NDA: Clustering

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Bibliography

 Introduction to Data Mining, 2nd Edition by Tan, Steinbach, Karpatne, Kumar Chapter 7.

Outline

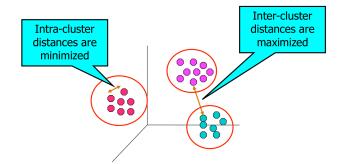
- What is clustering? Motivation?
- 2 k-means
- Other clustering algorithms
 - Mean-shift
 - GMM: Gaussian Mixture Model
 - Hierarchical clustering
- Validation

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- What is clustering? Motivation?
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What is clustering?

Finding groups of objects such that the objects in a group will be similar (or related) to one another and different from (or unrelated to) the objects in other groups



What is clustering?

- Clustering is an unsupervised learning method (i.e. no predefined classes)
- It is different from classification (supervised learning)

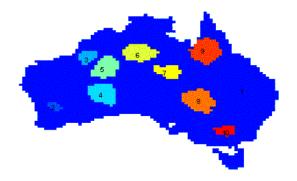
Why is it usefull (applications of cluster analysis)?

Understanding the data / get insights on the data:

- Group related documents for browsing
- Group genes and proteins that have similar functionality
- Group people sharing similar interest
- Group movies with similar genres or actors

Why is it usefull (applications of cluster analysis)?

Summarization: Reduce the size of large data sets



Clustering precipitation in Australia

Custering is an ill-defined problem

How many clusters?

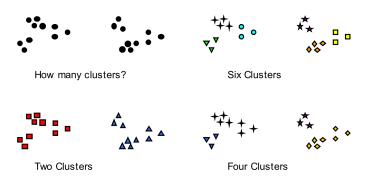


How many clusters?

► An Impossibility Theorem for Clustering, Jon Kleinberg, NIPS2003.

Custering is an ill-defined problem

How many clusters?



► An Impossibility Theorem for Clustering, Jon Kleinberg, NIPS2003.

Several types of clusters

- A well-separated cluster is a set of points such that any point in a cluster is closer to every other point in the cluster than to any point not in the cluster.
- A center-based cluster is a set of objects such that an object in a cluster is closer to the "center" of a cluster, than to the center of any other cluster
- A contiguous cluster is a set of points such that a point in a cluster is closer to one or more other points in the cluster than to any point not in the cluster.
- A density-based cluster is a dense region of points, which is separated by low-density regions, from other regions of high density.

Several types of clusterings

A clustering is a set of clusters. Important distinction between hierarchical and partitional sets of clusters.

- Partitional clustering: A division of data objects into non-overlapping subsets (clusters) such that each data object is in exactly one subset
- Hierarchical clustering: A set of nested clusters organized as a hierarchical tree

Several types of clusterings

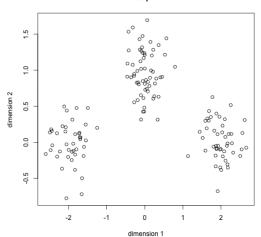
Other distinctions:

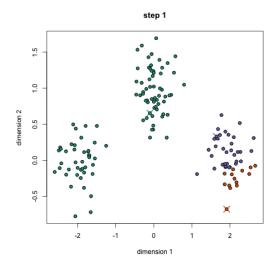
- Exclusive versus non-exclusive: In non-exclusive clusterings, points may belong to multiple clusters. Can represent multiple classes or 'border' points
- Fuzzy versus non-fuzzy: In fuzzy clustering, a point belongs to every cluster with some weight between 0 and 1. Weights must sum to 1
- Partial versus complete: In some cases, we only want to cluster some of the data
- Heterogeneous versus homogeneous: Clusters of widely different sizes, shapes, and densities

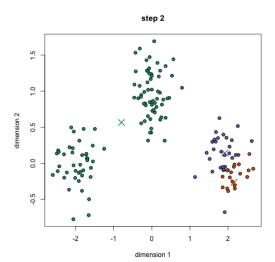
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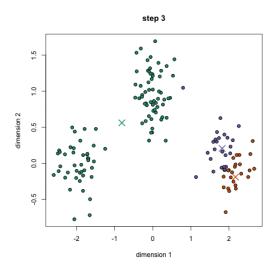
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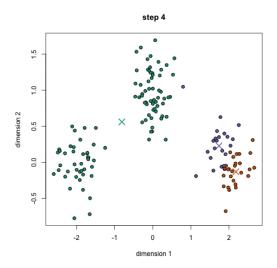


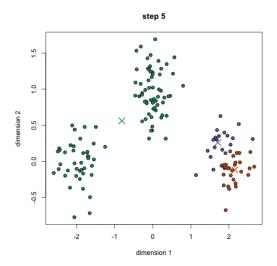


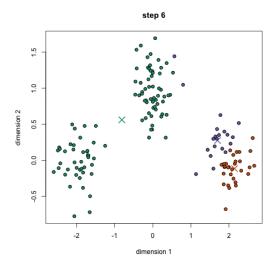


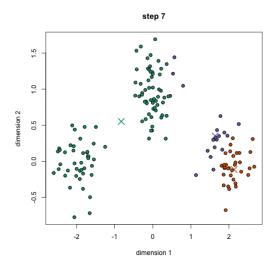


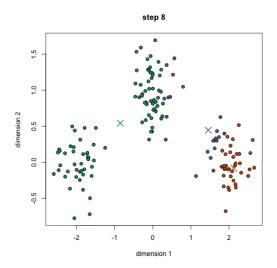


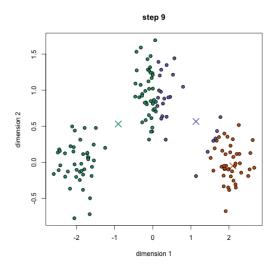


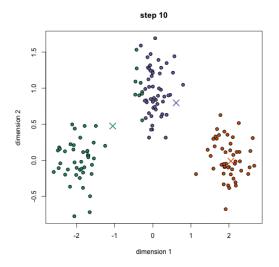




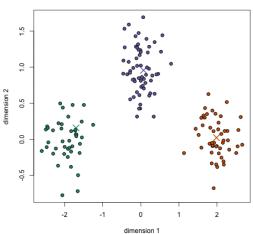


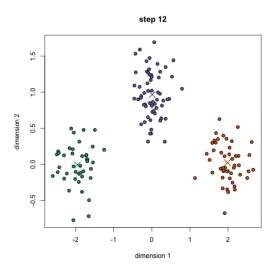












- Randomly chose k initial centroids
- While True:
 - Create k clusters by assigning each point to closest centroid
 - Compute k new centroids by averaging points in each clustering
 - If centroids don't change:
 - Break

k-means: distortion (a.k.a. SSE or SSD)

k-means can be seen as a heuristic to minimize the distortion:

distortion =
$$\sum_{i=1}^{n} \sum_{j=1}^{k} w_{i,j} ||x_i - \mu_j||_2^2$$

with

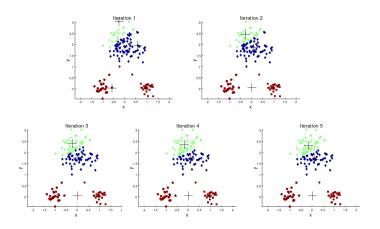
- μ_j the vector of centroid j and
- $w_{i,j} = 1$ if the sample x_i is in cluster j and 0 otherwise.

Limitation of k-means: clusters

- K-means has problems when clusters are of differing
 - Sizes
 - Densities
 - Non-globular shapes
- K-means has problems when the data contains outliers.

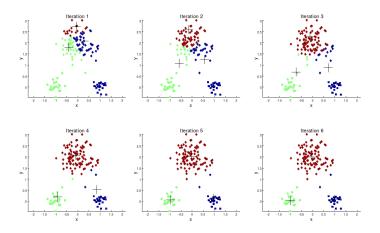
Normalising the data and removing outliers can help!

Limitation of k-means: initialisation



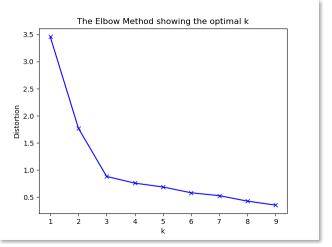
Using multiple runs or kmeans++ can help!

Limitation of k-means: initialisation



Using multiple runs or kmeans++ can help!

Limitation of k-means: How to choose k?



The elbow method: choose k=3, where the elbow is located

Limitation of k-means: How to choose k?

Silhouette value. A measure of how similar an object is to its own cluster (cohesion) compared to other clusters (separation).

For data point i in cluster C_k , let

$$a(i) = \frac{1}{|C_k| - 1} \sum_{j \in C_k, i \neq j} d(i, j)$$
 and $b(i) = \min_{l \neq k} \frac{1}{|C_l|} \sum_{j \in C_l} d(i, j)$

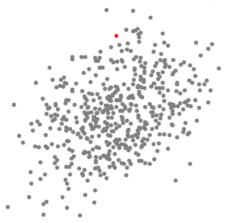
The silhouette score of one data point i: $s(i) = \frac{b(i) - a(i)}{\max(a(i), b(i))}$ Silouette score of a partition = average of the s(i)'s.

▶ Elbow method with silhouette score instead of distortion

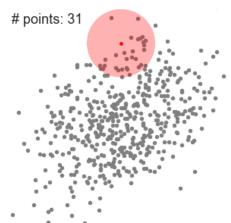
Outline

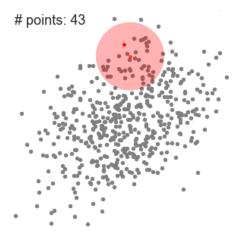
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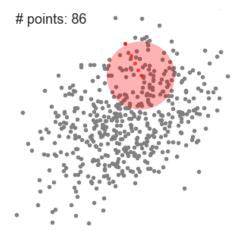
Mean-shift (with one centroid)

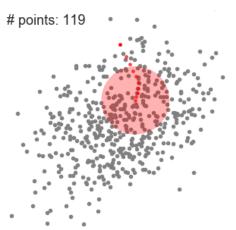


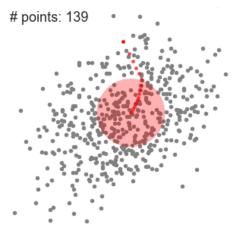
The centroid moves towards a higher density region

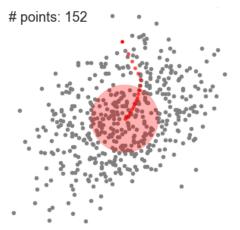


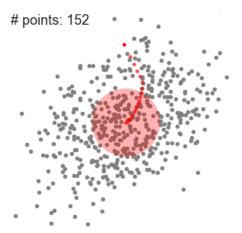




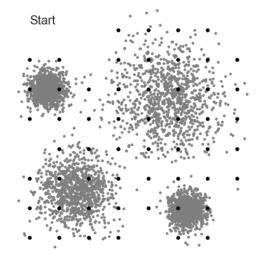




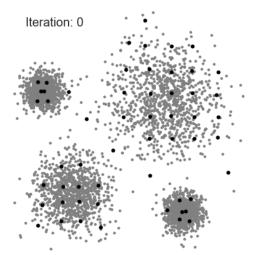




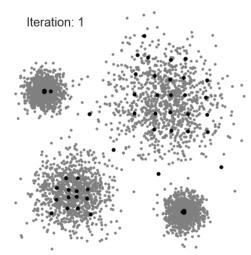
GMM: Gaussian Mixture Model Hierarchical clustering



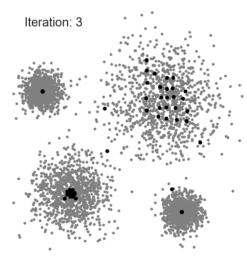
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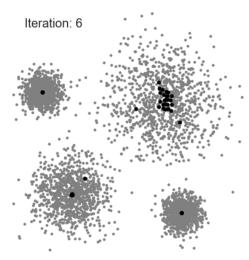
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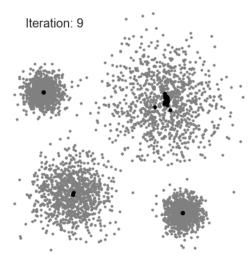
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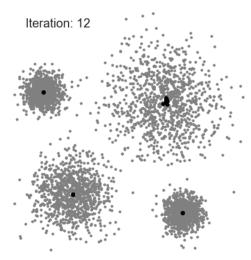
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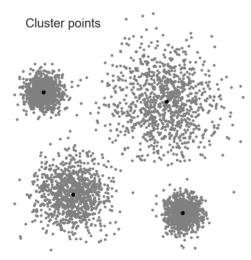
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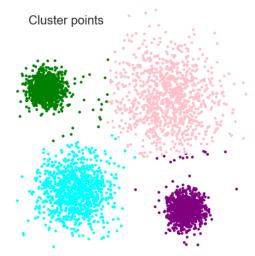
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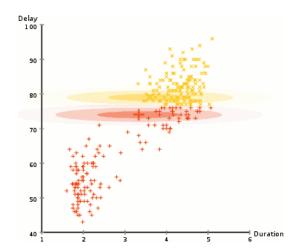
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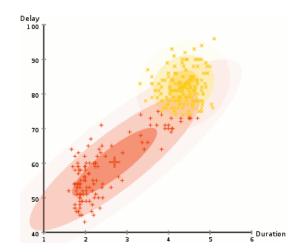


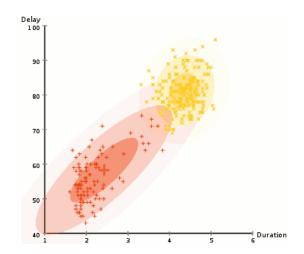
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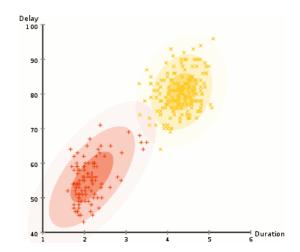


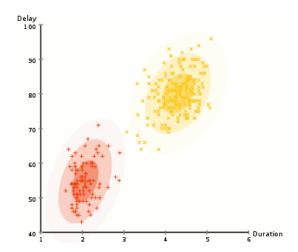
- Start with a given (large) number of circular sliding windows centered at randomly selected centroids and having radius r.
- While True;
 - Compute k new centroids by averaging examples in each sliding windows (the centroids are shifted towards regions of higher density)
 - If centroids don't change:
 - Break
- If multiple sliding windows overlap, then only the window containing the most points is preserved.
- Each data point is assigned to the nearest centroid.

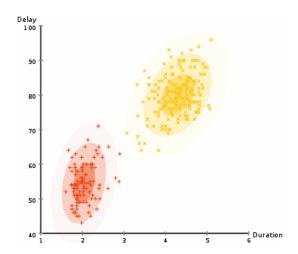












Model: K gaussians (one-dimentional):

•
$$p(x) = \sum_{k=1}^{K} \Phi_k N(x|\mu_k, \sigma_k)$$

•
$$N(x|\mu_k, \sigma_k) = \frac{1}{\sqrt{2\pi}\sigma_k} \exp\left(-\frac{(x-\mu_k)^2}{2\sigma_k^2}\right)$$

$$\bullet \sum_{k=1}^K \Phi_k = 1$$

- Want to find $(\Phi_k, \mu_k, \sigma_k)$ for all k maximizing $\prod_{i=1}^n p(x_i)$
- Chicken and egg problem:
 - need $(\Phi_k, \mu_k, \sigma_k)$ for all k to guess source of points
 - need to know source to estimate $(\Phi_k, \mu_k, \sigma_k)$

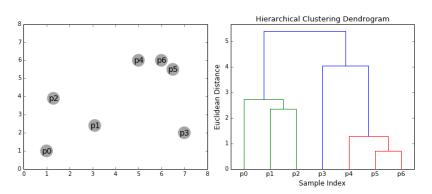
Expectation-Maximization algorithm:

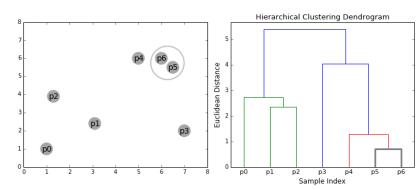
- start with randomly placed Gaussians $(\Phi_k, \mu_k, \sigma_k)$
- (E-step) for each i, k compute $\gamma_{i,k} \sim \text{did } x_i$ came from k?
- (M-step) adjust $(\Phi_k, \mu_k, \sigma_k)$ to fit points assigned to them
- iterate until convergence

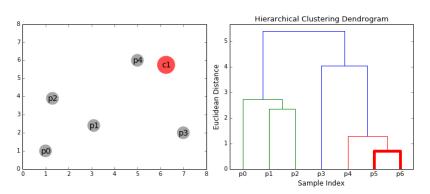
$$\gamma_{i,k} = \frac{\Phi_k \mathcal{N}(x_i | \mu_k, \sigma_k)}{\sum\limits_{k=1}^K \Phi_k(x | \mu_k, \sigma_k)}$$

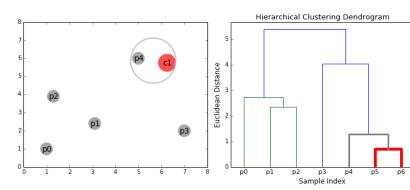
$$\Phi_k = \sum_{i=1}^N \frac{\gamma_{i,k}}{N}$$

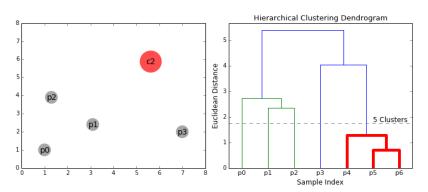
$$\mu_k = \frac{\sum_{i=1}^N \gamma_{i,k} x_i}{\sum_{i=1}^N \gamma_{i,k}} \quad \sigma_k^2 = \frac{\sum_{i=1}^N \gamma_{i,k} (x_i - \mu_k)^2}{\sum_{i=1}^N \gamma_{i,k}}$$

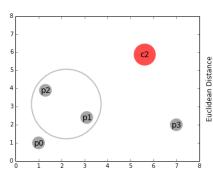


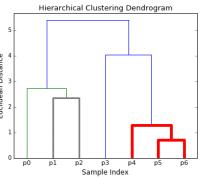




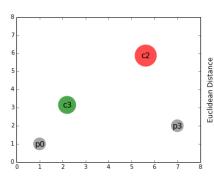


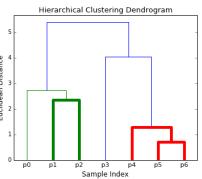


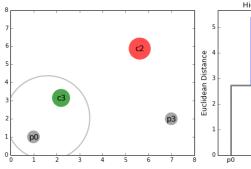


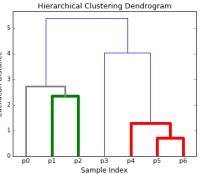


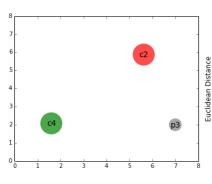
Mean-shift GMM: Gaussian Mixture Model Hierarchical clustering

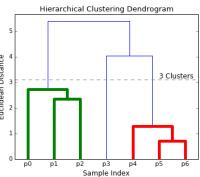


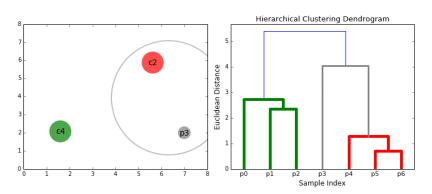


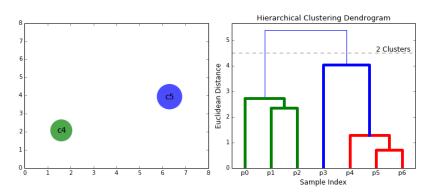


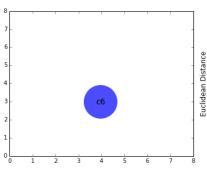


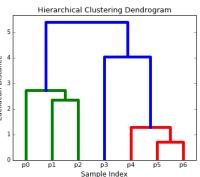












Two types of hierarchical clustering:

- Agglomerative. This is a "bottom-up" approach: each observation starts in its own cluster, and pairs of clusters are merged as one moves up the hierarchy.
- Divisive. This is a "top-down" approach: all observations start in one cluster, and splits are performed recursively as one moves down the hierarchy.

Aglomerative hierarchical clustering

Maximum (or complete) linkage clustering:

$$d(A,B) = \max\{d(a,b) : a \in A, b \in B\}$$

Minimum (or single) linkage clustering:

$$d(A,B) = \min\{d(a,b) : a \in A, b \in B\}$$

Average linkage clustering:

$$d(A,B) = \frac{1}{|A| \cdot |B|} \sum_{a \in A} \sum_{b \in B} d(a,b)$$

Divisive hierarchical clustering

Dasgupta's objective and algorithm: Repeatedly subdivides the elements using an approximation algorithm for the sparsest cut problem.

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Validation

- For supervised classification we have a variety of measures to evaluate how good our model is: Accuracy, precision, recall
- For clustering, the analogous question is how to evaluate the "goodness" of the resulting clusters?
- "Clusters are in the eye of the beholder"!
- We still want some tools to:
 - To avoid finding patterns in noise
 - To compare clustering algorithms
 - To compare two sets of clusters
 - To compare two clusters

Validation

Two types of numerical measures to judge cluster validity:

- Internal Index: Used to measure the goodness of a clustering structure without respect to external information. (e.g. distortion, silhoutte score)
- External Index: Used to measure the extent to which cluster labels match externally supplied class labels. (e.g. Entropy, Adjusted Rand Index)

Entropy

 Given a discrete random variable X with possible value {1,..,n}, entropy is defined as

$$H(X) = -\sum_{i=1}^{n} P(X = i) \log_2(P(X = i))$$

Entropy measures how uncertain is an event, the larger the entropy the more uncertain is the event.

Validation: External Index

Table 5.9. K-means Clustering Results for LA Document Data Set

Cluster	Entertainment	Financial	Foreign	Metro	National	Sports	Entropy	Purity
1	3	5	40	506	96	27	1.2270	0.7474
2	4	7	280	29	39	2	1.1472	0.7756
3	1	1	1	7	4	671	0.1813	0.9796
4	10	162	3	119	73	2	1.7487	0.4390
5	331	22	5	70	13	23	1.3976	0.7134
6	5	358	12	212	48	13	1.5523	0.5525
Total	354	555	341	943	273	738	1.1450	0.7203

 m_j = size of cluster j, m = number of documents p_{ij} = probability that a random document of cluster j belongs to topic i. For example, $p_{13} = 1/685$.

Entropy of a cluster:

$$e_j = -\sum_{i=1}^n p_{ij} \log_2(p_{ij})$$

• Entropy of a clustering: $\sum \frac{m_j}{m_i} e_i$

• Purity of a clustering: $\sum_{i=1}^{m_i} p_i w_i t_i v_i$

$$\sum_{i} \frac{m_{j}}{m} purity_{j}$$