# Complex Networks matrix algorithms and graph partitioning

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#### contents of this chapter

- eigenvector centrality
- finding the leading eigenvector of the adjacency matrix
- spectral partitioning method
- network community detection

### leading eigenvectors and eigenvector centrality

- eigenvector centrality of vertex i
  - ith element of the leading eigenvector of the adjacency matrix
- calculating the complete set of eigenvectors of the adjacency matrix is a wasteful approach
- power method: $\mathbf{x}(t) = \mathbf{A}^t \mathbf{x}(0)$ 
  - -x(t) will converge to the leading eigenvector as  $t \to \infty$
  - no faster method for calculating eigenvector centrality

#### caveats of power method

- the method will not work if the initial vector x(0)
  happens to be orthogonal to the leading eigenvector
  - one simple way to avoid this problem is to choose the initial vector to have all elements positive
- the elements of the vector have a tendency to grow on each iteration
  - we must periodically renormalize the vector by dividing all the elements by the same value
- how long do we need to go on multiplying by the adjacency matrix before the result converges to the leading eigenvalue?
  - this will depend on how accurate an answer we require.

### computational complexity of power method

- 1. how long each multiplication by the adjacency matrix takes?
- 2. how many multiplications are needed?
- 1.
- adjacency matrix : n<sup>2</sup> multiplication
- adjacency list : O(m)
  - $-\{u_j\}, j=1...k_i$ :set of neighbors of vertex i
  - $-[\mathbf{A}\mathbf{x}]_i$ : ith element of  $\mathbf{A}\mathbf{x}$   $[\mathbf{A}\mathbf{x}]_i = \sum_{j=1}^{k_i} x_{u_j}$
  - all elements can be completed in time proportional to  $\sum_{i} k_{i} = 2m$

k<sub>i</sub> operations

#### how many multiplications must we perform?

normalized ith

$$\mathbf{x}(0) = \sum_{i} c_{i} \mathbf{v}_{i}$$

$$\mathbf{x}(t) = \mathbf{A}^{t} \sum_{i} c_{i} \mathbf{v}_{i} = \sum_{i} c_{i} \kappa_{i}^{t} \mathbf{v}_{i} = \kappa_{1}^{t} \sum_{i} c_{i} \begin{bmatrix} \kappa_{i} \\ \kappa_{1} \end{bmatrix}^{t} \mathbf{v}_{i}$$
corresponding eigenvalue

$$\frac{\mathbf{x}(t)}{c_1 \kappa_1^t} = \mathbf{v}_1 + \frac{c_2}{c_1} \left(\frac{\kappa_2}{\kappa_1}\right)^t \mathbf{v}_2 + \dots,$$

$$\sqrt{\frac{\mathbf{x}(t)}{c_1 \kappa_1^t} - \mathbf{v}_1} = \frac{c_2}{c_1} \left(\frac{\kappa_2}{\kappa_1}\right)^t$$

• if we want this error to be at most ε n-1 spaces

$$t \geq \frac{\ln(1/\varepsilon) + \ln(c_1/c_2)}{\ln(\kappa_1/\kappa_2)} \qquad \begin{array}{c} \text{maximum} \\ \text{minimum} \end{array}$$

$$t \geq \frac{\ln(1/\varepsilon) + \ln(c_1/c_2)}{\ln(\kappa_1/\kappa_2)}$$
 maximum eigenvalue  $\kappa_1$  mean spacing minimum eigenvalue  $\kappa_n \geq -|\kappa_1|$   $\geq \kappa_1/(n-1)$ 

$$\ln \frac{\kappa_1}{\kappa_2} \approx -\ln \left(1 - \frac{a}{n}\right) = \frac{a}{n} + O(n^{-2})$$
  $t = O(n)$ 

$$\kappa_2 \approx \kappa_1 - a \, \kappa_1 / n$$

### computational complexity of power method

- 1. each multiplication : O(m)
   2. number of multiplication : O(n)
  - sparse network  $(m \infty n) : O(mn) \equiv O(n^2)$
  - dense network (m  $\propto$  n<sup>2</sup>): O(mn)  $\equiv$  O(n<sup>3</sup>)
- if adjacency matrix is used, multiplications take O(n²), so the total calculation takes O(n³)

### calculating other eigenvalues and eigenvectors

- power method calculates the largest eigenvalue of a matrix (and the corresponding eigenvector)
- method for finding the smallest eivenvalue
  - shifting all the eigenvalues by a constant amount
- eigenvalues of graph Laplacian L are non-negative
  - $\lambda_{1} \leq \lambda_{2} \leq ... \leq \lambda_{n} \quad \mathbf{v}_{1}, \mathbf{v}_{2}, ..., \mathbf{v}_{n}$   $(\lambda_{n} \mathbf{I} \mathbf{L}) \mathbf{v}_{i} = (\lambda_{n} \lambda_{i}) \mathbf{v}_{i}$   $\mathbf{v}_{i} \text{ is an eivenvector of } (\lambda_{n} \mathbf{I} \mathbf{L}) \mathbf{v}_{i} = (\lambda_{n} \lambda_{i}) \mathbf{v}_{i}$
  - their order is reversed from those of the original largest
  - smallest eigenvalue of L  $\leftarrow$  largest eigenvalue of  $\lambda_n \mathbf{I} \mathbf{L}$

## finding the second-largest eigenvalue (1)

- v<sub>1</sub>: normalized eigenvector corresponding to the largest eigenvalue of a matrix A
- define  $\mathbf{y} = \mathbf{x} (\mathbf{v}_1^T \mathbf{x}) \mathbf{v}_1$
- this vector has the property that

$$\mathbf{v}_{i}^{T}\mathbf{y} = \mathbf{v}_{i}^{T}\mathbf{x} - (\mathbf{v}_{1}^{T}\mathbf{x})(\mathbf{v}_{i}^{T}\mathbf{v}_{1}) = \mathbf{v}_{i}^{T}\mathbf{x} - \mathbf{v}_{1}^{T}\mathbf{x}\delta_{i1} = 0 \quad \text{if i = 1} \\ \mathbf{v}_{i}^{T}\mathbf{x} \quad \text{otherwise}$$

- y is equal to x along the direction of every eigenvector of A except the leading eigenvector
- $\mathbf{y} = \sum_{i=1}^{n} c_i \mathbf{v}_i$  with  $c_i = \mathbf{v}_i^T \mathbf{y}$  has no term in  $\mathbf{v_1}$  since  $c_1 = \mathbf{v}_1^T \mathbf{y} = 0$

$$\therefore \mathbf{y} = \sum_{i=2}^{n} c_i \mathbf{v}_i$$

## finding the second-largest eigenvalue

- use vector y as the starting vector for repeated multiplication by A
- after multiplying y\_by\_A t times

$$\mathbf{y}(t) = \mathbf{A}^{t} \mathbf{y}(0) = \kappa_{2}^{t} \sum_{i=2}^{n} c_{i} \left[ \frac{\kappa_{i}}{\kappa_{2}} \right]^{t} \mathbf{v}_{i}$$

$$\bullet \quad \text{as } t \to \infty, \ y(t) \to \kappa_{2}^{t} c_{2} v_{2}$$

$$\text{less than 1}$$
for all  $i > 2$ 

- caveats
  - y might have a very small component in the direction of V<sub>1</sub>
    - periodically perform a subtraction  $\mathbf{y}(t) = \mathbf{y}(t) (\mathbf{v}_1^T \mathbf{y}(t)) \mathbf{v}_1$
- not work well beyond the first couple of eigenvectors

## efficient algorithms for computing all eigenvalues and eigenvectors

- finding orthogonal matrix Q such that the similarity transform  $T = Q^T A Q$  gives either
  - a tridiagonal matrix (if A is symmetric) or
  - a Hessenberg matrix (if A is asymmetric)
- if we can find such Q, and if  $\mathbf{v}_i$  is an eigenvector of  $\mathbf{A}$  with eigenvalue  $\kappa_i$

$$\kappa_i \mathbf{Q}^T \mathbf{v}_i = \mathbf{Q}^T \mathbf{A} \mathbf{v}_i = \mathbf{T} \mathbf{Q}^T \mathbf{v}_i \qquad \qquad : \mathbf{Q}^{-1} = \mathbf{Q}^T$$

- the vector  $\mathbf{w}_i = \mathbf{Q}^T \mathbf{v}_i$  is an eigenvector of T with eigenvalue  $\kappa_i$
- eigenvectors of A are  $\mathbf{v}_i = \mathbf{Q}\mathbf{w}_i$

QL algorithm: efficient numerical method O(n) for a tridiagonal matrix O(n²) for a Heessenberg one

#### algorithms for finding Q

- for a symmetric matrix
  - Householder algorithm : O(n³)
- for a sparse symmetric matrix
  - Lanczos algorithm : O(mn)
- for an asymmetric matrix
  - Arnoldi algorithm
- combined Lanczos/QL algorithm : O(mn)

#### dividing networks into clusters

"graph partitioning", "community detection"

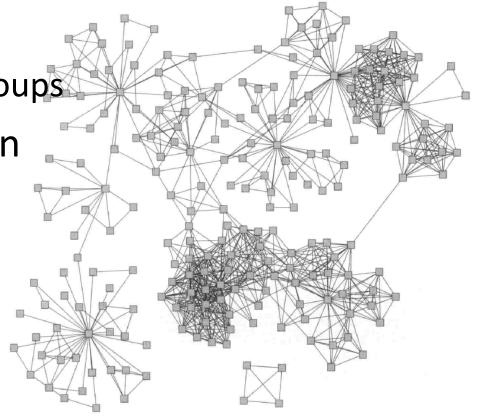
network of coauthorships in a university

department

densely connected groups

discovering groups can

be a useful tool for revealing structure and organization



#### graph partitioning

- the number and size of the groups is fixed
  - dividing a network into two groups of equal size, such that the number of edges between them is minimized
  - example: network processes on a parallel computer
    - message transmission between processors is slow

#### community detection

- the number and size of the group is unspecified
  - finding the natural divisions of a network into groups of vertices such that there are many edges within groups and few edges between groups

sizes of the groups might vary widely from one

group to another

a wide variety of definitions

a wide variety of algorithms

### difference between graph partitioning and community detection

- graph partitioning
  - the number and size of the groups is specified
  - dividing a network into smaller manageable pieces
- community detection
  - the number and size of the groups is unspecified
  - used as a tool for understanding the structure of a network

#### why partitioning is hard? (1)

- graph bisection : division into two parts
  - repeated bisection : division into arbitrary number of parts
- cut size : the number of edges between groups
- exhaustive search is prohibitively costly
  - the number of ways of dividing a network of n vertices into two groups of n<sub>1</sub> and n<sub>2</sub> vertices

$$\frac{n!}{n_1! n_2!} \cong \frac{\sqrt{2\pi n} (n/e)^n}{\sqrt{2\pi n_1} (n_1/e)^{n_1} \sqrt{2\pi n_2} (n_2/e)^{n_2}} = \frac{n^{n+1/2}}{n_1^{n_1+1/2} n_2^{n_2+1/2}} : n! \cong \sqrt{2\pi n} (n/e)^n$$

#### why partitioning is hard? (2)

• if we want to divide a network into two parts of equal size  $\frac{1}{2}n$ , the number of different ways to do it is roughly

to do it is roughly go up roughly exponentially with the size of the network with the size of the network

calculation becomes intractable for the networks beyond n=30

- Even algorithms that fail to find the very best division of a network may still find a pretty good one, and for many practical purposes pretty good is good enough.
  - heuristic algorithms

### The Kernighan-Lin algorithm 1. start with any division of the vertices into two groups

- 2. search for pairs of vertices whose interchange would reduce the cut size between the groups by the largest amount (or increase it by the smallest amount)
- 3. repeat the process (each vertex can only be moved once)
- 4. when all swaps have been completed, go back through every state and choose the state in which the cut size takes its smallest value
- 5. this entire procedure is performed repeatedly until no improvement in the cut size occurs

#### comments on the Kernighan-Lin algorithm

- if initial partitions are different, the results can be different
- the Kernighan-Lin algorithm is slow
  - # of swaps : O(n) < n/2

 $(n/2) \times (n/2) = n^2/4$ 

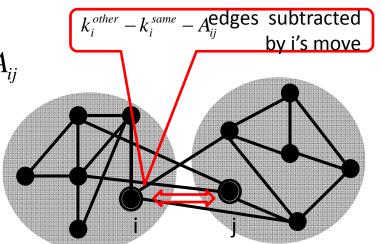
- for each swap, examine all pairs of vertices: O(n<sup>2</sup>)
- check the changes after swap
  - the change of cut size

$$\Delta = k_i^{other} - k_i^{same} + k_j^{other} - k_j^{same} - 2A_{ij}$$

• O(m/n)average
degree

- total :  $O(n \times n^2 \times m/n) = O(mn^2)$ 

O(n<sup>3</sup>) on a sparse network O(n4) on a dense network



#### spectral partitioning (1)

- a network of n vertices and m edges
- a division into group 1 and group 2
- cut size (the number of edges running between the two groups)  $R = \sum A_{ij}$

$$s_i = \begin{cases} +1 & \text{if vertex i belongs to group 1} \\ -1 & \text{if vertex i belongs to group 2} \end{cases}$$

$$\frac{1}{2}(1-s_is_j) = \begin{cases} 1 & \text{if i and j are in different groups} \\ 0 & \text{if i and j are in the same group} \end{cases}$$

$$R = \frac{1}{4} \sum_{ij} A_{ij} (1 - s_i s_j)$$

#### spectral partitioning (2)

the first term in the sum

The first term in the sum
$$\sum_{ij} A_{ij} = \sum_{i} k_{i} = \sum_{i} k_{i} S_{i}^{2} = \sum_{ij} k_{i} \delta_{ij} S_{i} S_{j} \qquad S_{i}^{2} = 1$$

$$R = \frac{1}{4} \sum_{ij} (k_{i} \delta_{ij} - A_{ij}) S_{i} S_{j} = \frac{1}{4} \sum_{ij} L_{ij} S_{i} S_{j}$$

$$R = \frac{1}{4} \mathbf{s}^{T} \mathbf{L} \mathbf{s}$$
division into groups
$$R = \frac{1}{4} \mathbf{s}^{T} \mathbf{L} \mathbf{s}$$
graph structure

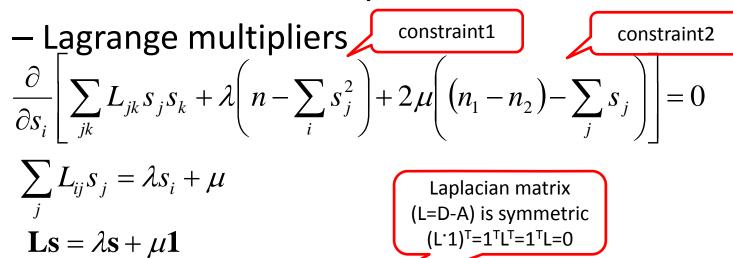
 goal: find the vector s that minimizes the cut size for given L

#### spectral partitioning (3)

- minimizing R is not easy
  - s<sub>i</sub> cannot take any values (+1 or -1)
- approximate approach: allow s<sub>i</sub> to take any values
   if no constraint, s=0 is trivial solution
  - constraint 1:  $|s| = \sqrt{n}$  or  $\sum_{i}^{\infty} s_i^2 = n$  the length of vector s should not be changed
  - constraint2:  $\sum_{i} S_{i} = n_{1} n_{2}$  or  $\mathbf{1}^{T} \mathbf{s} = n_{1} n_{2}$  the sizes of two groups should be  $n_{1}$  and  $n_{2}$

#### spectral partitioning (4)

differentiate with respect to the elements s<sub>i</sub>



$$\mathbf{L}\mathbf{s} = \lambda \mathbf{s} + \mu \mathbf{1}$$

• multiplying on the left by 
$$\mathbf{1}^{\mathsf{T}}$$

$$\mu = -\frac{n_1 - n_2}{n} \lambda \qquad \because \mathbf{L} \cdot \mathbf{1} = 0 \qquad \mathbf{1}^T \mathbf{s} = n_1 - n_2$$

$$\mathbf{x} = \mathbf{s} + \frac{\mu}{\lambda} \mathbf{1} = \mathbf{s} - \frac{n_1 - n_2}{n} \mathbf{1} \qquad \text{define new vector x}$$

$$\mathbf{L}\mathbf{x} = \mathbf{L}(\mathbf{s} + \frac{\mu}{\lambda} \mathbf{1}) = \mathbf{L}\mathbf{s} = \lambda \mathbf{s} + \mu \mathbf{1} = \lambda \mathbf{x} \qquad \text{x is an eigenvector of the Laplacian with eigenvalue } \lambda$$

(L=D-A) is symmetric (L·1) $^{T}$ =1 $^{T}$ L $^{T}$ =1 $^{T}$ L=0

#### spectral partitioning (5)

- which eigenvector should we choose?  $\mathbf{1}^{T}\mathbf{x} = \mathbf{1}^{T}\mathbf{s} + \frac{\mu}{\lambda}\mathbf{1}^{T}\mathbf{1} = (n_{1} n_{2}) \frac{n_{1} n_{2}}{n}n = 0$  **x** is orthogonal to  $\mathbf{1} \mathbf{x} \neq \mathbf{1}$   $\mathbf{1}^{T}\mathbf{x} = \mathbf{1}^{T}\mathbf{x} + \frac{\mu}{\lambda}\mathbf{1}^{T}\mathbf{1} = (n_{1} n_{2}) \frac{n_{1} n_{2}}{n}n = 0$  **x** is orthogonal to  $\mathbf{1} \mathbf{x} \neq \mathbf{1}$   $\mathbf{1}^{T}\mathbf{1} = \mathbf{1}^{T}\mathbf{1} + \mathbf{1}^{T}\mathbf{1} +$

$$R = \frac{1}{4}\mathbf{s}^{T}\mathbf{L}\mathbf{s} = \frac{1}{4}\mathbf{x}^{T}\mathbf{L}\mathbf{x} = \frac{1}{4}\lambda\mathbf{x}^{T}\mathbf{x}$$

$$\mathbf{x}^{T}\mathbf{x} = \mathbf{s}^{T}\mathbf{s} + \frac{\mu}{\lambda}(\mathbf{s}^{T}1 + \mathbf{1}^{T}\mathbf{s}) + \frac{\mu^{2}}{\lambda^{2}}\mathbf{1}^{T}\mathbf{1}$$

$$= n - 2\frac{n_{1} - n_{2}}{n_{1} - n_{2}}(n_{1} - n_{2}) + \frac{(n_{1} - n_{2})^{2}}{n} = 4\frac{n_{1}n_{2}}{n} \qquad R = \frac{n_{1}n_{2}}{n}\lambda$$

- cut size is proportional to the eigenvalue  $\lambda$ 
  - smallest eigenvalue is 0 that corresponds to eigenvector 1
  - choose x proportional to the eigenvector v<sub>2</sub> corresponding make this product to the second smallest eigenvalue  $\lambda_2$ as large as possible

$$\mathbf{s} = \mathbf{x} + \frac{n_1 - n_2}{n} \mathbf{1}$$

$$\mathbf{s}_i \text{ should be +1 or -1}$$

$$\mathbf{s}^T \left( \mathbf{x} + \frac{n_1 - n_2}{n} \mathbf{1} \right) = \sum_i s_i \left( x_i + \frac{n_1 - n_2}{n} \right)$$

#### algorithm of spectral partitioning

- 1. calculate the eigenvector  $v_2$  corresponding the second smallest eigenvalue  $\lambda_2$  of the graph Laplacian
- 2. sort the elements of the eigenvector in order from largest to smallest
- 3. put the vertices corresponding to the  $n_1$  largest elements in group 1, the rest in group 2, and calculate the cut size
- 4. then put the vertices corresponding to the n<sub>1</sub> smallest elements in group 1, the rest in group 2, and recalculate the cut size
- 5. between these two divisions of the network, choose the one that gives the smaller cut size

#### comments on spectral partitioning

- disadvantages
  - quality of partition: not quite as good as those returned by other methods
- advantages
  - speed :
    - calculation of the eigenvector v<sub>2</sub>: O(mn)
       (or O(n<sup>2</sup>) for sparse networks)

      Kernighan-Lin algorithm: O(n<sup>3</sup>)

#### community detection

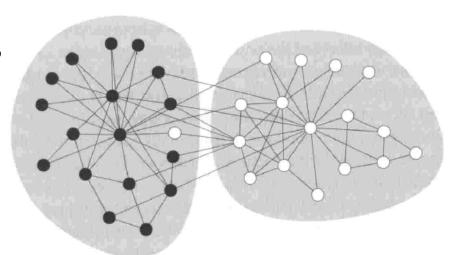
- difference from graph partitioning: the number or size of the groups is not fixed
- if no constraint, optimum division is trivial
  - group 1: all verticesgroup 2: no vertex
- loose constraint
  - minimize ratio  $\frac{R}{n_1}$  denominator has its largest value when  $n_1=n_2=n/2$
  - still biased towards equal partition
  - no principled rationale behind this definition

#### quality measure other than cut size

- a good division is one where there are fewer than expected such edges
- modularity (as the measure of assortative mixing)
- modularity maximization: look for the divisions that have the highest modularity scores
  - hard problem (takes exponential time)
  - heuristic algorithms are often used

#### simple modularity maximization

- analog of the Kernigham-Lin algorithm
  - starting from some initial (random) division
  - for each vertex check how much modularity would increase if it is moved to the other group
  - repeat the above process
  - identified communitiesare good
  - quite efficient : O(mn)



#### spectral modularity maximization (1)

analogous to spectral graph partitioning

$$Q = \frac{1}{2m} \sum_{ij} (A_{ij} - \frac{k_i k_j}{2m}) \delta(c_i, c_j) = \frac{1}{2m} \sum_{ij} B_{ij} \delta(c_i, c_j)$$

$$B_{ij} = A_{ij} - \frac{k_i k_j}{2m}$$

$$\sum_{j} B_{ij} = \sum_{j} A_{ij} - \frac{k_i}{2m} \sum_{j} k_j = k_i - \frac{k_i}{2m} 2m = 0$$

division of a network into two parts

$$s_{i} = \begin{cases} +1 & \text{if vertex i belongs to group 1} \\ -1 & \text{if vertex i belongs to group 2} \end{cases}$$

$$\delta(c_{i}, c_{j}) = \frac{1}{2}(s_{i}s_{j} + 1)$$

$$Q = \frac{1}{4m} \sum_{ij} B_{ij}(s_{i}s_{j} + 1) = \frac{1}{4m} \sum_{ij} B_{ij}s_{i}s_{j} \qquad Q = \frac{1}{4m} \mathbf{s}^{T} \mathbf{B} \mathbf{s}$$
modularity matrix
$$Q = \frac{1}{4m} \sum_{ij} B_{ij}(s_{i}s_{j} + 1) = \frac{1}{4m} \sum_{ij} B_{ij}s_{i}s_{j} \qquad Q = \frac{1}{4m} \mathbf{s}^{T} \mathbf{B} \mathbf{s}$$

#### spectral modularity maximization (2)

find the value of s that maximize Q

$$Q = \frac{1}{4m} \mathbf{s}^T \mathbf{B} \mathbf{s}^{-\frac{\text{elements of s}}{\text{are } + 1 \text{ or } -1}}$$

- relax the constraints on s
  - elements of s can take arbitrary value
  - but its length has to be the same  $\mathbf{s}^T\mathbf{s} = \sum_i s_i^2 = n$

$$\frac{\partial}{\partial s_i} \left[ \sum_{jk} B_{jk} s_j s_k + \beta \left( n - \sum_j s_j^2 \right) \right] = 0$$

$$\sum_j B_{ij} s_j = \beta s_i \qquad \mathbf{Bs} = \beta \mathbf{s} \qquad \text{$\mathbf{s}$ is one of the eigenvectors of the modularity matrix}$$

#### spectral modularity maximization (3)

$$Q = \frac{1}{4m} \beta \mathbf{s}^T \mathbf{s} = \frac{n}{4m} \beta$$

- choose  $\mathbf{s} = \mathbf{u}_1$  (eigenvector corresponding the largest eigenvalue of the modularity matrix)

• maximizing the product 
$$\mathbf{s}^{T}\mathbf{u}_{1} = \sum_{i} s_{i} [\mathbf{u}_{1}]_{i} \qquad s_{i} = \begin{cases} +1 & \text{if } [\mathbf{u}_{1}]_{i} > 0 \\ -1 & \text{if } [\mathbf{u}_{1}]_{i} < 0 \end{cases}$$

- <algorithm>
  - 1. calculate the eigenvector of the modularity matrix that corresponding to the largest eigenvalue
  - 2. assign vertices to communities according to the signs of the vector elements (positive/negative)

#### spectral modularity maximization (4)

- unlike Laplacian, modularity matrix is not sparse
  - finding the leading eigenvector:O(mn)
    - O(n³) for dense matrix
    - O(n²) for sparse matrix
  - by exploiting special properties of the modularity matrix, it is still possible to find the eigenvector in time  $O(n^2)$  on a sparse network

#### division into more than two groups

- repeating bipartitioning
  - even if each bipartition is optimal, repeated bipartitioning may not be optimal
- ullet consider the change  $\Delta Q$  in the modularity of the entire network

bisecting a community c of size n<sub>c</sub>

$$\Delta Q = \frac{1}{2m} \left[ \frac{1}{2} \sum_{i,j \in c} B_{ij} (s_i s_j + 1) - \sum_{i,j \in c} B_{ij} \right]$$

$$= \frac{1}{4m} \left[ \sum_{i,j \in c} B_{ij} s_i s_j - \sum_{i,j \in c} B_{ij} \right] = \frac{1}{4m} \sum_{i,j \in c} \left[ B_{ij} - \delta_{ij} \sum_{k \in c} B_{ik} \right] s_i s_j$$

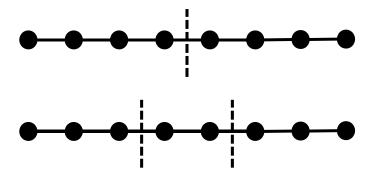
$$= \frac{1}{4m} \mathbf{s}^T \mathbf{B}^{(c)} \mathbf{s}$$

$$B_{ij}^{(c)} = B_{ij} - \delta_{ij} \sum_{k \in c} B_{ik}$$

$$\mathbf{n}_c \times \mathbf{n}_c \text{ matrix}$$

#### weakness of repeated bipartitioning

 repeated optimal bipartitioning may not be able to find optimal division



### other modularity maximization methods

- simulated annealing
  - analogy with the physics of slow cooling of solids
  - ground state: the state of the lowest energy
  - pros: high quality cons: slow
- genetic algorithm
  - analogy with biological evolution
  - assign "fitness" to each of the population of different divisions
  - pros: high quality cons: slow
- greedy algorithm
  - bottom up : choose the merge of the biggest increase of modularity
  - pros: fast cons: moderate quality

### other algorithms for community detection

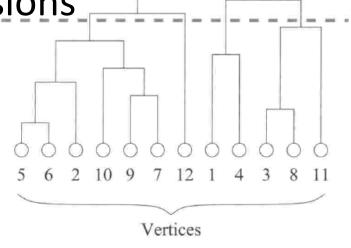
- there is no universally agreed definition of communities
  - the previous discussion focus on modularity maximization, but there are other definitions
- some algorithms are introduced in the following slides
  - betweenness-based method
  - hierarchical clustering
- if you want to learn more, read the following article
  - "Community detection in graphs" by Santo Fortunato
  - Physics Reports, Vol. 486, Issues 3-5, pp.75-174 February 2010
  - http://www.sciencedirect.com/science/article/pii/S0370157309 002841

#### betweenness-based methods

- look for the edges that lie between communities, and remove them
- edge betweenness: the number of geodesic (shortest) paths that run along the edge
  - takes time of order O(n(m+n))
- recalculation is required for each removal

#### dendrogram

- root(top): all vertices in one group
- leaves(bottom): each vertex in a one-vertex group
- the algorithm generates the dendrogram from top to bottom
- selection from different divisions
  - coarse division (top)
  - fine division (bottom)



### comments on betweenness-based method

- pros
  - hierarchical decomposition : dendrogram
- cons
  - slow: entire algorithm takes time O(mn(m+n))
- Radicchi's method

search for the edges that belong to short loops

- faster :  $O(n^2)$ 

"bridge" edges do not belong to short loops

O(n<sup>3</sup>) for sparse networks

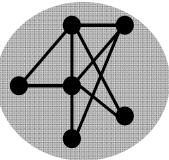
#### hierarchical clustering

- agglomerative (<-> divisive )
- define a measure of similarity, and join the most similar vertices to form groups
  - cosine similarity
  - correlation coefficient
  - Euclidean distance

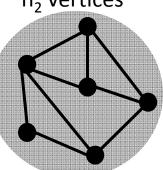
### similarity of vertices → similarity of groups

- there are n<sub>1</sub>n<sub>2</sub> pairs of vertices
- single-linkage clustering
  - similarity of the most similar pair
- complete-linkage clustering
  - similarity of the least similar pair
- average-linkage clustering
  - mean similarity of all pairs

group 1 n<sub>1</sub> vertices



group 2 n<sub>2</sub> vertices

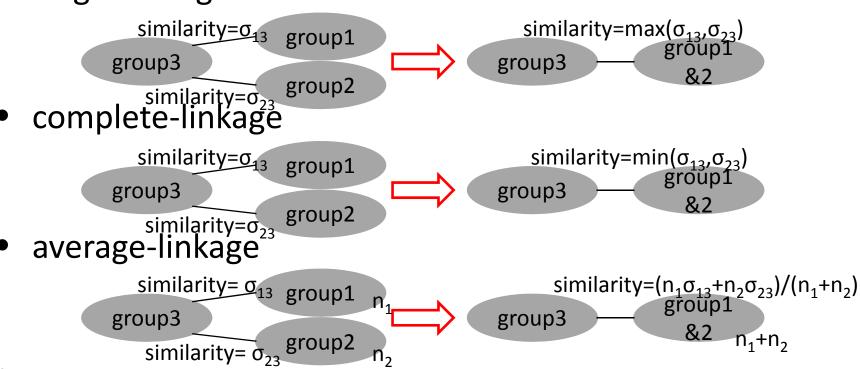


#### procedure of hierarchical clustering

- Choose a similarity measure and evaluate it for all vertex pairs.
- 2. Assign each vertex to a group of its own, consisting of just that one vertex. The initial similarities of the groups are simply the similarities of the vertices.
- 3. Find the pair of groups with the highest similarity and join them together into a single group.
- 4. Calculate the similarity between the new composite group and all others using one of the three methods (single-, complete-, or average-linkage clustering).
- 5. Repeat from step 3 until all vertices have been joined into a single group.

#### similarity of two groups after join

single-linkage



### computational complexity of hierarchical clustering

- O(n) for recalculation of similarities
  - O(n²) for naive approach (recalculation + search for the biggest one)
  - O(nlogn) for using a heap (recalculation + storing in a binary heap)
- joining groups has to be repeated n-1 times
- total
  - O(n³) for naive implementation
  - O(n<sup>2</sup>logn) for using a heap

#### comments on hierarchical clustering

- results depend on which similarity measure one chooses and which linkage method
- good at picking out the cores of groups, but less good at assigning peripheral vertices to appropriate groups