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Efficient basis selection for smoothing splines via rotated lattices

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Fitting a smoothing spline model on a large-scale dataset is daunting due to the high computational cost. In this study, we develop an efficient basis selection method for smoothing spline calculation. The key idea is to force a nonparametric function in an infinite-dimensional functional space to reside in a relatively small and finitedimensional model space without the loss of too much prediction accuracy. Such an approximation naturally allows for much faster numerical calculation, especially for large datasets. Among various basis selection methods, space-filling basis selection has been proven to be more efficient since its model space dimension is smaller than that of others. Despite algorithmic benefits, most of the space-filling selection methods only take the overall space-filling property into account. These methods may be less efficient when the underlying response surface is not isomorphic. To overcome this obstacle, we develop an efficient algorithm to improve projective uniformity for space-filling basis selection. It has been proved that the proposed estimator has the same convergence rate as the full bases estimator. Compared with the standard approach, the proposed method significantly reduces the computational cost. Simulation and real data studies demonstrate the efficiency and superiority of the proposed method.

KEYWORDS

massive data, projective uniformity, rotated lattice, smoothing spline, space-filling basis selection

INTRODUCTION 1

Consider the following nonparametric model:

$$\mathbf{y}_i = \eta(\mathbf{x}_i) + \boldsymbol{\epsilon}_i, \, i = 1, 2, \dots, N, \tag{1}$$

where y_i is the response for the *i*th observation, $\eta(\cdot)$ is an unknown function, $x_i \in [0,1]^p$ denotes the covariate, ϵ_i 's are independent and identically distributed errors with zero mean and finite variance, and N is the size of training data. In this study, we adopt the smoothing spline to fit $\eta(\cdot)$, which is one of the most pervasive choices (Grace, 1990; Gu, 2013). To be precise, $\eta(\cdot)$ in Model (1) is assumed to have resided in a reproducing kernel Hilbert space (RKHS), say \mathcal{H} . It can be estimated by minimizing the following penalized least squares

$$N^{-1} \sum_{i=1}^{N} \{ \mathbf{y}_{i} - \eta(\mathbf{x}_{i}) \}^{2} + \lambda J(\eta),$$
(2)

where $J(\eta)$ is a roughness penalty and λ is the tuning parameter to control the smoothness of the estimator.

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Despite impressive performance and solid theoretical foundations, solving the optimization problem (2) is a challenging task. Except for the univariate case, which can be solved in O(N), the computational cost for calculating the smoothing spline with $p \ge 2$ using the standard approach is $O(N^3)$. This huge computational cost renders the smoothing spline impractical for a large-scale dataset. Great efforts have been made to alleviate the computational burden. Typical works include but are not limited to divide-and-conquer kernel ridge regressions (Xu & Wang, 2018; Zhang et al., 2015) and basis selection algorithms (Gu & Kim, 2002; Luo & Wahba, 1997; Ma et al., 2015; Meng et al., 2020).

Among various techniques designed to reduce the computational cost of the smoothing spline, basis selection methods have gained the attention of data scientists due to the following two reasons. First, it is possible to combine parallel-based strategies with basis selection methods to reap more computational savings. Second, basis selection procedures are more flexible than divide-and-conquer kernel ridge regression strategies. The idea of basis selection is to approximate the minimizer of (2) by restricting the estimator $\hat{\eta}$ to a *n*-dimensional subspace $\mathcal{H}_{S} \subset \mathcal{H}$. When $n \ll N$, the computational cost can be efficiently reduced from $O(N^3)$ to $O(Nn^2)$. Numerous studies have been developed along this line of thinking. For example, Hastie (1996) suggested fixing n basis functions to approximate splines. Such a method differs from constructing basis functions in smoothing splines. Thus, it is also known as "pseudo splines" or P-splines. Ruppert (2002) considered basis selection for P-splines by adding a penalty to control the number of basis functions. He et al. (2001), Sklar et al. (2013), and Yuan et al. (2013) further considered cases where the regression function has non-homogeneous smoothness across the input space. As for smoothing splines, Luo and Wahba (1997) and Zhang et al. (2004) proposed approximating the minimizer of (2) using variable selection techniques. Due to the additional computational cost brought by variable screening, the computational benefits of the above two methods are unclear. Gu and Kim (2002) proposed a uniform basis selection (UBS) method by randomly selecting some basis functions among the N basis functions in $\{G_J(\mathbf{x}_i, \cdot)\}_{i=1}^N$, facilitating quick analysis of large datasets. Here, G₁ is the reproducing kernel. Ma et al. (2015) developed an adaptive basis selection (ABS) method that uses an adaptive sampling scheme according to the values of the response variable. Meng et al. (2020) proposed a space-filling basis selection (SBS) method that suggests selecting the basis functions $\{G_J(\mathbf{x}_i^*, \cdot)\}_{i=1}^n$ with $\{x_i^*\}_{i=1}^n$ being roughly equally-spaced. Meng et al. (2021) extended the basis selection method to cases where the covariates are nonuniformly distributed in $[0,1]^p$. As the space-filling property of the selected sub-data $\{x_i^*\}_{i=1}^n$ is achieved through Hilbert curves, this method is the so-called Hilbert curve basis selection (HBS) method.

A fundamental question regarding basis selection methods is how to choose the basis functions in order to ensure that the restricted estimator converges to the true function η at the identical rate as the full bases estimator $\hat{\eta}$. It has been shown that both the UBS and ABS methods require that *n* roughly be of the order $O(N^{2/(dr+1)})$, where $d \in [1,2]$ and $r \approx 4$ are constants, depending on the type of spline. In contrast, the SBS only requires that *n* roughly be of the order $O(N^{1/(dr+1)})$ when $p \leq dr + 1$ and the covariates are uniformly distributed in $[0,1]^p$. Thus, the SBS is regarded as a more efficient method. To relax the assumption that all covariates are uniformly distributed in $[0,1]^p$, HBS achieves a similar space-filling property as SBS by increasing the number of bases to $O(N^{2p/\{(p+2)(dr+1)\}})$. The additional basis functions are the price we pay for the irregular distribution of the covariates. Nevertheless, HBS is also more efficient than both UBS and ABS.

To the best of our knowledge, the space-filling type basis selection methods (i.e., SBS and HBS) are only focused on the uniformity of the sub-data $\{x_i^*\}_{i=1}^n$ with respect to the whole input space $[0,1]^p$. Unfortunately, this can result in poor projections of lower dimensional spaces, which is undesirable when the response curve has non-homogeneous smoothness across the input space. Consider a case where a researcher fits a smoothing spline on a two-dimensional space $[0,1]^2$ while the response curve is $\eta(\mathbf{x}) = \sin(15x_1^2)$, which depends solely on the first dimension of the covariate. The true function and the $\{\mathbf{x}_i^*\}_{i=1}^n$ generated via UBS, ABS, SBS, HBS, and the proposed method are demonstrated in Figure 1. It is observed that the covariates selected via the UBS and ABS methods are not uniformly distributed in the input space. In contrast, SBS, HBS, and the proposed method enjoy space-filling properties in the whole input space $[0,1]^2$. It is worth mentioning that the three methods have different behaviours when we only consider the projection on x_1 . One can conclude that the proposed RBS method has better uniformity when we consider the projection on the subspace of the design region compared with the other methods. Thus, the selected basis functions are more likely to capture the volatility in the first dimension in this case, which will undoubtedly lead to better estimation.

In this study, we propose a rotated lattice-based basis selection (RBS) method to improve the projection uniformity with space-filling properties of the selected sub-data. The contributions of this work are threefold as follows. First, a novel sub-data selection is proposed for basis selection when the covariates are not uniformly distributed in $[0,1]^p$, and the response curve may have non-homogeneous smoothness across the input space. Second, it has been proved that only $O(N^{2p/\{(dr+1)(p+2)\}})$ basis functions are needed to achieve the same convergence rate as the full bases estimator, which is the same as with HBS. This implies that our method increases the projective uniformity without involving additional computational costs. Third, numerical experiments show that the RBS method has comparative performance when compared to the HBS and UBS methods and that it outperforms UBS and ABS when the response curves are isomorphic. Moreover, it is uniformly better than the UBS, ABS, SBS, and HBS methods when the response curve has non-homogeneous smoothness.

The remainder of this paper is organized as follows. In Section 2, the preliminaries of smoothing splines and the sub-sampling of basis functions are reviewed. In Section 3, we present the RBS method and an asymptotic analysis. In Section 4, simulation and real data examples are provided to demonstrate the advantage of the proposed basis selection method. Several conclusions and remarks are given in Section 5. All proofs are provided in the supporting information.



FIGURE 1 Illustration of different basis selection methods. The leftmost panel at the top shows the heat map for the true function. The heat maps for the spline estimates based on uniform basis selection (UBS), adaptive basis selection (ABS), space-filling basis selection (SBS), Hilbert curve basis selection (HBS), and the proposed method are presented in the other five panels, respectively. Black dots are the sampled basis functions.

2 | SMOOTHING SPLINES ESTIMATION AND THEIR BASIS SELECTION

In this section, we briefly review the basic background of smoothing splines and introduce the general basis selection methods. For ease of presentation, all vectors are row vectors throughout this study.

Let $\mathcal{H} = \{\eta : J(\eta) < \infty\}$ be an RKHS equipped with a squared semi-norm $J(\cdot)$. Then, \mathcal{H} can be decomposed by $\mathcal{H} = \mathcal{N}_J \bigoplus \mathcal{H}_J$ with $\mathcal{N}_J = \{\eta : J(\eta) = 0\}$ being the null space of $J(\eta)$ and \mathcal{H}_J being the orthogonal complement of \mathcal{N}_J in \mathcal{H} . It is proved that the space \mathcal{H}_J is also a reproducing kernel space associated with a reproducing kernel $G_J(\cdot, \cdot)$ (Gu, 2013).

Assume that \mathcal{N}_{J} is a finite W-dimensional linear subspace of \mathcal{H} with the basis functions $\{\omega_{j}(\cdot)\}_{j=1}^{W}$. According to the well-known Wahba's representer theorem (Grace, 1990), the unique solution (usually called the smoothing spline estimator) to (2) can be represented by

$$\eta(\mathbf{x}) = \sum_{j=1}^{W} \alpha_j \omega_j(\mathbf{x}) + \sum_{i=1}^{N} \beta_i \mathsf{G}_J(\mathbf{x}_i, \mathbf{x}) \, \mathbf{x} \in [0, 1]^p.$$

Denote $\boldsymbol{\alpha} = (\alpha_1, ..., \alpha_W)$ and $\boldsymbol{\beta} = (\beta_1, ..., \beta_N)$. The computation of the smoothing spline estimator is reduced to computing the coefficients $\boldsymbol{\alpha}$ and $\boldsymbol{\beta}$. Denote Y as the response vector, Ω as the $N \times W$ matrix, with the (i, j)th element being $\omega_j(\mathbf{x}_i)$, and G as the $N \times N$ matrix, with the (i, j)th element being $G_j(\mathbf{x}_i, \mathbf{x}_j)$. Then, the original problem (2) becomes that of solving

$$(\hat{\boldsymbol{\alpha}}, \hat{\boldsymbol{\beta}}) = \arg\min_{\boldsymbol{\alpha} \in \mathcal{R}^{W}, \boldsymbol{\beta} \in \mathcal{R}^{N}} \{ N^{-1} (\boldsymbol{Y} - \boldsymbol{\Omega} \boldsymbol{\alpha}^{\mathsf{T}} - \boldsymbol{G} \boldsymbol{\beta}^{\mathsf{T}}) (\boldsymbol{Y} - \boldsymbol{\Omega} \boldsymbol{\alpha}^{\mathsf{T}} - \boldsymbol{G} \boldsymbol{\beta}^{\mathsf{T}})^{\mathsf{T}} + \lambda \boldsymbol{\beta} \boldsymbol{G} \boldsymbol{\beta}^{\mathsf{T}} \}.$$

The computational cost of the standard approach is $O(N^3)$, which is prohibitive when the observation size N is considerable. To reduce this computational burden, basis selection is adopted. The key idea is to further restrict the unknown function $\eta(\cdot)$ in a smaller functional space $\mathcal{H}_S = \mathcal{N}_J \bigoplus \text{span} \{G_J(\mathbf{x}_i^*, \cdot), i = 1, 2, ..., n\}$, where $\{\mathbf{x}_i^*\}_{i=1}^n$ is a subset of $\{\mathbf{x}_i\}_{i=1}^N$. Consequently, the computation of the smoothing spline estimator is reduced to the computation of coefficients α_S and β_S by solving

$$(\hat{\boldsymbol{\alpha}}_{S}, \hat{\boldsymbol{\beta}}_{S}) = \arg\min_{\boldsymbol{\alpha}_{S} \in \mathcal{R}^{W}, \boldsymbol{\beta}_{S} \in \mathcal{R}^{n}} \{ N^{-1} (\boldsymbol{Y} - \boldsymbol{\Omega} \boldsymbol{\alpha}_{S}^{\mathsf{T}} - \boldsymbol{G}_{*} \boldsymbol{\beta}_{S}^{\mathsf{T}}) (\boldsymbol{Y} - \boldsymbol{\Omega} \boldsymbol{\alpha}_{S}^{\mathsf{T}} - \boldsymbol{G}_{*} \boldsymbol{\beta}_{S}^{\mathsf{T}})^{\mathsf{T}} + \lambda \boldsymbol{\beta}_{S} \boldsymbol{G}_{**} \boldsymbol{\beta}_{S}^{\mathsf{T}} \},$$
(3)

where G_* is an $N \times n$ matrix, with the (i,j)th entry $G_J(\mathbf{x}_i, \mathbf{x}_j^*)$ and G_{**} is an $n \times n$ matrix, with the (i,j)th entry $G_J(\mathbf{x}_i^*, \mathbf{x}_j^*)$. Evaluation of the restricted estimator $\tilde{\eta}$ based on sample observations satisfies $\hat{H}_S^T = \Omega \hat{\alpha}_S^T + G_* \hat{\beta}_S^T$, where $\hat{H}_S = \{\tilde{\eta}(\mathbf{x}_1), ..., \tilde{\eta}(\mathbf{x}_n)\}$. Obviously, when $n \ll N$, the computational cost is only $O(N^2n)$, which is a significant reduction compared with $O(N^3)$.

3 | BASIS SELECTION USING ROTATED LATTICES

3.1 | Rotated thinnest covering lattice

The thinnest covering lattice entails finding the best placement of the identical *p*-dimensional spheres that jointly cover the input space. For $2 \le p \le 22$, one can systematically construct the rotated thinnest covering lattice (He, 2017) via

$$L = \{ laM_L R : a \in \mathcal{Z}^p \}, \tag{4}$$

where *I* is a positive constant giving the scale of the lattice cells, M_L is a generator matrix, and *R* is a $p \times p$ rotated matrix. Let *V* be the volume of the input space, that is, the covariate space. A reasonable scaling parameter is $l = n^{-1/p}(p+1)^{1/(2p)}V^{1/p}$, which yields *n* lattice cells in the covariate space $[0,1]^p$ (He, 2017, 2020). One of the most commonly used generator matrices is $M_L = I_p - [\{(1+p)^{1/2}+1+p\}/\{p(p+1)\}]\mathbf{1}_p^T\mathbf{1}_p$, where I_p is the $p \times p$ identity matrix and $\mathbf{1}_p$ is the *p*-dimensional row vector with all elements as one. The rotated matrix *R* can be implemented by any orthogonal matrix. More details can be referred to He (2017). Denote $\{c_i\}_{i=1}^n$ as the elements of the lattice *L* in $[0,1]^p$; then, the Voronoi cell of point c_i is defined as the region $\operatorname{Vor}(c_i) = \{\mathbf{z} | ||\mathbf{z} - c_i||_2 = \min_{c \in L} ||\mathbf{z} - c||_2\}$, where c_i is called the centre of the Voronoi cell $\operatorname{Vor}(c_i)$ and $\|\cdot\|$ denotes the Euclidean norm. Here, we focus on the best known thinnest coverings; other lattices such as the densest packings listed in Conway and Sloane (1998) can also be applied. One may expect to use some other types of lattices to achieve the desired results when *p* is beyond 22.

A two-dimensional case for a rotated thinnest covering lattice in $[0,1]^2$ is demonstrated in the left panel of Figure 2. The hexagons are the Voronoi cells with their centres marked by the crosses, which are the elements of the lattice *L*. Clearly, the hexagon partitioned the space $[0,1]^2$. For comparison, we also show the space partitioned by Hilbert's space-filling curve, which is the main ingredient in the HBS method. This is presented in the right panel of Figure 2. From Figure 2, one can easily see that both the two-dimensional rotated thinnest covering lattice and the second-order Hilbert's space-filling curve yield reasonable partitions of $[0,1]^2$. Clearly, the 16 crosses in the two panels are uniform over the whole covariate space; however, the uniformity in the left panel is better than that in the right panel. Moreover, all projections of the 16 crosses in the left panel onto each axis have a space-filling character. However, some projections of the 16 crosses in the right panel onto any axis are coincident, and only four different projection points are obtained. Judging from the projection uniformity of each subspace, the rotated thinnest covering lattice is far superior to Hilbert's space-filling curve. Thus, the $\{x_i^*\}_{i=1}^n$ selected by the rotated thinnest covering lattice will benefit cases



FIGURE 2 The hexagonal partitioning of $[0,1]^2$ with the rotated thinnest covering lattice on the left and the second-order Hilbert's space-filling curve on the right shows cells (blue squares or hexagons), centres (red crosses), and the second-order Hilbert's space-filling curve (black solid line).

where the response curve has non-homogeneous smoothness. Proposition 1 gives the space-filling property for the general rotated thinnest covering lattice.

Proposition 1. For any two covariates $\mathbf{x}, \mathbf{x}' \in [0,1]^p$ located in $Vor(\mathbf{c}_i), i = 1, 2, ..., n$, we have

$$\|\mathbf{x} - \mathbf{x}'\| \le 2(p+1)^{1/(2p)} \{p(p+2)\}^{1/2} \{12(p+1)\}^{-1/2} n^{-1/p}$$

3.2 | The rotated lattice-based basis selection algorithm

The proposed RBS algorithm includes the following three main steps:

Step 1. Generate a rotated thinnest covering lattice *L* with *n* Voronoi cells $\{Vor(\mathbf{c}_i)\}_{i=1}^n$ in $[0,1]^p$. Divide the covariates of *N* observations into *n* cells according to the distance information of *p*-dimensional covariates to Voronoi cells $\{Vor(\mathbf{c}_i)\}_{i=1}^n$;

Step 2. Adjust the estimated value of the volume of non-empty cells, so that the size of the non-empty cells \hat{n} is roughly equal to n;

Step 3. Randomly select one covariate from each non-empty cell to form $\{\mathbf{x}_i^*\}_{i=1}^n$, and then, estimate the smoothing spline on the space \mathcal{H}_S as in (3).

A rotated thinnest covering lattice *L* is generated in the first step. After achieving the lattice *L*, we will map the covariates to the Voronoi cells in *L*. Clearly, performing an exhaustive search of all the Voronoi cells in *L* and then deciding to which cell a particular covariate belongs requires $O(np^2)$ time.

In the second step, denote \hat{n} as the size of non-empty cells. Then, we should adjust the estimated value of the volume of non-empty cells so that \hat{n} is closer to *n*. For clarity, we denote the estimated value by \hat{V} . Initially, $\hat{V} = 1$, that is, the volume of the covariate space $[0,1]^p$. Then, *I* is estimated by $n^{-1/p}(p+1)^{1/(2p)}\hat{V}^{1/p}$. When the difference between \hat{n} and the target *n* is too large, we should adjust *I* according to a new estimated \hat{V} , so that the difference between \hat{n} and the target *n* loss not exceed 0.1*n*.

In the last step, we randomly select one covariate allocated in each non-empty cell. Then, the total \hat{n} covariates $\{\mathbf{x}_i^*\}_{i=1}^{\hat{n}}$ are collected. If $n > \hat{n}$, we randomly add $n - \hat{n}$ covariates from the remaining $N - \hat{n}$ covariates $\{\mathbf{x}_i^*\}_{i=\hat{n}+1}^N$. If $n < \hat{n}$, we randomly delete $\hat{n} - n$ covariates from the collected \hat{n} covariates $\{\mathbf{x}_i^*\}_{i=\hat{n}+1}^N$. Suppose that, without loss of generality, the selected covariates are denoted as $\{\mathbf{x}_i^*\}_{i=1}^n$.

Finally, the selected covariates $\{\mathbf{x}_i^*\}_{i=1}^n$ are used to construct the effective subspace \mathcal{H}_S . Then, the smoothing spline estimator $\tilde{\eta}$ of η is achieved through (3). We summarize the proposed method in Algorithm 1.

Algorithm 1 Basis selection method using a rotated lattice

1: Compute $l = n^{-1/p} (p+1)^{1/(2p)} \hat{V}^{1/p}$ with the initial $\hat{V} = 1$.

2: for $1 \le i \le N$ do

- 3: Determine to which cell x_i belongs.
- 4: end for
- 5: Obtain the number of non-empty cells, denoted as \hat{n} .

6: Adjust I according to a new estimated \hat{V} and repeat the above process until the difference between \hat{n} and n does not exceed 0.1n.

7: Randomly select one covariate from each non-empty cell; then, collect \hat{n} covariates $\{\mathbf{x}_i^*\}_{i=1}^{\hat{n}}$.

8: if $\hat{n} > n$ then

9: Randomly delete $\hat{n} - n$ covariates from the collected \hat{n} covariates $\{\mathbf{x}_{i}^{*}\}_{i=1}^{\hat{n}}$.

10: else

11: Randomly add $n - \hat{n}$ covariates from the remaining $N - \hat{n}$ covariates $\{\mathbf{x}_i^*\}_{i=\hat{n}+1}^N$.

12: end if

13: Denote the selected sub-data as $\{\mathbf{x}_{i}^{*}\}_{i=1}^{n}$.

14: Minimize the penalized least squares criterion (2) over the effective subspace $\mathcal{H}_{S} = \mathcal{N}_{J} \oplus \text{span}\{G_{J}(\mathbf{x}_{i}^{*}, \cdot), i = 1, 2, ..., n\}$.

The computational cost for Algorithm 1 mainly depends on dividing *N* covariates into *n* cells and estimating the smoothing spline estimator in the last step. Note that Algorithm 1 requires O(np) operations to obtain the cell closest to a given covariate. Therefore, the computational complexity of finding the cells closest to *N* covariates is of the order O(Nnp). As discussed in Meng et al. (2020, 2021), estimating the smoothing spline

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estimator $\tilde{\eta}$ of η over the effective subspace \mathcal{H}_S is of the order $O(Nn^2)$. Since *n* is much larger than *p*, O(Nnp) is negligible compared with $O(Nn^2)$. In summary, the overall computational cost of the RBS method in Algorithm 1 is in the order of $O(Nn^2)$.

3.3 | Theoretical analysis

In this section, the asymptotic properties of the smoothing spline estimator using the proposed RBS method are presented. Let f_X denote the probability density function of the covariate. Define the mean square error of the estimator $\hat{\eta}$ in estimating η as the quadratic functional

$$\Lambda(\hat{\eta} - \eta) = \int_{\mathcal{X}} (\hat{\eta}(\mathbf{x}) - \eta(\mathbf{x}))^2 f_X(\mathbf{x}) d\mathbf{x}$$

Conventionally, the rate of asymptotic convergence of $\hat{\eta}$ is described by eigenvalue analysis of $J(\eta)$ in (2) with respect to $\Lambda(\eta)$. In the following, we shall use $\Lambda(\cdot, \cdot)$ and $J(\cdot, \cdot)$ to represent the (semi) inner products associated with the square (semi) norms Λ and J.

Below, we introduce six primary regularity conditions, which are the same as in the HBS method. A more detailed discussion can be found in Meng et al. (2021).

Condition 1. The functional Λ is completely continuous with respect to *J*, where $\Lambda(\phi_{\gamma}, \phi_{\mu}) = \delta_{\gamma\mu}, J(\phi_{\gamma}, \phi_{\mu}) = \rho_{\gamma}\delta_{\gamma\mu}, \phi_{\gamma}$, and ϕ_{μ} are the eigenfunctions associated with Λ and *J* in $\mathcal{H}, \rho_{\gamma}$ is the nonnegative eigenvalue associated with ϕ_{γ} , and $\delta_{\gamma\mu}$ denotes the Kronecker delta;

Condition 2. For some $\beta > 0$ and r > 1, $\rho_{\gamma} > \beta \gamma^{r}$ for sufficiently large γ ;

Condition 3. For all μ, γ and a positive constant C, var $\{\phi_{\gamma}(\mathbf{x})\phi_{\mu}(\mathbf{x})\} \leq C$;

Condition 4. For all μ and γ , $\phi_{\gamma}(\mathbf{x})\phi_{\mu}(\mathbf{x})$ is Lipschitz continuous; that is, for any $\mathbf{x}, \mathbf{x}' \in \mathcal{X}$, there exists a positive constant B such that $|\phi_{\gamma}(\mathbf{x})\phi_{\mu}(\mathbf{x}) - \phi_{\gamma}(\mathbf{x}')\phi_{\mu}(\mathbf{x}')| \le B||\mathbf{x} - \mathbf{x}'||$;

Condition 5. Suppose $\max\{(nN_i)/N\}_{i=1}^n = O_P(1)$, where N_i is the number of covariates in the *i*th cell;

Condition 6. The number of bases *n* satisfy that $n^{1+2/p} = O(N)$.

The first four conditions are the regularity conditions for smoothing spline estimators even if the full bases are used. Condition 5 prevents the extreme case of the probability density function f_X , that is, when there is only one non-empty cell; then, we have $\max\{(nN_j)/N\}_{j=1}^n = nN/N = n$, which conflicts with Condition 5. Lastly, Condition 6 naturally holds when *n* is a manageable size. For example, when p = 2, Condition 6 holds for $n = O(N^{1/2})$.

Denote $\tilde{\eta}_R$ as the estimator of η obtained by the basis functions $\{G_J(\mathbf{x}_i^*, \cdot)\}_{i=1}^n$ selected using the RBS method. The following theorem shows that $\tilde{\eta}_R$ converges to the true function η_0 at the same rate as $\hat{\eta}$, whose proof is provided in the supporting information.

Theorem 1. Assume that $\sum \rho_{\gamma}^{d} \Lambda(\eta_{0}, \phi_{\gamma})^{2} < \infty$ for some $d \in [1,2]$. Under Conditions 1–6, as $\lambda \to 0$ and $n^{1+2/p} \lambda^{2/r} \to \infty$, we have $(\Lambda + \lambda J)(\tilde{\eta}_{R} - \eta_{0}) = O_{P}(N^{-1}\lambda^{-1/r} + \lambda^{d})$. Particularly, when $\lambda \simeq N^{-r/(dr+1)}, \tilde{\eta}_{R}$ achieves the optimal convergence rate $(\Lambda + \lambda J)(\tilde{\eta}_{R} - \eta_{0}) = O_{P}(N^{-dr/(dr+1)})$.

Denote $\tilde{\eta}_U, \tilde{\eta}_A, \tilde{\eta}_S$, and $\tilde{\eta}_H$ as the estimators of η obtained based on observations selected using the UBS, ABS, SBS, and HBS methods, respectively. Obviously, the convergence rates of $\tilde{\eta}_U, \tilde{\eta}_A, \tilde{\eta}_S$, and $\tilde{\eta}_H$ are identical to that of $\hat{\eta}$ with different essential choices of n. The convergence rates of $\tilde{\eta}_U$ and $\tilde{\eta}_A$ are the same as that of $\hat{\eta}$ when the size of the basis functions n is of the order $O(N^{2/(dr+1)})$. For the estimator $\tilde{\eta}_R, n$ is of the order $O(N^{2/(dr+1)})$, which is smaller than those of $\tilde{\eta}_U$ and $\tilde{\eta}_A$. The order of n in estimating $\tilde{\eta}_S$ is $O(N^{1/(dr+1)})$, which is smaller than that for the proposed $\tilde{\eta}_R$. However, the estimator $\tilde{\eta}_S$ requires that the covariates be uniformly distributed in the whole covariate space. Once the covariates are not uniformly distributed, $\tilde{\eta}_S$ may not perform well, compared with the proposed $\tilde{\eta}_R$. The order of n in $\tilde{\eta}_H$ is the same as $\tilde{\eta}_R$ for the response curve in general. When the response curve has homogeneous smoothness across the covariate space, our method performs similarly to the HBS method. Additionally, the selected sub-data enjoy a pretty projection property in lower dimensional subspaces through a well-designed rotation, instead of increasing the sub-data size. Thus, it has some advantages in cases where the response curves have non-homogeneous smoothness.

The values of *d* and *r* are determined by the roughness of η and the order of the fitted spline, respectively. More details can be found in Gu and Kim (2002), Ma et al. (2015), and Meng et al. (2020, 2021). In our numerical study, we set *r* to 4, as Ma et al. (2015) and Gu and Kim (2002) have suggested. Hence, we can obtain that *n* is roughly located in the interval $(O(N^{2p/\{9(p+2)\}}), O(N^{2p/\{5(p+2)\}}))$. In the following real data analysis, we take the dimension *n* of the effective subspace \mathcal{H}_5 to be between $15N^{2/9}$ and $30N^{2/9}$.

4 | SIMULATION AND REAL DATA STUDIES

In this section, we compare our proposed basis selection method to several other methods using five numerical examples and two real data examples in terms of prediction accuracy with the log mean absolute error (MAE) given by

$$\log(\mathsf{MAE}) = \log\left(N_{test}^{-1}\sum_{i=1}^{N_{test}} |\tilde{\eta}(\mathbf{x}_i) - \eta_0(\mathbf{x}_i)|\right)$$

where N_{test} is the size of the test data, $\{\mathbf{x}_i\}_{i=1}^{N_{test}}$ is an independent test dataset generated using the same probability density function as in the training data, and $\eta_0(\mathbf{x}_i)$ and $\tilde{\eta}(\mathbf{x}_i)$ denote the real and fitted response values from the *i*th test data point, respectively. The methods in the comparison are the UBS method in Gu and Kim (2002), the ABS method in Ma et al. (2015), the SBS method in Meng et al. (2020), and the HBS method in Meng et al. (2021).

4.1 | Simulation studies

We generate a synthetic training dataset with N = 2000 from each of the following four probability density functions, and then, the covariate is scaled to $[0,1]^p$. The signal-to-noise ratio, defined as $var(\eta(\mathbf{x}))/\sigma^2$, is set to two. We then use the five basis selection methods to obtain a subset of $n = \{40,60,80,100,120\}$ observations and fit the smoothing spline model.

We consider four distributions of the covariate variable x over the domain of interest, identical to those considered in Meng et al. (2021). The p in the following dataset are adapted to the functions f_1 to f_5 .

d₁: The *p*-dimensional covariate variable **x** is uniformly distributed in $[0,1]^p$;

d₂: The *p*-dimensional covariate variable **x** follows a mixture *t*-distribution $(T_1,...,T_p)$, where $\{T_i\}_{i=1}^p$ are independently generated from t(10, -5)/2 + t(10,5)/2;

d₃: The *p*-dimensional covariate variable **x** obeys a banana-shaped distribution, which is generated by $(Z_1, Z_2 + Z_1^2/1.2, ..., Z_p + Z_1^2/1.2)$, where $(Z_1, Z_2, ..., Z_p)$ is generated from the standard multivariate Gaussian distribution;

d₄: The *p*-dimensional covariate variable **x** obeys a multivariate Gaussian distribution $GP(0, \Sigma)$, where the (i, j)th element of Σ is $\Sigma_{i,i} = 0.9^{|i-j|}, i, j = 1, ..., p$.

We adopt five different regression function settings. The first four are the same as those considered in Meng et al. (2021), and the last one is similar to that in Ma et al. (2015):

f₁: A two-dimensional function given by

$$f_1(\mathbf{x}) = \sin\{10/(x_1 + x_2 + 0.15)\};$$

f₂: A two-dimensional function given by

$$f_2(\mathbf{x}) = h_1(x_1, x_2) + h_2(x_1, x_2),$$

where

$$h_1(x_1, x_2) = \{0.75/(\pi\sigma_1\sigma_2)\} \times \exp\{-(x_1 - 0.2)^2/\sigma_1^2 - (x_2 - 0.3)^2/\sigma_2^2\}, h_2(x_1, x_2) = \{0.75/(\pi\sigma_1\sigma_2)\} \times \exp\{-(x_1 - 0.7)^2/\sigma_1^2 - (x_2 - 0.5)^2/\sigma_2^2\},$$

 $\sigma_1 = 0.1$, and $\sigma_2 = 0.2$;

f₃: A three-dimensional function given by

$$f_3(\mathbf{x}) = \sin\{\pi(x_1 + x_2 + x_3)/3\} - x_1 - x_2^2;$$

f₄: A four-dimensional function given by

$$f_4(\mathbf{x}) = x_1 + (2x_2 - 1)^2 / 2 + [\sin(10\pi x_3) / \{2 - \sin(10\pi x_3)\}] / 3 + \{0.1\sin(2\pi x_4) + 0.2\cos(4\pi x_4) + 0.3\sin(6\pi x_4)^2 + 0.4\cos(8\pi x_4)^3 + 0.5\sin(10\pi x_4)^3 \} / 4;$$

f₅: A copula function with only two important covariates given by

$$f_5(\tilde{\mathbf{x}}) = (2\pi)^{-1} |\Sigma|^{-1/2} \exp\{-1/2(g_1(\tilde{\mathbf{x}}_1), g_2(\tilde{\mathbf{x}}_2))^T \Sigma^{-1}(g_1(\tilde{\mathbf{x}}_1), g_2(\tilde{\mathbf{x}}_2))\} |g_1'(\tilde{\mathbf{x}}_1)g_2'(\tilde{\mathbf{x}}_2)|$$

where $g_1(\tilde{x}_1) = 2 \operatorname{sign}(\tilde{x}_1) |\tilde{x}_1|^2$, $g_2(\tilde{x}_2) = 3 \widetilde{x}_2^3$, $\tilde{x}_j = 2.3(x_j - 0.5)$ for j = 1, 2, and Σ is a 2 × 2 matrix, with the (*i*, *j*)th entry being $0.5^{|i-j|}$, *i*, *j* = 1, 2.

Figure 3 predicts $N_{test} = 2000$ testing data, records the log(MAE), and calculates the standard errors shown in vertical bars, which are obtained from a hundred replicates. The full bases estimator is omitted here due to the high computation cost. Figure 3 illustrates that all the methods in the first column have similar performances, while the proposed RBS method performs slightly better and the UBS method performs slightly worse, in which cases the observed values are uniformly distributed in a hypercube. Meanwhile, in the right three columns, we observe that the proposed RBS method yields perfect performance compared with the UBS, ABS, and SBS methods, followed by the HBS method, in cases where the covariates are not uniformly distributed in a hypercube. Moreover, the UBS, ABS, and SBS methods perform unstably in the non-uniformly distributed case. Overall, our RBS method has the upper hand in all settings, yielding lower log(MAE) and smaller standard errors, which implies that the RBS method is feasible for a broad range of covariate distributions, surface functions, and dimensions and thus emerges as the best estimator of the true function, compared with other methods.

In the following, we consider a scenario in which the underlying response surface is not isomorphic to further illustrate the advantages of the proposed method. To be precise, we consider a smoothing spline on d_5 with *p* varying from three to eight. The underlying response surface is taken as f_5 , which only relies on the first two dimensions of d_5 . Note that there is at least one redundant covariate in this example, which makes



FIGURE 3 log(MAE) values from the smoothing spline model for simulation under five different regression functions (from upper to lower) and four different probability density functions (from left to right) are plotted versus different *n*. ABS, adaptive basis selection; HBS, Hilbert curve basis selection; MAE, mean absolute error; RBS, rotated lattice-based basis selection; SBS, space-filling basis selection; UBS, uniform basis selection.



FIGURE 4 $\log(MAE)$ values from the smoothing spline model for simulation under regression function f_5 and probability density function d_4 with one, two, three, four, five, and six redundant covariates (from upper to lower and left to right) are plotted versus different *n*. The response surface is not isomorphic due to the existence of redundant covariates. ABS, adaptive basis selection; HBS, Hilbert curve basis selection; MAE, mean absolute error; RBS, rotated lattice-based basis selection; SBS, space-filling basis selection; UBS, uniform basis selection.

the response surface not isomorphic. Figure 4 reports the log(MAE)s of all five basis selection methods when redundant covariates exist in the settings of regression functions and the dataset is not uniformly generated in a hypercube.

The results show that the performance of the RBS method is remarkably better when the function has non-homogeneous smoothness and that the performance of the RBS method is comparable with that of the HBS method when the function is isomorphic. This is because when the response surface is not isomorphic, the uniformity on the subspace with relatively large volatility is more important than other subspaces. Compared with the uniformity in full data space, the projective uniformity is much more appreciated in this scenario since it achieves a better uniformity on all k(< p) dimensional subspace than the uniform design in full space. Consequently, the proposed RBS method is more advantageous than the other four methods since it achieves better projective uniformity.

To clearly see the projective uniformity of the proposed method, we evaluate the proposed method together with the other four methods by the log maximum projection criterion proposed in Joseph et al. (2015). To be precise, the log maximum projection is calculated by

$$\log(\psi(D)) = \log\left(\frac{1}{n(n-1)}\sum_{1 \le i < j \le n}\prod_{k=1}^{p} (x_{i,k}^* - x_{j,k}^*)^{-2}\right)^{1/p},$$

with $D = {\mathbf{x}_i^*}_{i=1}^n$, which is the dataset we use to construct the basis functions in \mathcal{H}_5 . Clearly, a smaller value implies better projection uniformity since the data points in D will not be close to each other in any dimensions. For cases where $(\mathbf{x}_{i,k}^* - \mathbf{x}_{j,k}^*) = 0$ for some i, j, and k, we simply denote the results as NA, implying that projective uniformity is not available for the sub-data D. The projection uniformity as measured by $\log(\psi(D))$ for the five methods is shown in Table 1 under f_5 and d_4 with redundant covariates.

TABLE 1 $\log{\{\psi(D)\}}$ values of the five basis selection methods for simulation under regression function f₅ and probability density function d₄ with n = 80 basis functions.

p	UBS	ABS	SBS	HBS	RBS
3	9.551428	9.501454	NA	NA	8.357445
4	9.108542	8.869277	NA	NA	7.794891
5	8.710636	8.749605	NA	NA	7.684428
6	8.642750	8.738538	NA	8.865160	7.744047
7	8.358219	8.379835	NA	8.198035	7.439442
8	8.105443	7.993638	NA	7.989379	7.290941

Note: The best performance of each setting is highlighted in boldface.

Abbreviations: ABS, adaptive basis selection; HBS, Hilbert curve basis selection; NA, $\log\{\psi(D)\}\$ is not applicable to the sub-data *D*; RBS, rotated lattice-based basis selection; SBS, space-filling basis selection; UBS, uniform basis selection.

TABLE 2	Computing time (in second	ls) of the five basis se	lection methods and fu	Il data for simulation	under regression	function f ₅ and
probability de	ensity function d_4 with $n = -$	{40,60,80,100,120} I	pasis functions.			

n	UBS	ABS	SBS	HBS	RBS	Full data
40	1.2402	1.2221	1.2063	1.2610	1.5928	
60	1.5383	1.5937	1.3282	1.6333	1.9558	
80	1.7584	1.7986	1.5685	1.8367	2.2261	37.7657
100	1.9162	1.9991	1.7520	2.0945	2.5139	
120	2.0824	2.1675	1.9644	2.3453	2.7677	

Abbreviations: ABS, adaptive basis selection; HBS, Hilbert curve basis selection; RBS, rotated lattice-based basis selection; SBS, space-filling basis selection; UBS, uniform basis selection.

Based on the result presented in Table 1, one can clearly see that the $\log{\{\psi(D)\}}$ value of RBS is the smallest among the other four methods, which implies the data points selected via our methods have a better projective uniformity on the subspace spanned by the first two covariates.

In the following, we will evaluate the performance of all five methods in computing time. Since all the cases have similar performance, we only illustrate the results that the covariates are generated as d_4 and the regression function is f_5 with *n* varying from 40 to 120. We record the computing time for our desktop PC with 8 G memory and a 2.4 GHz Celeron processor. Each method has been repeated 100 times, and the corresponding results are reported in Table 2.

From Table 2, it can be clearly seen that all methods significantly reduce the computing time compared with the full data approach since the computational results are reduced from $O(N^3)$ to $O(Nn^2)$. As discussed in Section 3.3, all the basis selection methods have comparable performance since fitting the smoothing spline ANOVA model takes the most computing time.

4.2 | White River pollutant data

This dataset contains the pollutant concentrations at different locations in China's White River. The main task is to predict the pollutant concentration of unmonitored locations. The variable pollutant concentration is treated as the response, and the corresponding longitude and latitude are treated as two covariates. The dataset contains N = 3,965 training data and $N_{test} = 396$ test data. We select $n = N^{2/9} \{15,20,25,30\}$ basis functions from the N = 3,965 training data using the UBS, ABS, SBS, HBS, and RBS methods. We fit the cubic tensor product smoothing spline analysis of the variance model to the dataset and consider the following model settings:

M1: An additive smoothing spline regression model with all two main effects and the two-way interaction function of corresponding covariates,

$$y_i = \eta_{\emptyset} + \eta_1(x_{i1}) + \eta_2(x_{i2}) + \eta_{1,2}(x_{i1}, x_{i2}) + \epsilon_i, i = 1, 2, ..., N;$$

M₂: A redundant covariate x_{i3} , for example, the depth of the river, independently generated through a uniform distribution on [0,1], is added to the respected model. To be precise, an additive smoothing spline regression model with all three main effects and the two-way interaction of x_{i1} and x_{i2} ,



FIGURE 5 $\log(MAE)$ values versus different *n* under model setting M₁ and M₂ from the smoothing spline model for the pollutant data. The horizontal line represents the performance for all the training data. ABS, adaptive basis selection; HBS, Hilbert curve basis selection; MAE, mean absolute error; RBS, rotated lattice-based basis selection; SBS, space-filling basis selection; UBS, uniform basis selection.

$$y_i = \eta_{\emptyset} + \eta_1(x_{i1}) + \eta_2(x_{i2}) + \eta_3(x_{i3}) + \eta_{1,2}(x_{i1}, x_{i2}) + \epsilon_i, i = 1, 2, ..., N,$$

where y_i is the contaminant concentration measurement at the *i*th observation, x_{ij} is the value of the *j*th dimension at the *i*th covariate, η_{\emptyset} is a constant function, $\{\eta_j\}_{j=1}^3$ are main effect functions, $\eta_{1,2}$ is the two-way interaction function of the corresponding covariate, and $\{\epsilon_i\}_{i=1}^N$ represent the independent and identically distributed random errors with zero mean and unknown variance σ^2 . It is worth mentioning that the $R^2 = 0.7543$, which implies the fitness of M₁. By applying model checking tools in (Gu, 2013, Chapter 3.7), one can find a strong correlation between all three terms with the response. To show the effectiveness of the proposed estimator, we compare it with four mainstream competitors, as mentioned in the previous section, in terms of the prediction log(MAE), calculated on a holdout testing set, and summarize the results in Figure 5.

As shown in the left panel of Figure 5, the log(MAE)s of all five basis selection methods come closer with the increasing basis function size *n*. As expected, in models M₁ and M₂, RBS is superior to the other four basis selection methods, which echoes the results in Section 4.1.

4.3 | Debutanizer column data

This dataset comprises debutanizer distillation column data from the process of separating butane from gasoline. The task is to predict the butane content by using the conditions of the debutanizer columns and other relevant information measured using soft sensors in the process of petroleum refining. Seven relevant features are provided in the dataset, such as temperature, pressure, flow, and so forth. More details about the dataset can be found in Fortuna et al. (2007).

We first scale the dataset to [0,1]⁷ and then fit the cubic tensor product smoothing spline analysis of the variance model to the scaled dataset. We consider the following model setting:

M₃: An additive smoothing spline regression model with all main effects,

$$y_i = \eta_{\emptyset} + \sum_{j=1}^{7} \eta_j(x_{ij}) + \epsilon_i, i = 1, 2, ..., N,$$

where y_i is the butane concentration of the *i*th observation, x_{ij} is the value of the *j*th dimension of the *i*th covariate, η_{\emptyset} is a constant function, $\{\eta_i\}_{i=1}^7$ are main effect functions, and $\{\epsilon_i\}_{i=1}^N$ represent the independent and identically distributed random errors with zero mean and unknown variance. It is worth mentioning that the cosine values of all possible interaction terms are no more than 0.29. Thus, according to Gu (2013), we opt to consider this smoothing spline ANOVA model with main effects only for model brevity.

The simulation is performed under the setting that the size of the full data is 2395, 10% of the full data is randomly selected without repetition as test data, the remaining data are used as training data, and set $n = N^{2/9} \{15, 20, 25, 30\}$.



FIGURE 6 log(MAE) values versus different *n* under model setting M₃ from the smoothing spline model for the debutanizer distillation column data. The horizontal line represents the performance for all the training data. ABS, adaptive basis selection; HBS, Hilbert curve basis selection; MAE, mean absolute error; RBS, rotated lattice-based basis selection; SBS, space-filling basis selection; UBS, uniform basis selection.

As shown in Figure 6, with the increase of the size *n* of the basis functions, the log(MAE)s of all the five basis selection methods come closer to one another. Furthermore, as expected, in Figure 6, RBS is superior to the other four basis selection methods, which is identical to the results in Section 4.1 and Section 4.2.

5 | CONCLUSION AND DISCUSSION

This study introduces a new basis selection method called RBS for large *N* datasets. The RBS method is an efficient stratified basis selection method in which the strata are Voronoi cells of rotated thinnest covering lattices, which does not require the assumption that the covariates are uniformly distributed on a hypercube and that there is homogeneous smoothness across the covariate space.

Theoretically, we prove that for the proposed RBS method, when the size of the basis functions *n* is roughly in the order $O\{N^{2p/(dr+1)(p+2)}\}$, the convergence rate of $\tilde{\eta}_R$ is identical to that of $\hat{\eta}$, which is the same as that of the HBS method, and is also faster than that of the UBS and ABS methods. Furthermore, comprehensive simulation and real data studies were conducted to illustrate the performance of the RBS method, and the numerical results are consistent with our theoretical results. Compared with the other four existing methods, the RBS method is more effective, especially when the covariates are not uniformly distributed on a hypercube and the response is not isotropic since the RBS method improve the projective uniformity substantially via rotating the lattice.

The newly proposed basis selection method may be suitable for other nonparametric models, including but not limited to the Gaussian process model and the nearest neighbour model. However, more work needs to be done to verify its performance. It would also be interesting to explore whether the new method can be modified by considering the response information.

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DATA AVAILABILITY STATEMENT

The data that support the findings of this study are available on request from the corresponding author. The data are not publicly available due to privacy or ethical restrictions.

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SUPPORTING INFORMATION

Additional supporting information can be found online in the Supporting Information section at the end of this article.

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