Fast Approximate Spectral Clustering

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Algorithm 1. NJW spectral clustering algorithm

Input: Dataset $S = \{x_1, ..., x_n\}$ in \Re^1 and the number of clusters k

Output: *k*-way partition of the input data

(1) Construct the affinity matrix *A* by the following Gaussian kernel function:

$$A_{ij} = \begin{cases} \exp(\frac{-\|\mathbf{x}_i - \mathbf{x}_j\|^2}{\delta^2}) & \text{for } i \neq j, \\ 0 & \text{for } i = j, \end{cases}$$
 (1)

where δ is a scale parameter to control how fast the similarity attenuates with the distance between the data points x_i and x_j . (2) Compute the normalized affinity matrix $\mathbf{L} = \mathbf{D}^{-1/2}\mathbf{A} \ \mathbf{D}^{-1/2}$,

where \mathbf{D} is the diagonal matrix with $D_{ii} = \sum_{j=1}^{n} A_{ij}$.

- (3) Compute the k eigenvectors of L, v₁, v₂, ..., v_k, which are associated with the k largest eigenvalues, and form the matrix X = [v₁v₂, ..., v_k].
- (4) Renormalize each row to form a new matrix $Y \in \mathfrak{R}^{n \times k}$ with $Y_{ij} = X_{ij}/(\sum_j X_{ij}^2)^{1/2}$, so that each row of **Y** has a unit magnitude.
- (5) Treat each row of Y as a point in ℜ^k and partition the n points (n rows) into k clusters via a general cluster algorithm, such as the K-means algorithm.
- (6) Assign the original point x_i to the cluster c if and only if the corresponding row i of the matrix Y is assigned to the cluster c.







KASP

- ► Create a representative set
- Spectral cluster representative set
- Assign labels to all original data







Algorithm KASP $(\mathbf{x}_1, \dots, \mathbf{x}_n, k)$

Input: n data points $\{\mathbf{x}_i\}_{i=1}^n$, number of representative points k

Output: m-way partition of the input data

- 1. Perform k-means with k clusters on x_1, \ldots, x_n to:
 - a) Compute the cluster centroids y_1, \dots, y_k as the k representative points.
 - b) Build a correspondence table to associate each x_i with the nearest cluster centroid y_j .
- 2. Run a spectral clustering algorithm on y_1, \dots, y_k to obtain an m-way cluster membership for each of y_i .
- 3. Recover the cluster membership for each \mathbf{x}_i by looking up the cluster membership of the corresponding centroid \mathbf{y}_j in the correspondence table.





Perturbation Analysis

Assume x_1, \ldots, x_n i.i.d according to a probability distribution G

$$\tilde{x}_i = x_i + \epsilon_i \tag{1}$$

 \tilde{x} distributed by \tilde{G} .

- $ightharpoonup \epsilon_i$ independent of x_i
- $ightharpoonup \epsilon_i$ are i.i.d. according to a symmetric dist (mean zero, bounded support)
- ▶ $Var(\epsilon)$ small relative to Var(X).







Mis-clustering Rate

$$\rho = \frac{1}{n} \sum_{i=1}^{n} \mathbb{I}(I_i \neq \tilde{I}_i)$$
 (2)





Theorem

Under the assumptions(*), the mis-clustering rate ρ of a spectral bi-partitioning algorithm on the perturbed data satisfies

$$\rho \le \|\tilde{\mathbf{v}}_2 - \mathbf{v}_2\|^2 \tag{3}$$







Lemma

Let g denote the eigengap between the second and the third eigenvalues of L. Then the following holds:

$$\|\tilde{v}_2 - v_2\| \le \frac{1}{g} \|\tilde{L} - L\| + O(\|\tilde{L} - L\|^2).$$
 (4)





Theorem

Assume assumptions hold throughout recursive invocation of the Ncut algorithm, g_0 is bounded away from zero and Frobenius norm of perturbation of Laplacian matricies along the recursion is bounded by $c\|\tilde{L}-L\|_F^2$ for some constant $c\geq 1$. Then the mis-clustering rate for an m-way spectral clustering solution can be bounded by:

$$\rho \le \|\frac{m}{g_0^2} \cdot c\|\tilde{L} - L\|_F^2. \tag{5}$$



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$$r_1 \le n \cdot \frac{L_1^2}{g_1^2}$$

 $r_i \le (n - n_{i-1}) \cdot \frac{L_i^2}{g_i^2}, i = 2, \dots, m - 1.$





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Linking Laplacian to ϵ

Theorem

$$\|\tilde{L} - L\|_F^2 \le_p c_1 \sigma_{\epsilon}^{(2)} + c_2 \sigma_{\epsilon}^{(4)} \tag{6}$$







If we know the probability distribution of the original data, it is possible to characterise the exact amount of distortion.

f is the density function of G

Theorem

Let data be distributed with density f.

$$\rho = c \cdot b_{2,d} \cdot ||f||_{\frac{d}{d+2}} \cdot k^{-\frac{2}{d}} + O(k^{-\frac{4}{d}})$$

where c is constant determined by number of clusters, variance of original data, bandwidth of Gaussian kernel and the minimum eigengap of all affinity matrices used in Ncut.





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FAIL





















