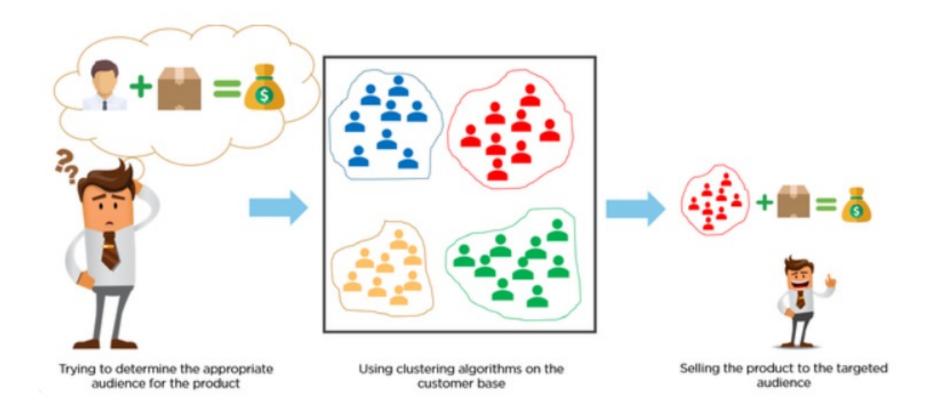
Clustering

ECE57000: Artificial Intelligence

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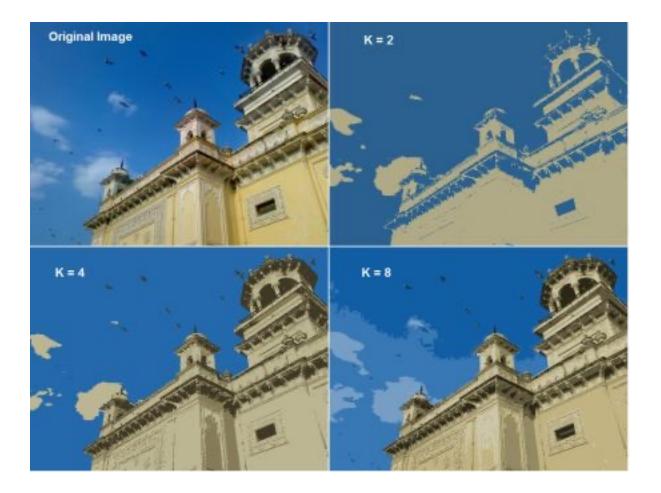
2021

Clustering application: Market segmentation to group customers



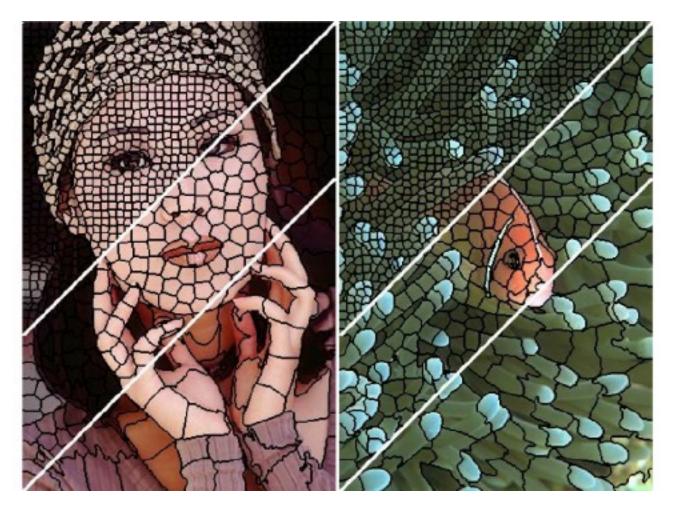
https://medium.com/analytics-vidhya/customer-segmentation-for-differentiatedtargeting-in-marketing-using-clustering-analysis-3ed0b883c18b

Clustering applications: Discretization of colors for compression



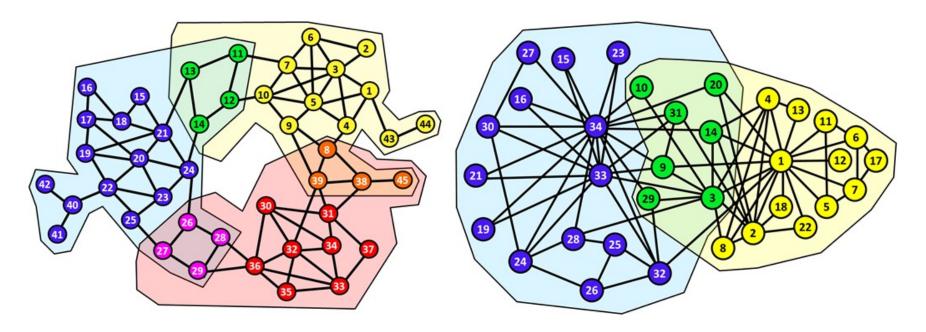
https://docs.opencv.org/3.4/d1/d5c/tutorial_py_kmeans_opencv.html

Clustering applications: Unsupervised image segmentation



R. Achanta, A. Shaji, K. Smith, A. Lucchi, P. Fua and S. Süsstrunk, "SLIC Superpixels Compared to State-of-the-Art Superpixel Methods," in *IEEE Transactions on Pattern Analysis and Machine Intelligence*, vol. 34, no. 11, pp. 2274-2282, Nov. 2012, doi: 10.1109/TPAMI.2012.120.

Another clustering application: Clustering people in social networks



Zachary's Karate Club Network

https://en.wikipedia.org/wiki/Zachary%27s_karate_club

https://bigdata.oden.utexas.edu/project/graph-clustering/

Outline

- Clustering applications
- K-means algorithm (Python demo)
- K-means relation to PCA
- Graph clustering
- Spectral clustering

The K-means algorithm is the most common clustering algorithm

(See demo of k-means algorithm and derivation)

Recap: Principal Component Analysis (PCA) can be formalized as minimizing the linear reconstruction error of the data using only $k \leq d$ dimensions

PCA can be formalized as

 $\min_{\mathbf{Z},\mathbf{W}} \| X_c - \mathbf{Z} \mathbf{W}^T \|_F^2$

▶ where

 $\begin{array}{l} X_{c} = X - \mathbf{1}_{n} \mu_{x}^{T} \in \mathbb{R}^{n \times d} \quad (\text{centered input data}) \\ Z \in \mathbb{R}^{n \times k} \quad (\text{latent representation or "scores"}) \\ W^{T} \in \mathbb{R}^{k \times d} \quad (\text{principal components}) \\ w_{s}^{T} w_{t} = 0, w_{s}^{T} w_{s} = \|w_{s}\|_{2} = 1, \forall s, t \\ (\text{orthogonal constraint}) \end{array}$

Solution

•
$$W^T = V_{1:k}^T$$
 where $X_c = USV^T$ is the SVD of X_c

$$\blacktriangleright Z = X_c W$$

K-means relation to PCA: One-hot vectors vs continuous vectors

- k-means clustering can be seen as reducing the dimensionality to k latent categories
 - Each category can be represented by a one-hot vector of length k
 - e.g., if $k = 3, z_i \in \{[1,0,0], [0,1,0], [0,0,1]\}, \forall i$
 - Every instance can only "belong" to one category
- In dimensionality reduction techniques, the latent vectors can have non-zeros for all k latent dimensions
 - ▶ e.g., if $k = 3, z_i \in \mathbb{R}^3, \forall i$

K-means objective can be reformulated as seeking the best approximation to X with low rank constraint (k < d)

Original k-means objective

 $\min_{\substack{\mathcal{C}_1, \dots, \mathcal{C}_k \\ \mu_1, \dots, \mu_k}} \sum_{j=1}^n \sum_{x \in \mathcal{C}_j} \left\| x - \mu_j \right\|_2^2$ • Equivalent to the following objective

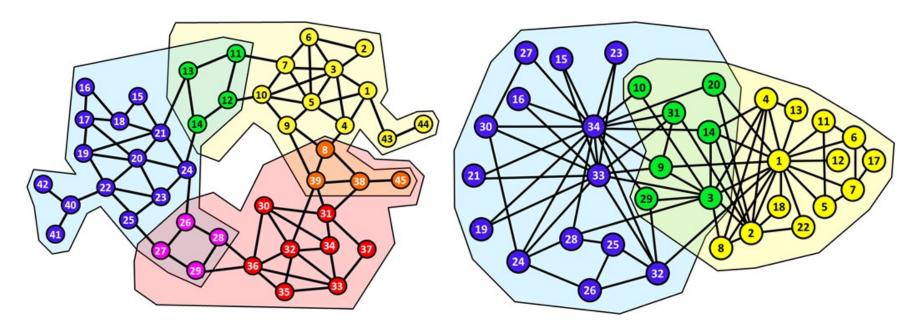
$$\begin{split} & \min_{Z,M} \|X - ZM\|_F^2 \\ & \text{where } Z \in \{0,1\}^{n \times k}, \sum_j z_{ij} = 1, \forall i \\ & \text{and } M \in \mathbb{R}^{k \times d} \end{split}$$

Notice the similarity and differences with PCA objective and constraints

Derivation of equivalence between two objectives for k-means

▶ $y_i \in \{1, ..., k\}$ is the cluster label for each instance • z_i is the corresponding one hot vector to y_i • $M = \begin{bmatrix} \mu_1 \\ \vdots \\ \mu_i \end{bmatrix}$ is the matrix of mean vectors $\sum_{j=1}^{k} \sum_{x \in C_{j}} \|x - \mu_{j}\|_{2}^{2}$ $= \sum_{i=1}^{n} \left\| x_i - \mu_{y_i} \right\|_{2}^{2}$ • = $\sum_{i=1}^{n} \left\| x_i^T - z_i^T M \right\|_2^2$ (row vector form) $= \sum_{i=1}^{n} \sum_{s=1}^{d} \left(x_{is} - \overline{z_i^T} m_s \right)^2 \quad (m_s \text{ is a column of } M)$ $= \left(\sqrt{\sum_{i=1}^{n} \sum_{s=1}^{d} \left(x_{is} - z_i^T m_s \right)^2} \right)^{-1}$ $\bullet = \|X - ZM\|_{F}^{2}$

How can we cluster the nodes of a network (a.k.a. a graph) instead of a set of points?



Zachary's Karate Club Network

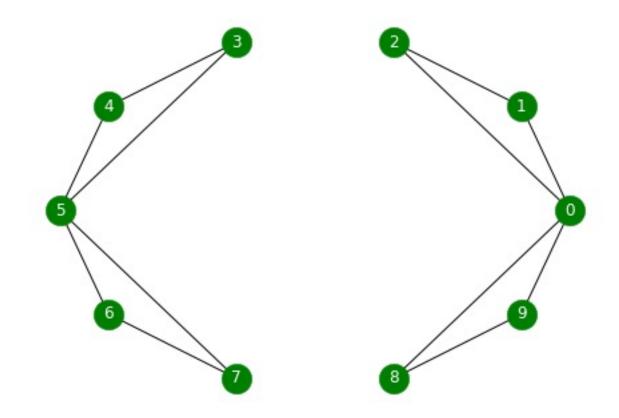
https://en.wikipedia.org/wiki/Zachary%27s_karate_club

https://bigdata.oden.utexas.edu/project/graph-clustering/

Graph clustering puts the nodes of a graph into clusters

- What is a graph?
- How do we represent a graph?
 - Adjacency matrix
 - Graph Laplacian
- How do we use graph Laplacian to cluster?

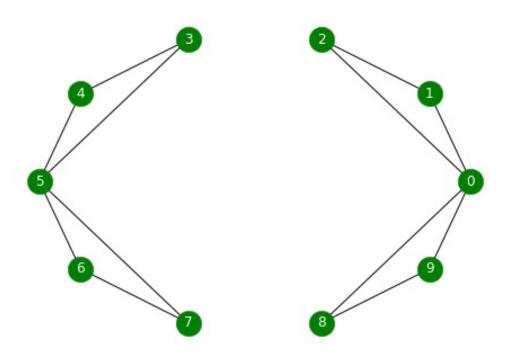
A graph/network is composed of nodes and weighted edges between the nodes

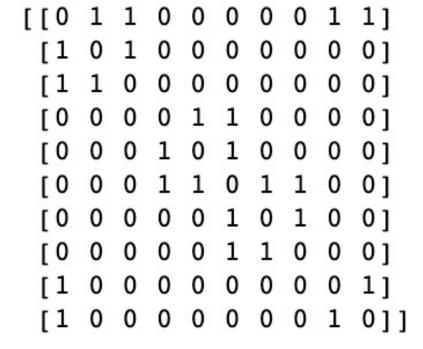


10 node graph with 2 disconnected components.

A graph can be represented as an **adjacency matrix**

- Nodes are represented by rows/columns
- Edges are encoded as 1s





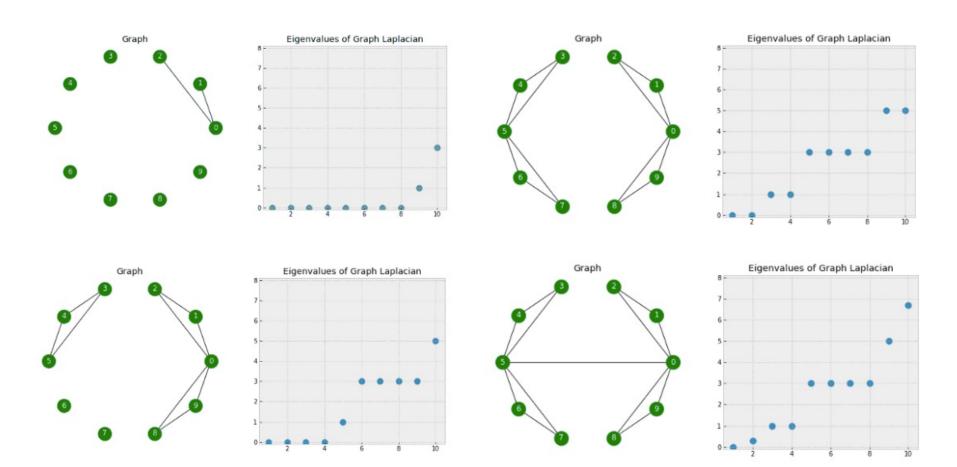
The **graph Laplacian** is formed by subtracting the adjacency from the degree matrix

The degree matrix is a diagonal matrix whose elements are the sum of the rows:
D = diag(A1)

• Graph Laplacian is defined as: L = D - A

[[4 -1 -	1 0 0 0	0 0 -1 -1]	[[4 0 0 0 0 0 0 0 0]	[[0 1 1 0 0 0 0 0 1 1]
[-1 2 -	1 0 0 0	0 0 0 0]	[0 2 0 0 0 0 0 0 0 0]	[1 0 1 0 0 0 0 0 0 0]
[-1 -1	2 0 0 0	0 0 0 0]	[0 0 2 0 0 0 0 0 0 0]	[1 1 0 0 0 0 0 0 0 0]
[0 0	0 2 -1 -1	0 0 0 0]	[0 0 0 2 0 0 0 0 0 0]	[0 0 0 0 1 1 0 0 0]
[0 0	0 -1 2 -1	0 0 0 0]	[0 0 0 0 2 0 0 0 0 0]	[0 0 0 1 0 1 0 0 0 0]
[0 0	0 -1 -1 4 -	-1 -1 0 0] —	[0 0 0 0 4 0 0 0 0]	[0 0 0 1 1 0 1 1 0 0]
[0 0	0 0 0 -1	2 -1 0 0]	[0 0 0 0 0 0 2 0 0 0]	[0 0 0 0 0 1 0 1 0 0]
[0 0	0 0 0 -1 -	-1 2 0 0]	[0 0 0 0 0 0 0 2 0 0]	[0 0 0 0 0 1 1 0 0 0]
[-1 0	0 0 0 0	0 0 2 -1]	[0 0 0 0 0 0 0 0 2 0]	[1 0 0 0 0 0 0 0 0 1]
[-1 0	0 0 0 0	0 0 -1 2]]	[0 0 0 0 0 0 0 0 0 2]]	[1 0 0 0 0 0 0 0 1 0]]

The number of 0 eigenvalues of the Laplacian is the number of connected components



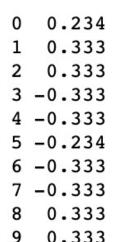
The Fiedler vector (2nd to last eigenvector) can be used to create 2 clusters

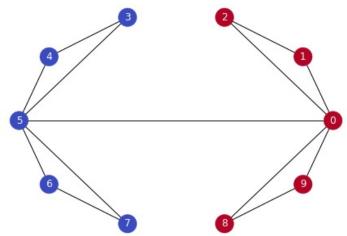
- Intuitively, we could 0 the 2nd to last eigenvalue to get 2 components instead
- Nodes are clustered based on whether their values in the Fiedler vector

y = 1(f > 0)

In theory, this is known as the <u>minimal cut</u>

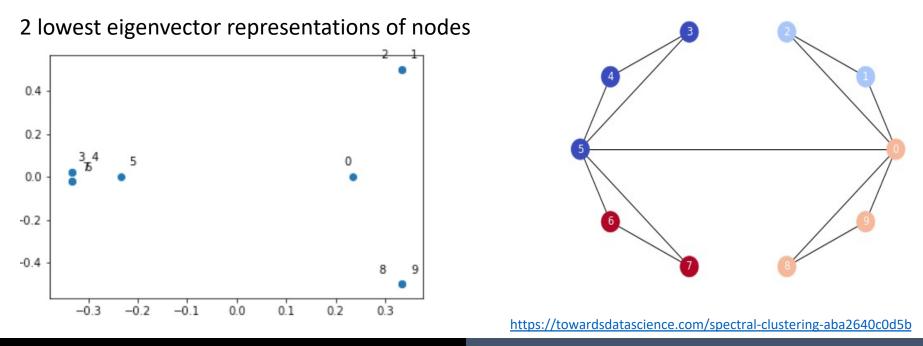






<u>Spectral clustering</u> generalizes to k > 2 clusters by taking the lowest eigenvectors as a new node representation and then doing K-means

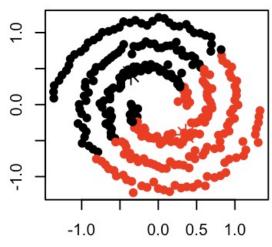
- We take the *m*-lowest eigenvectors to represent the data
- Then just run K-means



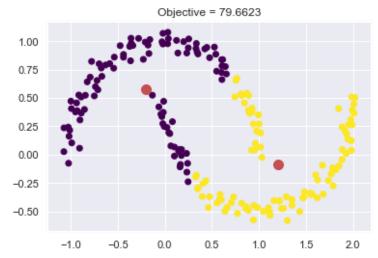
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Standard K-means clustering is limited to circular clusters with linear boundaries between clusters

- K-means is based on the clustering assumption of "compactness"
 - Points in a cluster are close to one another
 - Squared error objective within cluster
- This assumption may not be appropriate

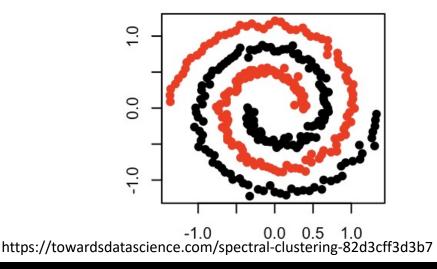


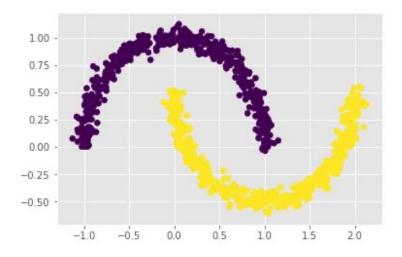




<u>Spectral clustering</u> applied to vector data can be used to learn clusters based on "connectivity"

- Points that are "connected" to each other are clustered together
- This allows non-circular clustering



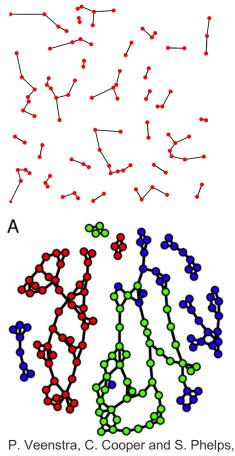


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Spectral clustering: First create similarity graph based on data

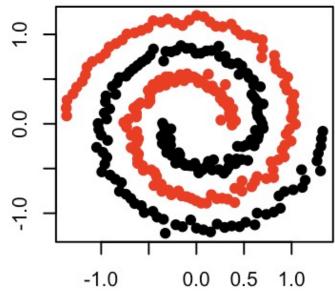
- K-nearest neighbor graph
 - Add edge for all k-nearest neighbors
- General similarity graph
 - Compute all pairwise similarity between points such as:

$$s(x,y) = \exp\left(-\frac{\|x-y\|^2}{2\sigma^2}\right)$$

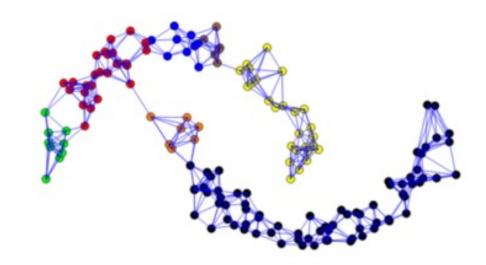


P. Veenstra, C. Cooper and S. Phelps, "Spectral clustering using the kNN-MST similarity graph," *2016 8th Computer Science and Electronic Engineering (CEEC)*, Colchester, 2016, pp. 222-227, doi: 10.1109/CEEC.2016.7835917.

Spectral clustering: Second, apply spectral clustering to resulting graph

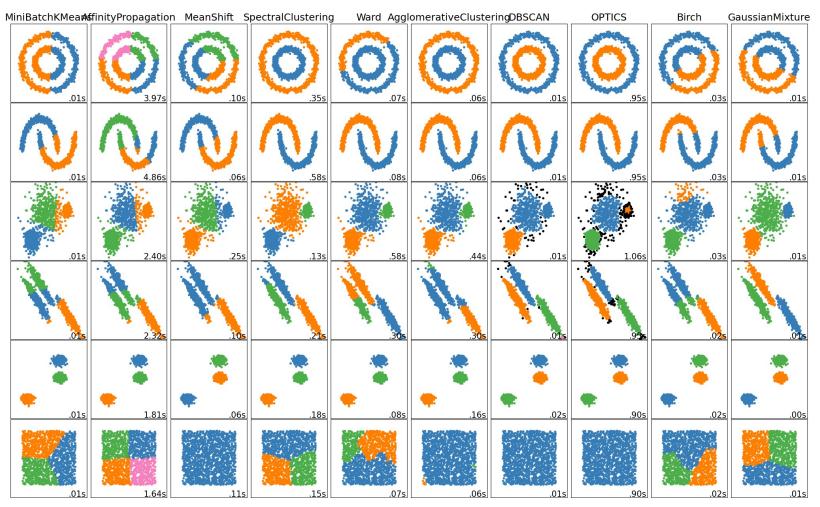


https://towardsdatascience.com/spectral-clustering-82d3cff3d3b7



http://math.ucdenver.edu/~sborgwardt/wiki/index.php/Spectral_clust ering

Many other clustering algorithms exist



https://scikit-learn.org/stable/modules/clustering.html

Simple demo of spectral clustering (time permitting)