# Kernel Principal Component Analysis

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**Abstract**

Principal component analysis (PCA) and kernel methods are tools often used in data science. The underlying theory of these tools depend on the properties of a special type of Hilbert space called a reproducing kernel Hilbert space (RKHS). This paper explores the essence of RKHSs using data science examples, in particular, PCA and kernel PCA. When kernel methods are applied to PCA, we can analyze nonlinear data in a high-dimensional feature space with some nice properties.

## 1 Introduction

Pattern recognition is an applied science that draws from a variety of mathematical fields. Principal component analysis (PCA) is an unsupervised learning technique that can be used to reveal patterns in high-dimensional data. The transformation used in the PCA algorithm resembles the singular value decomposition (SVD) in linear algebra and the Karhunen-Loève transform (KLT) in stochastic processes. Borrowing results from these different approaches, we can start to generalize the PCA algorithm. The kernel trick from machine learning can be used to create a nonlinear algorithm from a linear one. In particular,

the kernel trick can be applied to PCA. Kernel PCA is a nonlinear version of PCA that maps data to a suitable Hilbert space and attempts to find patterns in this new context.

We will begin this paper by deriving the PCA transform in Section 2. Part of this work depends on results from the SVD and is discussed in Section 2.2. The PCA section is concluded with examples. In Section 3, we justify the kernel trick. This prompts the discussion of reproducing kernel Hilbert spaces, Mercer's theorem, the Moore-Aronszajn theorem, and the Riesz representation theorem. Last, the kernel PCA algorithm is derived in Section 4.

# 2 Principal Component Analysis

When analyzing data, it can be convenient to transform the given input variables to produce new features. For a well-chosen transform, these features may be approximated using fewer dimensions than the original input space [10]. This is an example of a data preprocessing technique known as *dimension reduction* and can reveal low-dimensional structure.

Principal component analysis (PCA) is an orthogonal coordinate transform that is suitable for dimension reduction if some of the inputs are linearly correlated. In this case, PCA transforms redundant variables in the input space producing uncorrelated variables in the feature space.

There are a number of ways to derive the optimal PCA transform. One approach presented in [10] is based on finding uncorrelated features. It is straightforward to show that uncorrelated features have a diagonal covariance matrix. This can be used to solve for the covariance matrix $C$ of input variables. By asserting the orthogonality of the PCA transform, we obtain $V$ from the diagonalization of the covariance matrix $C = VDV^\top$. Given this PCA transform, we can show that $V$ minimizes projection residuals as in [18].

**Remark 2.1.** The following derivations make use of both inner products and outer products. If $x$ is a column vector, then $x^\top x$ represents an inner product[1] and the result is a scalar. But, if $x$ is a row vector, then $x^\top x$ represents an outer product and the result is a matrix. To avoid this ambiguity, we will use the notations $\langle \cdot, \cdot \rangle$ and $\otimes$ to indicate inner product and outer product, respectively.

Let $x = [x_i], y = [y_i] \in \mathbb{R}^n$ and define

$$\langle x, y \rangle = \sum_{i=1}^{n} x_i y_i \in \mathbb{R} \tag{1}$$

and

$$x \otimes y = [x_i y_j]_{ij}^{n \times n} = \begin{bmatrix} x_1y_1 & x_1y_2 & \cdots & x_1y_n \\ x_2y_1 & x_2y_2 & \cdots & x_2y_n \\ \vdots & \vdots & \ddots & \vdots \\ x_ny_1 & x_ny_2 & \cdots & x_ny_n \end{bmatrix} \in \mathbb{R}^{n \times n}. \tag{2}$$

[1] We will define inner product more precisely in Definition 3.2. Until then, $\langle \cdot, \cdot \rangle$ will only be used as a dot product of vectors.

## 2.1 Finding uncorrelated features

The correlation between two random variables $x$ and $y$ is defined as

$$\operatorname{corr}(x, y) = \frac{E[(x - \mu_x)(y - \mu_y)]}{\sigma_x \sigma_y}, \tag{3}$$

where $\mu_x$, $\mu_y$ and $\sigma_x$, $\sigma_y$ are the respective means and standard deviations of $x$ and $y$. We say $x$ and $y$ are uncorrelated when $\operatorname{corr}(x, y) = 0$. This happens if and only if

$$E[(x - \mu_x)(y - \mu_y)] = \operatorname{cov}(x, y) = 0. \tag{4}$$

For a multivariate random variable $x = [x_j] \in \mathbb{R}^{1\times d}$, the covariance matrix has the entry $\operatorname{cov}(x_i, x_j)$ in the $i$-th row and $j$-th column. That is,

$$E[(x - \mu_x) \otimes (x - \mu_x)] = \begin{bmatrix} \operatorname{cov}(x_1, x_1) & \operatorname{cov}(x_1, x_2) & \cdots & \operatorname{cov}(x_1, x_d) \\ \operatorname{cov}(x_2, x_1) & \operatorname{cov}(x_2, x_2) & \cdots & \operatorname{cov}(x_2, x_d) \\ \vdots & \vdots & \ddots & \vdots \\ \operatorname{cov}(x_d, x_1) & \operatorname{cov}(x_d, x_2) & \cdots & \operatorname{cov}(x_d, x_d) \end{bmatrix}. \tag{5}$$

If $x_1, x_2, \ldots, x_d$ are pairwise uncorrelated, then $\operatorname{cov}(x_i, x_j) = 0$ for all $i \neq j$. Hence, the covariance matrix of uncorrelated variables is diagonal.

Now let $a_1, a_2, \ldots, a_n \in \mathbb{R}^{1\times d}$ represent $n$ observations in $d$ variables with $n \geq d$. These observations can be considered points in $d$-dimensional space whose centroid is $\mu_a = \frac{1}{n}\sum_{i=1}^{n} a_i$. Our goal is to determine a coordinate transform whose image has uncorrelated variables. Accordingly, let $V \in \mathbb{R}^{d\times d}$ be the change of basis matrix and let

$$b_i = (a_i - \mu_a)V, \quad \text{for } i = 1, 2, \ldots, n \tag{6}$$

be observations with respect to the feature coordinates. Then

$$\mu_b = \frac{1}{n}\sum_{i=1}^{n} b_i = \frac{1}{n}\left(\sum_{i=1}^{n} a_i - \mu_a\right) V = \left[\left(\frac{1}{n}\sum_{i=1}^{n} a_i\right) - \mu_a\right] V = 0. \tag{7}$$

Using equation (5), we can compute the sample covariance matrices[2] as

$$C = \sum_{i=1}^{n} \frac{(a_i - \mu_a) \otimes (a_i - \mu_a)}{n - 1}, \qquad D = \sum_{i=1}^{n} \frac{b_i \otimes b_i}{n - 1}. \tag{8}$$

Combining these, we have

$$D = \sum_{i=1}^{n} \frac{b_i \otimes b_i}{n - 1} = \sum_{i=1}^{n} \frac{V^\top (a_i - \mu_a) \otimes (a_i - \mu_a) V}{n - 1} = V^\top C V. \tag{9}$$

[2]Population variance is scaled by $\frac{1}{n}$ while sample variance is scaled by $\frac{1}{n-1}$. This is known as Bessel's correction and is consistent with the NumPy `cov` function [5].

If we restrict $V$ to be orthogonal, then we can solve for $C$ to get

$$C = VDV^\top. \tag{10}$$

Since $D$ is the covariance matrix of uncorrelated features, by the argument above, it is diagonal and equation (10) is the diagonalization of $C$. Hence, the columns of $V$ are an orthonormal basis $(v_j)_{j=1}^d$ corresponding to eigenvalues $(\lambda_j)_{j=1}^d$ on the diagonal of $D$. When the eigenvalues and eigenvectors are ordered such that

$$\lambda_1 \geq \lambda_2 \geq \cdots \geq \lambda_d, \tag{11}$$

then $(v_j)_{j=1}^d$ are called the *principal components*.

Let $A \in \mathbb{R}^{n\times d}$ be the matrix whose rows are $a_1, a_2, \ldots, a_n$ and let $B \in \mathbb{R}^{n\times d}$ be the matrix whose rows are $b_1, b_2, \ldots, b_n$. Assume the matrix $A$ has been centered, i.e., $\mu_a$ is the zero vector. Then equation (6) implies $B = AV$ and equation (8) implies $C = \frac{1}{n-1}A^\top A$.

For $p \leq d$, the rank-$p$ projection matrix $V_p = \begin{bmatrix} v_1 & v_2 & \cdots & v_p \end{bmatrix} \in \mathbb{R}^{d\times p}$. In the following sections, we will show that $B_p = AV_p$ has the smallest possible projection error for all $1 \leq p \leq d$.

## 2.2 Singular value decomposition

The singular value decomposition (SVD) of a rectangular matrix generalizes the idea of diagonalization for square matrices. Moreover, SVD illustrates a connection between the matrices $A^\top A$ and $AA^\top$.

**Theorem 2.2** (Singular value decomposition)**.** *[8] Let $A \in \mathbb{R}^{n\times d}$. Then there exist orthogonal matrices $U \in \mathbb{R}^{n\times n}$ and $V \in \mathbb{R}^{d\times d}$ and a diagonal matrix $S \in \mathbb{R}^{n\times d}$ such that $A = USV^\top$. We say the columns of $U = \begin{bmatrix} u_1 & u_2 & \cdots & u_n \end{bmatrix}$ and $V = \begin{bmatrix} v_1 & v_2 & \cdots & v_d \end{bmatrix}$ are the left and right singular vectors of $A$, respectively The diagonal entries of $S$ are the called the singular values $\sigma_1, \sigma_2, \ldots, \sigma_r$, where $r = \operatorname{rank} A \leq \min\{n, d\}$. Then we can write*

$$A = USV^\top = \sum_{i=1}^{r} \sigma_i u_i v_i^\top. \tag{12}$$

The SVD of a matrix $A$ can be found by diagonalizing $A^\top A$ and $AA^\top$. If $A = USV^\top$, then

$$\begin{aligned} A^\top A &= (USV^\top)^\top (USV^\top) = VSU^\top USV^\top = VS^2V^\top \\ AA^\top &= (USV^\top)(USV^\top)^\top = USV^\top VSU^\top = US^2U^\top. \end{aligned}$$

So, $\{v_j\}_{j=1}^d$ are the eigenvectors of $A^\top A$, $\{u_j\}_{j=1}^n$ are the eigenvectors of $AA^\top$, and $\{\sigma_j^2\}_{j=1}^r$ are the eigenvalues of both $A^\top A$ and $AA^\top$. Notice that the SVD of $A$ will give us the projection matrix $V$ in equation (10), provided that $A$ is centered. In this way, we see that PCA is really just a special case of the SVD.

**Theorem 2.3** (Frobenius norm)**.** *[13, 8] The Frobenius norm (or Hilbert-Schmidt norm) of a matrix $A = [a_{ij}] \in \mathbb{R}^{n\times d}$ is given by*

$$\|A\|_F = \sqrt{\operatorname{tr}(A^\top A)} = \sqrt{\sum_{i=1}^{n}\sum_{j=1}^{d} a_{ij}^2}. \tag{13}$$

*Proof.* Let $A = [a_{ij}] \in \mathbb{R}^{n\times d}$. Then $A^\top A = [\sum_{k=1}^{n} a_{ki}a_{kj}]_{ij}$. It follows that $\|A\|_F^2 = \operatorname{tr}(A^\top A) = \sum_{i=1}^{n}\sum_{j=1}^{d} a_{ij}^2$. Clearly, $\|A\|_F > 0$ whenever $A$ is not the zero matrix and $\|A\|_F = 0$ whenever $A$ is the zero matrix.

For the triangle inequality, consider another matrix $B = [b_{ij}] \in \mathbb{R}^{d\times m}$. Then

$$\begin{aligned}
\|AB\|_F &= \sqrt{\sum_{i=1}^{n}\sum_{j=1}^{m}\sum_{k=1}^{d}(a_{ik}b_{kj})^2} \\
&\le \sqrt{\sum_{i=1}^{n}\sum_{j=1}^{m}\left(\sum_{k=1}^{d} a_{ik}^2\right)\left(\sum_{k=1}^{d} b_{kj}^2\right)} \\
&= \sqrt{\sum_{i=1}^{n}\sum_{j=1}^{d} a_{ij}^2}\sqrt{\sum_{i=1}^{d}\sum_{j=1}^{m} b_{ij}^2} \\
&= \|A\|_F\|B\|_F.
\end{aligned}$$

Thus, $\|\cdot\|_F$ is a matrix norm. □

Combining equations (12) and (13), we have

$$\|A\|_F = \sqrt{\sum_{i=1}^{r}\sigma_i^2} = \sqrt{\sum_{i=1}^{r}\lambda_i}, \tag{14}$$

where $\{\sigma_i\}_{i=1}^{r}$ are the singular values of $A$ and $\{\lambda_i\}_{i=1}^{r}$ are the eigenvalues of $A^\top A$ or $AA^\top$.

## 2.3 Minimizing projection residuals

In this section, we want to show that the PCA projection minimizes the residual error.

Suppose $a_1, a_2, \ldots, a_n \in \mathbb{R}^n$. Let $v_1, v_2, \ldots, v_d \in \mathbb{R}^d$ be the principal component vectors given in Section 2.1. Define the projections onto the subspace spanned by $v_1, v_2, \ldots, v_p$, for $p \le d$, as

$$\hat{a}_i = \sum_{j=1}^{p} \langle a_i, v_j\rangle\, v_j, \quad \text{for } i = 1, 2, \ldots, n. \tag{15}$$

Then each residual from the projection is given by

$$\begin{aligned}
\|a_i - \hat{a}_i\|^2 &= \langle a_i - \hat{a}_i, a_i - \hat{a}_i \rangle \\
&= \|a_i\|^2 - 2\langle a_i, \hat{a}_i \rangle + \|\hat{a}_i\|^2 \\
&= \|a_i\|^2 - 2\left\langle a_i, \sum_{j=1}^{p} \langle a_i, v_j \rangle v_j \right\rangle + \left\| \sum_{j=1}^{p} \langle a_i, v_j \rangle v_j \right\|^2 \\
&= \|a_i\|^2 - 2\sum_{j=1}^{p} \langle a_i, v_j \rangle \langle a_i, v_j \rangle + \sum_{j=1}^{p} \langle a_i, v_j \rangle^2 \|v_j\|^2 \\
&= \|a_i\|^2 - 2\sum_{j=1}^{p} \langle a_i, v_j \rangle^2 + \sum_{j=1}^{p} \langle a_i, v_j \rangle^2 \\
&= \|a_i\|^2 - \sum_{j=1}^{p} \langle a_i, v_j \rangle^2.
\end{aligned} \tag{16}$$

Then the mean squared error to be minimized is

$$\text{MSE} = \frac{1}{n}\sum_{i=1}^{n} \|a_i\|^2 - \frac{1}{n}\sum_{i=1}^{n}\sum_{j=1}^{p} \langle a_i, v_j \rangle^2. \tag{17}$$

Since the first term does not depend on $v_j$, the MSE is minimized whenever $\sum_{i=1}^{n}\sum_{j=1}^{p} \langle a_i, v_j \rangle^2$ is maximized. Let $A$ be the matrix whose rows are $a_1, a_2, \ldots, a_n$ and $V_p$ be the matrix whose columns are $v_1, v_2, \ldots, v_p$. By Theorem 2.3 and equation (14), this becomes

$$\sum_{i=1}^{n}\sum_{j=1}^{p} \langle a_i, v_j \rangle^2 = \text{tr}[(AV_p)^\top AV] = \|AV_p\|_F = \sum_{i=j}^{p} \lambda_j. \tag{18}$$

It follows that the vectors $v_1, v_2, \ldots, v_p$ which minimize projection error correspond to the $p$ largest eigenvalues given by equation (11).

## 2.4 PCA algorithm

Let $A$ be a data matrix whose $n$ rows correspond to observations and $d$ columns correspond to variables. The following algorithm demonstrates a simple method for computing the PCA of $A$:

1. Compute the centered matrix $A_0 = A - \text{col mean}(A)$.
2. Compute the covariance matrix $C = \frac{1}{n-1} A_0^\top A_0$.
3. Diagonalize the covariance matrix such that $C = VDV^\top$.
4. Order the eigenvalues and eigenvectors so that $\lambda_1 \geq \lambda_2 \geq \cdots \geq \lambda_d$. We call the ordered eigenvalues the *principal components*.

5. Choose the dimension of the subspace $p \leq d$.
6. Construct the rank-$p$ transformation matrix $V_p \in \mathbb{R}^{d \times p}$ using the first $p$ principal components $v_1, v_2, \ldots, v_p$.
7. The image of $A$ under the PCA transform is $B = A_0 V_p$.

**Example 2.4.** Consider the following matrix

$$A = \begin{bmatrix} 5 & 3 & 6 & 7 & 6 \\ 4 & 5 & 7 & 1 & 3 \\ 5 & 7 & 6 & 1 & 0 \\ 6 & 10 & 12 & 12 & 11 \\ 9 & 10 & 12 & 13 & 9 \end{bmatrix}. \tag{19}$$

The column means are $\mu = [5.8, 7, 8.6, 6.8, 5.8]$. Then the mean-centered data becomes

$$A_0 = A - \mu = \frac{1}{5} \begin{bmatrix} -4 & -20 & -13 & 1 & 1 \\ -9 & -10 & -8 & -29 & -14 \\ -4 & 0 & -13 & -29 & -29 \\ 1 & 15 & 17 & 26 & 26 \\ 16 & 15 & 17 & 31 & 16 \end{bmatrix}. \tag{20}$$

The covariance matrix is

$$C = A_0^\top A_0 = \frac{1}{5} \begin{bmatrix} 74 & 85 & 93 & 179 & 104 \\ 85 & 190 & 170 & 225 & 150 \\ 93 & 170 & 196 & 313 & 238 \\ 179 & 225 & 313 & 664 & 484 \\ 104 & 150 & 238 & 484 & 394 \end{bmatrix}. \tag{21}$$

Diagonalizing $C$ gives

$$V = \begin{bmatrix} 0.1888 & -0.2020 & -0.6366 & 0.5495 & -0.4651 \\ 0.2755 & -0.7886 & 0.1472 & -0.4502 & -0.2791 \\ 0.3606 & -0.3464 & 0.3128 & 0.5836 & 0.5582 \\ 0.6979 & 0.2522 & -0.4422 & -0.3707 & 0.3411 \\ 0.5209 & 0.3922 & 0.5288 & 0.1316 & -0.5271 \end{bmatrix},$$

$$D = \begin{bmatrix} 264.8458 & 0 & 0 & 0 & 0 \\ 0 & 27.9766 & 0 & 0 & 0 \\ 0 & 0 & 9.3198 & 0 & 0 \\ 0 & 0 & 0 & 1.4579 & 0 \\ 0 & 0 & 0 & 0 & 0 \end{bmatrix}.$$

If we keep all 5 principal component vectors, then $V_5 = V$ and the projection of $A_0$ along $V$ is

$$B = A_0 V = \begin{bmatrix} -1.9469 & 4.3453 & -0.8756 & -0.2039 & 0 \\ -6.9742 & -0.0660 & 1.4352 & 0.7590 & 0 \\ -8.1577 & -2.6752 & -0.8063 & -0.5704 & 0 \\ 8.4282 & -0.2330 & 1.8282 & -0.4996 & 0 \\ 8.6507 & -1.3711 & -1.5815 & 0.5149 & 0 \end{bmatrix}. \tag{22}$$

Here, the last column of $B$ is the zero vector because the last eigenvalue of $C$ is zero[3]. To perfectly reconstruct $A$, we need $k = 4$ principal components and the row vector $\mu$

$$A = BV^\top + \mu = BV_4^\top + \mu. \tag{23}$$

If we use $k = 3$ principal components, then the projection of $A_0$ onto $V_3$ is

$$B = A_0 V_3 = \begin{bmatrix} -1.9469 & 4.3453 & -0.8756 \\ -6.9742 & -0.0660 & 1.4352 \\ -8.1577 & -2.6752 & -0.8063 \\ 8.4282 & -0.2330 & 1.8282 \\ 8.6507 & -1.3711 & -1.5815 \end{bmatrix} \tag{24}$$

and $A$ is approximately reconstructed by

$$A \approx BV_3^\top + \mu = \begin{bmatrix} 5.1 & 2.9 & 6.1 & 6.9 & 6.0 \\ 3.6 & 5.3 & 6.6 & 1.3 & 2.9 \\ 5.3 & 6.7 & 6.3 & 0.8 & 0.1 \\ 6.3 & 9.8 & 12.3 & 11.8 & 11.1 \\ 8.7 & 10.2 & 11.7 & 13.2 & 8.9 \end{bmatrix}. \tag{25}$$

We can compute the reconstruction error using the Frobenius norm

$$E_k = \|A - (V_k^\top + \mu)\|_F.$$

By the SVD, we have $A_0 = USV^\top$, where $S = \sqrt{D}$. So, the projection of $A_0$ onto $V_p$ is

$$B = A_0 V_p = US_p, \tag{26}$$

where $S_p$ is the diagonal matrix of the first $p$ singular values. Then the reconstruction error becomes

$$\begin{aligned} \|A - (BV_p^\top + \mu)\|_F &= \|(A - \mu) - BV_p^\top\|_F \\ &= \|A_0 - BV_p^\top\|_F \\ &= \|USV^\top - US_pV^\top\|_F \\ &= \|U(S - S_p)V^\top\|_F \\ &= \|S - S_p\|_F \\ &= \sqrt{\sigma_{p+1}^2 + \cdots + \sigma_d^2}. \end{aligned}$$

Hence,

$$E_3 = \sigma_3 + \sigma_4 = \sqrt{1.4579^2 + 0^2} = 1.2074. \tag{27}$$

[3]Since we subtracted the column means from a square matrix $A$, the dimension of the row space was reduced to 4.

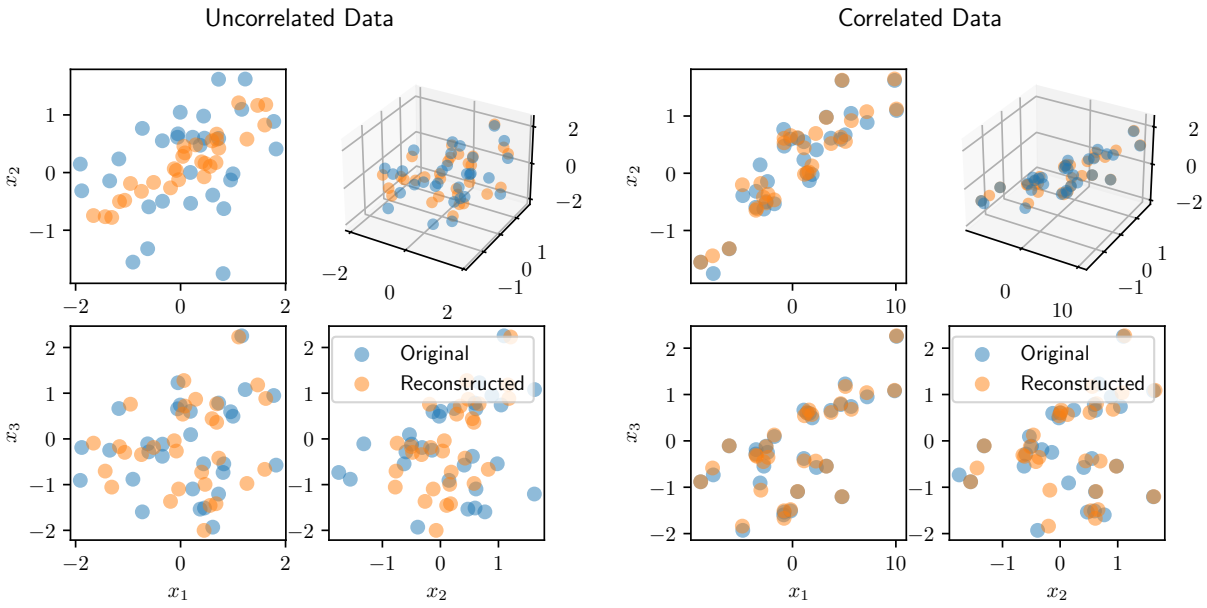


Figure 1: PCA projection of three-dimensional data onto two-dimensional subspace.

## 2.5 Applications of PCA

One of the most common applications of PCA is dimension reduction. When variables are correlated, the observations lie in some linear subspace of the original space. In this situation, PCA can be used to project down to the lower dimension and have the smallest possible projection error. This is particularly useful when the dimension of the input space is extremely large. A number challenges arise when analyzing high-dimensional data and are collectively referred to as the *curse of dimensionality* [10]. By working in the PCA feature space, these problems may be avoided.

**Example 2.5.** In the following experiment, we consider two sets of three-dimensional data. Each set of vectors $x_1, x_2, x_3 \in \mathbb{R}^n$ are sampled from a standard normal distribution with sample size $n = 30$. Suppose in the first set there is no apparent relationship among variables while, in second set, we have $x_1 = 4x_2 + 2x_3$. For simplicity, we say the first set is *uncorrelated* while the second set is *correlated*. See Figure 1. In the uncorrelated data, the reconstruction error is 3.92, which does not seem to indicate the presence of a pattern. The reconstruction error for the correlated data is 0.973. This error can be explained by the variation of $x_2$ with $x_3$. Meanwhile, the correlated pair plots indicate a pattern among $x_2$ vs $x_1$ and $x_3$ vs $x_1$.

# 3 Reproducing Kernel Hilbert Space

In this section, our goal is to establish properties of Hilbert spaces and kernel functions that can be used to modify the PCA algorithm. To begin, we will briefly cite some definitions and results from analysis [11], [15] and matrix theory [8].

## 3.1 Inner products and Hilbert spaces

**Definition 3.1** (Definite matrix)**.** [8] Let $A$ be an $n \times n$ matrix over the real numbers having the form

$$A = \begin{bmatrix} a_{11} & a_{12} & \cdots & a_{1n} \\ a_{21} & a_{22} & \cdots & a_{2n} \\ \vdots & \vdots & \ddots & \vdots \\ a_{n1} & a_{n2} & \cdots & a_{nn} \end{bmatrix} = [a_{ij}]. \tag{28}$$

The *transpose* of $A$ is $A^\top = [a_{ji}]$. We say that $A$ is *symmetric* whenever $A = A^\top$. A symmetric matrix $A$ is

1. *positive definite* if $x^\top Ax > 0$, for all nonzero $x \in \mathbb{R}^n$, or
2. *positive semidefinite* if $x^\top Ax \geq 0$, for all $x \in \mathbb{R}^n$.

Negative (semi)definite matrices can be defined in a similar fashion. A matrix is *definite* if it is either positive semidefinite or negative semidefinite. Otherwise, $A$ is an *indefinite matrix*.

Be aware that some authors use the terms positive definite ($>$) and nonnegative definite ($\geq$). Other authors use the modifier *strict* as in strict positive definite ($>$) and positive definite ($\geq$).

**Definition 3.2** (Inner product)**.** [20] Let $\langle \cdot, \cdot \rangle : \mathcal{X} \times \mathcal{X} \to \mathbb{R}$ be a function defined on the vector space $\mathcal{X}$. Then $\langle \cdot, \cdot \rangle$ is an *inner product* if the following properties hold for all $x, y, z \in \mathcal{X}$ and $\alpha, \beta \in \mathbb{R}$:

1. $\langle x, y \rangle = \langle y, x \rangle$; (symmetry)
2. $\langle \alpha x + \beta y, z \rangle = \alpha \langle x, z \rangle + \beta \langle y, z \rangle$; (bilinear)
3. $\langle x, x \rangle \geq 0$ and $\langle x, x \rangle = 0$ if and only if $x = 0$. (positive definite)

Note that linearity in the first argument with symmetry implies that the inner product is bilinear (linear in both arguments). Since inner products are positive definite, they induce a norm $\| \cdot \| : \mathcal{X} \to \mathbb{R}$ and metric $d : \mathcal{X} \times \mathcal{X} \to \mathbb{R}$ such that

$$\|x\| = \sqrt{\langle x, x \rangle} \quad \text{and} \quad d(x, y) = \|x - y\|. \tag{29}$$

An *inner product space*, *normed space*, and *metric space* are vector spaces along with an inner product, norm, and metric, respectively. It follows that an inner product space is also a normed space and a metric space. Then the induced norm has the following properties for all $x, y \in \mathcal{X}$ and $\alpha \in \mathbb{R}$:

1. $\|\alpha x\| = |\alpha|\|x\|$;
2. $\|x\| \geq 0$ and $\|x\| = 0$ if and only if $x = 0$;
3. $\|x + y\| \leq \|x\| + \|y\|$; (triangle inequality)
4. $\langle x, y\rangle^2 \leq \|x\|\|y\|$. (Cauchy-Schwarz inequality)

**Example 3.3.** The *Euclidean inner product* (or *dot product*) is the function $\langle \cdot, \cdot \rangle : \mathbb{R}^n \times \mathbb{R}^n \to \mathbb{R}$ such that

$$\langle x, y\rangle = \sum_{i=1}^{n} x_i y_i = x^\top y, \tag{30}$$

for all $x = [x_i], y = [y_i] \in \mathbb{R}^n$. Sometimes we write $x \cdot y$ to mean the Euclidean inner product. This induces the *Euclidean norm*

$$\|x\| = \sqrt{\sum_{i=1}^{n} x_i^2} \tag{31}$$

**Definition 3.4** (Hilbert space)**.** [11] A metric space is *complete* if the limit of every Cauchy sequence is in the space. A complete normed space is called a *Banach space*. A complete inner product space $\mathcal{H}$ is called a *Hilbert space*. We sometimes denote the inner product and norm of $\mathcal{H}$ as $\langle \cdot, \cdot \rangle_{\mathcal{H}}$ and $\| \cdot \|_{\mathcal{H}}$ to avoid ambiguity.

A Hilbert space is said to be *separable* if it contains a dense countable subset.

Two real Hilbert spaces $\mathcal{H}$ and $\mathcal{L}$ are said to be *isomorphic* if there is a linear bijection $T : \mathcal{H} \to \mathcal{L}$ such that $\langle x, y\rangle_{\mathcal{H}} = \langle Tx, Ty\rangle_{\mathcal{L}}$, for every $x, y \in \mathcal{H}$.

The *dimension* of a Hilbert space is the cardinality of its basis.

**Theorem 3.5** (Properties of Hilbert spaces)**.** *[11] Then the following properties hold for Hilbert spaces.*

1. *A Hilbert space is separable if and only if it has a countable orthonormal basis.*
2. *Two Hilbert spaces are isomorphic if and only if they have the same dimension.*
3. *Any inner product $\mathcal{H}$ space can be extended to a Hilbert space by completion and is unique up to isomorphism. We denote the completion as $\overline{\mathcal{H}}$.*
4. *A Hilbert space $\mathcal{H}$ can be decomposed into orthogonal subspaces $M$ and $M^\perp$ such that whenever $f \in M$ and $g \in M^\perp$, then $\langle f, g\rangle = 0$ [3]. We denote the* orthogonal decomposition *of a Hilbert space as the direct sum $\mathcal{H} = M \oplus M^\perp$.*

The following example demonstrates a useful property of separable Hilbert spaces.

**Example 3.6.** [15] Let $A$ be a nonempty index set. The space of square-summable indexed families is defined as

$$\ell^2(A) = \left\{ x : A \to \mathbb{R} \;\middle|\; \sum_{a \in A} x_a^2 < \infty \right\}. \tag{32}$$

Given the inner product

$$\langle x, y \rangle = \sum_{a \in A} x_a y_a, \tag{33}$$

$\ell^2(A)$ is a Hilbert space. Moreover, $\ell^2(A)$ is separable if and only if $A$ is countable. It follows that the sequence space $\ell^2 = \ell^2(\mathbb{N})$ is the separable Hilbert space of square-summable sequences. Due to the Riesz-Fischer theorem, every infinite-dimensional Hilbert space is isomorphic to $\ell^2$.

**Definition 3.7** (Gram matrix)**.** [8] Let $x_1, x_2, \ldots, x_n \in \mathcal{X}$ for some inner product space $\mathcal{X}$ equipped with $\langle \cdot, \cdot \rangle$. We say $G$ is a *Gram matrix* (or *Gramian*) for the sequence of vectors $x_1, x_2, \ldots, x_n$ with respect to $\langle \cdot, \cdot \rangle$ if $G = \big[\langle x_i, x_j \rangle\big]_{ij}$.

**Example 3.8.** Consider the vectors in $\mathbb{R}^3$:

$$v_1 = \begin{bmatrix} v_{11} \\ v_{21} \\ v_{31} \end{bmatrix}, \qquad v_2 = \begin{bmatrix} v_{12} \\ v_{22} \\ v_{32} \end{bmatrix}, \qquad v_3 = \begin{bmatrix} v_{13} \\ v_{23} \\ v_{33} \end{bmatrix}, \qquad v_4 = \begin{bmatrix} v_{14} \\ v_{24} \\ v_{34} \end{bmatrix}.$$

The Gram matrix for these vectors is

$$G = \begin{bmatrix} v_1^\top v_1 & v_1^\top v_2 & v_1^\top v_3 & v_1^\top v_4 \\ v_2^\top v_1 & v_2^\top v_2 & v_2^\top v_3 & v_2^\top v_4 \\ v_3^\top v_1 & v_3^\top v_2 & v_3^\top v_3 & v_3^\top v_4 \\ v_4^\top v_1 & v_4^\top v_2 & v_4^\top v_3 & v_4^\top v_4 \end{bmatrix}.$$

If $V$ is a matrix whose columns are $v_1$, $v_2$, $v_3$, $v_4$, then we can write $G = V^\top V$.

**Theorem 3.9.** *[8] A matrix $G$ is a Gram matrix if and only if $G$ is positive semidefinite.*

*Proof.* ($\Rightarrow$) Suppose $G$ is the Gram matrix of $x_1, x_2, \ldots, x_n$ with respect to $\langle \cdot, \cdot \rangle$. Let $c_1, c_2, \ldots, c_n \in \mathbb{R}$. Then $G$ is positive semidefinite because

$$\sum_{i=1}^{n} \sum_{j=1}^{n} c_i c_j \langle x_i, x_j \rangle = \left\langle \sum_{i=1}^{n} c_i x_i, \sum_{j=1}^{n} c_j x_j \right\rangle = \left\| \sum_{i=1}^{n} c_i x_i \right\|^2 \geq 0. \tag{34}$$

($\Leftarrow$) Suppose $G$ is positive semidefinite. Then $G$ can be factored as $G = B^\top B$. Let $b_1, b_2, \ldots, b_n$, be the columns of $B$. Then $G = [b_i^\top b_j]_{ij}$. Hence $G$ is the Gram matrix of $b_1, b_2, \ldots, b_n$ with respect to the dot product. $\square$

**Definition 3.10** (Symmetric bilinear form)**.** A *symmetric bilinear form* is a map $k : \mathcal{X} \times \mathcal{X} \to \mathbb{R}$ over a vector space $\mathcal{X}$ such that, for all $x, y, z \in \mathcal{X}$, $\alpha, \beta \in \mathbb{R}$,

1. $k(x, y) = k(y, x)$ and (symmetry)
2. $k(\alpha x + \beta y, z) = \alpha k(x, z) + \beta k(y, z)$. (bilinear)

This can be thought of as a generalization of an inner product which is symmetric and bilinear, but not necessarily positive definite.

**Example 3.11.** If $U = \{u_1, u_2, \ldots, u_n\}$ is a basis for $\mathcal{X}$, then we can define a matrix $K = \big[k(u_i, u_j)\big]_{ij}$. Clearly, $K$ is symmetric since $k(u_i, u_j) = k(u_j, u_i)$. Let $v = \sum_{i=1}^n \alpha_i u_i$ and $w = \sum_{i=1}^n \beta_i u_i$ be vectors with respect to $U$ and let $x = [\alpha_i]_{i=1}^n$ and $y = [\beta_i]_{i=1}^n$. Then

$$k(v, w) = k\left(\sum_{i=1}^n \alpha_i u_i, \sum_{j=1}^n \beta_j u_j\right) = \sum_{i=1}^n \sum_{j=1}^n \alpha_i \beta_j k\left(u_i, u_j\right) = x^\top K y. \tag{35}$$

If $K = I$, then $v = x$, $w = y$, and $k(v, w) = v^\top w$ is simply the dot product. Otherwise, if $K$ is positive semidefinite, then $K = B^\top B$ implies

$$k(v, w) = x^\top B^\top B y = (Bx)^\top (By) \tag{36}$$

In this case, $k(v, w)$ is just the dot product after the transformation under $B$. Notice that if $U$ is merely a subset of $\mathcal{X}$, then $v$ and $w$ no longer have unique representations, but equations (35) and (36) are still valid for all $v, w \in \operatorname{span} U$.

We say that $k$ is *positive semidefinite* if $K = [k(u_i, u_j)]_{ij}$ is a positive semidefinite matrix for any finite subset $U = \{u_1, u_2, \ldots, u_n\} \subseteq \mathcal{X}$. Then $K$ is a Gram matrix with respect to some set of transformed vectors related to $U$ and some inner product related to $k$. In the next subsection, we will show that $k$ still corresponds to some inner product even if $k$ is not bilinear.

## 3.2 Kernels

The development of *kernel functions* can be traced back to the beginning of the twentieth century when David Hilbert and James Mercer were studying integral equations [7]. Hilbert proved some important results in [6] about the eigenvalues of an integral operator whose kernel function is of *definite* type. Expanding on Hilbert's work, Mercer provided the necessary conditions in [12] that allow a kernel function to be written in terms of the eigenvalues and eigenfunctions of the integral operator. This result became known as Mercer's theorem. See **??**. A simplified version of Mercer's theorem states that a kernel function can be written as an inner product in a higher-dimensional space.

Hilbert space theory and Mercer's theorem led to a number of advances in functional analysis over the next few decades. Notably, in 1950, Nachman Aronszajn introduced reproducing kernel Hilbert spaces in [2]. This work expanded

on Mercer's theorem and shows that a kernel generates a Hilbert space whose inner product agrees with the kernel.

Later, the work of Mercer and Aronszajn inspired the application of kernels in machine learning. A *kernel method* is an adaptation of a machine learning algorithm that replaces a dot product with a kernel function. The earliest research involving kernel methods was in 1964 by Mark Aizerman et al. [1]. In the 1990s, Bernhard Schölkopf et al. used Aizerman's technique to develop kernel PCA and suggested the kernel trick could work in other cases too. In Section 4, we will look at the kernel method applied to the PCA algorithm. For now, we will examine the mathematics behind kernel methods.

**Definition 3.12** (Kernel)**.** [14] Let $k : \mathcal{X} \times \mathcal{X} \to \mathbb{R}$ be defined on a nonempty set $\mathcal{X}$. Similar to the Gram matrix, define a *kernel matrix* for a set of vectors $\{x_1, x_2, \ldots, x_n\} \subseteq X$ with respect to $k(\cdot, \cdot)$ as $K = [k(x_i, x_j)]_{ij}$. Then $k$ is a *kernel function* (or just *kernel*) if the following hold:

1. $k(x, y) = k(y, x)$, for all $x, y \in \mathcal{X}$ and (symmetry)
2. any kernel matrix $K$ generated by $k$ is positive semidefinite.

We can easily show some properties that kernels have in common with inner products.

**Lemma 3.13.** *[14] Let $k$ be a kernel. Then the following hold:*

1. *$k(x, x) \geq 0$ for all $x \in \mathcal{X}$ and* *(positive semidefinite)*
2. *$k(x, y)^2 \leq k(x, x)k(y, y)$.* *(Cauchy-Schwarz inequality)*

*Proof.* Let $x, y \in \mathcal{X}$.

1. The $1 \times 1$ kernel matrix $[k(x, x)]$ is positive semidefinite. So, $k(x, x) \geq 0$.
2. The $2 \times 2$ kernel matrix

$$K = \begin{bmatrix} k(x,x) & k(x,y) \\ k(y,x) & k(y,y) \end{bmatrix} \tag{37}$$

is positive semidefinite. Let $v = \begin{bmatrix} k(y,y) \\ -k(x,y) \end{bmatrix}$. Then

$$\begin{aligned} 0 &\leq v^\top K v \\ &= \begin{bmatrix} k(y,y) \\ -k(x,y) \end{bmatrix}^\top \begin{bmatrix} k(x,x)k(y,y) - k(x,y)^2 \\ 0 \end{bmatrix} \\ &= k(y,y)\left[k(x,x)k(y,y) - k(x,y)^2\right]. \end{aligned} \tag{38}$$

Then $v^\top K v \geq 0$ implies $k(x, y)^2 \leq k(x, x)k(y, y)$. □

**Definition 3.14** (Feature map)**.** [7] Let $\mathcal{X}$ be a nonempty set and let $\mathbb{R}^{\mathcal{X}}$ be the vector space of real-valued functions on $\mathcal{X}$. A *feature map* is a function $\Phi : \mathcal{X} \to \mathcal{H}$ for some subspace $\mathcal{H} \subseteq \mathbb{R}^{\mathcal{X}}$. In this context, $\mathcal{H}$ is referred to as the *feature space* and its elements $\Phi(x) \in \mathcal{H}$ are called *features*.

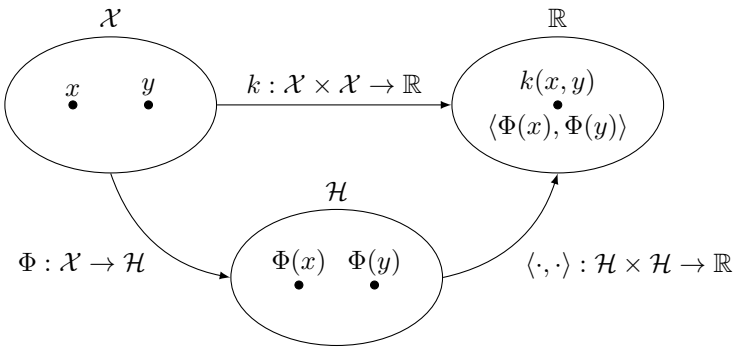


Figure 2: Kernel map diagram.

Starting with a kernel $k : \mathcal{X} \times \mathcal{X} \to \mathbb{R}$, we want to construct a feature map $\Phi : \mathcal{X} \to \mathcal{H}$ and inner product $\langle \cdot, \cdot \rangle : \mathcal{H} \times \mathcal{H} \to \mathbb{R}$ which satisfies

$$k(x, y) = \langle \Phi(x), \Phi(y) \rangle , \tag{39}$$

for all $x, y \in \mathcal{X}$. Then the linear span of $\Phi(\mathcal{X}) = \{\Phi(x) \mid x \in \mathcal{X}\}$ will be an inner product space. Since inner product spaces can be completed, we can make this a Hilbert space. See Figure 2.

**Constructing a feature map.** Consider the map[4] $\Phi(x) = k(x, \cdot)$, for all $x \in \mathcal{X}$. Note that by the symmetry of $k$, we can write $\Phi(x) = k(x, \cdot) = k(\cdot, x)$. Taking the linear span of $\Phi(\mathcal{X})$ gives us

$$H_0 = \operatorname{span} \Phi(\mathcal{X}) = \left\{ f = \sum_{i=1}^{n} \alpha_i k(x_i, \cdot) \;\middle|\; \begin{array}{r} \forall n \in \mathbb{N}, \\ x_1, x_2, \ldots, x_n \in \mathcal{X}, \\ \alpha_1, \alpha_2, \ldots, \alpha_n \in \mathbb{R} \end{array} \right\}, \tag{40}$$

which forms a subspace of $\mathbb{R}^{\mathcal{X}}$. By Definition 3.14, $H_0$ is a feature space.

**Constructing an inner product.** Let $f, g \in H_0$. Then there exist $n, m \in \mathbb{N}$, $(\alpha_i)_{i=1}^n$, $(\beta_j)_{j=1}^m$, $(x_i)_{i=1}^n$, $(y_j)_{j=1}^m$ such that

$$f = \sum_{i=1}^{n} \alpha_i k(x_i, \cdot) \quad \text{and} \quad g = \sum_{j=1}^{m} \beta_j k(y_j, \cdot). \tag{41}$$

Define $\langle \cdot, \cdot \rangle_{H_0} : H_0 \times H_0 \to \mathbb{R}$ as

$$\langle f, g \rangle_{H_0} = \sum_{i=1}^{n} \sum_{j=1}^{m} \alpha_i \beta_j k(x_i, y_j). \tag{42}$$

[4]Here, $\Phi : \mathcal{X} \to (\mathcal{X} \to \mathbb{R})$ is just the *curried* form of the binary function $k : \mathcal{X} \times \mathcal{X} \to \mathbb{R}$. For example, $\{k(x, \cdot) \mid x \in \mathcal{X}\}$ describes a set of unary functions curried from the binary function $k$.

Letting $m = 1$, $\beta_1 = 1$, $y_1 = y$ in equations (41) and (42), then $g = k(y, \cdot)$. This shows that $k$ has the *reproducing property*

$$\langle f, k(y, \cdot)\rangle_{H_0} = \sum_{i=1}^{n} \alpha_i k(x_i, y) = f(y), \tag{43}$$

for all $y \in \mathcal{X}$. Similarly, letting $n = 1$, $\alpha_1 = 1$, $x_1 = x$, we have

$$k(x, y) = \langle k(x, \cdot), k(y, \cdot)\rangle_{H_0} = \langle \Phi(x), \Phi(y)\rangle_{H_0}, \tag{44}$$

for all $x, y \in \mathcal{X}$. Now we will show that $\langle \cdot, \cdot \rangle_{H_0}$ is an inner product.

1. Since $k$ is symmetric, we have

$$\langle f, g\rangle_{H_0} = \sum_{i=1}^{n}\sum_{j=1}^{m} \alpha_i \beta_j k(x_i, y_j) = \sum_{j=1}^{m}\sum_{i=1}^{n} \beta_j \alpha_i k(y_j, x_i) = \langle g, f\rangle_{H_0}. \tag{45}$$

2. By rearrangement, equation (42) becomes

$$\langle f, g\rangle_{H_0} = \sum_{j=1}^{m} \beta_j \sum_{i=1}^{n} \alpha_i k(x_i, y_j) = \sum_{j=1}^{m} \beta_j f(y_j). \tag{46}$$

   Then for all $f_1, f_2 \in \operatorname{span} \Phi(\mathcal{X})$ and $\alpha, \gamma \in \mathbb{R}$,

$$\begin{aligned} \langle \alpha f_1 + \gamma f_2, g\rangle_{H_0} &= \sum_{j=1}^{m} \beta_j (\alpha f_1 + \gamma f_2)(y_j) \\ &= \alpha \sum_{j=1}^{m} \beta_j f_1(y_j) + \gamma \sum_{j=1}^{m} \beta_j f_2(y_j) \\ &= \alpha \langle f_1, g\rangle_{H_0} + \gamma \langle f_2, g\rangle_{H_0}. \end{aligned} \tag{47}$$

3. Since $k$ is positive semidefinite, by Lemma 3.13 part 1

$$\langle f, f\rangle_{H_0} = \sum_{i=1}^{n}\sum_{j=1}^{n} \alpha_i \alpha_j k(x_i, x_j) \geq 0. \tag{48}$$

   Now let $f_1, f_2, \ldots, f_p$ be functions and $\gamma_1, \gamma_2, \ldots, \gamma_p \in \mathbb{R}$. Then by bilinearity,

$$\sum_{i=1}^{p}\sum_{j=1}^{p} \gamma_i \gamma_j \langle f_i, f_j\rangle_{H_0} = \left\langle \sum_{i=1}^{p} \gamma_i f_i, \sum_{j=1}^{p} \gamma_j f_j \right\rangle_{H_0} \geq 0. \tag{49}$$

   Thus, $\langle \cdot, \cdot \rangle_{H_0}$ is a kernel. Then by equation (43) and Lemma 3.13 part 2,

$$f(x)^2 = \langle f, k(x, \cdot)\rangle_{H_0}^2 \leq \langle f, f\rangle_{H_0} \langle k(x, \cdot), k(x, \cdot)\rangle_{H_0}, \tag{50}$$

   for all $x \in \mathcal{X}$. If $\langle f, f\rangle_{H_0} = 0$, then $f(x)^2 = 0$ implies $f = 0$.

**Constructing a Hilbert space.** Now that we know $\langle \cdot, \cdot \rangle_{H_0}$ is an inner product, $H_0$ is an inner product space. By [11], this can be completed with respect to the induced metric. Then

$$\mathcal{H} = \overline{\operatorname{span} \Phi(\mathcal{X})} = \overline{\operatorname{span}\{k(x, \cdot) \mid x \in \mathcal{X}\}} \tag{51}$$

is a Hilbert space with inner product $\langle \cdot, \cdot \rangle_{\mathcal{H}} : \mathcal{H} \times \mathcal{H} \to \mathbb{R}$. Since $\mathcal{H}$ contains all its limit points, functions in this space have the form the sequences $(\alpha_i)_{i=1}^{\infty}$ and $(x_i)_{i=1}^{\infty}$ determine a function $f \in \mathcal{H}$ such that

$$f = \sum_{i=1}^{\infty} \alpha_i k(x_i, \cdot), \tag{52}$$

provided the series converges.

**Definition 3.15** (Reproducing kernel Hilbert space)**.** [7] Let $\mathcal{H}$ be a Hilbert space of functions $\mathcal{X} \to \mathbb{R}$ on some nonempty set $\mathcal{X}$. Then $\mathcal{H}$ is a *reproducing kernel Hilbert space* (RKHS) if there exists a function $k : \mathcal{X} \times \mathcal{X} \to \mathbb{R}$ such that for all $f \in \mathcal{H}$ and $x \in \mathcal{X}$,

1. $f(x) = \langle f, k(x, \cdot) \rangle$ and (reproducing property)
2. $\mathcal{H} = \overline{\operatorname{span}\{k(x, \cdot) \mid x \in \mathcal{X}\}}$. (spanning property)

Fix $y \in \mathcal{X}$ and treat $k(x, y)$ as a univariate function of $x \in \mathcal{X}$. By the reproducing property, $k(x, y) = \langle k(x, \cdot), k(y, \cdot) \rangle$. Define $\Phi(x) = k(x, \cdot)$ to give the desired result in equation (39).

## 3.3 Related notions of an RKHS

In the previous section, we showed that a (symmetric positive semidefinite) kernel defines an RKHS. Presently, we will look at three alternative methods for constructing an RKHS.

1. **Positive semidefinite kernels.** Due to Aronszajn [2], a reproducing kernel will generate a unique RKHS. Moreover, a kernel is unique to its RKHS. See Theorems 3.16 and 3.17.

2. **Continuous linear functionals.** By the Riesz representation theorem, if every evaluation functional is continuous, then every function in the Hilbert space can be reproduced at every point. In this way, a reproducing kernel can be defined.

3. **Feature maps.** A explicit feature map with an inner product can be used to define a kernel as in equation (39). Alternatively, by Mercer's theorem 3.21, a kernel has a series expansion which allows us to define a feature map in terms of eigenvalues and eigenfunctions.

### 3.3.1 Positive semidefinite kernels.

**Theorem 3.16** (Moore-Aronszajn theorem)**.** *[2] Let $\mathcal{X}$ be a nonempty set and $k : \mathcal{X} \times \mathcal{X} \to \mathbb{R}$ be a kernel. Then there exists a unique RKHS for which $k$ is a reproducing kernel.*

*Proof.* For existence, we summarize the construction provided in Section 3.2.

1. $k$ defines a feature map $\Phi : \mathcal{X} \to \mathbb{R}^{\mathcal{X}}$ such that $\Phi(x) = k(x, \cdot)$ for all $x \in \mathcal{X}$.
2. The linear span of $\Phi(\mathcal{X})$ is a feature space.
3. $\Phi$ defines an inner product $\langle \cdot, \cdot \rangle_{H_0} : H_0 \times H_0 \to \mathbb{R}$ in equation (42).
4. Completing the feature space yields a Hilbert space $\mathcal{H}$ with inner product $\langle \cdot, \cdot \rangle : \mathcal{H} \times \mathcal{H} \to \mathbb{R}$.
5. $k$ has the reproducing property $\langle f, k(x, \cdot) \rangle = f(x)$ shown by equation (43).

For uniqueness, suppose $k$ is a reproducing kernel for Hilbert spaces $\mathcal{H}$ and $\mathcal{L}$.

Let $f \in \mathcal{L}$. Then we can write $\mathcal{L} = \mathcal{H} \oplus \mathcal{H}^{\perp}$ as the orthogonal decomposition of $\mathcal{L}$. There exist $g \in \mathcal{H}$ and $g^{\perp} \in \mathcal{H}^{\perp}$ such that $f = g + g^{\perp}$. Let $x \in \mathcal{X}$. Then $k(x, \cdot) \in \mathcal{H}$ implies $\left\langle g^{\perp}, k(x, \cdot) \right\rangle_{\mathcal{L}} = 0$. Thus

$$f(x) = \langle g, k(x, \cdot) \rangle_{\mathcal{L}} + \left\langle g^{\perp}, k(x, \cdot) \right\rangle_{\mathcal{L}} = \langle g, k(x, \cdot) \rangle_{\mathcal{L}} = \langle g, k(x, \cdot) \rangle_{\mathcal{H}} = g(x). \quad (53)$$

Thus $f = g \in \mathcal{H}$ implies $\mathcal{L} \subseteq \mathcal{H}$. We can repeat this argument for $f \in \mathcal{H}$ to show $f \in \mathcal{L}$. □

**Theorem 3.17.** *[2] A reproducing kernel for an RKHS is unique.*

*Proof.* Let $\mathcal{H}$ be an RKHS of functions $\mathcal{X} \to \mathbb{R}$ for some set $\mathcal{X} \neq \emptyset$. Suppose $k$ and $\ell$ reproducing kernels for $\mathcal{H}$. Denote $k_x = k(x, \cdot)$ and $\ell_x = \ell(x, \cdot)$, for all $x \in \mathcal{X}$. By the reproducing property,

$$\begin{aligned} \|k_x - \ell_x\|^2 &= \langle k_x - \ell_x, k_x - \ell_x \rangle \\ &= \langle k_x - \ell_x, k_x \rangle - \langle k_x - \ell_x, \ell_x \rangle \\ &= k_x(x) - \ell_x(x) - k_x(x) + \ell_x(x) \\ &= 0. \end{aligned} \quad (54)$$

It follows that $k_x - \ell_x$ is the zero function. Hence, $k(x, \cdot) = \ell(x, \cdot)$ for all $x \in \mathcal{X}$. By symmetry, $k(\cdot, x) = \ell(\cdot, x)$. Therefore, $k = \ell$. □

### 3.3.2 Continuous linear functionals.

Suppose we have a Hilbert space and we want to know if it is an RKHS. To do this, we need to construct a kernel from the Hilbert space.

**Example 3.18.** Consider a Hilbert space $\mathbb{R}^n$ with the dot product. Note that the vectors in $\mathbb{R}^n$ are actually just sequences $\mathbb{N} \to \mathbb{R}$ with some vector operations. It is straightforward to show that the reproducing kernel is the Kroenecker delta $k(i,j) = \delta_{ij}$ for all $i, j \in \{1, 2, \ldots, n\}$. This generates the standard basis $\{e_i\}_{i=1}^n$, where $e_i = k(i, \cdot)$. So, $\mathbb{R}^n$ is an RKHS.

We can replace $\mathbb{N}$ with any other index set with cardinality $n$, say $I_n = \{\frac{0}{n}, \frac{1}{n}, \frac{2}{n}, \ldots, \frac{n-1}{n-1}\}$. Then an indexed family $f_n : I_n \to \mathbb{R}$ has a vector representation in $\mathbb{R}^n$. Letting $n$ tend to infinity, we have $I_\infty = \mathbb{Q} \cap [0,1]$. By completion, this is the space of functions $\{f \mid f : [0,1] \to \mathbb{R}\}$. In one sense, $f_n \in \mathbb{R}^n$ is a point in $n$-dimensional space and, in another sense, $f_n : I_n \to \mathbb{R}$ is the discretization of a function $f : [0,1] \to \mathbb{R}$. This way, real-valued functions on $[0,1]$ can be interpreted as infinite-dimensional vectors.

Now consider the Hilbert space $L^2([0,1]) = \{f \mid \int_{[0,1]} f^2 < \infty\}$ with inner product $\langle f, g \rangle = \int_{[0,1]} fg$. Then for all $x \in [0,1]$,

$$f(x) = \int_{[0,1]} \delta(x - t) f(t) \, \mathrm{d}\, t, \tag{55}$$

where $\delta$ is the Dirac delta. If $L^2([0,1])$ is an RKHS, then by Theorem 3.17, $k(x,t) = \delta(x - t)$ is the unique reproducing kernel. But $\int_{\mathcal{X}} \delta^2 = \infty$ implies $\delta \notin L^2([0,1])$. Therefore, $L^2([0,1])$ is not an RKHS.

The Kroenecker delta and Dirac delta in the example reproduce functions in the Hilbert space with the inner product.

**Definition 3.19** (Evaluation functional)**.** Let $\mathcal{X}$ be a nonempty set and $\mathcal{H}$ be a Hilbert space of functions $\mathcal{X} \to \mathbb{R}$. Then for all $x \in \mathcal{X}$, let $\delta_x : \mathcal{H} \to \mathbb{R}$ such that $\delta_x(f) = f(x)$, for each $f \in \mathcal{H}$. We call $\delta_x$ the *evaluation functional* at $x$.

**Theorem 3.20** (Riesz representation theorem)**.** *[20] Let $\delta : H \to \mathbb{R}$ be a continuous linear functional defined on a Hilbert space $\mathcal{H}$. Then there exists a unique element $g \in \mathcal{H}$ such that $\delta(g) = \langle f, g \rangle_H$ for all $g \in \mathcal{H}$.*

Suppose the evaluation functional $\delta_x$ is continuous on $\mathcal{H}$ for every $x \in \mathcal{X}$. If $x, y \in \mathcal{X}$, then, by the Riesz representation theorem, there exist $k_x, k_y \in \mathcal{H}$ such that for all $f \in \mathcal{H}$,

$$f(x) = \delta_x(f) = \langle f, k_x \rangle, \qquad f(y) = \delta_y(f) = \langle f, k_y \rangle. \tag{56}$$

Let $k : \mathcal{X} \times \mathcal{X} \to \mathbb{R}$ be defined as $k(x,y) = k_x(y)$. Then by symmetry of the inner product,

$$k(x,y) = k_x(y) = \langle k_x, k_y \rangle = \langle k_y, k_x \rangle = k_y(x) = k(y,x). \tag{57}$$

It follows that $k$ is a kernel, $\Phi(x) = k_x$ is a feature map, and $\mathcal{H}$ is an RKHS.

### 3.3.3 Feature maps

Mercer's theorem provides a result similar to Aronszajn's, but without the context of an RKHS. Rather, the focus is the decomposition of an integral operator.

**Theorem 3.21** (Mercer's theorem)**.** *[12] Let $k : \mathcal{X} \times \mathcal{X} \to \mathbb{R}$ be a continuous bounded kernel on a compact set $\mathcal{X}$. Define the Hilbert-Schmidt integral operator $T_k : L^2(\mathcal{X}) \to L^2(\mathcal{X})$ as*

$$(T_k f)(x) = \int_{\mathcal{X}} k(x,t) f(t) \, \mathrm{d}t. \tag{58}$$

*Then there exists an orthonormal basis $\{\psi_i\}_{i=1}^{\infty}$ of eigenfunctions of $T_k$ and corresponding eigenvalues $(\lambda_i)_{i=1}^{\infty}$ with $\lambda_i \geq 0$, for all $i \in \mathbb{N}$. Moreover, for all $x, y \in \mathcal{X}$,*

$$k(x,y) = \sum_{i=1}^{\infty} \lambda_i \psi_i(x) \psi_i(y), \tag{59}$$

*where convergence is uniform.*

See [4] for a sketch of this proof. Otherwise, this is main result proved in [12].

Using equation (59), we can define a feature map $\Phi : \mathcal{X} \to \mathcal{H}$ such that for all $x \in \mathcal{X}$,

$$\Phi(x) = \sum_{i=1}^{\infty} \sqrt{\lambda_i} \psi_i(x). \tag{60}$$

In the finite-dimensional case, we have $\Phi(x) = \left[\sqrt{\lambda_j}\psi_j(x)\right]_{j=1}^{d}$. It follows that $k(x,y) = \langle \Phi(x), \Phi(y) \rangle$.

## 3.4 Constructing kernels

**Theorem 3.22.** *[14, 19] Suppose $k_1$ and $k_2$ are kernels on $\mathcal{X} \times \mathcal{X}$. The following functions kernels.*

1. *$k(x,y) = a_1 k_1(x,y) + a_2 k_2(x,y)$ for all $a_1, a_2 \geq 0$.*
2. *$k(x,y) = k_1(x,y) k_2(x,y)$.*
3. *$k(x,y) = a_0 + a_1 k_1(x,y) + a_2 k_1(x,y)^2 + \cdots + a_n k_1(x,y)^n$ for all $n \in \mathbb{N}$ and $a_0, \ldots, a_n \geq 0$.*
4. *$k(x,y) = k_1(h(x), h(y))$ for all $h : \mathcal{X} \to \mathcal{X}$.*
5. *$k(x,y) = g(x) g(y)$ for all $g : \mathcal{X} \to \mathbb{R}$.*
6. *$k(x,y) = \exp(k_1(x,y))$.*

*Proof.* Let $x_1, \ldots, x_n \in \mathcal{X}$ and $c_1, \ldots, c_n \in \mathbb{R}$.

1. Let $k = a_1 k_1 + a_2 k_2$ for $a_1, a_2 \geq 0$. Since $k_1$ and $k_2$ are symmetric,

$$k(x, y) = a_1 k_1(x, y) + a_2 k_2(x, y) = a_1 k_1(y, x) + a_2 k_2(y, x) = k(y, x),$$

for all $x, y \in \mathcal{X}$. So, $k$ is symmetric.

Since $k_1$ and $k_2$ are positive semidefinite and $a_1, a_2 \geq 0$,

$$\begin{aligned}\sum_{i=1}^{n}\sum_{j=1}^{n} c_i c_j k(x_i, x_j) &= \sum_{i=1}^{n}\sum_{j=1}^{n} c_i c_j (a_1 k_1(x_i, x_j) + a_2 k_2(x_i, x_j)) \\ &= a_1 \sum_{i=1}^{n}\sum_{j=1}^{n} c_i c_j k_1(x_i, x_j) + a_2 \sum_{i=1}^{n}\sum_{j=1}^{n} c_i c_j k_2(x_i, x_j) \\ &\geq 0.\end{aligned}$$

So, $k$ is positive semidefinite.

2. Let $k = k_1 k_2$. Define $K$ so that $[K]_{ij} = k(x_i, x_j) = k_1(x_i, x_j) k_2(x_i, x_j)$. Let $K_1$ and $K_2$ be the Gram matrices for $k_1$ and $k_2$, respectively. Then $K_1, K_2$ have orthonormal eigenvectors and nonnegative eigenvalues such that

$$\begin{aligned}K_1 &= V L V^\top \\ &= \begin{bmatrix} v_{11} & \cdots & v_{1n} \\ \vdots & \ddots & \vdots \\ v_{n1} & \cdots & v_{nn} \end{bmatrix} \begin{bmatrix} \lambda_1 & \cdots & 0 \\ \vdots & \ddots & \vdots \\ 0 & \cdots & \lambda_n \end{bmatrix} \begin{bmatrix} v_{11} & \cdots & v_{n1} \\ \vdots & \ddots & \vdots \\ v_{1n} & \cdots & v_{nn} \end{bmatrix} \\ &= \begin{bmatrix} \sum_{j=1}^{n} \lambda_j v_{1j} v_{1j} & \cdots & \sum_{j=1}^{n} \lambda_j v_{nj} v_{1j} \\ \vdots & \ddots & \vdots \\ \sum_{j=1}^{n} \lambda_j v_{1j} v_{nj} & \cdots & \sum_{j=1}^{n} \lambda_j v_{nj} v_{nj} \end{bmatrix} \\ &= \sum_{j=1}^{n} \lambda_j \begin{bmatrix} v_{1j} v_{1j} & \cdots & v_{nj} v_{1j} \\ \vdots & \ddots & \vdots \\ v_{1j} v_{nj} & \cdots & v_{nj} v_{nj} \end{bmatrix}\end{aligned}$$

and

$$K_2 = U M U^\top = \sum_{j=1}^{n} \mu_j \begin{bmatrix} u_{1j} u_{1j} & \cdots & u_{nj} u_{1j} \\ \vdots & \ddots & \vdots \\ u_{1j} u_{nj} & \cdots & u_{nj} u_{nj} \end{bmatrix}.$$

Let $\mathbf{v}_i = \begin{bmatrix} v_{1i} & \cdots & v_{ni} \end{bmatrix}^\top$ and $\mathbf{u}_j = \begin{bmatrix} u_{1j} & \cdots & u_{nj} \end{bmatrix}$, for all $i, j =$

$1, 2, \ldots, n$. Then

$$\begin{aligned}
K &= K_1 \circ K_2 \\
&= \sum_{i=1}^{n} \lambda_i \begin{bmatrix} v_{1i}v_{1i} & \cdots & v_{ni}v_{1i} \\ \vdots & \ddots & \vdots \\ v_{1i}v_{ni} & \cdots & v_{ni}v_{ni} \end{bmatrix} \circ \sum_{j=1}^{n} \mu_j \begin{bmatrix} u_{1j}u_{1j} & \cdots & u_{nj}u_{1j} \\ \vdots & \ddots & \vdots \\ u_{1j}u_{nj} & \cdots & u_{nj}u_{nj} \end{bmatrix} \\
&= \sum_{i=1}^{n}\sum_{j=1}^{n} \lambda_i\mu_j \begin{bmatrix} v_{1i}u_{1j}v_{1i}u_{1j} & \cdots & v_{1i}u_{1j}v_{ni}u_{nj} \\ \vdots & \ddots & \vdots \\ v_{ni}u_{nj}v_{1i}u_{1j} & \cdots & v_{ni}u_{nj}v_{ni}u_{nj} \end{bmatrix} \\
&= \sum_{i=1}^{n}\sum_{j=1}^{n} \lambda_i\mu_j \begin{bmatrix} v_{1i}u_{1j} \\ \vdots \\ v_{ni}u_{nj} \end{bmatrix} \begin{bmatrix} v_{1i}u_{1j} & \cdots & v_{ni}u_{nj} \end{bmatrix} \\
&= \sum_{i=1}^{n}\sum_{j=1}^{n} \lambda_i\mu_j (\mathbf{v}_i \circ \mathbf{u}_j)(\mathbf{v}_i \circ \mathbf{u}_j)^\top,
\end{aligned}$$

where $\circ$ is the Hadamard product. Each $(\mathbf{v}_i \circ \mathbf{u}_j)(\mathbf{v}_i \circ \mathbf{u}_j)^\top$ is a symmetric positive semidefinite matrix. Since $K_1, K_2$ are positive semidefinite, we have $\lambda_i, \mu_i > 0$. Then $K$ is symmetric positive semidefinite.

3. By part 2, $k_1, k_1^2, \ldots, k_1^n$ are kernels. By part 1, $a_0 + a_1k_1 + a_2k_1^2 + \cdots + a_nk_1^n$ is a kernel.

4. Since $y_i = h(x_i) \in \mathcal{X}$ for all $i = 1, 2, \ldots, n$, we have

$$\begin{aligned}
\sum_{i=1}^{n}\sum_{j=1}^{n} c_ic_jk(x_i, x_j) &= \sum_{i=1}^{n}\sum_{j=1}^{n} c_ic_jk_1(h(x_i), h(x_j)) \\
&= \sum_{i=1}^{n}\sum_{j=1}^{n} c_ic_jk_1(y_i, y_j) \\
&\geq 0.
\end{aligned}$$

5. Let $g : \mathcal{X} \to \mathbb{R}$ and let $c_ig(x_i) = y_i \in \mathbb{R}$. If $k(x, y) = g(x)g(y)$, then

$$\begin{aligned}
\sum_{i=1}^{n}\sum_{j=1}^{n} c_ic_jk(x_i, x_j) &= \sum_{i=1}^{n}\sum_{j=1}^{n} c_ig(x_i)c_jg(x_j) \\
&= \sum_{i=1}^{n}\sum_{j=1}^{n} y_iy_j \\
&= \left(\sum_{i=1}^{n} y_i\right)^2 \\
&\geq 0.
\end{aligned}$$

6. Let $K_1$ be the Gram matrix for $k_1$. If $K_1 v = \lambda v$, then $K_1^m = \lambda^m v$ for all $m \in \mathbb{N}$. So,

$$(\exp K_1)v = \sum_{m=0}^{\infty} \frac{K_1^m v}{m!} = \sum_{m=0}^{\infty} \frac{\lambda^m v}{m!} = e^{\lambda} v.$$

Then $K = \exp K_1$ has eigenvalues $e^{\lambda}$. Since $K_1$ is positive semidefinite, it has real eigenvalues so that $e^{\lambda} > 0$. It follows that $K$ is positive definite.

□

**Theorem 3.23** (Gaussian kernel)**.** *The function $k : \mathbb{R}^n \times \mathbb{R}^n \to \mathbb{R}$ defined by*

$$k(x, y) = \exp\left(\frac{-\|x - y\|^2}{2\sigma^2}\right), \tag{61}$$

*is a kernel.*

*Proof.* Notice that

$$\begin{aligned} k(x, y) &= \exp\left(\frac{-\|x - y\|^2}{2\sigma^2}\right) \\ &= \exp\left(\frac{-(\|x\|^2 - 2\langle x, y\rangle + \|y\|^2)}{2\sigma^2}\right) \\ &= \exp\left(\frac{-\|x\|^2}{2\sigma^2}\right) \exp\left(\frac{-\|y\|^2}{2\sigma^2}\right) \exp\left(\frac{\langle x, y\rangle}{\sigma^2}\right). \end{aligned}$$

Let $g(x) = \exp\left(-\|x\|^2/2\sigma^2\right)$, $k_1(x, y) = g(x)g(y)$, and $k_2(x, y) = \langle x, y\rangle / \sigma^2$. By part 5, $k_1$ is a kernel. Since $\langle x, y\rangle$ is a kernel and $1/\sigma^2 > 0$, part 1 implies $k_2$ is a kernel and part 6 implies $\exp(k_2(x, y))$ is a kernel. Therefore, by part 2,

$$k(x, y) = g(x)g(y)\exp(k_2(x, y)) = k_1(x, y)\exp(k_2(x, y)) \tag{62}$$

is a kernel. □

# 4 Kernel PCA

Recall that linear PCA generates new features from a linear combination of the input variables. PCA is an orthogonal projection that rotates the data witin the original space of input variables. These components provide a new basis that may provide more information about the structure of high-dimensional data. The work of Schölkopf, Smola, and Müller [17] generalized PCA based on the successful application of kernel methods in support vector machines by Aizerman [1]. In kernel PCA, the inner product of the input space is replaced with the inner product of a feature space. As such, the principal components, or features, of kernel PCA are nonlinear transformations of input variables.

## 4.1 Covariance matrix and kernel matrix

Let $A \in \mathbb{R}^{n \times d}$ be a centered matrix and $(a_i)_{i=1}^n$ be the rows of $A$. The PCA algorithm computes the covariance matrix $C = [\mathrm{cov}(a_i, a_j)]_{ij}$ to find a basis transformation in the input space $\mathbb{R}^d$. If $k$ is a kernel on $\mathbb{R}^n$, then it corresponds to a valid inner product in some feature space $\mathcal{H}$. By replacing the covariance matrix $C$ with a kernel matrix $K$, we are implicitly performing PCA on features $\Phi(a_1), \Phi(a_2), \ldots, \Phi(a_n)$ under some feature map $\Phi : \mathbb{R}^d \to \mathcal{H}$. We call this method kernel PCA [17].

For now, assume that the features are centered, i.e., $\frac{1}{n}\sum_{i=1}^n \Phi(a_i) = 0$. Then we compute the covariance matrix in the feature space as

$$C = \frac{1}{n-1} \sum_{i=1}^{n} \Phi(a_i) \otimes \Phi(a_i). \tag{63}$$

At this point, we need to find the eigenvalues $(\lambda_i)_{i=1}^n$ of $C$. In section 2.2, we saw that $AA^\top$ and $A^\top A$ have the same eigenvalues. It follows that we can replace the outer product with an inner product and compute the same eigenvalues. To compute the eigenvectors, notice that the SVD equation $C = USV^\top$ implies $V = S^{-1}U^\top C$.

## 4.2 Centering in the feature space

Let $\Phi : \mathcal{X} \to \mathcal{H}_k$ be a feature map determined by a kernel $k$. Since $\Phi$ may be nonlinear, the image $\Phi(x)$ of a centered vector $x \in \mathcal{X}$ is not guaranteed to be centered. For an effective PCA algorithm, it is necessary to compute the kernel matrix of centered vectors in the feature space. [17]

Given $x_1, \ldots, x_n \in \mathcal{X}$, the points

$$\Phi_0(x_i) = \Phi(x_i) - \frac{1}{n}\sum_{i=1}^{n} \Phi(x_i), \quad \text{for } i = 1, \ldots, m \tag{64}$$

are the centered feature vectors in $H_k$. Then the centered kernel matrix becomes

$$\begin{aligned}
[K_0]_{ij} &= \langle \Phi_0(x_i), \Phi_0(x_j) \rangle \\
&= \left\langle \Phi(x_i) - \frac{1}{n}\sum_{p=1}^{n} \Phi(x_p), \Phi(x_j) - \frac{1}{n}\sum_{q=1}^{n} \Phi(x_q) \right\rangle \\
&= \langle \Phi(x_i), \Phi(x_j) \rangle - \frac{1}{n}\sum_{p=1}^{n} \langle \Phi(x_p), \Phi(x_j) \rangle \\
&\qquad - \frac{1}{n}\sum_{q=1}^{n} \langle \Phi(x_i), \Phi(x_q) \rangle \\
&\qquad + \frac{1}{n^2}\sum_{p=1}^{n}\sum_{q=1}^{n} \langle \Phi(x_p), \Phi(x_q) \rangle \\
&= [K]_{ij} - \frac{1}{n}\sum_{p=1}^{n} [K]_{pj} - \frac{1}{n}\sum_{q=1}^{n} [K]_{iq} + \frac{1}{n^2}\sum_{p=1}^{n}\sum_{q=1}^{n} [K]_{pq},
\end{aligned}$$

where $K$ is the uncentered kernel matrix given by $[K]_{ij} = \langle \Phi(x_i), \Phi(x_j) \rangle$. Then the formula for the centered kernel matrix can be written as

$$K_0 = K - \text{col mean}(K) - \text{row mean}(K) + \text{mean}(K). \tag{65}$$

See notes in Appendices A.1 and A.2.

## 4.3 Kernel PCA algorithm

Let $x_1, x_2, \ldots, x_n \in \mathbb{R}^d$ be input vectors and $k : \mathbb{R}^d \times \mathbb{R}^d \to \mathbb{R}$ be a kernel. The kernel PCA algorithm outputs the transformed input vectors $\tilde{x}_1, \tilde{x}_2, \ldots, \tilde{x}_n \in \mathbb{R}^n$.

1. Compute the kernel matrix $K = [k(x_i, x_j)]_{ij}^{n \times n}$.
2. Center kernel matrix $K_0 = K - \text{col mean}(K) - \text{row mean}(K) + \text{mean}(K)$.
3. Compute eigenvalues $(\lambda_j)_{j=1}^n$ and eigenvectors $(u_j)_{j=1}^n$ of $K_0$ so that $\lambda_1 \geq \lambda_2 \geq \cdots \geq \lambda_n$.
4. Scale the eigenvectors to obtain the principal components in the RKHS: $v_j = u_j / \sqrt{\lambda_j}$ for all $j = 1, 2, \ldots, n$.
5. Transformed points can be computed using $y = \left[ \langle v_j, [k(x_i, x)]_{i=1}^n \rangle \right]_{j=1}^n$.

**Example 4.1.** Consider the problem of classifying points based on their radii. These points cannot be separated using a linear classifier in the two dimensions. However, by mapping them to a three-dimensional space, they can be separated by planes. Applying kernel PCA, these points can be sent to the RKHS associated with a Gaussian kernel without using an explicit feature map. The points in this high-dimensional feature space can then be projected onto the first three principal components to find separation boundaries. See Figure 3.

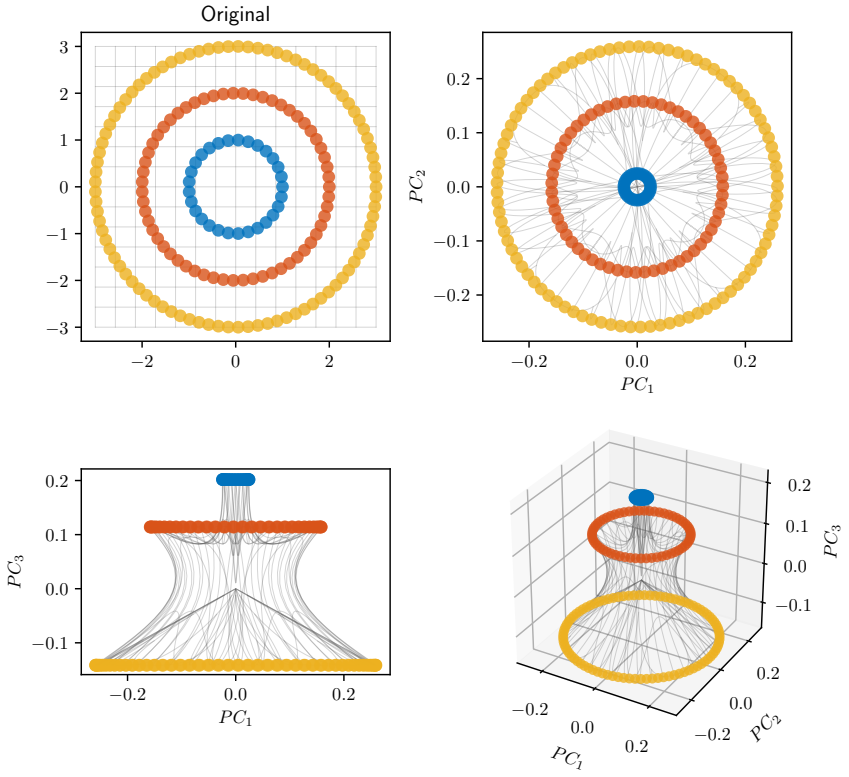


Figure 3: An idealized set of points in the plane are classified based on their radius. Kernel PCA with the Gaussian kernel is applied to find separation boundaries using the first three principal components.

# 5 Conclusion

Kernel PCA is a powerful tool that reveals nonlinear patterns in data. This is only one example of an entire class of kernel methods that extend the capabilities of linear algorithms. By exploring the theory behind kernel methods, we see how much structure exists due to symmetric positive semidefinite kernels.

# A Linear Algebra

Some useful properties of symmetric and positive definite matrices are listed below [8].

1. Symmetric matrices have orthogonal eigenvectors and real eigenvectors.
2. Positive semidefinite matrices have nonnegative eigenvalues.
3. Positive definite matrices have positive eigenvalues.
4. $A$ is positive semidefinite if and only if there exists a matrix $B$ such that $A = B^\top B$. We say $B$ is the *square root* of $A$ and write $A^{1/2} = B$.
5. $A^\top A$ and $AA^\top$ are symmetric positive semi-definite.

## A.1 Matrix operations and notation

First, we explain some notation used in this paper similar to that used in Horn and Johnson [8]. We write $(A)_{ij}$ to indicate the entry in the $i$-th row and the $j$-th column of $A$. It can be convenient to declare a matrix using index notation, such as $A = [a_{ij}] \in \mathbb{R}^{n\times d}$ or $A = [a_{ij}]^{n\times d}$ to mean $(A)_{ij} = a_{ij}$, for all $i = 1, 2, \ldots, n$; $j = 1, 2, \ldots, d$. Similar to how sets and sequences are indexed, e.g., $(c_i)_{i=1}^n$, we may indicate a vector as $[c_i]_{i=1}^d$. We may also use the notation $[1]^{n\times d}$ to mean the $n \times d$ matrix of ones.

**Definition A.1.** Let $A = [a_{ij}] \in \mathbb{R}^{n\times d}$. Define the *entry-wise mean* of $A$ as

$$\operatorname{mean}(A) = \frac{1}{nd}\sum_{i=1}^{n}\sum_{j=1}^{d} a_{ij}. \tag{66}$$

Define the *column-wise mean* of $A$ as a $1 \times d$ row vector whose $j$-th entry is the mean of column $j$ given by the formula

$$\operatorname{col\,mean}(A) = \left[\tfrac{1}{n}\textstyle\sum_{i=1}^{n} a_{ij}\right]_{j=1}^{d} = \frac{1}{n}\sum_{i=1}^{n}\left[a_{ij}\right]_{j=1}^{d} \in \mathbb{R}^{1\times d}. \tag{67}$$

Define the *row-wise mean* of $A$ as an $n \times 1$ column vector whose $i$-th entry is the mean of row $i$ given by the formula

$$\operatorname{row\,mean}(A) = \left[\tfrac{1}{d}\textstyle\sum_{j=1}^{n} a_{ij}\right]_{i=1}^{n} = \frac{1}{d}\sum_{j=1}^{d}\left[a_{ij}\right]_{i=1}^{n} \in \mathbb{R}^{n\times 1}. \tag{68}$$

Let $[a]^{p\times q}$ denote the $p \times q$ repeated matrix whose entries are all $a$. Then equations (66) to (68) can be written as

$$\text{mean}(A) = \left[\tfrac{1}{n}\right]^{1\times n} \cdot A \cdot \left[\tfrac{1}{d}\right]^{d\times 1} \tag{69}$$

$$\text{col mean}(A) = \left[\tfrac{1}{n}\right]^{1\times d} \cdot A \tag{70}$$

$$\text{row mean}(A) = A \cdot \left[\tfrac{1}{d}\right]^{d\times 1}. \tag{71}$$

## A.2 Broadcasting

Consider the sum of two real matrices $A+B$. By definition, $A$ and $B$ must both have size $n \times d$. This means we cannot add a $2 \times 3$ matrix $A$ and a $2 \times 1$ vector $b$. However, in many programming languages the sum $A + b$ would be handled using *broadcasting* [5]. In this case, $b$ is converted to a $2 \times 3$ matrix $\begin{bmatrix} b & b & b \end{bmatrix}$ so that normal matrix addition applies. Generally, broadcasting a vector $b \in \mathbb{R}^n$ to an $n \times d$ matrix can be represented as the matrix product

$$b \cdot [1]^{1\times d} = \begin{bmatrix} b_1 \\ b_2 \\ \vdots \\ b_n \end{bmatrix} \cdot \begin{bmatrix} 1 & 1 & \cdots & 1 \end{bmatrix} = \begin{bmatrix} b_1 & b_1 & \cdots & b_1 \\ b_2 & b_2 & \cdots & b_2 \\ \vdots & \vdots & & \vdots \\ b_n & b_n & \cdots & b_n \end{bmatrix}. \tag{72}$$

**Definition A.2.** For an $n \times d$ matrix $A$, we can define addition by an $n \times 1$ column vector $c$ as

$$A + c := A + c \cdot [1]^{1\times d}. \tag{73}$$

Similarly, addition by a $1 \times d$ row vector $r$ can be defined as

$$A + r := A + [1]^{n\times 1} \cdot r \tag{74}$$

and addition by a scalar $a$ can be defined as

$$A + a := A + a \cdot [1]^{n\times d}. \tag{75}$$

The left hand sides of equations (73) to (75) are more concise and intuitive than the right hand sides. Provided that the vector types are clearly defined and compatible, there should be no ambiguity when adding column vectors, row vectors, and scalars to matrices. Moreover, this method of broadcasting is consistent with scientific programming languages.

# B Code

Listing 1: PCA example

```
1 import numpy as np
2 from pca import *
3
```

```
# Random number generator for repeatability
rng = np.random.default_rng(12)
# Set data dimensions
d = 3  # number of columns (variables)
n = 30 # number of rows (observations)
# Generate uncorrelated and correlated data
A = rng.standard_normal((n,d)) # uncorrelated
B = A.copy() # correlated
B[:,0] += 4*B[:,1] + 2*B[:,2]  # correlated
# PCA projections
coeff, score, latent, mu = pca(A, n_components=2)
Ahat = score @ coeff.T + mu # reconstruct uncorrelated data
coeff, score, latent, mu = pca(B, n_components=2)
Bhat = score @ coeff.T + mu # reconstruct correlated data
# Reconstruction error
err_uncorr = np.linalg.norm(A - Ahat)
err_corr = np.linalg.norm(B - Bhat)
print(f"Uncorrelated reconstruction error: {err_uncorr}")
print(f"Correlated reconstruction error: {err_corr}")
# Plot Scatter
pairwise3dscatter(A, Ahat, "Uncorrelated Data")
pairwise3dscatter(B, Bhat, "Correlated Data")
```

Listing 2: PCA source functions

```
import numpy as np
import matplotlib.pyplot as plt

def pca(data, n_components=None):
    '''
    Perform principal component analysis.

    Parameters
    ----------
    data: array_like
        Input data.
    n_components: int
        Number of principal components to keep.

    Returns
    -------
    V: array_like
        Principal component vectors (aka coeff).
    A: array_like
        Transformed datta (aka score).
    D: array_like
        Explained variance for each principal component (aka latent).
    '''
    # Copy data as numpy array
    A = np.copy(data)
```

```
    # Center data matrix
    col_mean = A.mean(0)
    A -= col_mean
    # Compute covariance matrix
    C = A.T @ A / (A.shape[0]-1)       # C = A'A/(n-1)
    # Get eigenvalues D and eigenvectors V
    D, V = np.linalg.eigh(C)
    # Assert sign convention for eigenvectors
    V *= np.sign(V.min(0) + V.max(0)) # change sign where |min|>|max|
    # Sort eigenvalues and eigenvectors
    sort_index = D.argsort()[::-1]     # descending
    D = D[sort_index]                  # sort eigenvalues
    V = V[:,sort_index]                # sort eigenvectors by columns
    # Get principal component coefficient matrix
    V = V if n_components is None else V[:,:n_components]
    # Transform data
    A = A @ V
    return V, A, D, col_mean

def pairwise3dscatter(A, B, title=None):
    '''
    Helper function for pairwise plots.
    '''
    fig = plt.figure(figsize=(4,4))
    fig.suptitle(title)
    ax = [
        fig.add_subplot(2, 2, 2, projection='3d'),
        fig.add_subplot(2, 2, 1),
        fig.add_subplot(2, 2, 3),
        fig.add_subplot(2, 2, 4)
    ]
    a1, a2, a3 = A.T
    b1, b2, b3 = B.T
    ax[0].scatter(a1, a2, a3, alpha=.5)
    ax[0].scatter(b1, b2, b3, alpha=.5)
    ax[1].scatter(a1, a2, alpha=.5)
    ax[1].scatter(b1, b2, alpha=.5)
    ax[2].scatter(a1, a3, alpha=.5)
    ax[2].scatter(b1, b3, alpha=.5)
    ax[3].scatter(a2, a3, label="Original", alpha=.5)
    ax[3].scatter(b2, b3, label="Reconstructed", alpha=.5)
    ax[1].set_ylabel("$x_2$")
    ax[2].set_ylabel("$x_3$")
    ax[2].set_xlabel("$x_1$")
    ax[3].set_xlabel("$x_2$")
    ax[3].legend()
    plt.show()
```

Listing 3: Kernel PCA example

```
import numpy as np
from kpca import *

# Make test/train data sets
train = make_circles((1, 2, 3))
test = make_circles((1.5, 2.5), density=12, label=4)
circles = np.row_stack([train, test])
plot_circles(circles)

# Make kernel
sigma = 1 / np.sqrt(8)
k = gaussian_kernel(sigma)

# Do KPCA on training data
X = train[:,:-1]
X_label = train[:,-1]
coeff, Xhat, latent = kpca_fit(k, X, 3)

# Transform test data
Y = test[:,:-1]
Y_label = test[:,-1]
Yhat = kpca_transform(k, coeff, X, Y)

# Plot training and test data
points = np.row_stack([Xhat, Yhat])
labels = np.append(X_label, Y_label)
kpca_plot3(points, labels)
```

Listing 4: Kernel PCA source functions

```
import numpy as np
import matplotlib.pyplot as plt

def gaussian_kernel(sigma):
    sqnorm = lambda x: np.linalg.norm(x)**2
    gamma = 1 / (2 * sigma**2)
    return lambda x,y: np.exp(-sqnorm(x-y) * gamma)

def pairwise_kernels(kernel, X, Y=None):
    X = np.copy(X) # training data
    Y = X if Y is None else np.copy(Y) # test data
    # Compute pairwise kernel matrix
    K = np.array([[kernel(x,y) for x in X] for y in Y])
    # Kernel matrix centering
    cmean = K.mean(0)[:,np.newaxis].T # column mean
    rmean = K.mean(1)[:,np.newaxis]   # row mean
    Kmean = K.mean()                  # entry mean
    return K - cmean - rmean + Kmean

```

```
20 def kpca_fit(kernel, X, n_components=None):
21     X = np.copy(X)
22     # Make kernel matrix
23     K = pairwise_kernels(kernel, X)
24     # Get PC eigenvectors and eigenvalues
25     coeff, latent, _ = np.linalg.svd(K, hermitian=True)
26     # Dimension reduction
27     if n_components is not None:
28         latent = latent[:n_components]
29         coeff = coeff[:,:n_components]
30     # Coefficient scaling
31     coeff /= np.sqrt(latent)
32     # Transform data
33     score = latent * coeff
34     return coeff, score, latent
35
36 def kpca_transform(kernel, coeff, X, Y):
37     X = np.copy(X) # training data
38     Y = np.copy(Y) # test data
39     K = pairwise_kernels(kernel, X, Y)
40     return K.dot(coeff)
41
42 def kpca_plot3(xyz, l=None):
43     x, y, z = xyz.T
44     if l is None:
45         l = np.zeros_like(x)
46     fig = plt.figure()
47     ax1 = fig.add_subplot(221, aspect='equal')
48     ax2 = fig.add_subplot(222, projection='3d')
49     ax3 = fig.add_subplot(223, aspect='equal')
50     ax4 = fig.add_subplot(224, aspect='equal')
51     for lab in np.unique(l):
52         i = lab == l
53         ax1.scatter(x[i], y[i], alpha=.8)
54         ax2.scatter(x[i], y[i], z[i], alpha=.8)
55         ax3.scatter(x[i], z[i], alpha=.8)
56         ax4.scatter(y[i], z[i], alpha=.8)
57     ax1.set_ylabel("$x_2$")
58     ax3.set_ylabel("$x_3$")
59     ax3.set_xlabel("$x_1$")
60     ax4.set_xlabel("$x_2$")
61     plt.show()
62
63 def make_circle(npts=30, rad=1, label=0):
64     th = np.linspace(start=0,
65                      stop=2 * np.pi,
66                      num=npts,
67                      endpoint=False)
68     x = rad * np.cos(th)
69     y = rad * np.sin(th)
```

```
    l = np.repeat(label,npts)
    return np.column_stack([x, y, l])

def make_circles(rads, density=15, label=None):
    n = lambda r: int(density * r)
    if label is None:
        ret = [make_circle(n(r), r, l) for l, r in enumerate(rads)]
    else:
        ret = [make_circle(n(r), r, label) for r in rads]
    return np.row_stack(ret)

def plot_circles(circ):
    labs = np.unique(circ[:,2])
    ax = plt.axes(aspect='equal')
    for lab in labs:
        i = circ[:,2] == lab
        ax.scatter(circ[:,0][i], circ[:,1][i], alpha=.8)
    plt.show()
```

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