

Technische Universiteit
Eindhoven
University of Technology

Sheets are largely based on the those provided by Tan, Steinbach, and Kumar. *Introduction to Data Mining*

Where innovation starts

Classification: Definition

- Given a training set
 - Relation over attributes, one of the attributes is the class.
- Find a model for the class attribute
- That allows for:
 - predicting <u>previously unseen</u> records accurately.



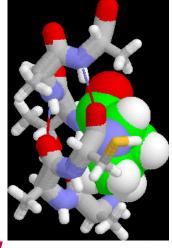
Examples of Classification Task

- Predicting tumor cells as benign or malignant
- Classifying credit card transactions as legitimate or fraudulent



 Classifying secondary structures of protein as alpha-helix, beta-sheet, or random coil

 Categorizing news stories as finance, weather, entertainment, sports, etc



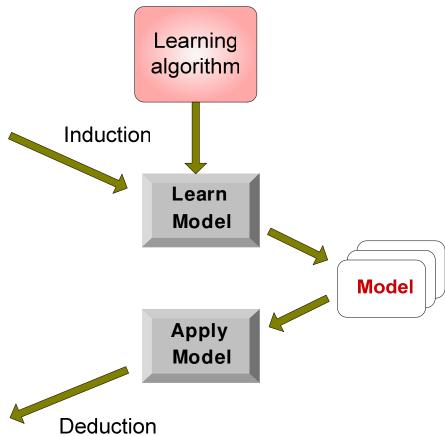
Illustrating Classification Task



Training Set

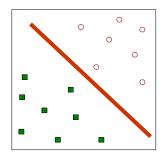
Tid	Attrib1	Attrib2	Attrib3	Class
11	No	Small	55K	?
12	Yes	Medium	80K	?
13	Yes	Large	110K	?
14	No	Small	95K	?
15	No	Large	67K	?

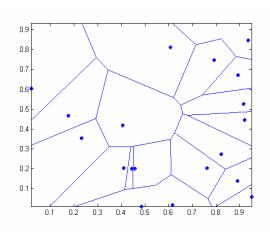
Test Set

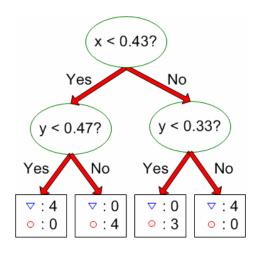




Many different types of models







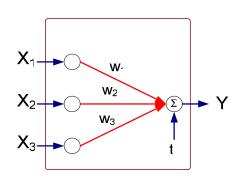
R1: (Give Birth = no) \land (Can Fly = yes) \rightarrow Birds

R2: (Give Birth = no) \land (Live in Water = yes) \rightarrow Fishes

R3: (Give Birth = yes) \land (Blood Type = warm) \rightarrow Mammals

R4: (Give Birth = no) \land (Can Fly = no) \rightarrow Reptiles

R5: (Live in Water = sometimes) → Amphibians





Outline

- K-nearest neighbors
 - Distance measures
- Decision trees
 - Induction of a decision tree
 - Hunt's algorithm
 - Issues with decision trees



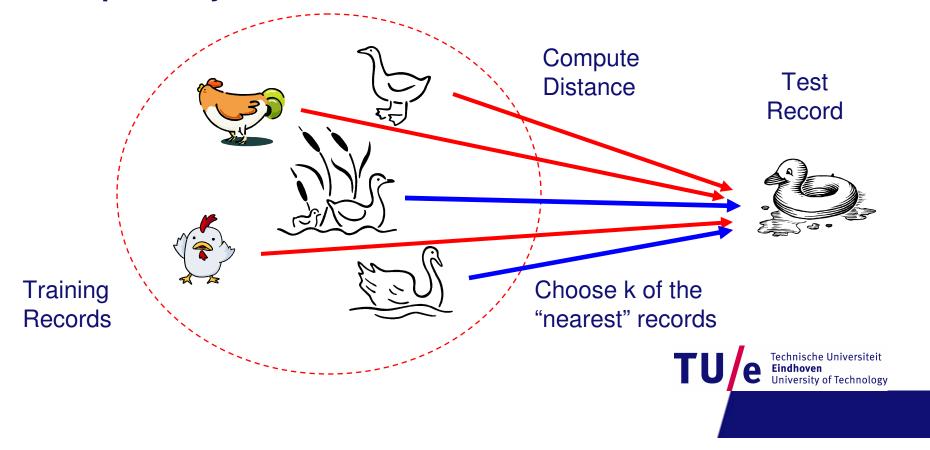
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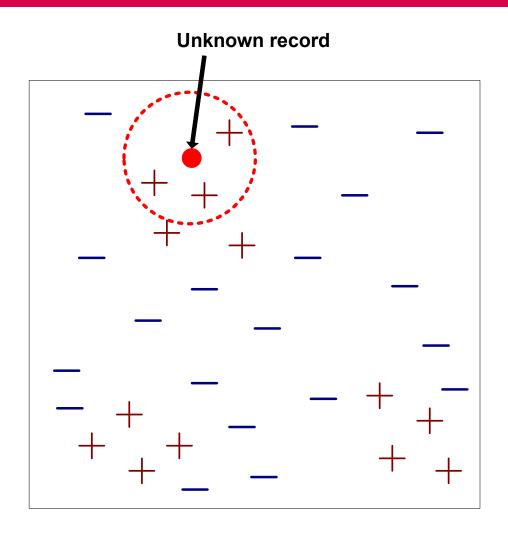


Nearest Neighbor Classifiers

- Basic idea:
 - If it walks like a duck, quacks like a duck, then it's probably a duck



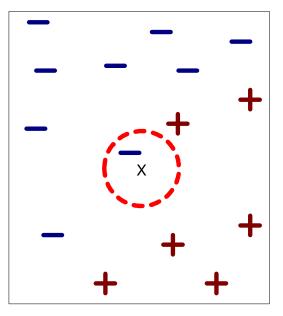
Nearest-Neighbor Classifiers

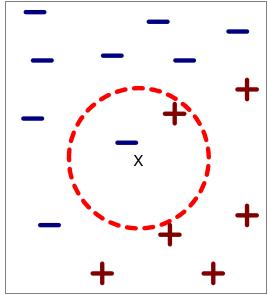


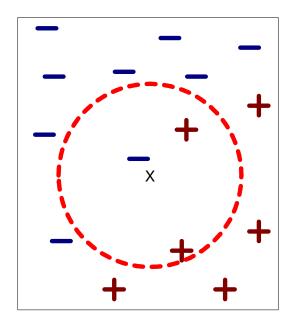
- Requires three things
 - The set of stored records
 - Distance Metric to compute distance between records
 - The value of k, the number of nearest neighbors to retrieve
- To classify an unknown record:
 - Compute distance to other training records
 - Identify k nearest neighbors
 - Use class labels of nearest neighbors to determine the class label of unknown record



Definition of Nearest Neighbor





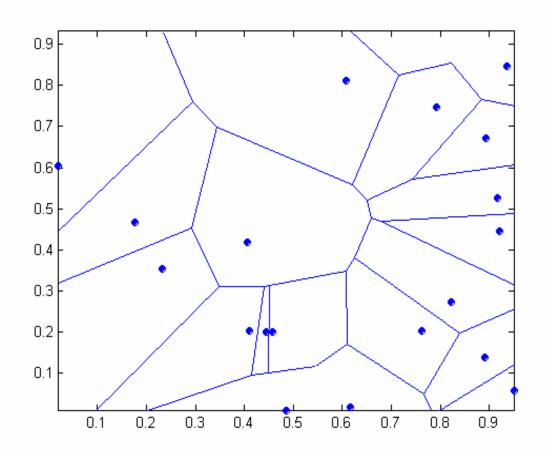


- (a) 1-nearest neighbor
- (b) 2-nearest neighbor
- (c) 3-nearest neighbor

K-nearest neighbors of a record x are data points that have the k smallest distance to x

1 nearest-neighbor

Voronoi Diagram





Nearest Neighbor Classification

- Compute distance between two points:
 - Euclidean distance

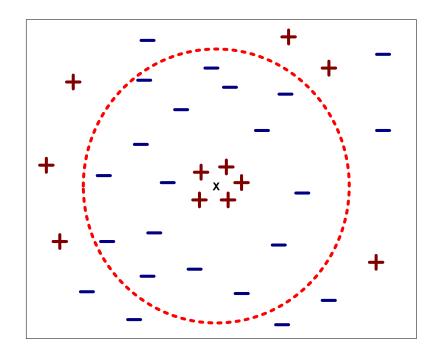
$$d(p,q) = \sqrt{\sum_{i} (p_{i} - q_{i})^{2}}$$

- Determine the class from nearest neighbor list
 - take the majority vote of class labels among the knearest neighbors
 - Weigh the vote according to distance
 - weight factor, $w = 1/d^2$



Nearest Neighbor Classification...

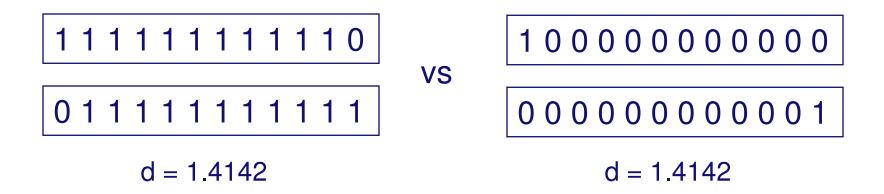
- Choosing the value of k:
 - If k is too small, sensitive to noise points
 - If k is too large, neighborhood may include points from other classes





Nearest Neighbor Classification...

- Problem with Euclidean measure:
 - High dimensional data
 - curse of dimensionality
 - Can produce counter-intuitive results





Outline

- K-nearest neighbors
 - Distance measures
 - Example: for strings and sequences
- Decision trees
 - Induction of a decision tree
 - Hunt's algorithm
 - Issues with Decision trees



Distance Measures ...

- Choosing the correct distance function is essential
 - Eucledian, Minkowski
 - Mahalanobis
 - Simple Matching Coefficient
 - Jaccard measure
 - Tanimoto Coefficient
 - Cosine Measure
- Example: distance measure for strings



Edit Distance

- Distance between two strings: minimal number of operations to transform one into another
 - Insert a character
 - Delete a character
 - Replace a character with aother
- Example:
 - paard → paad → parade distance = 3
 - eauivlaent → equivalent → equivalent distance = 3



Edit Distance: Algorithm

E

A A R P A R Fill a matrix entry i,j: edit distance between t[1..i] en s[1..j] A D

Edit distance: algoritme

```
A A
                                           R
P
Α
        2
R
        3
                 Filling the matrix: recursively
                 d[i,j] = min \{ d(i-1, j) + 1 \quad (del) \}
A
                               d(i,j-1) + 1 (ins)
        4
                               d(i-1,j-1) + cost }
                                   (match of subst.)
        5
D
E
        6
```

Edit distance: algoritme

	_	Р	Α	Α	R	D	
_	0	1	2	3	4	5	
Р	1	0	1	2	3	4	
Α	2	1	0	1	2	3	
R	3	2	1	1	1	2	
Α	4	3	2	1	2	2	
D	5	4	3	2	2	2	
Е	6	5	4	3	3 T	Technische Eindhoven University	Universiteit of Technology

Distance for DNA Sequences

- Matching in BLAST (Basic Local Alignment and Search Tool) is based on this type of match
- Similarity is defined as the maximal match

```
ATGGCGT
*** !**
ATG-AGT
```

- Not every replacement is equally likely
 - Evolutionary theory



BLOSUM62 Substitution Matrix

```
Y2-2-32-1-2-33-22-7-1
                                                   E-1002-42520-33123-10132-2
                                                                                G02013226244233202233
                                                                                                       H20113002833121212223
                                                                                                                                                                                                                                                                                                                                                      W33442232232311432123
C033393433112311221
                          Q11003522032103101212
                                                                                                                                                               L-23-4-12-3-4-220-32-12
                                                                                                                                                                                           K12013112132513101322
                                                                                                                                                                                                                     M -1 -2 3 -1 0 2 3 2 1 2 1 5 0 2 1 -1 1
                                                                                                                                                                                                                                                                           P12213112233124711432
                                                                                                                                                                                                                                                                                                                                                                                                                03331211220314
```



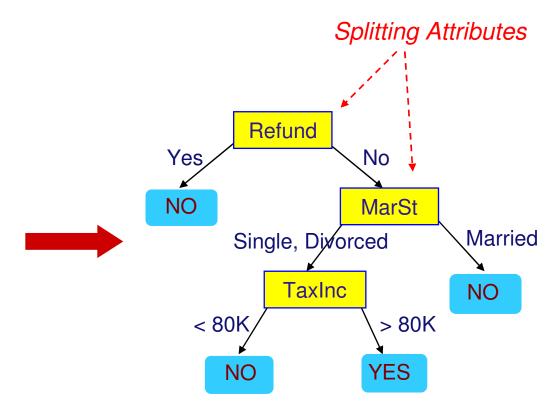
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Example of a Decision Tree

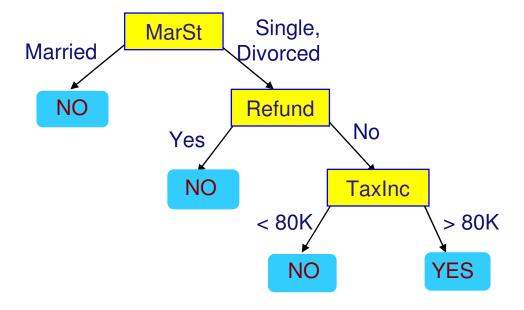
Tid	Refund	Marital Status	Taxable Income	Cheat
1	Yes	Single	125K	No
2	No	Married	100K	No
3	No	Single	70K	No
4	Yes	Married	120K	No
5	No	Divorced	95K	Yes
6	No	Married	60K	No
7	Yes	Divorced	220K	No
8	No	Single	85K	Yes
9	No	Married	75K	No
10	No	Single	90K	Yes





Another Example of Decision Tree

Tid	Refund	Marital Status	Taxable Income	Cheat
1	Yes	Single	125K	No
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3	No	Single	70K	No
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5	No	Divorced	95K	Yes
6	No	Married	60K	No
7	Yes	Divorced	220K	No
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10	No	Single	90K	Yes



There could be more than one tree that fits the same data!



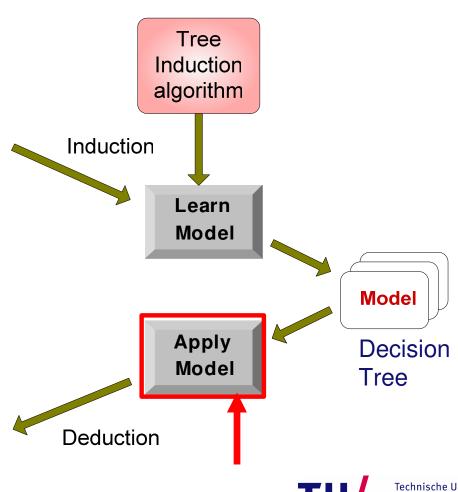
Decision Tree Classification Task



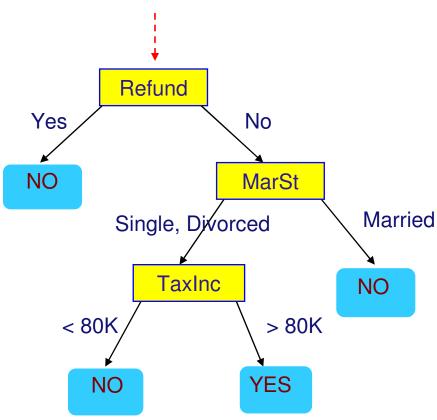
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Test Set

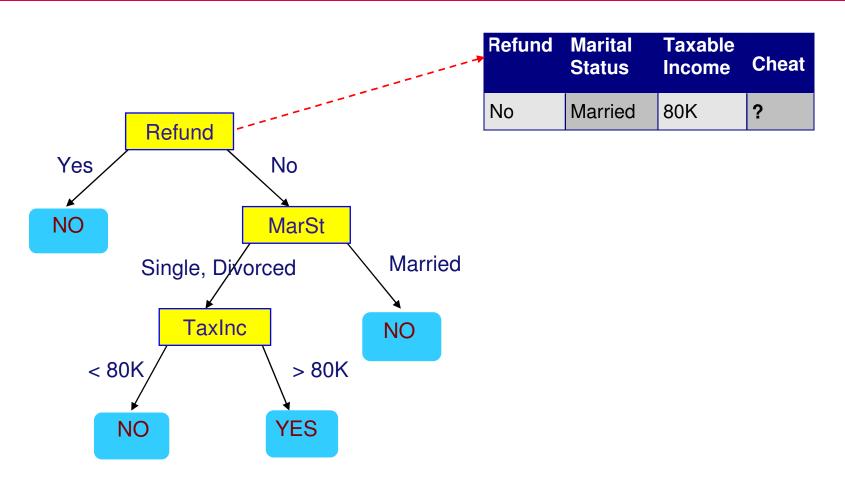


Start from the root of tree.

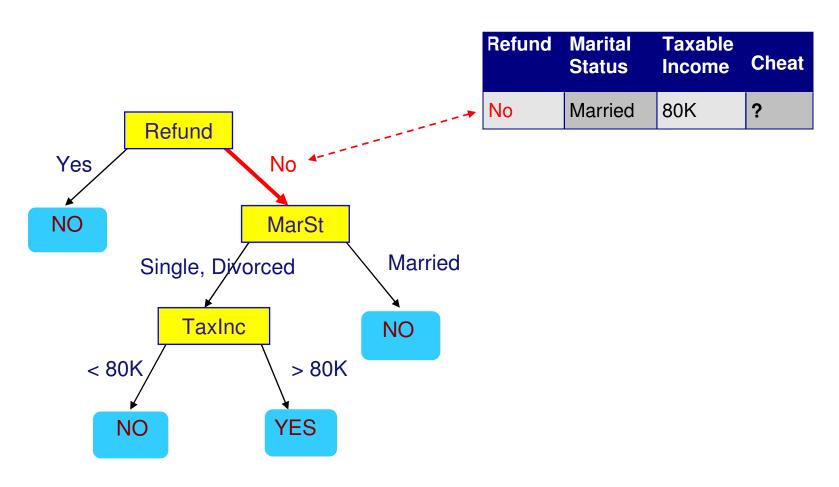


Refund		Taxable Income	Cheat
No	Married	80K	?

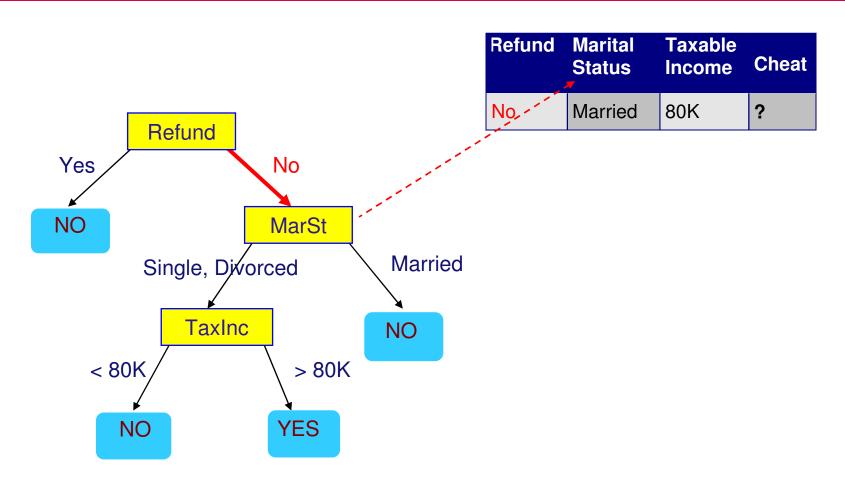




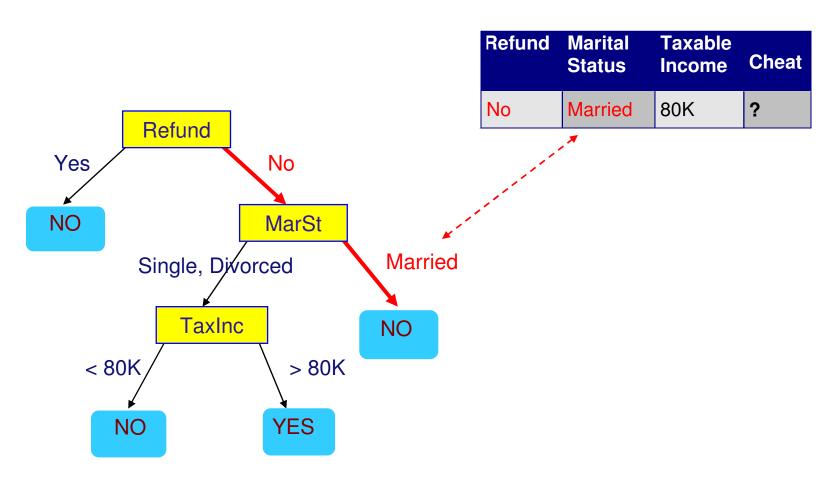




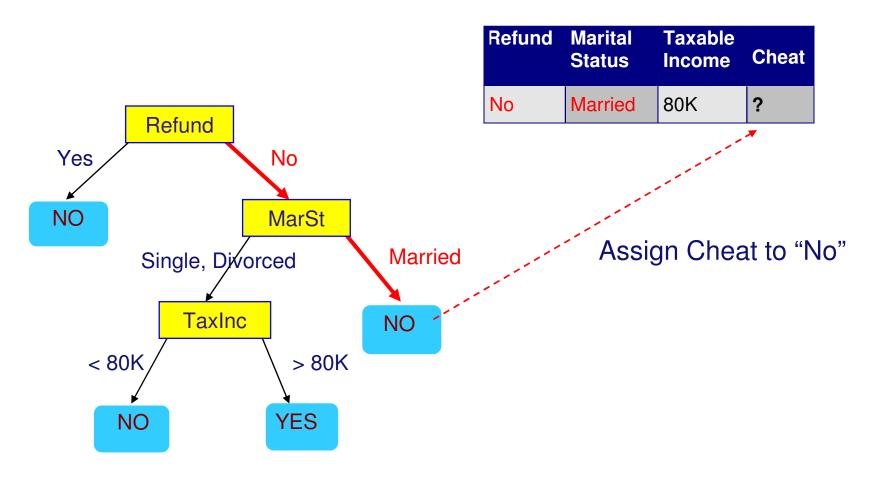














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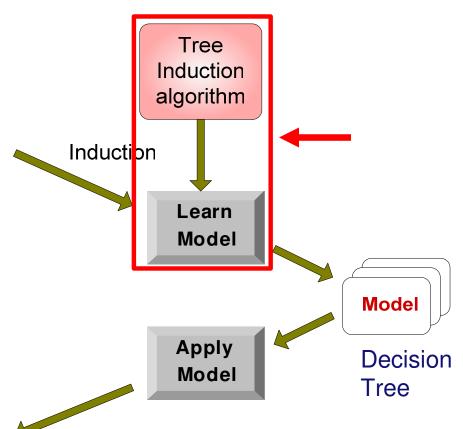
Decision Tree Classification Task



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Test Set



Deduction



General Structure of Hunt's Algorithm

Input: Dataset D

Output: Decision tree t

Induce(D):

If all tuples t in D have label + then

return



If all tuples t in D have label - then

return <



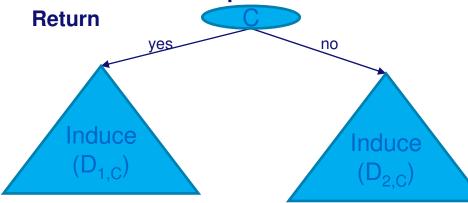
For all split criteria C:

 $D_{1,C} = \{ t \text{ in } D \mid t \text{ satisfies } C \}$

 $D_{2,C} = D - D_1$

Measure Quality(D_1 , D_2)

Let C be the best split



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Tree Induction

- Greedy strategy.
 - Split the records based on an attribute test that optimizes certain criterion.
- Issues
 - Determine how to split the records
 - How to specify the attribute test condition?
 - How to determine the best split?
 - Determine when to stop splitting



Tree Induction

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How to Specify Test Condition?

- Depends on attribute types
 - Nominal
 - Ordinal
 - Continuous
- Depends on number of ways to split
 - 2-way split
 - Multi-way split

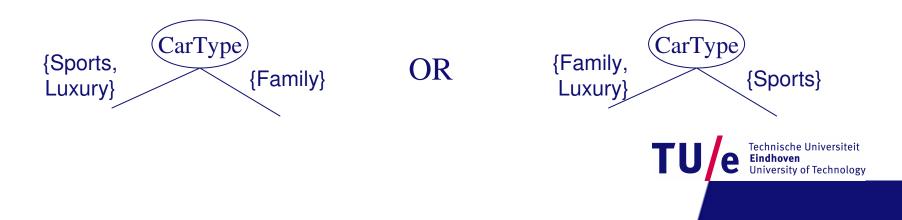


Splitting Based on Nominal Attributes

Multi-way split: Use as many partitions as distinct values.



Binary split: Divides values into two subsets.
 Need to find optimal partitioning.



Splitting Based on Ordinal Attributes

Multi-way split: Use as many partitions as distinct values.

Size

Large

Binary split: Divides values into two subsets.
 Need to find optimal partitioning.

Medium

Small



What about this split?



Splitting Based on Continuous Attributes

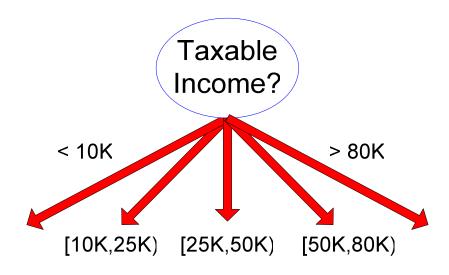
- Different ways of handling
 - Discretization to form an ordinal categorical attribute
 - Static discretize once at the beginning
 - Dynamic ranges can be found by equal interval bucketing, equal frequency bucketing (percentiles), or clustering.
 - Binary Decision: (A < v) or (A ≥ v)
 - consider all possible splits and finds the best cut
 - can be more compute intensive



Splitting Based on Continuous Attributes



(i) Binary split



(ii) Multi-way split



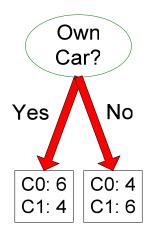
Tree Induction

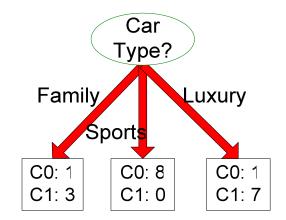
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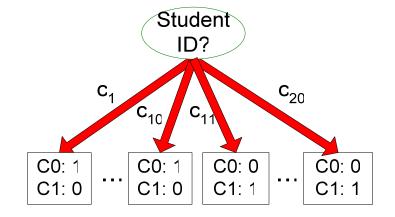


How to determine the Best Split

Before Splitting: 10 records of class 0, 10 records of class 1







Which test condition is the best?



How to determine the Best Split

- Greedy approach:
 - Nodes with homogeneous class distribution are preferred
- Need a measure of node impurity:

C0: 5

C1: 5

Non-homogeneous,

High degree of impurity

C0: 9

C1: 1

Homogeneous,

Low degree of impurity

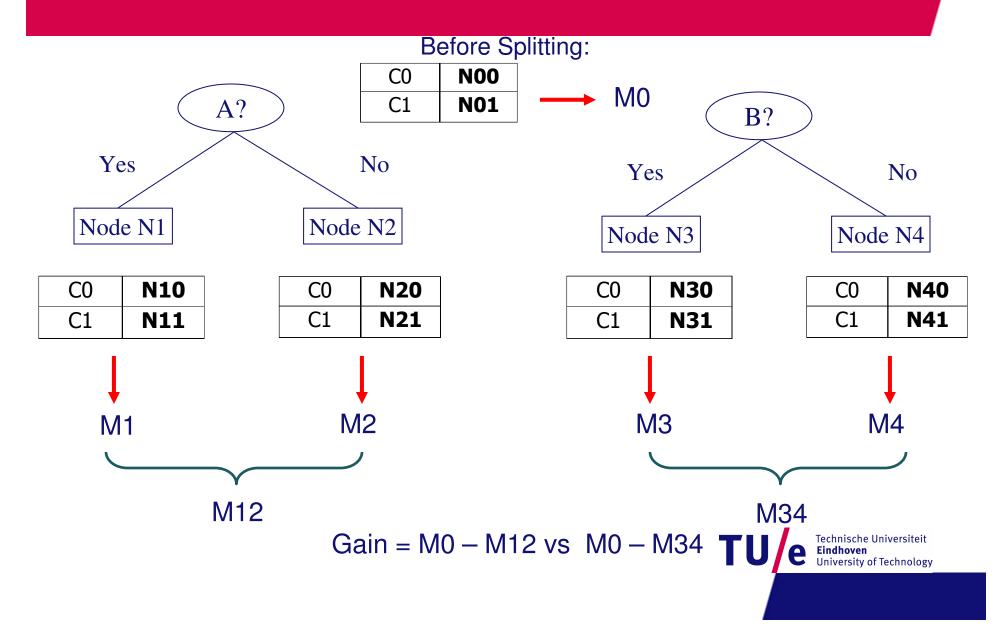


Measures of Node Impurity

- Gini Index
- Entropy
- Misclassification error



How to Find the Best Split



Measure of Impurity: GINI

Gini Index for a given node t :

$$GINI(t) = 1 - \sum_{j} [p(j|t)]^{2}$$

(NOTE: $p(j \mid t)$ is the relative frequency of class j at node t).

- Maximum (1 1/n_c) (records equally distributed)
- Minimum 0 (all records in one class)

C1	0	
C2	6	
Gini=0.000		

C1	3
C2	3
Gini=0.500	



Examples for computing GINI

$$GINI(t) = 1 - \sum_{j} [p(j|t)]^{2}$$

$$P(C1) = 0/6 = 0$$
 $P(C2) = 6/6 = 1$

Gini =
$$1 - P(C1)^2 - P(C2)^2 = 1 - 0 - 1 = 0$$

$$P(C1) = 1/6$$
 $P(C2) = 5/6$

Gini =
$$1 - (1/6)^2 - (5/6)^2 = 0.278$$

$$P(C1) = 2/6$$
 $P(C2) = 4/6$

Gini =
$$1 - (2/6)^2 - (4/6)^2 = 0.444$$



Splitting Based on GINI

- Used in CART, SLIQ, SPRINT.
- When a node p is split into k partitions (children), the quality of split is computed as,

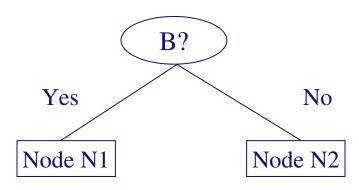
$$GINI_{split} = \sum_{i=1}^{k} \frac{n_i}{n} GINI(i)$$

where, n_i = number of records at child i, n_i = number of records at node p.



Binary Attributes: Computing GINI Index

- Splits into two partitions
- Effect of Weighing partitions:
 - Larger and Purer Partitions are sought for.



	Parent
C1	6
C2	6
Gini	= 0.500

Gini(N1)

$$= 1 - (5/7)^2 - (2/7)^2$$

= 0.408

Gini(N2)

$$= 1 - (1/5)^2 - (4/5)^2$$

= 0.32

	N1	N2
C1	5	1
C2	2	4
Gini=0.333		

Gini(Children)

= 7/12 * 0.408 +

5/12 * 0.32

= 0.371



Categorical Attributes: Computing Gini Index

- For each distinct value, gather counts for each class in the dataset
- Use the count matrix to make decisions

Multi-way split

Two-way split (find best partition of values)

	CarType		
	Family	Sports	Luxury
C1	1	2	1
C2	4	1	1
Gini	0.393		

	CarType	
	{Sports, Luxury}	{Family}
C1	3	1
C2	2	4
Gini	0.400	

	CarType	
	{Sports}	{Family, Luxury}
C1	2	2
C2	1 5	
Gini	0.419	



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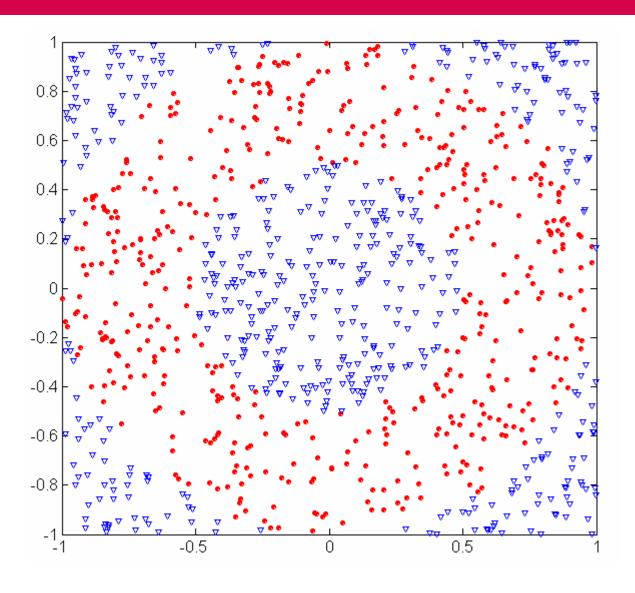


Practical Issues of Classification

- Underfitting and Overfitting
- Missing Values
- Costs of Classification



Underfitting and Overfitting (Example)



500 circular and 500 triangular data points.

Circular points:

$$0.5 \le \text{sqrt}(x_1^2 + x_2^2) \le 1$$

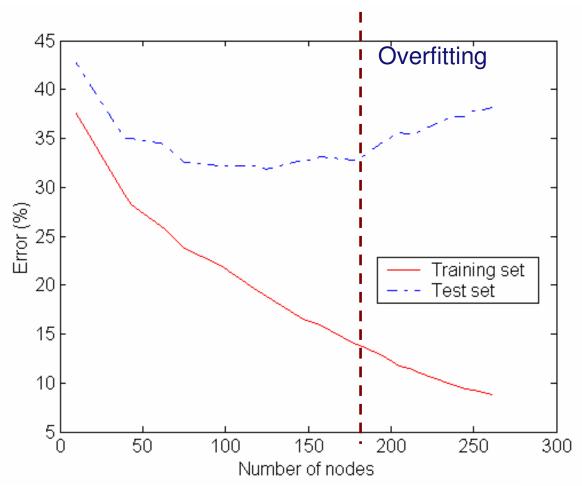
Triangular points:

$$sqrt(x_1^2 + x_2^2) > 0.5 or$$

$$sqrt(x_1^2 + x_2^2) < 1$$

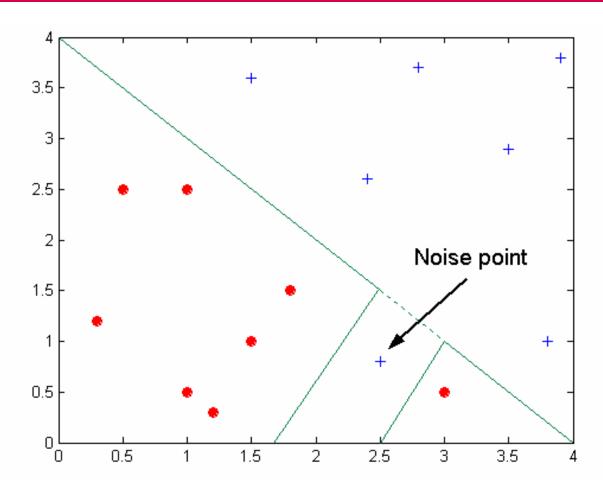


Underfitting and Overfitting





Overfitting due to Noise



Decision boundary is distorted by noise point



Notes on Overfitting

- Overfitting results in decision trees that are more complex than necessary
- Training error no longer provides a good estimate of how well the tree will perform on previously unseen records
- Need new ways for estimating errors



How to Address Overfitting

- Pre-Pruning (Early Stopping Rule)
 - Stop the algorithm before it becomes a fully-grown tree
 - Typical stopping conditions for a node:
 - all instances belong to the same class
 - all the attribute values are the same
 - More restrictive conditions:
 - if number of instances becomes too small
 - If class distribution becomes independent of attributes
 - If expanding the current node does not improve impurity measures.



How to Address Overfitting...

Post-pruning

- Grow decision tree to its entirety
- Trim the nodes of the decision tree in a bottom-up fashion
- If generalization error improves after trimming, replace sub-tree by a leaf node → use a validation set
- Class label of leaf node is determined from majority class of instances in the sub-tree



Conclusion

- Classification problem
 - Learning a model on labeled data
 - Model used to predict class of new examples
- K-nearest neighbor
 - Distance function essential
- Decision trees
 - Hunt's algorithm
 - Split criteria
 - Stopping condition

