CSE8803/CX4803 Machine Learning in Computational Biology

Lecture 10: Clustering II

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Course logistics

- Final exam: take home and timed
 - 24 hour time window: May 4, 10:00am to May 5, 10:00am
 - 4 hour duration including the time to scan and submit your solutions to Canavs quizzes.
 You must successfully submit your solutions to Canvas or it is considered as late submission.

Paper presentation comments

- Each student in the audience can submit optional comments on paper presentations
- Some comments will be considered when evaluating the presentation
- Reasonable and meaningful comments on the presentation are also counted towards your "participation" points. You can also submit insights on the paper (that is not discussed during the presentation) and that can also be considered towards participation points.

Hierarchical clustering

Idea: agglomerative clustering

- First merge very similar instances
- Incrementally build larger clusters out of smaller clusters

Algorithm: Maintain a set of clusters

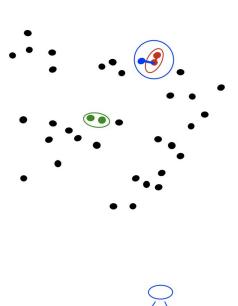
- Initially, each instance in its own cluster
- Repeat:
 - Pick the two closest clusters
 - Merge them into a new cluster

Stop when there's only one cluster left

Output:

Produces not one clustering, but a family of clusterings represented by a dendrogram

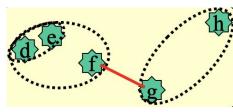
How to define?



Hierarchical clustering: define "closest" for clusters

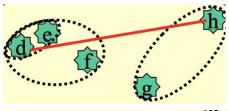
Closest pair (single-link)

 $CD(X,Y)=min_{x \in X, y \in Y} D(x,y)$



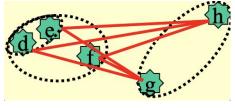
Furthest pair (complete-link)

 $CD(X,Y)=max_{x \in X, y \in Y}D(x,y)$



average-link

 $CD(X,Y)=avg_{x \in X, y \in Y}D(x,y)$



centroids method

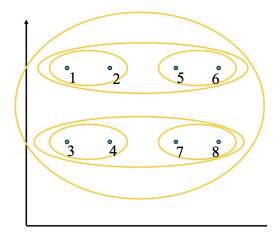
CD(X,Y)=D(avg(X), avg(Y))



Hierarchical clustering: define "closest" for clusters

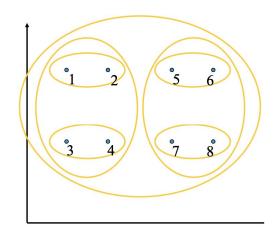
Cluster distance affects both results and runtime

Closest pair (single-link clustering)



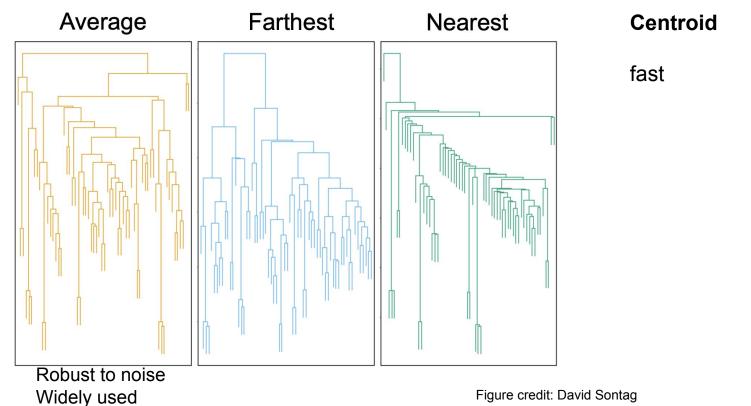
Potentially long and skinny clusters

Farthest pair (complete-link clustering)



tight clusters

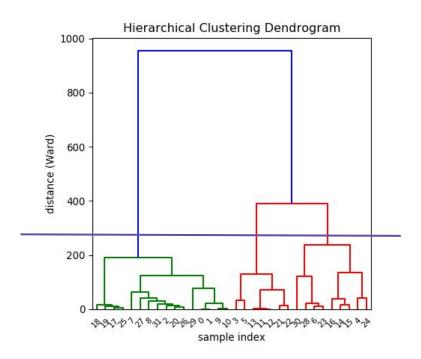
Hierarchical clustering: define "closest" for clusters



Hierarchical clustering: distance measures

Table 1 Gene expression similarity measure	es
Manhattan distance (city-block distance, L1 norm)	$d_{fg} = \sum_{c} \left e_{fc} - e_{gc} \right $
Euclidean distance (L2 norm)	$d_{fg} = \sqrt{\sum_c \left(e_{fc} - e_{gc}\right)^2}$
Mahalanobis distance	$d_{fg} = (\mathbf{e}_f - \mathbf{e}_g)^{T} \Sigma^{-1} (\mathbf{e}_f - \mathbf{e}_g)$, where Σ is the (full or within-cluster) covariance matrix of the data
Pearson correlation (centered correlation)	$d_{fg} = 1 - r_{fg}$, with $r_{fg} = \frac{\sum_{c} (e_{fc} - \bar{e}_f)(e_{gc} - \bar{e}_g)}{\sqrt{\sum_{c} (e_{fc} - \bar{e}_f)^2 \sum_{c} (e_{gc} - \bar{e}_g)^2}}$
Uncentered correlation (angular separation, cosine angle)	$d_{fg} = 1 - r_{fg}$, with $r_{fg} = \frac{\sum_{c} e_{fc} e_{gc}}{\sqrt{\sum_{c} e_{fc}^2 \sum_{c} e_{gc}^2}}$
Spellman rank correlation	As Pearson correlation, but replace e_{gc} with the rank of e_{gc} within the expression values of gene g across all conditions $c=1C$
Absolute or squared correlation	$d_{fg} = 1 - r_{fg} \text{ or } d_{fg} = 1 - r_{fg} ^2$
d_{fg} , distance between expression patterns for genes f and g . e_{l}	$_{ m gc}$, expression level of gene g under condition c .

Hierarchical clustering: obtaining the clusters



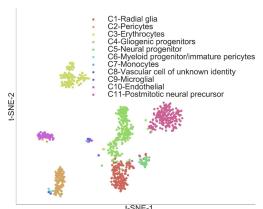
Hierarchical clustering: Summary

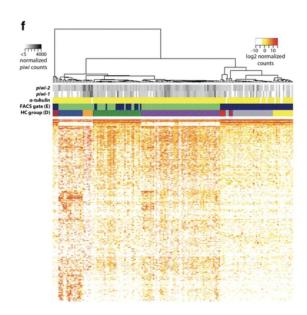
- No need to specify the number of clusters in advance.
- Hierarchical structure maps nicely onto human intuition for some domains
- They do not scale well: where n is the number of total objects.
- Like any heuristic search algorithms, local optima are a problem.
- Interpretation of results can be subjective

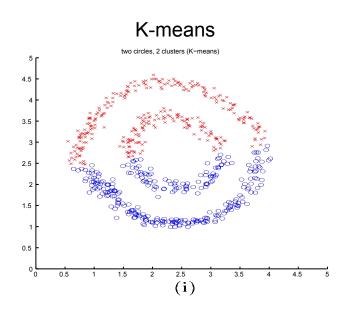
Clustering algorithms

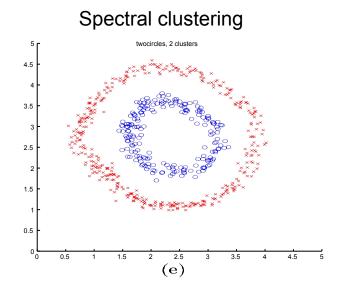
Clustering - unsupervised learning methods

- K-means (and fuzzy k-means)
- Hierarchical clustering
- Spectral clustering
- Louvain, Leiden clustering

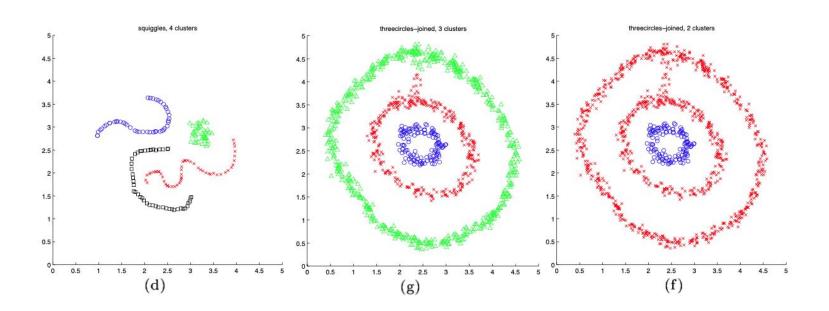




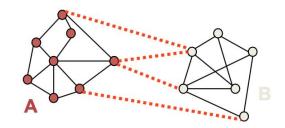




[Shi & Malik '00; Ng, Jordan, Weiss NIPS '01]



Idea: group points based on links in a graph



To construct the graph:

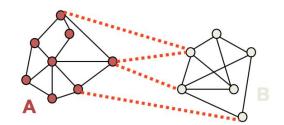
- Fully connected weighted graph
- K nearest neighbor (kNN) graph, where for each vertex, it's connected to its k nearest neighbors.

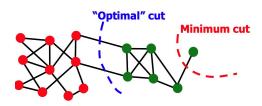
Question: degree of nodes in a kNN graph?

The *min-cut* problem

Partition graph into two sets A and B such that weight of edges connecting vertices in A to vertices in B is minimum.

For an unweighted graph, find a *cut* with the smallest number of edges.





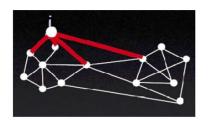
Problem with the min-cut criterion for clustering:

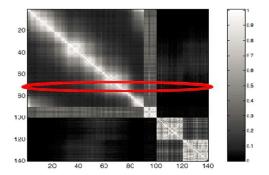
- Only considers external cluster connections
- Does not consider internal cluster connectivity

Spectral clustering: graph terminologies

• Degree of nodes

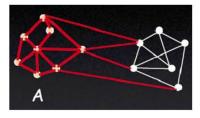
$$d_i = \sum_j w_{i,j}$$

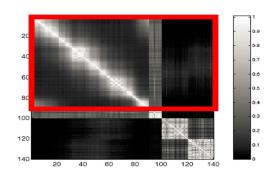




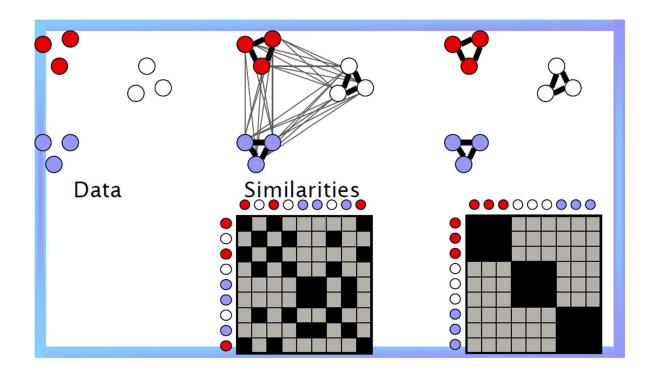
Volume of a set

$$vol(A) = \sum_{i \in A} d_i, A \subseteq V$$





Spectral Clustering: graph partitioning



Spectral clustering: objective

Objective of min-cut: minimize
$$\operatorname{cut}(A, B) = \sum_{i \in A, j \in B} w_{i,j}$$

Objective of spectral clustering: minimize normalized cut

$$Ncut(A, B) = cut(A, B) \left(\frac{1}{vol(A)} + \frac{1}{vol(B)} \right)$$

Which represents connectivity between groups relative to the density of each group

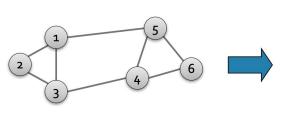
Advantage: produces more balanced clusters

Spectral clustering: objective

Objective of spectral clustering: minimize normalized cut

$$Ncut(A, B) = cut(A, B) \left(\frac{1}{vol(A)} + \frac{1}{vol(B)} \right)$$

Define D, which is a diagonal matrix with the degree of every node: $D(i,i) = \sum_{j} W(i,j)$



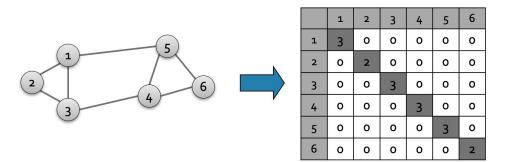
	1	2	3	4	5	6
1	3	0	0	0	0	0
2	0	2	0	0	0	0
3	0	0	3	0	0	0
4	0	0	0	3	0	0
5	0	0	0	0	3	0
6	0	0	0	0	0	2

Spectral clustering: objective

Objective of spectral clustering: minimize normalized cut

$$Ncut(A, B) = cut(A, B) \left(\frac{1}{vol(A)} + \frac{1}{vol(B)} \right)$$

Define L = D - W (graph Laplacian)



	1	2	3	4	5	6
1	3	-1	-1	o	-1	0
2	-1	2	-1	o	o	0
3	-1	-1	3	-1	o	0
4	o	o	-1	3	-1	-1
5	-1	o	o	-1	3	-1
6	o	o	o	-1	-1	2

The laplacian matrix L is positive semi-definite

- All eigenvalues are ≥ 0
- $x^T L x = \sum_{ij} L_{ij} x_i x_j \ge 0$, for every x
- $\bullet \quad L = N^T \cdot N$

Spectral clustering: steps

Objective of spectral clustering: minimize normalized cut

$$Ncut(A, B) = cut(A, B) \left(\frac{1}{vol(A)} + \frac{1}{vol(B)} \right)$$

Steps:

- 1. Calculate D, $D(i,i) = \sum_{j} W(i,j)$
- 2. Calculate L, L = D W, and L', L'= $D^{-1/2}LD^{-1/2}$
- 3. Calculate eigenvalues and eigenvectors of L'. Sort the eigenvalues from low to high values.
- 4. Take the corresponding first k eigenvectors
- 5. Perform k-means on these eigenvectors and obtain clusters

Objective of spectral clustering: minimize normalized cut

$$Ncut(A, B) = cut(A, B) \left(\frac{1}{vol(A)} + \frac{1}{vol(B)} \right)$$

$$\text{define} \quad f = \begin{bmatrix} f_1, f_2, \dots, f_n \end{bmatrix}^T \quad \text{where} \quad f_i = \begin{cases} \frac{1}{\text{vol}(A)} & \text{if } i \in A \\ -\frac{1}{\text{vol}(B)} & \text{if } i \in B \end{cases}$$

f is determined by the cluster assignment of nodes!

$$\mathbf{f}^T\mathbf{D}\mathbf{f} = \sum_i d_i \mathbf{f}_i^2 = \sum_{i \in A} \frac{d_i}{\operatorname{vol}(A)^2} + \sum_{j \in B} \frac{d_j}{\operatorname{vol}(B)^2} = \frac{1}{\operatorname{vol}(A)} + \frac{1}{\operatorname{vol}(B)}$$
Divide all nodes into set A and set B
$$\sum_{i \in A} d_i = \operatorname{vol}(A)$$

Objective of spectral clustering: minimize normalized cut

$$\operatorname{Ncut}(A, B) = \operatorname{cut}(A, B) \left(\frac{1}{\operatorname{vol}(A)} + \frac{1}{\operatorname{vol}(B)} \right)$$

$$\operatorname{define} \ f = \begin{bmatrix} f_1, f_2, \dots, f_n \end{bmatrix}^T \text{ where } f_i = \begin{cases} \frac{1}{\operatorname{vol}(A)} & \text{if } i \in A \\ -\frac{1}{\operatorname{vol}(B)} & \text{if } i \in B \end{cases}$$

$$\mathbf{f}^T \mathbf{L} \mathbf{f} = \sum_{ij} w_{ij} (\mathbf{f}_i - \mathbf{f}_j)^2 = \sum_{i \in A, j \in B} w_{ij} \left(\frac{1}{\text{vol}(A)} + \frac{1}{\text{vol}(B)} \right)^2$$

$$\downarrow^{\text{L=D-W}}_{\text{W}_{ij}=0}$$
If i,j in the same cluster, then $\mathbf{f}_i - \mathbf{f}_j = 0$

Objective of spectral clustering: minimize normalized cut

$$\operatorname{Ncut}(A, B) = \operatorname{cut}(A, B) \left(\frac{1}{\operatorname{vol}(A)} + \frac{1}{\operatorname{vol}(B)}\right)$$

$$= \sum_{i \in A, j \in B} w_{ij} \left(\frac{1}{\operatorname{vol}(A)} + \frac{1}{\operatorname{vol}(B)}\right)$$

$$\operatorname{Ncut}(A, B) = \frac{\mathbf{f}^T \mathbf{L} \mathbf{f}}{\mathbf{f}^T \mathbf{D} \mathbf{f}}$$

$$\mathbf{f}^T \mathbf{L} \mathbf{f} = \sum_{ij} w_{ij} (\mathbf{f}_i - \mathbf{f}_j)^2 = \sum_{i \in A, j \in B} w_{ij} \left(\frac{1}{\operatorname{vol}(A)} + \frac{1}{\operatorname{vol}(B)}\right)^2$$

$$\mathbf{f}^T \mathbf{D} \mathbf{f} = \sum_{j} d_i \mathbf{f}_i^2 = \sum_{i \in A} \frac{d_i}{\operatorname{vol}(A)^2} + \sum_{j \in B} \frac{d_i}{\operatorname{vol}(B)^2} = \frac{1}{\operatorname{vol}(A)} + \frac{1}{\operatorname{vol}(B)}$$

Objective of spectral clustering: minimize normalized cut

$$\min_{\mathbf{f}} \mathbf{Ncut}(A, B) = \min_{\mathbf{f}} \frac{\mathbf{f}^T \mathbf{Lf}}{\mathbf{f}^T \mathbf{Df}}$$

$$f^T \mathbf{D} 1 = 0, \text{ that is, } \Sigma_i \mathbf{d}_i f_i = 0$$

$$f = \begin{cases} \frac{1}{\operatorname{vol}(A)} & \text{if } i \in A \\ -\frac{1}{\operatorname{vol}(B)} & \text{if } i \in B \end{cases}$$

With relaxation on **f** from discrete to continuous space

Once this is solved, we can obtain cluster labels by the sign of f_i ($f_i > 0$ or $f_i < 0$)

Objective of spectral clustering: minimize normalized cut

min Ncut(
$$A$$
, B) = min $\frac{\mathbf{f}^T \mathbf{L} \mathbf{f}}{\mathbf{f}^T \mathbf{D} \mathbf{f}}$

$$f_{i} = \begin{cases} \frac{1}{\operatorname{vol}(A)} & \text{if } i \in A \\ -\frac{1}{\operatorname{vol}(B)} & \text{if } i \in B \end{cases}$$

$$\min_{f} f^{T} L f \quad \text{s.t.} f^{T} D f = 1$$

The objective thus becomes

$$f^{T}Lf = f^{T}D^{1/2}D^{-1/2}L D^{-1/2}D^{1/2}f$$

$$= u^{T}D^{-1/2}L D^{-1/2}u$$

$$f^TD1=0$$
, that is, $\Sigma_i d_i f_i = 0$

Now let $u=D^{1/2}f$, then $u^{T}=f^{T}D^{1/2}$, thus the constraint becomes:

$$\int_{-1}^{1} dt \frac{1}{2} D^{\frac{1}{2}} f = 1$$

$$\int_{-1}^{1} u^{T} u = 1$$

Let L'=D^{-1/2}LD^{-1/2}, the optimization problem becomes: $\min_{u} u^{T} L' u$ s.t. $u^{T} u = 1$

Objective of spectral clustering: minimize normalized cut

$$\begin{array}{c|c} \min_{u} u^T L' \ u \\ \text{s.t. } u^T u = 1 \end{array} \qquad \begin{array}{c} \text{Objective function:} \\ \max_{w: \|w\| \leq 1} \mathbf{w}^T \hat{\mathbf{R}} \mathbf{w} \end{array} \qquad \begin{array}{c} \text{PCA proof} \\ \end{array}$$

The eigen decomposition of L' will give us the solution.

As we are minimizing, we will sort eigenvalues from small to large.

However, can we use the smallest eigenvalue?

Observation. L has an eigenvalue (smallest) which is 0, corresponding to an eigenvector with all 1s.

$$(D-W) \overrightarrow{1} = D\overrightarrow{1} - W\overrightarrow{1} = \overrightarrow{0}$$

$$L\overrightarrow{1} = 0 \cdot \overrightarrow{1}$$

L' also has a 0 eigenvalue and the corresponding eigenvector is $D^{1/2}\vec{1}$

Objective of spectral clustering: minimize normalized cut

$$\min_{u} u^T L' u$$
 Objective function:
$$\sup_{\mathbf{x}.\mathbf{t}.\ u^T u = 1} \text{Looks familiar?} \qquad \max_{w: \|w\| \le 1} \mathbf{w}^T \hat{\mathbf{R}} \mathbf{w}$$
 PCA proof

The eigen decomposition of L' will give us the solution.

As we are minimizing, we will sort eigenvalues from small to large.

However, can we use the smallest eigenvalue?

If we take this eigenvalue,
$$u=D^{1/2}\vec{1}$$

What is f?

$$u=D^{1/2}f$$

 $f=D^{-1/2}u=D^{-1/2}D^{1/2}$ **1 = 1**

This f is not informative for us to find cluster assignment for the nodes!

L' also has a 0 eigenvalue and the corresponding eigenvector is $D^{1/2}\vec{1}$

Objective of spectral clustering: minimize normalized cut

$$\min_{u} u^{T} L' u$$
s.t. $u^{T} u = 1$

Looks familiar?

 $max w^T \hat{R} w$

 $w:||w|| \le 1$

PCA proof

The eigen decomposition of L' will give us the solution.

As we are minimizing, we will sort eigenvalues from small to large. However, can we use the smallest eigenvalue?

If we take this eigenvalue, $u=D^{1/2}1$ What is f?

$$u=D^{1/2}f$$

 $f=D^{-1/2}u=D^{-1/2}D^{1/2}$ **1 = 1**

This f is not informative for us to find cluster assignment for the nodes!



We will use the **2nd smallest** eigenvalue of L' and its corresponding eigenvector to obtain u and f, and then the cluster assignment.

Spectral clustering: partition a graph into 2 clusters

Objective of spectral clustering: minimize normalized cut

$$\operatorname{Ncut}(A, B) = \operatorname{cut}(A, B) \left(\frac{1}{\operatorname{vol}(A)} + \frac{1}{\operatorname{vol}(B)} \right)$$

$$= \sum_{i \in A, j \in B} w_{ij} \left(\frac{1}{\operatorname{vol}(A)} + \frac{1}{\operatorname{vol}(B)} \right)$$
1.

$$\min_{f} f^{T} L f \quad \text{s.t.} f^{T} D f = 1$$

$$\min_{u} u^{T} L' u$$
s.t. $u^{T} u = 1$

Solution:

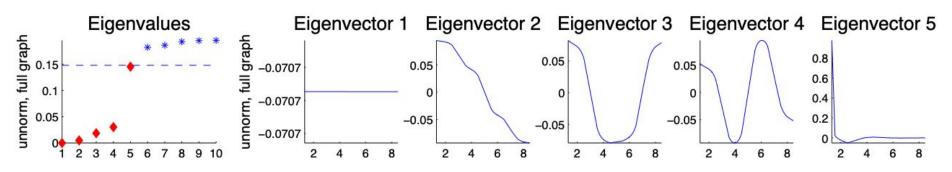
- 1. Calculate the eigenvalues and eigenvectors of L'
- 2. u = the 2nd smallest eigenvalue
- 3. $f = D^{-1/2}u$
- 4. Apply a threshold on f to determine the cluster assignment of nodes.

Other versions of spectral clustering: Perform eigen-decomposition on L or other versions of L.

From 2-way partition to k-way partition

Cluster multiple eigenvectors

- Build a reduced space from multiple eigenvectors.
- (its like doing dimension reduction then k-means)



- 1st Eigenvector is the all ones vector **1** (if graph is connected)
- 2nd Eigenvector thresholded at 0 separates first two clusters from last two
- · k-means clustering of the 4 eigenvectors identifies all clusters

Spectral clustering: steps

Objective of spectral clustering: minimize normalized cut

$$Ncut(A, B) = cut(A, B) \left(\frac{1}{vol(A)} + \frac{1}{vol(B)} \right)$$

Steps:

- 1. Calculate D, $D(i,i) = \sum_{j} W(i,j)$
- 2. Calculate L, L = D W, and L', L'= $D^{-1/2}LD^{-1/2}$

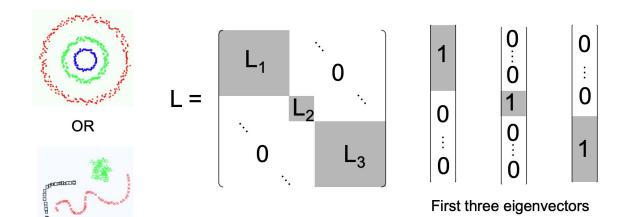
There are different normalized forms of L

- 3. Calculate eigenvalues and eigenvectors of L'. Sort the eigenvalues from low to high values.
- 4. Take the corresponding first k eigenvectors and obtain $U = \begin{bmatrix} u_1 & u_2 & \dots & u_k \end{bmatrix}_{nxk}$
- 5. Perform k-means on these eigenvectors and obtain clusters

Normalize each row of U

Spectral clustering: more fun facts

- If graph is connected, first Laplacian evec is constant (all 1s)
- If graph is disconnected (k connected components), Laplacian is block diagonal and first k Laplacian evecs are:



First 3 eigenvalues are 0s!

Spectral clustering: why it works

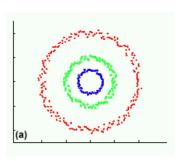
Steps:

- 1. Calculate D, $D(i,i) = \sum_{j} W(i,j)$
- 2. Calculate L, L = D W
- 3. Calculate eigenvalues and eigenvectors of L. Sort the eigenvalues from low to high values.
- 4. Take the corresponding first k eigenvectors

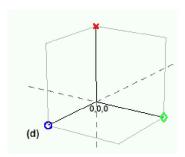
Non-linear embedding

5. Perform k-means on these eigenvectors and obtain clusters

Original data



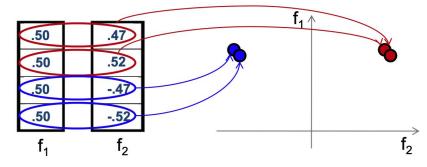
Projected data



Spectral clustering: why it works

Can put data points into blocks using eigenvectors:

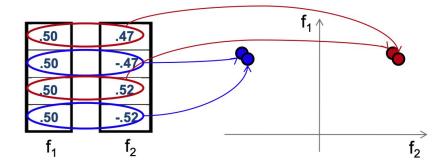
1	1	.2	0
1	1	0	.1
.2	0	1	1
0	.1	1	1
	V	V	



Non-linear embedding

Embedding is same regardless of data ordering:

1	.2	1	0
.2	0	1	1
1	1	0	.1
0	1	.1	1
	V	·——	

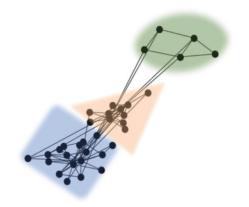


Graph based clustering methods

Spectral clustering

Louvain/Leiden clustering methods:

Optimize *Modularity*



References

Jianbo Shi and J. Malik, "Normalized cuts and image segmentation," Proceedings of IEEE Computer Society Conference on Computer Vision and Pattern Recognition, San Juan, PR, USA, 1997, pp. 731-737, doi: 10.1109/CVPR.1997.609407.

Barton, T., Bruna, T. & Kordik, P. Chameleon 2: An Improved Graph-Based Clustering Algorithm. *ACM Trans. Knowl. Discov. Data* **13**, 1–27 (2019)