# Data Mining Lecture 3: Discovering Groups

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#### Discovering Groups - Introduction

Understanding large datasets is hard, especially if it has high dimensional features

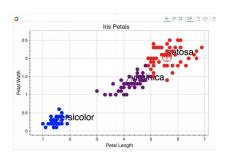
To help understand a dataset:

- Find similar data items
- Find similar features

#### Discovering Groups - Clustering

Grouping data, just using the feature vectors

- Unsupervised
- Similar feature vectors grouped together
- ► Can be
  - Soft (allow overlapping groups)
  - ► Hard (each item assigned to one group)



#### Discovering Groups - K Means

#### **Algorithm 1:** K Means clustering

Data: X, K

initialise K centroids;

while positions of centroids change do

for each data point do

assign to nearest centroid;

end

for each centroid do

move to average of assigned data points

end

end

return centroids, assignments;

A special case of Expectation Maximisation

K Means ipynb demo

K Means Java Demo

#### Hierarchical Clustering:

Creates a binary tree that recursively groups pairs of similar items or clusters

#### Can be:

- ► Agglomerative (bottom up)
- Divisive (top down)

# Algorithm 2: Hierarchical Agglomerative Clustering Data: N data points with feature vectors $X_i$ i = 1 ... NnumClusters = N; while numClusters > 1 do | cluster1, cluster2 = FindClosestClusters(); merge(cluster1, cluster2); end

The distance between the clusters is evaluated using a linkage criterion.

If each merge is recorded, a binary tree structure linking the clusters can be formed.

This gives a dendrogram

Linkage criterion: A measure of dissimilarity between clusters Centroid Based:

- ▶ Dissimilarity is equal to distance between centroids
- Needs numeric feature vectors

#### Distance-Based:

- Dissimilarity is a function of distance between items in clusters
- Only needs precomputed measure of similarity between items

#### Centroid based linkage:

- ▶ WPGMC: Weighted Pair Group Method with Centroids When two clusters are combined into a new cluster, the average of the two centroids is the new centroid
- ▶ UPGMC: Unweighted Pair Group Method with Centroids When two clusters are combined into a new cluster, the new centroid is recalculated based on the positions of the items

Distance based linkage:

► Minimum, or single-linkage clustering Distance between two closest members

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

Produces long, thin clusters

► Maximum, or complete-linkage clustering Distance between two most distant members

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

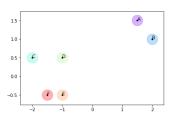
Finds compact clusters, approximately equal diameter

► Mean or Average Linkage Clustering (UPGMA: Unweighted Pairwise Group Method with Arithmetic Mean):

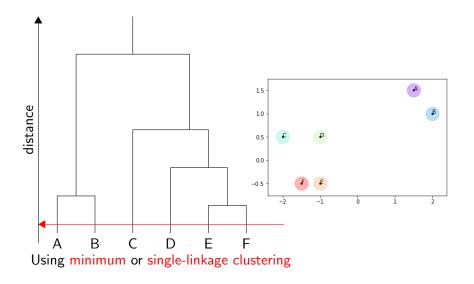
$$\frac{1}{|A||B|} \sum_{a \in A} \sum_{b \in B} d(a, b)$$

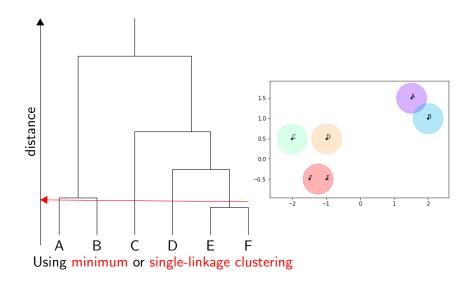
With sample data:

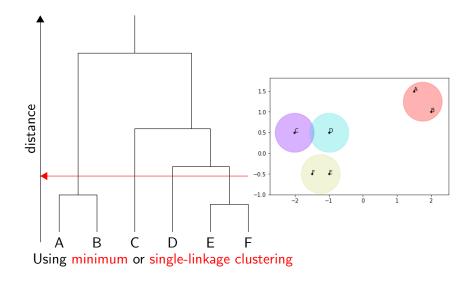
$$X = \begin{bmatrix} 1.5 & 1.5 \\ 2.0 & 1.0 \\ 2.0 & 0.5 \\ -1.0 & 0.5 \\ -1.5 & -0.5 \end{bmatrix}$$

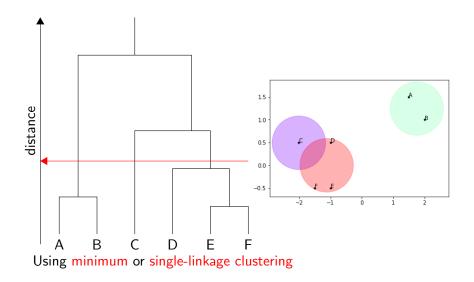


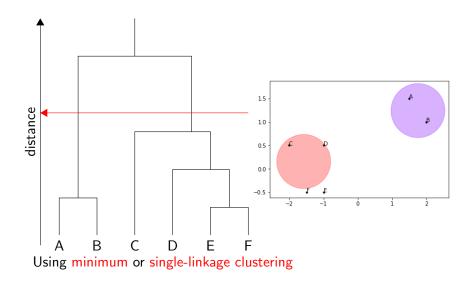
Using minimum or single-linkage clustering

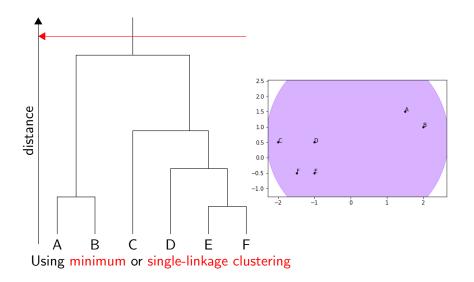


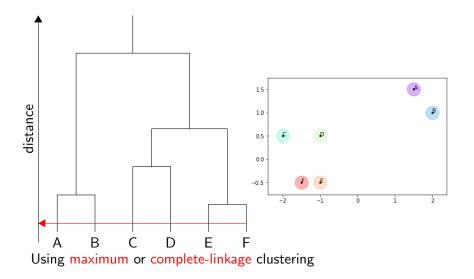


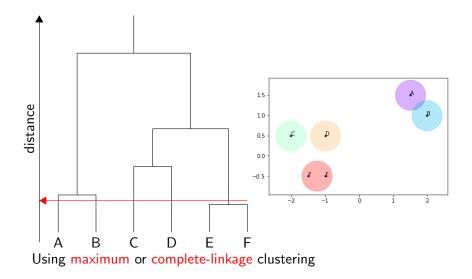


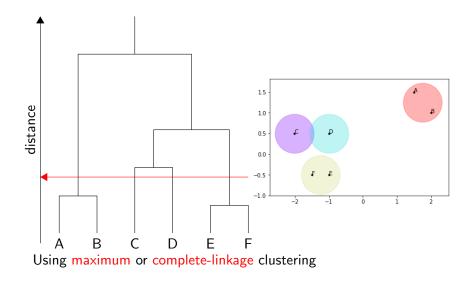


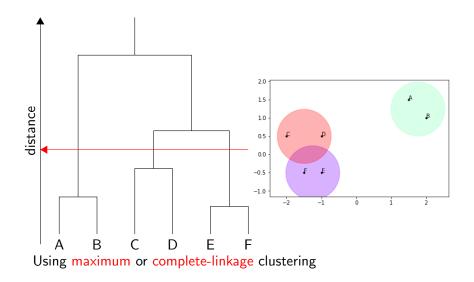


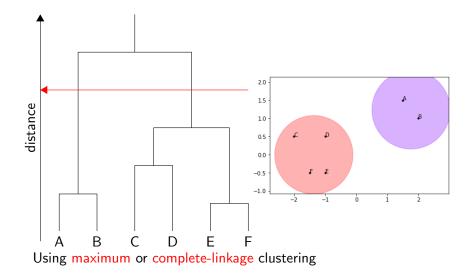


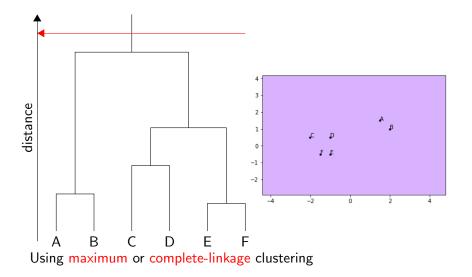












# Discovering Groups - Hierarchical Agglomerative Clustering

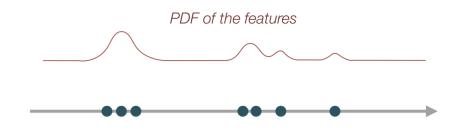
#### Java HAC Demo

Minimum distance linkage tends to give long thin clusters maximum distance linkage tends to give rounded clusters

Mean shift finds the *modes* of a probability density function.

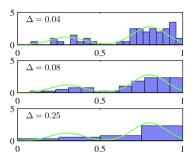
This means if finds the points in feature space with the highest feature density, i.e. are the most likely given the dataset Needs a kernel and a kernel bandwidth.

It is a hill climbing algorithm that



unbiased sample of 1D features from a dataset

How can we estimate the PDF? Could use a histogram, need to guess number of bins



Changing bin size affecting accuracy of probability density estimation<sup>1</sup>

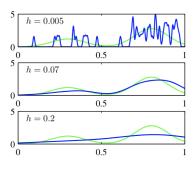
#### Can be too crude

<sup>&</sup>lt;sup>1</sup>C. Bishop, Pattern Recognition and Machine Learning

Kernel Density Estimation (aka Parzen Window) Gives a smooth continuous estimate

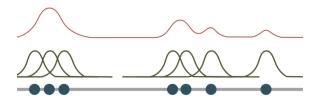
$$f(x) = \frac{1}{nh^d} \sum_{i=1}^n K(\frac{x - x_i}{h})$$

Where nh is the number of items, d is the dimensionality of the feature space, K is the kernel function, x is an arbitrary position in feature space, h is the kernel bandwidth



Changing bandwidth affecting accuracy of probability density estimation

Usually use a Gaussian kernel with  $\sigma=1$ If kernel is radially symmetric, then only need profile of kernel, k(x) that satisfies  $K(x)=C_{k,d}k(||x||^2)$ 



Find the modes of the probability density function (PDF), i.e. where the gradient is zero.  $\Delta f(x) = 0$ 



$$f(x) = \frac{1}{nh^d} \sum_{i=1}^n K\left(\frac{x - x_i}{h}\right)$$
$$K(x) = c_{k,d} k(||x||^2)$$

Where  $c_{k,d}$  is a normalisation constant

$$f(x) = \frac{c_{k,d}}{nh^d} \sum_{i=1}^{n} k(||\frac{x - x_i}{h}||^2)$$

Assuming a radially symmetric kernel:

$$\Delta f(x) = \frac{2c_{k,d}}{nh^{d+2}} \sum_{i=1}^{n} (x - x_i) g(||\frac{x - x_i}{h}||^2) \quad g(x) = -k'(x)$$

$$\Delta f(x) = \frac{2c_{k,d}}{nh^{d+2}} \sum_{i=1}^{n} g(||\frac{x - x_i}{h}||^2) \frac{\sum_{i=1}^{n} x_i g(||\frac{x - x_i}{h}||^2)}{\sum_{i=1}^{n} g(||\frac{x - x_i}{h}||^2)} - x$$

$$\Delta f(x) = \frac{\frac{2c_{k,d}}{nh^{d+2}} \sum_{i=1}^{n} g(||\frac{x-x_i}{h}||^2)}{\sum_{i=1}^{n} g(||\frac{x-x_i}{h}||^2)} \frac{\sum_{i=1}^{n} x_i g(||\frac{x-x_i}{h}||^2)}{\sum_{i=1}^{n} g(||\frac{x-x_i}{h}||^2)} - x$$

The first part is a probability density estimate with kernel  $G(x) = x_{g,d}g(||x||^2)$ 

$$\Delta f(x) = \frac{2c_{k,d}}{nh^{d+2}} \sum_{i=1}^{n} g(||\frac{x - x_i}{h}||^2) \frac{\sum_{i=1}^{n} x_i g(||\frac{x - x_i}{h}||^2)}{\sum_{i=1}^{n} g(||\frac{x - x_i}{h}||^2)} - x$$

The first part is a probability density estimate with kernel  $G(x) = x_{g,d}g(||x||^2)$ 

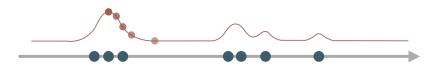
The second part is the mean shift, the vector that always points in the direction of maximum density

#### Mean shift algorithm:

#### Algorithm 3: Mean Shift Procedure

**Data:** N data points with feature vectors  $X_i$  i = 1...N while  $x_t not = x_{t+1}$  do  $m_h(x_t) = \text{computeMeanShiftVect}();$   $x_{t+1} = x_t + m_h(x_t);$ 

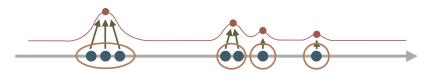
#### end



#### For each feature vector:

- ▶ apply mean shift procedure until convergence
- store resultant mode

Set of feature vectors that converge to the same mode define the basin of attraction of that mode



#### Discovering Groups - Summary

Clustering is a key way to understand your data.

There are many different approaches

- K Means Need to chose K
- Hierarchical Agglomerative Clustering -
- Mean Shift Clustering

They are a very good way to start exploring a dataset