

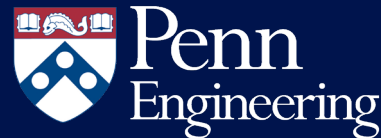


Decision Trees

Dan Roth

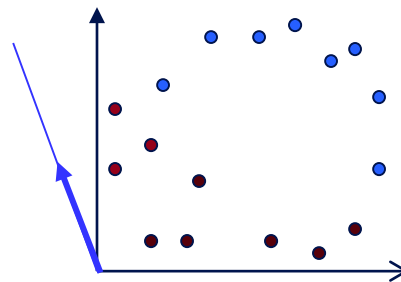
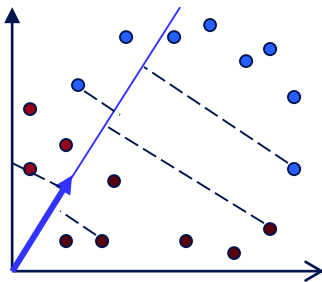
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Slides were created by Dan Roth (for CIS519/419 at Penn or CS446 at UIUC), Eric Eaton for CIS519/419 at Penn, or from other authors who have made their ML slides available.



Introduction - Summary

- We introduced the technical part of the class by giving two (very important) examples for learning approaches to linear discrimination.
- There are many other solutions.
- **Question 1:** Our solution learns a linear function; in principle, the target function may not be linear, and this will have implications on the performance of our learned hypothesis.
 - **Can we learn a function that is more flexible in terms of what it does with the feature space?**
- **Question 2:** Can we say something about the quality of what we learn (sample complexity, time complexity; quality)

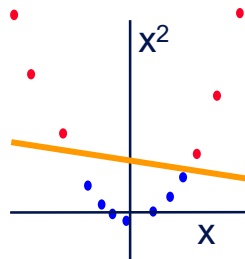


Decision Trees

- Earlier, we decoupled the generation of the feature space from the learning.
- Argued that we can map the given examples into another space, in which the target functions are linearly separable.
- Do we always want to do it?
- How do we determine what are good mappings?
- The study of **decision trees** may shed some light on this.
- Learning is done directly from the given data representation.
- The algorithm “transforms” the data itself.

Think about the Badges problem

What's the best learning algorithm?



This Lecture

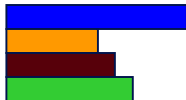
- Decision trees for (binary) classification
 - Non-linear classifiers
- Learning decision trees (ID3 algorithm)
 - Greedy heuristic (based on information gain)
Originally developed for discrete features
 - Some extensions to the basic algorithm
- Overfitting
 - Some experimental issues



Introduction of Decision trees

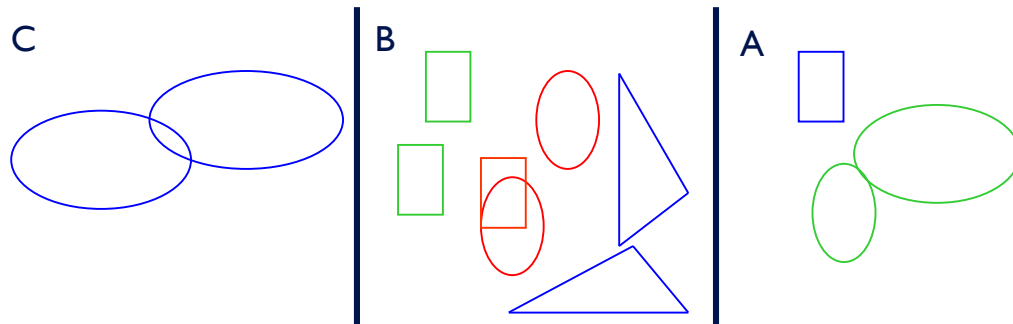
Representing Data

- Think about a large table, **N attributes**, and assume you want to know something about the people represented as entries in this table.
- E.g. **own** an expensive car or **not**;
- Simplest way: Histogram on the **first** attribute – **own**
- Then, histogram on **first and second (own & gender)**
- But, what if the # of attributes is larger: **N=16**
- How large are the **1-d histograms** (contingency tables) ? 16 numbers
- How large are the **2-d histograms**? 16-choose-2 = 120 numbers
- How many 3-d tables? 560 numbers
- With 100 attributes, the 3-d tables need 161,700 numbers
 - **We need to figure out a way to represent data in a better way, and figure out what are the important attributes to look at first.**
 - **Information theory has something to say about it – we will use it to better represent the data.**



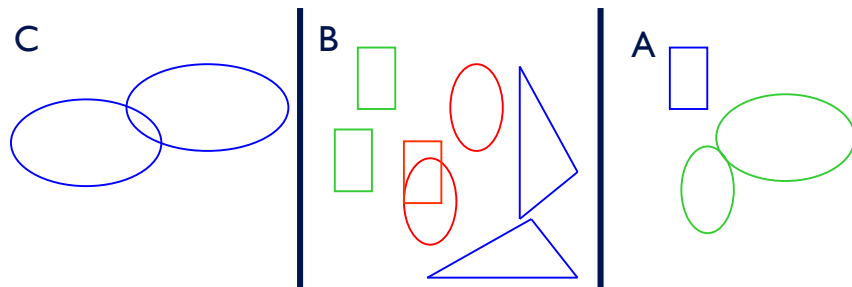
Decision Trees

- A hierarchical data structure that represents data by implementing a divide and conquer strategy
- Can be used as a non-parametric classification and regression method
- Given a collection of examples, learn a decision tree that represents it.
- Use this representation to classify new examples



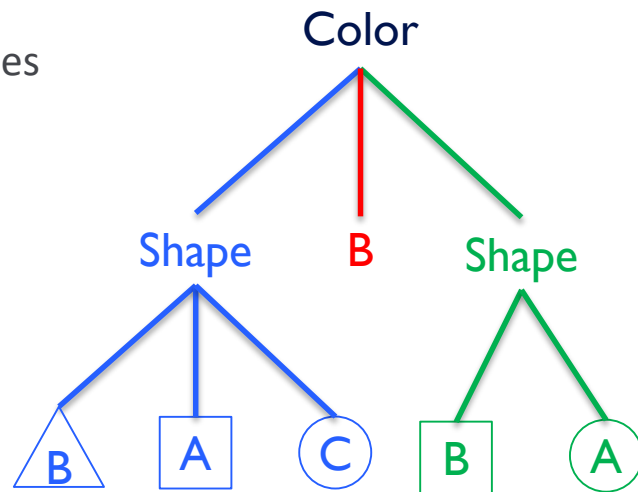
The Representation

- Decision Trees are classifiers for instances represented as feature vectors
 - color={red, blue, green} ; shape={circle, triangle, rectangle} ; label= {A, B, C}
- Nodes** are **tests** for feature values
- There is one branch for each value of the feature
- Leaves** specify the category (labels)
- Can categorize instances into multiple disjoint categories

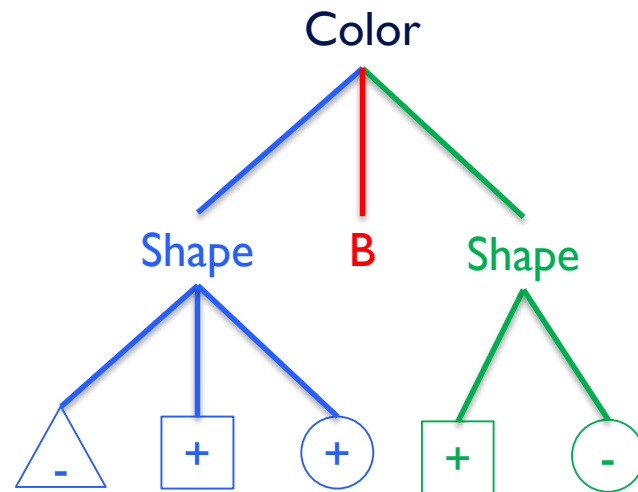


Evaluation of a
Decision Tree

Learning a
Decision Tree

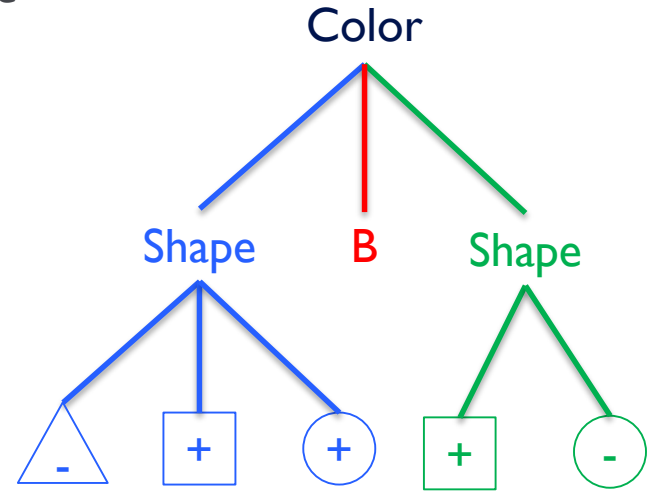


Expressivity of Decision Trees



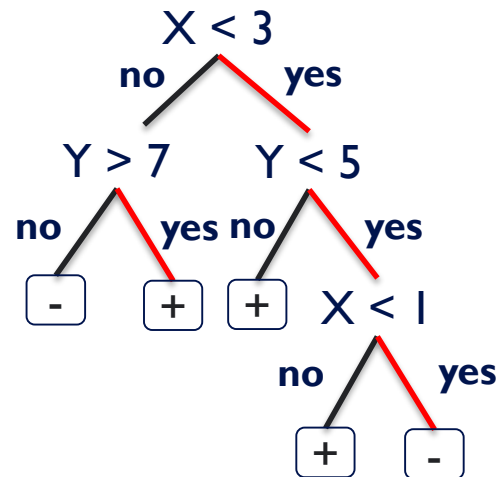
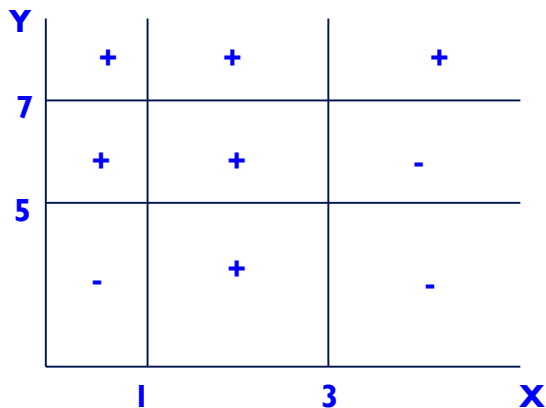
Decision Trees

- Output is a discrete category. Real valued outputs are possible (regression trees)
- There are efficient algorithms for processing large amounts of data (but not too many features)
- There are methods for handling **noisy data** (classification noise and attribute noise) and for handling missing attribute values



Decision Boundaries

- Usually, instances are represented as attribute-value pairs (color=blue, shape = square, +)
- Numerical values can be used either by discretizing or by using thresholds for splitting nodes
- In this case, the tree divides the features space into axis-parallel rectangles, each labeled with one of the labels



Today's key concepts

- Learning decision trees (ID3 algorithm)
 - Greedy heuristic (based on information gain)
Originally developed for discrete features
- Overfitting
 - What is it? How do we deal with it?
- Some extensions of DTs
- Principles of Experimental ML

Administration

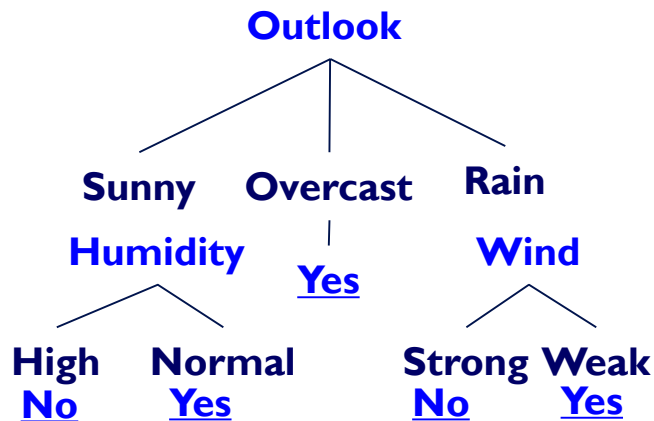
- Since there is no waiting list anymore; all people that wanted to be in are in.
- Everyone should have submitted HW0
- Recitations
- Quizzes
- HW 1 will be released on Monday.
 - Please start working on it as soon as you can. Don't wait until the last couple of days.
- Questions?
 - Please ask/comment during class.



Learning decision trees (ID3 algorithm)

Decision Trees

- Can represent any Boolean Function
- Can be viewed as a way to compactly represent a lot of data.
- Natural representation: (20 questions)
- The **evaluation** of the Decision Tree Classifier is easy
- Clearly, given data, there are many ways to represent it as a decision tree.
- Learning a **good** representation from data is the challenge.



Will I play tennis today?

- **Features**

- Outlook: {Sun, Overcast, Rain}
- Temperature: {Hot, Mild, Cool}
- Humidity: {High, Normal, Low}
- Wind: {Strong, Weak}

- **Labels**

- Binary classification task: $Y = \{+, -\}$

Will I play tennis today?

	O	T	H	W	Play?
1	S	H	H	W	-
2	S	H	H	S	-
3	O	H	H	W	+
4	R	M	H	W	+
5	R	C	N	W	+
6	R	C	N	S	-
7	O	C	N	S	+
8	S	M	H	W	-
9	S	C	N	W	+
10	R	M	N	W	+
11	S	M	N	S	+
12	O	M	H	S	+
13	O	H	N	W	+
14	R	M	H	S	-

Outlook: S(unny),
O(vercast),
R(ainy)

Temperature: H(ot),
M(edium),
C(ool)

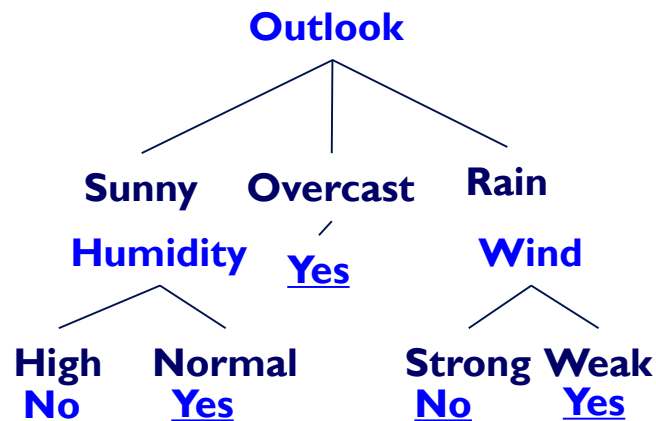
Humidity: H(igh),
N(ormal),
L(ow)

Wind: S(trong),
W(eak)

Basic Decision Trees Learning Algorithm

	O	T	H	W	Play?
1	S	H	H	W	-
2	S	H	H	S	-
3	O	H	H	W	+
4	R	M	H	W	+
5	R	C	N	W	+
6	R	C	N	S	-
7	O	C	N	S	+
8	S	M	H	W	-
9	S	C	N	W	+
10	R	M	N	W	+
11	S	M	N	S	+
12	O	M	H	S	+
13	O	H	N	W	+
14	R	M	H	S	-

- Data is processed in Batch (i.e. all the data available) Algorithm?
- Recursively build a decision tree top down.



Basic Decision Tree Algorithm

- Let S be the set of Examples
 - $Label$ is the target attribute (the prediction)
 - $Attributes$ is the set of measured attributes
- $ID3(S, Attributes, Label)$

If all examples are labeled the same return a single node tree with $Label$

Otherwise Begin

$A =$ attribute in $Attributes$ that best classifies S (Create a Root node for tree)

for each possible value v of A

Add a new tree branch corresponding to $A=v$

Let S_v be the subset of examples in S with $A=v$

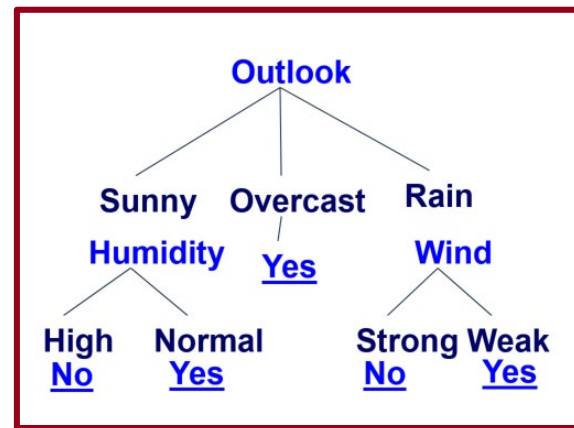
if S_v is empty: add leaf node with the common value of $Label$ in S

Else: below this branch add the subtree

$ID3(S_v, Attributes - \{A\}, Label)$

End

Return Root



why?

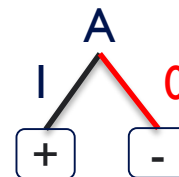
For evaluation time

Picking the Root Attribute

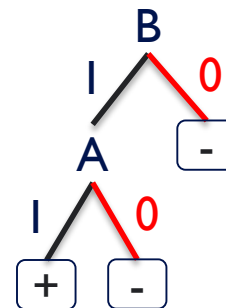
- The goal is to have the resulting decision tree as small as possible (Occam's Razor)
 - But, finding the minimal decision tree consistent with the data is NP-hard
- The recursive algorithm is a greedy heuristic search for a simple tree, but cannot guarantee optimality.
- The main decision in the algorithm is the selection of the next attribute to condition on.

Picking the Root Attribute

- Consider data with two Boolean attributes (A,B).
 - $\langle (A=0, B=0), - \rangle$: 50 examples
 - $\langle (A=0, B=1), - \rangle$: 50 examples
 - $\langle (A=1, B=0), - \rangle$: 0 examples
 - $\langle (A=1, B=1), + \rangle$: 100 examples
- What should be the first attribute we select?
 - Splitting on A: we get purely labeled nodes.
 - Splitting on B: we don't get purely labeled nodes.
 - What if we have: $\langle (A=1, B=0), - \rangle$: 3 examples?
- (one way to think about it: # of queries required to label a random data point)



splitting on A



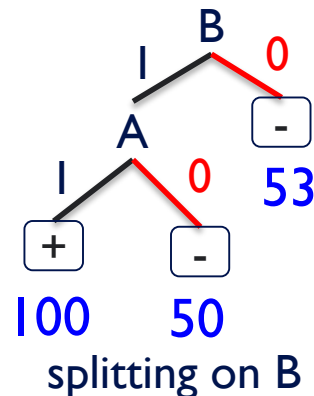
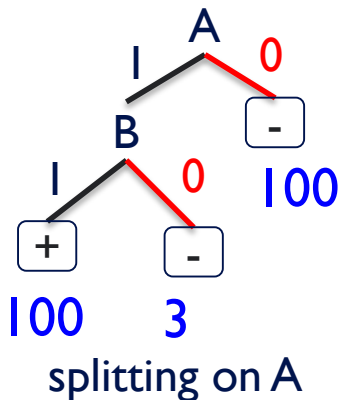
splitting on B

Picking the Root Attribute

- Consider data with two Boolean attributes (A,B).
 - $\langle A=0, B=0 \rangle, -$: 50 examples
 - $\langle A=0, B=1 \rangle, -$: 50 examples
 - $\langle A=1, B=0 \rangle, -$: 0 examples 3 examples
 - $\langle A=1, B=1 \rangle, +$: 100 examples
- What should be the first attribute we select?
- Trees looks structurally similar; which attribute should we choose?

Advantage A. But...
Need a way to quantify things

- One way to think about it: # of queries required to label a random data point.
- If we choose A we have less uncertainty about the labels.



Picking the Root Attribute

- The goal is to have the resulting decision tree as small as possible (Occam's Razor)
 - The main decision in the algorithm is the selection of the next attribute to condition on.
- We want attributes that split the examples to sets that are **relatively pure in one label**; this way we are closer to a leaf node.
 - The most popular heuristics is based on **information gain**, originated with the ID3 system of Quinlan.

Entropy

- Entropy (impurity, disorder) of a set of examples, S , relative to a binary classification is:

$$Entropy(S) = -p_+ \log(p_+) - p_- \log(p_-)$$

- p_+ is the proportion of positive examples in S and
- p_- is the proportion of negatives examples in S
 - If all the examples belong to the same category: Entropy = 0
 - If all the examples are equally mixed (0.5, 0.5): Entropy = 1
 - Entropy = Level of uncertainty.
- In general, when p_i is the fraction of examples labeled i :

$$Entropy(S[p_1, p_2, \dots, p_k]) = -\sum_{i=1}^k p_i \log(p_i)$$

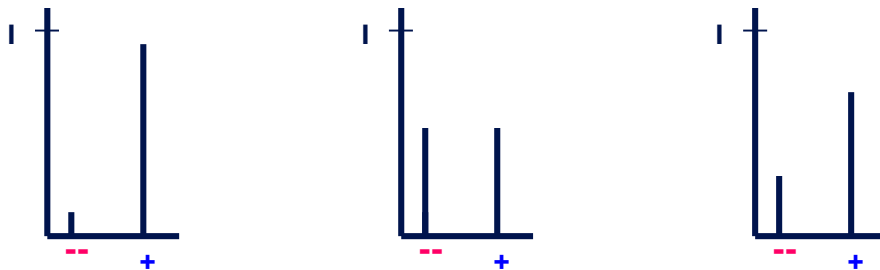
- Entropy can be viewed as the number of bits required, on average, to encode the class of labels. If the probability for + is 0.5, a single bit is required for each example; if it is 0.8 – can use less than 1 bit.

Entropy

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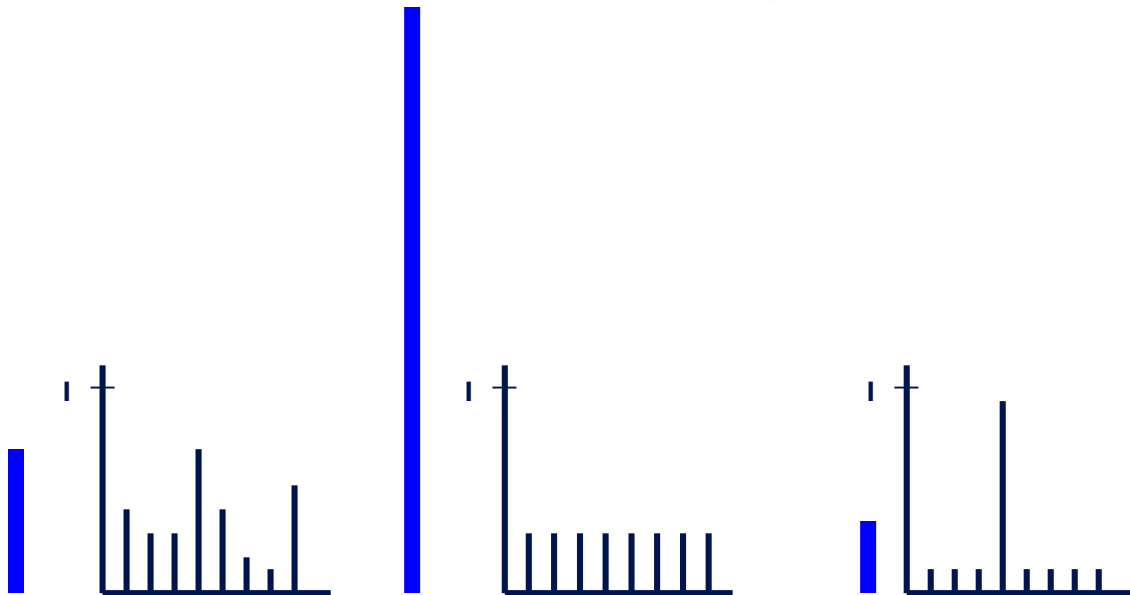


Entropy

(Convince yourself that the max value would be $\log(k)$)

(Also note that the base of the log only introduce a constant factor; therefore, we'll think about base 2)

$$\text{Entropy}(S[p_1, p_2, \dots, p_k]) = - \sum_1^k p_i \log(p_i)$$



Information Gain

High Entropy – High level of Uncertainty

Low Entropy – No Uncertainty.

- The information gain of an attribute **a** is the expected reduction in entropy caused by partitioning on this attribute

$$Gain(S, a) = Entropy(S) - \sum_{v \in values(S)} \frac{|S_v|}{|S|} Entropy(S_v)$$

- Where:

- S_v is the subset of **S** for which attribute **a** has value **v**, and
- the entropy of partitioning the data is calculated by **weighing the entropy of each partition** by its size relative to the original set



- Partitions of low entropy (imbalanced splits) lead to high gain
- Go back to check which of the A, B splits is better

Will I play tennis today?

	O	T	H	W	Play?
1	S	H	H	W	-
2	S	H	H	S	-
3	O	H	H	W	+
4	R	M	H	W	+
5	R	C	N	W	+
6	R	C	N	S	-
7	O	C	N	S	+
8	S	M	H	W	-
9	S	C	N	W	+
10	R	M	N	W	+
11	S	M	N	S	+
12	O	M	H	S	+
13	O	H	N	W	+
14	R	M	H	S	-

Outlook: S(unny),
O(vercast),
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Temperature: H(ot),
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Humidity: H(igh),
N(ormal),
L(ow)

Wind: S(trong),
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Will I play tennis today?

	O	T	H	W	Play?
1	S	H	H	W	-
2	S	H	H	S	-
3	O	H	H	W	+
4	R	M	H	W	+
5	R	C	N	W	+
6	R	C	N	S	-
7	O	C	N	S	+
8	S	M	H	W	-
9	S	C	N	W	+
10	R	M	N	W	+
11	S	M	N	S	+
12	O	M	H	S	+
13	O	H	N	W	+
14	R	M	H	S	-

calculate current entropy

- $p_+ = \frac{9}{14}$ $p_- = \frac{5}{14}$
- $Entropy(Play) = -p_+ \log_2(p_+) - p_- \log_2(p_-)$
 $= -\frac{9}{14} \log_2 \frac{9}{14} - \frac{5}{14} \log_2 \frac{5}{14}$
 ≈ 0.94

Information Gain: Outlook

	O	T	H	W	Play?
1	S	H	H	W	-
2	S	H	H	S	-
3	O	H	H	W	+
4	R	M	H	W	+
5	R	C	N	W	+
6	R	C	N	S	-
7	O	C	N	S	+
8	S	M	H	W	-
9	S	C	N	W	+
10	R	M	N	W	+
11	S	M	N	S	+
12	O	M	H	S	+
13	O	H	N	W	+
14	R	M	H	S	-

$$Gain(S, a) = Entropy(S) - \sum_{v \in values(S)} \frac{|S_v|}{|S|} Entropy(S_v)$$

Outlook = sunny:

$$p_+ = 2/5 \quad p_- = 3/5$$

$$Entropy(O = S) = 0.971$$

Outlook = overcast:

$$p_+ = 4/4 \quad p_- = 0$$

$$Entropy(O = O) = 0$$

Outlook = rainy:

$$p_+ = 3/5 \quad p_- = 2/5$$

$$Entropy(O = R) = 0.971$$

Expected entropy

$$= \sum_{v \in values(S)} \frac{|S_v|}{|S|} Entropy(S_v)$$

$$= (5/14) \times 0.971 + (4/14) \times 0 + (5/14) \times 0.971 = 0.694$$

$$Information\ gain = 0.940 - 0.694 = 0.246$$

Information Gain: Humidity

	O	T	H	W	Play?
1	S	H	H	W	-
2	S	H	H	S	-
3	O	H	H	W	+
4	R	M	H	W	+
5	R	C	N	W	+
6	R	C	N	S	-
7	O	C	N	S	+
8	S	M	H	W	-
9	S	C	N	W	+
10	R	M	N	W	+
11	S	M	N	S	+
12	O	M	H	S	+
13	O	H	N	W	+
14	R	M	H	S	-

$$Gain(S, a) = Entropy(S) - \sum_{v \in values(S)} \frac{|S_v|}{|S|} Entropy(S_v)$$

Humidity = high:

$$p_+ = 3/7 \quad p_- = 4/7$$

$$Entropy(H = H) = 0.985$$

Humidity = Normal:

$$p_+ = 6/7 \quad p_- = 1/7$$

$$Entropy(H = N) = 0.592$$

Expected entropy

$$= \sum_{v \in values(S)} \frac{|S_v|}{|S|} Entropy(S_v)$$

$$= (7/14) \times 0.985 + (7/14) \times 0.592 = 0.7785$$

$$Information\ gain = 0.940 - 0.694 = 0.246$$

Which feature to split on?

	O	T	H	W	Play?
1	S	H	H	W	-
2	S	H	H	S	-
3	O	H	H	W	+
4	R	M	H	W	+
5	R	C	N	W	+
6	R	C	N	S	-
7	O	C	N	S	+
8	S	M	H	W	-
9	S	C	N	W	+
10	R	M	N	W	+
11	S	M	N	S	+
12	O	M	H	S	+
13	O	H	N	W	+
14	R	M	H	S	-

Information gain:

Outlook: 0.246

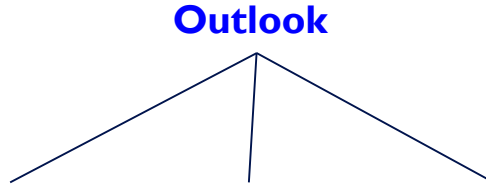
Humidity: 0.151

Wind: 0.048

Temperature: 0.029

→ Split on Outlook

An Illustrative Example (III)



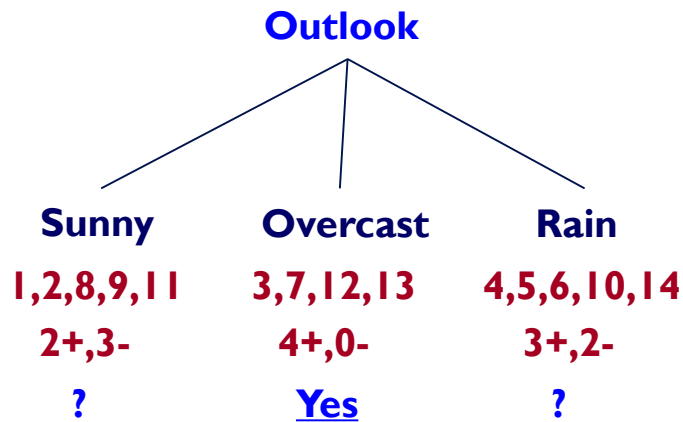
Gain(S, Humidity) = 0.151

Gain(S, Wind) = 0.048

Gain(S, Temperature) = 0.029

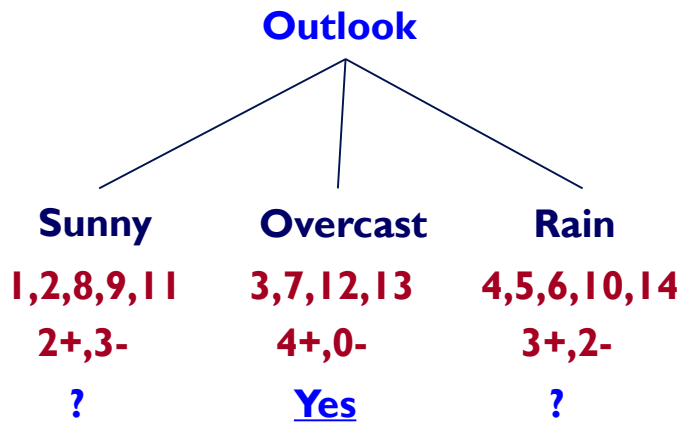
Gain(S, Outlook) = **0.246**

An Illustrative Example (III)



	O	T	H	W	Play?
1	S	H	H	W	-
2	S	H	H	S	-
3	O	H	H	W	+
4	R	M	H	W	+
5	R	C	N	W	+
6	R	C	N	S	-
7	O	C	N	S	+
8	S	M	H	W	-
9	S	C	N	W	+
10	R	M	N	W	+
11	S	M	N	S	+
12	O	M	H	S	+
13	O	H	N	W	+
14	R	M	H	S	-

An Illustrative Example (III)

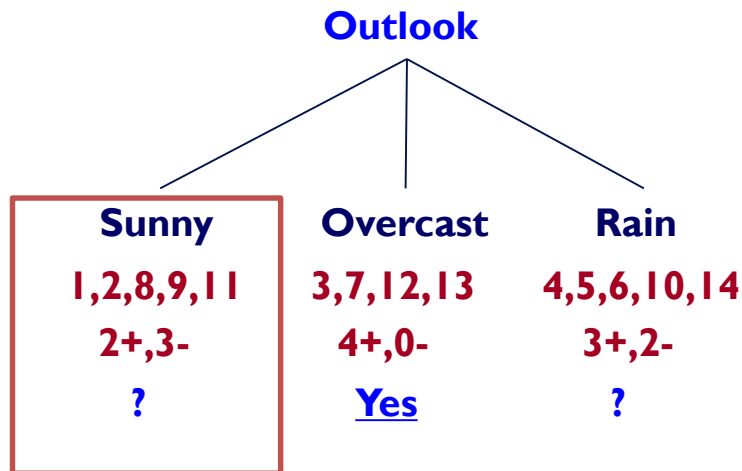


Continue until:

- Every attribute is included in **path**, or,
- All examples in the leaf have same label

	O	T	H	W	Play?
1	S	H	H	W	-
2	S	H	H	S	-
3	O	H	H	W	+
4	R	M	H	W	+
5	R	C	N	W	+
6	R	C	N	S	-
7	O	C	N	S	+
8	S	M	H	W	-
9	S	C	N	W	+
10	R	M	N	W	+
11	S	M	N	S	+
12	O	M	H	S	+
13	O	H	N	W	+
14	R	M	H	S	-

An Illustrative Example (IV)



$$\text{Gain}(S_{\text{sunny}}, \text{Humidity}) = .97 - (3/5) \cdot 0 - (2/5) \cdot 0 = .97$$

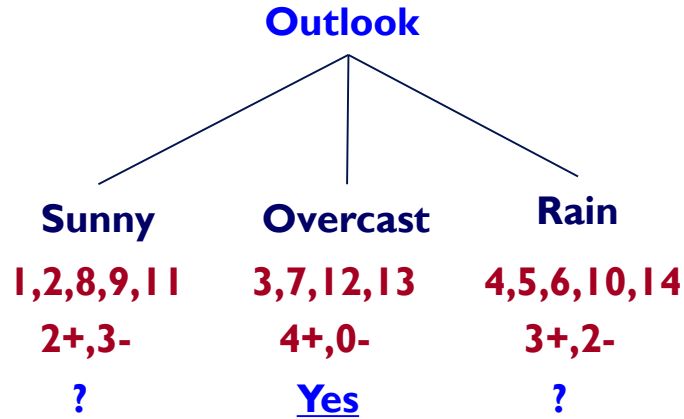
$$\text{Gain}(S_{\text{sunny}}, \text{Temp}) = .97 - 0 - (2/5) \cdot 1 = .57$$

$$\text{Gain}(S_{\text{sunny}}, \text{Wind}) = .97 - (2/5) \cdot 1 - (3/5) \cdot .92 = .02$$

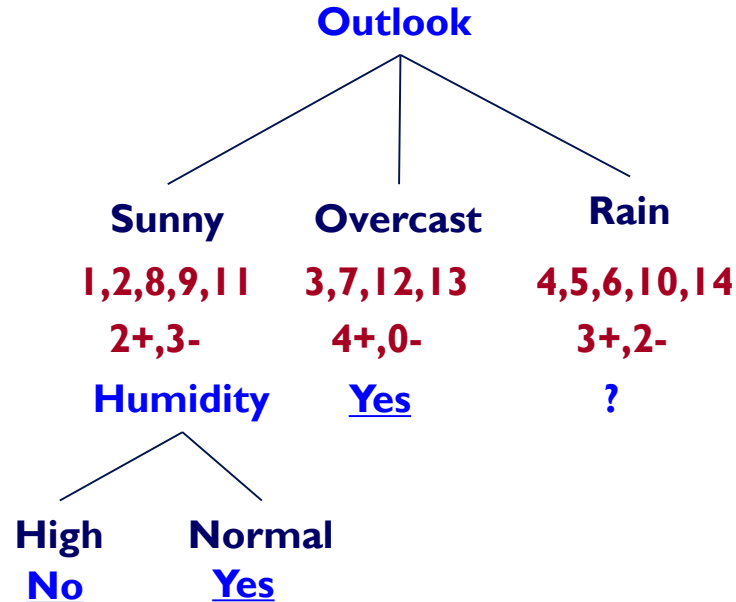
Split on Humidity

	O	T	H	W	Play?
1	S	H	H	W	-
2	S	H	H	S	-
4	R	M	H	W	+
5	R	C	N	W	+
6	R	C	N	S	-
7	O	C	N	S	+
8	S	M	H	W	-
9	S	C	N	W	+
10	R	M	N	W	+
11	S	M	N	S	+
12	O	M	H	S	+
13	O	H	N	W	+
14	R	M	H	S	-

An Illustrative Example (V)



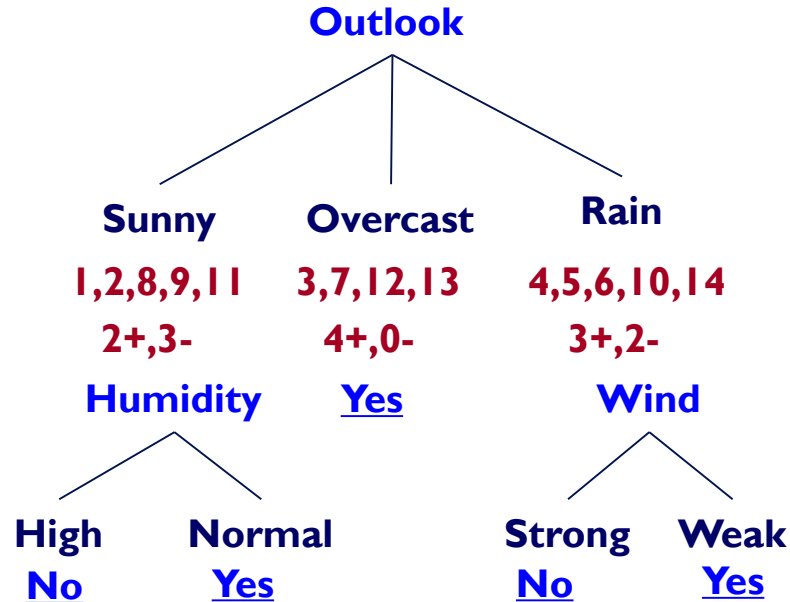
An Illustrative Example (V)



induceDecisionTree(S)

- 1. Does S uniquely define a class?
if all $s \in S$ have the same label y : **return** S ;
- 2. Find the feature with the most information gain:
 $i = \operatorname{argmax}_i \operatorname{Gain}(S, X_i)$
- 3. Add children to S :
for k in $\operatorname{Values}(X_i)$:
 $S_k = \{s \in S \mid x_i = k\}$
 addChild(S, S_k)
 induceDecisionTree(S_k)
return S ;

An Illustrative Example (VI)



Hypothesis Space in Decision Tree Induction

- Conduct a search of the space of decision trees which can represent all possible discrete functions. (**pros and cons**)
- Goal: to find the **best** decision tree
 - Best could be “smallest depth”
 - Best could be “minimizing the expected number of tests”
- Finding a minimal decision tree consistent with a set of data is **NP-hard**.
- Performs a greedy heuristic search: hill climbing **without backtracking**
- Makes statistically based decisions using **all data**

History of Decision Tree Research

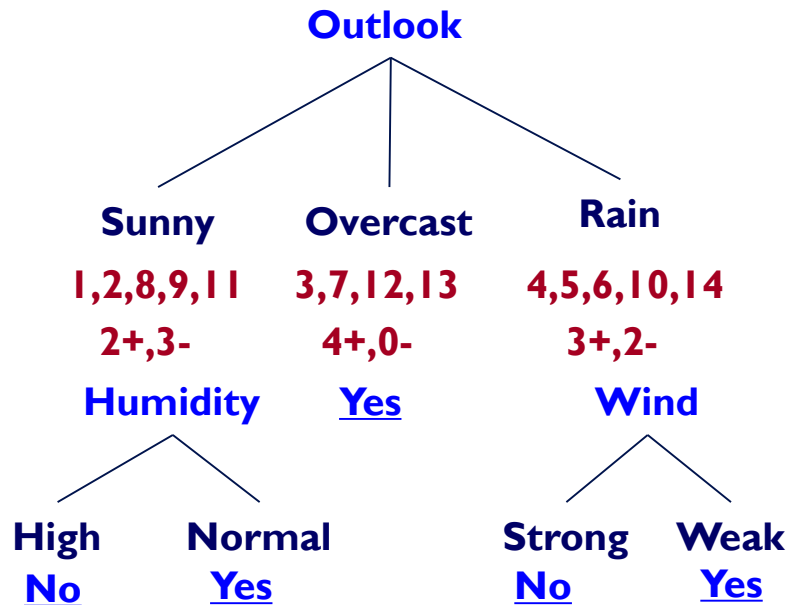
- Hunt and colleagues in Psychology used full search decision tree methods to model human concept learning in the 60s
 - Quinlan developed ID3, with the information gain heuristics in the late 70s to learn expert systems from examples
 - Breiman, Freidman and colleagues in statistics developed CART (classification and regression trees simultaneously)
- A variety of improvements in the 80s: coping with noise, continuous attributes, missing data, non-axis parallel etc.
 - Quinlan's updated algorithm, C4.5 (1993) is commonly used (New: C5)
- Boosting (or Bagging) over DTs is a very good general purpose algorithm



Overfitting

Example

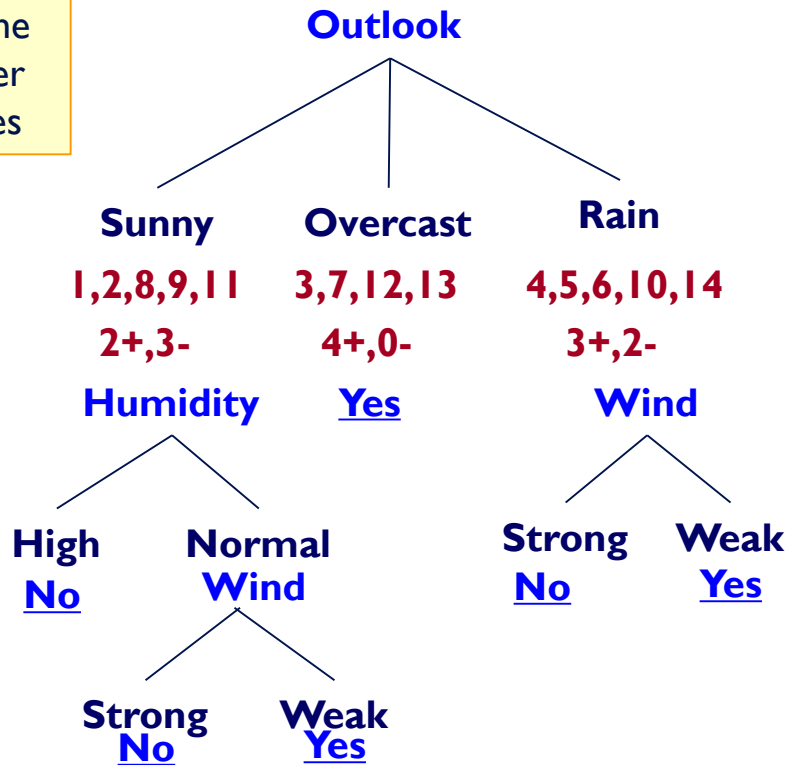
- Outlook = Sunny,
- Temp = Hot
- Humidity = Normal
- Wind = Strong
- label: NO
- this example doesn't exist in the tree



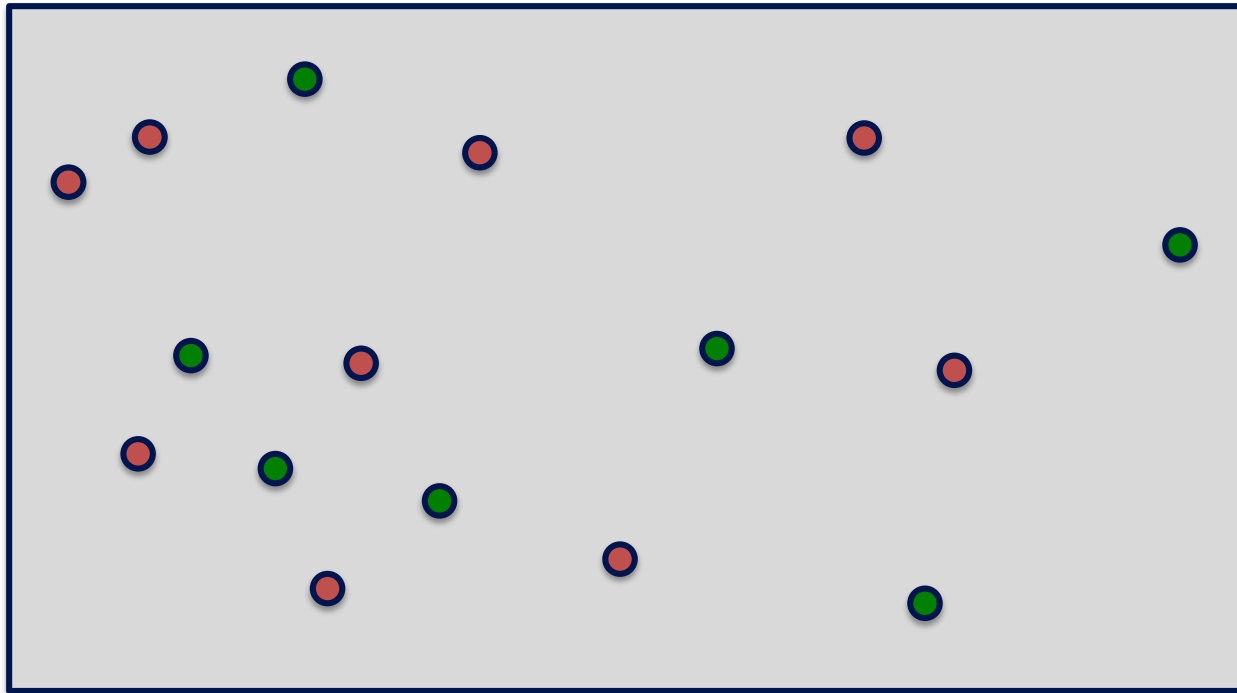
Overfitting - Example

This can always be done
– may fit noise or other
coincidental regularities

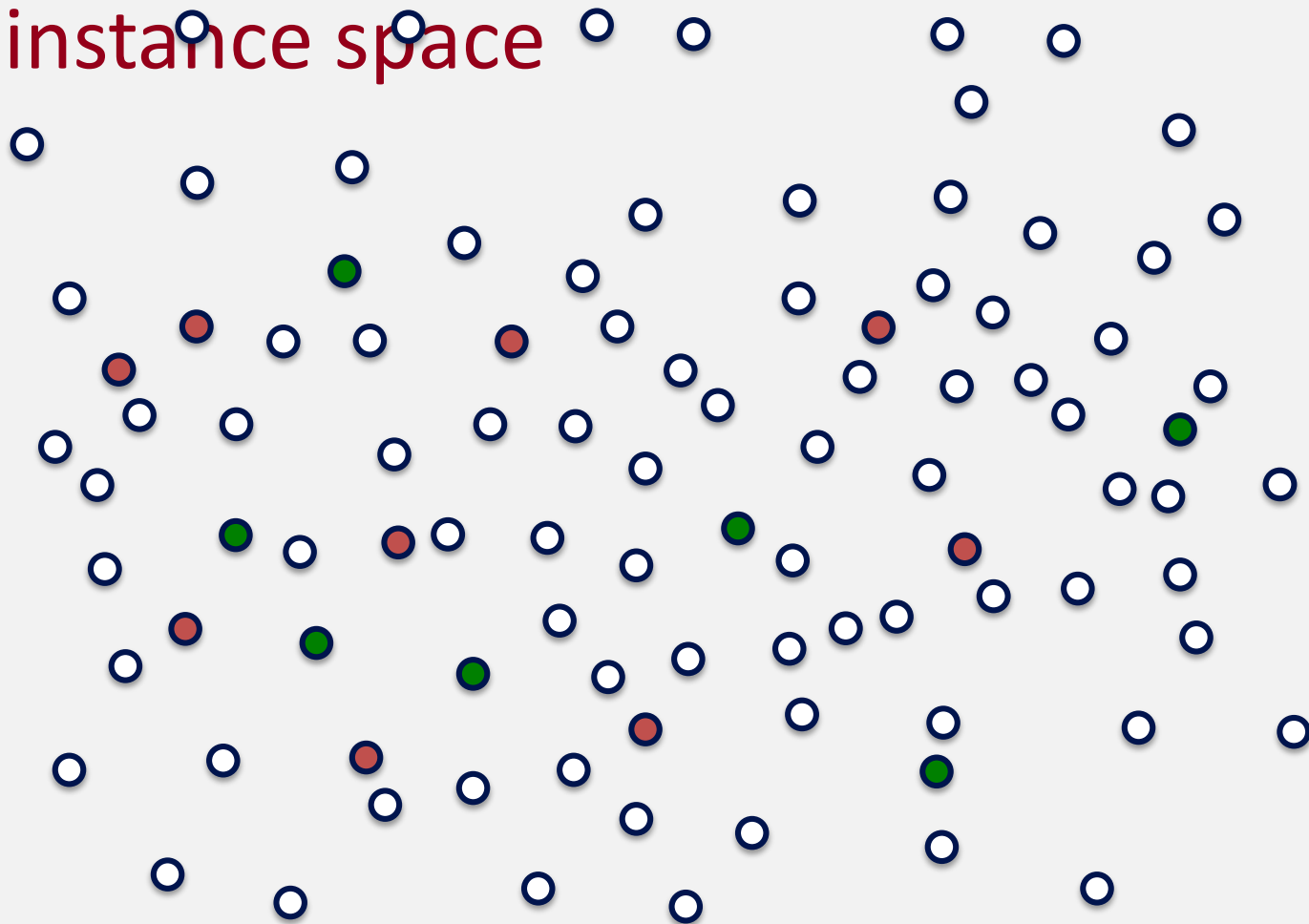
- Outlook = Sunny,
- Temp = Hot
- Humidity = Normal
- Wind = Strong
- label: NO
- this example doesn't exist in the tree



Our training data

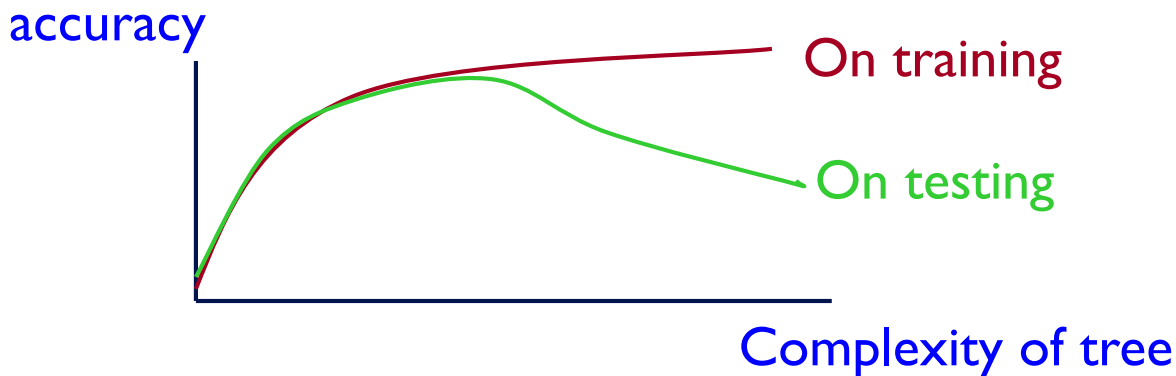


The instance space



Overfitting the Data

- Learning a tree that classifies the training data perfectly may not lead to the tree with the **best generalization performance**.
 - There may be noise in the training data the tree is fitting
 - The algorithm might be making decisions based on very little data
- A hypothesis h is said to **overfit the training data** if there is another hypothesis h' , such that h has a smaller error than h' on the **training data** but h has larger error on the **test data** than h' .



Reasons for overfitting

- **Too much variance** in the training data
 - Training data is not a representative sample of the instance space
 - We split on features that are actually irrelevant
- **Too much noise** in the training data
 - Noise = some feature values or class labels are incorrect
 - We learn to predict the noise
- In both cases, it is a result of our will to **minimize the empirical error** when we learn, and the **ability to do** it (with DTs)

Pruning a decision tree

- Prune = remove leaves and assign majority label of the parent to all items
- Prune the children of node s if:
 - all children are leaves, and
 - the accuracy on the **validation set** does not decrease if we assign the most frequent class label to all items at s .

Avoiding Overfitting

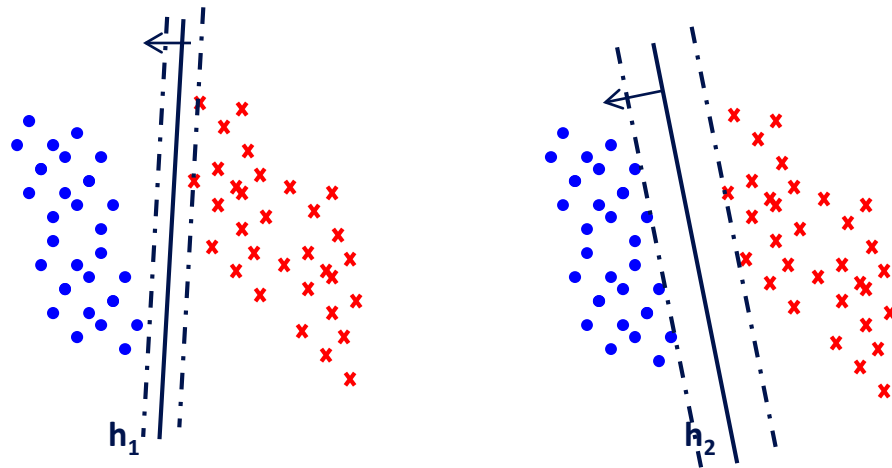
How can this be avoided with linear classifiers?

- Two basic approaches
 - **Pre-pruning**: Stop growing the tree at some point during construction when it is determined that there is not enough data to make reliable choices.
 - **Post-pruning**: Grow the full tree and then remove nodes that seem not to have sufficient evidence.
- Methods for evaluating subtrees to prune
 - **Cross-validation**: Reserve hold-out set to evaluate utility
 - **Statistical testing**: Test if the observed regularity can be dismissed as likely to occur by chance
 - **Minimum Description Length**: Is the additional complexity of the hypothesis smaller than remembering the exceptions?
- This is related to the notion of **regularization** that we will see in other contexts – **keep the hypothesis simple**.

Hand waving, for now.

Next: a brief detour into explaining generalization and overfitting

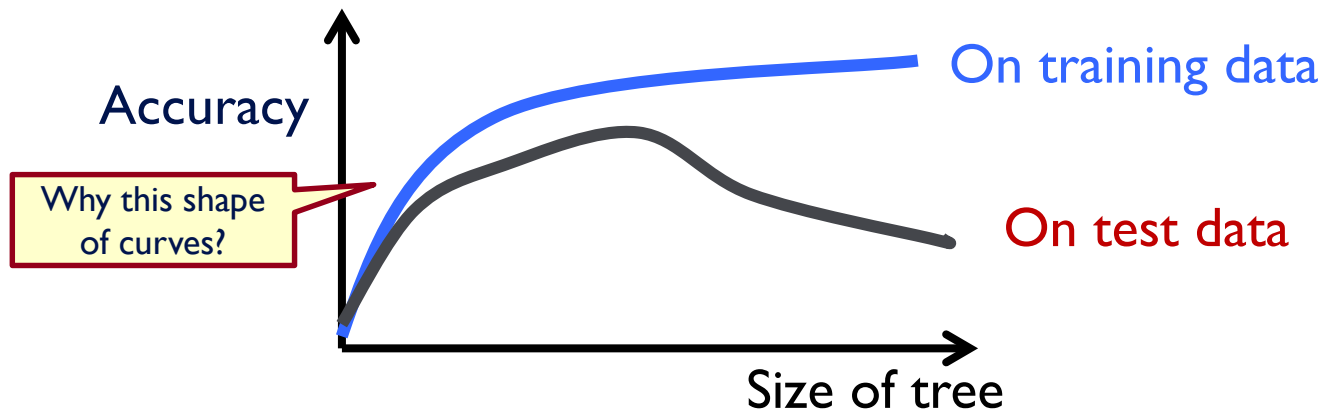
Preventing Overfitting



The i.i.d. assumption

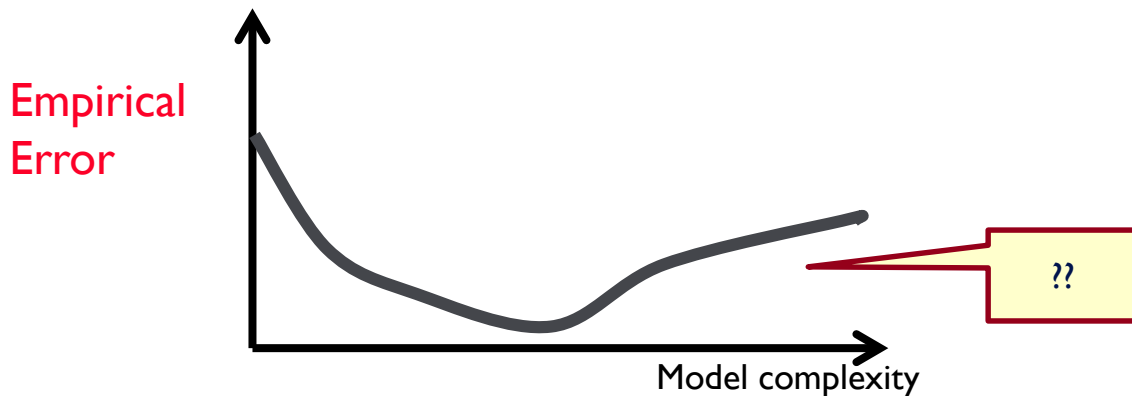
- Training and test items are **independently and identically distributed (i.i.d.)**:
 - There is a distribution $P(\mathbf{X}, Y)$ from which the data $\mathcal{D} = \{(\mathbf{x}, y)\}$ is generated.
 - Sometimes it's useful to rewrite $P(\mathbf{X}, Y)$ as $P(\mathbf{X})P(Y|\mathbf{X})$
Usually $P(\mathbf{X}, Y)$ is unknown to us (we just know it exists)
 - Training and test data are samples drawn from the *same* $P(\mathbf{X}, Y)$: they are **identically distributed**
 - Each (\mathbf{x}, y) is drawn **independently** from $P(\mathbf{X}, Y)$

Overfitting



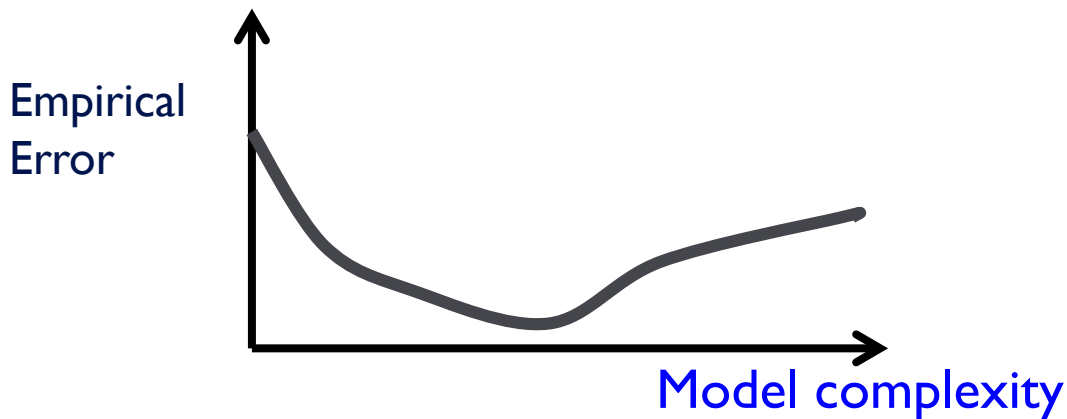
- A decision tree **overfits the training data** when its accuracy on the training data goes up but its accuracy on unseen data goes down

Overfitting



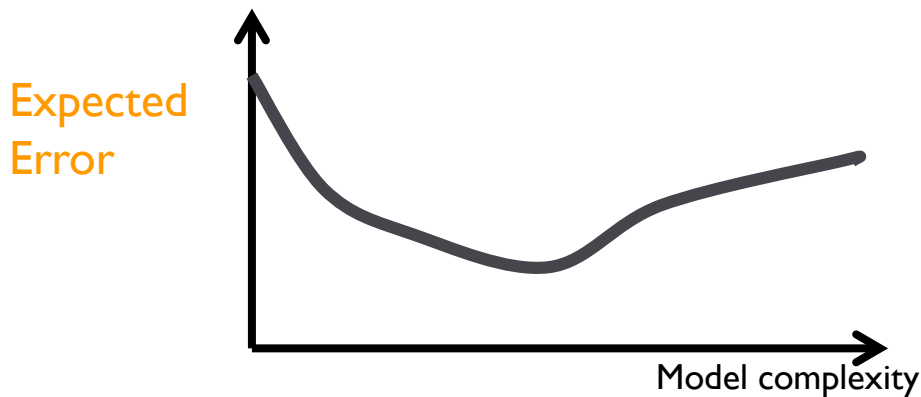
- **Empirical error** (= on a given data set):
The percentage of items in this data set are misclassified by the classifier f .

Overfitting



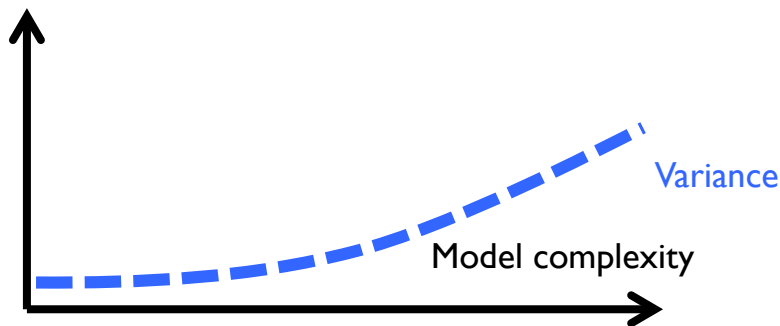
- **Model complexity** (informally):
How many parameters do we have to learn?
 - Decision trees: complexity = #nodes

Overfitting



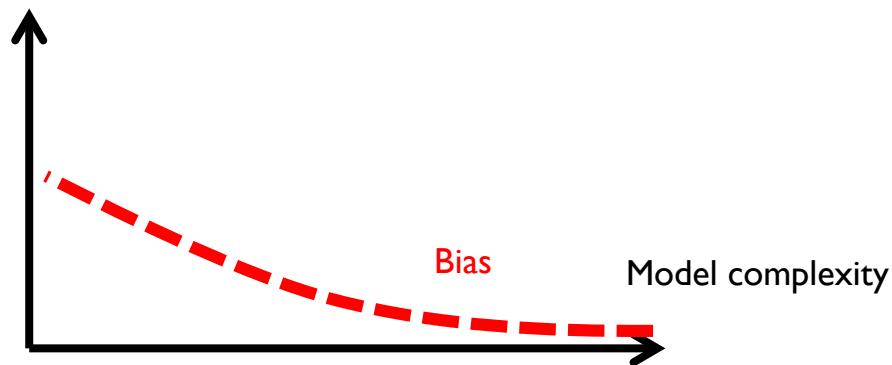
- **Expected error:**
What percentage of items drawn from $P(\mathbf{x}, y)$ do we expect to be misclassified by f ?
- (That's what we really care about – generalization)

Variance of a learner (informally)



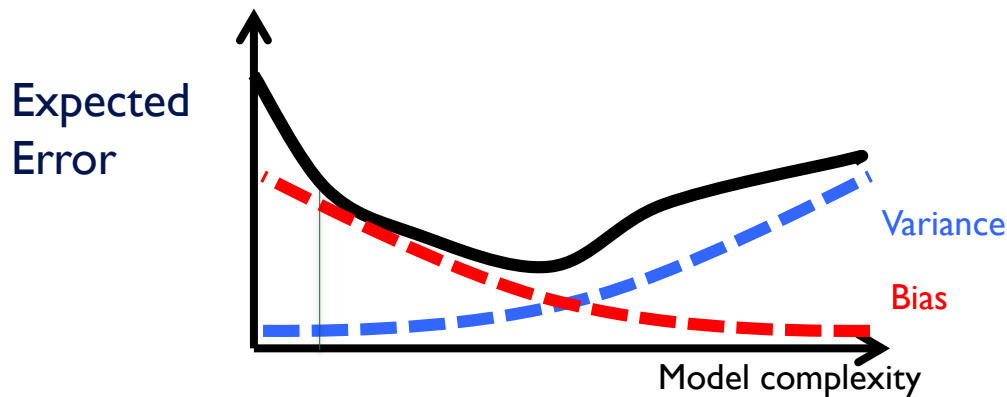
- How susceptible is the learner to minor changes in the training data?
 - (i.e. to different samples from $P(\mathbf{X}, Y)$)
- Variance increases with model complexity
 - Think about **extreme cases**: a hypothesis space with one function vs. all functions.
 - Or, adding the “wind” feature in the DT earlier.
 - The larger the hypothesis space is, the more flexible the selection of the chosen hypothesis is as a function of the data.
 - More accurately: for each data set D , you will learn a different hypothesis $h(D)$, that will have a different true error $e(h)$; we are looking here at the variance of this random variable.

Bias of a learner (informally)



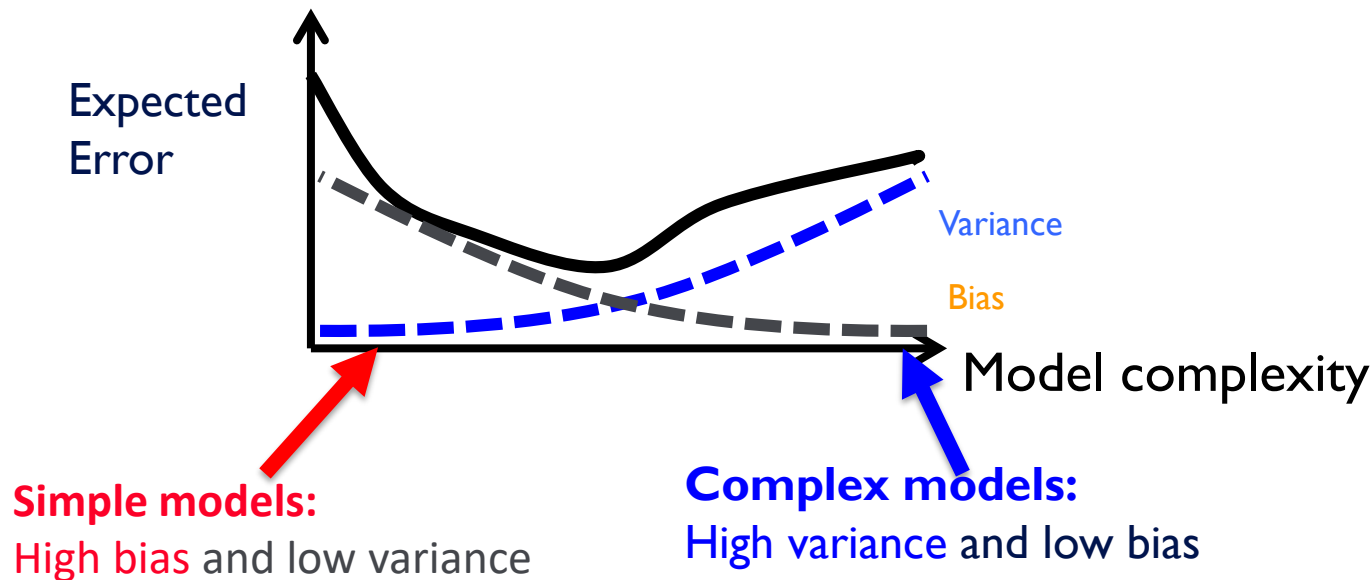
- How likely is the learner to identify the **target** hypothesis?
- Bias is **low** when the model is expressive (low empirical error)
- Bias is **high** when the model is (too) simple
 - The larger the hypothesis space is, the easiest it is to be close to the true hypothesis.
 - More accurately: for each data set D , you learn a different hypothesis $h(D)$, that has a different true error $e(h)$; we are looking here at the difference of the mean of this random variable from the true error.

Impact of bias and variance

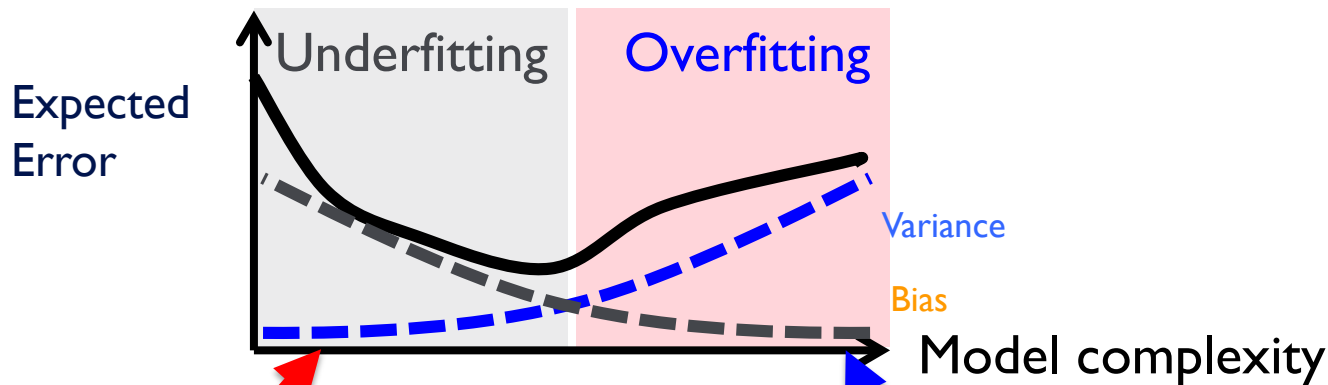


- Expected error \approx bias + variance

Model complexity



Underfitting and Overfitting



Simple models:

High bias and low variance

Complex models:

High variance and low bias

- This can be made more accurate for some loss functions.
- We will discuss a more precise and general theory that trades **expressivity of models** with **empirical error**

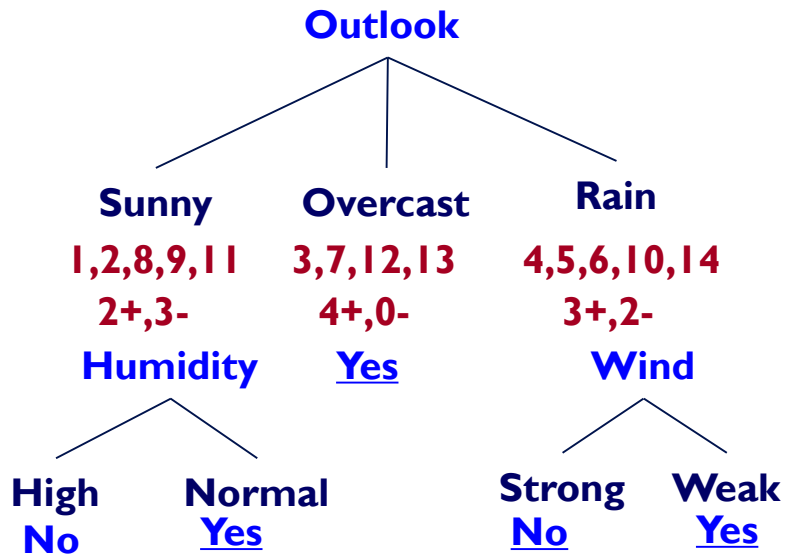
Avoiding Overfitting

How can this be avoided with linear classifiers?

- Two basic approaches
 - Pre-pruning: Stop growing the tree at some point during construction when it is determined that there is not enough data to make reliable choices.
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Trees and Rules

- Decision Trees can be represented as Rules
 - (outlook = sunny) and (humidity = normal) then YES
 - (outlook = rain) and (wind = strong) then NO
- Sometimes Pruning can be done at the **rules level**
 - Rules are generalized by erasing a condition (**different!**)





DT Extensions: continuous attributes and missing values

Continuous Attributes

- Real-valued attributes can, in advance, be discretized into ranges, such as big, medium, small
- Alternatively, one can develop splitting nodes based on thresholds of the form $A < c$ that partition the data into examples that satisfy $A < c$ and $A \geq c$.
 - The information gain for these splits is calculated in the same way and compared to the information gain of discrete splits.
- How to find the split with the highest gain?
- For each continuous feature A:
 - Sort examples according to the value of A
 - For each ordered pair (x,y) with different labels
 - Check the mid-point as a possible threshold, i.e.
 - $S_{a < x} S_{a \geq y}$

Continuous Attributes

- Example:
 - Length (L): 10 15 21 28 32 40 50
 - Class: - + + - + + -
 - Check thresholds: $L < 12.5$; $L < 24.5$; $L < 45$
 - Subset of Examples = {...}, Split = k+, j-
- How to find the split with the highest gain ?
 - For each continuous feature A:
 - Sort examples according to the value of A
 - For each ordered pair (x,y) with different labels
 - Check the mid-point as a possible threshold. I.e,
 - $S_{a < x}$, $S_{a \geq y}$

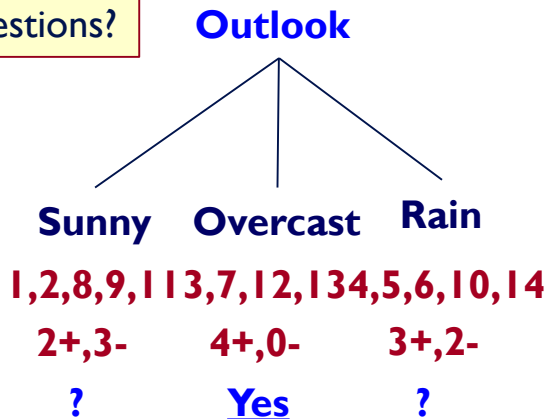
Missing Values

- Diagnosis = < fever, blood_pressure,..., blood_test=?,...>
- Many times values are not available for all attributes during training or testing (e.g., medical diagnosis)
- **Training**: evaluate $\text{Gain}(S,a)$ where in some of the examples a value for a is not given

Missing Values

$$Gain(S, a) = Entropy(S) - \sum \frac{|S_v|}{|S|} Entropy(S_v)$$

Other suggestions?



$$Gain(S_{\text{sunny}}, Temp) = .97 - 0 - (2/5) = .57$$

$$Gain(S_{\text{sunny}}, Humidity) =$$

- Fill in: assign the **most likely value of X_i** to **s**:
 $\text{argmax}_k P(X_i = k)$: **Normal**
 - $.97 - (3/5) Ent[+0, -3] - (2/5) Ent[+2, -0] = .97$
- Assign **fractional counts** $P(X_i = k)$
 for each value of X_i to **s**
 - $.97 - (2.5/5) Ent[+0, -2.5] - (2.5/5) Ent[+2, -.5] < .97$

Day	Outlook	Temperature	Humidity	Wind	PlayTennis
1	Sunny	Hot	High	Weak	No
2	Sunny	Hot	High	Strong	No
8	Sunny	Mild	???	Weak	No
9	Sunny	Cool	Normal	Weak	Yes
11	Sunny	Mild	Normal	Strong	Yes

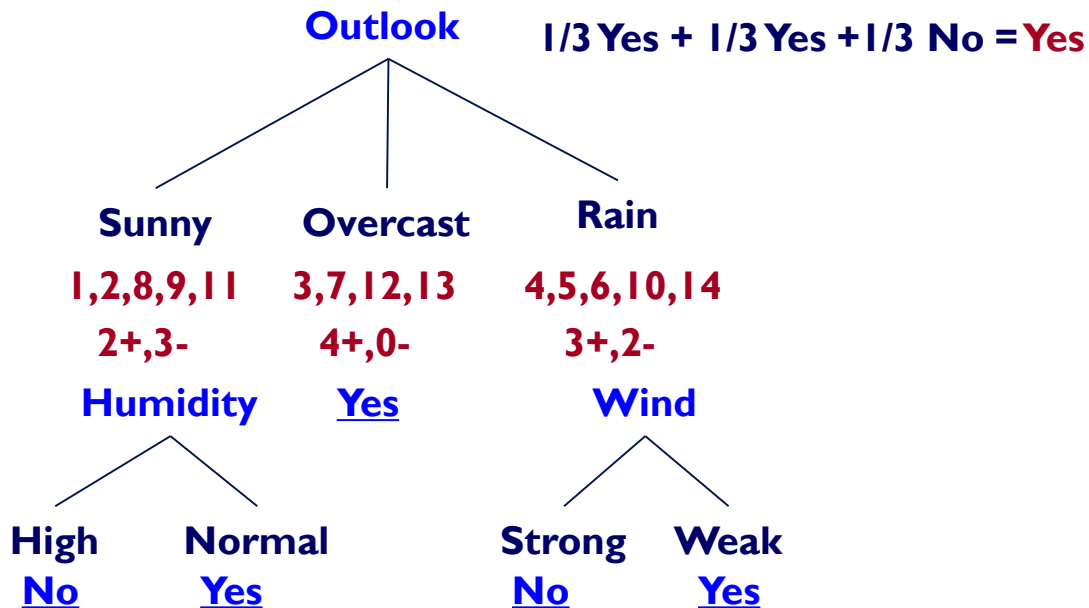
Missing Values

- Diagnosis = $\langle \text{fever, blood_pressure}, \dots, \text{blood_test} = ?, \dots \rangle$
- Many times values are not available for all attributes during training or testing (e.g., medical diagnosis)
- **Training:** evaluate $\text{Gain}(S, a)$ where in some of the examples a value for a is not given
- **Testing:** classify an example without knowing the value of a

Missing Values

Outlook = Sunny, Temp = Hot, Humidity = ???, Wind = Strong, label = ?? Normal/High

Outlook = ???, Temp = Hot, Humidity = Normal, Wind = Strong, label = ??



Other Issues

- Attributes with different costs
 - Change information gain so that low cost attribute are preferred
 - Dealing with features with different # of values
- Alternative measures for selecting attributes
 - When different attributes have different number of values information gain tends to prefer those with many values
- Oblique Decision Trees
 - Decisions are not axis-parallel
- Incremental Decision Trees induction
 - Update an existing decision tree to account for new examples incrementally (Maintain consistency?)

Summary: Decision Trees

- Presented the hypothesis class of Decision Trees
 - Very expressive, flexible, class of functions
- Presented a learning algorithm for Decision Trees
 - Recursive algorithm.
 - Key step is based on the notion of Entropy
- Discussed the notion of overfitting and ways to address it within DTs
 - In your problem set – look at the performance on the training vs. test
- Briefly discussed some extensions
 - Real valued attributes
 - Missing attributes
- Evaluation in machine learning
 - Cross validation
 - Statistical significance

Decision Trees as Features

- Rather than using decision trees to represent the target function it is becoming common to use small decision trees as **features**
- When learning over a large number of features, learning decision trees is difficult and the resulting tree may be very large
→ (over fitting)
- Instead, learn small decision trees, with limited depth.
- Treat them as “experts”; they are correct, but only on a small region in the domain. (**what DTs to learn? same every time?**)
- Then, learn another function, typically a linear function, over these as features.
- Boosting (but also other linear learners) are used on top of the small decision trees. (Either Boolean, or real valued features)
- In HW1 you learn a linear classifier over DTs.
 - Not learning the DTs sequentially; all are learned at once.
 - How can you learn multiple DTs?
 - Combining them using an SGD algorithm.

Experimental Machine Learning

- Machine Learning is an Experimental Field and we will spend some time (in Problem sets) learning how to run experiments and evaluate results
 - First hint: be organized; write scripts
- Basics:
 - Split your data into three sets:
 - Training data (often 70-90%)
 - Test data (often 10-20%)
 - Development data (10-20%)
- You need to report performance on test data, but you are not allowed to look at it.
 - You are allowed to look at the development data (and use it to tune parameters)



Metrics Methodologies Statistical Significance

Metrics

- We train on our training data $\text{Train} = \{x_i, y_i\}_{1,m}$
- We test on **Test data**.
- We often set aside part of the training data as a **development set**, especially when the algorithms require tuning.
 - In the HW we asked you to present results also on the Training; why?
- When we deal with binary classification we often measure performance simply using **Accuracy**:

$$\text{accuracy} = \frac{\# \text{ correct predictions}}{\# \text{ test instances}}$$

$$\text{error} = 1 - \text{accuracy} = \frac{\# \text{ incorrect predictions}}{\# \text{ test instances}}$$

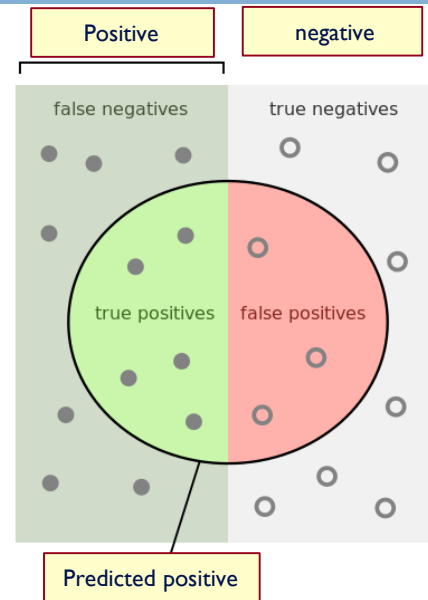
- Any possible problems with it?

Alternative Metrics

- If the Binary classification problem is biased
 - In many problems most examples are negative
- Or, in multiclass classification
 - The distribution over labels is often non-uniform
- Simple accuracy is not a useful metric.
 - Often we resort to task specific metrics
- However one important example that is being used often involves **Recall** and **Precision**

• **Recall:**
$$\frac{\text{\# (positive identified = true positives)}}{\text{\# (all positive)}}$$

• **Precision:**
$$\frac{\text{\# (positive identified = true positives)}}{\text{\# (predicted positive)}}$$



How many selected items are relevant?

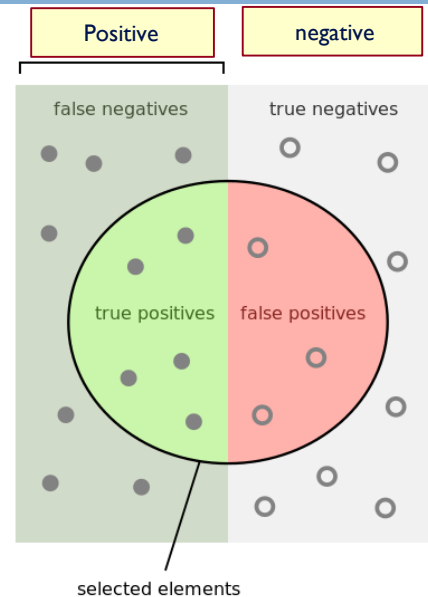
$$\text{Precision} = \frac{\text{true positives}}{\text{true positives} + \text{false positives}}$$

How many relevant items are selected?

$$\text{Recall} = \frac{\text{true positives}}{\text{true positives} + \text{false negatives}}$$

Example

- 100 examples, 5% are positive.
- Just say NO: your accuracy is 95%
 - Recall = precision = 0
- Predict 4+, 96-; 2 of the +s are indeed positive
 - Recall: 2/5; Precision: 2/4
- Recall: $\frac{\# \text{ (positive identified = true positives)}}{\# \text{ (all positive)}}$
- Precision: $\frac{\# \text{ (positive identified = true positives)}}{\# \text{ (predicted positive)}}$



How many selected items are relevant?

$$\text{Precision} = \frac{\text{true positives}}{\text{true positives} + \text{false positives}}$$

How many relevant items are selected?

$$\text{Recall} = \frac{\text{true positives}}{\text{true positives} + \text{false negatives}}$$

Confusion Matrix

- Given a dataset of P positive instances and N negative instances:

The notion of a confusion matrix can be usefully extended to the multiclass case
(i, j) cell indicate how many of the i -labeled examples were predicted to be j

		Predicted Class	
		Yes	No
Actual Class	Yes	TP	FN
	No	FP	TN

$$\text{accuracy} = \frac{TP + TN}{P + N}$$

- Imagine using classifier to identify positive cases (i.e., for information retrieval)

$$\text{precision} = \frac{TP}{TP + FP}$$

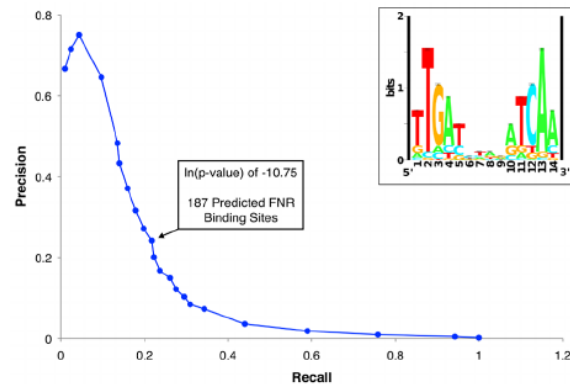
Probability that a randomly selected positive prediction is indeed positive

$$\text{recall} = \frac{TP}{TP + FN}$$

Probability that a randomly selected positive is identified

Relevant Metrics

- It makes sense to consider Recall and Precision together or combine them into a single metric.
- Recall-Precision Curve:
- F-Measure:
 - A measure that combines precision and recall is the harmonic mean of precision and recall.
 - F1 is the most commonly used metric.



$$F_{\beta} = (1 + \beta^2) \cdot \frac{\text{precision} \cdot \text{recall}}{\beta^2 \cdot \text{precision} + \text{recall}}$$

Comparing Classifiers

Say we have two classifiers, $C1$ and $C2$, and want to choose the best one to use for future predictions

Can we use training accuracy to choose between them?

- No!
- What about accuracy on test data?

N-fold cross validation

- Instead of a single test-training split:



- Split data into N equal-sized parts



- Train and test N different classifiers
- Report average accuracy and standard deviation of the accuracy

Evaluation: significance tests

- You have two different classifiers, A and B
- You train and test them on the same data set using N-fold cross-validation
- For the n -th fold:
 $\text{accuracy}(A, n), \text{accuracy}(B, n)$
 $p_n = \text{accuracy}(A, n) - \text{accuracy}(B, n)$
- Is the difference between A and B's accuracies significant?



Hypothesis testing

- You want to show that **hypothesis H is true**, based on your data
 - (e.g. H = “classifier A and B are different”)
- Define a **null hypothesis H_0**
 - (H_0 is the contrary of what you want to show)
- **H_0 defines a distribution $P(m / H_0)$ over some statistic**
 - e.g. a distribution over the difference in accuracy between A and B
- **Can you refute (reject) H_0 ?**

Rejecting H_0

- H_0 defines a distribution $P(M / H_0)$ over some statistic M
 - (e.g. M = the difference in accuracy between A and B)
- Select a significance value S
 - (e.g. 0.05, 0.01, etc.)
 - You can only reject H_0 if $P(m / H_0) \leq S$
- Compute the test statistic m from your data
 - e.g. the average difference in accuracy over your N folds
- Compute $P(m / H_0)$
- Refute H_0 with $p \leq S$ if $P(m / H_0) \leq S$

Paired t-test

- Null hypothesis (H_0 ; to be refuted):
 - There is no difference between A and B, i.e. the expected accuracies of A and B are the same
- That is, the expected difference (over all possible data sets) between their accuracies is 0:
 $H_0: E[p_D] = 0$
- We don't know the true $E[p_D]$
- N -fold cross-validation gives us N samples of p_D

Paired t-test

- Null hypothesis $H_0: E[\text{diff}_D] = \mu = 0$
- m : our estimate of μ based on N samples of diff_D
$$m = 1/N \sum_n \text{diff}_n$$
- The estimated variance S^2 :
$$S^2 = 1/(N-1) \sum_{1,N} (\text{diff}_n - m)^2$$
- **Accept Null hypothesis** at significance level α if the **following statistic** lies in $(-t_{\alpha/2, N-1}, +t_{\alpha/2, N-1})$

$$\frac{\sqrt{Nm}}{S} \sim t_{N-1}$$

Decision Trees - Summary

- Hypothesis Space:
 - Variable size (contains all functions)
 - Deterministic; Discrete and Continuous attributes
- Search Algorithm
 - ID3 - batch
 - Extensions: missing values
- Issues:
 - What is the goal?
 - When to stop? How to guarantee good generalization?
- Did not address:
 - How are we doing? (Correctness-wise, Complexity-wise)