Lecture 11: Decision Tree and Random Forest Statistical Methods for Data Science

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Today

- Decision trees
 - Classification tree
 - Regression tree
 - Pruning
 - ID3, C4.5, C5.0, CART
- Ensemble methods
 - Bias-variance trade-off
 - Bagging
 - Random forest
- Summary





Learning outcome

- Understand classification and regression decision trees
- Be able to compute the Gini score
- Be able to explain how to compute the split in the CART algorithm
- Be able to explain the random forest algorithm





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- - Bias-variance trade-off
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Classification tree with categorical variables

- Variable (categorical): $\mathbf{x} = [x_1, x_2, x_3]$, $x_1 \in \{\text{chonker, slim}\}, x_2 \in \{\text{exercises, never exercises}\}, x_3 \in \{\text{big-boned, small-boned}\}$
- Target (categorical): $y \in \{\text{give duck a treat, do not give duck a treat}\}$

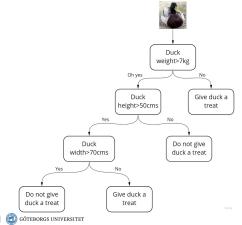






Classification tree with numerical variables

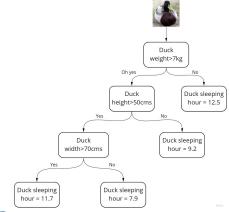
- Variable (numerical): x = [weight, height, width]
- Target (categorical): $y \in \{\text{give duck a treat, do not give duck a treat}\}\$





Regression tree with numerical variables

- Variable (numerical): x = [weight, height, width]
- Target (numerical): y = sleep hours per day







Decision trees Ensemble methods Summary Classification tree Regression tree Pruning ID3, C4.5, C5.0, CART

Classification tree





• Data x: a d dimensional continuous numerical or categorical feature vector $\mathbf{x} = [x_1, \cdots, x_d]$





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$$y = \sum_{m=1}^{M} c_m I_{R_m}(x)$$

where I_{R_m} is the indicater function:

$$I_{R_m}(\mathbf{x}) = \begin{cases} 1, & \text{if } \mathbf{x} \in R_m \\ 0, & \text{otherwise} \end{cases}$$

and $c_m \in \{0, \cdots, C-1\}$ is the class index and R_m are the decision regions (leaf nodes)

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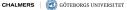
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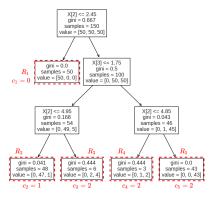
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- Hyperparameter: stopping criteria (e.g. maximum depth, minimum number of data points assigned to each region), hyperparameters for pruning, etc





Visualization - maximum depth 3

Example visualization using scikit-learn on the data set iris

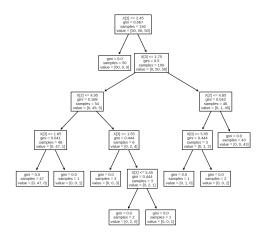


 R_1, \dots, R_5 are the decision regions (leaf nodes)





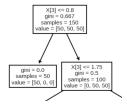
Visualization - maximum depth 5







Parameter estimation (CART)

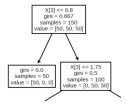


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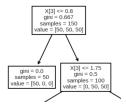


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- Once a tree is **trained**, an unseen data point x will be assigned to a region R_m (leaf node) by the decision tree to identify the class \hat{y}





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 i.e. parameter estimation or training
- Once a tree is **trained**, an unseen data point x will be assigned to a region R_m (leaf node) by the decision tree to identify the class \hat{y}
- To train a tree, we need to estimate the parameters, i.e. all the nodes and a class c_m associated with each leaf node R_m for $m = 1, \dots, M$





Parameter estimation (CART) (cont.)

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- 3. Is it possible to have multi-way splits instead of binary (left/right)?
 - Yes, but they are in general more complex. We focus on binary splits in this lecture





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Parameter estimation (CART) for categorical variables

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- There are alternative impurity measures
 - Gini score (used by CART): $\sum_{b=1}^{C} \hat{p}_b (1 \hat{p}_b)$
 - Entropy: $-\sum_{b=1}^{C} \hat{p}_b \log \hat{p}_b$

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• Previous example: left=[20, 15, 5] and right=[0, 35, 0]; in total,

$$N_{\text{left}} = 20 + 15 + 5 = 40 \text{ and } N_{\text{right}} = 0 + 35 + 0 = 35$$

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$$G_{\text{left}} = \frac{20}{40} (1 - \frac{20}{40}) + \frac{15}{40} (1 - \frac{15}{40}) + \frac{5}{40} (1 - \frac{5}{40}) = 0.59$$

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$$G_{\text{right}} = \frac{0}{35}(1 - \frac{0}{35}) + \frac{35}{35}(1 - \frac{35}{35}) + \frac{0}{35}(1 - \frac{0}{35}) = 0$$



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• Impurity of the split: $G = \frac{N_{\text{left}}}{N_{\text{left}} + N_{\text{right}}} G_{\text{left}} + \frac{N_{\text{right}}}{N_{\text{left}} + N_{\text{right}}} G_{\text{right}}$



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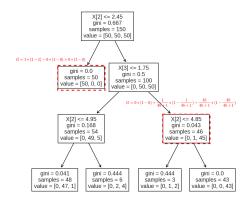
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•
$$G = \frac{40}{40+35}G_{\text{left}} + \frac{35}{40+35}G_{\text{right}} = 0.53 * 0.59 + 0.467 * 0 = 0.3127$$



Exercise: compute the Gini score for each node



Gini:
$$G = \sum_{c=1}^{C} \hat{p}_{mc} (1 - \hat{p}_{mc})$$





Choose the best feature *i* and category *c*

Use the impurity score to choose the best i and c

- Loop over all $i = 1, \dots, d$
- For each i, loop over all categories $c=0,\cdots,C_i-1$

For each choice in the loop, compute the impurity score and choose the i and c that minimizes the impurity.



Pseudocode for categorical variables (CART)

- For each node:
 - Choose the best feature index i
 - Choose the best category c
- 1: for i in $[1, \dots, d]$ do
- 2: **for** c in $[0, \dots, C_i 1]$ **do**
- 3: Compute the Gini score
- 4: end for
- 5: end for
- 6: Find the best i and c corresponding to the smallest Gini score

where C is the number of classes and C_i is the number of categories for feature x_i

· After the tree is constructed,

$$\hat{c}_m = \arg\max_{c} N_m^c, \ \forall c \in \{0, \cdots, C-1\}$$

where N_m^c is the number of data points that are 1) from class c and 2) assigned to region (leaf node) R_m

Example (C=3): in region 2, if $[N_2^0, N_2^1, N_2^2] = [10, 25, 0]$, then $\hat{c}_2 = 1$ since 25 is the largest count.





Similar to categorical variables with minor modifications

- Start with the root node that contains the entire training set
- Grow the tree by splitting each node until a stopping criterion is met, e.g.
 maximum depth, minimum number of data assigned to a leaf node, etc; this is a
 recursive procedure
- This split will create two child nodes (groups): left and right
- Each child node will contain a (sub) training set
 - $\mathcal{I}_{left} \leftarrow \{l \mid x_i^l \leq t\}$: all indices assigned to the left child node
 - $\mathcal{I}_{right} \leftarrow \{r \mid x_i^r > t\}$: all indices assigned to the right child node
- In order to split a node into the left and right child nodes, we need to estimate
 - The best feature x_i for the split
 - ullet The splitting threshold t (modification)

- To estimate x_i and t (modification)
 - Loop over all $i = 1, \dots, d$
 - For each feature index i, let $t \leftarrow x_i^j$, loop over all data in the (sub) training sample in the node





Pseudocode for numerical variables (CART)

- For each node:
 - Choose the best feature index i
 - Choose the best threshold t
- 1: for i in $[1, \dots, d]$ do
- for j in $[1, \dots, N_{node}]$ do
- $t \leftarrow x^{j}$
- Compute the Gini score
- end for
- 6: end for
- 7: Find the best i and t corresponding to the smallest Gini score

where N_{node} is the number of training data points in the current node

After the tree is constructed.

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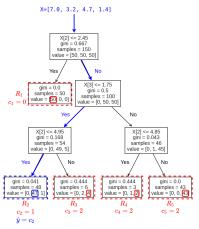
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Prediction with a trained decision tree

For a new data point X = [7.0, 3.2, 4.7, 1.4], $\hat{y} = \sum_{m=1}^{5} c_m I_{R_i}(x)$







Regression tree





$$y = g(x; \theta \mid h)$$



Modeling for regression

$$y = g(x; \theta \mid h)$$

Regression:



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- Parameter estimation: supervised learning





Regression decision tree

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where I_{R_i} is the indicater function:

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- Parameter estimation: the classification and regression tree (CART) algorithm



Parameter estimation

- Similar to parameter estimation for classification trees
- The difference is the score being calculated
- Instead of using the impurity (e.g. Gini score), which is defined for classification, we use the sum of squared error instead





Pseudocode for regression tree (CART)

- For each node:
 - Choose the best feature i
 - Choose the best threshold t
- 1: for i in $[1, \dots, d]$ do
- for j in $[1, \dots, N_{node}]$ do
- $\mathcal{I}_{left} \leftarrow \{I \mid x_i^I \leq t\}$: all indices assigned to the left child node 4:
- $\mathcal{I}_{right} \leftarrow \{r \mid x_i^r > t\}$: all indices assigned to the right child node
- Let $a_{\text{left}} \leftarrow \bar{y}_l$ and $a_{\text{right}} \leftarrow \bar{y}_r$, where \bar{y}_l and \bar{y}_r are the sample means of the left and right child nodes, respectively.
- Compute $s = \sum_{l \in \mathcal{I}_{left}} (y_l a_{left})^2 + \sum_{r \in \mathcal{I}_{sink}} (y_r a_{right})^2$
- end for
- 9: end for
- 10: Find the best i and t corresponding to the lowest score s

where N_{node} is the number of training data points in the current node

After the tree is constructed,

$$\hat{a}_m = \frac{1}{N_m} \sum_{i=1}^{N_m} y_{m_i}$$

where N_m is the number of data points assigned to region (leaf node) R_m and y_{m_i} denotes labels of data points assigned to R_m





Pruning





Overfitting

Without modification, decision trees suffer from overfitting





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 This can be done using, e.g. cross-validation at each node split during the tree construction process





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- Pre-pruning (top-down, early stopping):
 This can be done using, e.g. cross-validation at each node split during the tree construction process
- Post-pruning (bottom-up, remove leaves after the tree is constructed):
 - This can be done by, e.g. removing leaves one by one until the optimal tree is found
 - The **optimality** is defined by a loss function *L*, e.g.

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Typically, there are additional hyperparameters (e.g. α) to tune using, e.g. cross-validation; a commonly used technique is cost complexity pruning, where complexity is defined as the number of leaf nodes.





Classification tree Regression tree Pruning ID3, C4.5, C5.0, CART

ID3, C4.5, C5.0, CART





They are parameter estimation algorithms in the literature.

Algorithm	ID3	C4.5	C5.0	CART
Variable type	Categorical	Continuous and categorical	Continuous and categorical	Continuous and categorical
Target type	Categorical	Categorical	Categorical	Continuous and categorical
Splitting score	Information gain	Information gain	Information gain	Gini impurity
Computational efficiency	Low	Okay	High	Good





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- In summary, NO BIGGY! Read the documentation (and the literature) before using them
- For example, the scikit-learn documentation (December 2021): https://scikit-learn.org/stable/modules/tree.html
- The most important thing is to understand the different components in the algorithms, e.g. splitting score, splitting criteria, maximum tree depth, etc





Summary

- Pros:
 - Interpretable, visualizable, intuitive
 - Do not require preprocessing
 - Handle categorical and continuous variables
 - Fairly simple to implement
- Cons:
 - High variance, overfitting
 - Not robust





Today

- Decision trees
 - Classification tree
 - Regression tree
 - Pruning
 - ID3, C4.5, C5.0, CART
- 2 Ensemble methods
 - Bias-variance trade-off
 - Bagging
 - Random forest
- Summary





Bias-variance trade-off Bagging Random forest

Bias-variance trade-off





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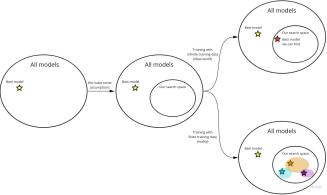
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Bias-variance trade-off (cont.)

Complex models tend to have low bias and high variance





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- Complex models tend to have low bias and high variance
- Simple models tend to have high bias and low variance





Bias-variance trade-off (cont.)

- Complex models tend to have low bias and high variance
- Simple models tend to have high bias and low variance
- Ensemble methods combine multiple models to produce a better result - with reduced bias or variance





Bias-variance trade-off Bagging Random forest





• Purpose: to reduce variance





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- General idea behind bagging





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 - Note: this variance cannot be reduced to zero



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a bootstrapped sample could be

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- Step 2: Build a tree based on the bootstrapped sample
- Step 3: Repeat step 1 and 2 B times, which results in B trees





Bagging: prediction

Aggregate the predictions from all B trees

• Regression tree (sample mean):

$$\hat{y} = \frac{1}{B} \sum_{b=1}^{B} \hat{y}_b^*$$

Classification tree (majority vote):

$$\hat{y} = \arg\max_{c} \operatorname{count}(\hat{y}_b^* == c)$$



Bias-variance trade-off Bagging Random forest

Random forest





Random forest

Similar to bagging with one addition +

- Step 1: Create a bootstrapped sample
- Step 2: Build a tree based on the bootstrapped sample
 - + Randomly choose $\tilde{d} \approx \sqrt{d}$ at each node and only use \tilde{d} features for the split (decorrelation)
- Step 3: Repeat step 1 and 2 B times, which results in B trees

Compared to decision trees, random forest results in significantly lower variance with slightly increased bias; less interpretable





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Summary

- Data types and data containers
- Descriptive data analysis: descriptive statistics, visualization
- Probability distributions, events, random variables, PMF, PDF, parameters
- CDF. Q-Q plot, how to compare two distributions (data vs theoretical, data vs data)
- Modeling
- Parameter estimation: maximum likelihood estimation (MLE) and maximum a posteriori estimation (MAP)
- Classification, multinomial naive Bayes classifier, Gaussian naive Bayes classifier
- Central limit theorem, interval estimation.
- Hypothesis tests, comparison of two classifiers
- Clustering, cluster tendency, k-means
- SSE and Silhouette score for cluster evaluation, one dimensional Gaussian Mixture Models, AIC/BIC, the EM algorithm, clustering validation
- Decision tree for classification and regression, random forest





Stay safe, happy, curious and enthusiastic!





