Nonlinear Regression Estimation Using Subset-Based Kernel Principal Components

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NONLINEAR REGRESSION ESTIMATION USING SUBSET-BASED KERNEL PRINCIPAL COMPONENTS

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Abstract: We study the estimation of conditional mean regression functions through the so-called subset-based kernel principal component analysis (KPCA). Instead of using one global kernel feature space, we project a target function into different localized kernel feature spaces at different parts of the sample space. Each localized kernel feature space reflects the relationship on a subset between the response and covariates more parsimoniously. When the observations are collected from a strictly stationary and weakly dependent process, the orthonormal eigenfunctions which span the kernel feature space are consistently estimated by implementing an eigenanalysis on the subset-based kernel Gram matrix, and the estimated eigenfunctions are then used to construct the estimation of the mean regression function. Under some regularity conditions, the developed estimator is shown to be uniformly consistent over the subset with a convergence rate faster than those of some well-known nonparametric estimation methods. In addition, we also discuss some generalizations of the KPCA approach, and consider using the same subset-based KPCA approach to estimate the conditional distribution.
function. The numerical studies including three simulated examples and two real data sets illustrate the reliable performance of the proposed method. In particular, the improvement over the global KPCA method is evident.

Key words and phrases: Conditional distribution function, eigenanalysis, kernel Gram matrix, KPCA, mean regression function, nonparametric regression.

1 Introduction

Let $Y$ be a scalar response variable and $X$ be a $p$-dimensional random vector. We are interested in estimating the conditional mean regression function defined by

$$ h(x) = E(Y|X = x), \quad x \in G, \quad (1.1) $$

where $G \subset \mathbb{R}^p$ is a measurable subset of the sample space of $X$, and $P(X \in G) > 0$. We allow that the mean regression function $h(\cdot)$ is not specified except certain smoothness conditions, which makes (1.1) more flexible than the traditional parametric linear and nonlinear regression. Nonparametric estimation of $h(\cdot)$ has been extensively studied in the existing literature such as Green and Silverman (1994), Wand and Jones (1995), Fan and Gijbels (1996), Fan and Yao (2003) and Teräsvirta, Tjøstheim and Granger (2010). When the dimension of random covariates $p$ is large, a direct use of the nonparametric regression estimation methods such as the spline and kernel-based smoothing typically perform poorly due to the so-called “curse of dimensionality”. Hence, some dimension-reduction techniques/assumptions (such as the additive models, single-index models and varying-coefficient models) have to be imposed when estimating the mean regression function. However, it is well known that some dimension reduction
techniques may result in systematic biases in estimation. For instance, the estimation based on an additive model may perform poorly when the data generation process deviates from the additive assumption.

In this paper we propose a data-driven dimension reduction approach through using a Kernel Principal Components Analysis (KPCA) for the random covariate $X$. The KPCA is a nonlinear version of the standard linear Principal Component Analysis (PCA) and overcomes the limitations of the linear PCA by conducting the eigendecomposition of the kernel Gram matrix, see, for example, Schölkopf, Smola and Müller (1999), Braun (2005) and Blanchard, Bousquet and Zwald (2007). See also Section 2.2 below for a detailed description on the KPCA and its relation to the standard PCA. The KPCA has been applied in, among others, feature extraction and de-noising in high-dimensional regression (Rosipal et al., 2001), density estimation (Girolami, 2002), robust regression (Wibowo and Desa, 2011), conditional density estimation (Fu, Shih and Wang, 2011; Izbicki and Lee, 2013), and regression estimation (Lee and Izbicki, 2013).

Unlike the existing literature on KPCA, we approximate the mean regression $h(x)$ on different subsets of the sample space of $X$ by linear combinations of different subset-based kernel principal components. The subset-based KPCA identifies nonlinear eigenfunctions in a subset, and thus reflects the relationship between $Y$ and $X$ on that set more parsimoniously than, for example, a global KPCA (see Proposition 1 in Section 2.2 below). The subsets may be defined according to some characteristics of $X$ and/or those on the relationship between $Y$ and $X$ (e.g., MACD for financial prices, different seasons/weekdays for electricity consumption, or adaptively by some change-point detection methods) and they are not necessarily connected sets. This is a marked difference from some conventional nonparametric regression techniques such as the kernel smoothing and nearest neighbour methods. Meanwhile, we assume that the
observations in the present paper are collected from a strictly stationary and weakly dependent process, which relaxes the independence and identical distribution assumption in the KPCA literature and makes the proposed methodology applicable to the time series data. Under some regularity conditions, we show that the estimated eigenvalues and eigenfunctions which are constructed through an eigenanalysis on the subset-based kernel Gram matrix are consistent. The conditional mean regression function $h(\cdot)$ is then estimated through the projection to the kernel spectral space which is spanned by a few estimated eigenfunctions whose number is determined by a simple ratio method. The developed conditional mean estimation is shown to be uniformly consistent over the subset with a convergence rate faster than those of some well-known nonparametric estimation methods. We further extend the subset-based KPCA method to estimation of the conditional distribution function:

$$F_{Y|X}(y|x) = P(Y \leq y | X = x), \ x \in G,$$

and establish the associated asymptotic property.

The rest of the paper is organized as follows. Section 2 introduces the subset-based KPCA and the estimation methodology for the mean regression function. Section 3 derives the main asymptotic theorems of the proposed estimation method. Section 4 extends the proposed subset-based KPCA for estimation of conditional distribution functions. Section 5 illustrates the finite sample performance of the proposed methods by simulation. Section 6 reports two real data applications. Section 7 concludes the paper. All the proofs of the theoretical results are available in an online supplementary material.
2 Methodology

Let \( \{(Y_i, X_i), 1 \leq i \leq n\} \) be observations from a strictly stationary process with the same marginal distribution as that of \( \{(Y, X)\} \). Our aim is to estimate the mean regression function \( h(x) \) for \( x \in G \), as specified in (1.1). We first introduce the kernel spectral decomposition in Section 2.1, followed by the illustration on the kernel feature space and the relationship between the KPCA and the standard PCA in Section 2.2, and finally propose an estimation method for the conditional mean regression function in Section 2.3.

2.1 Kernel spectral decomposition

Let \( \mathcal{L}_2(G) \) be the Hilbert space consisting of all the functions defined on \( G \) which satisfy the following conditions: for any \( f \in \mathcal{L}_2(G) \),

\[
\int_G f(x)P_X(dx) = E[f(X)I(X \in G)] = 0,
\]

and

\[
\int_G f^2(x)P_X(dx) = E[f^2(X)I(X \in G)] < \infty,
\]

where \( P_X(\cdot) \) denotes the probability measure of \( X \), and \( I(\cdot) \) is an indicator function. The inner product on \( \mathcal{L}_2(G) \) is defined as

\[
\langle f, g \rangle = \int_G f(x)g(x)P_X(dx) = \text{Cov} \{ f(X)I(X \in G), g(X)I(X \in G) \}, \quad f, g \in \mathcal{L}_2(G). \tag{2.1}
\]

Let \( K(\cdot, \cdot) \) be a Mercer kernel defined on \( G \times G \), i.e., \( K(\cdot, \cdot) \) is a bounded and symmetric function, and for any \( u_1, \cdots, u_k \in G \) and \( k \geq 1 \), the \( k \times k \) matrix with \( K(u_i, u_j) \) being its
(i, j)-th element is non-negative definite. For any fixed \( u \in \mathcal{G}, K(x, u) \) can be seen as a function of \( x \). A Mercer kernel \( K(\cdot, \cdot) \) defines an operator on \( L_2(\mathcal{G}) \) as follows:

\[
K(x, u) \rightarrow \int_{\mathcal{G}} K(x, u) f(u) P_X(du).
\]

It follows from Mercer’s Theorem (Mercer, 1909) that a Mercer kernel admits the following spectral decomposition:

\[
K(u, v) = \sum_{k=1}^{d} \lambda_k \varphi_k(u) \varphi_k(v), \quad u, v \in \mathcal{G}, \tag{2.2}
\]

where \( \lambda_1 \geq \lambda_2 \geq \cdots \geq \lambda_d > 0 \) are the positive eigenvalues of \( K(\cdot, \cdot) \), and \( \varphi_1, \varphi_2, \cdots \) are the orthonormal eigenfunctions in the sense that

\[
\int_{\mathcal{G}} K(x, u) \varphi_k(u) P_X(du) = \lambda_k \varphi_k(x), \quad x \in \mathcal{G}, \tag{2.3}
\]

and

\[
\langle \varphi_i, \varphi_j \rangle = \int_{\mathcal{G}} \varphi_i(u) \varphi_j(u) P_X(du) = \begin{cases} 
1 & i = j, \\
0 & i \neq j.
\end{cases} \tag{2.4}
\]

As we can see from the spectral decomposition (2.2), \( d = \max\{k: \lambda_k > 0\} \) and is possible to be infinity. We say that the Mercer kernel is of finite-dimension when \( d \) is finite, and of infinite-dimension when \( d = \infty \). To simplify the discussion, in this section and Section 3 below, we assume \( d \) is finite. This restriction will be relaxed in Section 4. We refer to Ferreira and Menegatto (2009) for Mercer’s Theorem for metric spaces. The eigenvalues \( \lambda_k \) and the associated eigenfunctions \( \varphi_k \) are usually unknown, and they need to be estimated in practice. To this end,
we construct the sample eigenvalues and eigenvectors through an eigenanalysis of the kernel Gram matrix which is defined in (2.6) below, and then obtain the estimate of the eigenfunction \( \varphi_k \) by the Nyström extension (Drineas and Mahoney, 2005).

Define
\[
\{(Y^j_i, X^j_i), j = 1, \ldots, m \} = \{(Y_i, X_i) \mid 1 \leq i \leq n, X_i \in G\},
\]
where \( m \) is the number of observations satisfying \( X_i \in G \), and define the subset-based kernel Gram matrix:
\[
K_G = \begin{pmatrix}
K(X_1^G, X_1^G) & K(X_1^G, X_2^G) & \cdots & K(X_1^G, X_m^G) \\
K(X_2^G, X_1^G) & K(X_2^G, X_2^G) & \cdots & K(X_2^G, X_m^G) \\
\vdots & \vdots & \ddots & \vdots \\
K(X_m^G, X_1^G) & K(X_m^G, X_2^G) & \cdots & K(X_m^G, X_m^G)
\end{pmatrix},
\]
(2.6)

Let \( \hat{\lambda}_1 \geq \cdots \geq \hat{\lambda}_m \geq 0 \) be the eigenvalues of \( K_G \), and \( \hat{\varphi}_1, \cdots, \hat{\varphi}_m \) be the corresponding \( m \) orthonormal eigenvectors. Write
\[
\hat{\varphi}_k = [\hat{\varphi}_k(X_1^G), \cdots, \hat{\varphi}_k(X_m^G)]^T.
\]
(2.7)

We next use the so-called Nyström extension to obtain the estimate of the eigenfunction. The Nyström method is originally introduced to get the approximate numerical solution of an integral equation by replacing the integral with a representative weighted sum. The integral in (2.3) can be approximated by \( \frac{1}{m} \sum_{i=1}^{m} K(x, X_i^G) \varphi_k(X_i^G) \). Under some mild conditions (e.g., Assumption 3 in Section 3), and using the Law of Large Numbers, such an approximation is sensible. Hence, the eigenfunction \( \varphi_k(x) \) can be approximated by \( \frac{1}{m \lambda_k} \sum_{i=1}^{m} K(x, X_i^G) \varphi_k(X_i^G) \).
Replacing \( \lambda_k \) and \( \varphi_k(X^\varphi) \) by \( \hat{\lambda}_k/m \) and \( \sqrt{m}\hat{\varphi}_k(X^\varphi) \), respectively, we may define the Nyström extension of the eigenvector \( \hat{\varphi}_k \) as

\[
\hat{\varphi}_k(x) = \frac{\sqrt{m}}{\hat{\lambda}_k} \cdot \sum_{i=1}^{m} K(x, X^\varphi_i) \hat{\varphi}_k(X^\varphi_i), \quad x \in \mathcal{G}, \quad k = 1, \ldots, d. \tag{2.8}
\]

Let

\[
\tilde{\lambda}_k = \frac{\hat{\lambda}_k}{m}, \quad k = 1, \ldots, d. \tag{2.9}
\]

Proposition 3 in Section 3 below shows that, for any \( x \in \mathcal{G}, \) \( \tilde{\lambda}_k \) and \( \hat{\varphi}_k(x) \) are consistent estimators of \( \lambda_k \) and \( \varphi_k(x) \), respectively.

Another critical issue in practical application is to estimate the dimension of the Mercer kernel \( K(\cdot, \cdot) \). When the dimension of \( K(\cdot, \cdot) \) is \( d \) and \( d \ll m \), we may estimate \( d \) by the following ratio method (Lam and Yao, 2012):

\[
d = \arg \min_{1 \leq k \leq \lceil mc_0 \rceil} \frac{\hat{\lambda}_{k+1}}{\hat{\lambda}_k} = \arg \min_{1 \leq k \leq \lceil mc_0 \rceil} \frac{\tilde{\lambda}_{k+1}}{\tilde{\lambda}_k}, \tag{2.10}
\]

where \( c_0 \in (0, 1) \) is a pre-specified constant such as \( c_0 = 0.5 \) and \( \lceil z \rceil \) denotes the integer part of the number \( z \). The numerical results in Sections 5 and 6 show that this ratio method works well in finite sample cases.

### 2.2 Kernel feature space and KPCA

Let \( \mathcal{M}(K) \) be a \( d \)-dimensional linear space spanned by the eigenfunctions \( \varphi_1, \ldots, \varphi_d \), and

\[
\dim \{ \mathcal{M}(K) \} = d = \max \{ k : \lambda_k > 0 \}.
\]
By the spectral decomposition (2.2), $M(K)$ can also be viewed as a linear space spanned by functions $g_u(.) \equiv K(.,u)$ for all $u \in G$. Thus we call $M(K)$ the kernel feature space as it consists of the feature functions extracted by the kernel function $K(.,.)$, and call $\varphi_1, \cdots, \varphi_d$ the characteristic features determined by $K(.,.)$ and the distribution of $X$ on set $G$. In addition, we call $\varphi_1(X), \varphi_2(X), \cdots$ the kernel principal components of $X$ on set $G$, and one can see they are nonlinear functions of $X$ in general. We next give an interpretation to see how the KPCA is connected to the standard PCA.

Any $f \in M(K)$ whose mean is zero on set $G$ admits the following expression:

$$f(x) = \sum_{j=1}^{d} \langle f, \varphi_j \rangle \varphi_j(x) \quad \text{for} \ x \in G.$$ 

Furthermore,

$$||f||^2 = \langle f, f \rangle = \text{Var}\{f(X)I(X \in G)\} = \sum_{j=1}^{d} \langle f, \varphi_j \rangle^2.$$ 

Now we introduce a generalized variance incited by the kernel function $K(.,.)$:

$$\text{Var}_K\{f(X)I(X \in G)\} = \sum_{j=1}^{d} \lambda_j \langle f, \varphi_j \rangle^2,$$  \hspace{0.5cm} (2.11)

where $\lambda_j$ is assigned as the weight on the “direction” of $\varphi_j$ for $j = 1, \cdots, d$. Then it follows from (2.2) and (2.3) that

$$\varphi_1 = \arg \max_{f \in M(K), ||f||=1} \int_{G \times G} f(u)f(v)K(u,v)P_X(du)P_X(dv)$$

$$= \arg \max_{f \in M(K), ||f||=1} \sum_{j=1}^{d} \lambda_j \langle f, \varphi_j \rangle^2$$

$$= \arg \max_{f \in M(K), ||f||=1} \text{Var}_K\{f(X)I(X \in G)\},$$
which indicates that the function \( \varphi_1 \) is the “direction” which maximizes the generalized variance \( \text{Var}_K \{ f(X) | (X \in G) \} \). Similarly it can be shown that \( \varphi_k \) is the solution of the above maximization problem with additional constraints \( \langle \varphi_k, \varphi_j \rangle = 0 \) for \( 1 \leq j < k \). Hence, the kernel principal components are the orthonormal functions in the feature space \( M(K) \) with the maximal kernel induced variances defined in (2.11). In other words, the kernel principal components \( \varphi_1, \varphi_2, \cdots \) can be treated as “directions” while their corresponding eigenvalues \( \lambda_1, \lambda_2, \cdots \) can be considered as the importance of these “directions”.

A related but different approach is to view \( M(K) \) as a reproducing kernel Hilbert space, for which the inner product is defined different from (2.1) to serve as a penalty in estimating functions via regularization; see section 5.8 of Hastie, Tibshirani and Friedman (2009) and Wahba (1990). Since the reproducing property is irrelevant in our context, we adopt the more natural inner product (2.1). For the detailed interpretation of KPCA in a reproducing kernel space, we refer to section 14.5.4 of Hastie, Tibshirani and Friedman (2009).

We end this subsection by stating a proposition which shows that the smaller \( G \) is, the lower the dimension of \( M(K) \) is. This indicates that a more parsimonious representation can be obtained by using the subset-based KPCA instead of the global KPCA. The proof of the proposition follows immediately from (2.2) and Proposition 2 in Section 2.3 below.

**Proposition 1.** Let \( \bar{G} \) be a measurable subset of the sample space of \( X \) such that \( G \subset \bar{G} \), and \( K(\cdot, \cdot) \) be a Mercer kernel on \( \bar{G} \times \bar{G} \). The kernel feature spaces defined with sets \( G \) and \( \bar{G} \) are denoted, respectively by \( M(K) \) and \( \bar{M}(K) \). Then \( \text{dim}\{M(K)\} \leq \text{dim}\{\bar{M}(K)\} \).
2.3 Estimation for conditional mean regression

For the simplicity of presentation, we assume that the mean of random variate \( h(X) = E(Y|X) \) on set \( G \) is 0, i.e.

\[
E[h(X)I(X \in G)] = E[E(Y|X)I(X \in G)] = E[YI(X \in G)] = 0.
\]

This amounts to replacing \( Y_i^G \) by \( Y_i^G - \bar{Y}^G \) in (2.5) with \( \bar{Y}^G = m^{-1} \sum_{i \in G} Y_i^G \). In general \( M(K) \) is a genuine subspace of \( L_2(G) \). Suppose that on set \( G \), \( h(x) = E(Y|X = x) \in M(K) \), i.e., \( h(x) \) may be expressed as

\[
h(x) = \int y f_{Y|X}(y|x)dy = \sum_{k=1}^{d} \beta_k \varphi_k(x), \quad x \in G,
\]

where \( f_{Y|X}(y|x) \) denotes the conditional density function of \( Y \) given \( X = x \), and

\[
\beta_k = \langle \varphi_k, h \rangle = \int_{x \in G} \varphi_k(x)P_X(dx) \int y f_{Y|X}(y|x)dy = E[Y \varphi_k(X) I(X \in \mathcal{G})].
\]

This leads to the estimator for \( \beta_k \) which is constructed as

\[
\tilde{\beta}_k = \frac{1}{m} \sum_{i=1}^{m} Y_i^G \varphi_k(X_i^G), \quad k = 1, \ldots, d,
\]
where \((Y_i^G, X_i^G), i = 1, \cdots, m\), are defined in (2.5), and \(\tilde{\varphi}_k(\cdot)\) are given in (2.8). Consequently the estimator for \(h(\cdot)\) is defined as

\[
\hat{h}(x) = \sum_{k=1}^{d} \beta_k \tilde{\varphi}_k(x), \quad x \in G. \tag{2.14}
\]

When the dimension of the kernel \(K(\cdot, \cdot)\) is unknown, the sum on the right hand side of the above expression runs from \(j = 1\) to \(d\) with \(d\) determined via (2.10).

The estimator in (2.14) is derived under the assumption that on set \(G\), \(h(x) \in M(K)\). When this condition is unfulfilled, (2.14) is an estimator for the projection of \(h(\cdot)\) on \(M(K)\). Hence the goodness of \(\hat{h}(\cdot)\) as an estimator for \(h(\cdot)\) depends critically on (i) kernel function \(K(\cdot, \cdot)\), (ii) set \(G\) and \(P_X(\cdot)\) on \(G\). In the simulation studies in Section 5 below, we will illustrate an approach to specify \(G\). Ideally we would like to choose a \(K(\cdot, \cdot)\) that induces a large enough \(M(K)\) such that \(h \in M(K)\). Some frequently used kernel functions include

- Gaussian kernel: \(K(u, v) = \exp(-\|u - v\|^2/c)\),

- Thin-plate spline kernel: \(K(u, v) = \|u - v\|^2 \log(\|u - v\|)\),

- Polynomial kernel (Fu, Shih and Wang, 2011):

\[
K(u, v) = \begin{cases} 
[1 - (u'v)^{\ell+1}]/(1 - u'v), & \text{if } u'v \neq 1, \\
\ell + 1, & \text{otherwise,}
\end{cases}
\]

where \(\|\cdot\|\) denotes the Euclidean norm, \(c\) is a positive constant, and \(\ell \geq 1\) is an integer. Also note that for any functions in \(\psi_1, \cdots, \psi_d \in L_2(G)\),

\[
K(u, v) = \sum_{k=1}^{d} \psi_k(u)\psi_k(v) \tag{2.15}
\]
is a well-defined Mercer kernel. A possible choice of the kernel function is to let \( \{ \psi_1(u), \ldots, \psi_d(u) \} \) be a set of basis functions of \( u \), e.g., Fourier series, polynomial series, wavelets, B-spline, etc.

The numerical studies in Sections 5 and 6 use (2.15) with appropriately chosen functions \( \psi_k \) in the estimation and dimension reduction procedure, which performs reasonably well. The following proposition shows that the dimension of \( M(K) \) with \( K(\cdot, \cdot) \) defined above is controlled by \( d \).

**PROPOSITION 2.** For the kernel function \( K(\cdot, \cdot) \) defined in (2.15), \( \dim(M(K)) \leq d \).

### 3 Large sample theory

In this section, we study the asymptotic properties for the estimators of the eigenvalues and eigenfunctions of the Mercer kernel as well as the mean regression estimation. We start with some regularity conditions which are sufficient to derive our asymptotic theory.

**ASSUMPTION 1.** The process \( \{ (Y_i, X_i) \} \) is strictly stationary and \( \alpha \)-mixing (or strongly mixing) dependent with the mixing coefficient satisfying

\[
\alpha_t = O(t^{-\kappa}), \quad \kappa > 2\delta_\star + p + \frac{3}{2},
\]

where \( p \) is the dimension of the random covariate, \( 0 \leq \delta_\star < \infty \) such that the volume of the set \( \mathcal{G} \) has the order \( m^{\delta_\star} \).

**ASSUMPTION 2.** The positive eigenvalues of the Mercer kernel \( K(\cdot, \cdot) \) are distinct and satisfy

\[
0 < \lambda_d < \cdots < \lambda_2 < \lambda_1 < \infty.
\]
Assumption 3. The eigenfunctions $\varphi_j$, $j = 1, \cdots, d$, are Lipschitz continuous and bounded on the set $\mathcal{G}$. Furthermore, the kernel $K(\cdot, \mathbf{x})$ is Lipschitz continuous and bounded on the set $\mathcal{G}$ for any $\mathbf{x} \in \mathcal{G}$.

Remark 1. In Assumption 1, we allow the process to be stationary and $\alpha$-mixing dependent, which is mild and can be satisfied by some commonly-used time series models; see e.g., Section 2.6 of Fan and Yao (2003) and the references within. For example the causal ARMA processes with continuous innovations are $\alpha$-mixing with exponentially decaying mixing coefficients. Note that for the processes with exponentially decaying mixing coefficients, (3.1) is fulfilled automatically, and the technical arguments in the proofs can be simplified. We allow set $\mathcal{G}$ to expand with the size of the sub-sample in $\mathcal{G}$ in the order of $m^{\delta_0}$, and $\delta_0$ would be 0 if $\mathcal{G}$ is bounded. Assumptions 2 and 3 impose mild restrictions on the eigenvalues and eigenfunctions of the Mercer kernel, respectively. They are crucial to ensure the consistency of the sample eigenvalues and eigenvectors constructed in Section 2.1. The boundedness condition on $\varphi_j$ and $K(\cdot, \mathbf{x})$ in Assumption 3 can be replaced by the $2(2 + \delta)$-order moment conditions for some $\delta > 0$, and Proposition 3 below still holds at the cost of more lengthy arguments. Furthermore, by the smoothness condition on the kernel function and using (3.2) in Proposition 3 below, we may easily show that $\tilde{\varphi}_j(\cdot)$ defined in (2.8), $j = 1, \cdots, d$, are Lipschitz continuous and bounded with probability tending to one.

Proposition 3. Suppose that Assumptions 1–3 are satisfied. Then we have

$$\max_{1 \leq k \leq d} |\tilde{\lambda}_k - \lambda_k| = \max_{1 \leq k \leq d} \left| \frac{1}{m} \tilde{\lambda}_k - \lambda_k \right| = O_P \left( m^{-1/2} \right)$$

and

$$\max_{1 \leq k \leq d} \sup_{\mathbf{x} \in \mathcal{G}} |\tilde{\varphi}_k(\mathbf{x}) - \varphi_k(\mathbf{x})| = O_P \left( \xi_m \right),$$

where

14
where $\xi_m = m^{-1/2} \log^{1/2} m$.

**Remark 2.** Proposition 3 presents the convergence rates of the estimated eigenvalues and eigenfunctions of the Mercer kernel $K(\cdot, \cdot)$. The result is of independent interest. It complements some statistical properties of the KPCA in the literature such as Braun (2005) and Blanchard, Bousquet and Zwald (2007). Note that $P(X \in G)$ can be consistently estimated by $m/n$. If it is assumed that $P(X \in G) = c_0 > 0$, $m$ would be of the same order as the full sample size $n$ (with probability tending to one). As a consequence, the convergence rates in (3.2) and (3.3) would be equivalent to $O_P \left( n^{-1/2} \right)$ and $O_P \left( n^{-1/2} \log^{1/2} n \right)$, respectively, which are not uncommon in the context of functional principal component analysis (Bosq, 2000; Horváth and Kokoszka, 2012). Based on Proposition 3, we can easily derive the following uniform consistency result for $\tilde{h}(\cdot)$.

**Theorem 1.** Suppose that Assumptions 1–3 are satisfied, $E[|Y|^{2+\delta}] < \infty$ for some $\delta > 0$ and $h(\cdot) \in M(K)$. Then it holds that

$$\sup_{x \in G} \left| \tilde{h}(x) - h(x) \right| = O_P \left( \xi_m \right),$$

where $\xi_m$ is defined in Proposition 3.

**Remark 3.** As stated above, the uniform convergence rate in (3.4) is equivalent to $O_P \left( n^{-1/2} \log^{1/2} n \right)$, which is faster than the well-known uniform convergence rate $O_P \left( (nb)^{-1/2} \log^{1/2} n \right)$ in the kernel smoothing method (Fan and Yao, 2003), where $b$ is a bandwidth which converges to zero as $n$ tends to $\infty$. The intrinsic reason of the faster rate in (3.4) is that we assume the dimension of the subset-based kernel feature space is finite, and thus the number of the unknown elements in (2.12) is also finite. Section 4 below shows that the increasing dimension of the kernel feature space would slow down the convergence rates.
4 Extensions of the estimation methodology

In this section, we consider two extensions of the methodology proposed in Section 2: the estimation for the conditional distribution function, and the case when the dimension of a kernel feature space diverges together with the sample size.

4.1 Estimation for conditional distribution functions

Estimation of the conditional distribution function defined in (1.2) is a key aspect in various statistical topics (such as the quantile regression), as the conditional mean regression may be not informative enough in many situations. Nonparametric estimation of the conditional distribution has been extensively studied in the literature including Hall, Wolff and Yao (1999), Hansen (2004) and Hall and Yao (2005). In this section, we use the subset-based KPCA approach discussed above to estimate a conditional distribution function in low-dimensional kernel feature space when the random covariates are multi-dimensional.

Let $F^*_y(x) = F_{Y|X}(y|x) - c_*$, where $c_* = P(Y \leq y, X \in \mathcal{G})$. Then $E[F^*_y(y|X)] = 0$. In practice $c_*$ can be easily estimated by the relative frequency. Suppose that $F^*_y(y|X) \in \mathcal{M}(K)$, i.e.,

$$F^*_y(y|x) = F_{Y|X}(y|x) - c_* = \int_{x} f_{Y|X}(z|x)dz - c_* = \sum_{k=1}^{d} \beta^*_k \varphi_k(x), \quad x \in \mathcal{G}. \quad (4.1)$$

Note that the coefficients $\beta^*_k$ in the above decomposition depend on $y$. The orthonormality of $\varphi_i$ implies that

$$\beta^*_k = \langle F^*_y(y|X), \varphi_k \rangle = \int_{\mathcal{G}} \varphi_k(x)P_X(dx) \left[ \int_{-\infty}^{y} f_{Y|X}(z|x)dz - c_* \right]$$

$$= \int_{\mathcal{G}} I(z \leq y, x \in \mathcal{G}) \varphi_k(x) f_{Y|X}(z|x)dzP_X(dx) - c_* \int_{\mathcal{G}} \varphi_k(x)P_X(dx)$$

$$= E[I(Y \leq y, X \in \mathcal{G})\varphi_k(X)] - c_* E[I(X \in \mathcal{G})\varphi_k(X)].$$
This leads to the following estimator for $\beta_k^*$:

$$
\hat{\beta}^*_k = \frac{1}{m} \sum_{i=1}^{m} I(Y_i \leq y) \tilde{\varphi}_k(X_i^y) - \frac{\tilde{c}_*}{m} \sum_{i=1}^{m} \tilde{\varphi}_k(X_i^y),
$$

(4.2)

where $(Y_i^y, X_i^y)$ are defined in (2.5), $\tilde{\varphi}_k()$ are defined in (2.8), and

$$
\tilde{c}_* = \frac{1}{n} \sum_{i=1}^{n} I(Y_i \leq y, X_i \in \mathcal{G}),
$$

(4.3)

$n$ is the full sample size. Consequently, we obtain the estimator for the conditional distribution:

$$
\hat{F}_{Y|X}(y|x) = \sum_{k=1}^{d} \hat{\beta}^*_k \tilde{\varphi}_k(x) + \tilde{c}_*.
$$

(4.4)

The estimator $\hat{F}_{Y|X}(\cdot|x)$ is not necessarily a bona fide distribution function. Some further normalization may be required to make the estimator non-negative, non-decreasing and between 0 and 1 (Glad, Hjort and Ushakov, 2003).

By the classic result for the $\alpha$-mixing sequence, we may show that $\tilde{c}_*$ is a consistent estimator of $c_*$ with a root-$n$ convergence rate. Then, by Proposition 3 and following the proof of Theorem 1, we have the following convergence result for $\hat{F}_{Y|X}(y|x)$.

**Theorem 2.** Suppose that Assumptions 1–3 are satisfied and $F_* (\cdot | \cdot) \in \mathcal{M}(K)$. Then it holds that

$$
\sup_{x \in \mathcal{X}} \left| \hat{F}_{Y|X}(y|x) - F_{Y|X}(y|x) \right| = O_P (\xi_m)
$$

(4.5)

for any given $y$, where $\xi_m$ is defined in Proposition 3.
4.2 Kernel feature spaces with diverging dimensions

We next consider the case when the dimension of the kernel feature space \( d_m = \max \{ k : \lambda_k > 0 \} \) depends on \( m \), and may diverge to infinity as \( m \) tends to infinity. Let

\[
\rho_m = \min \{ \lambda_k - \lambda_{k+1}, \ k = 1, \cdots, d_m \}.
\]

In order to derive a more general asymptotic theory, we need to slightly modify Assumption 2.

**Assumption 2*. The positive eigenvalues of the Mercer kernel \( K(\cdot, \cdot) \) are distinct and satisfy \( 0 < \lambda_{d_m} < \cdots < \lambda_2 < \lambda_1 < \infty, \sum_{k=1}^{d_m} \lambda_k < \infty \).**

The following proposition shows that the diverging \( d_m \) would slow down the convergence rates in Proposition 3.

**Proposition 4.** Suppose that Assumptions 1, 2*, and 3 are satisfied, \( d_m = o \left( m \rho_m^2 \lambda_{d_m}^2 / \log m \right) \), and the \( \alpha \)-mixing coefficient decays to zero at an exponential rate. Then it holds that

\[
\max_{1 \leq k \leq d_m} \left| \hat{\lambda}_k - \lambda_k \right| = \max_{1 \leq k \leq d_m} \left| \frac{1}{m} \hat{\lambda}_k - \lambda_k \right| = O_p \left( \frac{d_m^{1/2} \xi_m}{\rho_m} \right) \tag{4.6}
\]

and

\[
\max_{1 \leq k \leq d_m} \sup_{x \in \mathcal{G}} |\hat{\varphi}_k(x) - \varphi_k(x)| = O_p \left( \frac{d_m^{1/2} \xi_m}{\rho_m \lambda_{d_m}} \right), \tag{4.7}
\]

**Remark 4.** When \( d_m \) is fixed, it may be reasonable to assume both \( \rho_m \) and \( \lambda_{d_m} \) are bounded away from zero, and consequently the convergence rates in Proposition 4 would be simplified.

When \( d_m \to \infty \) as \( m \to \infty \), we usually have \( \rho_m \to 0 \) and \( \lambda_{d_m} \to 0 \). This implies that the
convergence rates in (4.6) and (4.7) would be generally slower than those in (3.2) and (3.3). Let $c_i, i = 1, \cdots, 5,$ be five positive constants. For any two sequences $a_m$ and $b_m$, $a_m \asymp b_m$ means that $0 < c_4 \leq a_m/b_m \leq c_5 < \infty$ when $m$ is sufficiently large. If $d_m = c_1 \log m$, $\rho_m = c_2 \log^{-1} m$ and $\lambda d_m = c_3 \log^{-1} m$, we have

$$d_m^{1/2} \xi_m \asymp m^{-1/2} \log m, \quad d_m^{1/2} \xi_m / (\rho_m \lambda d_m) \asymp m^{-1/2} \log^3 m.$$  

Using Proposition 4 and following the proof of Theorem 1, we can easily obtain the uniform convergence rate for the conditional mean regression estimation when $d_m$ is diverging.

In practice, we may encounter the more challenging case when the dimension of the Mercer kernel is infinite (e.g., $\lambda_k \asymp k^{-\iota_1}$ with $\iota_1 > 0$ or $\lambda_k \asymp k^{\iota_2}$ with $0 < \iota_2 < 1$). In this case, the convergence result in Proposition 4 is not directly applicable as the rates in (4.6) and (4.7) become divergent when the dimension is infinite. However, the proposed subset-based KPCA approach can still be used to estimate the conditional mean regression function. Assuming that the mean regression function $h(x) \in \mathcal{M}(K)$ and noting that the dimension of $\mathcal{M}(K)$ is infinite, we have

$$h(x) = \sum_{k=1}^{d_m} \beta_k \varphi_k(x) + \sum_{k=d_m+1}^{\infty} \beta_k \varphi_k(x) \equiv h_1(x) + h_2(x),$$  

where $\beta_k$ and $\varphi_k(x)$ are defined as in Section 2, and $d_m$ is a divergent number satisfying the condition in Proposition 4. Let $\mathcal{M}_1(K)$ be a $d_m$-dimensional kernel feature space spanned by $\varphi_1, \cdots, \varphi_{d_m}$. From (4.8), the mean regression function $h(x)$ can be well approximated by its projection onto the $d_m$-dimensional kernel feature space $\mathcal{M}_1(K)$ as long as the approximation error $h_2(x)$ uniformly converges to zero at certain rate. The latter usually holds if we impose
some smoothness condition on $h(x)$ and let $d_m$ divergent at an appropriate rate, which is similar to the conditions on sieve approximation accuracy (Chen, 2007).

Let $b^*_m = \sup_{x \in \mathcal{G}} |h_2(x)|$. We may estimate $\beta_k$ and $\varphi_k$, $k = 1, \ldots, d_m$, in the same manner as in Sections 2.2 and 2.3. Denote the estimates by $\tilde{\beta}_k$ and $\tilde{\varphi}_k$, and let $\tilde{h}_m(x) = \sum_{k=1}^{d_m} \tilde{\beta}_k \tilde{\varphi}_k(x)$. By Proposition 4 and following the proof of Theorem 1, we can establish the following uniform convergence rate:

$$
\sup_{x \in \mathcal{G}} |\tilde{h}_m(x) - h_1(x)| = O_P(\nu_m^*),
$$

where $\nu_m^* = d_m^{3/2} \xi_m / (\rho_m \lambda_d)$. Furthermore, we can prove, via the decomposition in (4.8), that

$$
\sup_{x \in \mathcal{G}} |\tilde{h}_m(x) - h(x)| = \sup_{x \in \mathcal{G}} |\tilde{h}_m(x) - h_1(x)| + \sup_{x \in \mathcal{G}} |h_2(x)| = O_P(\nu_m^* + b^*_m).
$$

5 Simulation Studies

In this section, we use three simulated examples to illustrate the finite sample performance of the proposed subset-based KPCA method and compare it with the global KPCA and other existing nonparametric estimation methods, i.e., cubic spline, local linear regression and kernel ridge regression. We start with an example to assess the out-of-sample estimation performance of conditional mean function based on a multivariate nonlinear regression model. Then, in the second example, we examine the one-step ahead out-of-sample forecast performance based on a multivariate nonlinear time series model. Finally, in the third example, we examine the finite sample performance of the estimation of conditional distribution function.

Throughout this section, the kernel function is either the Gaussian kernel or formulated
as in (2.15) with \( \{\psi_1(u), \ldots, \psi_d(u)\} \) being a set of normalized polynomial basis functions (with the unit norm) of \( u = (u_1, \ldots, u_p)^T \) of order 2 and 3, i.e., \( \{1, u_k, \ldots, u_k^r, \ k = 1, \ldots, p\} \), where \( r = 2, 3 \) and \( d = rp + 1 \). For the latter case, we call the kernel as the \textit{quadratic kernel} when \( r = 2 \) and the \textit{cubic kernel} when \( r = 3 \). In practice, \( d \) is estimated by the ratio method as in (2.10). The simulation results show that (2.10) can correctly estimate \( \hat{d} = d \) with frequency close to 1. The subset is chosen to be the \( \lceil \kappa n \rceil \) nearest neighbors, where \( n \) is the sample size and \( \kappa \in (0, 1) \) is a constant bandwidth. The bandwidth \( \kappa \) and the tuning parameter \( c \) in the Gaussian kernel are selected by a 5-fold cross validation.

**Example 5.1.** Consider the following model:

\[
y_i = g(x_{2i}) + \sin(\pi(x_{3i} + x_{4i})) + x_{5i} + \log(1 + x_{6i}^2) + \epsilon_i,
\]

where \( x_{1i}, \ldots, x_{6i} \) and \( \epsilon_i \) are i.i.d. \( N(0, 1) \), \( g(x) = e^{-2x^2} \) for \( x \geq 0 \), and \( g(x) = e^{-x^2} \) for \( x < 0 \).

In the model, the covariate \( x_{1i} \) is irrelevant to \( y_i \).

We draw a training sample of size \( n = 500 \) or \( 1000 \) and a testing sample of size 200. We estimate the conditional mean regression function using the training sample, and then calculate the mean squared errors (MSE) and out-of-sample \( R^2 \)'s over the testing sample as follows:

\[
\text{MSE} = \frac{1}{200} \sum_{i=1}^{200} \left( y_i - \tilde{h}(x_i) \right)^2, \quad R^2 = 1 - \frac{\sum_{i=1}^{200} \left( y_i - \bar{y} \right)^2}{\sum_{i=1}^{200} \left( y_i - \bar{y} \right)^2},
\]

where \( \tilde{h}(\cdot) \) is defined as in (2.14), \( x_i = (x_{1i}, \ldots, x_{6i})^T \) and \( \bar{y} \) is the sample mean of \( y_i \) over the training sample.

By repeating this procedure over 200 replications, we obtain a sample of MSE and \( R^2 \) with size 200. The estimation performance is assessed by the sample mean, median and variance.
of MSE and $R^2$. The simulation results are reported in Table 1. In this simulation, for the quadratic kernel and cubic kernel, the ratio method in (2.10) can always correctly estimate $\tilde{d} = rp + 1$. According to the results in Table 1, the subset-based KPCA with the quadratic kernel outperforms the global KPCA methods and other nonparametric methods as it has the smallest sample mean, median and variance of MSE and the highest $R^2$. In addition, both the quadratic kernel and cubic kernel perform better than the Gaussian kernel due to the fact that they can better capture different degree of smoothness on different directions.

To assess the the bandwidth choice for subset-based KPCA, we set $n = 500$, let $\kappa$ vary from 0.05 to 0.8 and calculate the sample mean of MSE over 100 replications. The results are plotted in Figure 1. According to Figure 1, the subset-based KPCA method is not sensitive to the choice of $\kappa$. The smallest MSE is achieved at $\kappa = 0.27$, and any $\kappa$ between 0.15 and 0.45 yields similar result.

Furthermore, we compare the computational costs between the subset-based KPCA and global KPCA approaches with quadratic kernel function. The computational cost of subset-based KPCA includes the selection of bandwidth $\kappa$. The bandwidth $\kappa$ is selected by 5-fold cross validation over 10 grid points equally spaced between 0.1 and 0.5. We let the sample size increase from 400 to 1000 with a step size of 20 and record the computational time over 100 replications for both approaches. The comparison results are presented in Figure 2. The major computational cost of the global KPCA methods is the eigen-decomposition of the $n \times n$ gram matrix which is of computational complexity $O(n^\omega)$ for some $\omega > 2$. To see this, we calculate the empirical order of growth for both approaches:

$$\hat{\omega} = \frac{1}{30} \sum_{l-1}^{30} \frac{\log(T_{l+1}/T_l)}{\log(n_{l+1}/n_l)}.$$
where $n_l$ is the $l$-th component in the set $\{400, 420, \ldots, 980, 1000\}$ and $T_l$ is the corresponding computational cost over 100 replications. The empirical order of growth for the global KPCA is 2.69, whereas that for the subset-based KPCA is 2.17. Hence, both Figure 2 and the calculation of empirical order of growth show the subset-based KPCA method scales with sample size much better than the global counterpart.

### Table 1: Out-of-sample estimation performance in Example 5.1

<table>
<thead>
<tr>
<th>Method</th>
<th>$n = 500$ MSE (the smaller the better)</th>
<th>$R^2$ (the larger the better)</th>
</tr>
</thead>
<tbody>
<tr>
<td></td>
<td>Mean</td>
<td>Median</td>
</tr>
<tr>
<td>sKPCA+Quadratic</td>
<td>1.300</td>
<td>1.294</td>
</tr>
<tr>
<td>sKPCA+Cubic</td>
<td>1.337</td>
<td>1.335</td>
</tr>
<tr>
<td>sKPCA+Gaussian</td>
<td>1.573</td>
<td>1.586</td>
</tr>
<tr>
<td>gKPCA+Quadratic</td>
<td>2.389</td>
<td>1.871</td>
</tr>
<tr>
<td>gKPCA+Cubic</td>
<td>2.586</td>
<td>1.937</td>
</tr>
<tr>
<td>gKPCA+Gaussian</td>
<td>3.023</td>
<td>2.021</td>
</tr>
<tr>
<td>Cubic Spline</td>
<td>1.386</td>
<td>1.383</td>
</tr>
<tr>
<td>Local Linear</td>
<td>1.429</td>
<td>1.431</td>
</tr>
<tr>
<td>Kernel Ridge</td>
<td>1.897</td>
<td>1.866</td>
</tr>
</tbody>
</table>

<table>
<thead>
<tr>
<th>Method</th>
<th>$n = 1000$ MSE (the smaller the better)</th>
<th>$R^2$ (the larger the better)</th>
</tr>
</thead>
<tbody>
<tr>
<td></td>
<td>Mean</td>
<td>Median</td>
</tr>
<tr>
<td>sKPCA+Quadratic</td>
<td>1.243</td>
<td>1.236</td>
</tr>
<tr>
<td>sKPCA+Cubic</td>
<td>1.278</td>
<td>1.271</td>
</tr>
<tr>
<td>sKPCA+Gaussian</td>
<td>1.531</td>
<td>1.528</td>
</tr>
<tr>
<td>gKPCA+Quadratic</td>
<td>2.380</td>
<td>2.371</td>
</tr>
<tr>
<td>gKPCA+Cubic</td>
<td>2.541</td>
<td>2.508</td>
</tr>
<tr>
<td>gKPCA+Gaussian</td>
<td>3.015</td>
<td>2.790</td>
</tr>
<tr>
<td>Cubic Spline</td>
<td>1.371</td>
<td>1.372</td>
</tr>
<tr>
<td>Local Linear</td>
<td>1.404</td>
<td>1.418</td>
</tr>
<tr>
<td>Kernel Ridge</td>
<td>1.858</td>
<td>1.831</td>
</tr>
</tbody>
</table>

“sKPCA” and “gKPCA” stand for the subset-based KPCA and global KPCA; “Quadratic”, “Cubic” and “Gaussian” stand for the quadratic kernel, cubic kernel and Gaussian kernel, respectively; “Cubic Spline”, “Local Linear” and “Kernel Ridge” stand for non-parametric estimation methods based on cubic spline, local linear regression and kernel ridge regression.

**Example 5.2.** Consider the following time series model:

$$y_t = \sin(0.02\pi y_{t-1}) + \exp(-y_{t-2}^2) + \ln(1 + |y_{t-3}|) - 0.3|y_{t-4}| + 0.2\epsilon_t.$$
Figure 1: The out-of-sample estimation performance of the subset-based KPCA approach with the quadratic kernel with respect to the bandwidth $\kappa$ when $n = 500$.

Figure 2: The computation costs for global KPCA method and subset-based KPCA method (with $\kappa$ selected by 5-fold cross validation) with respect to the sample size.

where $y_0 = 0$ and $\{\epsilon_t\}$ is a sequence of independent $\text{N}(0, 1)$ random variables. We next estimate the conditional mean $\mathbb{E}(y_t | y_{t-1}, y_{t-2}, y_{t-3}, y_{t-4})$ and denote the estimator as $\hat{y}_t$ which is to be used as the one-step-ahead predictor of $y_t$.

We generate a time series sample from the above model with the length $T + 100$. For
\( k = 1, \cdots, 100 \), we use the \( T \) observations right before time \( T + k \) as the training set to predict \( y_{T+k} \). The performance is measured by MSE and out-of-sample \( R^2 \):

\[
\text{MSPE} = \frac{1}{100} \sum_{k=1}^{100} (y_{T+k} - \hat{y}_{T+k})^2, \quad R^2 = 1 - \frac{\sum_{k=1}^{100} (y_{T+k} - \bar{y}_{T+k})^2}{\sum_{k=1}^{100} (y_{T+k} - \bar{y})^2},
\]

where \( \bar{y} \) is the sample mean of \( y_t \) over the training sample.

We set \( T = 500 \), and repeat the experiment 200 times for each method. The sample means, medians and variances of MSE and \( R^2 \) are presented in Table 2. Similar to Example 5.1, the subset-based KPCA method with the quadratic kernel provides the most accurate prediction. The subset-based KPCA method with the cubic kernel is a close second best in terms of both MSE and \( R^2 \). Figure 3 plots a typical path together with their one-step-ahead forecasts for each method. The typical path is the replication with median \( R^2 \). Figure 3 shows that the forecasted path from the subset-based KPCA method with the quadratic kernel follows the true path closely. A similar pattern can also be found for other subset-based KPCA method with the cubic and Gaussian kernel and the three nonparametric methods (cubic spline, local linear and kernel ridge). However the global KPCA methods fail to capture the variation of the series and tend to forecast the future values by the overall mean value, which is not satisfactory.

**Example 5.3.** Consider the model:

\[
X_1 \sim N(0, 1), \quad X_2 \sim N(0, 1),
\]

\[
Y \mid (X_1, X_2) \sim N(X_1, 1 + X_2^2).
\]

The conditional distribution of \( Y \) given \( X = (X_1, X_2)^T \) is a normal distribution with mean \( X_1 \) and variance \( 1 + X_2^2 \). According to the method proposed in Section 4.1, we estimate the
Figure 3: One-step ahead out-of-sample forecasting performance based on the replication with median $R^2$ for each method. The black solid line is the true value and the red dashed line is the predicted value.
Table 2: One-step ahead forecasting performance in Example 5.2

<table>
<thead>
<tr>
<th>Method</th>
<th>MSE (the smaller the better)</th>
<th>$R^2$ (the larger the better)</th>
</tr>
</thead>
<tbody>
<tr>
<td></td>
<td>Mean</td>
<td>Median</td>
</tr>
<tr>
<td>sKPCA+Quadratic</td>
<td>0.0435</td>
<td>0.0428</td>
</tr>
<tr>
<td>sKPCA+Cubic</td>
<td>0.0445</td>
<td>0.0437</td>
</tr>
<tr>
<td>sKPCA+Gaussian</td>
<td>0.0756</td>
<td>0.0751</td>
</tr>
<tr>
<td>gKPCA+Quadratic</td>
<td>0.1900</td>
<td>0.1908</td>
</tr>
<tr>
<td>gKPCA+Cubic</td>
<td>0.2042</td>
<td>0.2058</td>
</tr>
<tr>
<td>gKPCA+Gaussian</td>
<td>0.2172</td>
<td>0.2211</td>
</tr>
<tr>
<td>Cubic Spline</td>
<td>0.0516</td>
<td>0.0512</td>
</tr>
<tr>
<td>Local Linear</td>
<td>0.0522</td>
<td>0.0515</td>
</tr>
<tr>
<td>Kernel Ridge</td>
<td>0.0721</td>
<td>0.0717</td>
</tr>
</tbody>
</table>

conditional distribution function $F_{Y|X}(y|x)$ using the subset-based KPCA with the quadratic kernel.

We draw a training sample of size $n = 300$ or 500 and a testing sample of size 100. The estimated conditional distribution $\hat{F}_{Y|X}(y_i|x_i)$ is obtained using the training data. We repeat the simulation 200 times and measure the performance by MSE as well as largest absolute error (LAE) over the testing sample:

$$
\text{MSE} = \frac{1}{100} \sum_{i=1}^{100} \left( \hat{F}_{Y|X}(y_i|x_i) - F_{Y|X}(y_i|x_i) \right)^2,
$$

$$
\text{LAE} = \sup_{(y,x) \in \Omega^*} \left| \hat{F}_{Y|X}(y|x) - F_{Y|X}(y|x) \right|,
$$

where $\Omega^*$ is the union of all validation sets. The results are reported in Table 3. As the values of MSE and LAE in Table 3 are very small, the proposed method provides very accurate estimation for the conditional distribution function.
Table 3: Estimation of the conditional distribution function

<table>
<thead>
<tr>
<th>n</th>
<th>MSE Mean</th>
<th>LAE</th>
<th>MSE Median</th>
<th>MSE Variance</th>
</tr>
</thead>
<tbody>
<tr>
<td>300</td>
<td>$6.0 \times 10^{-4}$</td>
<td>0.098</td>
<td></td>
<td></td>
</tr>
<tr>
<td>500</td>
<td>$3.7 \times 10^{-4}$</td>
<td>0.080</td>
<td></td>
<td></td>
</tr>
</tbody>
</table>

6 Real data analysis

In this section, we apply the proposed subset-based KPCA method to two real data examples. Throughout this section, the kernel function is set to be either Gaussian or Quadratic kernel. The subset is chosen to be the $[\kappa n]$ nearest neighbors, where $n$ is the sample size and $\kappa \in (0, 1)$. The bandwidth $\kappa$ is selected by 5-fold cross validation.

6.1 Circulatory and respiratory problem in Hong Kong

We study the circulatory and respiratory problem in Hong Kong via an environmental data set. This data set contains 730 observations and was collected between January 1, 1994 and December 31, 1995. The response variable is the number of daily total hospital admissions for circulatory and respiratory problems in Hong Kong, and the covariates are daily measurements of seven pollutants and environmental factors: $SO_2$, $NO_2$, dust, temperature, change of temperature, humidity and ozone. We standardize the data so that all the covariates have zero sample mean and unit sample variance. To check the stationarity, we apply the augmented Dickey-Fuller test (e.g., Dickey and Fuller, 1981) to each variable in the data set. The tests are applied using the “urca” package in R and the lags included are selected by AIC. For each variable, the test result suggests us to reject the unit root null hypothesis. Therefore, we consider the variables in the dataset to be stationary.

The objective of this study is to estimate the number of daily total hospital admissions for
circulatory and respiratory problem using the collected environmental data, i.e., estimate the conditional mean regression function. The estimation performance is measured by the mean and variance of the out-of-sample $R^2$, which are calculated by a bootstrap method described as follows. We first randomly divide the data set into a training set of 700 observations and a testing set of 30 observations. For each observation in the testing set, we use the training set to estimate its conditional mean regression function. Then we calculate out-of-sample $R^2$ for the testing set as in Example 5.1. By repeating this re-sampling and estimation procedure 1000 times, we obtain a bootstrap sample of $R^2$s, and calculate its sample mean and variance.

Table 4: Estimation performance for the Hong Kong environmental data

<table>
<thead>
<tr>
<th>Method</th>
<th>$R^2$ (the larger the better)</th>
<th>Mean</th>
<th>Variance</th>
</tr>
</thead>
<tbody>
<tr>
<td>sKPCA + Quadratic</td>
<td>0.1544</td>
<td>0.0025</td>
<td></td>
</tr>
<tr>
<td>sKPCA + Gaussian</td>
<td>0.1262</td>
<td>0.0027</td>
<td></td>
</tr>
<tr>
<td>gKPCA + Quadratic</td>
<td>-0.3613</td>
<td>0.2232</td>
<td></td>
</tr>
<tr>
<td>gKPCA + Gaussian</td>
<td>-3.7058</td>
<td>1.6653</td>
<td></td>
</tr>
<tr>
<td>Cubic spline</td>
<td>0.0687</td>
<td>0.0042</td>
<td></td>
</tr>
</tbody>
</table>

We compare the performance among the following five methods: the subset-based KPCA with the quadratic kernel, the subset-based KPCA with the Gaussian kernel, the global KPCA with the quadratic kernel, the global KPCA with the Gaussian kernel and the cubic spline. The cubic spline is fitted with 10 knots using the “splines” package in R. The results are presented in Table 4. According to the results in Table 4, the subset-based KPCA with the quadratic kernel has the best estimation performance and the subset-based KPCA method outperforms the global KPCA method and cubic spline.
6.2 Forecasting the log return of CPI

The CPI is a statistical estimate that measures the average change in the price paid to a market basket of goods and services. The CPI is often used as an important economic indicator in macroeconomic and financial studies. For example, in economics, CPI is considered as closely related to the cost-of-living index and used to adjust the income eligibility levels for government assistance. In finance, CPI is considered as an indicator of inflation and used as the deflator to translate other financial series to inflation-free ones. Hence, it is always of interest to forecast the CPI. We next perform one-step-ahead forecasting for the monthly log return of CPI in USA based on the proposed subset-based KPCA method with the quadratic kernel. The data span from January 1970 to December 2014 with 540 observations. The augmented Dickey-Fuller test suggests the monthly log return of CPI over this time span is stationary.

Instead of using the traditional linear time series models, we consider that the log return of CPI follows a nonlinear AR(3) model:

\[ y_t = g(y_{t-1}, y_{t-2}, y_{t-3}) + \epsilon_t, \]

where \( g(\cdot) \) is an unknown function and \( \epsilon_t \) denotes an unobservable noise at time \( t \). The regression function \( g(\cdot) \) is estimated by the subset-based KPCA method with the quadratic kernel.

For the comparison purpose, we also forecast \( y_t \) based on a linear AR(\( p \)) model with the order \( p \) determined by AIC. Suppose the testing set starts from time \( T \) and ends at time \( T + S \), the forecast performance is measured by the out-of-sample \( R^2 \) as

\[ R^2 = 1 - \frac{\sum_{s=1}^{S} (y_{T+s} - \hat{y}_{T+s})^2}{\sum_{s=1}^{S} (y_{T+s} - \bar{y})^2}. \]
where $\hat{y}_{t+s}$ is the estimator of $y_{t+s}$, and $\bar{y}$ is the sample mean of $y_t$ over the training set.

We set the data from January 2005 to December 2014 as the testing set, which contains 120 observations. We forecast each observation in the testing set with the data up to its previous month. The out-of-sample $R^2$ is calculated over the testing set. The out-of-sample $R^2$ of the nonlinear AR(3) model is 0.2318 while the $R^2$ of the linear AR model is 0.0412. The detailed forecasting result is plotted in Figure 4, which shows clearly that the forecast based on the subset-based KPCA method is more accurate as it captures the variations much better than the linear AR modelling method.

![One step ahead out-of-sample forecast for the log return of CPI](image)

**Figure 4:** One step ahead out-of-sample forecast for the log return of CPI from January 2005 to December 2014. The black solid line is the true value, the red dashed line is the forecast value obtained by the subset-based KPCA, and the blue dotted line is the forecast value obtained by the linear AR model.
7 Conclusion

In this paper, we have developed a new subset-based KPCA method for estimating nonparametric regression functions. In contrast to the conventional (global) KPCA method which builds on a global kernel feature space, we use different lower-dimensional subset-based kernel feature spaces at different locations of the sample space. Consequently the resulting localized kernel principal components provide more parsimonious representation for the target regression function, which is also reflected by the faster uniform convergence rates presented in Theorem 1, see also the discussions immediately below Theorem 1. The reported numerical results with both simulated and real data sets illustrate clearly the advantages of using the subset-based KPCA method over its global counterpart. It also outperforms some popular nonparametric regression methods such as the cubic spline and kernel regression (the results on kernel regression are not reported to save the space). It is also worth mentioning that the quadratic kernel constructed based on (2.15) using normalized univariate linear and quadratic basis functions performs better than the more conventional Gaussian kernel for all the examples reported in Sections 5 and 6.

Supplementary materials

The online supplementary material contains the detailed proofs of Propositions 1–4 and Theorem 1.

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