

Spectral / K-means clustering

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Special Course in Computer and Information Science

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Outline

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- Summary



- *"Clustering is a process of organizing objects into groups whose members are similar in some way."*
- <u>Spectral clustering</u>: data points as nodes of a connected graph and clusters are found by partitioning this graph, based on its spectral decomposition, into subgraphs.
- <u>K-means clustering</u>: divide the objects into *k* clusters such that some metric relative to the centroids of the clusters is minimized.



Introduction

Spectral clustering

- How to define a graph for spectral methods?
- How to partition a graph into subgraphs?

K-means clustering

- How to choose K?
- Which metric to use?



By Ng, Jordan and Weiss

- Given a data set $S = \{s_1, \dots, s_n\}$ to be clustered
- 1. Calculate the affinity matrix $A_{ij} = \exp(-||s_i s_j||^2/2\sigma^2)$, if $i \neq j$ and $A_{ii} = 0$ where σ^2 is the scaling parameter
- 2. Define *D* to be the diagonal matrix whose (i,i)-element is the sum of *A*'s *i*-th row, and construct the matrix $L = D^{-1/2}AD^{-1/2}$
- 3. Find *k* largest eigenvectors of *L* and form the matrix $X = [x_1 x_2 \dots x_k]$
- 4. Form the matrix *Y* from *X* by normalizing each of *X*'s rows to have unit length, $Y_{ij} = X_{ij} / (\sum_j X_{ij}^2)^{1/2}$
- 5. Treating each row of Y as a point, cluster them into k clusters via K-means or any other algorithm
- 6. Assign the original point s_i to cluster j if and only if row i of the matrix Y was assigned to cluster j



Spectral clustering

Main difference between algorithms is the definition of *L* Meila & Shi: *L* = *D*⁻¹*A*, use largest eigenvectors, no normalization
Shi & Malik: *L* = *D* - *A*, use smallest eigenvectors, no normalization



Graph of two groups...can you believe?



The adjacency matrix



The second smallest eigenvector, unsorted



The second smallest eigenvector is the Fiedler vector, i.e. algebraic connectivity



The second smallest eigenvector, sorted





Permutated graph...looks like there is two groups



~0.5 s



K-means clustering / Algorithm

Standard version

- Given an integer k and a set of n data points $X \subset \mathbb{R}^d$
- 1. Arbitrarily choose initial *k* centers $C = \{c_1, c_2, ..., c_k\}, C \in \mathbb{R}^d$
- 2. For each $i \in \{1,...,k\}$, set the cluster C_i to be the set of points X that are closer to c_i than they are to c_j for all $j \neq i$
- 3. For each $i \in \{1,...,k\}$, set c_i to be the center of mass of all points in C_i , i.e. $c_i = (1/|C_i|) \sum_{x \in C_i} x$
- 4. Repeat Steps 2 and 3 until *C* no longer changes



- Standard practice to choose the initial centers uniformly at random from *X*
- For Step 2, ties may be broken arbitrarily, as long as the method is consistent
- Steps 2 and 3 guarantee to decrease the intra-cluster variance, i.e. to minimize the potential function $\phi = \sum_{x \in X} \min_{c \in C} ||x c||^2$, until it is no longer possible to do so
- Some extension:
 - Different ways to choose *k* initial centers, e.g. K-means++
 - Force the center point of each cluster to be one of the actual points, i.e. K-medoids
- All in all: fast, simple, no approximation guarantees



K-means clustering / Example

Two circles, originally





Two circles, K-means clustered (standard Matlab kmeans)



~0.01 s



Summary

- If successful, algorithm by Ng, Jordan & Weiss will dramatically improve the results of the standard K-means
- Tip: eigs(xxx,xxx,'opt'); , where opt has essential role
- Remember to normalize *Y*



References

Andrew Y. Ng, Michael I. Jordan, and Yair Weiss. *On spectral clustering: Analysis and an algorithm*. In: Advances in Neural Information Processing Systems 14, 2002.

D. Arthur, S. Vassilvitskii. k-means++: *The Advantages of Careful Seeding*. Symposium on Discrete Algorithms (SODA), 2007.