Data Mining and Machine Learning: Fundamental Concepts and Algorithms dataminingbook.info

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Chapter 5: Kernel Methods

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For mining and analysis, it is important to find a suitable data representation. For example, for complex data such as text, sequences, images, and so on, we must typically extract or construct a set of attributes or features, so that we can represent the data instances as multivariate vectors.

Given a data instance x (e.g., a sequence), we need to find a mapping ϕ , so that $\phi(x)$ is the vector representation of x.

Even when the input data is a numeric data matrix a nonlinear mapping ϕ may be used to discover nonlinear relationships.

The term *input space* refers to the data space for the input data \mathbf{x} and *feature* space refers to the space of mapped vectors $\phi(\mathbf{x})$.

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Consider a dataset of DNA sequences over the alphabet $\Sigma = \{A, C, G, T\}$.

One simple feature space is to represent each sequence in terms of the probability distribution over symbols in Σ . That is, given a sequence \mathbf{x} with length $|\mathbf{x}| = m$, the mapping into feature space is given as

$$\phi(\mathbf{x}) = \{P(A), P(C), P(G), P(T)\}$$

where $P(s) = \frac{n_s}{m}$ is the probability of observing symbol $s \in \Sigma$, and n_s is the number of times s appears in sequence x.

For example, if $\mathbf{x} = ACAGCAGTA$, with $m = |\mathbf{x}| = 9$, since A occurs four times, C and G occur twice, and T occurs once, we have

$$\phi(\mathbf{x}) = (4/9, 2/9, 2/9, 1/9) = (0.44, 0.22, 0.22, 0.11)$$

We can compute larger feature spaces by considering, for example, the probability distribution over all substrings or words of size up to k over the alphabet Σ .

Consider the mapping ϕ that takes as input a vector $\mathbf{x} = (x_1, x_2)^T \in \mathbb{R}^2$ and maps it to a "quadratic" feature space via the nonlinear mapping

$$\phi(\mathbf{x}) = (x_1^2, x_2^2, \sqrt{2}x_1x_2)^T \in \mathbb{R}^3$$

For example, the point $\mathbf{x} = (5.9,3)^T$ is mapped to the vector

$$\phi(\mathbf{x}) = (5.9^2, 3^2, \sqrt{2} \cdot 5.9 \cdot 3)^T = (34.81, 9, 25.03)^T$$

We can then apply well-known linear analysis methods in the feature space.

Kernel Method

Let \mathcal{I} denote the input space, which can comprise any arbitrary set of objects, and let $D = \{x_i\}_{i=1}^n \subset \mathcal{I}$ be a dataset comprising *n* objects in the input space. Let $\phi: \mathcal{I} \to \mathcal{F}$ be a mapping from the input space \mathcal{I} to the feature space \mathcal{F} .

Kernel methods avoid explicitly transforming each point x in the input space into the mapped point $\phi(x)$ in the feature space. Instead, the input objects are represented via their pairwise similarity values comprising the $n \times n$ kernel matrix, defined as

$$\boldsymbol{K} = \begin{pmatrix} \mathcal{K}(\boldsymbol{x}_1, \boldsymbol{x}_1) & \mathcal{K}(\boldsymbol{x}_1, \boldsymbol{x}_2) & \cdots & \mathcal{K}(\boldsymbol{x}_1, \boldsymbol{x}_n) \\ \mathcal{K}(\boldsymbol{x}_2, \boldsymbol{x}_1) & \mathcal{K}(\boldsymbol{x}_2, \boldsymbol{x}_2) & \cdots & \mathcal{K}(\boldsymbol{x}_2, \boldsymbol{x}_n) \\ \vdots & \vdots & \ddots & \vdots \\ \mathcal{K}(\boldsymbol{x}_n, \boldsymbol{x}_1) & \mathcal{K}(\boldsymbol{x}_n, \boldsymbol{x}_2) & \cdots & \mathcal{K}(\boldsymbol{x}_n, \boldsymbol{x}_n) \end{pmatrix}$$

 $K:\mathcal{I}\times\mathcal{I}\to\mathbb{R}$ is a *kernel function* on any two points in input space, which should satisfy the condition

$$\mathcal{K}(\boldsymbol{x}_i, \boldsymbol{x}_j) = \phi(\boldsymbol{x}_i)^T \phi(\boldsymbol{x}_j)$$

Intuitively, we need to be able to compute the value of the dot product using the original input representation x, without having recourse to the mapping $\phi(x)$.

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Linear Kernel

Let $\phi(\mathbf{x}) \rightarrow \mathbf{x}$ be the *identity kernel*. This leads to the *linear kernel*, which is simply the dot product between two input vectors:

$$\phi(\boldsymbol{x})^{\mathsf{T}}\phi(\boldsymbol{y}) = \boldsymbol{x}^{\mathsf{T}}\boldsymbol{y} = K(\boldsymbol{x},\boldsymbol{y})$$

For example, if $\boldsymbol{x}_1 = \begin{pmatrix} 5.9 & 3 \end{pmatrix}^T$ and $\boldsymbol{x}_2 = \begin{pmatrix} 6.9 & 3.1 \end{pmatrix}^T$, then we have

 $K(\mathbf{x}_1, \mathbf{x}_2) = \mathbf{x}_1^T \mathbf{x}_2 = 5.9 \times 6.9 + 3 \times 3.1 = 40.71 + 9.3 = 50.01$



Many data mining methods can be *kernelized* that is, instead of mapping the input points into feature space, the data can be represented via the $n \times n$ kernel matrix K, and all relevant analysis can be performed over K.

This is done via the *kernel trick*, that is, show that the analysis task requires only dot products $\phi(\mathbf{x}_i)^T \phi(\mathbf{x}_j)$ in feature space, which can be replaced by the corresponding kernel $K(\mathbf{x}_i, \mathbf{x}_j) = \phi(\mathbf{x}_i)^T \phi(\mathbf{x}_j)$ that can be computed efficiently in input space.

Once the kernel matrix has been computed, we no longer even need the input points x_i , as all operations involving only dot products in the feature space can be performed over the $n \times n$ kernel matrix K.

A function K is called a **positive semidefinite kernel** if and only if it is symmetric:

$$K(\boldsymbol{x}_i, \boldsymbol{x}_j) = K(\boldsymbol{x}_j, \boldsymbol{x}_i)$$

and the corresponding kernel matrix \pmb{K} for any subset $\pmb{D} \subset \mathcal{I}$ is positive semidefinite, that is,

$$\boldsymbol{a}^{\mathsf{T}}\boldsymbol{K}\boldsymbol{a} \geq 0, \text{ for all vectors } \boldsymbol{a} \in \mathbb{R}^{n}$$

which implies that

$$\sum_{i=1}^{n}\sum_{j=1}^{n}a_{i}a_{j}\mathcal{K}(\boldsymbol{x}_{i},\boldsymbol{x}_{j})\geq0,\text{ for all }a_{i}\in\mathbb{R},i\in[1,n]$$

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Dot Products and Positive Semi-definite Kernels

Positive Semidefinite Kernel

If $K(\mathbf{x}_i, \mathbf{x}_j)$ represents the dot product $\phi(\mathbf{x}_i)^T \phi(\mathbf{x}_j)$ in some feature space, then K is a positive semidefinite kernel.

First, K is symmetric since the dot product is symmetric, which also implies that K is symmetric.

Second, K is positive semidefinite because

$$\mathbf{a}^{T} \mathbf{K} \mathbf{a} = \sum_{i=1}^{n} \sum_{j=1}^{n} a_{i} a_{j} \mathbf{K}(\mathbf{x}_{i}, \mathbf{x}_{j})$$
$$= \sum_{i=1}^{n} \sum_{j=1}^{n} a_{i} a_{j} \phi(\mathbf{x}_{i})^{T} \phi(\mathbf{x}_{j})$$
$$= \left(\sum_{i=1}^{n} a_{i} \phi(\mathbf{x}_{i})\right)^{T} \left(\sum_{j=1}^{n} a_{j} \phi(\mathbf{x}_{j})\right)$$
$$= \left\|\sum_{i=1}^{n} a_{i} \phi(\mathbf{x}_{i})\right\|^{2} \ge 0$$

Empirical Kernel Map

We now show that if we are given a positive semidefinite kernel $K: \mathcal{I} \times \mathcal{I} \to \mathbb{R}$, then it corresponds to a dot product in some feature space \mathcal{F} .

Define the map ϕ as follows:

$$\phi(\mathbf{x}) = \left(\left(K(\mathbf{x}_1, \mathbf{x}), K(\mathbf{x}_2, \mathbf{x}), \dots, K(\mathbf{x}_n, \mathbf{x}) \right)^T \in \mathbb{R}^n \right)$$

The empirical kernel map is defined as

$$\phi(\mathbf{x}) = \mathbf{K}^{-1/2} \cdot \left((K(\mathbf{x}_1, \mathbf{x}), K(\mathbf{x}_2, \mathbf{x}), \dots, K(\mathbf{x}_n, \mathbf{x})) \right)^T \in \mathbb{R}^n$$

so that the dot product yields

$$\phi(\mathbf{x}_i)^T \phi(\mathbf{x}_j) = \left(\mathbf{K}^{-1/2} \, \mathbf{K}_i \right)^T \left(\mathbf{K}^{-1/2} \, \mathbf{K}_j \right)$$
$$= \mathbf{K}_i^T \left(\mathbf{K}^{-1/2} \mathbf{K}^{-1/2} \right) \mathbf{K}_j$$
$$= \mathbf{K}_i^T \, \mathbf{K}^{-1} \, \mathbf{K}_j$$

where K_i is the *i*th column of K. Over all pairs of mapped points, we have

$$\left\{\boldsymbol{K}_{i}^{T}\boldsymbol{K}^{-1}\boldsymbol{K}_{j}\right\}_{i,j=1}^{n} = \boldsymbol{K}\boldsymbol{K}^{-1}\boldsymbol{K} = \boldsymbol{K}$$

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The Mercer kernel map also corresponds to a dot product in feature space.

Since K is a symmetric positive semidefinite matrix, it has real and non-negative eigenvalues. It can be decomposed as follows:

$$K = U \wedge U^T$$

where \boldsymbol{U} is the orthonormal matrix of eigenvectors $\boldsymbol{u}_i = (u_{i1}, u_{i2}, \dots, u_{in})^T \in \mathbb{R}^n$ (for $i = 1, \dots, n$), and Λ is the diagonal matrix of eigenvalues, with both arranged in non-increasing order of the eigenvalues $\lambda_1 \ge \lambda_2 \ge \dots \ge \lambda_n \ge 0$: The Mercer map ϕ is given as

$$\phi(\boldsymbol{x}_i) = \sqrt{\Lambda} \boldsymbol{U}_i$$

where \boldsymbol{U}_i is the *i*th row of \boldsymbol{U} .

The kernel value is simply the dot product between scaled rows of U:

$$\phi(\mathbf{x}_i)^T \phi(\mathbf{x}_j) = \left(\sqrt{\Lambda} \mathbf{U}_i\right)^T \left(\sqrt{\Lambda} \mathbf{U}_j\right) = \mathbf{U}_i^T \wedge \mathbf{U}_j$$

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Polynomial Kernel

Polynomial kernels are of two types: homogeneous or inhomogeneous.

Let $x, y \in \mathbb{R}^d$. The (inhomogeneous) *polynomial kernel* is defined as

$$K_q(\mathbf{x},\mathbf{y}) = \phi(\mathbf{x})^T \phi(\mathbf{y}) = (\mathbf{c} + \mathbf{x}^T \mathbf{y})^q$$

where q is the degree of the polynomial, and $c \ge 0$ is some constant. When c = 0 we obtain the homogeneous kernel, comprising only degree q terms. When c > 0, the feature space is spanned by all products of at most q attributes. This can be seen from the binomial expansion

$$\mathcal{K}_q(\mathbf{x}, \mathbf{y}) = (c + \mathbf{x}^T \mathbf{y})^q = \sum_{k=1}^q \binom{q}{k} c^{q-k} \left(\mathbf{x}^T \mathbf{y} \right)^k$$

The most typical cases are the *linear* (with q = 1) and *quadratic* (with q = 2) kernels, given as

$$K_1(\mathbf{x}, \mathbf{y}) = c + \mathbf{x}^T \mathbf{y}$$
$$K_2(\mathbf{x}, \mathbf{y}) = (c + \mathbf{x}^T \mathbf{y})^2$$

The Gaussian kernel, also called the Gaussian radial basis function (RBF) kernel, is defined as

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

where $\sigma > 0$ is the spread parameter that plays the same role as the standard deviation in a normal density function.

Note that $K(\mathbf{x}, \mathbf{x}) = 1$, and further that the kernel value is inversely related to the distance between the two points \mathbf{x} and \mathbf{y} .

A feature space for the Gaussian kernel has infinite dimensionality.

Basic Kernel Operations in Feature Space

Basic data analysis tasks that can be performed solely via kernels, without instantiating $\phi(\mathbf{x})$.

Norm of a Point: We can compute the norm of a point $\phi(\mathbf{x})$ in feature space as follows:

$$\|\phi(\boldsymbol{x})\|^2 = \phi(\boldsymbol{x})^T \phi(\boldsymbol{x}) = K(\boldsymbol{x}, \boldsymbol{x})$$

which implies that $\|\phi(\mathbf{x})\| = \sqrt{K(\mathbf{x}, \mathbf{x})}$.

Distance between Points: The distance between $\phi(\mathbf{x}_i)$ and $\phi(\mathbf{x}_j)$ is

$$\|\phi(\mathbf{x}_i) - \phi(\mathbf{x}_j)\|^2 = \|\phi(\mathbf{x}_i)\|^2 + \|\phi(\mathbf{x}_j)\|^2 - 2\phi(\mathbf{x}_i)^T \phi(\mathbf{x}_j)$$
$$= K(\mathbf{x}_i, \mathbf{x}_i) + K(\mathbf{x}_j, \mathbf{x}_j) - 2K(\mathbf{x}_i, \mathbf{x}_j)$$

which implies that

$$\|\phi(\mathbf{x}_i) - \phi(\mathbf{x}_j)\| = \sqrt{K(\mathbf{x}_i, \mathbf{x}_i) + K(\mathbf{x}_j, \mathbf{x}_j) - 2K(\mathbf{x}_i, \mathbf{x}_j)}$$

Basic Kernel Operations in Feature Space

Kernel Value as Similarity: We can rearrange the terms in

$$\|\phi(\mathbf{x}_i) - \phi(\mathbf{x}_j)\|^2 = K(\mathbf{x}_i, \mathbf{x}_i) + K(\mathbf{x}_j, \mathbf{x}_j) - 2K(\mathbf{x}_i, \mathbf{x}_j)$$

to obtain

$$\frac{1}{2}\left(\|\phi(\mathbf{x}_i)\|^2 + \|\phi(\mathbf{x}_j)\|^2 - \|\phi(\mathbf{x}_i) - \phi(\mathbf{x}_j)\|^2\right) = K(\mathbf{x}_i, \mathbf{x}_j) = \phi(\mathbf{x}_i)^T \phi(\mathbf{x}_j)$$

The more the distance $\|\phi(\mathbf{x}_i) - \phi(\mathbf{x}_i)\|$ between the two points in feature space, the less the kernel value, that is, the less the similarity.

Mean in Feature Space: The mean of the points in feature space is given as $\mu_{\phi} = 1/n \sum_{i=1}^{n} \phi(\mathbf{x}_i)$. Thus, we cannot compute it explicitly. However, the the squared norm of the mean is:

$$\|\boldsymbol{\mu}_{\phi}\|^2 = \boldsymbol{\mu}_{\phi}^{\mathsf{T}} \boldsymbol{\mu}_{\phi} = \frac{1}{n^2} \sum_{i=1}^n \sum_{j=1}^n \mathcal{K}(\boldsymbol{x}_i, \boldsymbol{x}_j)$$

The squared norm of the mean in feature space is simply the average of the values in the kernel matrix **K**. Chapter 5: Kernel Methods

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Basic Kernel Operations in Feature Space

Total Variance in Feature Space: The total variance in feature space is obtained by taking the average squared deviation of points from the mean in feature space:

$$\sigma_{\phi}^{2} = \frac{1}{n} \sum_{i=1}^{n} \|\phi(\mathbf{x}_{i}) - \boldsymbol{\mu}_{\phi}\|^{2} = \frac{1}{n} \sum_{i=1}^{n} \mathcal{K}(\mathbf{x}_{i}, \mathbf{x}_{i}) - \frac{1}{n^{2}} \sum_{i=1}^{n} \sum_{j=1}^{n} \mathcal{K}(\mathbf{x}_{i}, \mathbf{x}_{j})$$

Centering in Feature Space We can center each point in feature space by subtracting the mean from it, as follows:

$$\hat{\phi}(\boldsymbol{x}_i) = \phi(\boldsymbol{x}_i) - \boldsymbol{\mu}_{\phi}$$

The kernel between centered points is given as

$$\hat{\mathcal{K}}(\mathbf{x}_{i}, \mathbf{x}_{j}) = \hat{\phi}(\mathbf{x}_{i})^{T} \hat{\phi}(\mathbf{x}_{j})$$
$$= \mathcal{K}(\mathbf{x}_{i}, \mathbf{x}_{j}) - \frac{1}{n} \sum_{k=1}^{n} \mathcal{K}(\mathbf{x}_{i}, \mathbf{x}_{k}) - \frac{1}{n} \sum_{k=1}^{n} \mathcal{K}(\mathbf{x}_{j}, \mathbf{x}_{k}) + \frac{1}{n^{2}} \sum_{a=1}^{n} \sum_{b=1}^{n} \mathcal{K}(\mathbf{x}_{a}, \mathbf{x}_{b})$$

More compactly, we have:

$$\hat{\boldsymbol{K}} = \left(\boldsymbol{I} - \frac{1}{n} \boldsymbol{1}_{n \times n} \right) \boldsymbol{K} \left(\boldsymbol{I} - \frac{1}{n} \boldsymbol{1}_{n \times n} \right)$$

where $1_{n \times n}$ is the $n \times n$ matrix of ones.

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Normalizing in Feature Space: The dot product between normalized points in feature space corresponds to the cosine of the angle between them

$$\phi_n(\mathbf{x}_i)^T \phi_n(\mathbf{x}_j) = \frac{\phi(\mathbf{x}_i)^T \phi(\mathbf{x}_j)}{\|\phi(\mathbf{x}_i)\| \cdot \|\phi(\mathbf{x}_j)\|} = \cos\theta$$

If the mapped points are both centered and normalized, then a dot product corresponds to the correlation between the two points in feature space.

The normalized kernel matrix, K_n , can be computed using only the kernel function K, as

$$\boldsymbol{K}_{n}(\boldsymbol{x}_{i},\boldsymbol{x}_{j}) = \frac{\phi(\boldsymbol{x}_{i})^{T}\phi(\boldsymbol{x}_{j})}{\|\phi(\boldsymbol{x}_{i})\| \cdot \|\phi(\boldsymbol{x}_{j})\|} = \frac{K(\boldsymbol{x}_{i},\boldsymbol{x}_{j})}{\sqrt{K(\boldsymbol{x}_{i},\boldsymbol{x}_{i}) \cdot K(\boldsymbol{x}_{j},\boldsymbol{x}_{j})}}$$

 K_n has all diagonal elements as 1.

Spectrum Kernel for Strings

Given alphabet Σ , the *I*-spectrum feature map is the mapping $\phi: \Sigma^* \to \mathbb{R}^{|\Sigma|^I}$ from the set of substrings over Σ to the $|\Sigma|^I$ -dimensional space representing the number of occurrences of all possible substrings of length *I*, defined as

$$\phi(\mathbf{x}) = \left(\cdots, \#(\alpha), \cdots\right)_{\alpha \in \Sigma}^{T}$$

where $\#(\alpha)$ is the number of occurrences of the *I*-length string α in *x*.

The (full) spectrum map considers all lengths from l = 0 to $l = \infty$, leading to an infinite dimensional feature map $\phi : \Sigma^* \to \mathbb{R}^\infty$:

$$\phi(\mathbf{x}) = \left(\cdots, \#(\alpha), \cdots\right)_{\alpha \in \Sigma^*}^T$$

where $\#(\alpha)$ is the number of occurrences of the string α in x.

The (*I*-)spectrum kernel between two strings x_i, x_j is simply the dot product between their (*I*-)spectrum maps:

$$K(\mathbf{x}_i, \mathbf{x}_j) = \phi(\mathbf{x}_i)^T \phi(\mathbf{x}_j)$$

The (full) spectrum kernel can be computed efficiently via suffix trees in O(n+m) time for two strings of length *n* and *m*.

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Diffusion Kernels on Graph Nodes

Let **S** be some symmetric similarity matrix between nodes of a graph G = (V, E). For instance, **S** can be the (weighted) adjacency matrix **A** or the Laplacian matrix $\mathbf{L} = \mathbf{A} - \Delta$ (or its negation), where Δ is the degree matrix for an undirected graph G, defined as $\Delta(i,i) = d_i$ and $\Delta(i,j) = 0$ for all $i \neq j$, and d_i is the degree of node *i*.

Power Kernels: Summing up the product of the base similarities over all *l*-length paths between two nodes, we obtain the *l*-length similarity matrix $S^{(l)}$, which is simply the *l*th power of S, that is,

$$\boldsymbol{S}^{(I)} = \boldsymbol{S}^{I}$$

Even path lengths lead to positive semidefinite kernels, but odd path lengths are not guaranteed to do so, unless the base matrix \boldsymbol{S} is itself a positive semidefinite matrix.

Power kernel K can be obtained via the eigen-decomposition of S':

$$\boldsymbol{K} = \boldsymbol{S}' = \left(\boldsymbol{U} \wedge \boldsymbol{U}^T \right)' = \boldsymbol{U} \left(\wedge' \right) \boldsymbol{U}^T$$

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Exponential Diffusion Kernel

The exponential diffusion kernel we can obtain a new kernel between nodes of a graph by paths of all possible lengths, but damps the contribution of longer paths

$$\begin{aligned} \boldsymbol{\kappa} &= \sum_{l=0}^{\infty} \frac{1}{l!} \beta^l \boldsymbol{S}^l \\ &= \boldsymbol{I} + \beta \boldsymbol{S} + \frac{1}{2!} \beta^2 \boldsymbol{S}^2 + \frac{1}{3!} \beta^3 \boldsymbol{S}^3 + \cdots \\ &= \exp\{\beta \boldsymbol{S}\} \end{aligned}$$

where β is a damping factor, and exp{ βS } is the matrix exponential. The series on the right hand side above converges for all $\beta \ge 0$.

Substituting $\boldsymbol{S} = \boldsymbol{U} \wedge \boldsymbol{U}^{T}$ the kernel can be computed as

$$\begin{aligned} \boldsymbol{\mathcal{K}} &= \boldsymbol{\mathcal{I}} + \beta \boldsymbol{\mathcal{S}} + \frac{1}{2!} \beta^2 \boldsymbol{\mathcal{S}}^2 + \cdots \\ &= \boldsymbol{\mathcal{U}} \begin{pmatrix} \exp\{\beta\lambda_1\} & 0 & \cdots & 0 \\ 0 & \exp\{\beta\lambda_2\} & \cdots & 0 \\ \vdots & \vdots & \ddots & 0 \\ 0 & 0 & \cdots & \exp\{\beta\lambda_n\} \end{pmatrix} \boldsymbol{\mathcal{U}}^T \end{aligned}$$

where λ_i is an eigenvalue of **S**.

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The von Neumann diffusion kernel is defined as

$$oldsymbol{\mathcal{K}} = \sum_{l=0}^{\infty} eta^l oldsymbol{\mathcal{S}}^l$$

where $\beta \ge 0$. Expanding and rearranging the terms, we obtain

$$\boldsymbol{K} = (\boldsymbol{I} - \beta \boldsymbol{S})^{-1}$$

The kernel is guaranteed to be positive semidefinite if $|\beta| < 1/\rho(\mathbf{S})$, where $\rho(\mathbf{S}) = \max_i \{|\lambda_i|\}$ is called the *spectral radius* of \mathbf{S} , defined as the largest eigenvalue of \mathbf{S} in absolute value.

Graph Diffusion Kernel: Example



Graph Diffusion Kernel: Example

Let the base similarity matrix \boldsymbol{S} be the negated Laplacian matrix

$$\boldsymbol{S} = -\boldsymbol{L} = \boldsymbol{A} - \boldsymbol{D} = \begin{pmatrix} -2 & 0 & 1 & 1 & 0 \\ 0 & -2 & 1 & 0 & 1 \\ 1 & 1 & -3 & 1 & 0 \\ 1 & 0 & 1 & -3 & 1 \\ 0 & 1 & 0 & 1 & -2 \end{pmatrix}$$

The eigenvalues of **S** are as follows:

 $\lambda_1 = 0$ $\lambda_2 = -1.38$ $\lambda_3 = -2.38$ $\lambda_4 = -3.62$ $\lambda_5 = -4.62$

and the eigenvectors of \boldsymbol{S} are

$$\boldsymbol{U} = \begin{pmatrix} \boldsymbol{u}_1 & \boldsymbol{u}_2 & \boldsymbol{u}_3 & \boldsymbol{u}_4 & \boldsymbol{u}_5 \\ 0.45 & -0.63 & 0.00 & 0.63 & 0.00 \\ 0.45 & 0.51 & -0.60 & 0.20 & -0.37 \\ 0.45 & -0.20 & -0.37 & -0.51 & 0.60 \\ 0.45 & -0.20 & 0.37 & -0.51 & -0.60 \\ 0.45 & 0.51 & 0.60 & 0.20 & 0.37 \end{pmatrix}$$

Graph Diffusion Kernel: Example

Assuming $\beta = 0.2$, the exponential diffusion kernel matrix is given as

$$\boldsymbol{\mathcal{K}} = \exp\{0.2\boldsymbol{\mathcal{S}}\} = \boldsymbol{\mathcal{U}} \begin{pmatrix} \exp\{0.2\lambda_1\} & 0 & \cdots & 0 \\ 0 & \exp\{0.2\lambda_2\} & \cdots & 0 \\ \vdots & \vdots & \ddots & 0 \\ 0 & 0 & \cdots & \exp\{0.2\lambda_n\} \end{pmatrix} \boldsymbol{\mathcal{U}}^{\mathsf{T}}$$
$$= \begin{pmatrix} 0.70 & 0.01 & 0.14 & 0.14 & 0.01 \\ 0.01 & 0.70 & 0.13 & 0.03 & 0.14 \\ 0.14 & 0.13 & 0.59 & 0.13 & 0.03 \\ 0.14 & 0.03 & 0.13 & 0.59 & 0.13 \\ 0.01 & 0.14 & 0.03 & 0.13 & 0.70 \end{pmatrix}$$

Assuming $\beta = 0.2$, the von Neumann kernel is given as

$$\boldsymbol{K} = \boldsymbol{U}(\boldsymbol{I} - 0.2\Lambda)^{-1} \boldsymbol{U}^{\mathsf{T}} = \begin{pmatrix} 0.75 & 0.02 & 0.11 & 0.11 & 0.02 \\ 0.02 & 0.74 & 0.10 & 0.03 & 0.11 \\ 0.11 & 0.10 & 0.66 & 0.10 & 0.03 \\ 0.11 & 0.03 & 0.10 & 0.66 & 0.10 \\ 0.02 & 0.11 & 0.03 & 0.10 & 0.74 \end{pmatrix}$$

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