

10 Spectral Clustering

Another perspective on clustering is that there are three main types: (1) *Bottom-Up*, (2) *Assignment-Based*, and (3) *Top-Down*. The bottom-up variety was like the hierarchical agglomerative clustering where we start with very small clusters and build bigger clusters. The assignment based clustering was like the k -center or the k -means variety where we “assign” each object to a center. Given the centers, there is no need to build or carve the clusters. The third type, top-down clustering, is what we will be discussing here. It starts from one big cluster and gradually divides the big clusters into smaller and smaller clusters.

At a high level the idea of top down clustering can be described very easily.

- Find the best cut of the data into two pieces.
- Recur on both pieces until that data should not be split anymore.

What remains is to determine the best way to split a set into two pieces. Then finding a threshold has similar options as with Hierarchical clustering.

Also we will need to discuss graphs, and perform clustering on graphs.

10.1 Graphs

A graph is an *abstract data type* that may seem very natural once you are familiar with and used to it. But if it is new, it may take a while to sink in. We will revisit them many times in the class.

A graph $G = (V, E)$ is defined by a set of vertices $V = \{v_1, v_2, \dots, v_n\}$ and a set of edges $E = \{e_1, e_2, \dots, e_m\}$ where each edge e_j is an unordered (or ordered in a directed graph) pair of edges: $e_j = \{v_i, v_{i'}\}$.

Two vertices v_1 and v_k are *connected* if there is a sequence of edges $\langle e_1, \dots, e_{k-1} \rangle$ such that e_1 contains v_1 , e_{k-1} contains v_k , and each consecutive edges can be ordered so $e_j = \{v_i, v_{i+1}\}$ and $e_{j+1} = \{v_{i+1}, v_{i+2}\}$ where the second element in e_j is the same as the first in e_{j+1} .

Consider an example graph portrayed three ways.

Mathematically: $G = (V, E)$ where

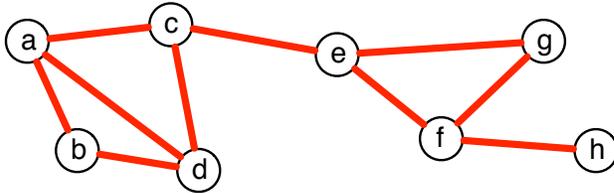
$$V = \{a, b, c, d, e, f, g\} \text{ and}$$

$$E = \left\{ \{a, b\}, \{a, c\}, \{a, d\}, \{b, d\}, \{c, d\}, \{c, e\}, \{e, f\}, \{e, g\}, \{f, g\}, \{f, h\} \right\}.$$

Matrix-Style: As a matrix with 1 if there is an edge, and 0 otherwise. (For a directed graph, it may not be symmetric).

$$G = \begin{array}{c|cccccccc} & a & b & c & d & e & f & g & h \\ \hline a & 0 & 1 & 1 & 1 & 0 & 0 & 0 & 0 \\ b & 1 & 0 & 0 & 1 & 0 & 0 & 0 & 0 \\ c & 1 & 0 & 0 & 1 & 1 & 0 & 0 & 0 \\ d & 1 & 1 & 1 & 0 & 0 & 0 & 0 & 0 \\ e & 0 & 0 & 1 & 0 & 0 & 1 & 1 & 0 \\ f & 0 & 0 & 0 & 0 & 1 & 0 & 1 & 1 \\ g & 0 & 0 & 0 & 0 & 1 & 1 & 0 & 0 \\ h & 0 & 0 & 0 & 0 & 0 & 1 & 0 & 0 \end{array} = \begin{pmatrix} 0 & 1 & 1 & 1 & 0 & 0 & 0 & 0 \\ 1 & 0 & 0 & 1 & 0 & 0 & 0 & 0 \\ 1 & 0 & 0 & 1 & 1 & 0 & 0 & 0 \\ 1 & 1 & 1 & 0 & 0 & 0 & 0 & 0 \\ 0 & 0 & 1 & 0 & 0 & 1 & 1 & 0 \\ 0 & 0 & 0 & 0 & 1 & 0 & 1 & 1 \\ 0 & 0 & 0 & 0 & 1 & 1 & 0 & 0 \\ 0 & 0 & 0 & 0 & 0 & 1 & 0 & 0 \end{pmatrix}$$

Pictorially: A ball stick model of a graph.



10.2 Clustering on Graphs

So how to cluster a graph? A cluster is a subset $S \subset V$. We are performing top down clustering, so we only need to consider a subset S and its compliment $\bar{S} = V \setminus S$.

In general, we want many edges within a cluster (small width), and few edges between clusters (large split).

- The *volume* of a cluster is $\text{Vol}(S) =$ the number of edges with at least one vertex in V .
- The *cut* between two clusters S, T is $\text{Cut}(S, T) =$ the number of edges with one vertex in S and the other in T .

Then we want a large $\text{Vol}(S)$ for each cluster and a small $\text{Cut}(S, T)$ for each pair of clusters.

Specifically, the *normalized cut* between S and T is $\text{NCut}(S, T) = \frac{\text{Cut}(S, T)}{\text{Vol}(S)} + \frac{\text{Cut}(S, T)}{\text{Vol}(T)}$. And we want to find the cluster S (and compliment $T = V \setminus S$) that has the *minimum* $\text{NCut}(S, T)$. Dividing by $\text{Vol}(S)$ and $\text{Vol}(T)$ prevents us from finding either S or T that is too small, and the $\text{Cut}(S, T)$ on top will force a large split.

For instance, in the above example, the minimum normalized cut is $S = \{a, b, c, d\}$, but the cluster with $S' = \{h\}$ has just as small $\text{Cut}(S', T')$ value. But its normalized cut is $1 + \frac{1}{10} = 1.1$, where as $\text{NCut}(S, T) = \frac{1}{6} + \frac{1}{5} = 0.367$.

10.2.1 Spectral Clustering

Start with an *adjacency* matrix

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

and the *degree* matrix, which along the diagonal stores the degree of each vertex. The *degree* of a vertex is the number of edges that contain that vertex.

$$D = \begin{pmatrix} 3 & 0 & 0 & 0 & 0 & 0 & 0 & 0 \\ 0 & 2 & 0 & 0 & 0 & 0 & 0 & 0 \\ 0 & 0 & 3 & 0 & 0 & 0 & 0 & 0 \\ 0 & 0 & 0 & 3 & 0 & 0 & 0 & 0 \\ 0 & 0 & 0 & 0 & 3 & 0 & 0 & 0 \\ 0 & 0 & 0 & 0 & 0 & 3 & 0 & 0 \\ 0 & 0 & 0 & 0 & 0 & 0 & 2 & 0 \\ 0 & 0 & 0 & 0 & 0 & 0 & 0 & 1 \end{pmatrix}.$$

Before we proceed we will want to use the matrix $D^{-1/2}$; for a diagonal matrix, a power p just takes every element $d_{i,i}$ on the diagonal to that power $d_{i,i}^p$. So

$$D^{-1/2} = \begin{pmatrix} 0.577 & 0 & 0 & 0 & 0 & 0 & 0 & 0 \\ 0 & 0.707 & 0 & 0 & 0 & 0 & 0 & 0 \\ 0 & 0 & 0.577 & 0 & 0 & 0 & 0 & 0 \\ 0 & 0 & 0 & 0.577 & 0 & 0 & 0 & 0 \\ 0 & 0 & 0 & 0 & 0.577 & 0 & 0 & 0 \\ 0 & 0 & 0 & 0 & 0 & 0.577 & 0 & 0 \\ 0 & 0 & 0 & 0 & 0 & 0 & 0.707 & 0 \\ 0 & 0 & 0 & 0 & 0 & 0 & 0 & 1 \end{pmatrix}.$$

Then the normalized *Laplacian* matrix is now the product of $D^{-1/2}$ and A

$$I - D^{-1/2}AD^{-1/2} = \begin{pmatrix} 1 & -0.408 & -0.333 & -0.333 & 0 & 0 & 0 & 0 \\ -0.408 & 1 & 0 & -0.408 & 0 & 0 & 0 & 0 \\ -0.333 & 0 & 1 & -0.333 & -0.333 & 0 & 0 & 0 \\ -0.333 & -0.408 & -0.333 & 1 & 0 & 0 & 0 & 0 \\ 0 & 0 & -0.333 & 0 & 1 & -0.333 & -0.408 & 0 \\ 0 & 0 & 0 & 0 & -0.333 & 1 & -0.408 & -0.577 \\ 0 & 0 & 0 & 0 & -0.408 & -0.408 & 1 & 0 \\ 0 & 0 & 0 & 0 & 0 & -0.577 & 0 & 1 \end{pmatrix}.$$

The *eigenvector* of a matrix M is the the vector v such that

$$Mv = \lambda v,$$

where λ is a scalar. Then λ is the corresponding *eigenvalue*. We usually restrict that $\|v\| = 1$.

There are (typically) several eigenvectors of L (the normalized Laplacian): We list them here sorted by λ .

λ	0	0.125	0.724	1.00	1.33	1.42	1.66	1.73
V	-.39	0.38	-.09	0.00	0.71	0.26	-.32	0.16
	-.32	0.36	-.27	0.50	0.00	-.51	0.38	-.18
	-.39	0.18	0.36	-.61	0.00	0.03	0.47	-.29
	-.39	0.38	-.09	0.00	-.71	0.26	-.32	0.16
	-.39	-.28	0.48	0.00	0.00	-.57	0.31	0.33
	-.39	-.48	-.29	0.00	0.00	0.05	-.31	-.65
	-.31	-.36	0.27	0.50	0.00	0.51	0.38	-.18
	-.22	-.32	-.61	-.35	0.00	-.07	0.27	0.51

This can be calculated easily in matlab using the $[V, \Lambda] = \text{eig}(L)$ command.

The first eigenvalue of the Laplacian is always 0, up to numerical error.

The second eigenvector of the normalized Laplacian is a *very important* descriptor of a graph. In the example it is $v_2 = (0.38, 0.36, 0.18, 0.38, -.28, -.48, -.36, -.32)$ as read off the second column of the above chart.

- It tells us how to best cut the graph.
- It tells us how “best” to put all of the vertices on a single line (this sometimes works better for the unnormalized Laplacian, see below. There it is called the *Fiedler vector*.)

- We can set $S = \{v_i \in V \mid u_2(v_i) < 0\}$ and $T = V \setminus S$.

Then $S = \{a, b, c, d\}$ and $T = \{e, f, g, h\}$.

- Can sometimes do better by checking all possible cuts along v_2 (use any threshold, not only 0). Take one with best $\text{NCut}(S, T)$.

The third eigenvector can be useful too. It can be used (with the second eigenvector) to lay out the vertices in \mathbb{R}^2 , and can then be used to make a 4-way cut.

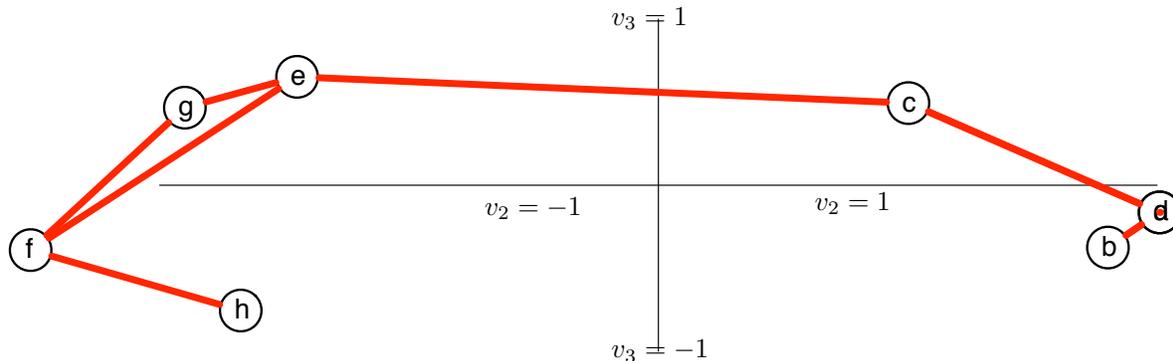
[++] $S = \{c\}$ defined as $v_2 > 0$ and $v_3 > 0$

[+-] $T = \{a, b, d\}$ defined as $v_2 > 0$ and $v_3 < 0$

[-+] $U = \{g, e\}$ defined as $v_2 < 0$ and $v_3 > 0$

[--] $R = \{f, h\}$ defined as $v_2 < 0$ and $v_3 < 0$.

When drawing the graph using v_2 and v_3 its good to scale the values by $1/\sqrt{\lambda_i}$ along each axis. Note that in the drawing below points a and d are directly on top of each other. From the perspective of the graph, they are indistinguishable. The eigenstructure does not separate them until v_5 .



Alternatively, we can use the first d eigenvectors (scaled by eigenvalues) to embed the vertices in \mathbb{R}^d . Then we can use any Euclidean clustering algorithm (such as Lloyds for k -means clustering). The smaller the eigenvalue, the more important the direction. So the larger the index of the eigenvalue, the smaller the $1/\sqrt{\lambda_i}$ will be. So the top 5 or so (depending on data) may be all required. Notice here how the second eigenvector provides much better separation than the third one.

This embedding into \mathbb{R}^2 is a form of *non-linear dimensionality reduction*.

Affinity matrix. More generally, the adjacency matrix need not be 0 – 1. It can be filled with the *similarity* value defined by some similarity between element; then A stands for *affinity*. The diagonal is defined as the sum of elements in a row (or column — it must be symmetric). Then spectral clustering can be run as before. When the similarity of a pair is very small, it is a good heuristic to round the values down to 0 in the matrix to make the algorithm run faster.

10.2.2 (Unnormalized) Laplacian

In some contexts it will be suitable to use the (regular, unnormalized) Laplacian defined:

$$L_0 = D - A = \begin{pmatrix} 3 & -1 & -1 & -1 & 0 & 0 & 0 & 0 \\ -1 & 2 & 0 & -1 & 0 & 0 & 0 & 0 \\ -1 & 0 & 3 & -1 & -1 & 0 & 0 & 0 \\ -1 & -1 & -1 & 3 & 0 & 0 & 0 & 0 \\ 0 & 0 & -1 & 0 & 3 & -1 & -1 & 0 \\ 0 & 0 & 0 & 0 & -1 & 3 & -1 & -1 \\ 0 & 0 & 0 & 0 & -1 & -1 & 2 & 0 \\ 0 & 0 & 0 & 0 & 0 & -1 & 0 & 1 \end{pmatrix}.$$

Note that the entries in each row and column of L_0 sum up to 0.

- think of D as the *flow* into a vertex, and
- think of A as the *flow* out of the vertex.

The water keeps flowing, so it does not get stuck anywhere. That is, as much flows in as flows out.

We can also convert L_0 to the normalized Laplacian L using the $D^{-1/2}$ matrix as

$$L = I - D^{-1/2}AD^{-1/2} = D^{-1/2}L_0D^{-1/2}.$$

The left- and right-multiplication by $D^{-1/2}$ can be thought of as normalizing by the degrees. That is each entry $P_{i,j}$ of $P = D^{-1/2}AD^{-1/2}$ (and edge (i,j)) is normalized by the amount of flow in and out of the nodes v_i and v_j of its corresponding edge (i,j) .

There are (typically) several eigenvectors of L_0 (the Laplacian). We list them here sorted by λ .

λ	0	0.278	1.11	2.31	3.46	4	4.82
V	$1/\sqrt{8}$	-.36	0.08	0.10	0.28	0.25	$1/\sqrt{2}$
	$1/\sqrt{8}$	-.42	0.18	0.64	-.38	0.25	0
	$1/\sqrt{8}$	-.20	-.11	0.61	0.03	-.25	0
	$1/\sqrt{8}$	-.36	0.08	0.10	0.28	0.25	$-1/\sqrt{2}$
	$1/\sqrt{8}$	0.17	-.37	0.21	-.54	-.25	0
	$1/\sqrt{8}$	0.36	-.08	-.10	-.28	0.75	0
	$1/\sqrt{8}$	0.31	-.51	-.36	-.56	0.56	0
	$1/\sqrt{8}$	0.50	0.73	0.08	0.11	0.11	0

And here is the drawing of the vertices according to v_2 and v_3 , scaled by $1/\sqrt{\lambda_i}$ along each axis. Again the drawing below points a and d are directly on top of each other. From the perspective of the graph, they are indistinguishable. The eigenstructure does not separate them until v_7 .

