Machine Learning Decision Trees



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STAT/CS 5810/6655



Supervised Learning



- Let's come back to supervised learning
- Remember that we have labeled data

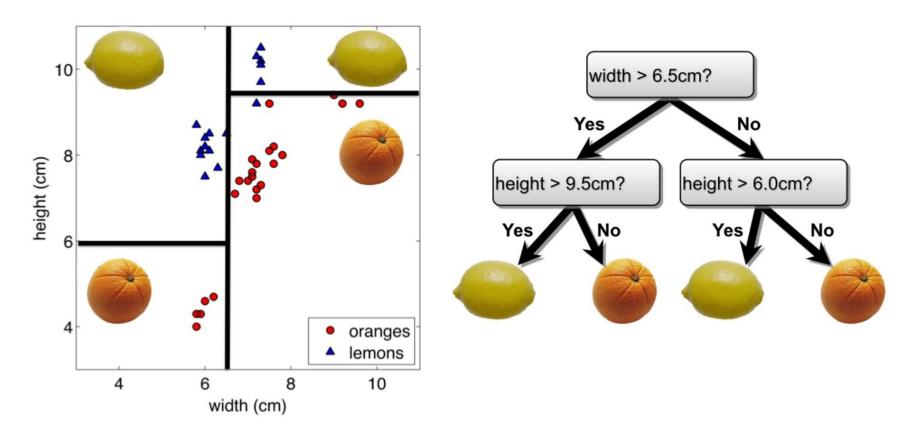
$$(\boldsymbol{x}_i, y_i)_{i=1}^n$$

- y_i finite \Rightarrow classification
- $y_i \in \mathbb{R} \Rightarrow$ regression
- We'll consider classification and regression trees (CART)

Example



 Classify a fruit as lemon or orange based on width and height



Example: Object Detecion





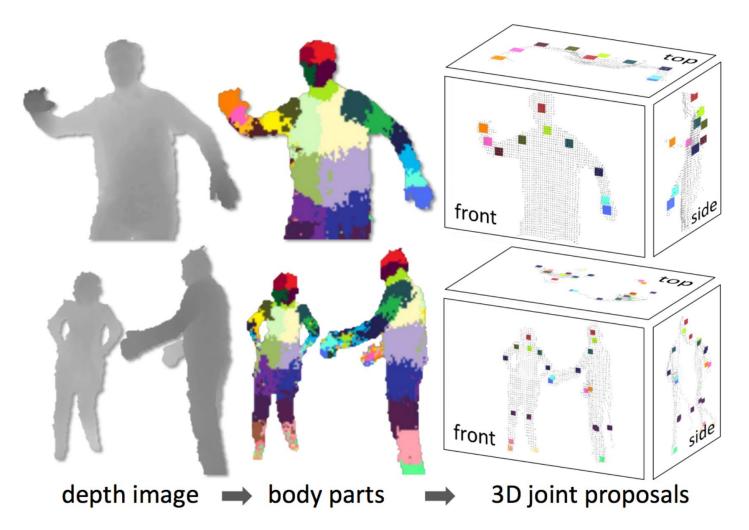




[J. Shotton, A. Fitzgibbon, M. Cook, T. Sharp, M. Finocchio, R. Moore, A. Kipman, A. Blake. Real-Time Human Pose

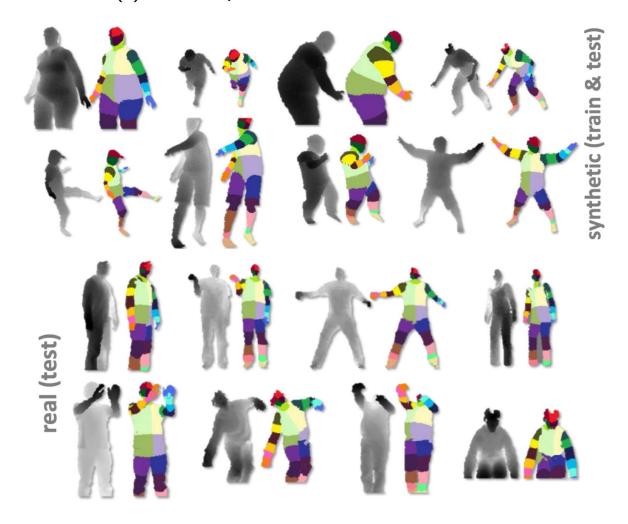


Classifying body parts



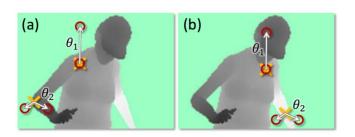


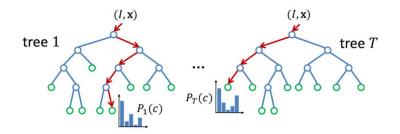
Trained on million(s) of examples



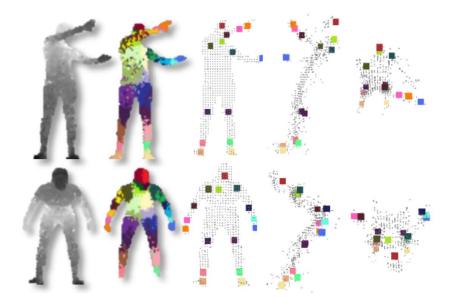


• Trained on million(s) of examples





Results:



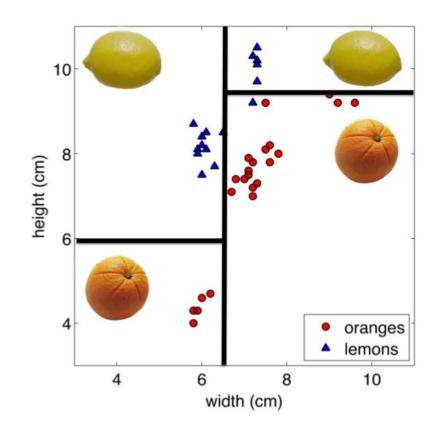
Decision Trees



- In short, a decision tree asks a series of simple questions about \boldsymbol{x} and then predicts a label for \boldsymbol{x}
- Decision trees can also be applied to regression
 - Labels at the "leaf nodes" are replaced with real numbers
- Other generalizations
 - Splits that involve more than one feature
 - Splits with more than two outcomes
 - The above splits can more easily lead to overfitting
- For simplicity, we'll assume only binary splits on a single feature

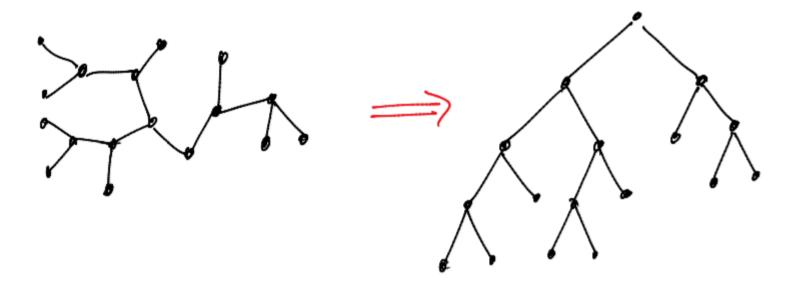


- Each decision tree is associated with a <u>partition</u> of the feature space
 - The elements of this partition are called <u>cells</u>
- Recall that a graph is a collection of <u>nodes</u>, some of which are connected by <u>edges</u>
 - The <u>degree</u> of a node is the number of edges incident on that node



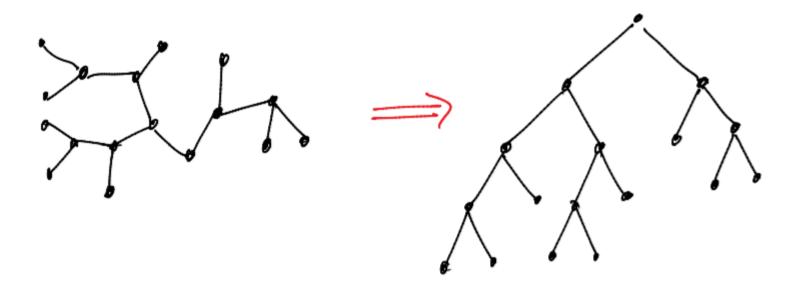


- A <u>tree</u> is a connected graph with no cycles
- A rooted binary tree is a tree where one node, called the root, has degree 2 and all other nodes have either degree 1 or degree 3
 - Degree 1 nodes are called <u>leaf</u> or <u>terminal</u> nodes
 - All other nodes are called internal nodes



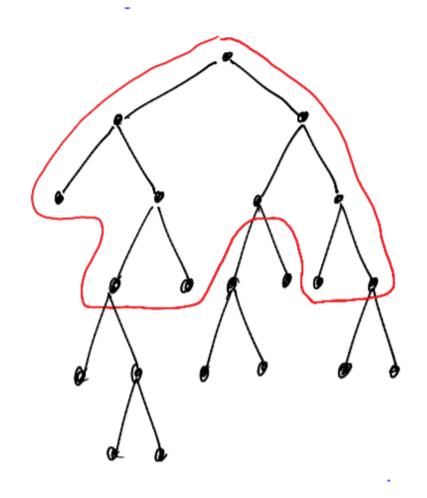


- The <u>depth</u> of a node is the number of edges between that node and the root
- The <u>parent</u> of a node is the neighbor of the node whose depth is one less
- Two nodes are siblings if they have the same parent





- A <u>subtree</u> is a subgraph that is also a tree
- A <u>rooted binary subtree</u> is a subtree that contains the root and is such that if any non-root node is included, so is its sibling
- A binary decision tree is a rooted binary tree where each internal node is associated with a binary classifier and each leaf node with a label



Learning Decision Trees



- Let T = set of all binary decision trees
- Basic strategy: regularized ERM

$$\min_{T \in \mathcal{T}} \frac{1}{n} \sum_{i=1}^{n} L(y_i, T(x_i)) + \lambda |T|$$

- L is an appropriate loss, e.g. 0-1 loss for classification or squared error loss for regression
- |T| = number of leaf nodes
- Unfortunately, this is intractable and so a different procedure is typically employed (next slide)

Learning Procedure



- 1. Grow a very large tree $T_{
 m 0}$ using a greedy approach
- 2. Prune T_0 by solving

$$\min_{T \in \mathcal{T}_0} \frac{1}{n} \sum_{i=1}^n L(y_i, T(\boldsymbol{x}_i)) + \lambda |T|$$

where \mathcal{T}_0 is the set of all binary decision trees based on rooted binary subtrees of T_0

Let's talk about each of these steps in detail



- A greedy algorithm is an algorithm that attempts to solve a global optimization problem by taking the locally optimal choice at each stage
- T_0 is constructed using a greedy approach
- 1. Start at root node (entire feature space)
- Decide whether to stop growing tree. If yes, assign a label. Stop
- 3. If no, consider all possible ways to split the data reaching the current node and select the best one
- 4. For each branch of the split, create a new node and go to Step 2



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- To implement this strategy, we need:
- a) List of possible splits
- b) Labelling rule
- c) Rule for stopping splitting
- d) Rule for selecting the best split



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 To implement this strategy, we need:

a) List of possible splits

- For continuous features, splits have the form $X^{(j)} \le t$?, j = 1, ..., d
- n data points \Rightarrow only n-1 values need to be considered for each j
- For categorical or discrete features, other simple splits can be used $X^{(j)} = \text{blue}$?



- Start at root node (entire feature space)
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- 3. If no, consider all possible ways to split the data reaching the current node and select the best one
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 To implement this strategy, we need:

b) Labelling rule

- Majority vote over the data reaching the given node for classification
- For regression, take the average y_i over the node



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- 4. For each branch of the split, create a new node and go to Step 2

- To implement this strategy, we need:
- c) Rule for stopping splitting
- Common strategy: split until each leaf node contains a single data point



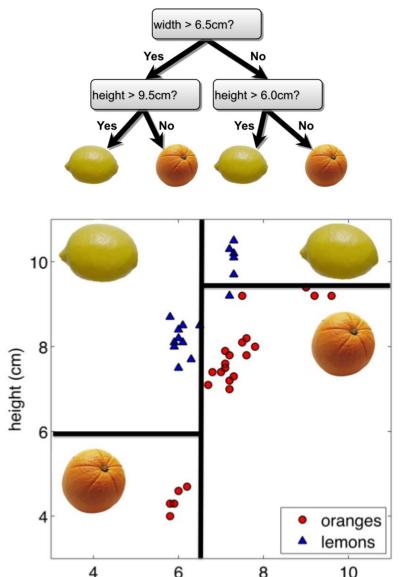
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- To implement this strategy, we need:
- d) Rule for selecting the best split
- Let's go over this in detail

Split Selection with Impurity Measures



- Let's focus on binary classification
- Suppose N is a leaf node at some stage in the growing process
 - Think of N as a cell in a partition of the feature space or the training points in that cell
- Intuitively, a good split leads to children that are more <u>homogeneous</u> or <u>pure</u> than their parent



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Impurity Measure



- Assume class labels are $\{0,1\}$
- Denote

$$q := \frac{|\{i: x_i \in N, y_i = 0\}|}{|\{i: x_i \in N\}|}$$

- This ratio defines a probability distribution of the labels
- An impurity measure is a function i(N) s.t.
 - $i(N) \ge 0$ with equality iff N consists of a single class
 - A larger value of i(N) indicates that the distribution defined by q is closer to the uniform distribution

Impurity Measure Examples



1. Entropy

$$i(N) = -[q \log q + (1 - q) \log(1 - q)]$$

2. Gini index (typically the default)

$$i(N) = 2q(1-q)$$

3. Misclassification rate of majority vote classifier

$$i(N) = \min(q, 1 - q)$$

Impurity Measure Examples

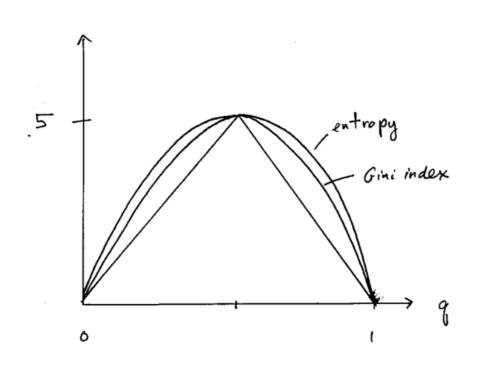


1. Entropy

$$i(N)$$

= $-[q \log q + (1-q) \log(1-q)]$

- 2. Gini index i(N) = 2q(1-q)
- 3. Misclassification rate of majority vote classifier $i(N) = \min(q, 1 q)$



Selecting the best split



- Let's maximize the decrease in impurity to choose the best split
- Let N_1 and N_2 be two children of N
- Define $p(N_1) = |N_1|/|N|$ and $p(N_2) = |N_2|/|N|$
 - |N| = # of training points in the cell corresponding to N
- The decrease in expected impurity:

$$i(N) - [p(N_1)i(N_1) + p(N_2)i(N_2)]$$

• When i is entropy, this is called the information gain

Concave impurity measures



Proposition: If i is concave as a function of q, then $i(N) - [p(N_1)i(N_1) + p(N_2)i(N_2)] \ge 0 \ \forall N_1, N_2$. If i is strictly concave and $N_1 \ne \phi \ne N_2$ (ϕ is the empty set) then equality holds iff $q = q_1 = q_2$ where q_i is the proportion of training points of class 0 in N_i .

- This result explains why strictly concave impurity measures are preferred
- The result generalizes easily to multiclass classification

Proof



Let $i(N) = \psi(q)$ where ψ is concave. Then

$$i(N) = \psi(q) = \psi(p(N_1)q_1 + p(N_2)q_2)$$

$$\geq p(N_1)\psi(q_1) + p(N_2)\psi(q_2)$$

$$= p(N_1)i(N_1) + p(N_2)i(N_2).$$

The inequality comes from Jensen's inequality applied to ψ which holds with equality iff the q_i terms are equal and $p(N_i) > 0$ for i = 1,2 (this follows from the fact that the N_i are nonempty). Therefore, $q = q_1 = q_2$.

Pruning



To solve

$$\min_{T \in \mathcal{T}_0} J(T) \coloneqq \frac{1}{n} \sum_{i=1}^n L(y_i, T(x_i)) + \lambda |T|$$

we rely on the fact that the objective function is additive:

• Let $\pi(T)=\{A_1,A_2\}$ be the partition of the feature space corresponding to the children of the root node. Then

$$J(T) = \sum_{A \in \pi(T)} J(T_A)$$

where T_A is the subtree rooted at A

- This suggests a recursive algorithm
- Alternatively, an efficient bottom-up dynamic programming algorithm can be used to solve the pruning problem.

Question

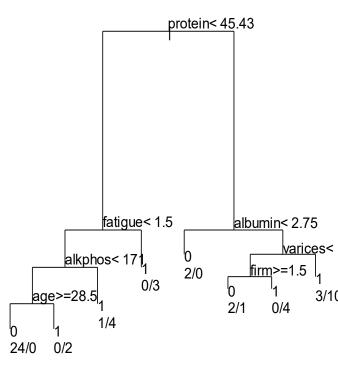


- Why should we grow a tree and then prune it as opposed to growing and just stopping when the decrease in impurity is negligible?
- Answer: because of ancillary splits
 - These are splits that have no value by themselves but that enable useful splits later on.

Final Thoughts



- Advantages of decision trees
 - Interpretable
 - Automatic variable selection
 - Can handle nonlinear interactions
 - Rapid evaluation
 - Easily handle categorical data and multiple classes
- Disadvantages of decision trees
 - Greedy growing is suboptimal
 - Unstable: a slight perturbation of training data could drastically change the learned tree
 - Nonsmooth decision boundaries
- The latter two issues can be addressed using ensemble methods (i.e. random forests)



Further Reading



- Ripley, <u>Pattern Recognition and Neural Networks</u>, 1996
- ISL Section 8.1
- ESL Section 9.2