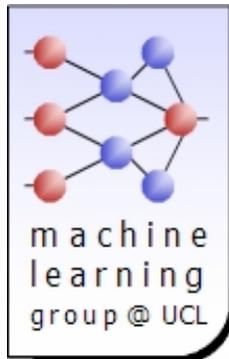


Collaborative filtering based on a random walk model on a graph

Marco Saerens, Francois Fouss, Alain Pirotte, Luh Yen, Pierre Dupont (UCL)

Jean-Michel Renders (Xerox Research Europe)



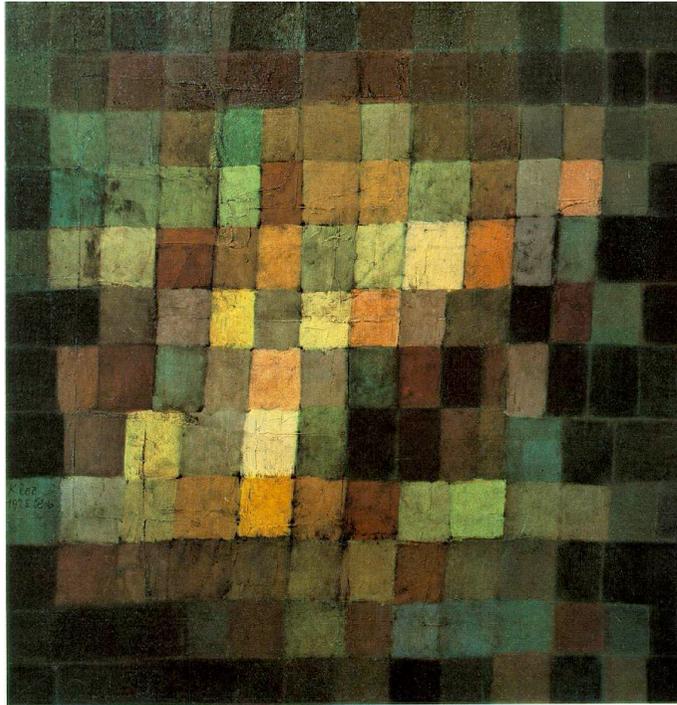
Université catholique de Louvain

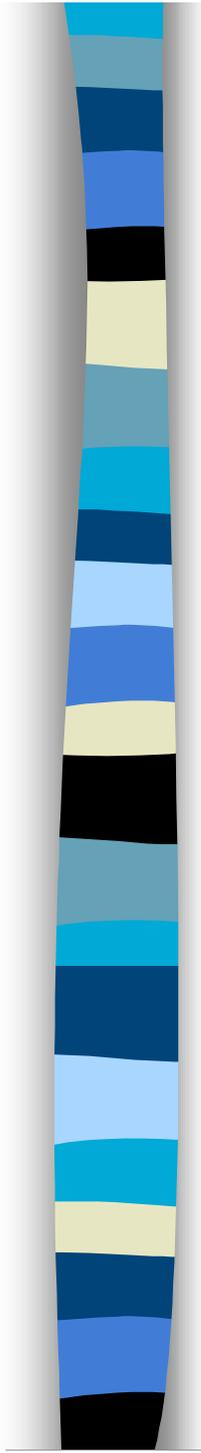
Université partenaire de l'Académie universitaire 'Louvain'



Some recent methods: web link analysis

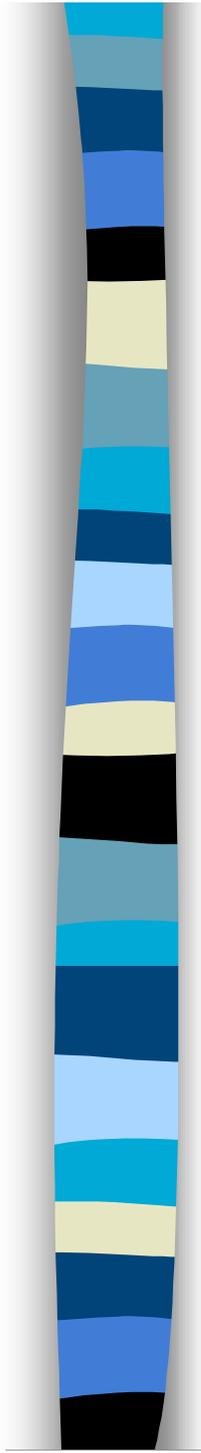
Exploiting links between documents





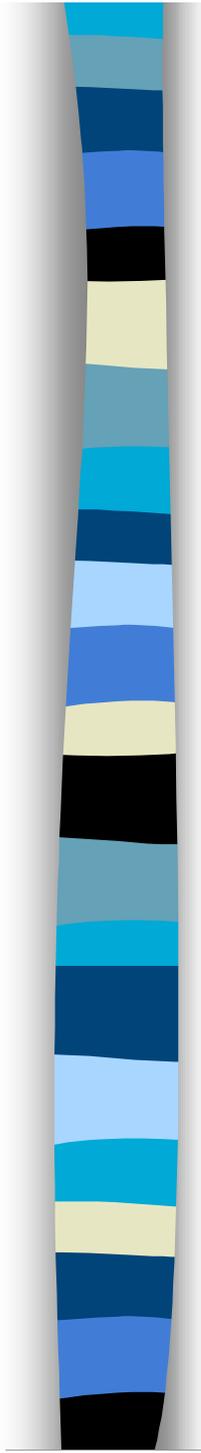
Web link analysis

- Suppose we performed a web search with a search engine
- **Objective:**
 - To improve the (content-based) ranking of the search engine
 - Based on the graph structure of the web hyperlinks



Web link analysis

- The objective here is to exploit the **links between documents** (hyperlinks)
- Documents/hyperlinks can be viewed as a **directed graph**
- Two algorithms will be introduced:
 - PageRank
 - HITS
- Then, we will introduce a more general algorithm for **collaborative recommendation**



Web link analysis

- A set of techniques

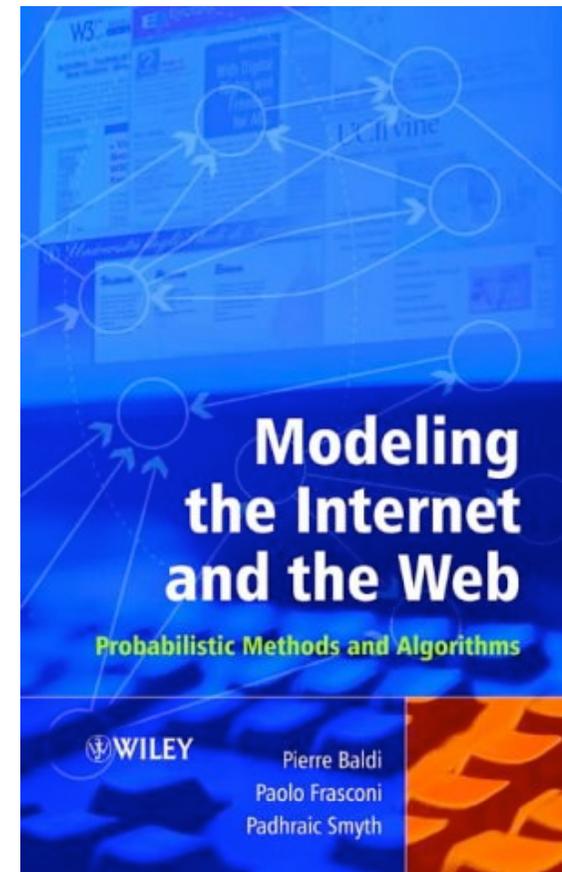
- Applied to: Hyperlink document repositories
- Typically web pages

- Objective:

- To exploit the link structure of the documents
- In order to extract interesting information
- Viewing the document repository as a graph where
 - Nodes are documents
 - Edges are directed links between documents

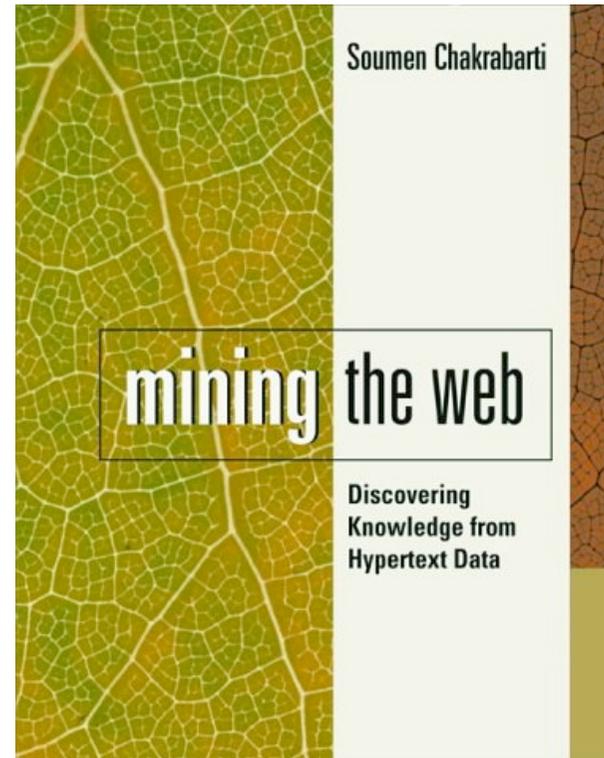
(1) The PageRank algorithm

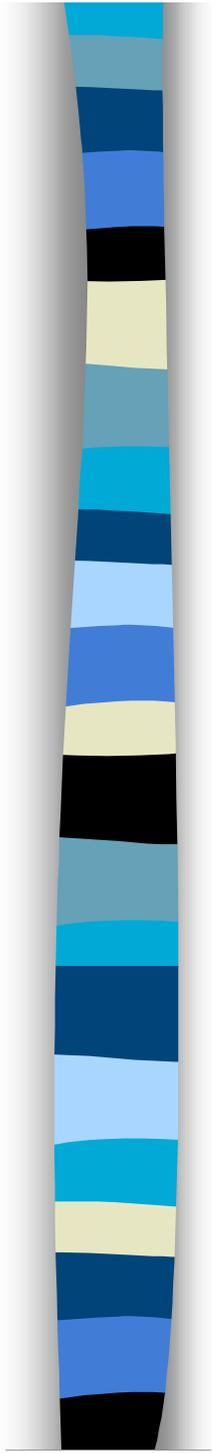
- P. Baldi, P. Frasconi & P. Smyth (2003)
 - Modeling the Internet and the Web
 - J. Wiley & Sons



The PageRank algorithm

- S. Chakrabarti (2003)
 - Mining the Web
 - Morgan Kaufmann



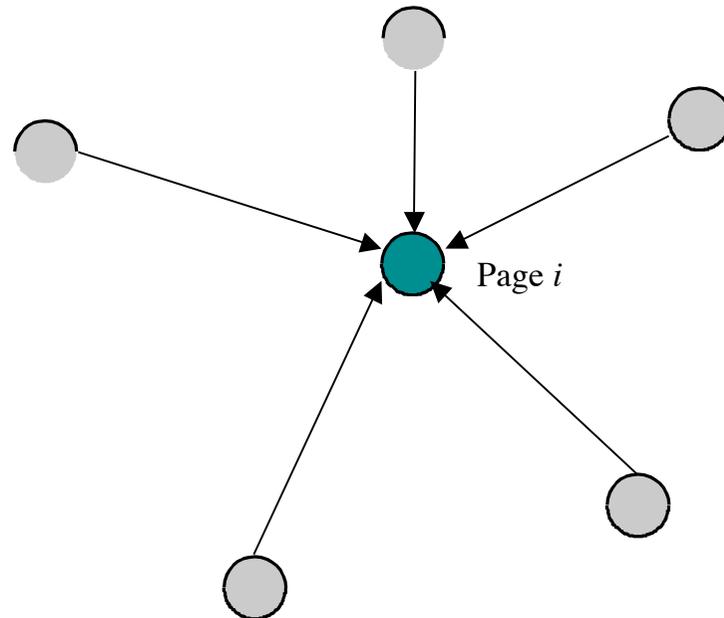


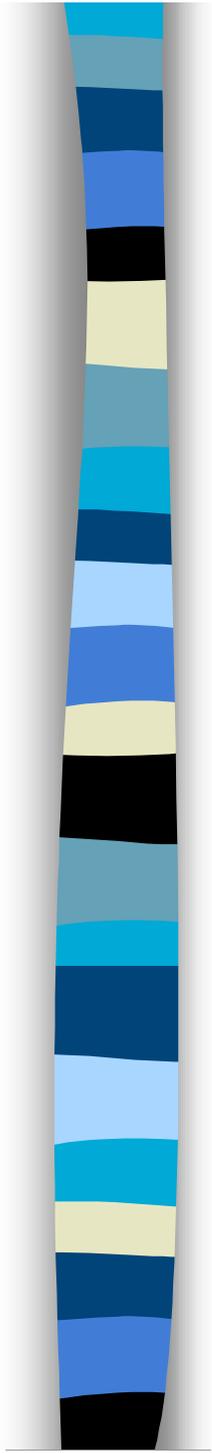
The PageRank algorithm

- Introduced by Page, Brin, Motwani & Winograd in 1998
- Partly implemented in Google

The PageRank algorithm

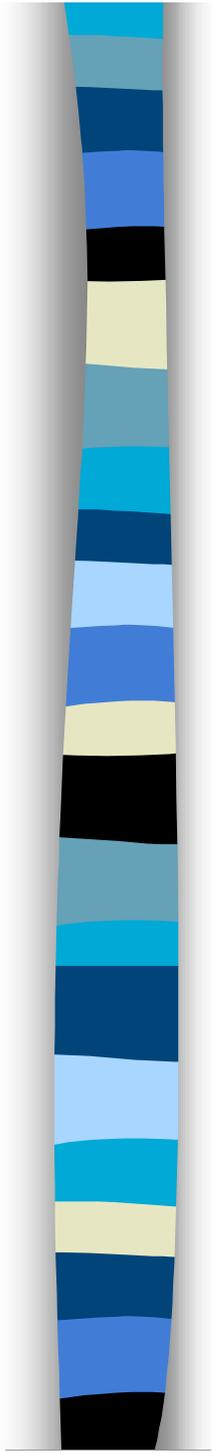
- To each web page we associate a score, x_i
 - The score of page i , x_i , is proportional to the weighted averaged score of the pages pointing to page i





The PageRank algorithm

- Let w_{ij} be the weight of the link connecting page i to page j
 - Usually, it is simply 0 or 1
 - Thus, $w_{ij} = 1$ if page i has a link to page j ; $w_{ij} = 0$ otherwise



The PageRank algorithm

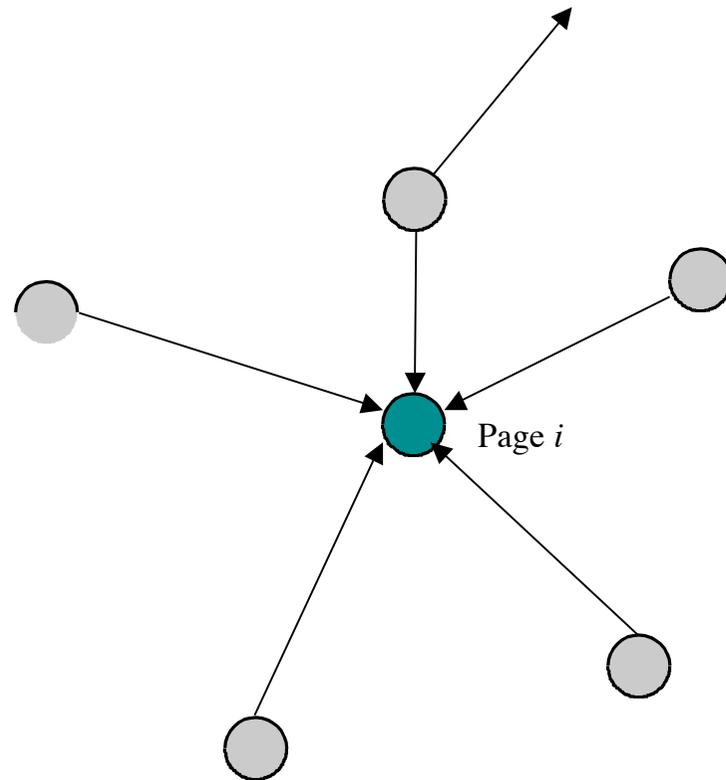
- Let \mathbf{W} be the adjacency matrix made of the elements w_{ij}
 - Notice that this matrix is not symmetric
 - We suppose that the graph is strongly connected

The PageRank algorithm

- In other words

$$x_i \propto \sum_{j=1}^n \frac{w_{ji} x_j}{w_j}$$

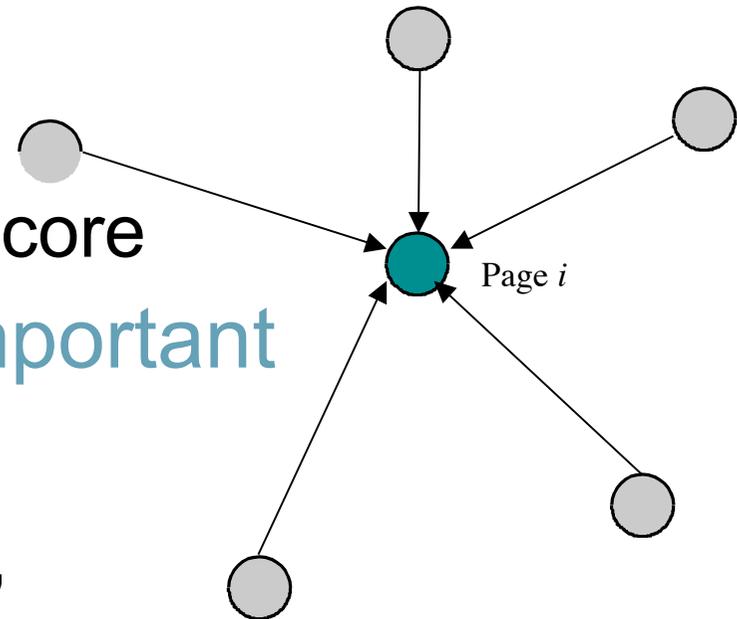
$$w_j = \sum_{i=1}^n w_{ji}$$

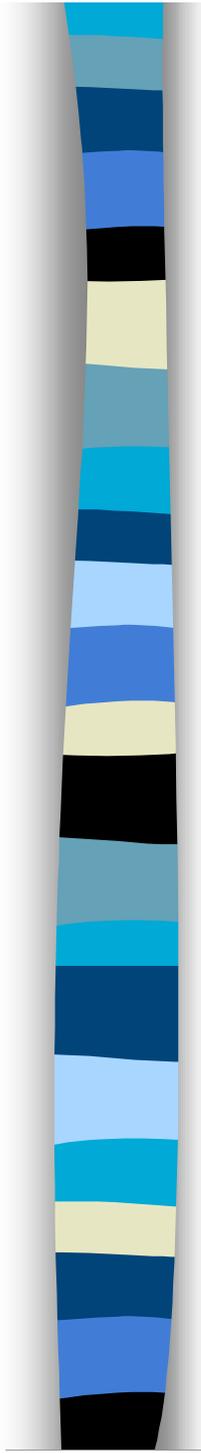


- where w_j is the outdegree of page j

The PageRank algorithm

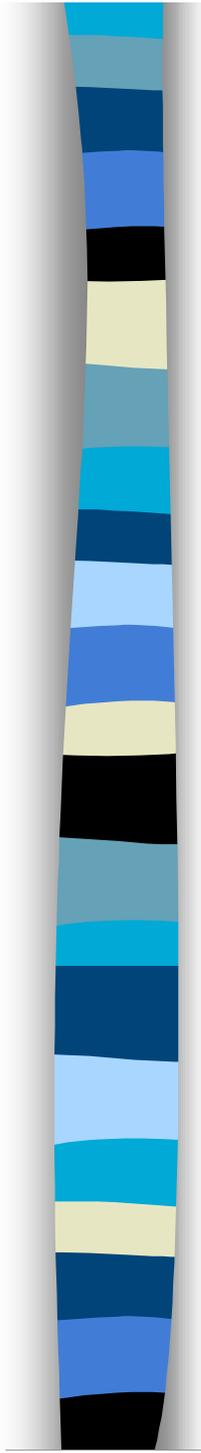
- In other words,
- A page with a **high score** is a page that is pointed by
 - Many pages
 - Having each a high score
- Thus a page is an **important page** if
 - It is pointed by **many, important, pages**





The PageRank algorithm

- These equations can be updated **iteratively** until convergence
- In order to obtain the scores, x_i
 - We normalize the vector \mathbf{x} at each iteration
- The pages are then **ranked** according to their score

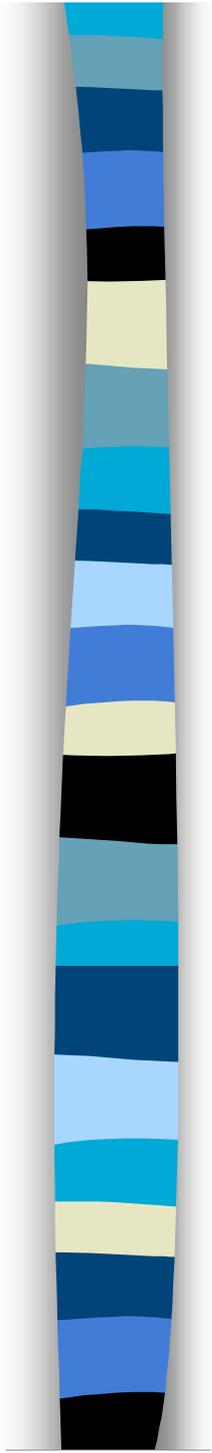


The PageRank algorithm

- This definition has a nice interpretation in terms of **random surfing**
- If we define the probability of following the link between page i to page j as

$$P(\text{page}(k+1) = i | \text{page}(k) = j) = \frac{w_{ji}}{w_j}$$

$$w_j = \sum_{i=1}^n w_{ji}$$



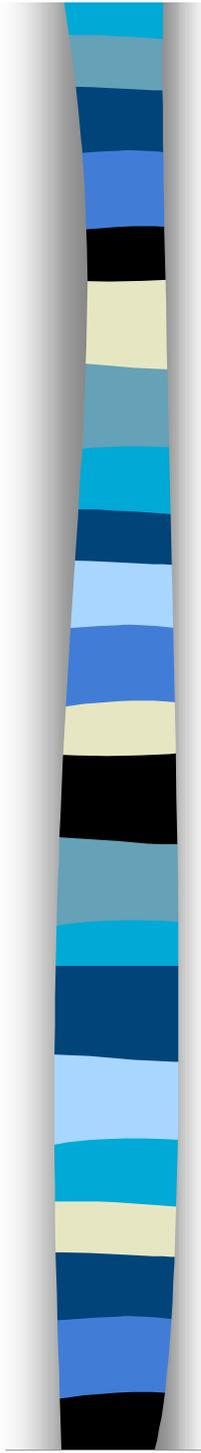
The PageRank algorithm

- We can write the updating equation as

$$\begin{aligned}x_i(k+1) &= P(\text{page}(k+1) = i) \\ &= \sum_{j=1}^n P(\text{page}(k+1) = i | \text{page}(k) = j) x_j(k)\end{aligned}$$

- And thus we can define a **random surfer** following the links according to the **transition probabilities**

$$P(\text{page}(k+1) = i | \text{page}(k) = j) = \frac{w_{ji}}{w_j}.$$

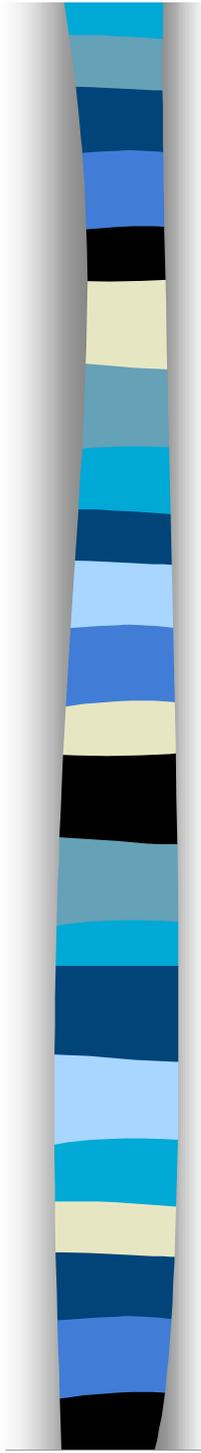


The PageRank algorithm

- This is the equation of a Markov model of random surf through the web
- This is exactly the same equation as before:

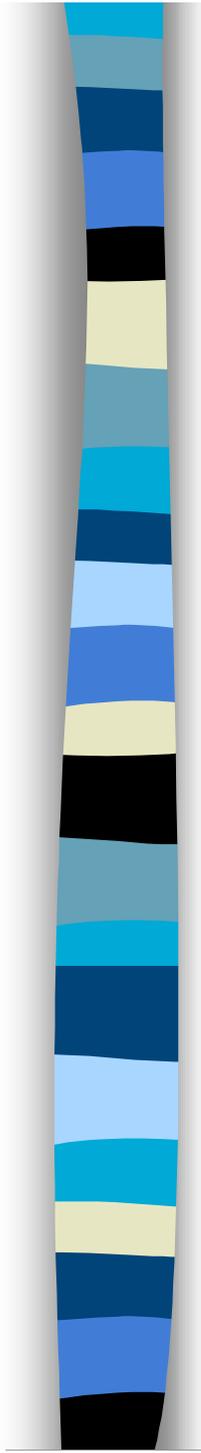
$$x_i \propto \sum_{j=1}^n \frac{w_{ji} x_j}{w_{j.}}$$

$$w_{j.} = \sum_{i=1}^n w_{ji}$$



The PageRank algorithm

- x_i can then be viewed as the probability of being at page i
 - One can show that the solution to these equations is the **stationary distribution** of the random surf
 - Which is the probability of finding the surfer on page i on the **long-term behaviour**
- The most probable page is the best ranked

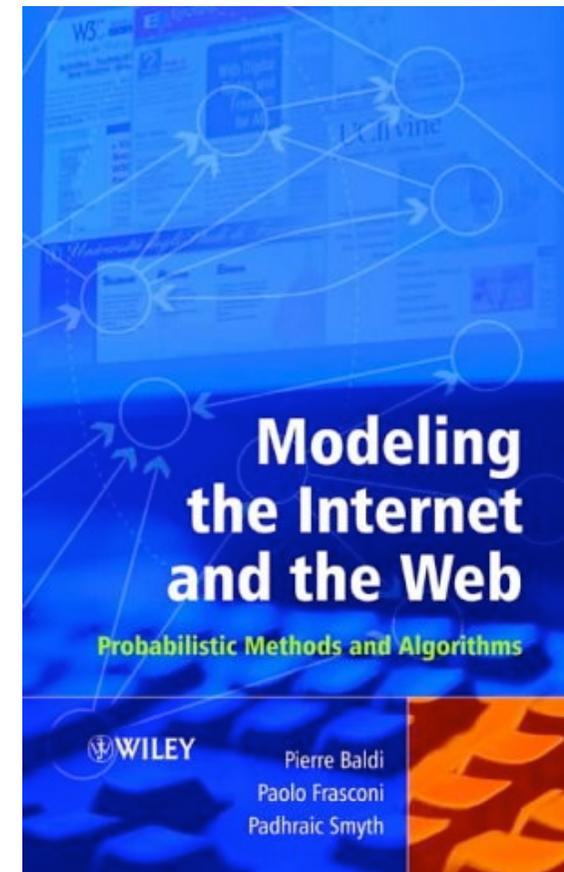


The PageRank algorithm

- One can show that the scores can also be obtained
 - By computing the principal **left eigenvector** of the matrix \mathbf{P} ,
 - The **probability transition matrix** of the Markov process
 - Containing the transition probabilities

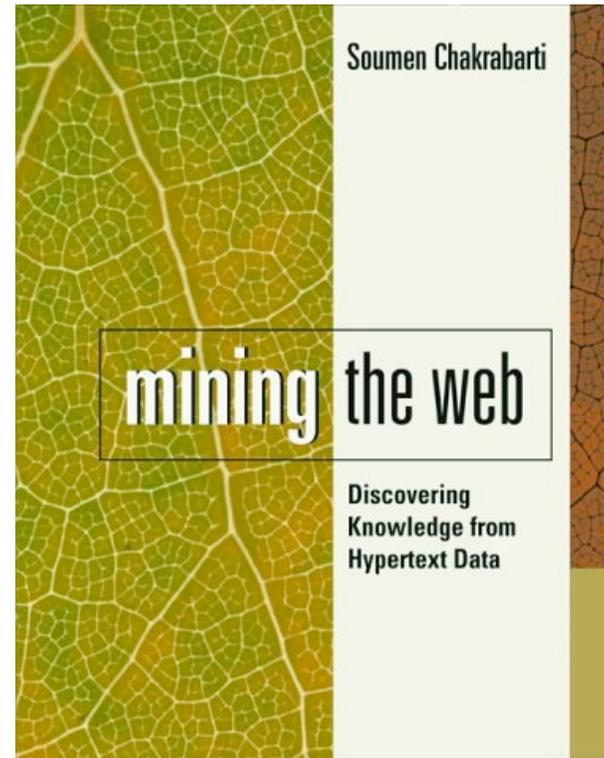
(2) The HITS algorithm

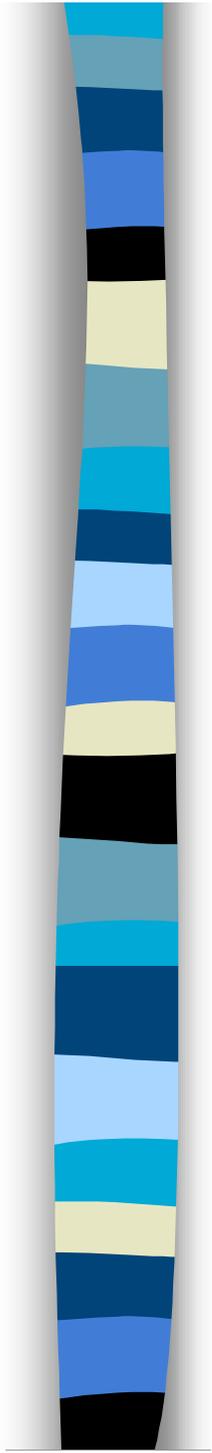
- P. Baldi, P. Frasconi & P. Smyth (2003)
 - Modeling the Internet and the Web
 - John Wiley & Sons



The HITS algorithm

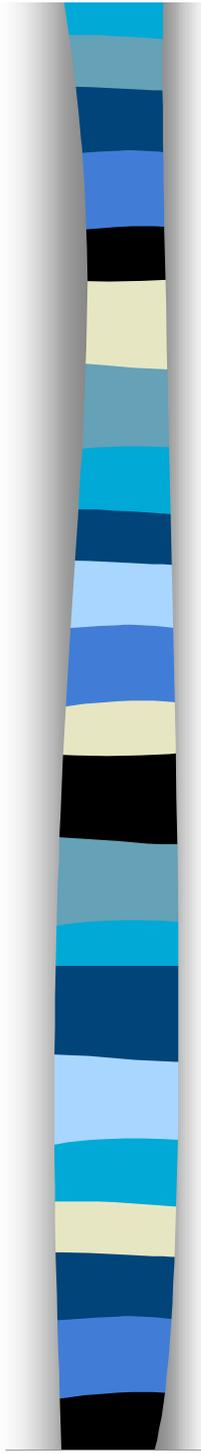
- S. Chakrabarti (2003)
 - Mining the web
 - Morgan Kaufmann





The HITS algorithm

- Introduced by Kleinberg in 1998/1999



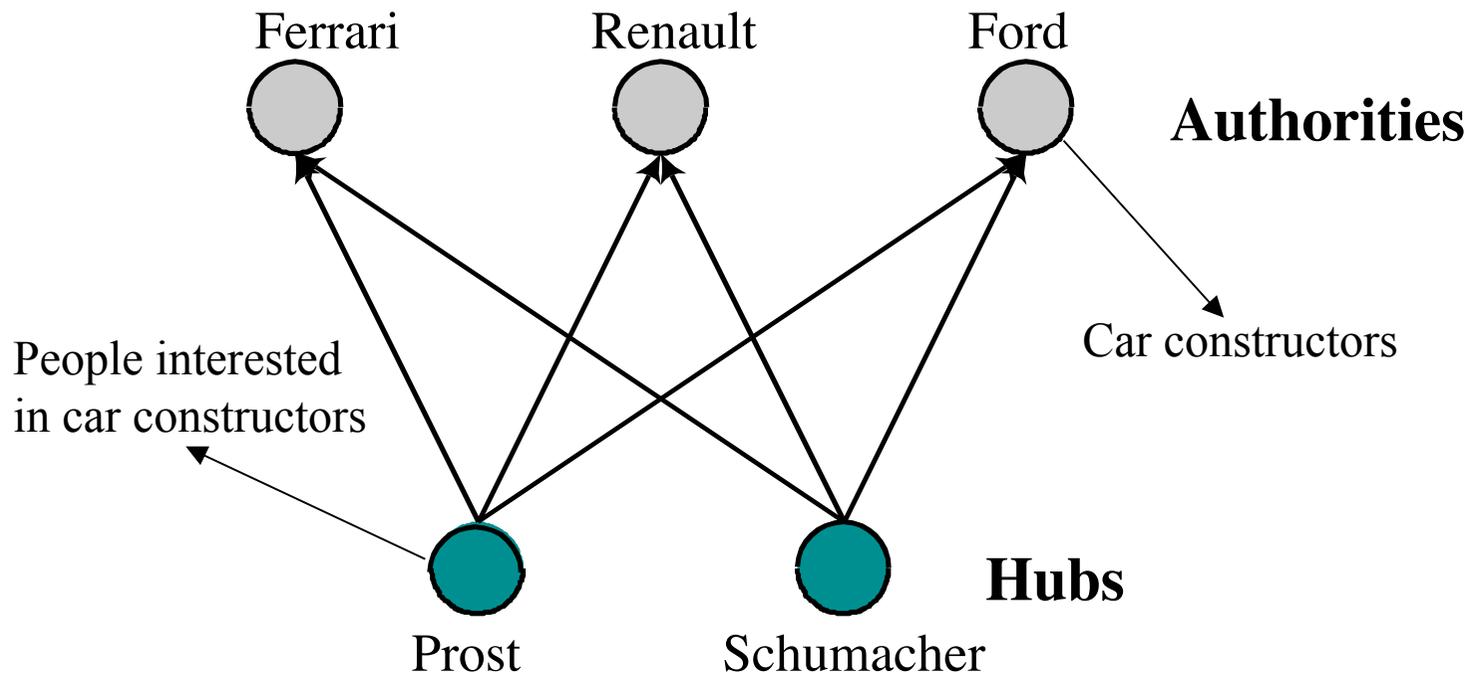
The HITS algorithm

- The model proposed by Kleinberg is based on two concepts
 - **Hub** pages
 - **Authorities** pages
- We thus assume that these are two categories of web pages
- These two concepts are strongly connected

The HITS algorithm

- Example:

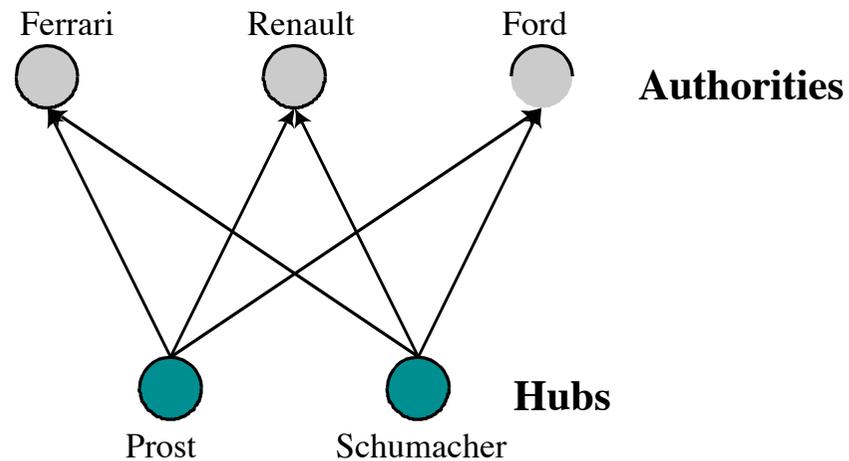
- Suppose we introduced the query “Car constructors”



The HITS algorithm

■ Hubs

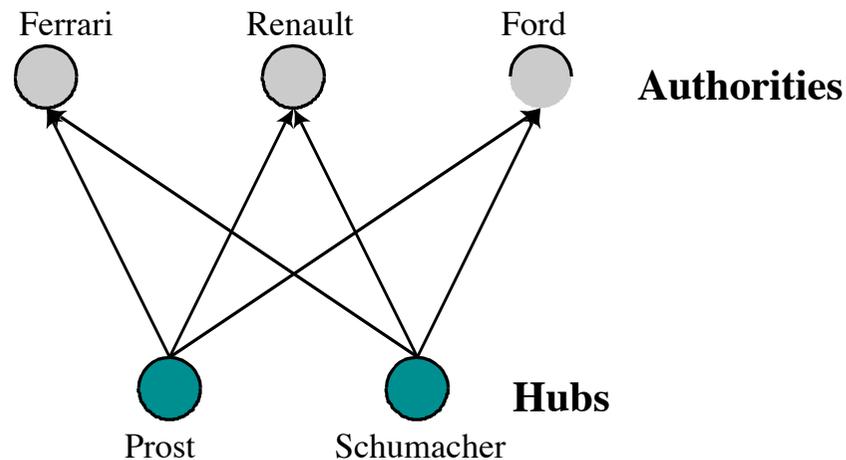
- Link heavily to authorities
- A good hub points to many good authorities
- Hubs have very few incoming links

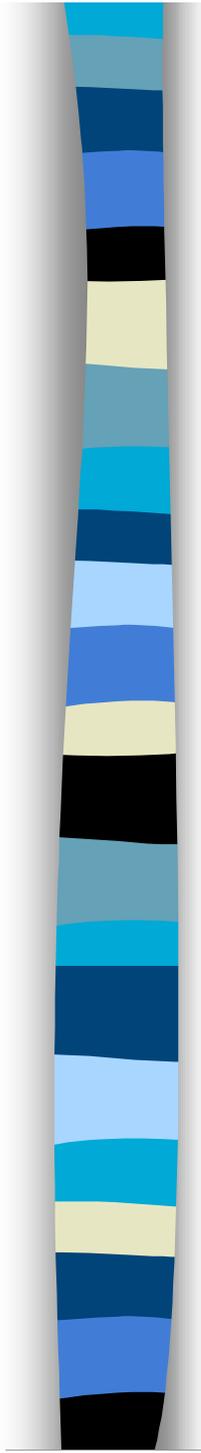


The HITS algorithm

■ Authorities

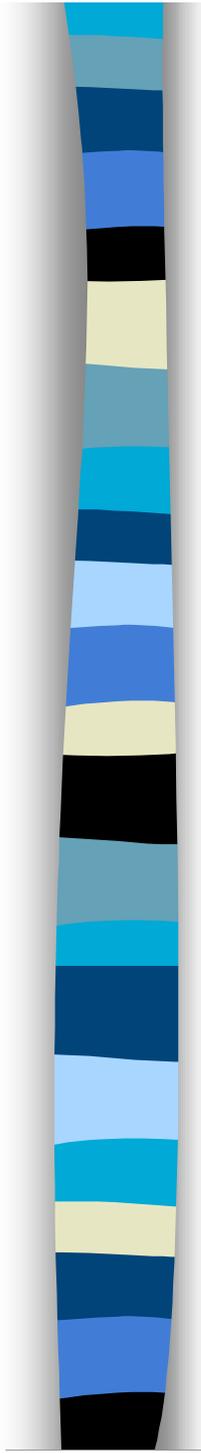
- Do not link to other authorities
- A good authority is pointed to by many good hubs
- The main authorities on a topic are often in competition with one another





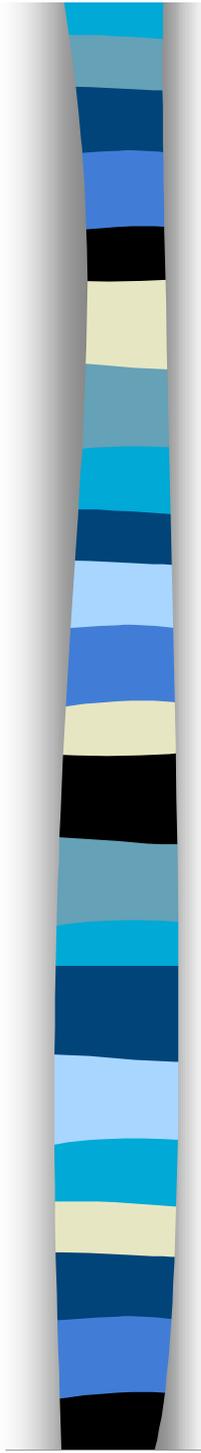
The HITS algorithm

- The objective is to detect **good hubs** and **good authorities**
 - from the results of the search engine
- We therefore assign two numbers to each returned page i :
 - A **hub score**, x^h_i
 - An **authority score**, x^a_i



The HITS algorithm

- Let w_{ij} be the weight of the link connecting page i to page j
 - Usually, it is simply 0 or 1
 - Thus, $w_{ij} = 1$ if page i has a link to page j ; $w_{ij} = 0$ otherwise
- Let \mathbf{W} be the matrix made of elements w_{ij}
 - Notice that this matrix is not symmetric
 - We suppose that the graph is strongly connected



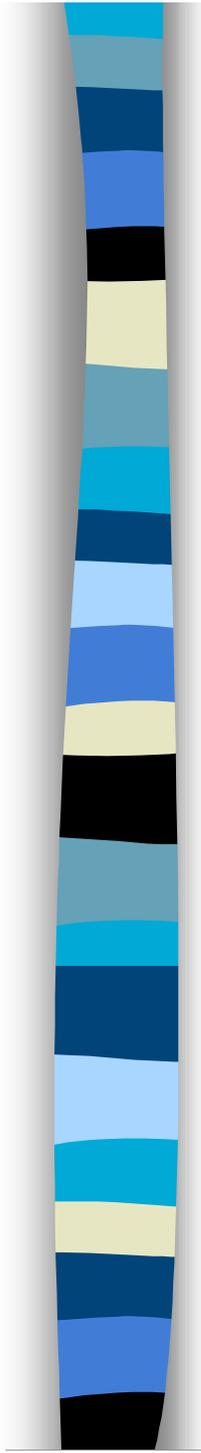
The HITS algorithm

- A possible procedure for computing hub/authorities scores (Kleinberg)
 - A page's **authority score** is proportional to the sum of the hub scores that link to it

$$x_j^a = \eta \sum_{i=1}^n w_{ij} x_i^h$$

- A page's **hub score** is proportional to the sum of the authority scores that it links to

$$x_i^h = \mu \sum_{j=1}^n w_{ij} x_j^a$$



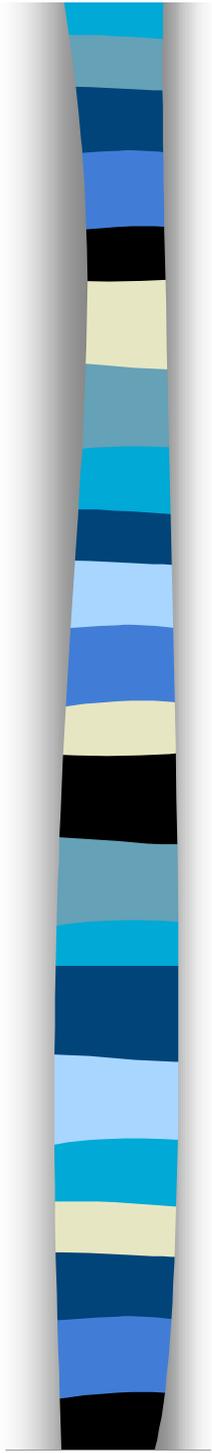
The HITS algorithm

- Kleinberg used this iterative procedure in order to estimate the scores (with a normalization)
 - He showed that this is equivalent to computing the eigenvectors of the following matrices

$$WW^T$$

$$W^TW$$

- To obtain respectively the vector of hubs scores and the vector of authorities scores
- We showed that this is exactly **uncentered principal components analysis (PCA)**

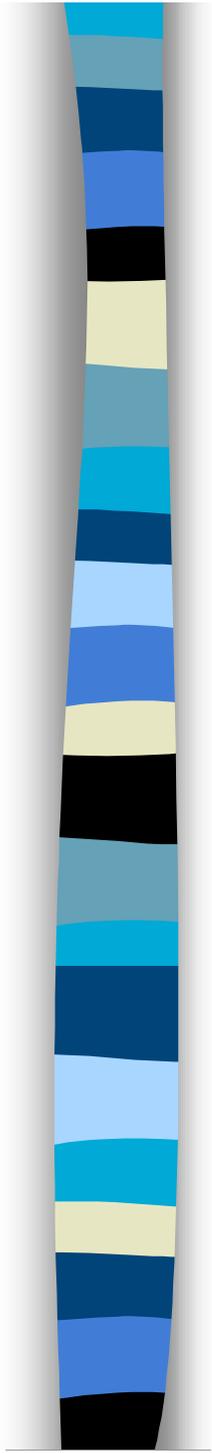


The HITS algorithm

- We further showed that this procedure is also related to both
 - Correspondence analysis
 - A random walk model through the graph

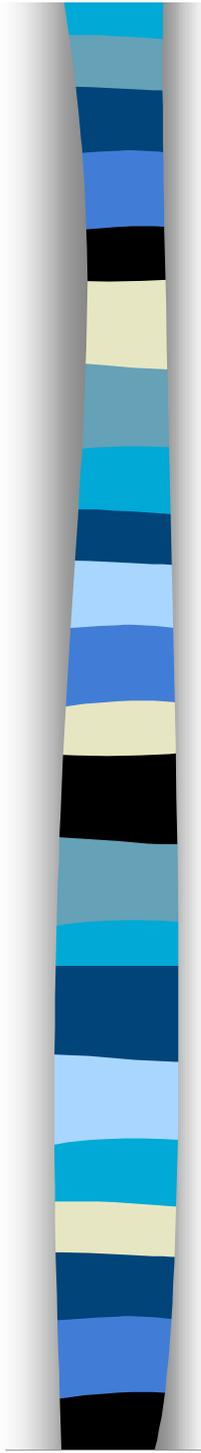
The multivariate analysis of undirected graphs





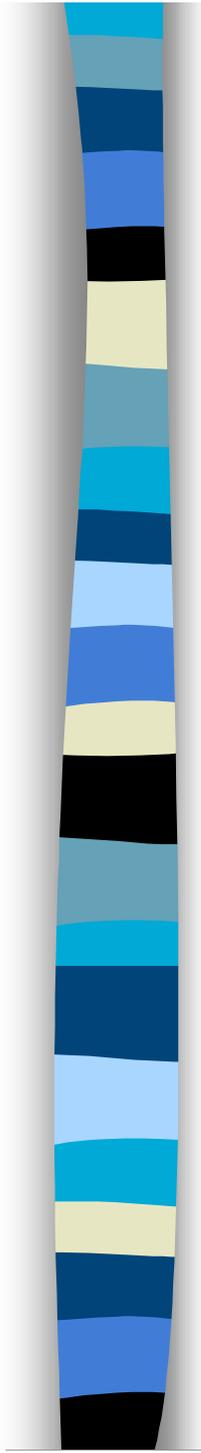
Context

- Main purpose: To exploit the **graph structure** of large repositories
 - Web environment
 - Digital documents repositories
 - Databases with metadata
- We will focus on **databases** and **collaborative recommendation**



Main goal

- To exploit and analyse
 - New **similarity measures** between the nodes of a graph
- To use these similarities for
 - Collaborative filtering
 - Clustering
 - Graph visualization
 - Etc...



Main point

- These similarity measures between two nodes not only depend on
 - The **weights** of the edges (like the « shortest path » distance)
- But also on
 - The **number of paths** connecting the two edges
- They take **high connectivity** into account
≠ shortest-path or geodesic (Dijkstra) distance

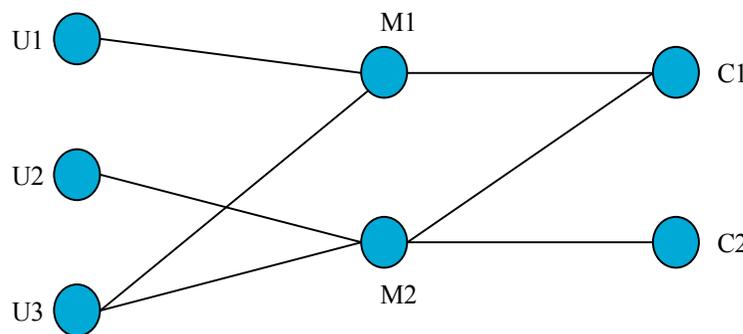
Graph: The adjacency matrix

- The elements a_{ij} of the adjacency matrix \mathbf{A} of a weighted, undirected, graph are defined as

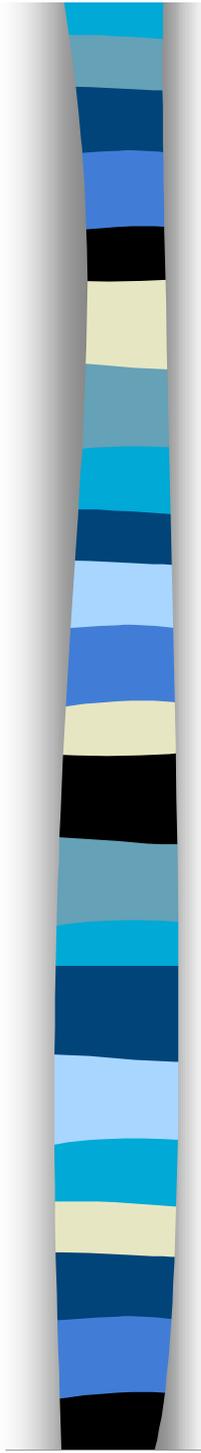
$$a_{ij} = \begin{cases} w_{ij} & \text{if node } i \text{ is connected to node } j \\ 0 & \text{otherwise} \end{cases}$$

where \mathbf{A} is symmetric

- The $w_{ij} \geq 0$ represent the strength of relationship between node i and node j



$$\mathbf{A} = \begin{pmatrix} 0 & 0 & 0 & 1 & 0 & 0 & 0 \\ 0 & 0 & 0 & 0 & 1 & 0 & 0 \\ 0 & 0 & 0 & 1 & 1 & 0 & 0 \\ 1 & 0 & 1 & 0 & 0 & 1 & 0 \\ 0 & 1 & 1 & 0 & 0 & 1 & 1 \\ 0 & 0 & 0 & 1 & 1 & 0 & 0 \\ 0 & 0 & 0 & 0 & 1 & 0 & 0 \end{pmatrix}$$



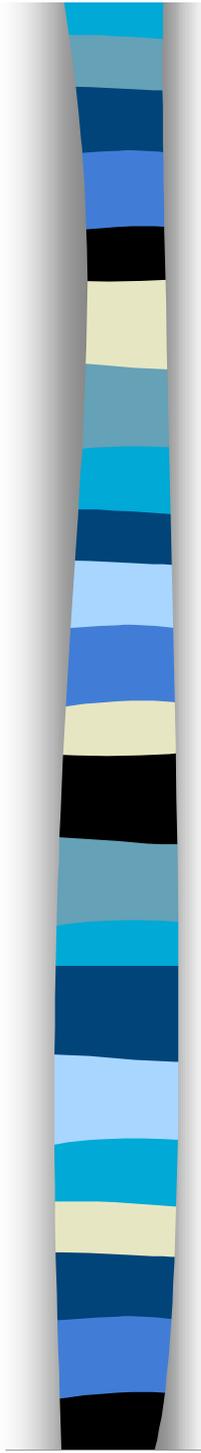
Graph: The Laplacian matrix

- The Laplacian matrix \mathbf{L} of the graph is defined by

$$\mathbf{L} = \mathbf{D} - \mathbf{A}$$

where $\mathbf{D} = \text{diag}(a_i.)$ with $d_{ii} = [\mathbf{D}]_{ii} = a_i. = \sum_{j=1}^n a_{ij}$
(the outdegree of each node)

- \mathbf{L} is doubly centered
- If the graph is connected, the rank of \mathbf{L} is $n - 1$, where n is the number of nodes
- \mathbf{L} is symmetric
- \mathbf{L} is positive semidefinite



A random walk model on the graph

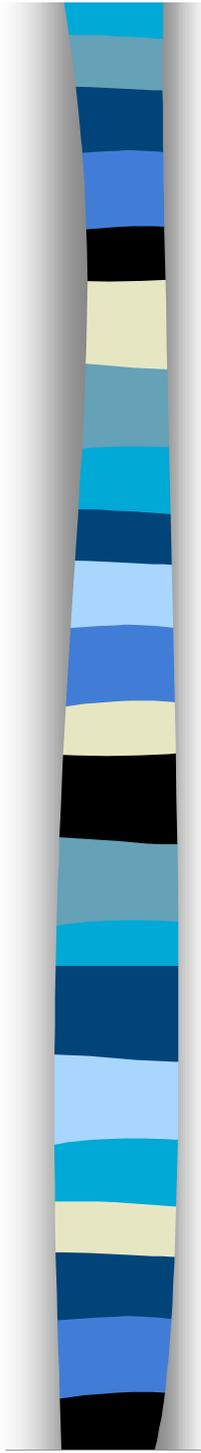
- As for PageRank, every node is associated to a **state** of a **Markov chain**
- The random walk is defined by the single-step **transition probabilities**

$$P(s(t+1) = j | s(t) = i) = p_{ij} = \frac{a_{ij}}{a_{i.}}$$

where

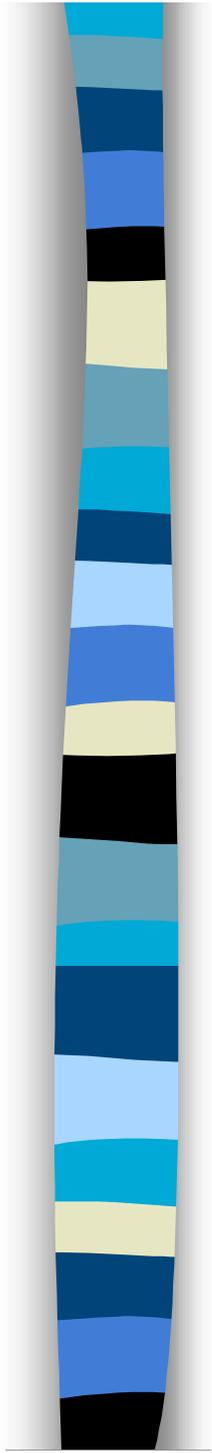
$$a_{i.} = \sum_{j=1}^n a_{ij}$$

- In other words, to any state or node i , we associate a probability of jumping to an adjacent node, $s(t+1) = j$
 - which is **proportional to** the weight w_{ij} of the edge connecting i and j



Two main quantities

- We then compute two main quantities from this Markov chain:
 - The *average first passage time*
 - The *average commute time*

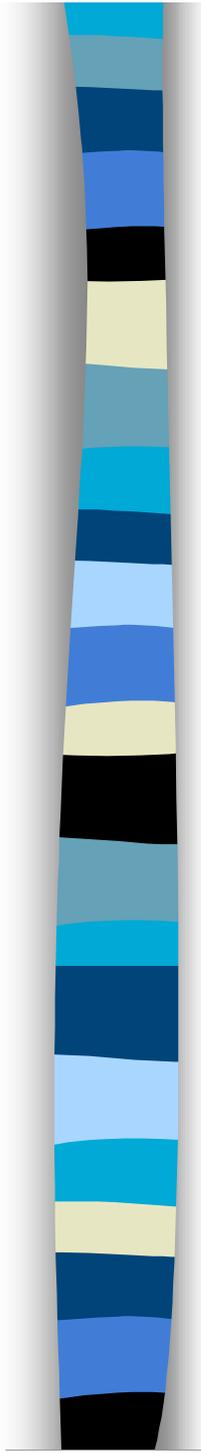


Average first-passage time

- $m(k|i)$ = average number of steps a random walker, starting in state i , will take to enter state k for the first time

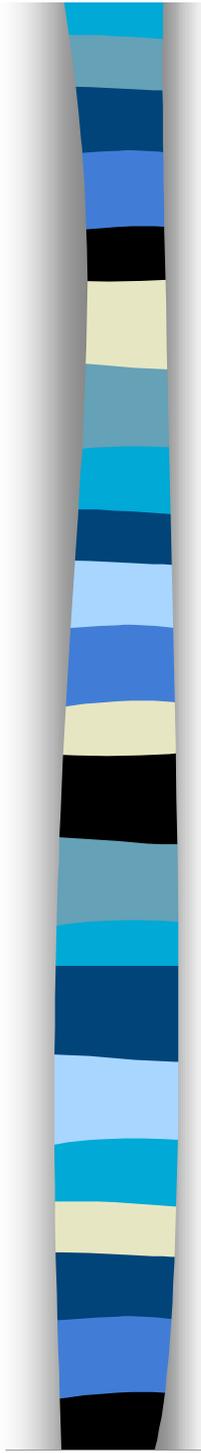
$$\begin{cases} m(k|i) = 1 + \sum_{\substack{j=1 \\ j \neq k}}^n p_{ij} m(k|j), \text{ for } i \neq k \\ m(k|k) = 0 \end{cases}$$

- These equations can be used in order to iteratively compute the first-passage times.



Average commute time

- $n(i,j) = m(j|i) + m(i|j)$
 - = average number of steps a random walker, starting in state $i \neq j$, will take before entering a given state j for the first time, and go back to i
- Note: while $n(i,j)$ is symmetric by definition, $m(i|j)$ is not.



Computation of the basic quantities by means of \mathbf{L}^+

- If we further define \mathbf{e}_i as the i th column of \mathbf{I}

$$\mathbf{e}_i = [0, \dots, 0, 1, 0, \dots, 0]^T$$

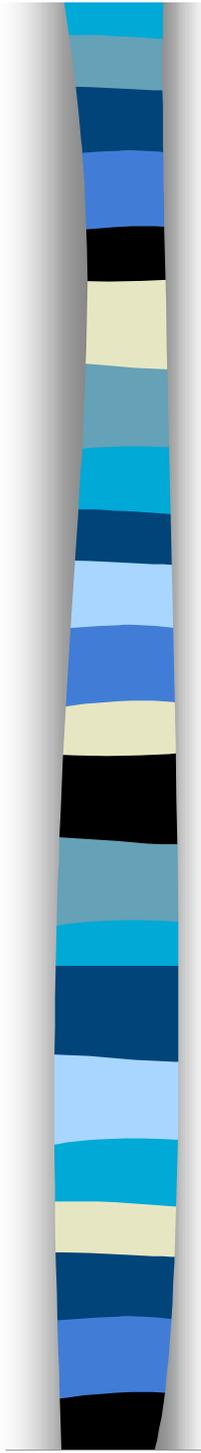
1 $i-1$ i $i+1$ n

- we obtain the remarkable form

$$n(i, j) = 2N_e(\mathbf{e}_i - \mathbf{e}_j)^T \mathbf{L}^+ (\mathbf{e}_i - \mathbf{e}_j)$$

where each node i is represented by a unit basis vector, \mathbf{e}_i , in the node space

- \mathbf{L}^+ is the Moore-Penrose pseudoinverse of the Laplacian matrix of the graph

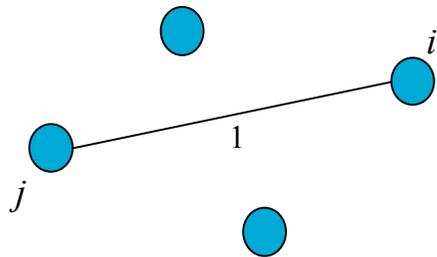


Computation of the basic quantities by means of \mathbf{L}^+

- Thus, $n(i,j)$ is a Mahalanobis distance
= Commute Time Distance
- Indeed, one can show that \mathbf{L}^+ is
 - (1) Symmetric
 - (2) Positive semidefinite
 - (3) Doubly centered

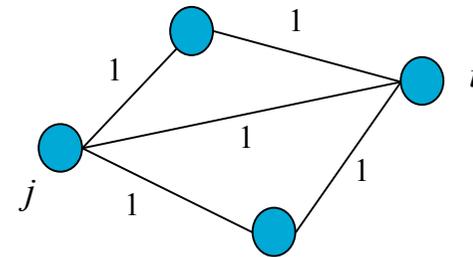
Commute time distance

- It takes high connectivity into account:



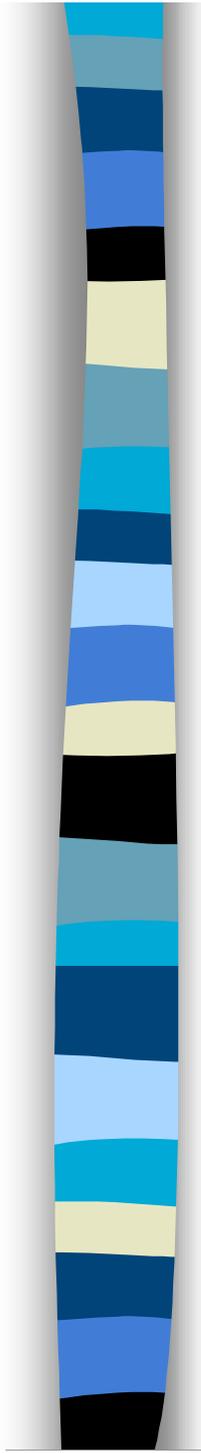
$$n(i,j) \text{ distance} = Cst \cdot 1.0$$

$$\text{Shortest_path} = 1$$



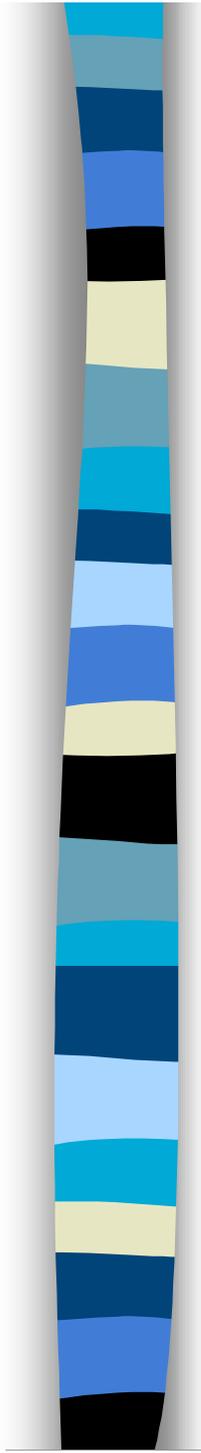
$$n(i,j) \text{ distance} = Cst \cdot 0.5$$

$$\text{Shortest_path} = 1$$



Embedding in an Euclidean space

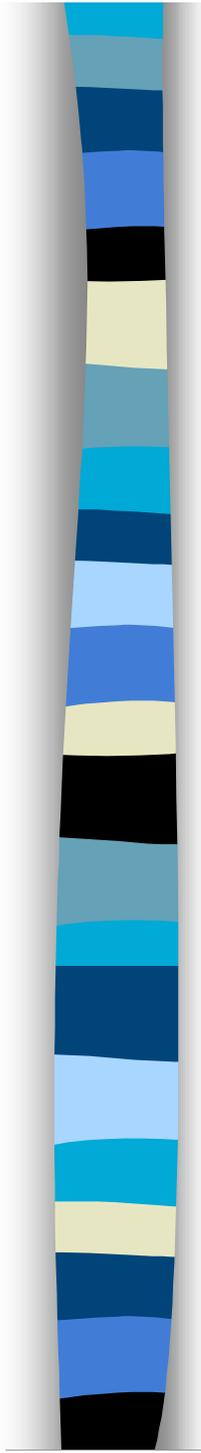
- The node vectors can be mapped into an Euclidean space preserving the commute time distance
 - In this space, the node vectors are exactly separated by commute time distances
- The node vectors form a cloud of points, each point being a node
- So that any multivariate statistical analysis tool can be applied to analyse the graph



Embedding in an Euclidean space

■ For instance:

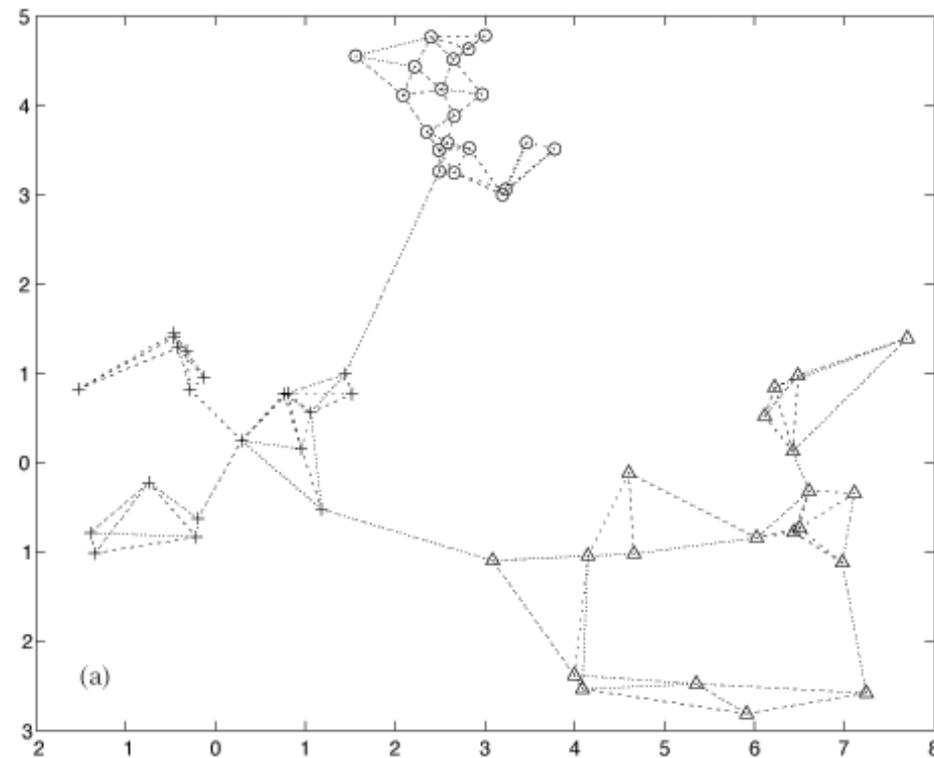
- Clustering of the nodes
- Finding dense regions in the graph
- Finding outlier nodes
- Representing the graph in a low-dimensional space (principal components analysis)
- Representing the graph in function of the similarity with some reference nodes (discriminant analysis)
- Finding central nodes in the graph
- Find the most similar node (nearest neighbour)
- Etc...



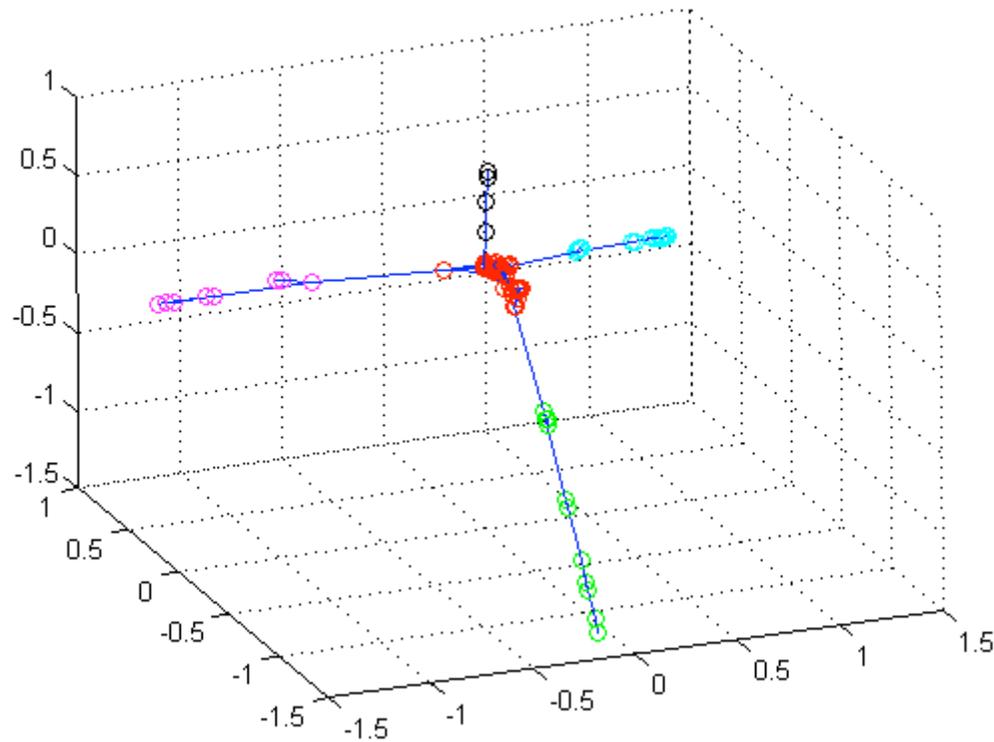
Maximum variance subspace projection of the nodes vectors

- This decomposition is similar to **principal component analysis (PCA)**
 - The projected node vectors has maximal variance
 - In terms of Euclidean commute time distance
 - Among all the possible candidate projections
- It allows us to **visualize** the graph

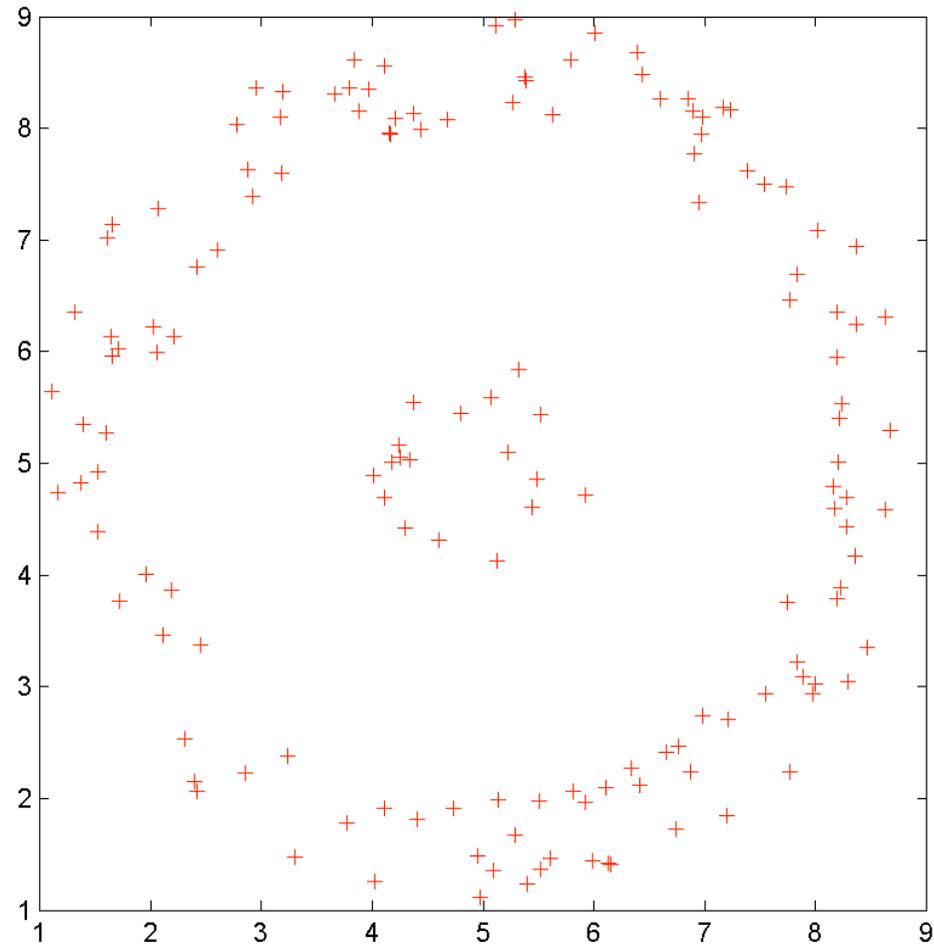
An example of PCA: Original graph



PCA: Application to the visualization of a network of criminals

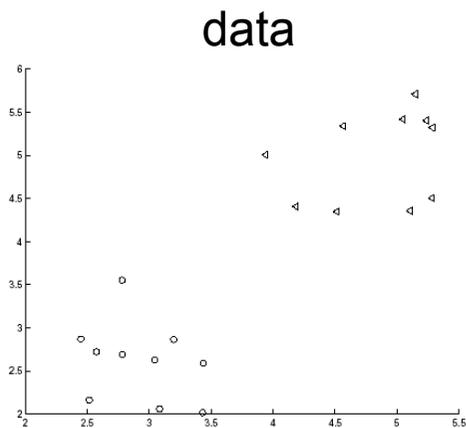


Another example : Application to clustering



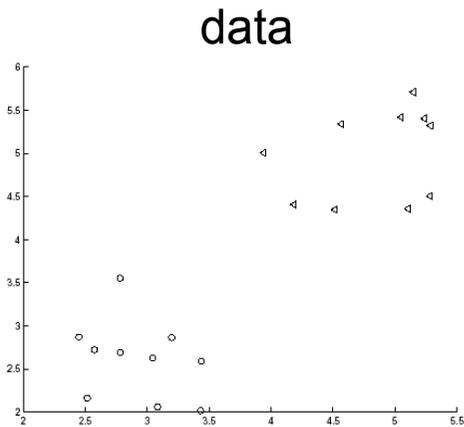
Clustering with ECT distance

Graph construction :

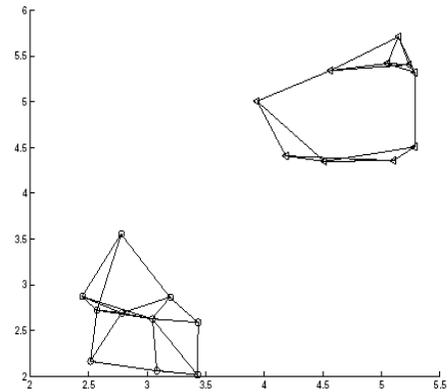


Clustering with ECT distance

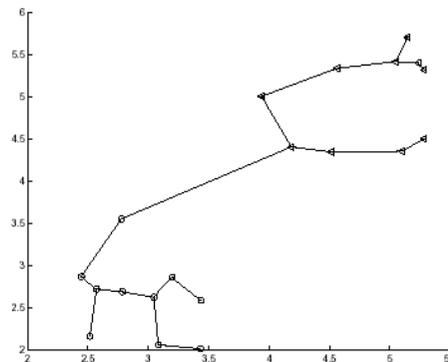
Graph construction :



3 nearest neighbors

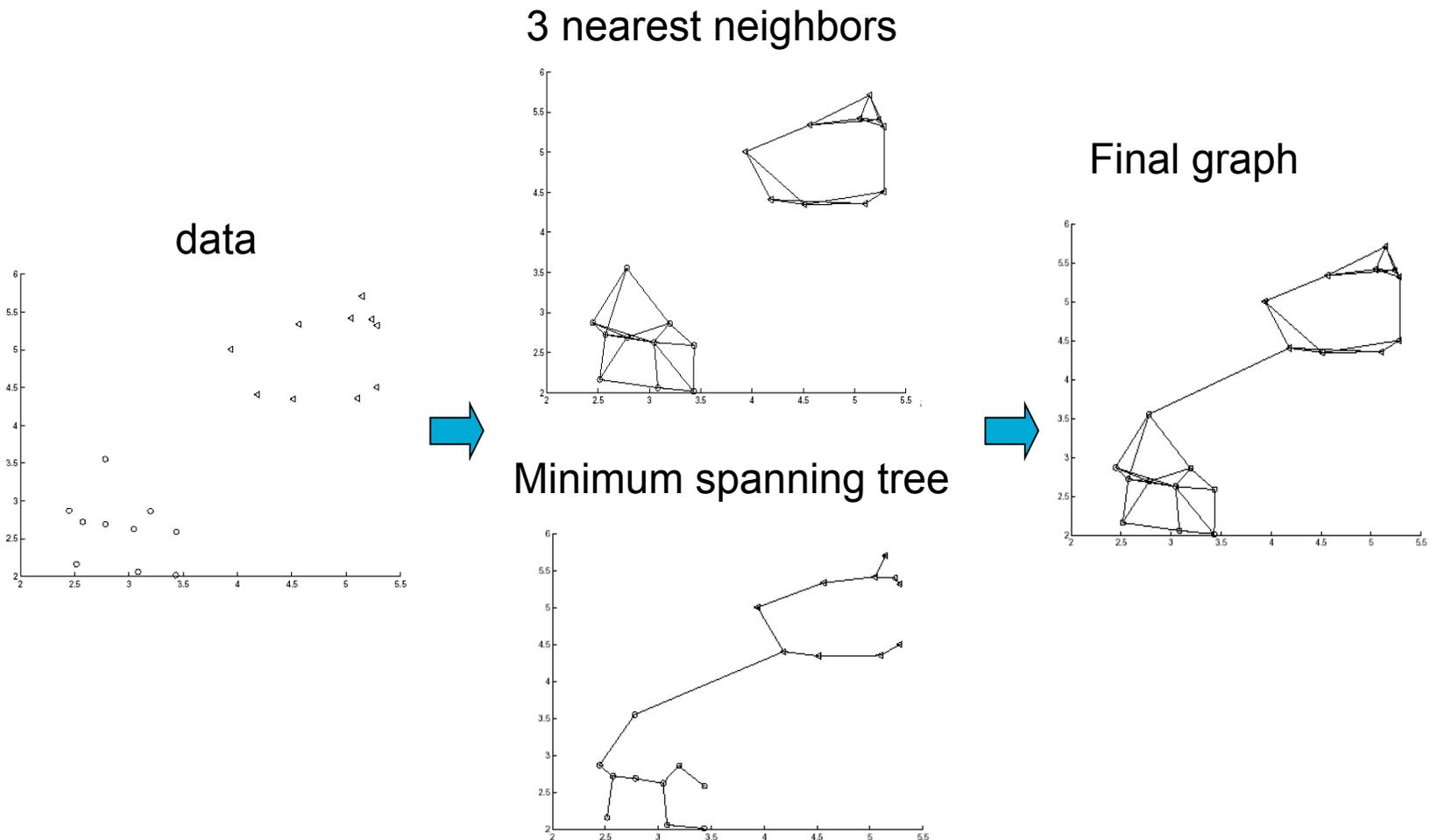


Minimum spanning tree

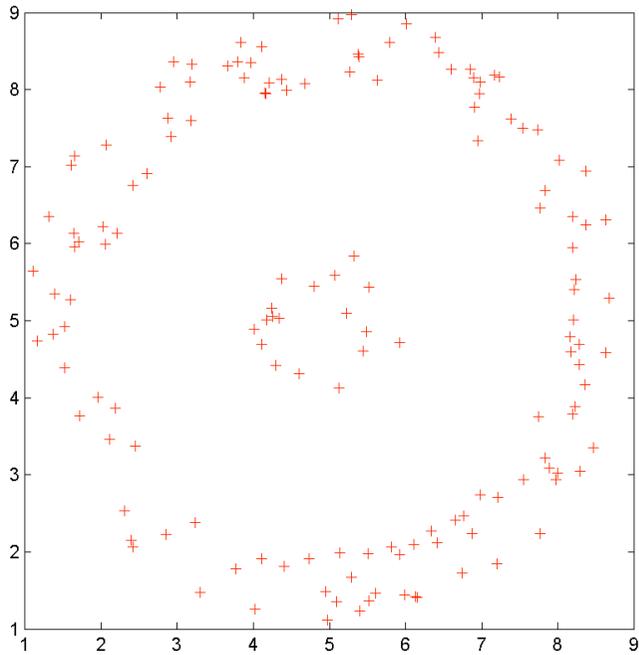


Clustering with ECT distance

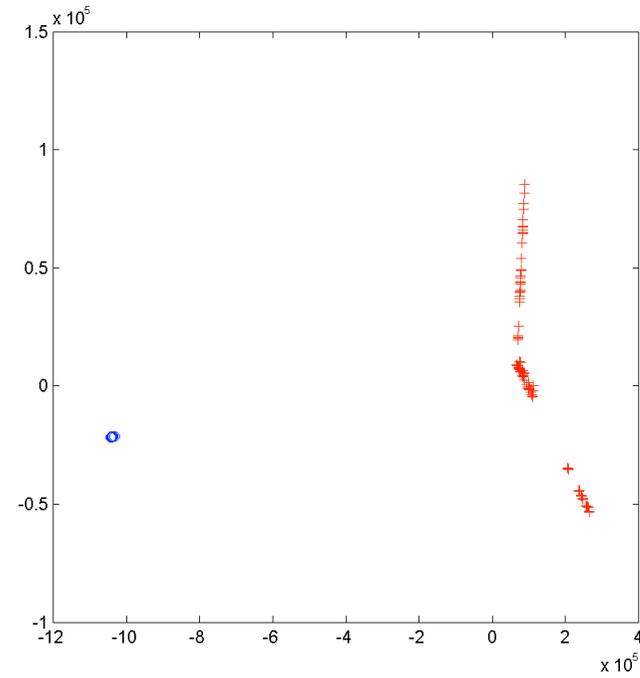
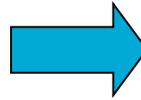
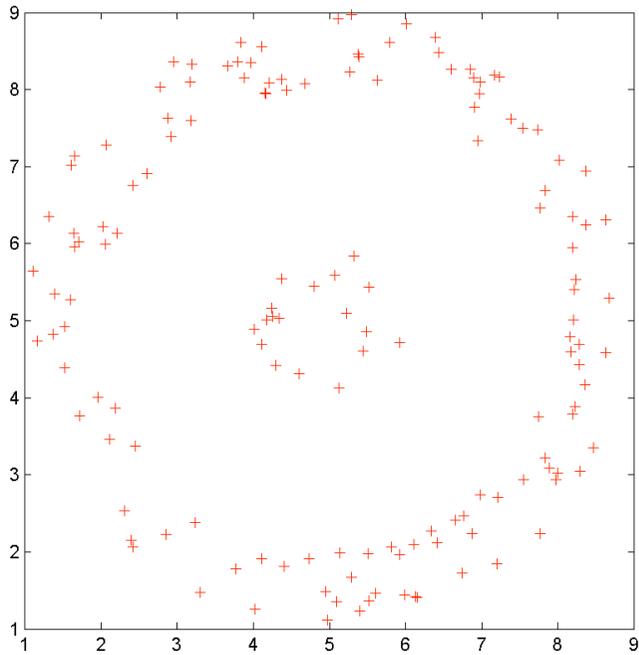
Graph construction :



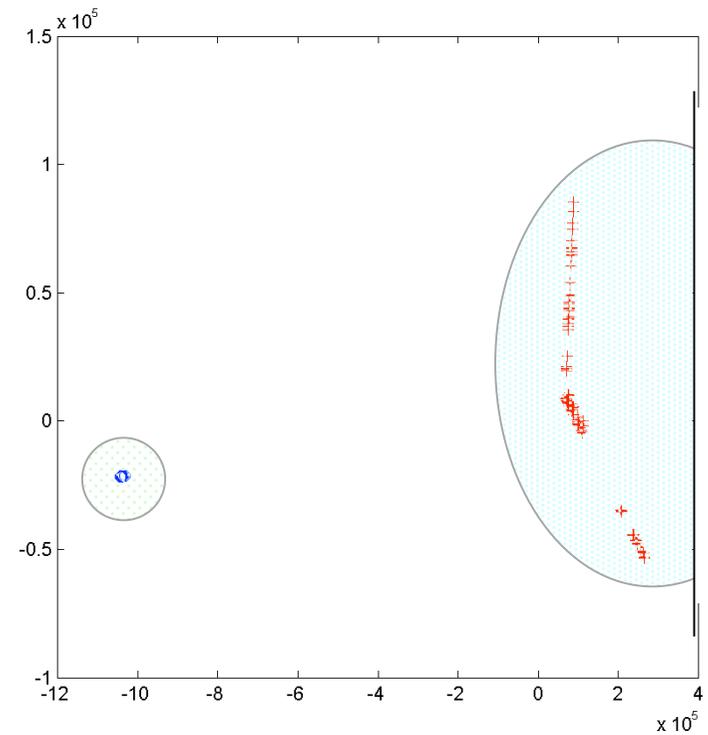
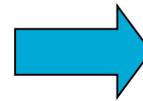
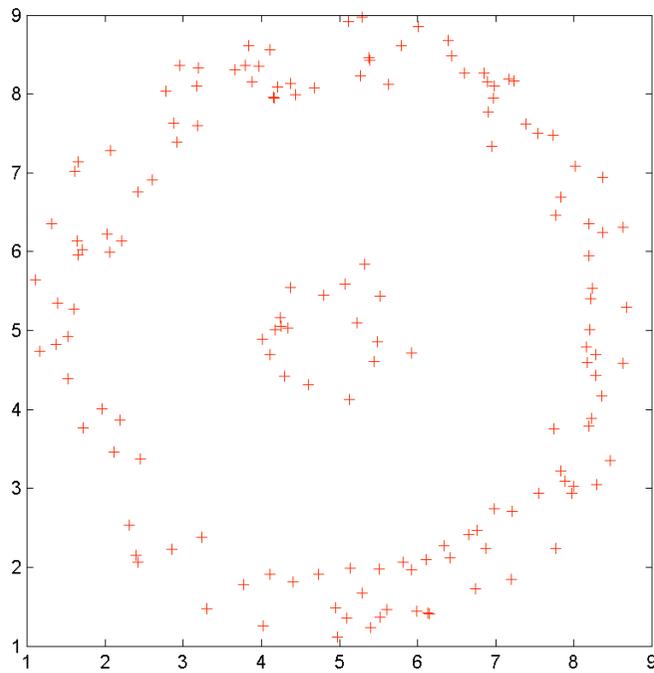
Clustering with ECT distance



Clustering with ECT distance

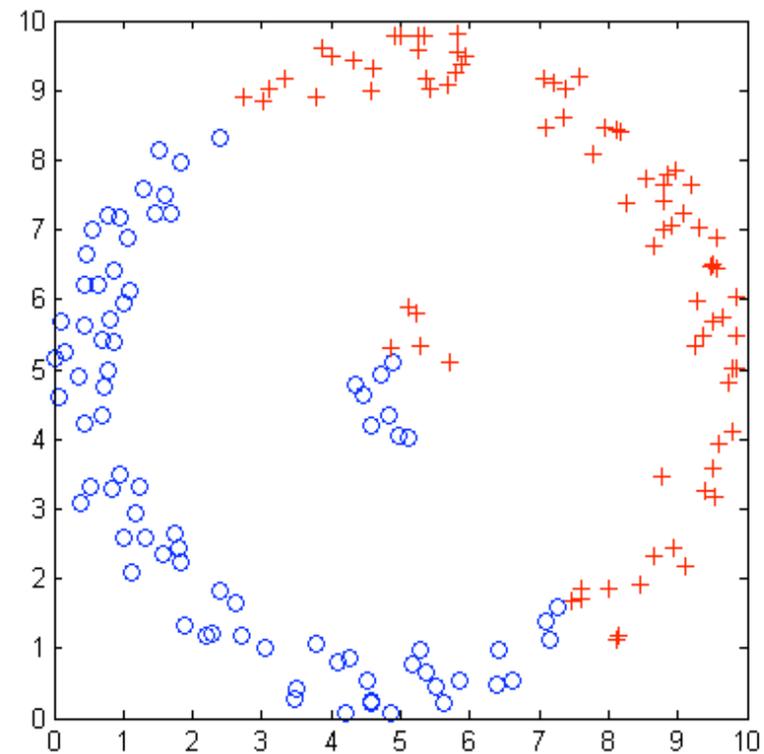
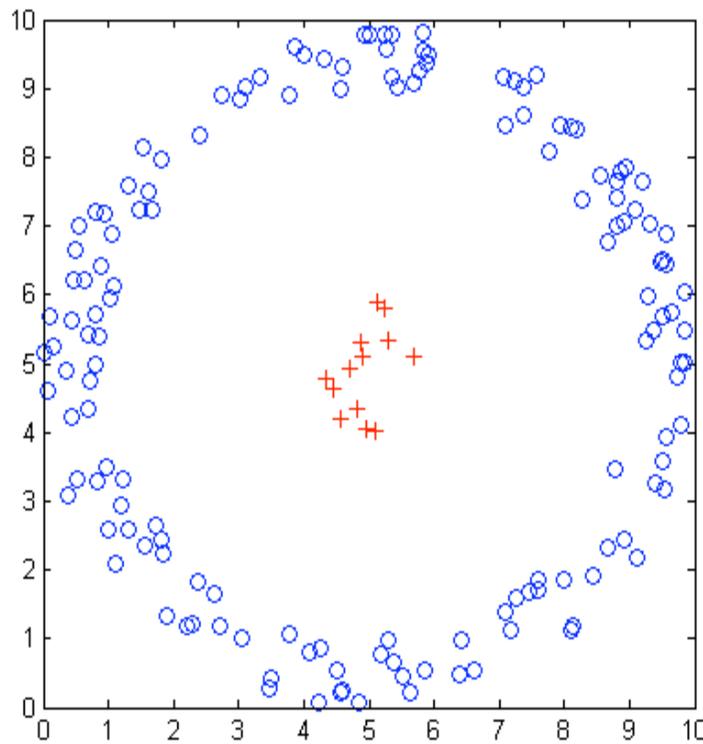


Clustering with ECT distance

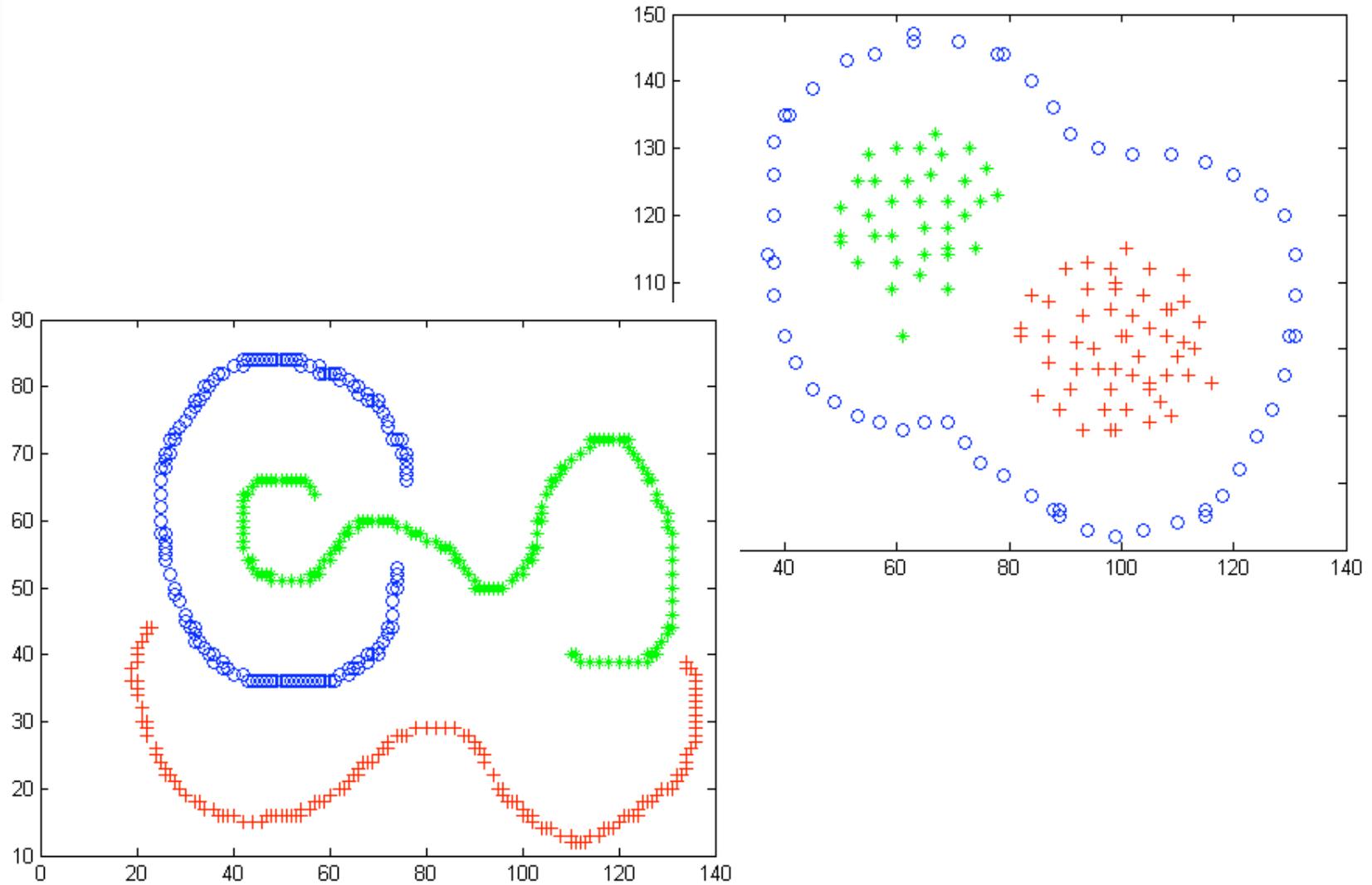


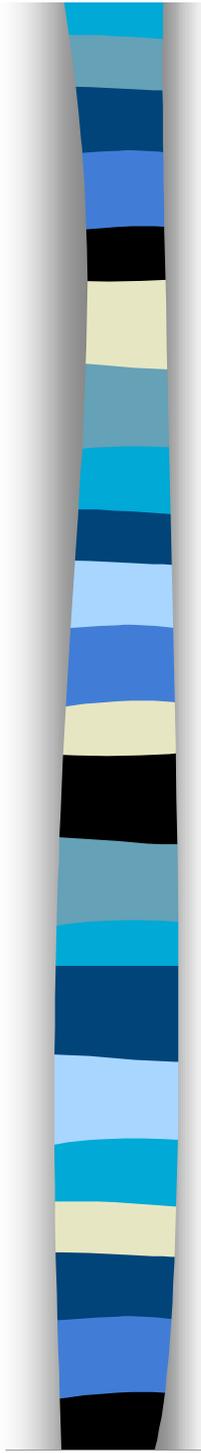
Clustering with ECT distance

- Clustering results using ECT distance k-means, in comparison with the classical k-means



Autres exemples



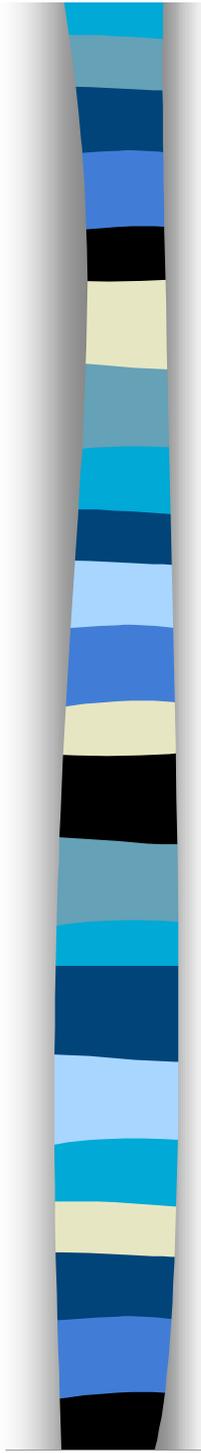


Links with other methods

- Very interesting links with
 - Spectral clustering
 - Graph visualization algorithms
 - Electrical networks
 - The commute time distance is equivalent to the effective resistance
 - Experimental design

Application to collaborative recommendation





Experimental results: Application to collaborative recommendation

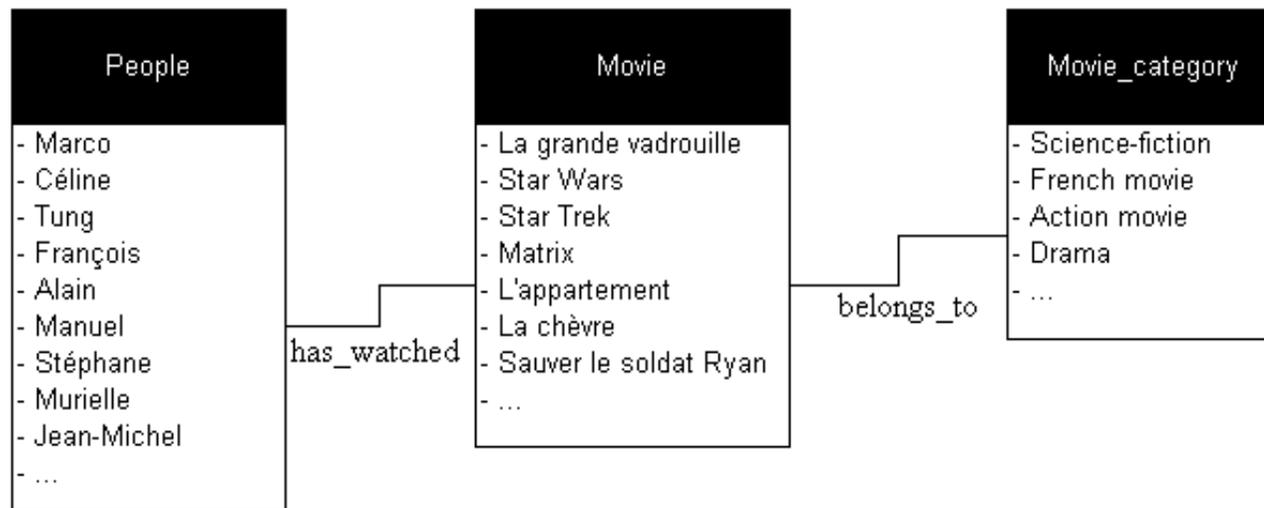
- Results on a real movie database: a sample of the **MovieLens** database
- 943 users
- 1682 movies
- 19 movie categories
- 100,000 ratings
- Divided into a **training set** and a **test set**

- **Experiment: suggest movies to people**

Experimental results: Application to collaborative recommendation

■ Tables connected by relationships

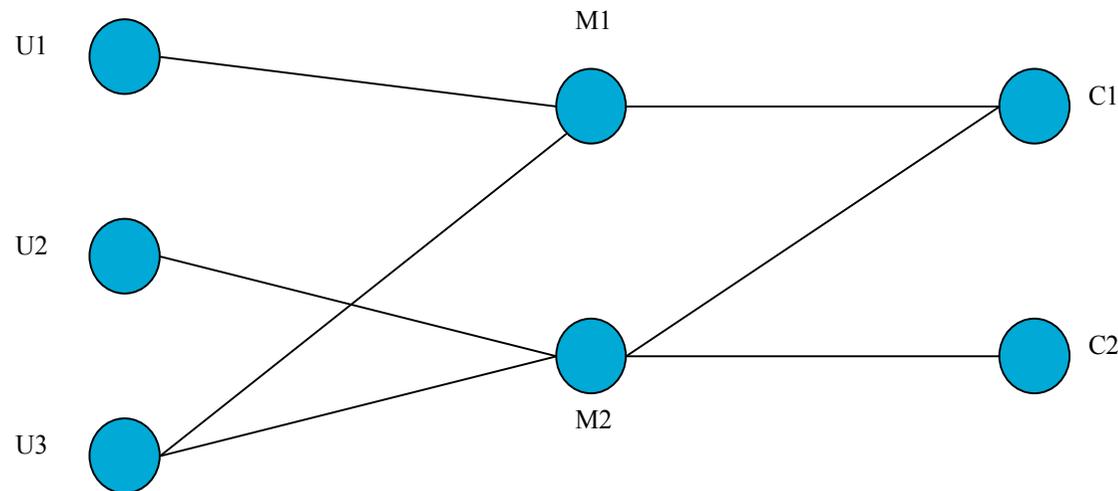
Example: A movie database



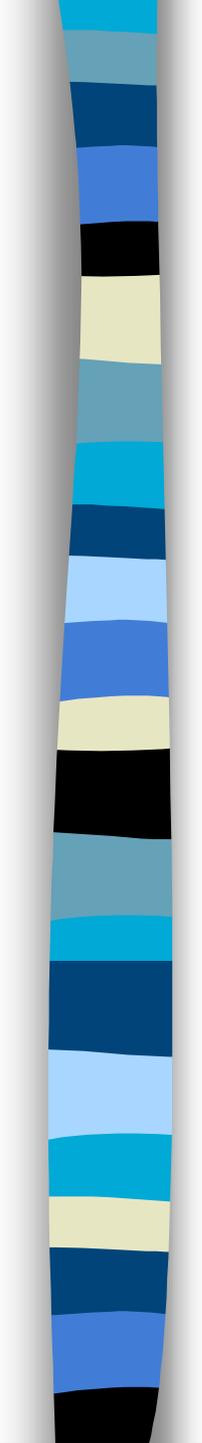
→ Computing similarities between **people** and **movies** allows to suggest movies to watch or not to watch (**collaborative recommendation**)

Experimental results: Application to collaborative recommendation

- Experiment: suggest unwatched movies to people



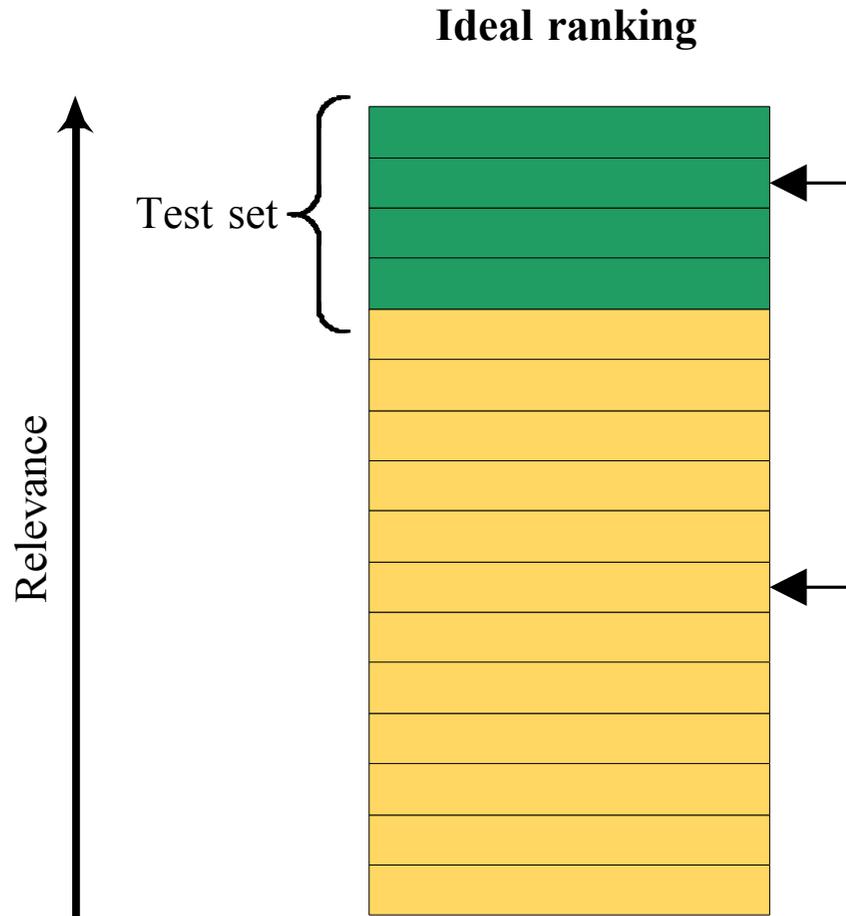
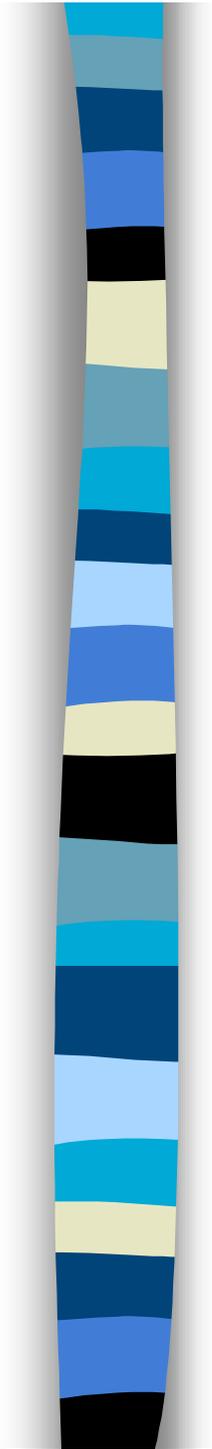
- The test set contains 10 movies for each user
= 9430 movies



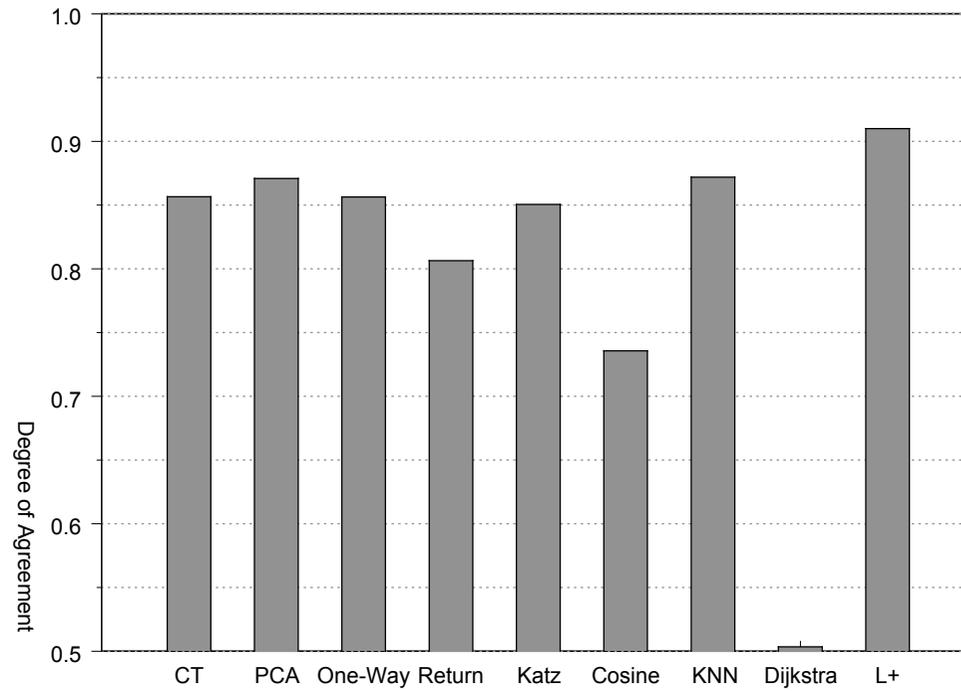
Scoring algorithms

- Average commute time (CT)
- Average first-passage time (One-way)
- Average first-passage time (Return)
- L^+ (pseudoinverse of the Laplacian matrix of the graph)
- Katz
- K-nearest neighbours (KNN) (Standard technique)
- Dijkstra (Standard technique)
- Cosine (Standard technique)

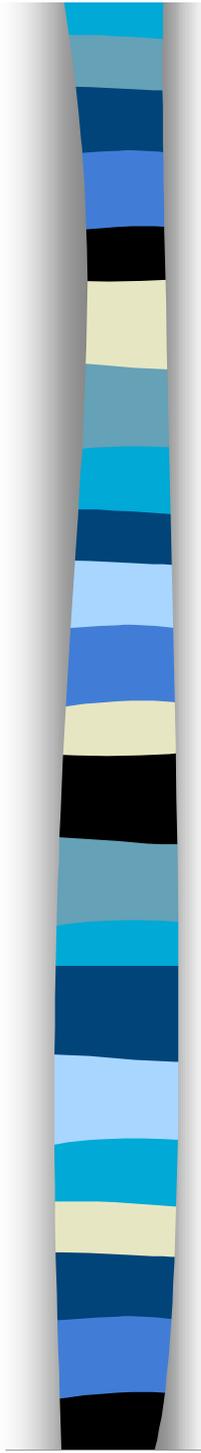
Performance evaluation: degree of agreement (a variant of Somers' D)



Results



CT	PCA CT	One-way	Return	Katz	KNN	Dijkstra	Cosine	\mathbf{L}^+
0.8566	0.8710	0.8564	0.8065	0.8506	0.8720	0.5034	0.7358	0.9101



Conclusion

- We introduced a general procedure for computing **dissimilarities** between any pair of elements
- The commute time is a **distance metric** in an Euclidean space
- This allows the application of **data analysis methods** (PCA, discriminant analysis, clustering) for graph analysis