



The k -means Lecture..

The Kernel Matrix

Defining for $D \in \mathbb{R}^{n \times d}$ the row-wise applied feature transformation

$$\phi(D) = \begin{pmatrix} \text{---} & \phi(D_{1\cdot}) & \text{---} \\ & \vdots & \\ \text{---} & \phi(D_{n\cdot}) & \text{---} \end{pmatrix},$$

the kernel matrix is given by

$$K = \phi(D)\phi(D)^\top \in \mathbb{R}^{n \times n}.$$

The Kernel k -means Objective

Given: a data matrix $D \in \mathbb{R}^{n \times d}$, a feature transformation $\phi : \mathbb{R}^d \rightarrow \mathbb{R}^p$ mapping into a p -dimensional feature space, where $p \in \mathbb{N} \cup \{\infty\}$, and the number of clusters r .

Find: clusters indicated by the matrix $Y \in \mathbb{1}^{n \times r}$ which minimize the within cluster scatter in the transformed feature space

$$\min_Y \|\phi(D) - YX^\top\|^2 \text{ s.t. } X = \phi(D^\top)Y(Y^\top Y)^{-1}, Y \in \mathbb{1}^{n \times r} \quad (4)$$

3

Optimization

If we want to apply the **kernel trick**, then we need to state the kernel k -means objective with respect to the **inner product of data points**.

Representing Data by the Inner Product Only

Theorem (*k*-means trace objective)

The *k*-means objective in Eq. (1) is equivalent to

$$\max_Y \operatorname{tr}(Z^\top D D^\top Z) \quad \text{s.t. } Z = Y(Y^\top Y)^{-1/2}, Y \in \mathbb{1}^{n \times r} \quad (5)$$

Interpretation: Clusters are now defined with respect to the inner product similarity:

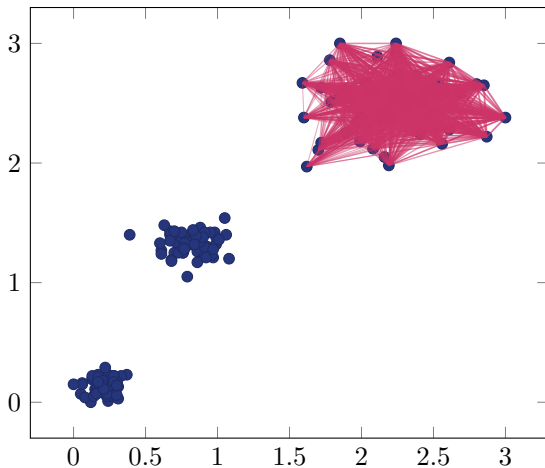
$$\operatorname{sim}(i, j) = D_i \cdot D_j^\top = \cos(\angle(D_{i\cdot}, D_{j\cdot})) \|D_{i\cdot}\| \|D_{j\cdot}\|$$

Points within one cluster need to be similar:

$$\operatorname{tr}(Z^\top D D^\top Z) = \sum_{s=1}^r \frac{Y_{\cdot s}^\top D D^\top Y_{\cdot s}}{|Y_{\cdot s}|} = \sum_{s=1}^r \frac{1}{|C_s|} \sum_{i, j \in C_s} D_i \cdot D_j^\top$$

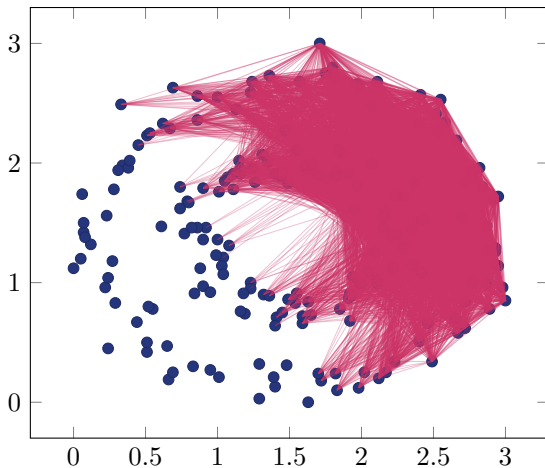


The Inner Product Similarity and Convex Clusters





The Inner Product Similarity and Nonconvex Clusters



Kernel k -means

Theorem (Equivalent kernel k -means objectives)

Given the kernel matrix $K = \phi(D)\phi(D)^\top$, the following objectives are equivalent:

$$\min_Y \|\phi(D) - YX^\top\|^2 \quad \text{s.t.} \quad X = \phi(D^\top)Y(Y^\top Y)^{-1}, Y \in \mathbb{1}^{n \times r} \quad (6)$$

$$\max_Y \text{tr}(Z^\top KZ) \quad \text{s.t.} \quad Z = Y(Y^\top Y)^{-1/2}, Y \in \mathbb{1}^{n \times r} \quad (7)$$

Problem: We do not know how to optimize Eq. (7), we only know how to optimize Eq. (6), but we do not want to compute ϕ !

Idea: We go the other way round: from the kernel matrix to the inner product.

Eigendecomposition of Symmetric Matrices

Theorem (Eigendecomposition of symmetric matrices)

For every symmetric matrix $K = K^T \in \mathbb{R}^{n \times n}$ there exists an orthogonal matrix $V \in \mathbb{R}^{n \times n}$ and a diagonal matrix $\Lambda = \text{diag}(\lambda_1, \dots, \lambda_n)$ where $|\lambda_1| \geq \dots \geq |\lambda_n|$ such that

$$K = V\Lambda V^T$$

Every symmetric matrix $K \in \mathbb{R}^{n \times n}$ has a **symmetric decomposition** $K = A^T A$ if and only if the **eigenvalues of K are nonnegative**. This is equivalent to K being **positive semi-definite**.

Kernel matrices are positive semi-definite!

Kernel k -means Inside Out

Theorem (Equivalent kernel k -means objectives)

Given a kernel matrix and its symmetric decomposition $K = AA^T$, the following objectives are equivalent:

$$\min_Y \|A - YX^T\|^2 \quad \text{s.t. } X = A^T Y(Y^T Y)^{-1}, Y \in \mathbb{1}^{n \times r} \quad (8)$$

$$\max_Y \text{tr}(Z^T K Z) \quad \text{s.t. } Z = Y(Y^T Y)^{-1/2}, Y \in \mathbb{1}^{n \times r} \quad (9)$$

Algorithm Idea: Use the objective in Eq. (8): compute a symmetric decomposition $AA^T = K$ by means of the eigendecomposition $A = V\Lambda^{1/2}$ and run k -means on A .

The Kernel k -means Algorithm

```
1: function KERNELKMEANS( $r, K$ )
2:    $(V, \Lambda) \leftarrow$  EIGENDECOMPOSITION( $K$ )
3:    $A \leftarrow V\Lambda^{1/2}$ 
4:    $(X, Y) \leftarrow$  KMEANS( $A, r$ )
5:   return  $Y$ 
6: end function
```

$$\triangleright AA^T = K$$

Let's try this kernel k -means idea on the two circles dataset.

Ok, so **in theory** we have a **method** to solve kernel k -means, but **in practice** this method is not often employed.

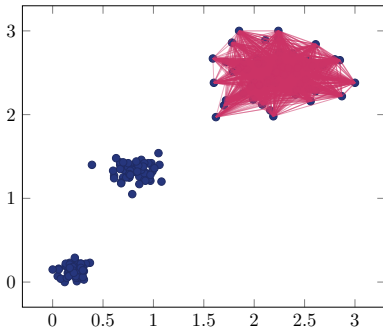
Drawbacks of kernel k -means is a **lack of robustness** and the requirement of a **full eigendecomposition**.

A related method based on a graph representation of the data facilitates nonconvex clustering based on a **truncated eigendecomposition**.

1

Informal Problem Description

Interpretation of the Data as a Graph

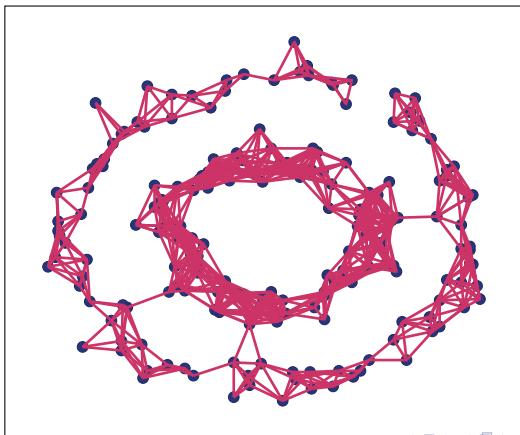


Every data **point** is a **node**.

The weight of an **edge** reflects the **similarity** between connected nodes.

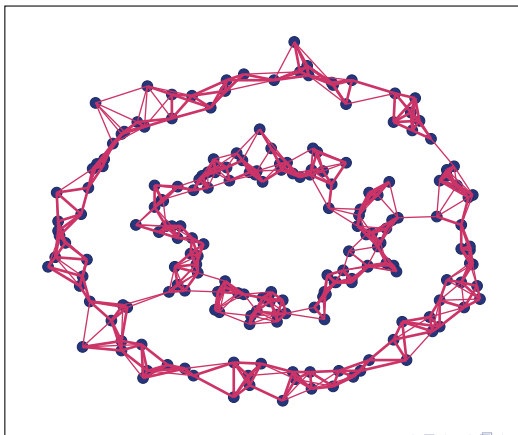
Similarity Measures: Epsilon Neighborhood

$$W_{ij} = \begin{cases} 1 & \text{if } \|D_{i\cdot} - D_{j\cdot}\| < \epsilon \\ 0 & \text{otherwise} \end{cases}$$



Similarity Measures: K-nearest neighbors (K=5)

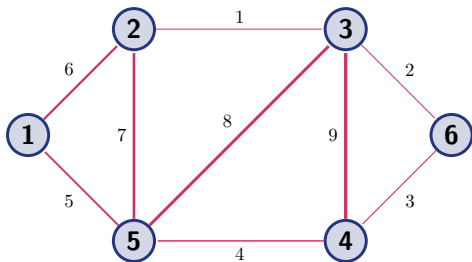
$$N_{ij} = \begin{cases} 1 & \text{if } D_i \in KNN(D_j) \\ 0 & \text{otherwise} \end{cases}, \quad W = \frac{1}{2}(N + N^T)$$



2

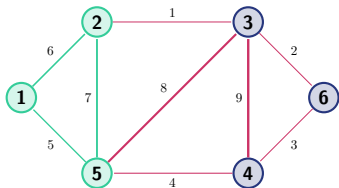
Derive the Formal Problem Definition

The Weighted Adjacency Matrix



$$W = \begin{pmatrix} 0 & 6 & 0 & 0 & 5 & 0 \\ 6 & 0 & 1 & 0 & 7 & 0 \\ 0 & 1 & 0 & 9 & 8 & 2 \\ 0 & 0 & 9 & 0 & 4 & 3 \\ 5 & 7 & 8 & 4 & 0 & 0 \\ 0 & 0 & 2 & 3 & 0 & 0 \end{pmatrix}$$

Computing the Similarity Within a Cluster



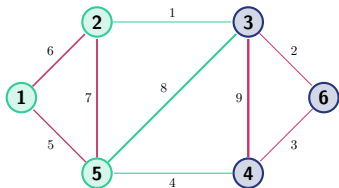
$$Y_{\cdot s}^{\top} = (1 \ 1 \ 0 \ 0 \ 1 \ 0)$$

$$Y_{\cdot s}^{\top} W Y_{\cdot s} = 2(5 + 6 + 7)$$

$$\text{Sim}(Y; W) = \text{tr}(Y^{\top} W Y (Y^{\top} Y)^{-1})$$

$$= \sum_{s=1}^r \frac{Y_{\cdot s}^{\top} W Y_{\cdot s}}{|Y_{\cdot s}|} = \sum_{s=1}^r \frac{1}{|C_s|} \sum_{i,j \in C_s} W_{ji}$$

Computing the Cut of a Cluster



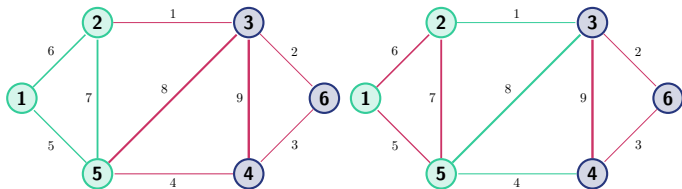
$$Y_{\cdot s}^\top = (1 \quad 1 \quad 0 \quad 0 \quad 1 \quad 0)$$

$$Y_{\cdot s}^\top W(1 - Y_{\cdot s}) = 1 + 8 + 4$$

$$\text{Cut}(Y; W) = \text{tr}((1 - Y)^\top WY(Y^\top Y)^{-1})$$

$$= \sum_{s=1}^r \frac{(1 - Y_{\cdot s})^\top WY_{\cdot s}}{|Y_{\cdot s}|} = \sum_{s=1}^r \frac{1}{|C_s|} \sum_{i \notin C_s} \sum_{j \in C_s} W_{ij}$$

Maximum Similarity vs. Minimum Cut



There are principally two ways to define clusters of graphs:

- 1 maximize the sum of weights **within** clusters
- 2 minimize the sum of weights **between** clusters

Maximum Similarity Graph Clustering

Given: a graph indicated by a symmetric, nonnegative similarity matrix $W \in \mathbb{R}_+^{n \times n}$, and the number of clusters r .

Find: clusters indicated by the matrix $Y \in \mathbb{1}^{n \times r}$ which maximize the similarity of points within a cluster

$$\max_Y \text{Sim}(Y; W) = \text{tr}(Y^\top W Y (Y^\top Y)^{-1}) \quad \text{s.t. } Y \in \mathbb{1}^{n \times r}$$

Minimum Cut Graph Clustering

Given: a graph indicated by a symmetric, nonnegative similarity matrix $W \in \mathbb{R}_+^{n \times n}$, and the number of clusters r .

Find: clusters indicated by the matrix $Y \in \mathbb{1}^{n \times r}$ which minimize the cut of all clusters

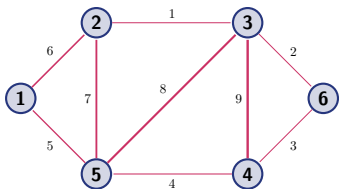
$$\min_Y \text{Cut}(Y; W) = \text{tr}((1 - Y)^\top W Y (Y^\top Y)^{-1}) \quad \text{s.t. } Y \in \mathbb{1}^{n \times r}$$

3

Optimization

The Degree Matrix

We have $Y_{.s}^T W Y_{.s} \leq Y_{.s}^T I_W Y_{.s}$ where I_W is the degree matrix:



$$I_W = \begin{pmatrix} 11 & 0 & 0 & 0 & 0 & 0 \\ 0 & 14 & 0 & 0 & 0 & 0 \\ 0 & 0 & 20 & 0 & 0 & 0 \\ 0 & 0 & 0 & 16 & 0 & 0 \\ 0 & 0 & 0 & 0 & 24 & 0 \\ 0 & 0 & 0 & 0 & 0 & 5 \end{pmatrix}$$

$Y_{.s}^T W Y_{.s} = Y_{.s}^T I_W Y_{.s}$ if and only if $Y_{.s}$ indicates a connected component. This is equivalent to

$$Y_{.s}^T \underbrace{(I_W - W)}_{=L} Y_{.s} = 0$$

The matrix $L = I_W - W$ is called **graph Laplacian**.

Relation of Minimum Cut and Maximum Similarity

Theorem (Minimum Cut and Maximum Similarity)

Given a symmetric similarity matrix $W \in \mathbb{R}_+^{n \times n}$, the degree matrix I_W and the Graph Laplacian $L = I_W - W$, then the following objectives are equivalent:

$$\min_Y \text{Cut}(Y; W) = \text{tr}((1 - Y)^T W Y (Y^T Y)^{-1}) \quad \text{s.t. } Y \in \mathbb{1}^{n \times r}$$

$$\max_Y \text{Sim}(Y; -L) = \text{tr}(Y^T (-L) Y (Y^T Y)^{-1}) \quad \text{s.t. } Y \in \mathbb{1}^{n \times r}$$

The **maximum similarity** objective is equal to the **kernel k -means** objective. However, note that $-L$ is not a kernel matrix (it's negative semi-definite).

The Spectral Clustering Algorithm

Requirement

The parameters of the similarity measure should be chosen such that the graph is connected!

- 1: **function** SPECTRALCLUSTERING(r, D, SIM)
- 2: $W \leftarrow \text{SIM}(D)$ ▷ Compute Similarity matrix
- 3: $L \leftarrow I_W - W$ ▷ Compute Graph Laplacian
- 4: $(V, \Lambda) \leftarrow \text{TRUNCATED EIGENDECOMPOSITION}(L, r + 1)$
- 5: $A \leftarrow V_{\cdot\{2, \dots, r+1\}}$ ▷ Remove connected component
- 6: $(X, Y) \leftarrow \text{KMEANS}(A, r)$
- 7: **return** Y
- 8: **end function**

Spectral Clustering with 10NN Similarity Matrix and L_{sym}

In practice, the weighted adjacency matrix is often normalized.
The corresponding Graph Laplacian is often denoted by

$$L_{sym} = I - I_W^{-1/2} W I_W^{-1/2}$$

Spect. Clustering

