Topics in Artificial Intelligence:

Machine Learning Techniques for Data Mining

B Semester 2000

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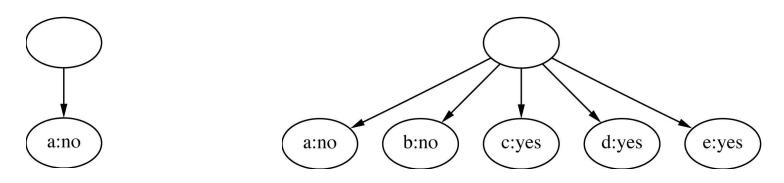
Incremental clustering

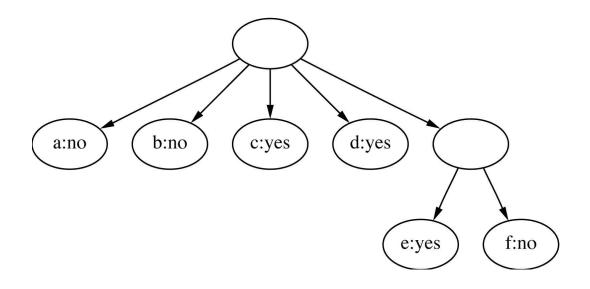
- COBWEB/CLASSIT: incrementally forms a hierarchy of clusters
- In the beginning tree consists of empty root node
- Instances are added one by one, and the tree is updated appropriately at each stage
- Updating involves finding the right leaf for an instance (possibly restructuring the tree)
- Updating decisions are based on category utility

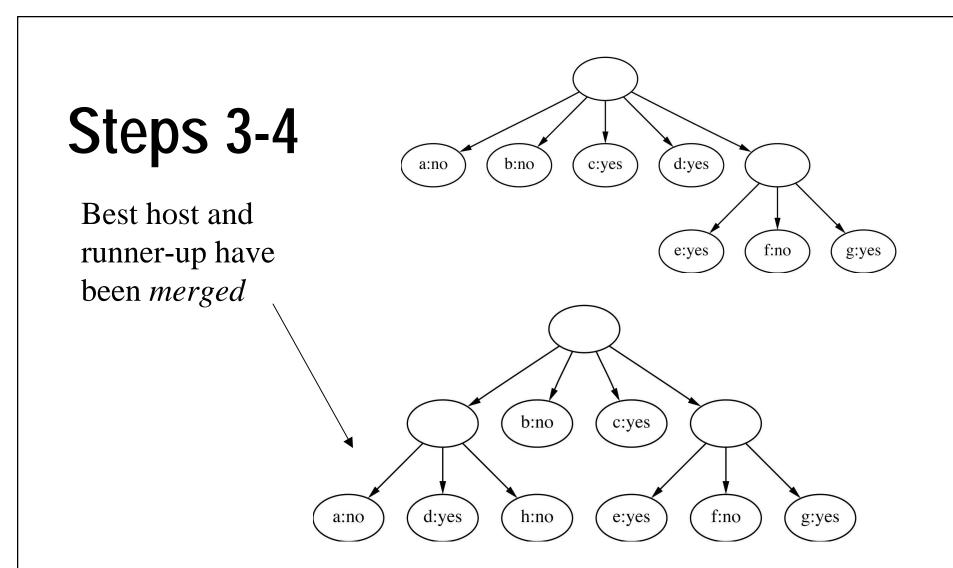
Clustering the weather data

ID code	Outlook	Temp.	Humidity	Windy
А	Sunny	Hot	High	False
В	Sunny	Hot	High	True
С	Overcast	Hot	High	False
D	Rainy	Mild	High	False
Е	Rainy	Cool	Normal	False
F	Rainy	Cool	Normal	True
G	Overcast	Cool	Normal	True
Н	Sunny	Mild	High	False
1	Sunny	Cool	Normal	False
J	Rainy	Mild	Normal	False
K	Sunny	Mild	Normal	True
L	Overcast	Mild	High	True
М	Overcast	Hot	Normal	False
N	Rainy	Mild	High	True

Steps 1-3

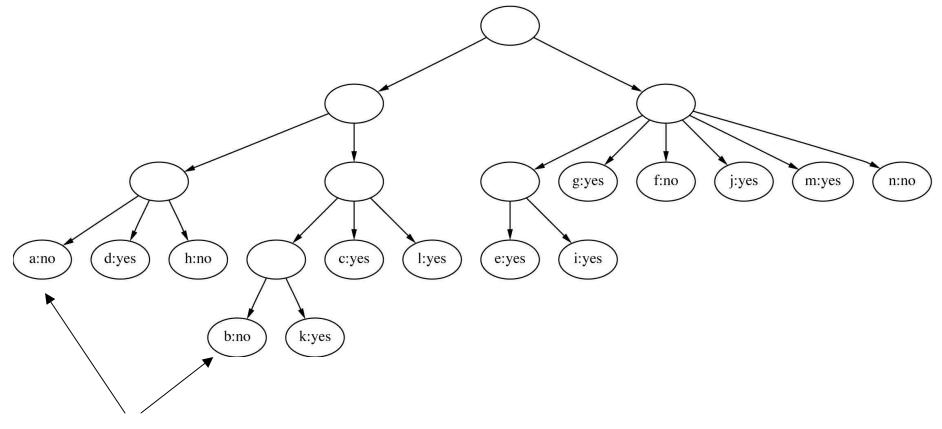






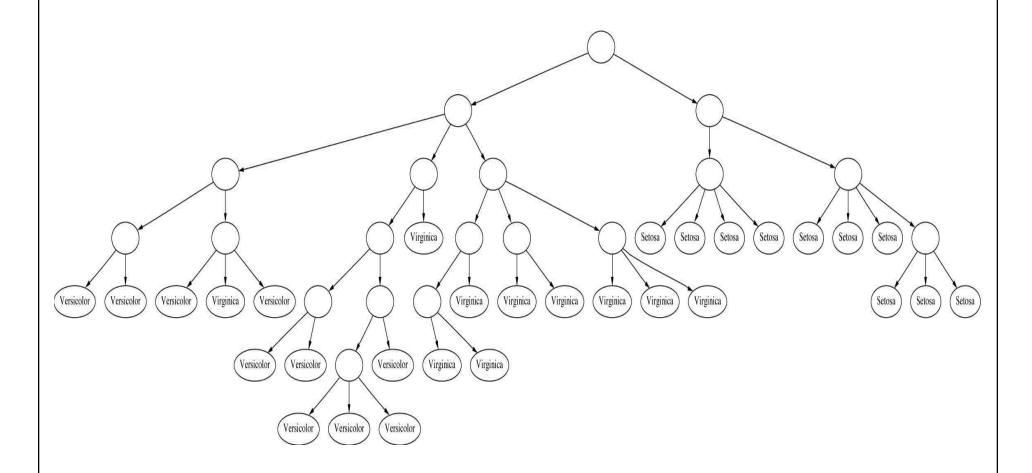
Note: splitting the best host is considered if merging doesn't help

The final hierarchy



a and b are actually very similar

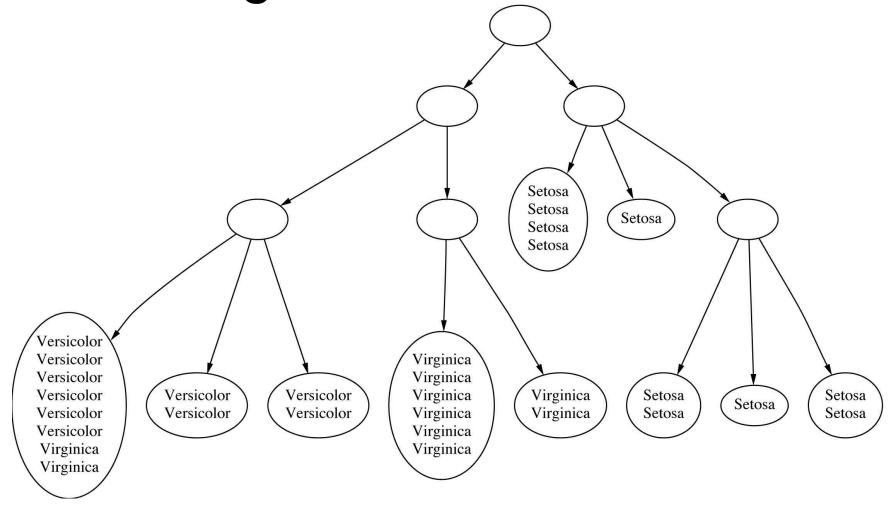
Clustering (parts) of the iris data



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Clustering the iris data with cutoff



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Category utility

Category utility is a kind of quadratic loss function defined on conditional probabilities:

$$CU(C_1, C_2, ..., C_k) = \frac{\sum_{l} \Pr[C_l] \sum_{i} \sum_{j} (\Pr[a_i = v_{ij} \mid C_l]^2 - \Pr[a_i = v_{ij}]^2)}{k}$$

If every instance gets put into a different category the numerator becomes (m = #attributes):

$$m - \Pr[a_i = v_{ij}]^2$$
 — maximum

Numeric attributes

$$f(a) = \frac{1}{\sqrt{2\pi}\sigma} e^{\frac{(a-\mu)^2}{2\sigma^2}}$$

We assume normal distribution: $f(a) = \frac{1}{\sqrt{2\pi}\sigma} e^{\frac{(a-\mu)^2}{2\sigma^2}}$ Then we get: $\sum_{i} \Pr[a_i = v_{ij}]^2 \Leftrightarrow \int f(a_i)^2 da_i = \frac{1}{2\sqrt{\pi}\sigma_i}$

Thus

$$CU = \frac{\sum_{l} \Pr[C_{l}] \sum_{i} \sum_{j} (\Pr[a_{i} = v_{ij} | C_{l}]^{2} - \Pr[a_{i} = v_{ij}]^{2})}{k}$$

is

$$CU = \frac{\sum_{l} \Pr[C_{l}] \frac{1}{2\sqrt{\pi}} \sum_{i} \left(\frac{1}{\sigma_{il}} - \frac{1}{\sigma_{i}} \right)}{k}$$

Acuity parameter: prespecified minimum variance

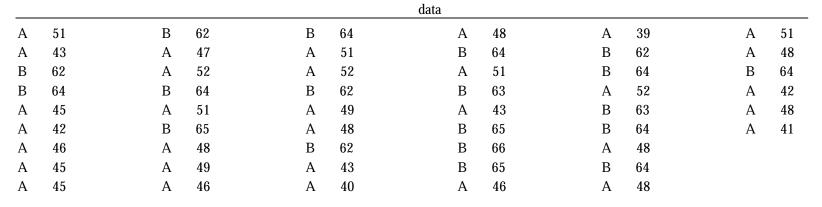
Probability-based clustering

- Problems with above heuristic approach:
 - ◆ Division by k?
 - Order of examples?
 - ◆ Are restructuring operations sufficient?
 - ◆ Is result at least *local* minimum of category utility?
- From a probabilistic perspective, we want to find the most likely clusters given the data
- Also: instance only has certain probability of belonging to a particular cluster

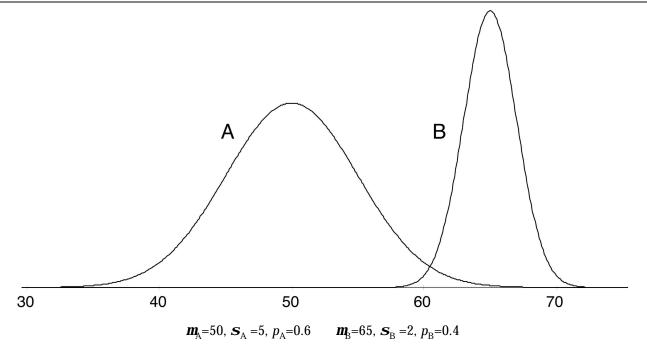
Finite mixtures

- Probabilistic clustering algorithms model the data using a *mixture* of distributions
- Each cluster is represented by one distribution
 - ◆ The distribution governs the probabilities of attributes values in the corresponding cluster
- They are called finite mixtures because there is only a finite number of clusters being represented
- Usually individual distributions are normal distribut.
- Distributions are combined using cluster weights

A two-class mixture model



model



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Using the mixture model

The probability of an instance x belonging to cluster A is:

$$\Pr[A \mid x] = \frac{\Pr[x \mid A] \Pr[A]}{\Pr[x]} = \frac{f(x; \mu_A, \sigma_A) p_A}{\Pr[x]}$$

with

$$f(x;\mu,\sigma) = \frac{1}{\sqrt{2\pi}\sigma} e^{\frac{(x-\mu)^2}{2\sigma^2}}$$

The likelihood of an instance given the clusters is:

$$Pr[x | \text{the distributions}] = \sum_{i} Pr[x | \text{cluster}_{i}] Pr[\text{cluster}_{i}]$$

Learning the clusters

- Assume we know that there are k clusters
- To learn the clusters we need to determine their parameters
 - ◆ I.e. their means and standard deviations
- We actually have a performance criterion: the likelihood of the training data given the clusters
- Fortunately, there exists an algorithm that finds a local maximum of the likelihood

The EM algorithm

- EM algorithm: expectation-maximization algorithm
 - ◆ Generalization of k-means to probabilistic setting
- Similar iterative procedure:
 - Calculate cluster probability for each instance (expectation step)
 - 2. Estimate distribution parameters based on the cluster probabilities (maximization step)
- Cluster probabilities are stored as instance weights

More on EM

Estimating parameters from weighted instances:

$$\mu_A = \frac{w_1 x_1 + w_2 x_2 + \dots + w_n x_n}{w_1 + w_2 + \dots + w_n}$$

$$\sigma_A^2 = \frac{w_1(x_1 - \mu)^2 + w_2(x_2 - \mu)^2 + \dots + w_n(x_n - \mu)^2}{w_1 + w_2 + \dots + w_n}$$

- Procedure stops when log-likelihood saturates
- Log-likelihood:

$$\sum_{i} \log(p_A \Pr[x_i \mid A] + p_B \Pr[x_i \mid B])$$

Extending the mixture model

- Using more then two distributions: easy
- Several attributes: easy if independence is assumed
- Correlated attributes: difficult
 - Modeled jointly using a bivariate normal distribution with a (symmetric) covariance matrix
 - ♦ With n attributes this requires estimating n+n(n+1)/2 parameters
- Nominal attributes: easy if independent

More on extensions

- Correlated nominal attributes: difficult
 - ullet Two correlated attributes result in $v_1 v_2$ parameters
- Missing values: easy
- Distributions other than the normal distribution can be used:
 - ◆ "log-normal" if predetermined minimum is given
 - ◆ "log-odds" if bounded from above and below
 - Poisson for attributes that are integer counts
- Cross-validation can be used to estimate k!!

Bayesian clustering

- Problem: overfitting possible if number of parameters gets large
- Bayesian approach: every parameter has a prior probability distribution
 - Gets incorporated into the overall likelihood figure and thereby penalizes introduction of parameters
- Example: Laplace estimator for nominal attributes
- Can also have prior on number of clusters!
- Actual implementation: NASA's AUTOCLASS

Discussion

- Clusters can be interpreted by using supervised learning in a post-processing step
- Can be used to fill in missing values
- May be advantageous to make attributes more independent in pre-processing step
 - ♦ I.e. using principal component analysis
- Big advantage of probabilistic clustering schemes:
 - Likelihood of data can be estimated and used to compare different clustering models