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Large scale Gaussian processes with Matheron's update rule and Karhunen-Loève expansion

Hassan Maatouk and Didier Rullière and Xavier Bay

Abstract Gaussian processes have become essential for non-parametric function estimation and widely used in many fields like machine learning. In this paper, large scale Gaussian process regression (GPR) is investigated. This problem is related to the simulation of high dimensional Gaussian vectors truncated on the intersection of a set of hyperplanes. The main idea is to combine both Matheron's update rule (MUR) and Karhunen-Lovève expansion (KLE). First, by the MUR we show how simulating from the posterior distribution is possible without computing the posterior covariance matrix and its decomposition. Second, by splitting the input domain in smallest nonoverlapping subdomains, the KLE coefficients are conditioned in order to guarantee the correlation structure in the entire domain. The parallelization of this technique is developed and the advantages are highlighted. By this, the computational complexity is drastically reduced. The mean-square *block* error is computed. It provides accurate results when using a family of covariance functions with compact support. Some numerical examples to study the performance of the proposed approach are included.

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1 Introduction

Due to their flexibility, Gaussian processes (GPs) are widely used in many fields like geostatistics [5, 9], finance [1, 7] and econometrics [4] and have become very popular in the context of machine learning [16, 17]. In this paper, large scale GPR is investigated. This problem is related to the simulation of high dimensional Gaussian vectors constrained on a set of intersection of hyperplanes. The direct approach is based on computing the posterior distribution and using the location scale transformation of the posterior covariance matrix (i.e., the *scaling* matrix). After computing the *scaling* matrix, we sample from a standard multivariate normal (MVN) distribution [11]. Computing the *scaling* matrix is possible via for example eigendecomposition or Cholesky factorization. When the dimension of the Gaussian vector is high, this approach becomes numerically heavy. This is because the computational complexity scales cubically with the dimension of the random vector [6].

The methodology presented in this paper is quite different. It is based on combining both Matheron's update rule (MUR) and Karhunen-Loève expansion (KLE). In the first hand, the MUR which had been discovered in geostatistics [9] and then in astrophysics [8] is developed. Contrarily to the direct approach, it is based on generating from the prior distribution and adding the update part to obtain the target posterior distribution. The advantage of this approach is that we sample before conditioning rather than after. So, we do not need to compute the posterior covariance matrix and its decomposition. In [3], the MUR is used to efficiently sampling multivariate normal distribution whose covariance (precision) matrix can be decomposed as a positive-definite matrix minus (plus) a low-rank symmetric matrix. The main idea is to sample from a block diagonal covariance matrix. Recently, in the context of machine learning, the authors in [17] use the MUR for GPR. This approach is denoted *pathwise conditioning*. In the second hand, the KLE can be seen as an efficient way to sample random fields, which is based on computing the eigendecomposition of the covariance operator [10]. In high dimensions, the eigendecomposition becomes numerically heavy. To handle this problem, the idea is to split the input domain into smallest nonoverlapping subdomains and to condition the KLE coefficients in order to respect the given correlation in the entire domain [2, 13].

In the present paper, we first investigate the advantage of the MUR in the context of GPR. Then, we develop the large scale KLE for sampling the prior distribution. The parallelization of this approach is studied which drastically reduces the computational complexity. Finally, the *truncation* and *block* errors induced from the proposed approach are computed.

The article is structured as follows: in Sect. 2 GPR is briefly reviewed. Sect. 3 is devoted to the MUR in the context of GPR. In Sect. 4, the large scale KLE is developed where the parallelization is investigated and the induced errors are computed. In Sect. 5, the performance of the proposed approach is shown through numerical examples.

2 Gaussian process regression

For any $x \in \mathbb{R}^d$, suppose $(Z(x))$ is a GP with mean function μ and covariance function k . Then

$$Z(x) = \mu(x) + Y(x), \quad \forall x \in \mathbb{R}^d,$$

where $(Y(x))$ is a zero-mean GP with covariance function k ,

$$k(x, x') = \text{Cov}(Y(x), Y(x')), \quad \forall x, x' \in \mathbb{R}^d.$$

Given a training data set $D = \{(x_i, y_i), i = 1, \dots, n\}$ of n input-output pairs, where x_i denotes an input vector of dimension d and y_i denotes a scalar output. The input vectors are aggregated in the $d \times n$ design matrix \mathbb{X} and the outputs are collected in the vector \mathbf{y} , so we can write $D = \{(\mathbb{X}, \mathbf{y})\}$. The following problem is considered

$$y_i = f(x_i) + \epsilon_i, \quad \epsilon_i \sim \mathcal{N}(0, \sigma_{\text{noise}}^2), \quad (1)$$

where f is an unknown latent function generates the outputs and ϵ_i is an additive independent identically distributed Gaussian noise with constant noise variance σ_{noise}^2 estimated via the maximum likelihood [16]. A GP prior distribution on the unknown function f is assumed. Conditionally to the observations $\mathbf{y} = (y_1, \dots, y_n)^\top$, the conditional process remains a GP

$$\{Y(x)|Y(\mathbb{X}) = \mathbf{y}\} \sim \mathcal{GP}(\mu_c, c), \quad (2)$$

where the conditional mean μ_c and covariance functions c are given as follows:

$$\begin{aligned} \mu_c(x) &= \text{E}[Y(x)|\mathbf{y}] = k(x, \mathbb{X})^\top (k(\mathbb{X}, \mathbb{X}) + \sigma_{\text{noise}}^2 \mathbf{I}_n)^{-1} \mathbf{y}; \\ c(x, x') &= k(x, x') - k(x, \mathbb{X})^\top (k(\mathbb{X}, \mathbb{X}) + \sigma_{\text{noise}}^2 \mathbf{I}_n)^{-1} k(x', \mathbb{X}); \end{aligned} \quad (3)$$

with \mathbf{I}_n the $n \times n$ identity matrix. Let us recall that $k(\mathbb{X}, \mathbb{X})$ is the covariance matrix of $Y(\mathbb{X})$ and $k(x, \mathbb{X})$ is the vector of covariance between $Y(x)$ and $Y(\mathbb{X})$.

In the simple special case where the observations are noise-free [14], that is we know $\{(x_i, f_i)|i = 1, \dots, n\}$, with $f_i = f(x_i)$, the predictive equations for GPR (3) are conserved where we replace σ_{noise}^2 by zero and \mathbf{y} by \mathbf{f} , with $\mathbf{f} = (f_1, \dots, f_n)^\top$.

Table 1 Some popular covariance functions with their degree of smoothness [16].

Name	Expression	Class
Squared Exponential	$\exp\left(-\frac{(x-x')^2}{2\theta^2}\right)$	\mathcal{C}^∞
Matérn $\nu = 5/2$	$\left(1 + \frac{\sqrt{5} x-x' }{\theta} + \frac{5(x-x')^2}{3\theta^2}\right) \exp\left(-\frac{\sqrt{5} x-x' }{\theta}\right)$	\mathcal{C}^2
Matérn $\nu = 3/2$	$\left(1 + \frac{\sqrt{3} x-x' }{\theta}\right) \exp\left(-\frac{\sqrt{3} x-x' }{\theta}\right)$	\mathcal{C}^1
Exponential	$\exp\left(-\frac{ x-x' }{\theta}\right)$	\mathcal{C}^0

Table 1 shows some popular covariance functions used in machine learning community [16], ordered by decreasing degree of smoothness.

In the next section, the MUR is briefly reviewed, as well as, the advantages of this method for the GPR are highlighted.

3 Matheron's update rule for GP regression

As said in the introduction, the MUR presented in this section has first appeared in geostatistics [9]. Let us briefly recall this method. Suppose \mathbf{X}_1 and \mathbf{X}_2 jointly Gaussian random variables. Then, the random vector \mathbf{X}_1 conditional on $\mathbf{X}_2 = \mathbf{x}_2$ can be expressed as

$$\{\mathbf{X}_1 | \mathbf{X}_2 = \mathbf{x}_2\} \stackrel{d}{=} \mathbf{X}_1 + \Sigma_{\mathbf{X}_1, \mathbf{X}_2} \Sigma_{\mathbf{X}_2, \mathbf{X}_2}^{-1} (\mathbf{x}_2 - \mathbf{X}_2),$$

where $\Sigma_{\mathbf{X}_1, \mathbf{X}_2} = \text{Cov}(\mathbf{X}_1, \mathbf{X}_2)$ is the covariance between \mathbf{X}_1 and \mathbf{X}_2 . As mentioned in [17], a key difference with the direct approach is that we now sample before conditioning, rather than after which is the key of the main idea developed in the present paper.

In the following proposition, we give the well-known MUR result for GPR.

Proposition 1 (MUR for GPR)

Using previous notations, $(Y(x))$ is a zero-mean GP with covariance function k . Then, Y conditioned on the observations $\{Y(\mathbb{X}) = \mathbf{y}\}$ can be expressed as follows:

$$\{Y | Y(\mathbb{X}) = \mathbf{y}\}(\cdot) \stackrel{d}{=} \underbrace{Y(\cdot)}_{\text{prior}} + \underbrace{k(\cdot, \mathbb{X})^\top (k(\mathbb{X}, \mathbb{X}) + \sigma_{\text{noise}}^2 \mathbf{I}_n)^{-1} (\mathbf{y} - Y(\mathbb{X}))}_{\text{update}}, \quad (4)$$

where $\mathbf{y} - Y(\mathbb{X})$ represents the residual.

The sampling scheme of the posterior distribution $\{Y | Y(\mathbb{X}) = \mathbf{y}\}(\cdot)$ using the MUR is given in the following algorithm.

Algorithm 1: Sampling scheme by MUR of $\{Y | Y(\mathbb{X}) = \mathbf{y}\}(\cdot)$.

- sample $Y(\cdot) \sim \mathcal{GP}(0, k)$;
 - return $Y(\cdot) + k(\cdot, \mathbb{X})^\top (k(\mathbb{X}, \mathbb{X}) + \sigma_{\text{noise}}^2 \mathbf{I}_n)^{-1} (\mathbf{y} - Y(\mathbb{X}))$.
-

Corollary 1 Suppose Y is simulated with Algorithm 1, then it is distributed as $\{Y | Y(\mathbb{X}) = \mathbf{y}\}$.

Proof The proof is a simple consequence of Proposition 1.

Let us give some remarks:

- By the MUR, we sample from the prior (first step in Algorithm 1) which is an advantage especially when the corresponding unconstrained (precision) matrix is diagonal or low-rank.
- By the MUR, the stationary property is preserved in the sampling procedure unlike the direct approach where the posterior covariance matrix must be computed which is not stationary anymore, cf. Sect. 4 below.
- The MUR provides numerical stability compared to standard approaches. This is because after sampling the prior, we directly map it on the set of observations, cf. Fig. 2 below.
- As seen in Proposition 1, the MUR can be used in the GPR framework when the data are observed with or without noise.

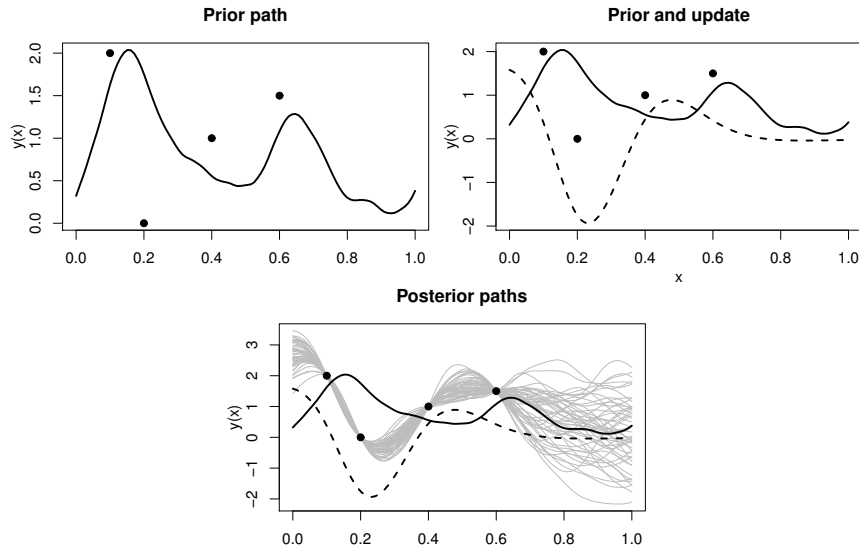


Fig. 1 Visual guide of the MUR. *Top left*: one path^x of unconstrained Gaussian prior together with the observations (black dots). *Top right*: the corresponding update path from (4) has been added (black dashed-curve). *Bottom*: the posterior paths (gray solid curves) obtained by adding the prior and the update as in (4).

In Fig. 1, the visual guide of the MUR is presented. The simple case where the observations are noise-free is considered. The Matérn covariance function with regularity parameter $\nu = 5/2$ and length parameter $\theta = 0.2$ (cf. Table 1) has been used. *Top left*: the prior (black solid curve) zero-mean GP $Y \sim \mathcal{GP}(0, k)$ together with the observations (black dots) have been shown. Let us mention that the prior does not interpolate the observations. *Top right*: the update part (black dashed-curve) of the MUR (4) has been added. As expected, the update part follows the trend of the observations. *Bottom*: the posterior paths (gray solid curves) have been illustrated by

adding the prior to the update as in (4). Each posterior sample path is obtained by adding a prior sample path to the update one. The posterior sample paths verify the interpolation conditions. One sample path of the prior and of the update is shown for clarity. However, fifty sample paths of the posterior distribution are shown in the bottom panel of Fig. 1.

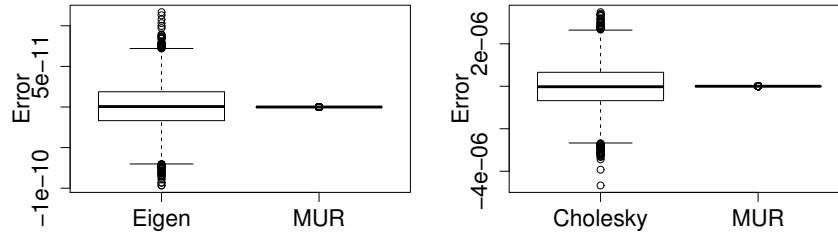


Fig. 2 The numerical error of the residual $Y(X) - \mathbf{f}$ using different approaches. For the Cholesky approach, a nugget effect of order 10^{-12} has been added in order to avoid numerical problems.

The stability of the MUR is now investigated. In Fig. 2, the boxplot of the numerical residual error $Y(\mathcal{X}) - \mathbf{f}$ using the MUR and different standard approaches are shown for ten thousand replicates. Let us mention that for the Cholesky factorization, a nugget effect of order 10^{-12} has been added in order to avoid numerical problems. As expected, the MUR provides accurate results. Additionally, based on numerical experiments, the MUR outperforms eigendecomposition and Cholesky factorization in terms of stability. This is because in the second step of the MUR algorithm, we map directly into the set of observations.

When the input domain \mathcal{D} is large and a fine distretization is used, simulating the prior (first step in Algorithm 1) using standard approaches (eigendecomposition and Cholesky factorization) becomes numerically heavy. The computational complexity grows cubically with the dimension of the associated Gaussian vector [6]. To sidestep this problem, we show in the next section how the prior GP can be simulated efficiently using the KLE update.

4 Karhunen-Loève expansion update

Let us first briefly recall the standard KLE [10]. In this section, the random field $(Y(x))_{x \in \mathcal{D}}$ is assumed stationary. Without loss of generality, the input domain \mathcal{D} is supposed to be the interval $[0, 1]$.

4.1 Standard Karhunen-Loève expansion

According to previous notations, $(Y(x))_{x \in \mathcal{D}}$ is a zero-mean Gaussian random field, whose stationary covariance function is $k(|x - x'|)$, for any $x, x' \in \mathcal{D}$

$$k(|x - x'|) = \text{Cov}(Y(x), Y(x')) = \mathbb{E}[Y(x)Y(x')].$$

The eigendecomposition of the covariance function on the domain \mathcal{D} is:

$$\int_0^1 k(|x - x'|)\varphi_i(x)dx = \lambda_i\varphi_i(x'), \quad \forall i, \forall x, x' \in \mathcal{D}.$$

The deterministic functions $\varphi_i(\cdot)$ and the coefficients λ_i are respectively the eigenfunctions and eigen-values of the covariance function $k(|\cdot|)$ on the domain \mathcal{D} . Let us recall that the eigen-values are real and non-negative since the covariance is symmetric and positive semi-definite:

$$\int_0^1 \int_0^1 k(|x - x'|)g(x)g(x')dx dx' \geq 0$$

for any g having finite L^2 norm on $[0, 1]$. Let us recall also that the eigen-functions $\varphi_i(\cdot)$ form a complete orthonormal basis functions set [15]. This means that

$$\int_0^1 \varphi_i(x)\varphi_j(x)dx = \delta_{ij},$$

where δ_{ij} represents the Kronecker delta (equal one if $i = j$ and zero otherwise).

By the KLE, the Gaussian random process $(Y(x))$ can be written as:

$$Y(x) = \sum_{i=1}^{+\infty} \sqrt{\lambda_i}\varphi_i(x)\zeta_i, \quad \forall x \in \mathcal{D} \quad (5)$$

where the KLE coefficients $(\zeta_i)_i$ are zero-mean uncorrelated Gaussian random variables (independent) with unit variance. The KLE coefficients are defined as the projection of the Gaussian random process onto the KLE eigen-functions:

$$\begin{aligned} \zeta_i &= \frac{1}{\sqrt{\lambda_i}} \int_0^1 \varphi_i(x)Y(x)dx, \\ \mathbb{E}[\zeta_i] &= 0 \quad \text{and} \quad \mathbb{E}[\zeta_i\zeta_j] = \delta_{ij}. \end{aligned} \quad (6)$$

The KLE is based on an approximation of the Gaussian random process $(Y(x))$ by a truncated sum of p terms on the domain \mathcal{D} :

$$Y(x) \approx \sum_{i=1}^p \sqrt{\lambda_i}\varphi_i(x)\zeta_i =: Y^p(x), \quad \forall x \in \mathcal{D}.$$

This approximation with a finite number of terms is called *truncated KLE*. Let us finish this section by recalling that the mean-square truncation error ϵ is related to the sum of the eigen-values:

$$\epsilon_{\text{KL}}^2 = \frac{\mathbb{E} \left[\int_{\mathcal{D}} (Y(x) - Y^p(x))^2 dx \right]}{\mathbb{E} \left[\int_{\mathcal{D}} Y(x)^2 dx \right]} = 1 - \frac{\sum_{i=1}^p \lambda_i}{\sum_{i=1}^{+\infty} \lambda_i}. \quad (7)$$

This mean-square truncation error decreases when the number of terms retained in the expansion increases.

When the domain \mathcal{D} is discretized into N equally spaced points, the eigendecomposition leads to a $N \times N$ eigen-value problem. When the domain is huge and a fine discretization is used, the eigendecomposition becomes expensive with computational complexity of order $\mathcal{O}(N^3)$. The idea in the following sections is to split the input domain \mathcal{D} in smallest nonoverlapping subdomains. The KLE coefficients are conditioned in order to follow the given correlation structure. By the stationary property of the Gaussian random field, we show that only the eigendecomposition of the first subdomain is needed. The parallelization of this approach is studied which drastically reduces the computational complexity.

4.2 Large scale Karhunen-Loève expansion

In this section, the Gaussian random field $(Y(x))_{x \in \mathcal{D}}$ is assumed stationary. The approach developed in [12, 13] is considered. For simplicity, the domain $\mathcal{D} = [0, MS]$ is split in M equal sized subdomains $\mathcal{D}^{(m)} = ((m-1)S, mS]$ for any block parameter $m \in \{1, \dots, M\}$. The extension to subdomains with different lengths has been investigated (cf. Sect. 2.2 in [12]). Let us recall this method. For any $m \in \{1, \dots, M\}$ and $x \in \mathcal{D}^{(m)}$, let

$$Y_m(x) := \sum_{i=1}^{+\infty} \sqrt{\gamma_i} \phi_i(x - (m-1)S) \zeta_i^{(m)} \quad (8)$$

be the standard KLE covering the m^{th} subdomain $\mathcal{D}^{(m)}$, where the deterministic functions $\phi_i(\cdot)$ and the coefficients γ_i are respectively the eigen-functions and eigen-values of the covariance function $k(|\cdot|)$ on the first subdomain $\mathcal{D}^{(1)} = [0, S]$. This means that only the eigendecomposition of the first subdomain is needed in the construction of the proposed approach. Before showing how the conditional coefficients are constructed, let us give the following notation:

$$(Y_m \cup Y_{m'}) (x) = Y_m(x) \mathbb{1}_{\mathcal{D}^{(m)}}(x) + Y_{m'}(x) \mathbb{1}_{\mathcal{D}^{(m')}}(x),$$

for any $x \in \mathcal{D}$ and $m, m' \in \{1, \dots, M\}$, where $\mathbb{1}_{\mathcal{D}^{(m)}}$ is the indicator function equal 1 if $x \in \mathcal{D}^{(m)}$ and zero otherwise. In practice, we suppose that p terms are retained in the expansion:

$$Y_m^p(x) = \sum_{i=1}^p \sqrt{\gamma_i} \phi_i(x - (m-1)S) \zeta_i^{(m)}, \quad \forall x \in \mathcal{D}^{(m)}. \quad (9)$$

Let us recall the following result proved in [12].

Proposition 2 (Distribution on blocks)

Under the stationary property of the Gaussian random process Y , we suppose that the m^{th} conditional coefficients set $\zeta^{(m)}$ is computed as follows

$$\zeta^{(m)} = \mathbf{K}^\top \zeta^{(m-1)} + \mathbf{L} \xi^{(m)}, \quad \forall m \in \{2, \dots, M\}, \quad (10)$$

where \mathbf{K} and \mathbf{L} are defined as

$$\begin{aligned} \mathbf{K}_{i,j} &:= \frac{1}{\sqrt{\gamma_i \gamma_j}} \int_0^S \int_0^S k(|x - x' - S|) \phi_i(x) \phi_j(x') dx dx'; \\ \mathbf{L} \mathbf{L}^\top &= \mathbf{I}_p - \mathbf{K}^\top \mathbf{K}, \end{aligned} \quad (11)$$

with \mathbf{I}_p the $p \times p$ identity matrix. Then

- the two processes Y and $(Y_{m-1} \cup Y_m)$ have the same distribution on $\mathcal{D}^{(m-1)} \cup \mathcal{D}^{(m)}$, for any $m \in \{2, \dots, M\}$.
- For any $m' \geq m$

$$\begin{cases} \mathbf{K}^{m,m'} := \text{Cov}(\zeta^{(m)}, \zeta^{(m')}) = \mathbf{K}^{m'-m}, \\ \mathbf{K}^{m,m} = \text{Cov}(\zeta^{(m)}) = \mathbf{I}_p, \end{cases} \quad (12)$$

where \mathbf{K}^0 is the identity matrix and \mathbf{K} is the coupling matrix defined in Eq. (11).

Let us give some comments on these results: the coupling matrix \mathbf{K} ensures the equality of distribution between the original GP random field Y and the approximated one Y_m at any two connected subdomains $\mathcal{D}^{(m-1)} \cup \mathcal{D}^{(m)}$. From the stationary of the GP random field, the matrix \mathbf{K} is computed once for any arbitrary number of subdomains.

In Fig. 3, we illustrate the method presented in this section for sampling the prior (left panel) and we applied it to the MUR to get the target posterior distribution (right panel). The same parameters used in Fig. 1 are also used in this figure. The black dots represent the observations. In the left panel of Fig. 3, the domain \mathcal{D} is split in three subdomains, i.e. $M = 3$. The solid curves represent the sample paths of the prior before conditioning which are uncorrelated. The dashed curves represent those after conditioning. Let us mention that the dashed curves follow the given correlation structure. In the right panel, the black solid curve represents the prior

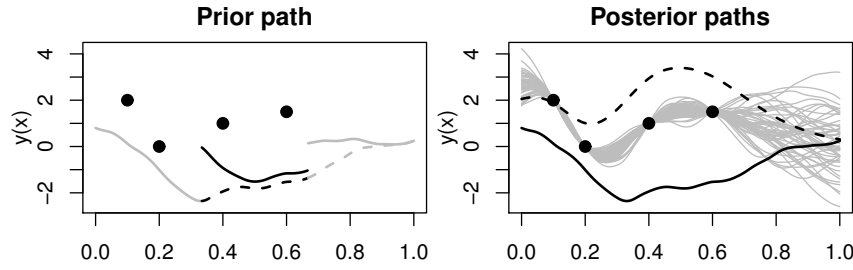


Fig. 3 *Left:* GP sample paths \hat{x} prior where the domain is split in three subdomains. Solid curves (before) and dashed-curves (after) conditioning. *Right:* the posterior paths (gray solid curves) obtained using the MUR as in (4) by adding the prior (black solid curve) to the update (black dashed curve).

sample path obtained in the left panel. However, the black dashed curve represents the corresponding update sample path as in (4). As in Fig. 1, the update sample path follows the trend of the observations. The posterior sample paths (gray solid curves) have been obtained by applying the MUR to the prior as in (4). As expected, they verify the interpolation condition.

In the next section, the parallelization of the proposed approach is developed and the advantages are highlighted. Additionally, the mean-square global *block* error is computed.

4.3 Parallel computing large scale KLE

In this section, the parallelization of the technique presented in Sect. 4.2 developed in [13] is investigated. This section presents the main result of the paper where the error due to the approach is provided and the efficiency for some special cases is highlighted. Without loss of generality, suppose that the domain \mathcal{D} is split into M odd subdomains. The case when M is even is discussed at the end of this section, cf. Remark 1. The same model developed in the previous section (9) will be used in this section

$$Y_m^p(x) = \sum_{i=1}^p \sqrt{\gamma_i} \phi_i(x - (m-1)S) \zeta_i^{(m)}, \quad \forall x \in \mathcal{D}^{(m)}.$$

However, we will show in this section, how the conditional KLE coefficients sets $\{\zeta^{(m)}\}$ can be generated using a parallelization technique. First, we sample independently M sets of KLE coefficients $\xi^{(1)}, \dots, \xi^{(M)}$. The KLE coefficients corresponding to an even part m is conditioned by the parts at the left and at the right as follows:

$$\zeta^{(m)} = \mathbf{K}^\top \xi^{(m-1)} + \mathbf{K} \xi^{(m+1)} + \mathbf{H} \xi^{(m)}, \quad (13)$$

where $m \in \{2, 4, \dots, (M-1)\}$ and \mathbf{H} is the lower triangular matrix such that

$$\mathbf{I}_p - \mathbf{K}^\top \mathbf{K} - \mathbf{K} \mathbf{K}^\top = \mathbf{H} \mathbf{H}^\top. \quad (14)$$

From equation (13), one can deduce that the coefficients sets $\{\zeta^{(m)}\}$ can be generated in parallel. To summarize, first, the prior KLE coefficients sets $\xi^{(1)}, \dots, \xi^{(M)}$ are generated in parallel. Second, the $M/2$ conditional coefficients $\zeta^{(m)}$ are computed in parallel too. By this strategy, the conditional prior sample paths are generated by parallelization. This leads to improve significantly the performance of the proposed approach in terms of average time of sampling.

Proposition 3 According to Eqs. (13) and (14),

- the two processes Y and $(Y_{m-1} \cup Y_m)$ have the same distribution on $\mathcal{D}^{(m-1)} \cup \mathcal{D}^{(m)}$, for any $m \in \{2, 4, \dots, (M-1)\}$.
- Additionally, we have the following results:

$$\begin{aligned} \text{Cov} \left(\zeta^{(m)} \right) &= \mathbf{I}_p, \quad \forall m \in \{2, 4, \dots, (M-1)\}; \\ \text{Cov} \left(\zeta^{(m)}, \zeta^{(m+2)} \right) &= \mathbf{K}^2 \quad \text{and} \quad \text{Cov} \left(\zeta^{(m)}, \zeta^{(m')} \right) = \mathbf{0}; \end{aligned}$$

for any $m \in \{2, 4, \dots, (M-3)\}$ and any $m' > m+2$ an even number.

Proof

- From Eq. (13), the correlation between $\xi^{(m-1)}$ and $\zeta^{(m)}$ is imposed by construction in order to satisfy the given correlation structure between the connected subdomains $\mathcal{D}^{(m-1)} \cup \mathcal{D}^{(m)}$

$$\text{Cov} \left(\xi^{(m-1)}, \zeta^{(m)} \right) = \mathbf{K} \quad \text{and} \quad \text{Cov} \left(\zeta^{(m)}, \xi^{(m+1)} \right) = \mathbf{K}.$$

For instance, when $m = 2$, we get

$$\begin{aligned} \text{Cov}(\xi^{(1)}, \zeta^{(2)}) &= \text{Cov}(\xi^{(1)}, \mathbf{K}^\top \xi^{(1)} + \mathbf{K} \xi^{(3)} + \mathbf{H} \xi^{(2)}) = \mathbf{K}; \\ \text{Cov}(\zeta^{(2)}, \xi^{(3)}) &= \text{Cov}(\mathbf{K}^\top \xi^{(1)} + \mathbf{K} \xi^{(3)} + \mathbf{H} \xi^{(2)}, \xi^{(3)}) = \mathbf{K}. \end{aligned}$$

Proposition 2 in [12] concludes the proof of the first item.

- According to Eqs. (13) and (14), we have for any $m \in \{2, 4, \dots, (M-1)\}$

$$\text{Cov} \left(\zeta^{(m)} \right) = \mathbf{K}^\top \mathbf{K} + \mathbf{K} \mathbf{K}^\top + \mathbf{H} \mathbf{H}^\top = \mathbf{I}_p.$$

Additionally, $\text{Cov} \left(\zeta^{(m)}, \zeta^{(m+2)} \right) = \mathbf{K}^2$. In particular, when $m = 2$, we get

$$\text{Cov}(\zeta^{(2)}, \zeta^{(4)}) = \text{Cov}(\mathbf{K} \xi^{(3)}, \mathbf{K}^\top \xi^{(3)}) = \mathbf{K}^2.$$

This concludes the proof of the second item and thus of the proposition.

Let us give some comments on these results:

- At each two connected subdomains, the proposed approach and the original random process have the same distribution.
- By the parallelization technique, the KLE coefficients corresponding to the odd parts are not conditioned. They are uncorrelated. This is because they are generated independently. Consequently, only $(M - 1)/2$ coefficients sets are conditioned. By this technique, the computational complexity of the sampling procedure is drastically reduced. However, as in Sect. 4.2, the eigendecomposition of the first subdomain to compute the coupling matrix \mathbf{K} is required.
- As we can see in Fig. 4, the domain $\mathcal{D} = [0, 1]$ is split in $M = 3$ subdomains. However, just one dashed conditioned part has been used to follow the given correlation structure.
- The parallelization technique works well in the low correlation structure cases. This is because the KLE sets corresponding to the odd part are uncorrelated by construction.

Remark 1 When the number of the subdomains M is even, only the KLE coefficients set corresponding to the last subdomain $m = M$ is conditioned from the left-hand side (as in Sect. 4.2). In that case, the number of conditional KLE coefficients sets is equal to $M/2$.

In the following corollary, the correlation between blocks is computed. This leads to compare the approximated correlation obtained by the proposed model and the true correlation function.

Corollary 2 (Correlation between blocks) For any $(x, x') \in \mathcal{D}^{(m)} \times \mathcal{D}^{(m+1)}$

$$\text{Cov}(Y_m^p(x), Y_{m+1}^p(x')) = \sum_{i,j=1}^p \sqrt{\gamma_i \gamma_j} \phi_i(x - (m-1)S) \phi_j(x' - mS) \mathbf{K}_{ij}, \quad (15)$$

for any $m \in \{1, 2, \dots, M\}$. However, for any $m \in \{2, 4, \dots, M-3\}$ and any $(x, x') \in \mathcal{D}^{(m)} \times \mathcal{D}^{(m+2)}$, we have

$$\text{Cov}(Y_m^p(x), Y_{m+2}^p(x')) = \sum_{i,j=1}^p \sqrt{\gamma_i \gamma_j} \phi_i(x - (m-1)S) \phi_j(x' - (m+1)S) (\mathbf{K}^2)_{ij}.$$

The correlation is equal to zero for the other situations.

Proof The proof is a simple consequence of Proposition 3.

In the following proposition, the mean-square global *block* error is computed. This error will be used to compare the proposed approach with versus without parallelization for different type of covariance functions.

Proposition 4 (Mean-square global block error)

In the setting of Proposition 3, we have the following global block error:

$$\epsilon_{B,M}^2 = \frac{\mathbb{E} \left[\int_{\mathcal{D}} (Y(x) - Y_{1:M}(x))^2 dG(x) \right]}{\mathbb{E} \left[\int_{\mathcal{D}} Y(x)^2 dG(x) \right]} = \frac{\text{Trace}((S_Y - S_{Y_{1:M}})(S_Y - S_{Y_{1:M}})^\top)}{\text{Trace}(S_Y S_Y^\top)} \quad (16)$$

where $Y_{1:M} = \cup_{m=1}^M Y_m$ and $(S_Y; S_{Y_{1:M}})$ are the Cholesky matrices of the covariance functions of Y and $Y_{1:M}$ on the grid $\mathcal{G} = \{x_1, \dots, x_N\}$ respectively and G is the cumulative distribution function (CDF) of the Uniform discrete random variable on \mathcal{G} .

Proof We know from [11] that any zero-mean Gaussian vector can be written as

$$\begin{pmatrix} Y(x_1) \\ \vdots \\ Y(x_N) \end{pmatrix} = S_Y \times \epsilon \quad \text{and} \quad \begin{pmatrix} Y_{1:M}(x_1) \\ \vdots \\ Y_{1:M}(x_N) \end{pmatrix} = S_{Y_{1:M}} \times \epsilon,$$

where ϵ is a N -dimensional standard Gaussian vector chosen the same for Y and $Y_{1:M}$ to get a specific dependence structure. The two matrices S_Y and $S_{Y_{1:M}}$ are the Cholesky factorization of the covariance of Y and $Y_{1:M}$ on the grid $\mathcal{G} = \{x_1, \dots, x_N\}$ respectively. If we denote by C the covariance matrix of the Gaussian vector $(Y(x_1), \dots, Y(x_N))^\top$, then $S_Y S_Y^\top = C$. Thus,

$$\mathbb{E} \left[\sum_{i=1}^N Y(x_i)^2 \right] = \mathbb{E}[\epsilon^\top S_Y^\top S_Y \epsilon] = \mathbb{E}[\epsilon^\top C \epsilon] = \text{Trace}(C) = \text{Trace}(S_Y S_Y^\top).$$

The result holds by following the same way.

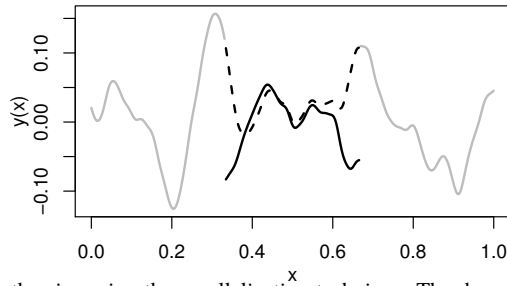


Fig. 4 GP sample path prior using the parallelization technique. The domain \mathcal{D} is split in three subdomains. The solid curves (resp. dashed curves) represent the paths before (resp. after) conditioning.

Figure 4 shows one GP sample path prior using the Matérn covariance function with the regularity parameter $\nu = 5/2$ by applying the parallelization technique

described in this section. The correlation length θ is fixed at 0.05. The domain \mathcal{D} is split in three subdomains. However, only one conditional KLE sets (gray dashed curve) is needed to follow the correlation structure in the entire domain. The solid curves (resp. dashed curves) represent the GP sample paths before (resp. after) conditioning. The black dashed curve follows the given correlation on the left and on the right hand sides as expected.

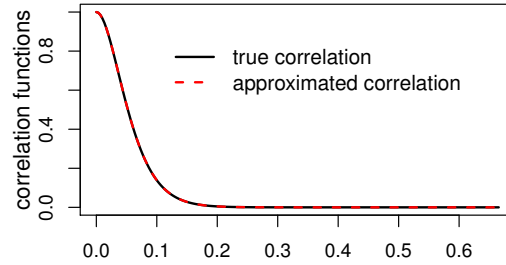


Fig. 5 The correlation functions between $t = 0$ and $t \in \mathcal{D}^{(2)}$ when the domain is split into three subdomains. The approximated correlation using the proposed approach is computed from Eq. (15). The true correlation is the Matérn 5/2 function.

Figure 5 shows the correlation functions between $t = 0$ and any $t \in \mathcal{D}^{(2)}$ when the domain \mathcal{D} is split into three subdomains. The black solid curve represents the true Matérn covariance function with regularity parameter $\nu = 5/2$ and length parameter $\theta = 0.05$. However, the red dashed curve represents the proposed correlation function obtained from Eq. (15). Let us mention that the correlation between the first and the last subdomain is zero. In that case, the root-mean-square error (RMSE) between the true correlation function and the proposed one on the entire domain is of order 3.8×10^{-12} . This is an expected result since the length parameter θ has been chosen small which leads to a correlation function decreasing rapidly to zero. This is a suitable situation for the parallelization technique developed in this section where the correlation between unconnected odd subdomains is zero by construction.

Remark 2 The parallelization technique developed in this section can be applied to the family of covariance functions with compact support (cf. Sect. 4.2 in [16]). As said in [16], ‘compact support means that the covariance between points become exactly zero when their distance exceeds a certain threshold’. This is an interesting class of covariance function which is suitable for the parallelization technique investigated in this section, since the correlation between unconnected odd parts is exactly zero by construction, cf. Fig. 6 below. However, the error due to the approximation still exists.

Example: compact support

In this example, the triangle correlation function (corresponding to the class of covariance functions with compact support) is used

$$k(|h|) = \max\left(1 - \frac{|h|}{\theta}; 0\right), \quad (17)$$

where θ is the correlation length parameter. In Fig. 6, θ is fixed at 0.3. This means that $k(|h|)$ is equal to zero for any $h \geq 0.3$. As before, the domain \mathcal{D} is split into three subdomains. The black solid curve represents the true triangle correlation function on \mathcal{D} . However, the red dashed curve represents the approximation correlation function using the proposed approach with the parallelization technique developed in this section. Let us mention that the root-mean-square error on \mathcal{D} between these two functions is of order 3.6×10^{-16} .

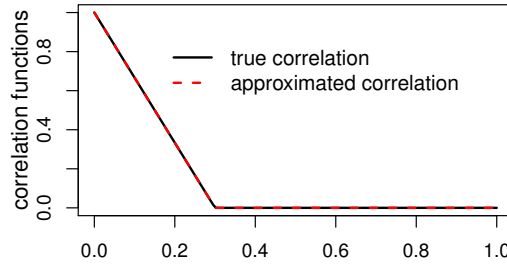


Fig. 6 The true correlation function and the approximated one on \mathcal{D} are shown together. The true one corresponding to the triangle correlation function in (17). The RMSE is equal to 3.6×10^{-16} .

Table 2 The mean-square global *block* error for different type of covariance functions using the proposed approach with and without parallelization. The domain \mathcal{D} is split into $M = 3$ subdomains.

covariance function	Mean-square global <i>block</i> error	
	without parallelization	with parallelization
Triangle, $\theta = 0.3$	3.82×10^{-2}	1.79×10^{-27}
Triangle, $\theta = 0.05$	1.49×10^{-5}	2.69×10^{-28}
Matérn 5/2, $\theta = 0.05$	2.99×10^{-24}	3.95×10^{-7}
Matérn 3/2, $\theta = 0.05$	6.42×10^{-27}	1.18×10^{-7}

Table 2 shows the mean-square global *block* error defined in (16) for different type of covariance functions using the proposed approach with and without parallelization. Based on the numerical experiments, the parallelization technique outperforms the case without parallelization when the covariance function with compact support

is used. This is because the correlation between odd subdomains is equal to zero by construction which is not the case for the classical approach without parallelization. However, for the Matérn family of covariance function, the parallelization has no advantage over the proposed approach without parallelization. However, let us mention that for example with the Matérn covariance function with regularity parameter $\nu = 5/2$, if the length correlation parameter θ is equal to 0.03 (resp. 0.01), we get a mean-square global *block* error of order 9.27×10^{-17} (resp. 1.81×10^{-28}) with the parallelization technique. So, it provides an accurate result. This is because, when the length parameter θ tends to zero, the Matérn covariance function decreases rapidly to zero. In that case, it can be considered to be close to a correlation function with compact support.

5 Computational illustrations

In this section the performance of the proposed approach is investigated. The problem of simulating a N -dimensional MVN vector $\eta \sim \mathcal{N}(\boldsymbol{\mu}, \boldsymbol{\Gamma})$ truncated on the intersection of a set of $n < N$ hyperplanes, denoted

$$\eta \sim \mathcal{N}_T(\boldsymbol{\mu}, \boldsymbol{\Gamma}), \quad T = \{\boldsymbol{x} \in \mathbb{R}^N \mid \boldsymbol{A}\boldsymbol{x} = \boldsymbol{y}\},$$

where $\boldsymbol{A} \in \mathbb{R}^{n \times N}$, $\boldsymbol{y} \in \mathbb{R}^n$ and $\text{rank}(\boldsymbol{A}) = n$ is considered. This problem is called hyperplane-truncated MVN distribution [3, 11]. The unconditional covariance matrix $\boldsymbol{\Gamma}$ is generated using the Matérn $\nu = 5/2$ covariance function. In that case, the covariance matrix admits no special structure. The elements of \boldsymbol{y} and \boldsymbol{A} are generated from a standard normal distribution $\mathcal{N}(0, 1)$.

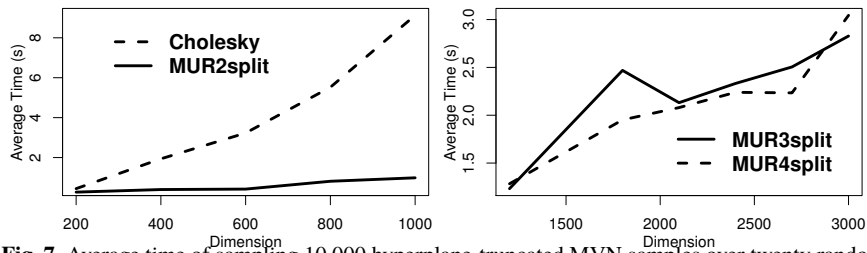


Fig. 7 Average time of sampling 10,000 hyperplane-truncated MVN samples over twenty random trials, when the number of data dimension N increases and of observations is fixed at $n = 10$.

In Fig. 7, the computation time of sampling 10,000 hyperplane-truncated MVN distributions averaged over twenty random trials is shown. The number of observations is fixed at $n = 10$ and the data dimension (i.e., dimension of η) N increases. *Left panel*: the Cholesky factorization has been compared to the proposed approach when the domain is split in only two subdomains. It is clear that the proposed ap-

proach outperforms the Cholesky factorization. Contrarily to Cholesky factorization, the computational time of the proposed approach grows linearly with the dimension of the vector η . *Right panel*: the black solid curve (resp. dashed curve) represents the average time in second of simulation 10,000 hyperplane-truncated MVN distribution using the proposed approach when the domain is split in three subdomains (resp. four subdomains). The average time is slightly different between these two approaches.

6 Conclusion

In this paper, a new methodology to sample large scale Gaussian process regression is developed. This problem is related to the simulation of high dimensional Gaussian vector truncated on a set of intersection of hyperplanes. The main idea is to combine both Matheron's update rule (MUR) and Karhunen Loève expansion (KLE). First, by the MUR we sample the target distribution without computing the posterior covariance matrix. Second, by splitting the input domain in smallest subdomains, the KLE coefficients are conditioned in order to follow the correlation structure on the entire domain. The parallelization of this approach has been developed. The mean-square global *block* error has been computed as well. The advantage of the parallelization technique has been highlighted through numerical examples and an illustration of the hyperplanes-truncated multivariate normal distribution problem is included. Based on the numerical experiments, the parallelization technique is especially efficient when using the class of covariance functions with compact support.

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