

Unsupervised learning

Clustering

Statistical methods for data analysis – Machine learning

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Motivation (1/2)

- **Supervised methods** (learning with a teacher):
 - **Input:** predictor variables $X^T = (X_1, \dots, X_p)$
 - **Output:** Y
 - Predictions based on the **training set** $(x_1, y_1), \dots, (x_N, y_N)$ of previously solved cases
 - **Loss function** $L(y, \hat{y})$, such as $L(y, \hat{y}) = (y - \hat{y})^2$
 - Supposing (X, Y) random variables supervised learning is a **density estimation problem**:

Determining the properties of the conditional density $\Pr(Y|X)$

E.g.: location parameters that minimize the expected error

$$\mu(x) = \operatorname{argmin}_{\theta} E_{Y|X} L(Y, \theta).$$

Motivation (2/2)

- **Unsupervised methods** (learning **without** a teacher):
 - **Input:** predictor variables $X^T = (X_1, \dots, X_p)$
 - **Output:** not available
 - **Goal:** infer the **properties of the joint probability** $\Pr(X)$ without the help of a supervisor/teacher
 - In **low dimensional problems** ($p \leq 3$) $\Pr(X)$ can be directly estimated and graphically represented
 - In **high dimensional problems** descriptive statistics methods are used to characterize $\Pr(X)$
 - **Low dimensional manifolds** representing high data density may be identified by **PCA** or other **dimensionality reduction** methods
 - **Cluster analysis** attempts to find multiple convex regions of the X -space that contain **modes** of $\Pr(X)$
 - No direct measure of success (as loss function)

Unsupervised learning methods

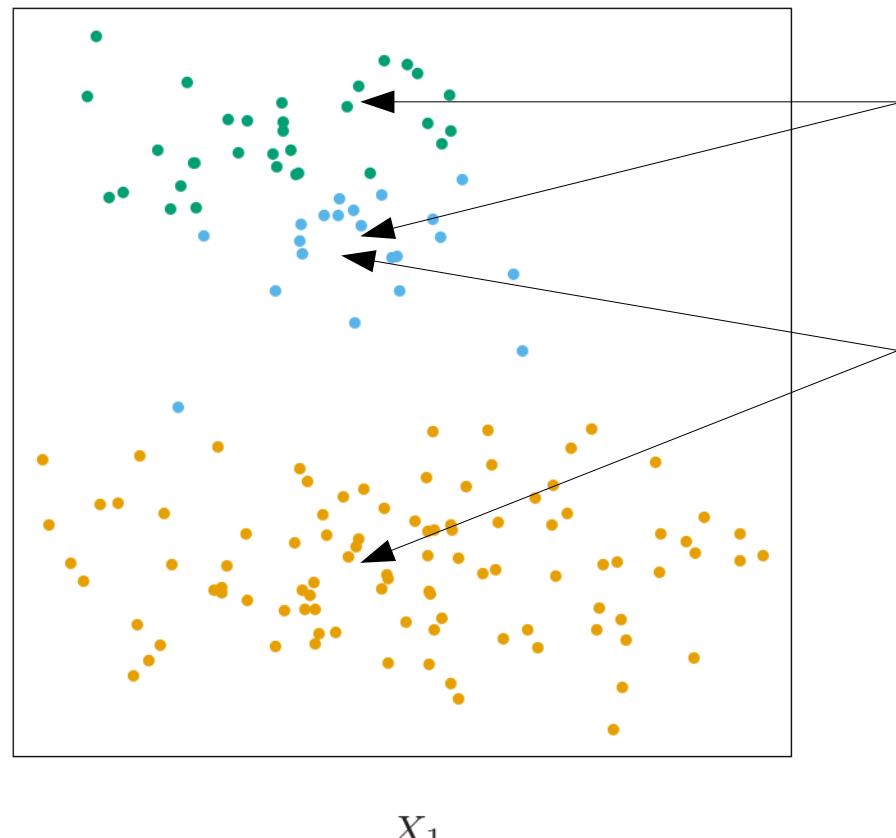
- Association rules
- Clustering analysis
 - K-means
 - K-medoids
 - Gaussian Mixture Models
 - Hierarchical clustering
- Self-organizing maps
- Principal components, curves and subspaces
 - Spectral clustering
- Matrix factorization
- ...

Unsupervised learning methods

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Cluster analysis

Grouping a collection of **objects** into **subsets (clusters)** such that objects **within** each clusters are **more closely related** to one another **than** objects assigned to **different clusters**



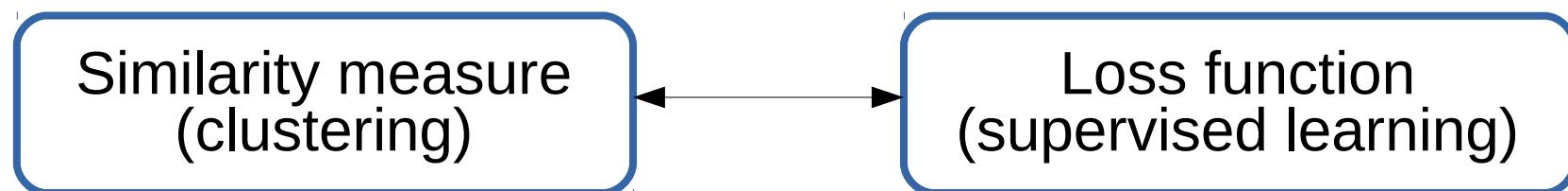
Not well separated
(similar properties)

Well separated
(different properties)

Why?

Measures of similarity/dissimilarity

- Central to clustering analysis is the notion of **similarity/dissimilarity between individual objects**
- **Clustering methods** attempt to **group** the objects according to the definition of **similarity** supplied to it.



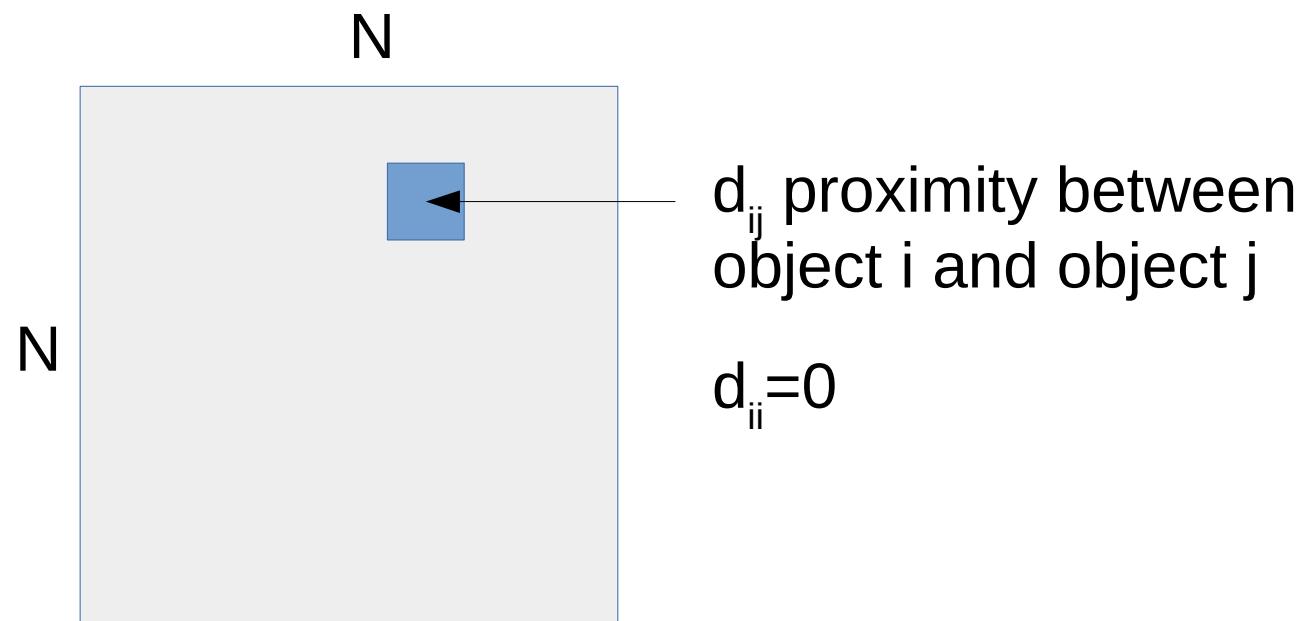
- **Examples of similarity/dissimilarity measures:**

- Euclidean distance
- Manhattan distance
- Mahalanobis distance
- Correlation
- Jaccard distance

$$d(\mathbf{p}, \mathbf{q}) = \sqrt{\sum_{i=1}^n (q_i - p_i)^2}$$
$$d_1(\mathbf{p}, \mathbf{q}) = \|\mathbf{p} - \mathbf{q}\|_1 = \sum_{i=1}^n |p_i - q_i|$$
$$D_M(\vec{x}) = \sqrt{(\vec{x} - \vec{\mu})^T S^{-1} (\vec{x} - \vec{\mu})}$$
$$\rho_{X,Y} = \frac{E[(X - \mu_X)(Y - \mu_Y)]}{\sigma_X \sigma_Y}$$
$$J(A, B) = \frac{|A \cap B|}{|A \cup B|} = \frac{|A \cap B|}{|A| + |B| - |A \cap B|}$$

Proximity matrices

- Sometimes the data is represented directly in terms of proximity between pairs of objects (similarities or dissimilarities).
- $N \times N$ matrix



- Dissimilarities are *distances* in the strict sense only if the triangle inequality $d_{ii'} \leq d_{ik} + d_{i'k}$, for all $k \in \{1, \dots, N\}$ holds.

Dissimilarities based on attributes (1/2)

- Usually we have measurements x_{ij} $i=1, \dots, N$, on variables $j=1, \dots, p$ (**attributes**).
- Then **pairwise dissimilarities** between **observations** can be expressed in terms of **attribute** values, that is

$$D(x_i, x_{i'}) = \sum_{j=1}^p d_j(x_{ij}, x_{i'j})$$

where $d_j(x_{ij}, x_{i'j})$ is the dissimilarity between values of the j -th attribute.

- Most common **D** function is the **squared distance**

$$d_j(x_{ij}, x_{i'j}) = (x_{ij} - x_{i'j})^2$$

Dissimilarities based on attributes (2/2)

- Other choices are possible depending on attribute types

- Quantitative variables

- Absolute difference $d(x_i, x_{i'}) = l(|x_i - x_{i'}|)$

- Correlation $\rho(x_i, x_{i'}) = \frac{\sum_j (x_{ij} - \bar{x}_i)(x_{i'j} - \bar{x}_{i'})}{\sqrt{\sum_j (x_{ij} - \bar{x}_i)^2 \sum_j (x_{i'j} - \bar{x}_{i'})^2}}$

- ...

- Ordinal variables

- Categorical variables

Object dissimilarity (1/2)

- Dissimilarities of p attributes are then combined into a single overall measure of dissimilarity $D(x_i, x_{i'})$ between objects
- Weighted average (convex combination):

$$D(x_i, x_{i'}) = \sum_{j=1}^p w_j \cdot d_j(x_{ij}, x_{i'j}); \quad \sum_{j=1}^p w_j = 1$$

Weight of j-th attribute

- Weight w_j regulates the relative influence of variable j in determining the overall dissimilarity between objects
- All $w_j=1$ does NOT give all attributes equal influence
- The relative influence of the j -th variable is $w_j * \text{avg}(d_j)$

where

$$\bar{d}_j = \frac{1}{N^2} \sum_{i=1}^N \sum_{i'=1}^N d_j(x_{ij}, x_{i'j})$$

Object dissimilarity (2/2)

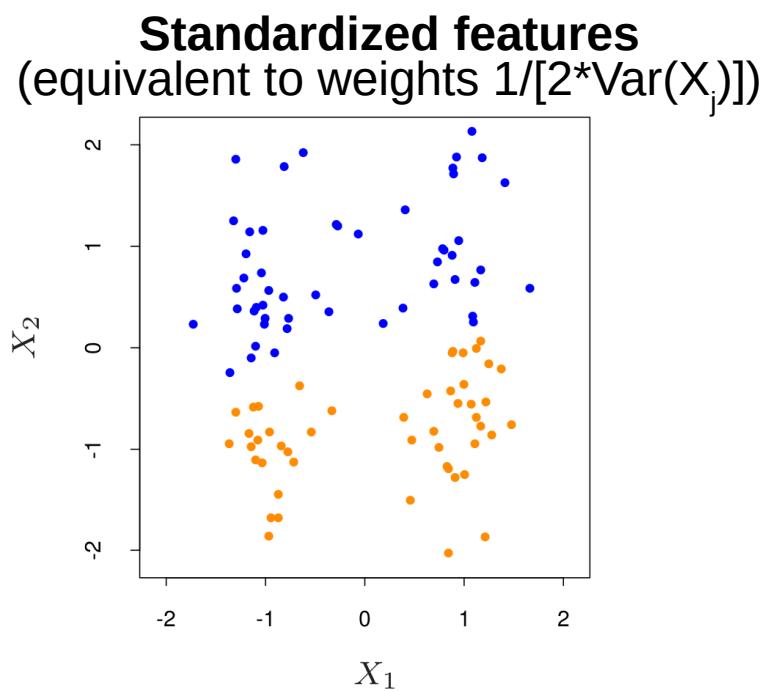
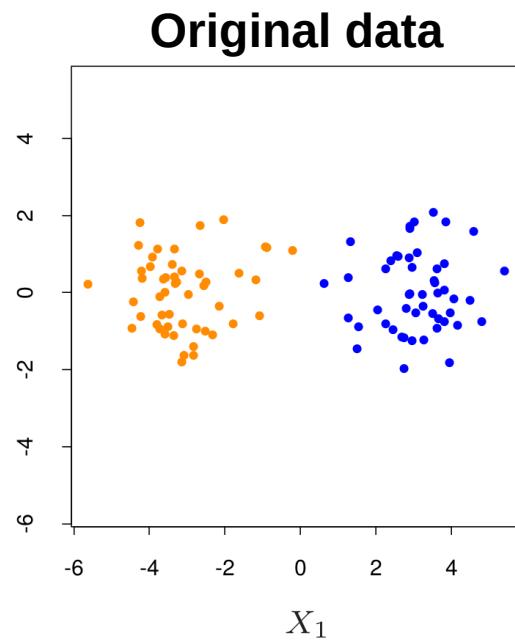
- Hence, setting $w_j \sim 1/\text{avg}(d_j)$ gives all attributes **equal influence** on the overall dissimilarity
- This is related to **data standardization** in supervised learning
- E.g., for squared error distance

$$D_I(x_i, x_{i'}) = \sum_{j=1}^p w_j \cdot (x_{ij} - x_{i'j})^2$$
$$\bar{d}_j = \frac{1}{N^2} \sum_{i=1}^N \sum_{i'=1}^N (x_{ij} - x_{i'j})^2 = 2 \cdot \text{var}_j$$

the **relative importance** of each attribute is **proportional** to its **variance** over the data

Attribute relative importance

- If the goal is **discovering natural grouping**, forcing equal influence among attributes can be **counterproductive**.
- **More relevant variables should have higher influence** in the object dissimilarity
- Giving all attributes equal influence tend to obscure the clusters to clustering algorithms



The choice of **appropriate dissimilarity measures** is often more important than the choice of the clustering algorithm

Goal of clustering algorithms: to partition observations into groups such that pairwise dissimilarities between observations assigned to the same cluster tend to be smaller than those in different clusters

Algorithm types:

- Combinatorial
- Mixture modeling based
- Mode seekers

Combinatorial algorithms

- Most popular
- No probability model
- Prespecified number of clusters $K < N$
- Each observation labeled by an integer k in $\{1, \dots, K\}$
- Assignments characterized by a many-to-one mapping (**encoder**):

$$k = C(i)$$

that assigns the i -th observation to the k -th cluster

Goal: seek the encoder $C^*(i)$ that minimizes a loss (or energy) function which depends on pairwise dissimilarities

$$W(C) = \frac{1}{2} \sum_{k=1}^K \sum_{C(i)=k} \sum_{C(i')=k} d(x_i, x_{i'})$$

Within cluster point scatter

- **Total point scatter**

$$T = \frac{1}{2} \sum_{i=1}^N \sum_{i'=1}^N d_{ii'} = \frac{1}{2} \sum_{k=1}^K \sum_{C(i)=k} \left(\sum_{C(i')=k} d_{ii'} + \sum_{C(i') \neq k} d_{ii'} \right)$$

- **Between-cluster point scatter**

$$B(C) = \frac{1}{2} \sum_{k=1}^K \sum_{C(i)=k} \sum_{C(i') \neq k} d_{ii'}$$

- **Relationships**

$$T = W(C) + B(C)$$

Minimizing $W(C)$ is equivalent to maximizing $B(C)$

- Number of possible assignments (*Jain ad Dubes, 1988*)

$$S(N, K) = \frac{1}{K!} \sum_{k=1}^K (-1)^{K-k} \binom{K}{k} k^N$$

- E.g., $S(10,4)=34105$, $S(19,4)=10^{10}$
- Heuristic strategies: iterative greedy descent
- Initial partition
- Iterative steps for reducing the loss
- Local optima

- **Squared error distance**

$$d(x_i, x_{i'}) = \sum_{j=1}^p (x_{ij} - x_{i'j})^2 = \|x_i - x_{i'}\|^2$$

- **Within point scatter**

$$\begin{aligned} W(C) &= \frac{1}{2} \sum_{k=1}^K \sum_{C(i)=k} \sum_{C(i')=k} \|x_i - x_{i'}\|^2 \\ &= \sum_{k=1}^K N_k \sum_{C(i)=k} \|x_i - \bar{x}_k\|^2, \end{aligned}$$

↑

Euclidean distance from the centroid
(mean vector) associated to the k-th cluster

K-means algorithm

1) Given a cluster assignment C , the total cluster variance

$$\min_{C, \{m_k\}_1^K} \sum_{k=1}^K N_k \sum_{C(i)=k} \|x_i - m_k\|^2$$

is minimized with respect to $\{m_1, \dots, m_K\}$ yielding the means of the currently assigned clusters.

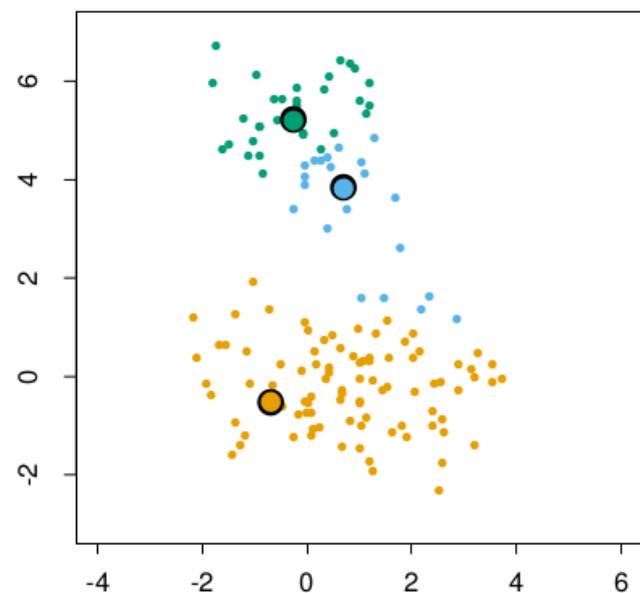
2) Given the current set of means $\{m_1, \dots, m_K\}$ the total cluster variance is minimized by assigning each observation to the closest (current) cluster mean:

$$C(i) = \operatorname{argmin}_{1 \leq k \leq K} \|x_i - m_k\|^2$$

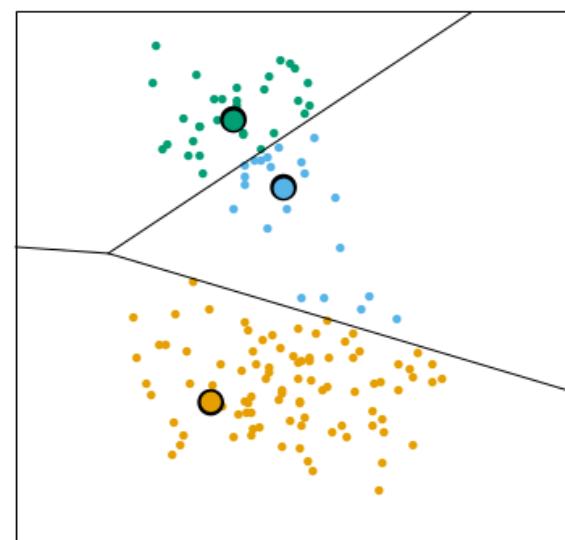
3) Iterate steps 1 and 2 until the assignments do not change.

Successive iterations of K-means

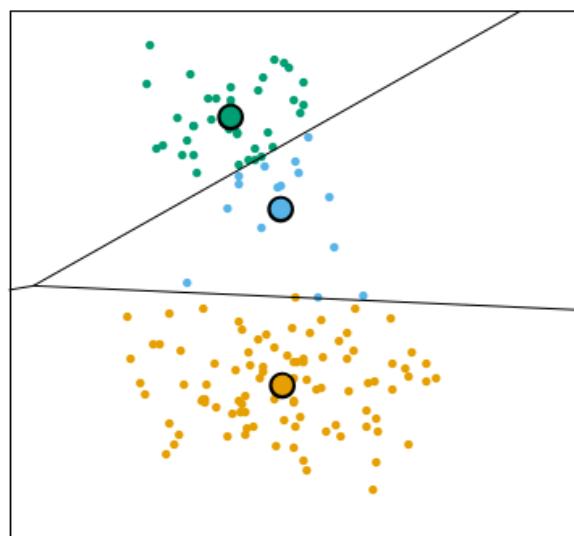
Initial Centroids



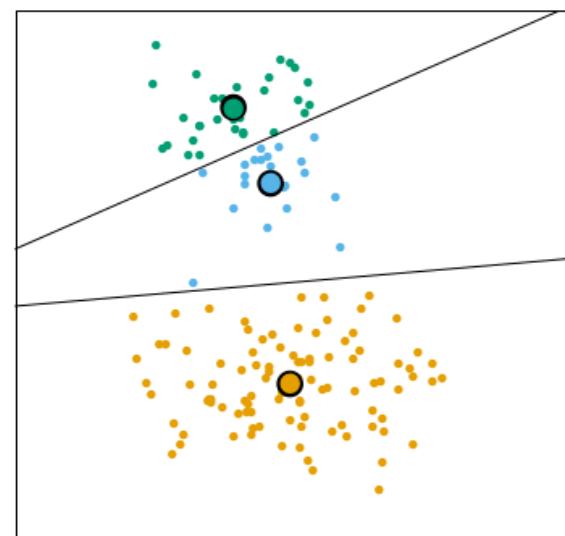
Initial Partition



Iteration Number 2



Iteration Number 20



Exercise: Clustering of Human tumor microarray data

See text of Exercise 6

Human tumor dataset



Experiments (samples)

Genes

References

[Hastie 2009] Trevor Hastie, Robert Tibshirani, Jerome Friedman. The Elements of Statistical Learning: Data Mining, Inference, and Prediction (second edition). Springer. 2009.