CS434b/641a: Pattern Recognition Prof. Olga Veksler

Lecture 15

Unsupervised Learning and Clustering

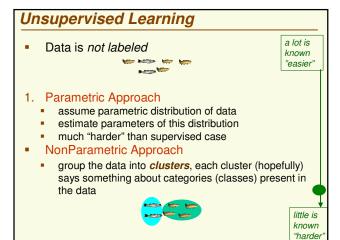
Supervised vs. Unsupervised Learning

- Up to now we considered supervised learning scenario, where we are given
 - 1. samples $x_1, ..., x_n$
 - 2. class labels for all samples $x_1, ..., x_n$
 - This is also called learning with teacher, since correct answer (the true class) is provided
- In the next few lectures we consider unsupervised learning scenario, where we are only given
 - 1. samples $x_1, ..., x_n$
 - This is also called learning without teacher, since correct answer is not provided
 - do not split data into training and test sets

Today

- New Topic: Unsupervised Learning
 - Supervised vs. unsupervised learning
 - Unsupervised learning
 - Next Time: parametric unsupervised learning
 - Today: nonparametric unsupervised learning = clustering
 - Proximity Measures Criterion Functions

 - Flat Clustering
 - k-means
 - Hierarchical Clustering
 - Divisive
 - Agglomerative



Why Unsupervised Learning?

- Unsupervised learning is harder
 - How do we know if results are meaningful? No answer labels are available.
 - Let the expert look at the results (external evaluation)
 - Define an objective function on clustering (internal evaluation)
- We nevertheless need it because
 - 1. Labeling large datasets is very costly (speech recognition)
 - sometimes can label only a few examples by hand
 - May have no idea what/how many classes there are (data mining)
 - 3. May want to use clustering to gain some insight into the structure of the data before designing a classifier
 - Clustering as data description

What we Need for Clustering

- 1. Proximity measure, either
 - similarity measure $s(x_i, x_k)$: large if x_i, x_k are similar
 - dissimilarity(or distance) measure $d(x_i,x_k)$: small if x_i,x_k are similar

large **d**, small **s**

large **s**, small **d**

2. Criterion function to evaluate a clustering





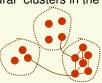
good clustering

bad cluster

Algorithm to compute clustering
 For example, by optimizing the criterion function

Clustering

Seek "natural" clusters in the data



- What is a good clustering?
 - internal (within the cluster) distances should be small
 - external (intra-cluster) should be large
- Clustering is a way to discover new categories (classes)

How Many Clusters?









3 clusters or 2 clusters?

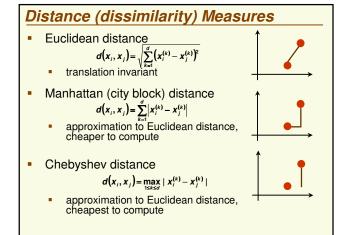
- Possible approaches
 - 1. fix the number of clusters to \boldsymbol{k}
 - 2. find the best clustering according to the criterion function (number of clusters may vary)

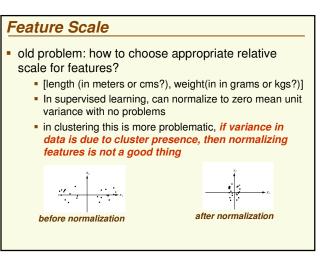
Proximity Measures good proximity measure is VERY application dependent Clusters should be invariant under the transformations "natural" to the problem For example for object recognition, should have invariance to rotation

For character recognition, no invariance to rotation

distance

Similarity Measures Cosine similarity: $s(x_i, x_j) = \frac{x_i^T x_j}{\|x_i\| \|x_j\|}$ • the smaller the angle, the larger the similarity • scale invariant measure • popular in text retrieval Correlation coefficient • popular in image processing $s(x_i, x_j) = \frac{\sum_{k=1}^{d} (x_i^{(k)} - \overline{x}_i)(x_i^{(k)} - \overline{x}_i)}{\left[\sum_{k=1}^{d} (x_i^{(k)} - \overline{x}_i)(x_i^{(k)} - \overline{x}_i)\right]^{1/2}}$

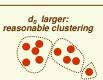




Simplest Clustering Algorithm

- Having defined a proximity function, can develop a simple clustering algorithm
 - go over all sample pairs, and put them in the same cluster if the distance between them is less then some threshold distance d_0 (or if similarity is larger than s_0)
 - Pros: simple to understand and implement
 - Cons: very dependent on d_0 (or s_0), automatic choice of d_0 (or s_0) is not an easily solved issue







SSE Criterion Function

Let n_i be the number of samples in D_i , and define the mean of samples in in D_i

$$\mu_i = \frac{1}{n_i} \sum_{x \in D_i} x$$

Then the sum-of-squared errors criterion function (to minimize) is: $J_{SSE} = \sum_{i=1}^{c} \sum_{\mathbf{x} \in \mathcal{D}} ||\mathbf{x} - \boldsymbol{\mu}_i||^2$

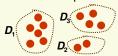




Note that the number of clusters, c, is fixed

Criterion Functions for Clustering

- Have samples $x_1, ..., x_n$
- Suppose partitioned samples into c subsets $D_1,...,D_c$



- There are approximately $c^n/c!$ distinct partitions
- Can define a criterion function $J(D_1,...,D_c)$ which measures the quality of a partitioning D_1, \dots, D_c
- Then the clustering problem is a well defined problem
 - the optimal clustering is the partition which optimizes the criterion function

SSE Criterion Function

$$\boldsymbol{J}_{\mathrm{SSE}} = \sum_{i=1}^{c} \sum_{\boldsymbol{x} \in D_i} \lvert \mid \boldsymbol{x} - \boldsymbol{\mu}_i \mid \rvert^2$$

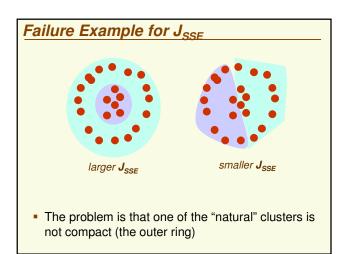
SSE criterion appropriate when data forms compact clouds that are relatively well separated

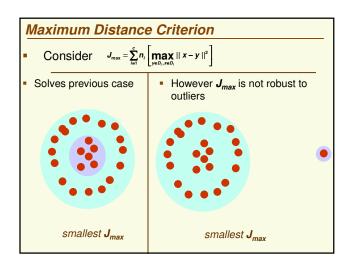


SSE criterion favors equally sized clusters, and may not be appropriate when "natural" groupings have very different sizes









Other Minimum Variance Criterion Functions

• We can eliminate constant terms from

$$J_{SSE} = \sum_{i=1}^{c} \sum_{x \in D_i} ||x - \mu_i||^2$$

We get an equivalent criterion function:

$$J_{E} = \frac{1}{2} \sum_{i=1}^{c} n_{i} \left[\frac{1}{n_{i}^{2}} \sum_{y \in D_{i}} \sum_{x \in D_{i}} || x - y ||^{2} \right]$$

d_i = average Euclidian distance between all pairs of samples in **D**_i

- Can obtain other criterion functions by replacing ||x - y||² by any other measure of distance between points in D_i
- Alternatively can replace d_i by the median, maximum, etc. instead of the average distance

Other Criterion Functions

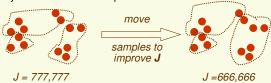
- Recall definition of scatter matrices
 - scatter matrix for ith cluster $S_i = \sum_{x \in D} (x \mu_i)(x \mu_i)^t$
 - within the cluster scatter matrix $S_w = \sum_{i=1}^{c} S_i$
- Determinant of S_w roughly measures the square of the volume
- Assuming S_w is nonsingular, define determinant criterion function:

 $J_d = |S_W| = \left| \sum_{i=1}^c S_i \right|$

 J_a is invariant to scaling of the axis, and is useful if there are unknown irrelevant linear transformations of the data

Iterative Optimization Algorithms

- Now have both proximity measure and criterion function, need algorithm to find the optimal clustering
- Exhaustive search is impossible, since there are approximately cⁿ/c! possible partitions
- Usually some iterative algorithm is used
 - 1. Find a reasonable initial partition
 - 2. Repeat: move samples from one group to another s.t. the objective function **J** is improved



K-means Clustering

 We now consider an example of iterative optimization algorithm for the special case of J_{SSE} objective function

$$J_{SSE} = \sum_{i=1}^{k} \sum_{x=0}^{k} ||x - \mu_i||^2$$

- for a different objective function, we need a different optimization algorithm, of course
- Fix number of clusters to \mathbf{k} ($\mathbf{c} = \mathbf{k}$)
- k-means is probably the most famous clustering algorithm
 - it has a smart way of moving from current partitioning to the next one

Iterative Optimization Algorithms

- Iterative optimization algorithms are similar to gradient descent
 - move in the direction of descent (ascent), but not in the steepest descent direction since have no derivative of the objective function
 - solution depends on the initial point
 - cannot find global minimum
- Main Issue
 - How to move from current partitioning to the one which improves the objective function

K-means Clustering Initialize pick k cluster centers arbitrary assign each example to closest center compute sample means for each cluster reassign all samples to the closest mean

if clusters changed at step 3, go to step 2

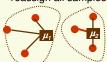
K-means Clustering

- Consider steps 2 and 3 of the algorithm
 - 2. compute sample means for each cluster



 $J_{SSE} = \sum_{i=1}^{k} \sum_{x \in D_i} ||x - \mu_i||^2$ = sum of

3. reassign all samples to the closest mean



If we represent clusters by their old means, the error has gotten smaller



K-means Clustering

- We just proved that by doing steps 2 and 3, the objective function goes down
 - in two step, we found a "smart " move which decreases the objective function
- Thus the algorithm converges after a finite number of iterations of steps 2 and 3
- However the algorithm is not guaranteed to find a global minimum



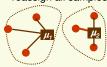
2-means gets stuck here



global minimum of **J**_{SSE}

K-means Clustering

3. reassign all samples to the closest mean



If we represent clusters by their old means, the error has gotten smaller



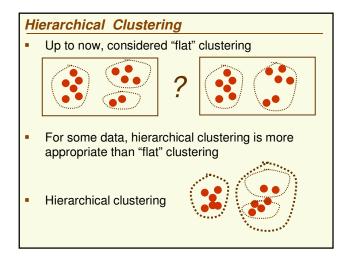
 However we represent clusters by their new means, and mean is always the smallest representation of a cluster

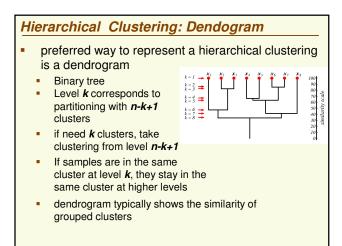
$$\frac{\partial}{\partial z} \sum_{x \in D_{i}} \frac{1}{2} ||x - z||^{2} = \frac{\partial}{\partial z} \sum_{x \in D_{i}} \frac{1}{2} (||x||^{2} - 2x^{t}z + ||z||^{2}) = \sum_{x \in D_{i}} (-x + z) = 0$$

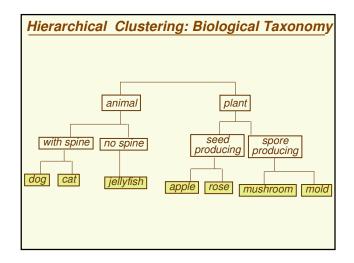
$$\Rightarrow z = \frac{1}{n_{i}} \sum_{x \in D_{i}} x$$

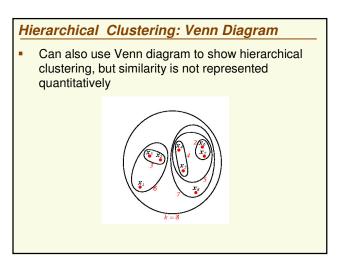
K-means Clustering

- Finding the optimum of J_{SSE} is NP-hard
- In practice, k-means clustering performs usually well
- It is very efficient
- Its solution can be used as a starting point for other clustering algorithms
- Still 100's of papers on variants and improvements of k-means clustering every year









Hierarchical Clustering

- Algorithms for hierarchical clustering can be divided into two types:
 - 1. Agglomerative (bottom up) procedures
 - Start with *n* singleton clusters
 - Form hierarchy by merging most similar clusters



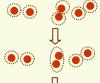
- 2. Divisive (top bottom) procedures
 - Start with all samples in one cluster
 - Form hierarchy by splitting the "worst" clusters

Agglomerative Hierarchical Clustering

initialize with each example in singleton cluster

while there is more than 1 cluster

- 1. find 2 nearest clusters
- 2. merge them

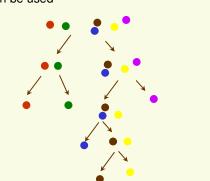


- Four common ways to measure cluster distance
 - 1. minimum distance $d_{\min}(D_i, D_j) = \min_{x \in \mathcal{X}} ||x y||$
 - 2. maximum distance $d_{\text{max}}(D_i, D_j) = \max_{\mathbf{x} \in D_i, \mathbf{y} \in D_j} ||\mathbf{x} \mathbf{y}||$
 - 3. average distance $d_{avg}(D_i, D_j) = \frac{1}{n_i n_j} \sum_{x \in D_i} \sum_{y \in D_i} ||x y||$
 - . mean distance $d_{mean}(D_i, D_j) = || \mu_i \mu_j ||$

Divisive Hierarchical Clustering

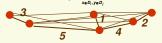
 Any "flat" algorithm which produces a fixed number of clusters can be used

• set **c** = **2**



Single Linkage or Nearest Neighbor

Agglomerative clustering with minimum distance
 d_{min}(ρ, ρ_i)= min || x - y ||



- generates minimum spanning tree
- encourages growth of elongated clusters
- disadvantage: very sensitive to noise

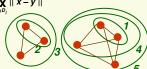
what we want at level with c=3

what we get at level with c=3

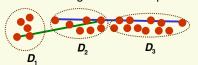
noisy sample

Complete Linkage or Farthest Neighbor

- Agglomerative clustering with maximum distance $d_{\max}(D_i, D_j) = \max_{i \in \mathcal{A}} ||x - y||$
- encourages compact clusters



Does not work well if elongated clusters present



- $d_{\max}(D_1, D_2) < d_{\max}(D_2, D_3)$
- thus D_1 and D_2 are merged instead of D_2 and D_3

Agglomerative vs. Divisive

- Agglomerative is faster to compute, in general
- Divisive may be less "blind" to the global structure of the data

Divisive

when taking the first step (split), have access to all the data; can find the best possible split in 2 parts



Agglomerative

when taking the first step merging, do not consider the global structure of the data, only look at pairwise structure



Average and Mean Agglomerative Clustering

Agglomerative clustering is more robust under the average or the mean cluster distance

$$d_{avg}(D_i, D_j) = \frac{1}{n_i n_j} \sum_{x \in D_i} \sum_{y \in D_j} ||x - y||$$

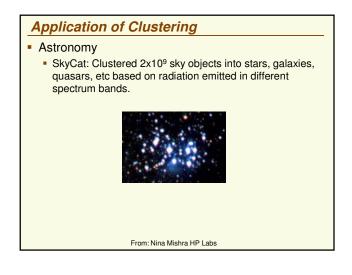
$$d_{mean}(D_i, D_i) = ||\mu_i - \mu_i||$$

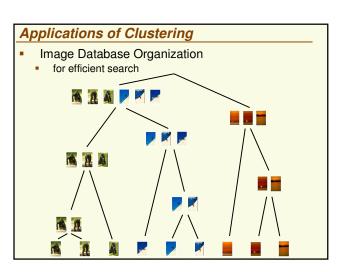
- mean distance is cheaper to compute than the average distance
- unfortunately, there is not much to say about agglomerative clustering theoretically, but it does work reasonably well in practice

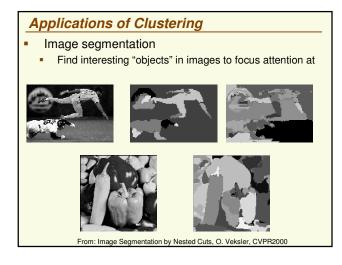
First (?) Application of Clustering

- John Snow, a London physician plotted the location of cholera deaths on a map during an outbreak in the 1850s.
- The locations indicated that cases were clustered around certain intersections where there were polluted wells -- thus exposing both the problem and the solution.









Applications of Clustering Data Mining Technology watch Derwent Database, contains all patents filed in the last 10 years worldwide Searching by keywords leads to thousands of documents Find clusters in the database and find if there are any emerging technologies and what competition is up to Marketing Customer database Find clusters of customers and tailor marketing schemes to them

■ gene expression profile clustering ■ similar expressions, expect similar function U1887 4CL -0.151 -0.207 0.128 0.339 0.208 0.091 -0.083 -0.209 MM4887 +71U8 0.188 0.030 0.111 0.093 -0.093 -0.173 -0.119 -0.138 MM5885 ACC2 0.000 0.011 0.003 -0.003 -0.003 0.000 N86719 ACC1 0.058 0.158 0.082 0.240 0.240 0.058 -0.159 -0.010 U41898 ACT 0.098 -0.198 -0.198 -0.197 0.098 -0.070 AFD87044 ACX1 0.258 0.403 0.679 0.788 0.586 0.249 0.233 0.252 AFD87043 ACX2 0.258 0.403 0.679 0.788 0.586 0.249 0.233 0.252 AFD87043 ACX2 0.188 0.009 -0.31 0.14 0.208 0.242 0.090 0.330 U40856 AIGI 0.098 -0.198 -0.027 -0.029 -0.055 -0.052 0.054 0.008 U40857 AIGZ 0.311 0.149 0.257 0.054 0.188 0.098 -0.498 0.058 AF123253 AIM1 -0.040 0.002 -0.202 -0.040 0.077 0.081 0.008 0.224 N2510 AOS 0.473 0.580 0.514 0.052 0.379 0.387 0.019 0.141 From:De Smet F., Mathys J., Marchal K., Thijs G., De Moor B. & Moreau Y. 2002. Adaptive Quality-based clustering of gene expression profiles, Bioinformatics, 18(6), 735-746.

Summary

- Clustering (nonparametric unsupervised learning) is useful for discovering inherent structure in data
- Clustering is immensely useful in different fields
- Clustering comes naturally to humans (in up to 3 dimensions), but not so to computers
- It is very easy to design a clustering algorithm, but it is very hard to say if it does anything good
- General purpose clustering does not exist, for best results, clustering should be tuned to application at hand

Applications of Clustering

- Profiling Web Users
 - Use web access logs to generate a feature vector for each user
 - Cluster users based on their feature vectors
 - Identify common goals for users
 - Shopping
 - Job Seekers
 - Product Seekers
 - Tutorials Seekers
 - Can use clustering results to improving web content and design

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