14.3 – Cluster analysis in the book "The elements of statistical learning"

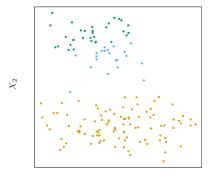
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Clustering



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FIGURE 14.4. Simulated data in the plane, clustered into three classes (represented by orange, blue and green) by the K-means clustering algorithm

Outline



2 Clustering algorithms

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14.1 - Proximity matrices

Data: $\{x_i\}_{i=1}^N$ Proximity matrix: $\mathbf{D} \in \mathbb{R}^{N \times N}$, $d_{ii} = 0$ Symmetrization: If \mathbf{D} is not symmetric, we can take $(\mathbf{D} + \mathbf{D}^T)/2$ Dissimilarities based on attributes:

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

where x_{ij} is the *j*th attribute of the *i*th data.

Dissimilarities

Euclidean distance

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

More general

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

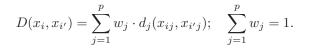
where $l(\cdot)$ is a monotone-increasing function.

Based on correlation

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

where $\bar{x}_i = \sum_j x_{ij}/p$

Weighted average



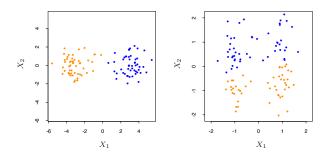


FIGURE 14.5. Simulated data: on the left, K-means clustering (with K=2) has been applied to the raw data. The two colors indicate the cluster memberships. On the right, the features were first standardized before clustering. This is equivalent to using feature weights $1/[2 \cdot var(X_j)]$. The standardization has obscured the two well-separated groups. Note that each plot uses the same units in the horizontal and vertical axes.

Outline





Clustering algorithms

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- *N* points: $x_i, i \in \{1, ..., N\}$
- K clusters $k \in \{1, \ldots, K\}$

• Encoder k = C(i), that assigns the *i*th point to the *k*th cluster **Energy functions:**

$$W(C) = \frac{1}{2} \sum_{k=1}^{K} \sum_{C(i)=k} \sum_{C(i')=k} d(x_i, x_{i'}).$$
$$B(C) = \frac{1}{2} \sum_{k=1}^{K} \sum_{C(i)=k} \sum_{C(i')\neq k} d_{ii'}$$
$$= \frac{1}{2} \sum_{i=1}^{N} \sum_{i'=1}^{N} d_{ii'} = \frac{1}{2} \sum_{k=1}^{K} \sum_{C(i)=k} \left(\sum_{C(i')=k} d_{ii'} + \sum_{C(i')\neq k} d_{ii'} \right)$$
$$T = W(C) + B(C),$$

W is the in-cluster point scatters, and W is the between-cluster point scatters. T is the total point scatters, which is a constant given the data, independent of cluster assignment.

Clustering algorithms

Clustering: maximize W(C) or minimize B(C)

Problem: solving this optimization problem directly is combinatorial. The number of distinct assignments is

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

S(10,4) = 34,105 and $S(19,4) \approx 10^{10}$

K-means

Euclidean distance:

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

Within-point scatter:

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

where $\bar{x}_k = (\bar{x}_{1k}, \dots, \bar{x}_{pk})$ is the mean vector associated with the *k*th cluster and $N_k = \sum_{i=1}^N I(C(i) = k)$.

Iterative descent algorithm

Minimization:

$$C^* = \min_{C} \sum_{k=1}^{K} N_k \sum_{C(i)=k} ||x_i - \bar{x}_k||^2$$

where for any subset S

$$\bar{x}_S = \underset{m}{\operatorname{argmin}} \sum_{i \in S} ||x_i - m||^2.$$

Enlarged minimization:

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

K-means algorithm

Algorithm 14.1 K-means Clustering.

- 1. For a given cluster assignment C, the total cluster variance (14.33) is minimized with respect to $\{m_1, \ldots, m_K\}$ yielding the means of the currently assigned clusters (14.32).
- 2. Given a current set of means $\{m_1, \ldots, m_K\}$, (14.33) is minimized by assigning each observation to the closest (current) cluster mean. That is,

$$C(i) = \underset{1 \le k \le K}{\operatorname{argmin}} ||x_i - m_k||^2.$$
(14.34)

3. Steps 1 and 2 are iterated until the assignments do not change.

Algorithm iteration

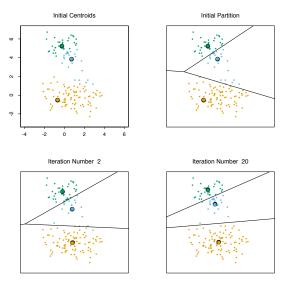


FIGURE 14.6. Successive iterations of the K-means clustering algorithm for $\mathbb{P} \to \mathbb{P} = \mathbb{P} =$

K-means on human gene data

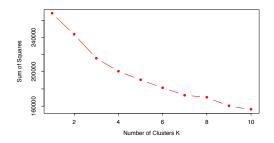


FIGURE 14.8. Total within-cluster sum of squares for K-means clustering applied to the human tumor microarray data.

TABLE 14.2. Human tumor data: number of cancer cases of each type, in each of the three clusters from K-means clustering.

Cluster	Breast	CNS	Colon	K562	Leukemia	MCF7
1	3	5	0	0	0	0
2	2	0	0	2	6	2
3	2	0	7	0	0	0
Cluster	Melanoma	NSCLC	Ovarian	Prostate	Renal	Unknown
1	1	7	6	2	9	1
2	7	2	0	0	0	0

Reference

Chapter 14: Trevor Hastie, Robert Tibshirani, The Elements of Statistical Learning, Second Edition.