitivity of the model to the fluctuations in the data

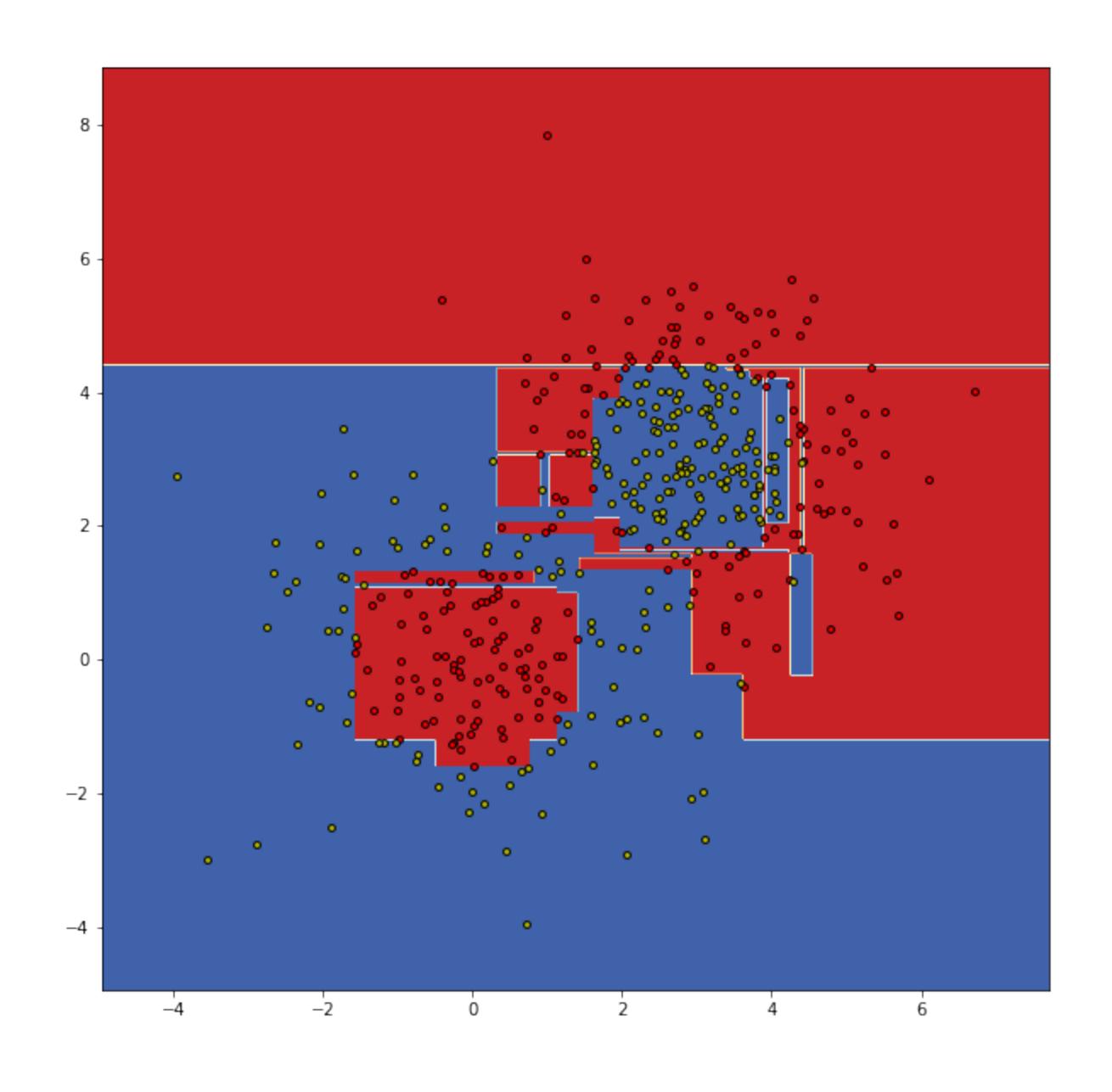


### Stability of a Model

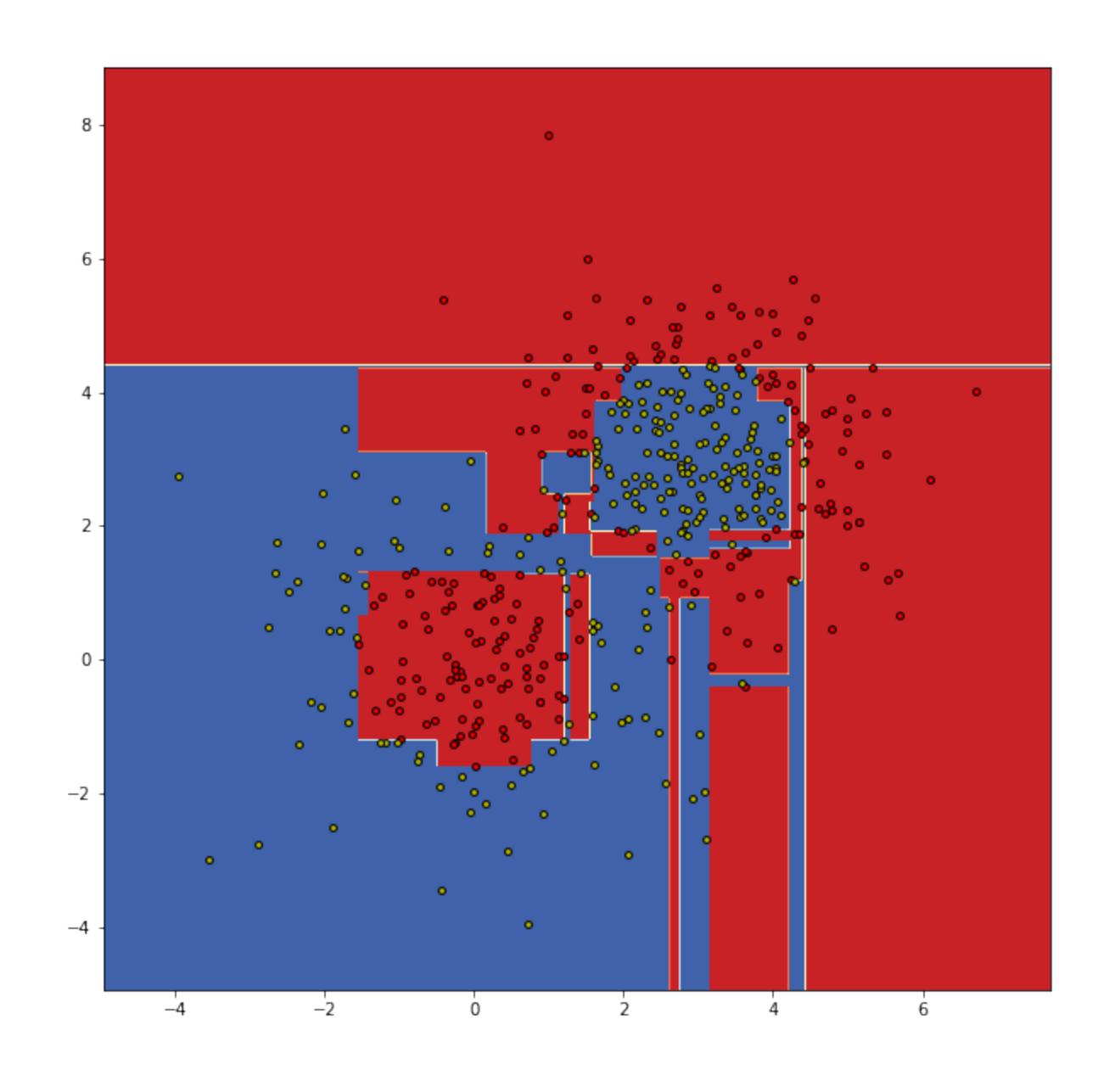
- $X = (x_i, y_i)_{i=1}^{\ell}$  training dataset
- If the model has low variance, we expect is to look similar if we use different subset of the initial training dataset

- $\widetilde{X}$  random subsample, approx. 90% from the initial
- Let us train decision trees on different  $\widetilde{X}$

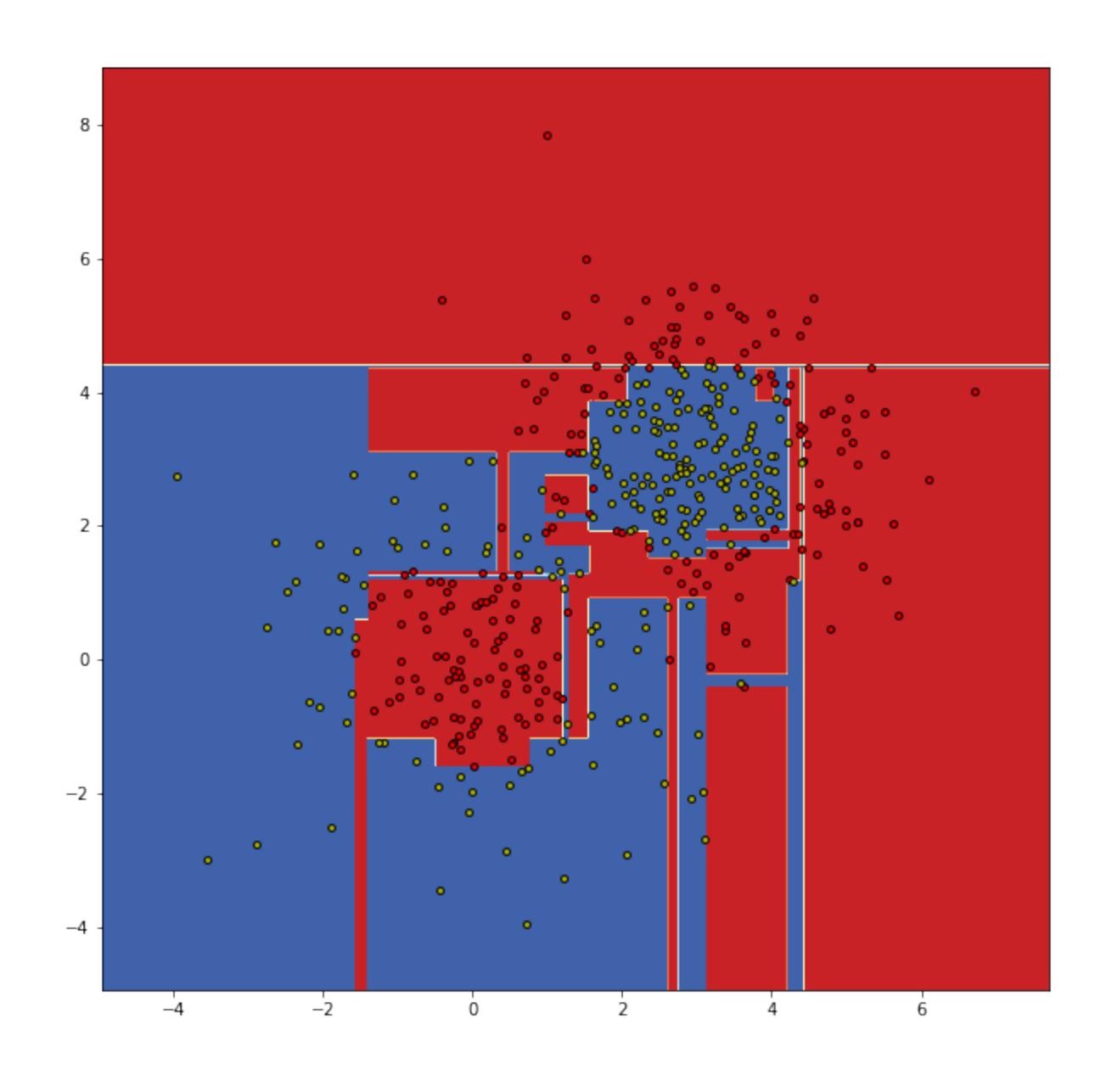
### Training Decision Tree on Subsamples



### Training Decision Tree on Subsamples

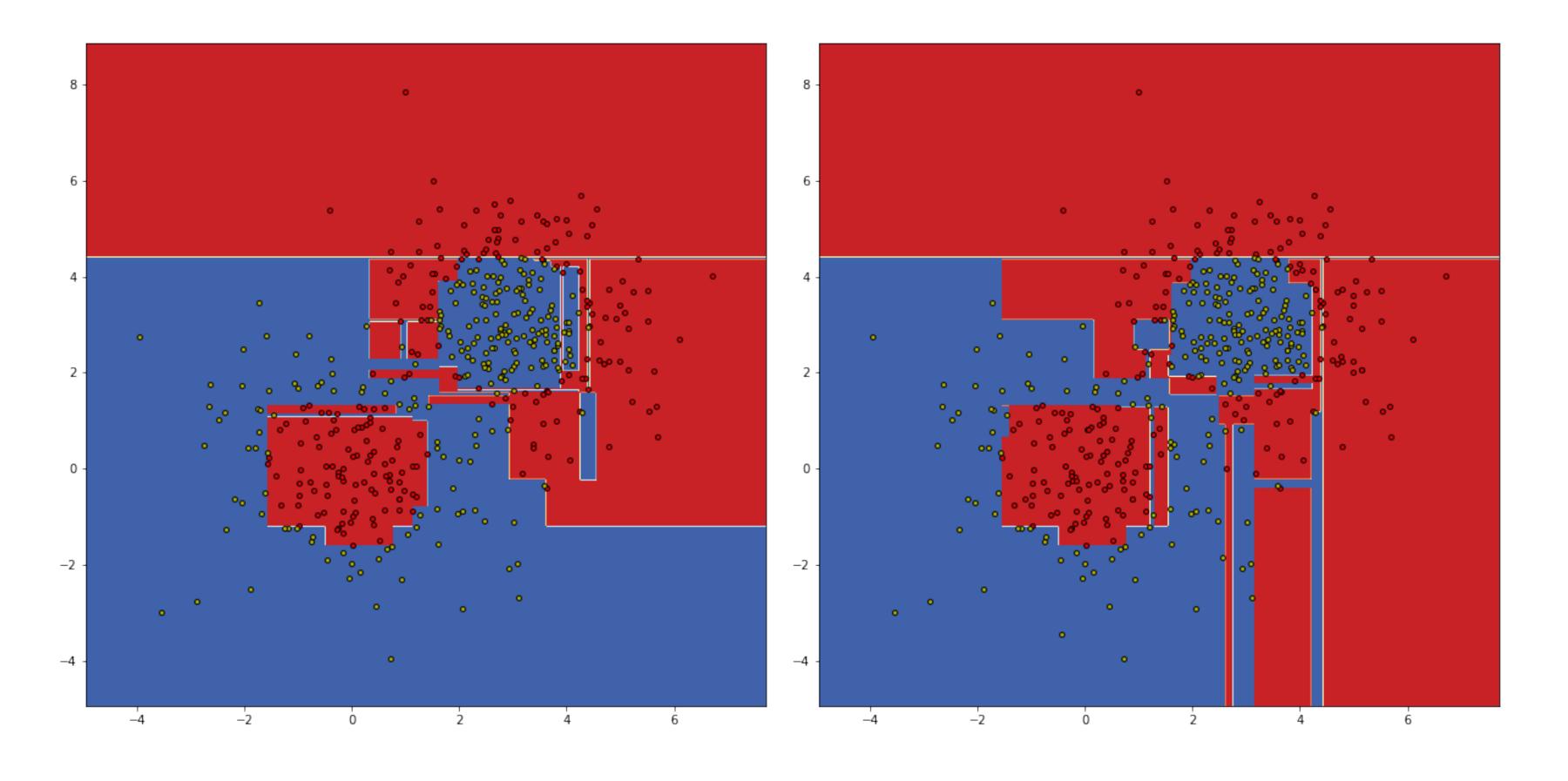


### Training Decision Tree on Subsamples

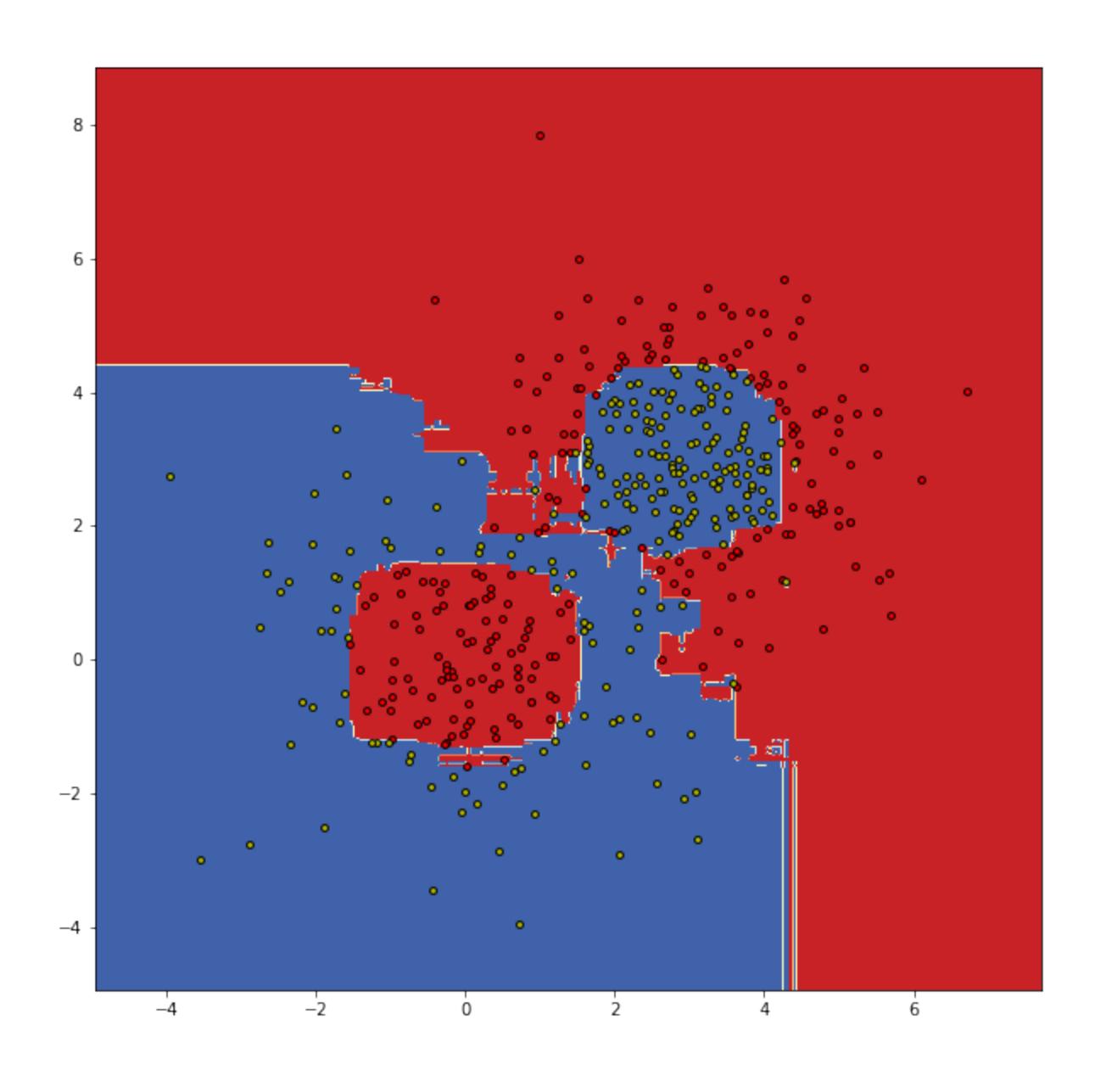


#### **Decision Trees**

- High variance
- Lack of smoothness



### Composition of Models



### Summary

• Decision trees are very sensitive to the changes in the dataset

Prediction surface of decision trees in very non-smooth

• In practice composition of decision trees is used