Chapter ML:VI

VI. Decision Trees

- Decision Trees Basics
- □ Impurity Functions
- □ Decision Tree Algorithms
- Decision Tree Pruning

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ID3 Algorithm [Quinlan 1986] [CART Algorithm]

Setting:

- \Box X is a multiset of feature vectors.
- \Box *C* is a set of classes.
- $D = \{(\mathbf{x}_1, c_1), \dots, (\mathbf{x}_n, c_n)\} \subseteq X \times C$ is a multiset of examples.

Learning task:

 \Box Fit D using a decision tree T.

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Learning task:

 \Box Fit *D* using a decision tree *T*.

Characteristics of the ID3 algorithm:

1. Each splitting is based on one nominal feature and considers its complete domain. Splitting based on feature A with domain $dom(A) = \{a_1, \ldots, a_m\}$:

$$X = \{ \mathbf{x} \in X : \mathbf{x}|_A = a_1 \} \cup \ldots \cup \{ \mathbf{x} \in X : \mathbf{x}|_A = a_m \}$$

2. Splitting criterion is information gain.

ID3 Algorithm (continued) [Mitchell 1997 version] [algorithm template]

ID3(D, Features)

- 1. Create a node t for the tree.
- Label t with the most common class in D.
- 3. If all examples in D have the same class, return the single-node tree t.
- 4. If Features is empty, return the single-node tree t.

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ID3 Algorithm (continued) [Mitchell 1997 version] [algorithm template]

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- 2. Label t with the most common class in D.
- 3. If all examples in D have the same class, return the single-node tree t.
- 4. If Features is empty, return the single-node tree t.

Otherwise:

- Let A* be the feature from Features that best classifies examples in D.
 Assign t the decision feature A*.
- 6. For each possible value "a" in dom(A*) do:
 - \Box Add a new tree branch below t, corresponding to the test $A^* = a^*$.
 - \Box Let D_a be the subset of D that has value "a" for A*.
 - If D_a is empty: Then add a leaf node with the label of the most common class in D. Else add the subtree ID3(D_a , Features \ {A*}).

Return t.

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ID3 Algorithm (continued) [algorithm template]

ID3(*D*, *Features*)

```
1. t = createNode()
```

```
2. label(t) = mostCommonClass(D)
```

- 3. IF $\forall (\mathbf{x}, c) \in D : c = label(t)$ THEN return(t) ENDIF // D is pure.
- 4. IF *Features* = \emptyset THEN *return*(t) ENDIF // We are running out of features.
- 5.
- 6.

7.

ID3 Algorithm (continued) [algorithm template]

ID3(*D*, *Features*)

- 1. t = createNode()
- 2. label(t) = mostCommonClass(D)
- 3. IF $\forall (\mathbf{x}, c) \in D : c = label(t)$ THEN return(t) ENDIF // D is pure.
- 4. IF *Features* = \emptyset THEN *return*(t) ENDIF // We are running out of features.
- 5. $A^* = \operatorname{argmax}_{A \in Features}(\operatorname{informationGain}(D, A))$

6.

7.

7 . return(t)

ID3 Algorithm (continued) [algorithm template]

```
ID3(D, Features)
  1. t = createNode()
  2. label(t) = mostCommonClass(D)
  3. IF \forall (\mathbf{x}, c) \in D : c = label(t) THEN return(t) ENDIF // D is pure.
      IF Features = \emptyset THEN return(t) ENDIF // We are running out of features.
  5. A^* = \operatorname{argmax}_{A \in Features}(\operatorname{informationGain}(D, A))
  6. FOREACH a \in dom(A^*) DO
          D_a = \{(\mathbf{x}, c) \in D : \mathbf{x}|_{A^*} = a\}
          IF D_a = \emptyset THEN
          ELSE
             createEdge(t, a, ID3(D_a, Features \setminus \{A^*\}))
          ENDIF
       ENDDO
```

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ID3 Algorithm (continued) [algorithm template]

```
ID3(D, Features)
  1. t = createNode()
  2. label(t) = mostCommonClass(D)
  3. IF \forall (\mathbf{x}, c) \in D : c = label(t) THEN return(t) ENDIF // D is pure.
      IF Features = \emptyset THEN return(t) ENDIF // We are running out of features.
  5. A^* = \operatorname{argmax}_{A \in Features}(\operatorname{informationGain}(D, A))
  6. FOREACH a \in dom(A^*) DO
         D_a = \{(\mathbf{x}, c) \in D : \mathbf{x}|_{A^*} = a\}
          IF D_a = \emptyset THEN // We are running out of data.
            t' = createNode()
            label(t') = label(t)
            createEdge(t, a, t')
         ELSE
            createEdge(t, a, ID3(D_a, Features \setminus \{A^*\}))
         ENDIF
       ENDDO
  7 . return(t)
```

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Remarks:

Step 3 of of the ID3 algorithm checks the purity of D and, given this case, assigns the unique
class to the respective node.

☐ The ID3 (Iterative Dichotomiser 3) was published by Ross Quinlan in 1986.

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ID3 Algorithm: Example

Example set D for mushrooms, drawn from a set of feature vectors X over the three dimensions color, size, and points:

	Color	Size	Points	Edibility
1	red	small	yes	toxic
2	brown	small	no	edible
3	brown	large	yes	edible
4	green	small	no	edible
5	red	large	no	edible



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ID3 Algorithm: Example (continued)

Top-level call of ID3. Analyze a splitting with regard to the feature "color":

$$D|_{ ext{color}} = egin{array}{c|c} \hline toxic & edible \\ \hline red & 1 & 1 \\ brown & 0 & 2 \\ green & 0 & 1 \\ \hline \end{array}$$

$$\rightarrow \quad |D_{\rm red}| = 2, \quad |D_{\rm brown}| = 2, \quad |D_{\rm green}| = 1$$

ID3 Algorithm: Example (continued)

Top-level call of ID3. Analyze a splitting with regard to the feature "color":

$$D|_{ ext{color}} = egin{array}{c|c} \hline toxic & edible \\ \hline red & 1 & 1 \\ brown & 0 & 2 \\ green & 0 & 1 \\ \hline \end{array}$$

$$\begin{array}{cccc} \mathbf{1} & & \mathbf{1} \\ \mathbf{0} & & \mathbf{2} \end{array} \qquad \sim & |D_{\text{red}}| = 2, \quad |D_{\text{brown}}| = 2, \quad |D_{\text{green}}| = 1 \end{array}$$

Estimated prior probabilities:

$$\hat{P}(\textit{Color} = \text{red}) = \frac{2}{5} = 0.4, \quad \hat{P}(\textit{Color} = \text{brown}) = \frac{2}{5} = 0.4, \quad \hat{P}(\textit{Color} = \text{green}) = \frac{1}{5} = 0.2$$

ID3 Algorithm: Example (continued)

Top-level call of ID3. Analyze a splitting with regard to the feature "color":

$$D|_{\rm color} = \begin{array}{c|cccc} & & & & \\ \hline toxic & {\rm edible} \\ \hline {\rm red} & {\rm 1} & {\rm 1} \\ {\rm brown} & {\rm 0} & {\rm 2} \\ {\rm green} & {\rm 0} & {\rm 1} \\ \end{array} \\ & \sim & |D_{\rm red}| = 2, \quad |D_{\rm brown}| = 2, \quad |D_{\rm green}| = 1$$

Estimated prior probabilities:

$$\hat{P}(\textit{Color} = \text{red}) = \frac{2}{5} = 0.4, \quad \hat{P}(\textit{Color} = \text{brown}) = \frac{2}{5} = 0.4, \quad \hat{P}(\textit{Color} = \text{green}) = \frac{1}{5} = 0.2$$

Conditional entropy:

$$\begin{split} H(\mathcal{A} \mid \mathcal{B}_1) &= H(\ \{A_1, A_2\} \mid \{B_{1,1}, B_{1,2}, B_{1,3}\}\) \\ &= H(\ \{\textit{\textbf{C}} = \mathsf{toxic}, \textit{\textbf{C}} = \mathsf{edible}\} \mid \{\textit{\textbf{Color}} = \mathsf{red}, \textit{\textbf{Color}} = \mathsf{brown}, \textit{\textbf{Color}} = \mathsf{green}\}\) \\ &= -(0.4 \cdot (\frac{1}{2} \cdot \log_2 \frac{1}{2} + \frac{1}{2} \cdot \log_2 \frac{1}{2}) + 0.4 \cdot (\frac{0}{2} \cdot \log_2 \frac{0}{2} + \frac{1}{2} \cdot \log_2 \frac{1}{2}) + 0.2 \cdot (\frac{0}{1} \cdot \log_2 \frac{0}{1} + \frac{1}{1} \cdot \log_2 \frac{1}{1})) = \ 0.4 \end{split}$$

ID3 Algorithm: Example (continued)

Top-level call of ID3. Analyze a splitting with regard to the feature "color":

$$D|_{\text{color}} \quad = \quad \begin{array}{c|cccc} & \text{toxic} & \text{edible} \\ \hline \text{red} & \mathbf{1} & \mathbf{1} \\ \text{brown} & \mathbf{0} & \mathbf{2} \\ \text{green} & \mathbf{0} & \mathbf{1} \end{array} \quad \rightsquigarrow \quad |D_{\text{red}}| = 2, \quad |D_{\text{brown}}| = 2, \quad |D_{\text{green}}| = 1$$

Estimated prior probabilities:

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$$H(A \mid \mathcal{B}_2) = H(\{C = \text{toxic}, C = \text{edible}\} \mid \{Size = \text{small}, Size = \text{large}\}) = \dots \approx 0.55$$

$$H(A \mid B_3) = H(\{C = \text{toxic}, C = \text{edible}\} \mid \{Points = \text{yes}, Points = \text{no}\}) = \dots = 0.4$$

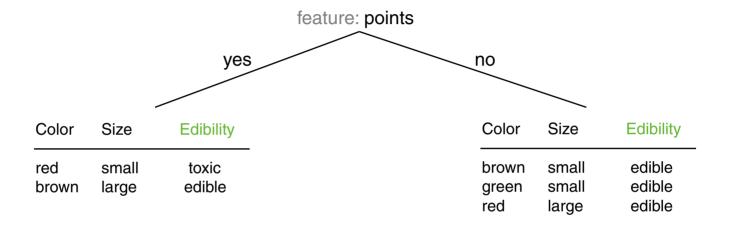


- The smaller $H(A \mid B)$ is, the larger becomes the information gain. Hence, the difference $H(A) H(A \mid B)$ needs not to be computed since H(A) is constant within each recursion step.
- ☐ In the example, the information gain in the first recursion step becomes maximum for the features "color" and "points".
- □ Notation. When used in the role of a random variable (here: in the argument of a probability P), features are written in italics and capitalized.
- □ Notation. The probabilities, denoted as $P(\cdot)$, are unknown and estimated by the relative frequencies, denoted as $\hat{P}(\cdot)$.

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ID3 Algorithm: Example (continued)

Decision tree before the first recursion step:

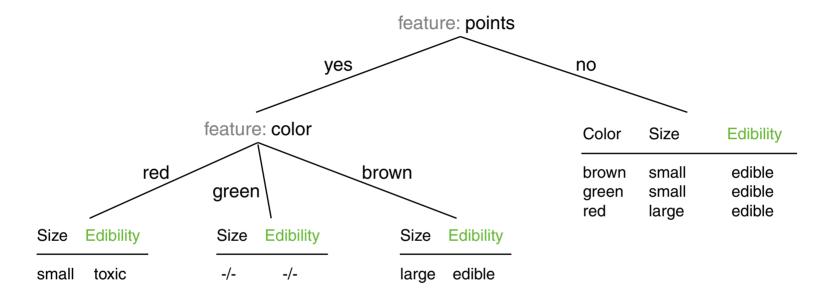


Choosing the feature "points" in Step 5 of the ID3 algorithm.

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ID3 Algorithm: Example (continued)

Decision tree before the second recursion step:

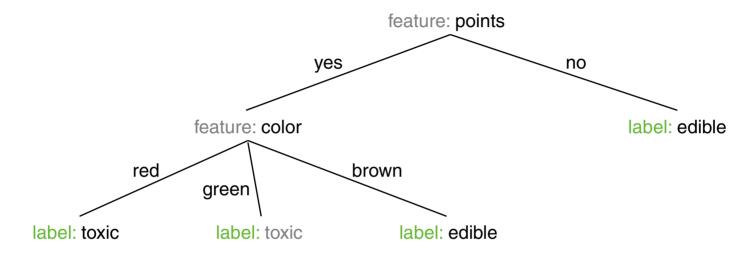


Choosing the feature "color" in Step 5 of the ID3 algorithm.

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ID3 Algorithm: Example (continued)

Final decision tree after second recursion step:

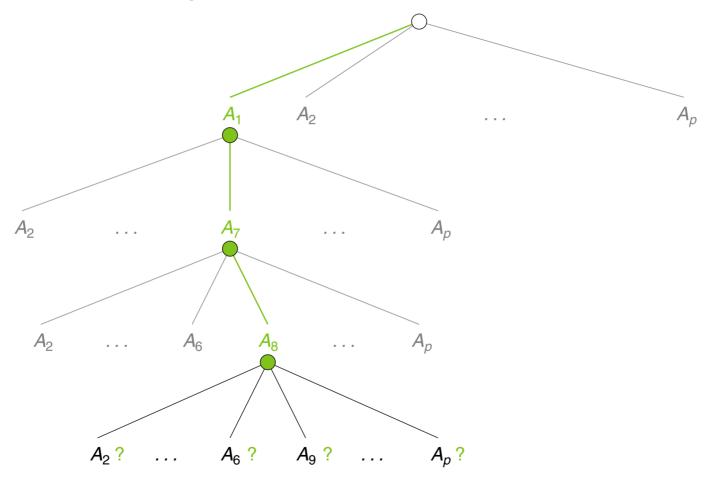


Break of a tie: choosing the class "toxic" for D_{green} in Step 6 of the ID3 algorithm.

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ID3 Algorithm: Search Space

Features =
$$\{A_1, A_2, ..., A_p\}$$



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Remarks (search space versus hypothesis space):

- The underlying search space of an algorithm that samples without replacement a single feature in each step (= monothetic splitting) consists of all permutations of the features in the feature set. In particular, if the number of features (= dimensionality of a feature vector \mathbf{x}) is p, then the search space contains p! elements.
- The set of possible decision trees over D forms the hypothesis space H. The maximum size of H, i.e., the maximum number of decision trees for a data set D in a binary classification setting, is $2^{|D|}$: If the feature vectors are pairwise distinct, every subset of D can form a class while the complement of the subset will form the other class. The set of possible subsets of D is $\mathcal{P}(D)$, where $|\mathcal{P}(D)| = 2^{|D|}$.
- Observe that either $p! < 2^{|D|}$ or $p! > 2^{|D|}$ can hold. I.e., the search space due to feature ordering can be smaller or larger than its underlying hypothesis space. The former characterizes the typical situation; also note that both the search space and the hypothesis space grow exponentially in the number of features and examples respectively.
- □ The difference between search space size and hypothesis space size results from Step 6 of the ID3 algorithm: the same feature selection order will lead to different decision trees when given different data sets. However, since the splitting operation in Step 6 is deterministic it has no effect on the search space.
- The runtime of the ID3 algorithm is in $O(p^2 \cdot n)$, i.e., significantly below p! since only a small part of the search space is explored. At each split, the algorithm greedily (in fact, irrevocably) selects the most informative feature by applying information gain as a heuristic for feature selection.

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ID3 Algorithm: Inductive Bias

Inductive bias is the rigidity in applying the (little bit of) knowledge learned from a training set for the classification of unseen feature vectors.

Observations:

Decision tree search happens in the space of all hypotheses.

 To generate a decision tree, the ID3 algorithm needs per branch at most as many decisions as features are given.

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ID3 Algorithm: Inductive Bias

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Observations:

- Decision tree search happens in the space of all hypotheses.
 - → The target concept is a member of the hypothesis space.
- To generate a decision tree, the ID3 algorithm needs per branch at most as many decisions as features are given.
 - → no backtracking takes place
 - → the decision tree is a result of *local optimization*

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- To generate a decision tree, the ID3 algorithm needs per branch at most as many decisions as features are given.
 - → no backtracking takes place
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Where the inductive bias of the ID3 algorithm becomes manifest:

- 1. Small decision trees are preferred.
- 2. Highly discriminative features tend to be closer to the root.

Is this justified?

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Remarks (inductive bias):

- ☐ The inductive bias of the ID3 algorithm is of a different kind than the inductive bias of the candidate elimination algorithm (or version space algorithm):
 - The underlying hypothesis space H of the candidate elimination algorithm is incomplete.
 H corresponds to a coarsened view onto the space of all hypotheses since H contains
 only conjunctions of feature-value pairs as hypotheses.
 However, this restricted hypothesis space is searched completely by the candidate
 elimination algorithm. Keyword: restriction bias
 - 2. The underlying hypothesis space *H* of the ID3 algorithm is complete since it contains all decision trees that can be constructed over *D*.
 - However, this complete hypothesis space is searched incompletely, but following a preference. Keyword: preference bias or search bias

☐ The inductive bias of the ID3 algorithm renders the algorithm robust wrt. noise.

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CART Algorithm [Breiman 1984] [ID3 Algorithm]

Setting:

- $\ \square$ X is a multiset of feature vectors. No restrictions are presumed for the features' measurement scales.
- \Box *C* is a set of classes.
- $D = \{(\mathbf{x}_1, c_1), \dots, (\mathbf{x}_n, c_n)\} \subseteq X \times C$ is a multiset of examples.

Learning task:

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Learning task:

 \Box Fit *D* using a decision tree *T*.

Characteristics of the CART algorithm:

- 1. Each splitting is binary and considers one feature at a time.
- 2. Splitting criterion is the information gain or the Gini index.

CART Algorithm (continued)

Let A be a feature with domain dom(A). Apply (probably multiple times) the respective rule to induce a finite number of binary splittings of X:

- 1. If A is nominal, choose $B \subset dom(A)$ such that $0 < |B| \le |dom(A) \setminus B|$.
- 2. If A is ordinal, choose $a \in dom(A)$ such that $x_{min} < a < x_{max}$, where x_{min} , x_{max} are the minimum and maximum values of feature A in D.
- 3. If A is numeric, choose $a \in dom(A)$ such that $a = 0.5 \cdot (x_{l_1} + x_{l_2})$, where x_{l_1} , x_{l_2} are consecutive elements in the ordered value list of feature A in D.

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CART Algorithm (continued)

Let A be a feature with domain dom(A). Apply (probably multiple times) the respective rule to induce a finite number of binary splittings of X:

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- 3. If A is numeric, choose $a \in dom(A)$ such that $a = 0.5 \cdot (x_{l_1} + x_{l_2})$, where x_{l_1} , x_{l_2} are consecutive elements in the ordered value list of feature A in D.

Adapt Step 5+6 to turn the ID3 into the CART algorithm:

- \Box For all $A \in Features$ generate with the above rules all splittings of D(t).
- fill Choose a splitting that maximizes the impurity reduction $\Delta\iota$:

figspace Recursively call CART to process $D(t_L)$ and $D(t_R)$.

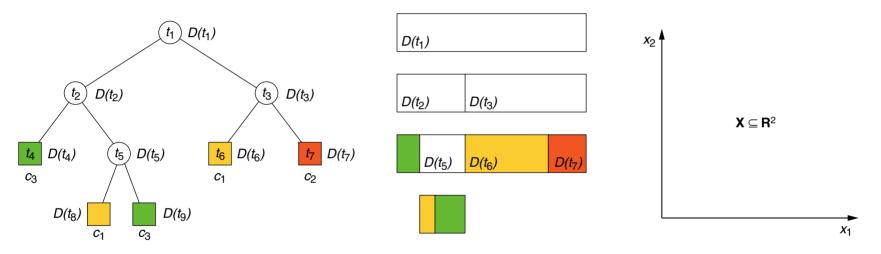
Remarks:

- \Box t_L and t_R denote the left and right successor of t in the decision tree. These nodes are returned by the calls of the CART algorithm and connected to t via createEdge().
- Since the CART algorithm creates binary splittings only, the feature A^* chosen in Step 5 can be chosen again later on. Hence, a call of CART to process $D(t_L)$ (or $D(t_R)$) in Step 6 passes the complete set of features as second parameter (and not: $Features \setminus \{A^*\}$).

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CART Algorithm (continued)

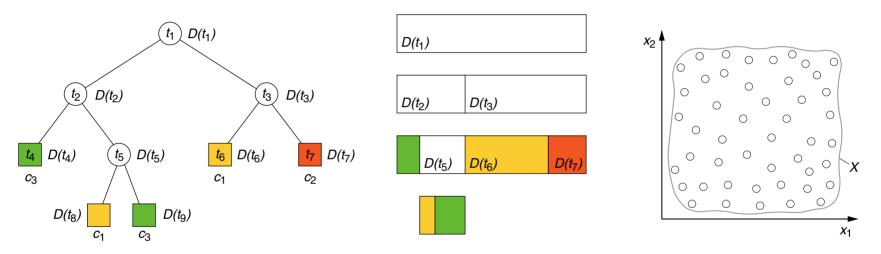
Illustration for two numeric features; i.e., the feature space X underlying X corresponds to a two-dimensional plane such as the \mathbf{R}^2 :



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CART Algorithm (continued)

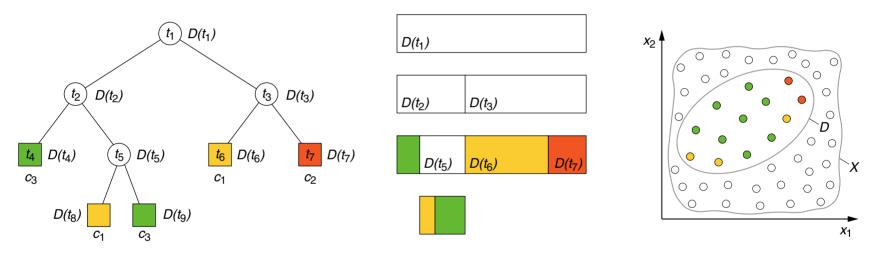
Illustration for two numeric features; i.e., the feature space \mathbf{X} underlying X corresponds to a two-dimensional plane such as the \mathbf{R}^2 :



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CART Algorithm (continued)

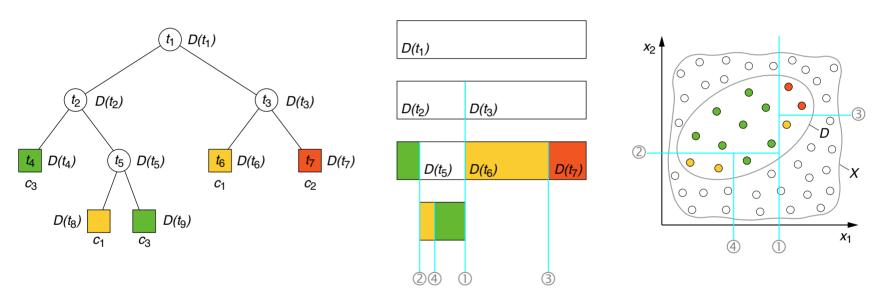
Illustration for two numeric features; i.e., the feature space \mathbf{X} underlying X corresponds to a two-dimensional plane such as the \mathbf{R}^2 :



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CART Algorithm (continued)

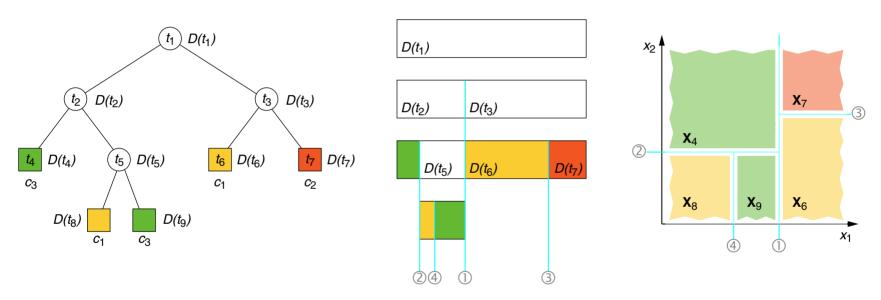
Illustration for two numeric features; i.e., the feature space \mathbf{X} underlying X corresponds to a two-dimensional plane such as the \mathbf{R}^2 :



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CART Algorithm (continued)

Illustration for two numeric features; i.e., the feature space X underlying X corresponds to a two-dimensional plane such as the \mathbf{R}^2 :



By the sequence of (here: four) splittings of D the feature space \mathbf{X} is cut into rectangular areas that are parallel to the two axes. Keyword: guillotine cuts

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