Greedy approximation algorithm

- $C = \emptyset$
- Gain of S in the context of  $\mathcal{C}$  is  $f(\mathcal{C} \cup \{S\}) f(\mathcal{C})$
- ullet Select the concept S that has the highest gain
- $\bullet \ \mathcal{C} := \mathcal{C} \cup \{S\}$
- ullet Repeat until  ${\mathcal C}$  has k elements

#### Basic theorem

Let  $\mathcal{C}_k^*$  be the optimal set of k concepts

Let  $C_i$  be the *i*th set formed by the greedy algorithm.

Assume

$$f(\mathcal{C}_i) - f(\mathcal{C}_{i-1}) \ge \frac{1}{k} (f(\mathcal{C}_k^*) - f(\mathcal{C}_{i-1}))$$

Then

$$f(\mathcal{C}_k) \ge \frac{e-1}{e} f(\mathcal{C}_k^*)$$

Proof. Separate.

## Why does the assumption hold?

$$f(\mathcal{C}_i) - f(\mathcal{C}_{i-1}) \ge \frac{1}{k} (f(\mathcal{C}_k^*) - f(\mathcal{C}_{i-1}))$$

f is submodular

the greedy approximation algorithm

The concept  $C_i \setminus C_{i-1}$  is the one that maximizes the gain.

(Something open here.)

# **Applications**

- Which functions are submodular?
- The concepts should not have interaction
- Variable selection?

Algorithmic Methods of Data Mining, Fall 2003, Chapter 11: Clustering2

Chapter 11: Clustering

# Clustering

- Task: group observations into groups so that the observations belonging to the same group are similar, whereas observations in different groups are different
- Lots and lots of research in various areas
- Just scratching the surface here

Basic questions?

- What does "similar" mean?
- What is a good partition of the objects? I.e., how is the quality of a solution measured?
- How to find a good partition of observations?

### What does "similar" mean?

- Some function of the attribute values of the observations
- Usual approach:  $L_p$  distance

$$L((x_1,\ldots,x_n),(y_1,\ldots,y_n))=(\sum_i(x_i-y_i)^p)^{1/p}$$

- Easy in 1-dimensional real case
- Already 2 dimensions cause problems: how to weigh the different dimensions?
- Lots of problems

## Partition-based clustering

- Data mining algorithms: task; model; score function; search
- ullet Task: partition the data into K disjoint sets of points
- The points within each set are as homogeneous as possible
- Measured by score function
- Often no clear model

## Score functions for clustering

- d(x,y): distance between points  $x,y \in D$
- $\bullet$  Assume d is a metric
- $\bullet \ \mathcal{C} = (C_1, C_2, \dots, C_K)$
- Clusters should be compact
- Clusters should be as far from each other as possible
- ullet Within cluster variation  $wc(\mathcal{C})$  and between cluster variation  $bc(\mathcal{C})$

- Cluster centers  $r_1, \ldots, r_K$ : representative points from each cluster, e.g., the centroid of the points
- Simple measure for within cluster variation

$$wc(C) = \sum_{k=1}^{K} wc(C_k) = \sum_{k=1}^{K} \sum_{x \in C_k} d(x, r_k)^2$$

Between cluster variation

$$bc(\mathcal{C}) = \sum_{1 \le j \le k \le K} d(r_j, r_k)^2$$

•  $wc(\mathcal{C})$  leads to spherical clusters

- Evaluation of  $bc(\mathcal{C})$  and  $wc(\mathcal{C})$ ?
- O(n) and  $O(K^2)$  operations
- ullet Variations abound: define  $wc(C_k)$  as the maximum of the minimum distance to another point in the same cluster
- Leads to elongated clusters

### The K-means algorithm

- ullet randomly pick K cluster centers
- assign each point to the cluster whose mean is closest in a Euclidean distance sense
- compute the mean vectors of the points assigned to each cluster
- use these as new centers
- repeat until convergence

#### As an algorithm

```
data points D = \{\mathbf{x}_1, \dots, \mathbf{x}_n\}
find K clusters \{C_1, \ldots, C_K\}:
      for k = 1, ..., K let \mathbf{r}_k be a randomly chosen point from D;
      while changes in clusters C_k happen do
             form clusters:
             for k = 1, \ldots, K do
                    C_k = \{ \mathbf{x} \in D \mid d(\mathbf{r}_k, \mathbf{x}) \leq d(\mathbf{r}_j, \mathbf{x}) \text{ for all } j = 1, \dots, K, j \}
              od;
             compute new cluster centers:
             for k = 1, \dots, K do
                    {\bf r}_k = the vector mean of the points in C_k
              od;
      od;
```

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Properties of the K-means algorithm

- Finds a local optimum
- Converges often quite quickly
- Sometimes slow convergence
- For high dimensions the initial points can have a large influence

## Hierarchical clustering

- Merge sets of points or divide sets of points
- Agglomerative or divisive
- Dendrograms (figures)

## Agglomerative clustering

```
\begin{aligned} &\textbf{for } i=1,\dots,n \text{ let } C_i=\{\mathbf{x}(i)\};\\ &\textbf{while there is more than one cluster left do}\\ &\textbf{let } C_i \text{ and } C_j \text{ be the clusters}\\ &\text{minimizing the distance } \mathcal{D}(C_k,C_h) \text{ between any two clust rs;}\\ &C_i=C_i\cup C_j;\\ &\text{remove cluster } C_j;\\ &\textbf{od;} \end{aligned}
```

# Complexity

- Quadratic, at least, in the number of points
- Not usable for large sets of data

# What is the distance?

- How to define the distance between two clusters?
- Two sets of points
- Lots of alternatives
- Actually quite difficult to define a metric

## Single-link distance

 $d(\mathbf{x}, \mathbf{y})$  the distance between objects  $\mathbf{x}$  and  $\mathbf{y}$ 

$$\mathcal{D}_{sl}(C_i, C_j) = \min_{\mathbf{x}, \mathbf{y}} \{ d(\mathbf{x}, \mathbf{y}) \mid \mathbf{x} \in C_i, \mathbf{y} \in C_j \}, \tag{1}$$

chaining: long, elongated clusters

## Complete link

Furthest distance

$$\mathcal{D}_{fl}(C_i, C_j) = \max_{\mathbf{x}, \mathbf{y}} \{ d(\mathbf{x}, \mathbf{y}) \mid \mathbf{x} \in C_i, \mathbf{y} \in C_j \},$$
 (2)

leads to equal volume (or at least diameter)

#### Other measures

- For vectors
- the centroid measure (the distance between two clusters is the distance between their centroids)
- the group average measure (the distance between two clusters is the average of all the distances between pairs of points, one from each cluster)
- Ward's measure for vector data (the distance between two clusters is the difference between the total within cluster sum of squares for the two clusters separately, and the within cluster sum of squares resulting from merging the two clusters discussed above)

#### Divisive methods

- Start with a single cluster composed of all of the data points
- split this into components
- continue recursively
- Monothetic divisive methods split clusters using one variable at a time
- Polythetic divisive methods make splits on the basis of all of the variables together
- Any intercluster distance measure can be used
- computationally intensive, less widely used than agglomerative methods

Kleinberg's impossibility theorem

- Jon Kleinberg, An impossibility theorem for clustering, NIPS 2002
- Clustering methods based on pairwise distances
- Three properties for clustering methods
- No algorithm can have all three

#### Computational task

- ullet A clustering function operates on a set S of n points
- No underlying space;  $S = \{1, 2, \dots, n\}$
- Distance function:  $d: S \times S \to \mathbf{R}$  with  $d(i,j) \geq 0$ , d(i,j) = d(j,i), and d(i,j) = 0 only if i=j
- (Metric: additionally have  $d(i,j) + d(j,k) \ge d(i,k)$ )
- Clustering function f:  $f(S,d) = \Gamma$ , where  $\Gamma$  is a partition of S
- (A partition)

## Scale invariance

 $\alpha>0$ ; distance function  $\alpha d$  has values  $(\alpha d)(i,j)=\alpha d(i,j)$ 

For any d and for any  $\alpha > 0$  we have  $f(d) = f(\alpha d)$ 

Richness

The range of f is equal to the set of partitions of S

I.e., for any S and any partition  $\Gamma$  of S there is a distance function d on S such that  $f(S,d)=\Gamma$ 

## Consistency

Shrinking distances between points inside a cluster and expanding distances between points in different clusters does not change the result.

 $\Gamma$  a partition of S

d, d' two distance functions on S

d' is a  $\Gamma$ -transformation of d, if

- for all  $i, j \in S$  in the same cluster of  $\Gamma$  we have  $d'(i, j) \leq d(i, j)$
- $\bullet$  for all  $i,j\in S$  in the different clusters of  $\Gamma$  we have  $d'(i,j)\geq d(i,j)$

Consistency: if  $f(S,d) = \Gamma$  and d' is a  $\Gamma$ -transformation of d, then  $f(S,d') = \Gamma$ 

# Examples

- Agglomerative clustering with single-link
- Repeatedly merge cluster whose distance is minimum
- Continue until a stopping criterion is met
  - k-cluster stopping criterion: continue until there are k connected components
  - distance-r stopping criterion: continue until all distances between clusters are larger than r
  - scale- $\alpha$  stopping criterion: let  $\rho^*$  be the maximum pairwise distance; contine until all distances are larger than  $\alpha \rho^*$

Examples, cont.

- Single link with k-cluster stopping criterion satisfies scale-invariance and consistency
- ullet Single link with distance-r stopping criterion satisfies richness and consistency
- ullet Single link with scale-lpha stopping criterion satisfies richness and scale-invariance

Theorem

For each  $n \geq 2$  there is no clustering function that satisfies scale-invariance, richness, and consistency

#### Proof of theorem

A partition  $\Gamma'$  is a refinement of partition  $\Gamma$ , if each set  $C' \in \Gamma'$  is included in some set  $C \in \Gamma$ 

A partial order between partitions:  $\Gamma' \leq \Gamma$ 

Antichain of partitions: collection of partitions such than no one is a refinement of others

Theorem: If a clustering function f satisfies scale-invariance and concistency, then the range of f is an antichain

# $\Gamma$ -forcing

- ullet partition  $\Gamma$
- d (a,b)-conforms to  $\Gamma$ , if for all points i,j in the same cluster of  $\Gamma$   $d(i,j) \leq a$ , and for all points i,j in different clusters of  $\Gamma$   $d(i,j) \geq b$
- ullet given a clustering function f
- (a,b) is  $\Gamma$ -forcing, if for all d that (a,b)-conform to  $\Gamma$  we have  $f(S,d)=\Gamma$

#### Forcing, cont.

- Assume f satisfies consistency; let  $\Gamma \in Range(f)$
- Claim: there are a < b such that (a,b) is  $\Gamma$ -forcing
- ullet  $\Gamma$  in range of f: there is d such that  $f(S,d)=\Gamma$
- a'= minimum distance of points in the same cluster in  $\Gamma$
- b'= maximum distance of points in different clusters in  $\Gamma$
- Choose a < b such that a < a' and b < b'
- If d'(a,b)-conforms to  $\Gamma$ , then d is a  $\Gamma$ -transformation of d
- By consistency  $f(d') = \Gamma$
- Thus (a,b) is  $\Gamma$ -forcing

#### Antichains

- Assume f satisfies scale-invariance
- Let  $\Gamma_0$  and  $\Gamma_1$  be possible results of f, and let  $\Gamma_0$  be a refinement of  $\Gamma_1$
- Show that this leads to contradiction
- $(a_0,b_0)$   $\Gamma_0$ -forcing,  $(a_1,b_1)$   $\Gamma_1$ -forcing
- Let  $a_2 < a_1$ , choose  $\epsilon$  so that  $0 < \epsilon < a_0 a_2 b_0^{-1}$
- Construct a d such that
  - For i, j in same cluster of  $\Gamma_0$  we have  $d(i, j) \leq \epsilon$
  - For i,j in same cluster of  $\Gamma_1$  but not in  $\Gamma_0$  we have  $a_2 \leq d(i,j) \leq a_1$
  - For i, j in different clusters of  $\Gamma_1$   $d(i, j) \geq b_1$

- d (a,b)-conforms to  $\Gamma_1$ , and thus  $f(S,d)=\Gamma_1$
- $\alpha = b_0 a_2^{-1}$ , and let  $d' = \alpha d$
- scale-invariance:  $f(d') = f(d) = \Gamma_1$
- i, j in same cluster of  $\Gamma_0$  we have

$$d'(i,j) \le \epsilon b_0 a_2^{-1} < a_0$$

• i, j in different clusters of  $\Gamma_0$  we have

$$d'(i,j) \ge a_2 b_0 a_2^{-1} = b_0$$

• d'  $(a_0,b_0$  conforms to  $\Gamma_0$ , and thus  $f(S,d')=\Gamma_0 \neq \Gamma_1$ , contradiction