



数据挖掘导论

Introduction to Data Mining

Advanced Clustering



数据智能实验室
DATA INTELLIGENCE LABORATORY



浙江大学
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Agenda

- **Soft Clustering**
- Gaussian Mixture Model
- Spectral Clustering
- Bi-Clustering
- Summary

Soft Clustering

- ❑ In hard clustering (e.g, k-means), each point belongs to only one cluster
- ❑ In soft clustering, each point can belong to multiple classes
 - ✓ An apple could be red or green
 - ✓ In Marketing, each user may belong to multiple target audience groups
- ❑ There is a weight or probability between each point and each cluster

$$SSE = \sum_{j=1}^k \sum_{i=1}^N w_{ij}^m (x_i - c_j)^2, \quad \sum_{j=1}^k w_{ij} = 1$$

- ✓ m is any real number greater than 1

Fuzzy C-means Clustering

□ Extends k-means clustering with the following idea:

✓ In the iterations, repeat the following steps:

- Fix c_j and determine the best w_{ij}

$$w_{ij} = \frac{1}{\sum_{k=1}^C \left(\frac{\|x_i - c_j\|}{\|x_i - c_k\|} \right)^{\frac{2}{m-1}}}$$

- Fix w_{ij} and recompute c_j

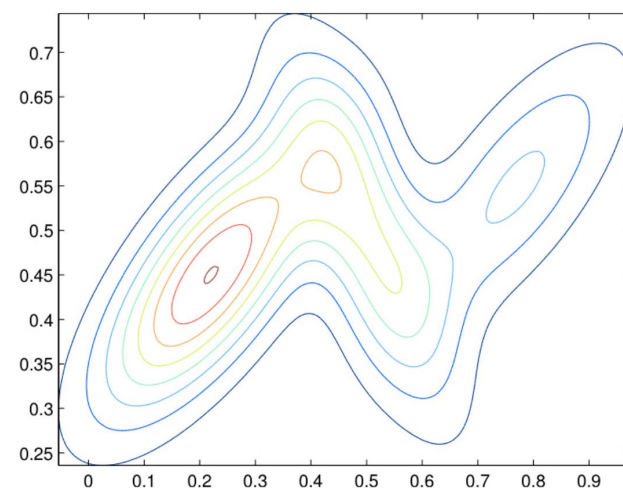
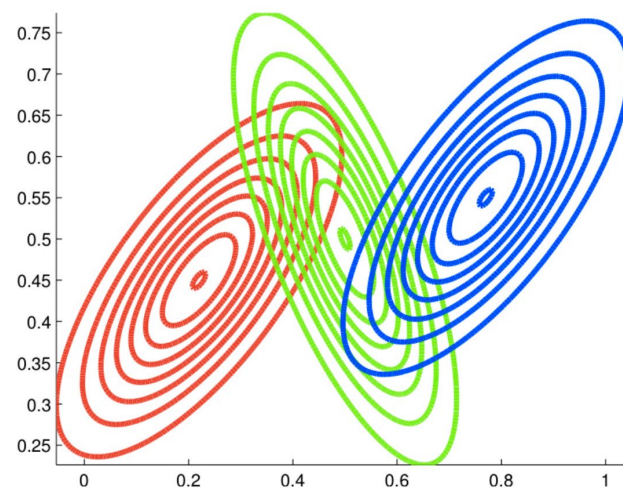
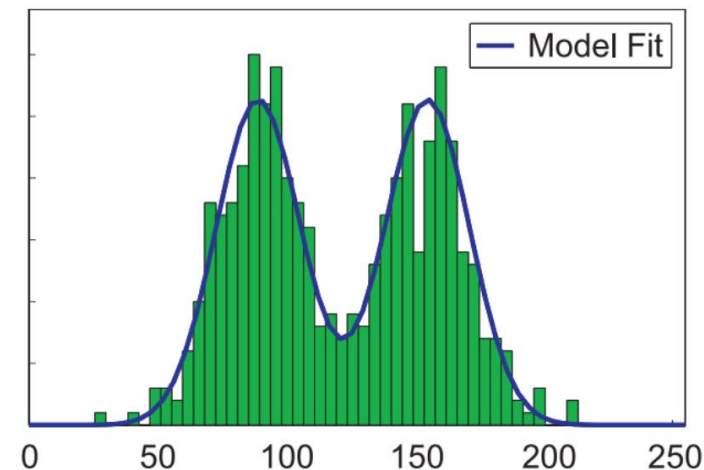
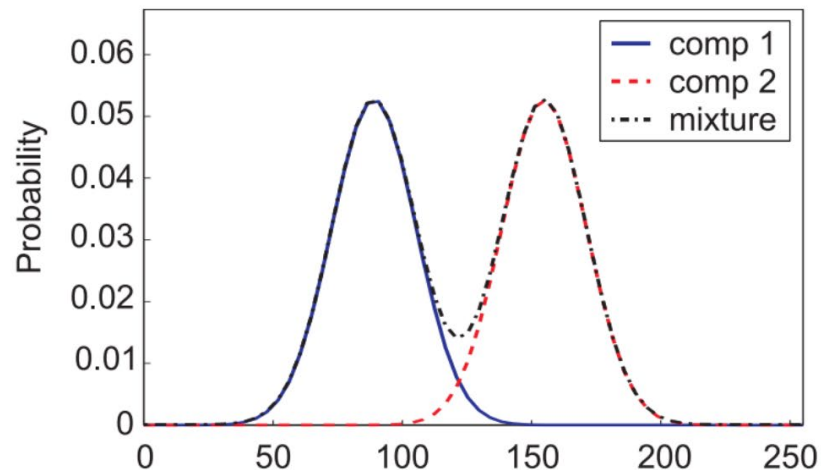
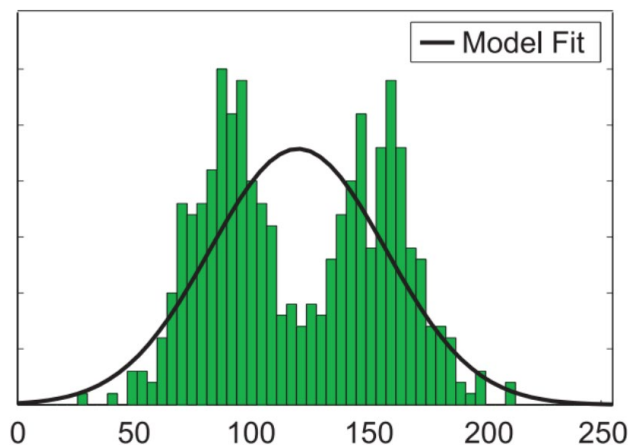
$$c_j = \frac{\sum_{i=1}^N w_{ij}^m \cdot x_i}{\sum_{i=1}^N w_{ij}^m}$$

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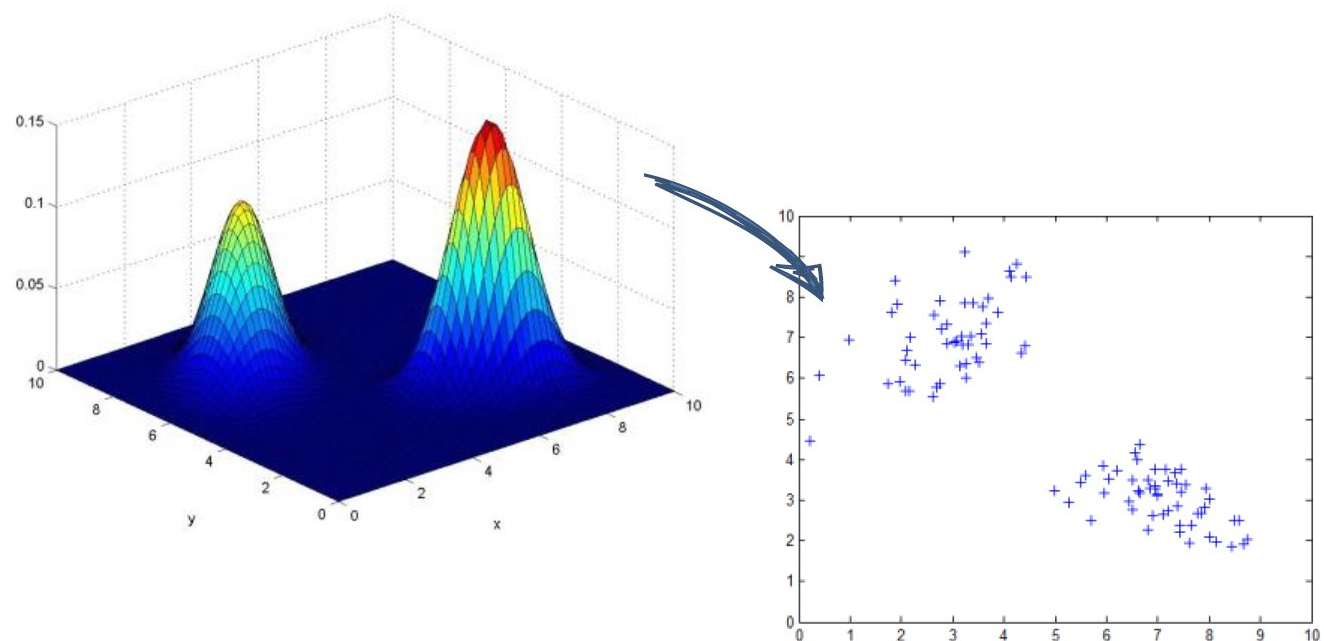
Clustering based on Mixture Model

□ Fit data with mixture of Gaussian distributions



Main Idea

- ❑ The data points are generated by the underlying mixture of Gaussian models
- ❑ Treat each cluster as one component of the mixture distribution, **whose mean is the cluster center**
- ❑ Each point has a certain probability of belonging to each cluster



EM Algorithm

□ Expectation Maximization Algorithm

- ✓ **E-step:** Compute the posterior probability over z given our current model - i.e. how much do we think each Gaussian generates each datapoint

- Estimate the probability of each point to a cluster

$$\gamma_{ik} = \frac{\pi_k \mathcal{N}(\mathbf{x}_i | \boldsymbol{\mu}_k, \Sigma_k)}{\sum_{j=1}^K \pi_j \mathcal{N}(\mathbf{x}_i | \boldsymbol{\mu}_j, \Sigma_j)}$$

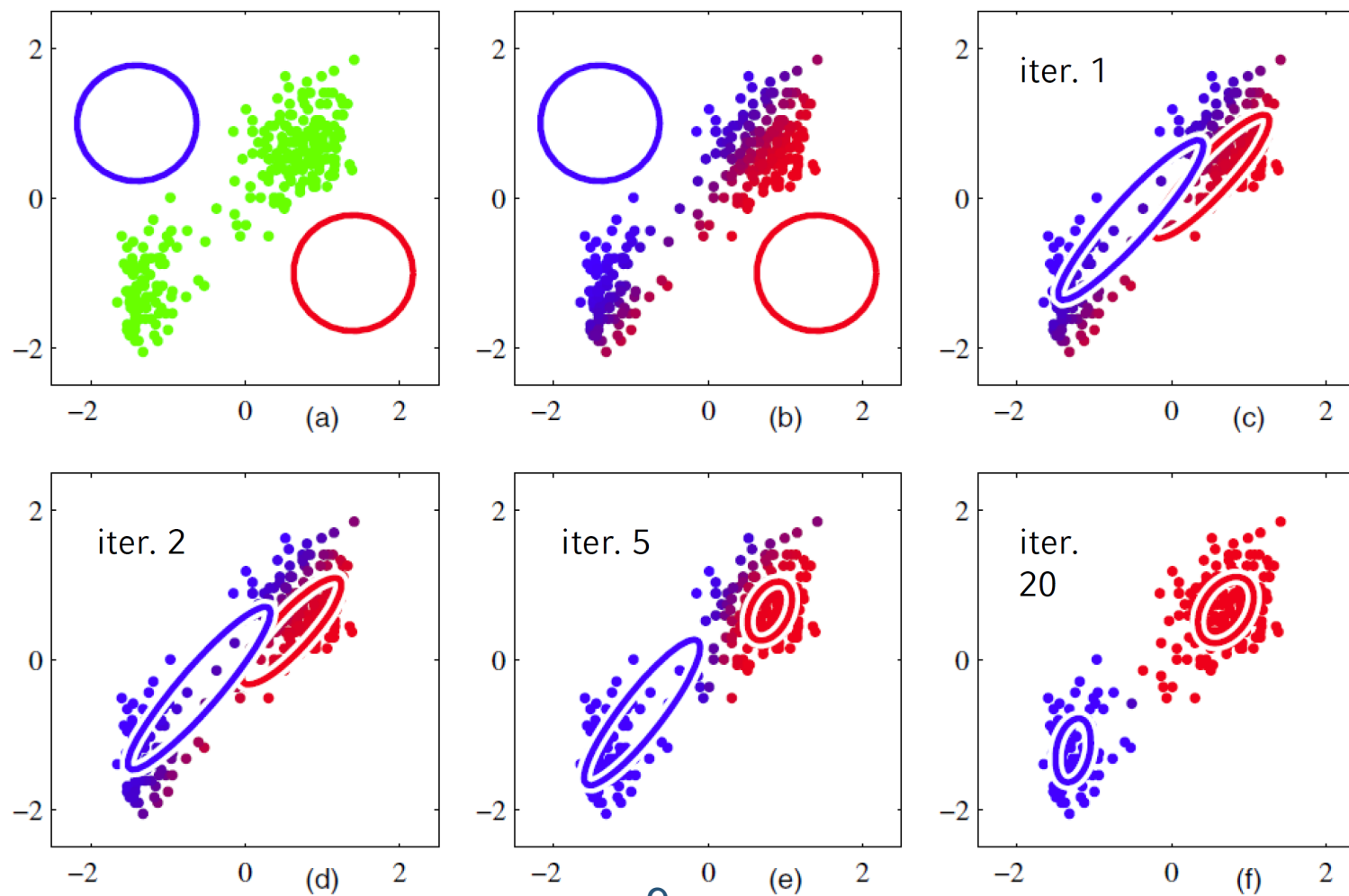
- ✓ **M-step:** Assuming that the data really was generated this way, change the parameters of each Gaussian to maximize the probability that it would generate the data it is currently responsible for

- ✓ Update the parameters of Gaussian distributions

$$\begin{aligned}\boldsymbol{\mu}_k &= \frac{1}{N_k} \sum_{i=1}^N \gamma_{ik} \mathbf{x}_i \\ \Sigma_k &= \frac{1}{N_k} \sum_{i=1}^N \gamma_{ik} (\mathbf{x}_i - \boldsymbol{\mu}_k)(\mathbf{x}_i - \boldsymbol{\mu}_k)^\top \\ \pi_k &= \frac{N_k}{N} \quad \text{where } N_k = \sum_{i=1}^N \gamma_{ik}\end{aligned}$$

Main Idea

- Iteratively improve the parameters of each distribution until some quality threshold is reached



Agenda

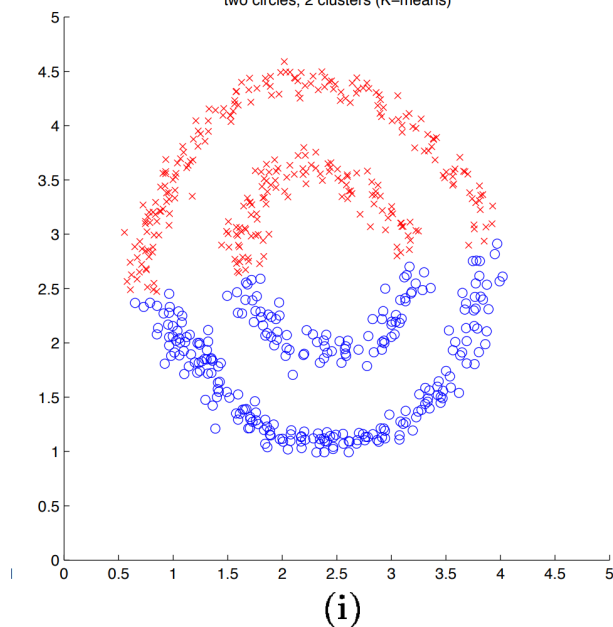
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Spectral Clustering

- ❑ Clusters are generated based on pairwise proximity/similarity/affinity
- ❑ Clusters do not have to be Gaussian or compact

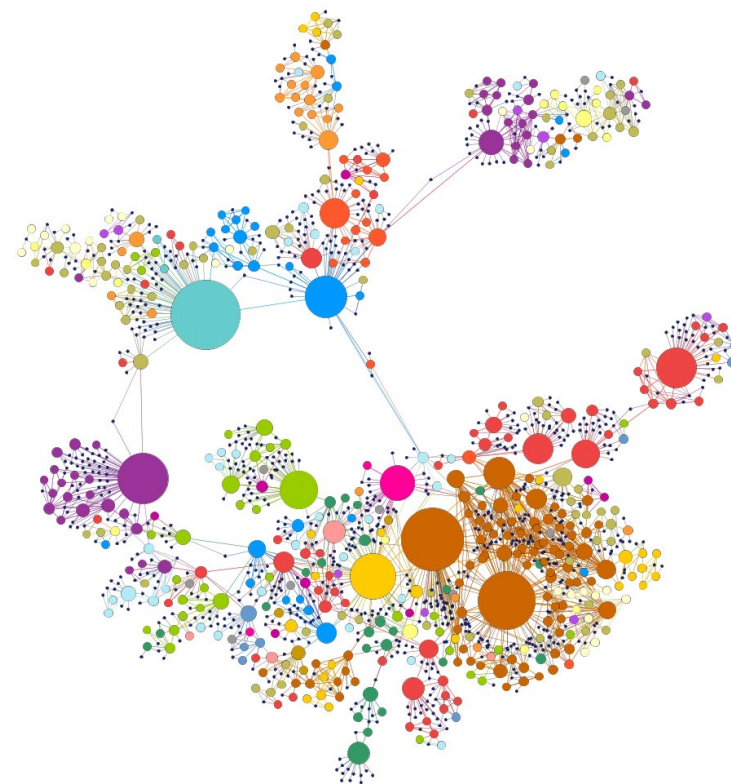
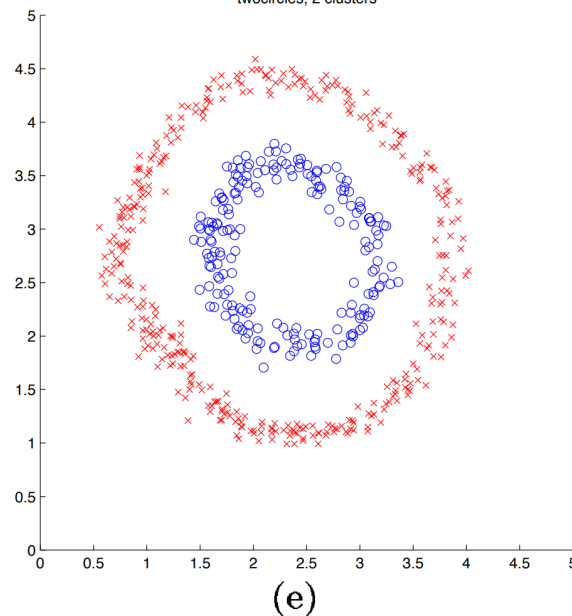
K-means

two circles, 2 clusters (K-means)



Spectral clustering

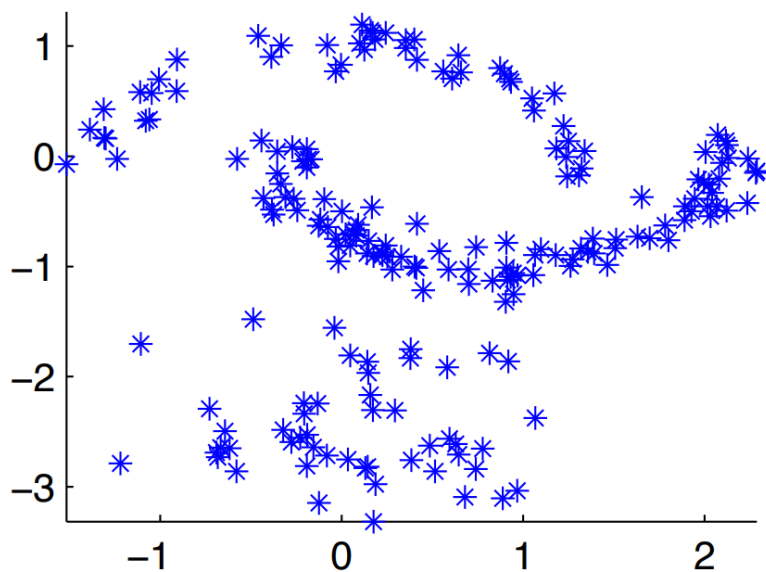
twocircles, 2 clusters



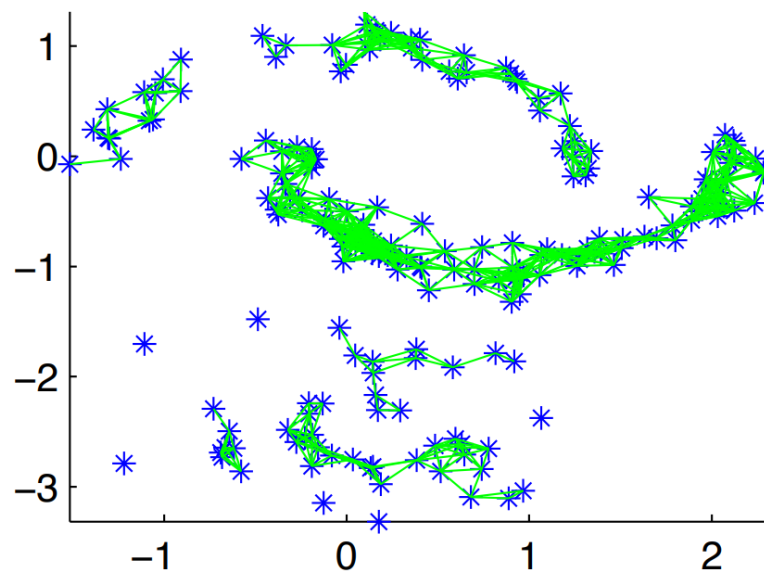
Similarity Graph Construction

- Compute the similarity between two objects
- Can use a threshold or kNN to reduce the number of edges

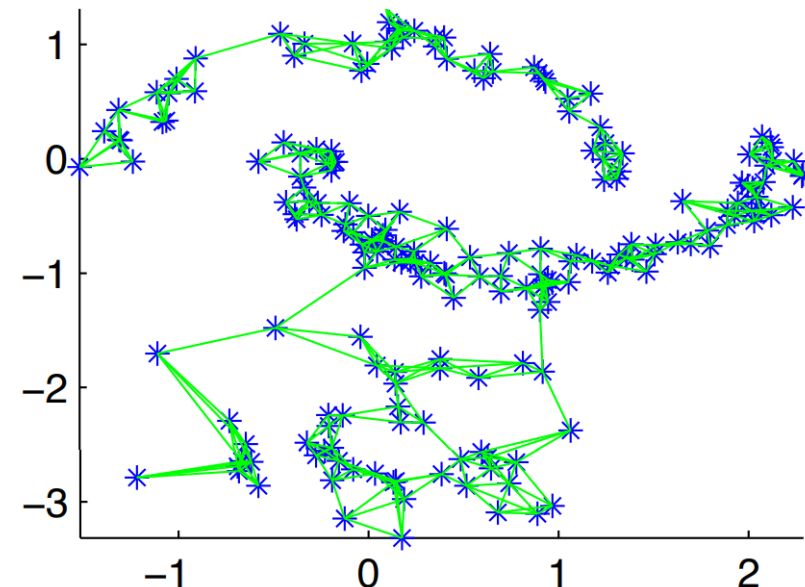
Data points



epsilon-graph, epsilon=0.3



kNN graph, k = 5

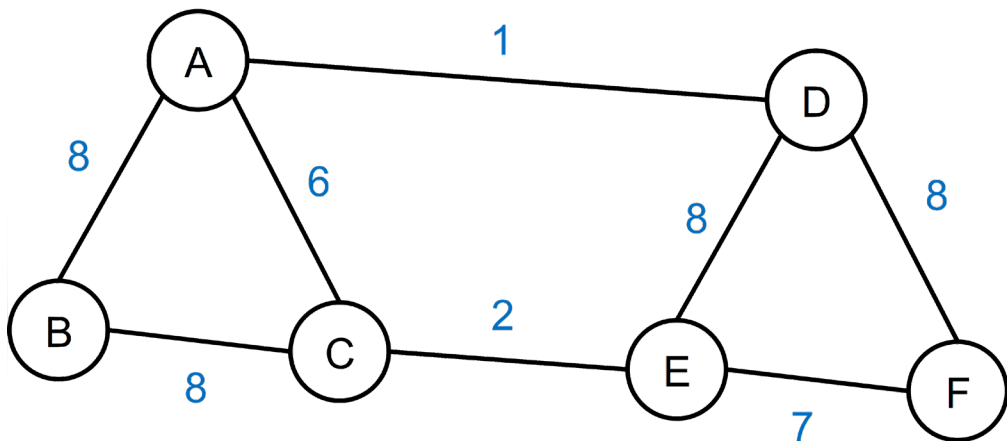


Spectral Clustering Algorithm

1. Create a similarity graph between our N objects to cluster.
2. Compute the first k eigenvectors of its Laplacian matrix to define a feature vector for each object.
3. Run k-means on these features to separate objects into k classes.

An Example of Spectral Clustering

□ Get Similarity Matrix



$$S = \begin{bmatrix} 0 & 8 & 6 & 1 & 0 & 0 \\ 8 & 0 & 8 & 0 & 0 & 0 \\ 6 & 8 & 0 & 0 & 2 & 0 \\ 1 & 0 & 0 & 0 & 8 & 8 \\ 0 & 0 & 2 & 8 & 0 & 7 \\ 0 & 0 & 0 & 8 & 7 & 0 \end{bmatrix}.$$

An Example of Spectral Clustering

□ Get Laplacian matrix

$$L = \begin{bmatrix} 15 & -8 & -6 & -1 & 0 & 0 \\ -8 & 16 & -8 & 0 & 0 & 0 \\ -6 & -8 & 16 & 0 & -2 & 0 \\ -1 & 0 & 0 & 17 & -8 & -8 \\ 0 & 0 & -2 & -8 & 17 & -7 \\ 0 & 0 & 0 & -8 & -7 & 15 \end{bmatrix}.$$

$$L_{\text{sym}} = D^{-1/2} L D^{1/2}$$

$$L_{\text{sym}} = \begin{bmatrix} 0.130 & -0.069 & -0.051 & -0.008 & 0. & 0. \\ -0.069 & 0.137 & -0.068 & 0. & 0. & 0. \\ -0.051 & -0.068 & 0.137 & 0. & -0.017 & 0. \\ -0.008 & 0. & 0. & 0.145 & -0.068 & -0.068 \\ 0. & 0. & -0.017 & -0.068 & 0.145 & -0.060 \\ 0. & 0. & 0. & -0.068 & -0.060 & 0.130 \end{bmatrix}$$

$L = D - S$

D is a simple diagonal matrix

D_{ii} equals the sum of i th row of S

An Example of Spectral Clustering

- Take the first k_1 eigenvectors, and to form a $N \times k_1$ matrix U
- Form a matrix T from U by normalizing the rows of U to norm 1
- Perform k-means clustering on the points in matrix T . Each row is considered as a point.

$$T = \begin{bmatrix} 0.706 & -0.708 \\ 0.677 & -0.735 \\ 0.738 & -0.674 \\ 0.710 & 0.703 \\ 0.740 & 0.672 \\ 0.677 & 0.735 \end{bmatrix}$$

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Motivation of Bi-Clustering

□ Simultaneous clustering of both rows and columns of a data matrix

- ✓ Bob is planning a housewarming party for his new 3-room house
- ✓ He owns 30 albums and wants to play different music in each room
- ✓ He has invited 50 guests and sent out a survey to each guest asking if they like each album
- ✓ He collects the data into a 50×30 binary matrix M , where $M_{ij}=1$ if guest i likes album j
- ✓ Bob wants to distribute people and albums evenly among the rooms of his house

$$s(\mathbf{M}, \mathbf{r}, \mathbf{c}) = b(\mathbf{r}, \mathbf{c}) \cdot \sum_{i,j,k} M_{ij} r_{ki} c_{kj} \quad b(\mathbf{r}, \mathbf{c}) = \exp\left(\frac{-(\max(\mathcal{S}) - \min(\mathcal{S}))}{\epsilon}\right)$$

- ✓ $b(\mathbf{r}, \mathbf{c})$ penalizes unbalanced solutions
- ✓ $r_{ki}=1$ if guest i is assigned to room k ; $c_{kj}=1$ if room k contains album j

Motivation of Bi-Clustering

- Starting with a random assignment of rows and columns to clusters, Bob **reassigns row and columns to improve the objective function** until convergence
 - ✓ Can use heuristic algorithms (simulated annealing etc) in operation research



Bi-clustering
Re-organize the order of
guests and albums



Room 3

Bi-clustering on Gene Data

	Conditions							
	A	B	C	D	E	F	G	H
Gene 1	Red	Green	White	Red	Green	Green	Red	Red
Gene 2	White	White	White	White	White	White	White	White
Gene 3	White	White	White	White	White	White	White	White
Gene 4	Red	Green	White	Red	Green	Green	Red	Red
Gene 5	White	White	White	White	White	White	White	White
Gene 6	White	Green	White	White	Green	Green	White	White
Gene 7	White	Green	White	White	Green	Green	White	White
Gene 8	White	White	White	White	White	White	White	White
Gene 9	Red	Green	White	Red	Green	Green	Red	Red

Clustering

	A	B	C	D	E	F	G	H
Gene 1	Red	Green	White	Red	Green	Green	Red	Red
Gene 4	Red	Green	White	Red	Green	Green	Red	Red
Gene 9	Red	Green	White	Red	Green	Green	Red	Red

Biclustering

Biclustering discovers local coherences over a subset of conditions

	A	B	D	E	F	G	H
Gene 1	Red	Green	Red	Green	Green	Red	Red
Gene 4	Red	Green	Red	Green	Green	Red	Red
Gene 9	Red	Green	Red	Green	Green	Red	Red

	B	E	F
Gene 1	Green	Green	Green
Gene 4	Green	Green	Green
Gene 6	Green	Green	Green
Gene 7	Green	Green	Green
Gene 9	Green	Green	Green

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