Scalable Gaussian Processes, with Guarantees: Kernel Approximations and Deep Feature Extraction

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Abstract

We provide approximation guarantees for a linear-time inferential framework for Gaussian processes, using two low-rank kernel approximations based on random Fourier features and truncation of Mercer expansions. In particular, we bound the Kullback–Leibler divergence between the idealized Gaussian process and the one resulting from a low-rank approximation to its kernel. Additionally, we present strong evidence that these two approximations, enhanced by an initial automatic feature extraction through deep neural networks, outperform a broad range of state-of-the-art methods in terms of time efficiency, negative log-predictive density, and root mean squared error.

1 Introduction

Gaussian Processes (GPs) have long been studied in probability and statistics; e.g. \cite{36}. In Bayesian inference, they provide a canonical way to define a probability distribution over functions, which can be used as a prior to build probabilistic frameworks for quantifying uncertainty in prediction. Among many applications, they have been a method of choice for hyperparameter tuning in deep learning.

In the simplest setting, a zero-mean probability distribution over functions $f : \mathbf{x} \mapsto y$ is defined as follows. For any collection $X = (\mathbf{x}_1, \ldots, \mathbf{x}_N)$ of feature vectors, it is assumed that their corresponding responses $y = (y_1, \ldots, y_N)$ are jointly Gaussian, with zero mean and covariance matrix $K(k_{\theta}, X) := (k_{\theta}(\mathbf{x}_i, \mathbf{x}_j))_{ij}$, where $k_{\theta}(\cdot, \cdot)$ is a positive semidefinite kernel indexed by a parameter vector $\theta$. 

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The usual inferential practice is to assume that we do not observe the Gaussian sample directly but additional noise drawn from a zero-mean isotropic Gaussian distribution is added to it prior to our observation. Bayesian inference then proceeds by focusing on estimation of $\theta$ and the noise variance distribution as well as to compute predictive distributions of unobserved responses $y^*$ corresponding to a collection of new feature vectors $X^*$ of interest. These inference tasks require computing the inverse and determinant of the covariance matrix $K(\theta, X)$, which naively costs $O(N^3)$ operations (or more precisely matrix multiplication time), making the inferential framework hard to scale computationally beyond a few thousand observations.

The challenge of scaling up GPs is a well-recognized and well-explored challenge in the GP literature. We provide theoretical grounding for the use of low-rank approximations to the covariance matrix $K(\theta, X)$, a widely used framework that improves the running time of GP regression from cubic to linear at a cost of performing approximate rather than exact inference. In particular, we provide bounds for the Kullback–Leibler divergence between the idealized $K(\theta, X)$ and low-rank approximations based on random Fourier features and Mercer expansions that can become smaller than a desired $\varepsilon N$ for moderate values of the rank of $K(\theta, X)$. These results immediately provide bounds for the corresponding Kullback–Leibler divergence between the idealized and approximate low rank predictive densities of unobserved responses. Guided by this theoretical understanding, we further propose how to build an implementation architecture to obtain quite substantial improvement in terms of both precision and speed against state-of-the-art methods. To achieve this, we adopt tools such as feature discovery through Deep Neural Networks (DNN), a semi-stochastic asynchronous gradient descent optimisation and an appropriate standardisation of the DNN output in the case of Mercer expansion-based low-rank approximations which enables nice analytical expressions for the eigenfunctions and eigenvalues.

At a high level, we propose to maximize the likelihood of our data $y$ conditioning on $X$, under the assumption that $y = (y_i)_{i=1}^N$ is sampled from a Gaussian distribution with covariance matrix that can be written as $\Sigma + \sigma^2 I_N$, with $\Sigma$ being a low-rank matrix that $\varepsilon$-approximates an ideal kernel matrix $K(\theta, Z) = (k_{\theta}(z_i, z_j))_{ij}$, computed on embedded feature vectors $Z = (z_i = g_w(x_i))_{i}$ obtained by passing the original feature vectors $x_i$ through a DNN $g_w$. A key ingredient in our inference framework is that we estimate jointly the parameters $w$ and $\theta$.

Our main implementation insights are (i) that inference in GPs whose kernel matrix has constant rank can be performed in linear time, avoiding the need for $N \times N$ matrix inversions/determinant computations for the purposes of maximum likelihood estimation/prediction; and (ii) while (i) can be used on its own, whenever the kernel is low-rank, we can also combine (i) with feature extraction using DNNs and low-rank kernel approximation methods into an end-to-end differentiable framework, which similarly does not require $N \times N$ matrix inversion/determinant computation in every step of the back-propagation, or to perform a prediction once the model is trained. The end-to-end differentiability
The rest of the paper proceeds as follows. Section 2 provides our theoretical grounding for the low-rank approximations together with an implementation architecture based on DNNs. In particular, in Sections 2.1.2 and 2.1.3 we detail two instantiations of our framework resulting from two different low-rank kernel approximations based on random Fourier features (Section 2.1.2) and truncating the Mercer expansion of the kernel (Section 2.1.3). In both cases, we provide bounds in the KL divergence between the idealized and the approximate data generation process resulting from kernel approximation. In Section 3 we provide a literature review and in Section 4 we test extensively our proposed methodology against many state-of-the-art methods, in a broad collection of real-world datasets.

2 Methodology

2.1 Linear-time inference in Gaussian processes with low-rank kernels

In GP regression, we are given a response vector $y = (y_i)_{i=1}^N \in \mathbb{R}^N$ whose entries are noisy evaluations of some random function $f(\cdot)$ on a collection of $D$-dimensional feature vectors $X = (x_i)_{i=1}^N \in \mathbb{R}^{N \times D}$, i.e. $y_i$ is a noisy observation of $f(x_i)$. We take the noise, $y_i - f(x_i)$, for each data entry $i$ to be independent Gaussian with mean 0 and variance $\sigma^2$. Moreover, we place a GP prior over $f(\cdot)$, with zero mean and kernel $k_0(\cdot, \cdot)$, so that the collection of function values $f(X) := (f(x_i))_{i=1}^N$ has a joint Gaussian distribution with zero mean and covariance matrix $K(k_0, X)$. Setting $A = K(k_0, X) + \sigma^2 I_N$, the log-marginal likelihood of the data becomes $\log p(y|X) = -\frac{1}{2}y^T A^{-1} y - \frac{1}{2} \log |A| - \frac{N}{2} \log (2\pi)$. Future observations $y^* \in \mathbb{R}^{N^*}$ corresponding to feature vectors $X^* = (x_i^*)_{i=1}^{N^*}$ have a conditional normal distribution with mean and variance given by $E(y^*|y) = K(k_0, X^*, X)A^{-1} y$ and $\text{Var}(y^*|y) = K(k_0, X^*, X^*) A^{-1} K(k_0, X^*, X) A^{-1}$. As such, we get that the covariance matrix of the data $y$ is $A = \Xi \Xi^T + \sigma^2 I_N$. We can then use the Woodbury matrix inversion

\footnote{We commonly view a collection $X = (x_i)_{i=1}^N$ of vectors as a matrix whose rows are the $x_i$'s. Similarly a collection $y = (y_i)_{i=1}^N$ of scalars is viewed as a column vector.}

and the light-weight computation required in each back-propagation step, allows for fast training of GP-based models that perform really well compared to state-of-the-art baselines in both prediction power and training/prediction time, as we show in Section 4. In particular, we have provided results for datasets up to 1.8 million points.
lemma and the Sylvester determinant theorem to obtain explicit forms for the inverse of $A$ and its determinant: $A^{-1} = \sigma^{-2}I_N - \sigma^{-2}\Xi(\sigma^2 I_r + \Xi^\top\Xi)^{-1}\Xi^\top$, and $|A| = \sigma^{2(N-r)}|\sigma^2 I_r + \Xi^\top\Xi|$. Since these identities involve inversion or determinant calculations of $r \times r$ matrices, by plugging them into the expressions for the log-marginal likelihood of observations $y$ and the mean and variance of the predictive density of future observations $y^*$, we can, with the right ordering of operations, compute the log-likelihood and the predictive density in $O(r^3 + r^2N)$ time, i.e. linear in $N$, when $r$ is a constant.

### 2.1.1 Approximation guarantees

We define a distribution over functions mapping feature vectors $x \in \mathbb{R}^D$ to responses $y$ through a random function $h : x \mapsto y$ which is sampled from a GP with noisy observations, as described in Section 2.1. In particular, $f(\cdot)$ is sampled from a GP with mean zero and kernel function $k_\theta : \mathbb{R}^D \times \mathbb{R}^D \rightarrow \mathbb{R}$, and then $y \sim \mathcal{N}(f(x), \sigma^2)$. Thus, a collection $X = (x_i)_{i=1}^N$ of feature vectors maps to a collection of responses $y = (y_i)_{i=1}^N$ sampled as

$$y \sim \mathcal{N}(0, K(k_\theta, X) + \sigma^2 I_N).$$

A well-studied topic in mathematics, statistics, and machine learning is approximating kernels with low-rank kernels. Given a kernel function $k_\theta$, the goal is to identify a feature map $\phi_{\theta,\varepsilon} : \mathbb{R}^D \rightarrow \mathbb{R}^r$, providing a guarantee of the following form for a collection $X = (x_i)_{i=1}^N$ of feature vectors in $\mathbb{R}^D$:

$$K(k_\theta, X) \approx \varepsilon \Sigma(\phi_{\theta,\varepsilon}, X),$$

where $\Sigma(\phi_{\theta,\varepsilon}, X) = (\phi_{\theta,\varepsilon}(x_i)^\top \phi_{\theta,\varepsilon}(x_j))_{ij}$. Equivalently $\Sigma(\phi_{\theta,\varepsilon}, X)$ can be written as $\Sigma = \Xi\Xi^\top$, where $\Xi$ is a $N \times r$ matrix whose rows are $\phi_{\theta,\varepsilon}(x_i)$ for $i = 1, \ldots, N$. In particular, $\Sigma$ is a rank $r$ matrix. Coming back to the setting of Eq. (1), a collection $X = (x_i)_{i=1}^N$ of feature vectors maps to a collection of responses $y = (y_i)_{i=1}^N$, which are approximately sampled as follows:

$$y \sim \mathcal{N}(0, \Sigma + \sigma^2 I_N).$$

We now show that (3) can be made to approximate (1) in a precise sense. First, we provide the following general result which will allow us to bound the KL divergence between (1) and (3), whenever we instantiate the latter with some low-rank kernel $\Sigma$:

**Theorem 1** (Proof in the supplementary material). *Suppose that $\Sigma_1$ and $\Sigma_2$ are $N \times N$ positive definite (symmetric) matrices, such that $(1 + \gamma)\Sigma_1 - \Sigma_2$ is positive semi-definite for some $\gamma \geq 0$. Then*

$$\text{KL} \left[ \mathcal{N}(0, \Sigma_1) \ || \ \mathcal{N}(0, \Sigma_2) \right] \leq \frac{1}{2} \text{Tr}(\Sigma_2^{-1/2}(\Sigma_1 - (1 - \gamma)\Sigma_2)\Sigma_2^{-1/2}).$$

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If additionally $\Sigma_2 \succeq (1 + \gamma)^{-1} \Sigma_1$, then we obtain
\[
\text{KL}[\mathcal{N}(0, \Sigma_1) \mid \mid \mathcal{N}(0, \Sigma_2)] \leq \gamma N. \tag{5}
\]

If $\Sigma_1 = \sigma^2 I_N + K_1$ and $\Sigma_2 = \sigma^2 I_N + K_2$, where $K_1$ and $K_2$ are positive semi-definite, $\sigma^2 > 0$, and $(1 + \gamma)\Sigma_1 - \Sigma_2$ is positive semi-definite, then
\[
\text{KL}[\mathcal{N}(0, \Sigma_1) \mid \mid \mathcal{N}(0, \Sigma_2)] \leq \frac{1}{2\sigma^2} \text{Tr}(K_1 - (1 - \gamma)K_2 + \gamma \sigma^2 I_N). \tag{6}
\]

Let us instantiate Theorem 1 by taking $K_1 = K(k_\theta, X)$ and $K_2 = \Sigma(\phi_{\theta, \phi}, X)$ (which has rank $r$). $K_1$ determines the idealized data generation process of [4], while $K_2$ determines the approximate one of [3]. Our theorem states that the KL divergence between these two processes is controlled by (1)–(6), which as we will see in the next sections can become smaller than any desired $\varepsilon N$ for relatively modest values of the rank $r$, namely poly-logarithmic in $N$ (Theorem 3), or even an absolute constant (Theorem 5), whenever the dimension $D$ is an absolute constant.

2.1.2 Instantiation No. 1: Random Fourier Gaussian Processes

A well-studied method for obtaining low-rank kernel approximations is by defining a parametrized family of functions $e_\eta : \mathbb{R}^D \to \mathbb{R}$ as well as a distribution $p(\eta)$ over $\eta$, defining the feature map $\phi(\mathbf{x}) = (e_{\eta_1}(\mathbf{x}), \ldots, e_{\eta_p}(\mathbf{x}))$ by sampling random $\eta_1, \ldots, \eta_p \sim p(\eta)$. For example, in a celebrated paper [34], Bochner’s theorem for shift-invariant kernels $k_\theta$ is used to define a kernel-specific density $p_\theta(\eta)$ such that $e_\eta(\cdot)$ is a cosine function with frequency and phase determined by $\eta \sim p_\theta(\eta)$ (derived from a random Fourier feature with spectral frequency $\eta$; see also [2]). The guarantees obtained by [34] for random Fourier features bound the pointwise distance between $k_\theta(\mathbf{x}_i, \mathbf{x}_j)$ and $\phi(\mathbf{x}_i)^\top \phi(\mathbf{x}_j)$, for arbitrary $\mathbf{x}_i, \mathbf{x}_j$. To be able to bound the KL divergence between [4] and [3] we need a spectral, rather than an entry-wise, approximation of $\sigma^2 I + K(k_\theta, X)$ by $\sigma^2 I + \Sigma(\phi, X)$. These types of results can be obtained as well, as exemplified by the following:

**Theorem 2** (Theorem 12 of [2]). Consider the $D$-dimensional Gaussian kernel $k(\mathbf{x}, \mathbf{x}') = \exp(-2\pi^2 ||\mathbf{x} - \mathbf{x}'||_2^2)$, and the kernel matrix $K = K(k, X) = (k(\mathbf{x}_i, \mathbf{x}_j))_{ij}$, where $X = (\mathbf{x}_1, \ldots, \mathbf{x}_N)$ is a collection of points in $\mathbb{R}^D$ such that, for some $R > 0$, $||\mathbf{x}_i - \mathbf{x}_j||_{\infty} \leq R$, $\forall i,j$. Suppose $D \leq 5\log(N/\sigma^2) + 1$ and $\varepsilon \in (0, 1)$. There exists (a samplable in $O(D)$ time) distribution $p(\eta)$ and a parametrized family $e_\eta(\cdot)$ of modified Fourier Features such that, if $r \geq \Omega(\sqrt{\frac{D}{\varepsilon^2}} \log N)\frac{\log(ans(K))}{\delta}$, where $s_{\sigma^2}(K) = \text{Tr}((\sigma^2 I + K)^{-1} K)$ and $\delta \in (0, 1)$, then the feature map $\phi(\mathbf{x}) = (e_{\eta_1}(\mathbf{x}), \ldots, e_{\eta_p}(\mathbf{x}))$ where $\eta_1, \ldots, \eta_r \sim p(\eta)$ satis-
fies the following with probability at least $1 - \delta$:

$$
(1 - \epsilon)(\sigma^2 I_N + K) \preceq (\sigma^2 I_N + \Sigma) \preceq (1 + \epsilon)(\sigma^2 I_N + K),
$$

(7)

where $\Sigma = (\phi(x_i)^\top \phi(x_j))_{ij}$, and $\preceq$ denotes semi-definite domination.

Using Theorems 1 and 2 we get the following theorem. We state it for the Gaussian kernel with the same fixed scaling in every direction for notational simplicity. It extends to the general Gaussian kernel with different scaling per direction in an obvious way (rescaling coordinates).

**Theorem 3** (Proof in the supplementary material). Consider the setting of Theorem 2, with the same kernel $k(\cdot, \cdot)$, matrix $K$, dataset $X$, radius $R$, constraint $D \leq 5 \log(N/\sigma^2) + 1$, and the same distribution $p(\eta)$ and parametric family $e_{\eta}(\cdot)$ of modified Fourier Features used in that theorem. Take $\epsilon \in (0, \frac{1}{2}]$. If we take $r \geq \Omega(\frac{R^2}{\epsilon^2}(\log \frac{N}{\sigma^2})^2 \log(\frac{N}{\delta}))$ random $\eta_1, \ldots, \eta_r \sim p(\eta)$ and define the rank $r$ matrix $\Sigma$ as in Theorem 2 then with probability at least $1 - \delta$, the KL divergence from distribution (3) to distribution (1) is at most $\epsilon N$.

### 2.1.3 Instantiation No. 2: Mercer Gaussian Processes

In this section we present an alternative approach for obtaining low-rank approximations to the kernel $K(k_\theta, X)$, namely truncating the Mercer expansion of the kernel [32]. Suppose that $k_\theta$ is a Mercer kernel on some probability space $\mathcal{X} \subseteq \mathbb{R}^D$ with probability measure $\mu$, which means that $k_\theta(\cdot, \cdot)$ can be written as:

$$
k_\theta(x, x') = \sum_{t=1}^{\infty} \lambda_t e_t(x)e_t(x'),
$$

(8)
where \((\lambda_t)_{t \in \mathbb{N}}\) is a sequence of summable non-negative, non-increasing numbers, i.e. eigenvalues, and \((\epsilon_t)_{t \in \mathbb{N}}\) is a family of mutually orthogonal unit-norm functions with respect to the inner product \(\langle f, g \rangle = \int_X f(x)g(x)d\mu(x)\), defined by \(\mu\), i.e. eigenfunctions. Now suppose that \(X = (x_i)_{i=1}^N\) is a collection of vectors \(x_i \in X\). It follows from Eq. (8) that the kernel matrix \(K(k_\theta, X)\) can be written:

\[
K(k_\theta, X) \equiv \sum_{t=1}^{\infty} \lambda_t \omega_t^T, \tag{9}
\]

where \(\omega_t = (\epsilon_t(x_1), \epsilon_t(x_2), \ldots, \epsilon_t(x_N))\), for all \(t \in \mathbb{N}\). Recall that the sequence \((\lambda_t)\) is summable so \(\lambda_t \to 0\) as \(t \to \infty\). The rate of convergence is very fast for many kernels. For example, we illustrate below the Mercer expansion of the multi-dimensional Gaussian kernel

\[
k_{\sigma^2, \Delta}(z_i, z_j) = \sigma^2 \exp(-\frac{1}{2}(z_i - z_j)^T \Delta (z_i - z_j)), \tag{10}
\]

where \(\Delta = \text{diag}(\epsilon_1^2, \ldots, \epsilon_D^2)\) contains the length scales along the \(D\) dimensions of the covariates, and \(\sigma^2\) is the variance. The parameters of the kernel are \(\theta = (\sigma^2, \Delta)\).

We view \(k_{\sigma^2, \Delta}(z_i, z_j)\) as a kernel over \(\mathbb{R}^D\) equipped with an axis aligned Gaussian measure \(\rho(z) = \rho(z^1, \ldots, z^D)\), whose density in dimension \(j\) is given by

\[
\rho_j(z^j) = \alpha_j \pi^{-1/2} e^{\exp(-\alpha_j^2 z^j)^2}, \quad \forall j = 1, \ldots, D. \tag{11}
\]

Mercer’s expansion theorem \cite{32} allows us to write

\[
k_{\sigma^2, \Delta}(z_i, z_j) = \sum_{n \in \mathbb{N}^D} \lambda_n e_n(z_i)e_n(z_j), \tag{12}
\]

where \((e_n)_{n \in \mathbb{N}^D}\) is an orthonormal basis of \(L_2(\mathbb{R}^D, \rho)\), wherein inner products are computed using \(\rho(z)\). It is well-known \cite{11, 12, 36, 49} that such an orthonormal basis \((e_n)_{n \in \mathbb{N}^D}\) can be constructed as a tensor product of the orthonormal bases of \(L_2(\mathbb{R}^{D_j}, \rho_j)\) for all \(j\), as follows. Setting \(\beta_j = (1 + (2\epsilon_j / \alpha_j)^2)^{1/4}\), \(\gamma_{n_j} = \beta_j^{-1/2} \Gamma(n_j)^{-1/2}\), and \(\delta_j = \alpha_j^2 (\beta_j^2 - 1)/2\) the orthonormal eigenvectors are defined as

\[
e_n(z) = \prod_{j=1}^{D} e_{n_j}(z^j) = \prod_{j=1}^{D} \left\{ \gamma_{n_j} \exp(-\delta_j^2 z^j) H_{n_j-1}(\alpha_j \beta_j z^j) \right\}, \tag{13}
\]

where \(H_n\) are the Hermite polynomials of degree \(n\) and the corresponding eigenvalues are
\[ \lambda_n = \sigma_j^2 \prod_{j=1}^{D} \lambda_{n_j} = \]
\[ \sigma_j^2 \prod_{j=1}^{D} \left\{ \left( \frac{\alpha_j^2}{\alpha_j^2 + \delta_j^2 + \epsilon_j^2} \right)^{1/2} \left( \frac{\epsilon_j^2}{\alpha_j^2 + \delta_j^2 + \epsilon_j^2} \right)^{n_j-1} \right\}. \] (14)

Note that \( \lambda_{n_j} \to 0 \) as \( n_j \to \infty \). Indeed, as long as \( \alpha_j^2/\epsilon_j^2 \) is bounded away from 0, this decay is exponentially fast. This motivates approximating \( K(k_0, X) \) by keeping the first few terms of (9), as motivated by the following theorem of [4].

**Theorem 4** (Proof of Theorem 4 in [4]). Let \( k(\cdot, \cdot) \) be a Mercer kernel on probability space \( (X, \mu) \) with \( k(x, x) \leq B \), for all \( x \in X \). Let \( X = (x_1, \ldots, x_N) \) comprise samples from \( \mu \), let \( K = K(k, X) \) (which satisfies (9)), and let \( \Sigma = \sum_{t=1}^{r} \lambda_t \omega_t \omega_t^\top \), for some \( r \in \mathbb{N} \) (which has rank \( r \)). With probability at least \( 1 - \delta \) over the samples \( X \):

\[ \text{Tr}(K - \Sigma) \leq N \cdot \left( \Lambda_{>r} + \sqrt{\frac{BA_{>r}}{N\delta}} \right), \]  

(15)

where \( \Lambda_{>r} = \sum_{t>r} \lambda_t \).

Using Theorems [1] and [4] we get the following theorem.

**Theorem 5** (Proof in the supplementary material). Consider the setting of Theorem [4]. Under event (15) which occurs with probability at least \( 1 - \delta \), the KL divergence from distribution (3) to distribution (1) is at most

\[ \frac{N}{2\sigma^2} \cdot \left( \Lambda_{>r} + \sqrt{\frac{BA_{>r}}{N\delta}} \right). \]  

(16)

For example, suppose \( k(x, x') = \exp(-2\pi^2 \lVert x - x' \rVert_2^2) \) is the multi-variate Gaussian kernel over \( \mathbb{R}^D \), endowed with a Gaussian density \( \mu(x) = (2\pi)^{-D/2} \exp(-\pi^2 \lVert x \rVert_2^2) \).

Then choosing \( r = (D \log D + \log \frac{1}{\varepsilon \delta})^{1/2} \) makes (16) at most \( \varepsilon N \).

### 2.2 Predictive densities

It is straightforward to establish bounds on the Kullback–Leibler divergence between idealized, based on \( K(k_0, X) \), and approximating via a low-rank approximation, based on \( \Sigma \), predictive densities of unobserved observations \( y^* \) given new features \( X^* \). Suppose that these Gaussian densities are denoted by \( P(y; y^*) \) and \( Q(y; y^*) \) respectively. Our Theorems 3 and 5 establish bounds on \( \text{KL}[P || Q] \). By the chain rule of KL divergence we obtain that

\[ \text{KL}[P || Q] = \text{KL}[P(y) || Q(y)] + \text{KL}[P(y^* | y) || Q(y^* | y)]. \]
Because of the non-negativity of Kullback–Leibler divergence, a bound on $\text{KL}[P || Q]$ implies a bound on $\text{KL}[P(y^*|y) || Q(y^*|y)]$, so in expectation, with respect to $y \sim P$, the predictive densities of the true and the approximating distributions on the test data are close.

### 2.3 DNNs for feature extraction

We describe here an implementation architecture that exploits our theoretical guarantees with the enhancement of feature extraction through a DNN. Instead of defining a direct mapping from $x \in \mathbb{R}^D$ to $y$ through a GP, we define a composition of a random function with a deterministic function as follows. First, a deterministic function $g_w : x \mapsto z$ embeds a feature vector $x$ to a feature vector $z \in \mathbb{R}^d$; we assume that $g_w$ is parametric, e.g. expressible by a DNN. Next, a random function $h : z \mapsto y$ is sampled from a GP with noisy observations exactly as described in Section 2.1.1, so

$$f(\cdot) \sim \mathcal{N}(0, K_{\theta}(\cdot) + \sigma^2 I_N),$$

where $Z = (g_w(x_i) \equiv z_i)^N_{i=1}$. Clearly, by taking the neural network to be trivial (i.e. the identity function) we obtain the setting of the Section 2.1.1. The goal now is to identify a feature map $\phi_{\theta, \epsilon} : \mathbb{R}^d \mapsto \mathbb{R}^r$, providing a guarantee of the form

$$K(\theta, Z) \approx \epsilon \Sigma(\phi_{\theta, \epsilon}, Z),$$

where $\Sigma(\phi_{\theta, \epsilon}, Z) = (\phi_{\theta, \epsilon}(z_i)^\top \phi_{\theta, \epsilon}(z_j))_{i,j}$. With this DNN enhancement, using Random Fourier Features [34], Modified Random Fourier Features [2], or other random feature-based methods to obtain a low-rank approximation to the kernel $K(\theta, Z)$ gives rise to our family of Deep Fourier Gaussian Processes (DFGP). Using Mercer approximations gives rise to our family of Deep Mercer Gaussian Processes (DMGP).

### 3 Related work

The computational burden of cubic (or more accurately matrix multiplication) time complexity of GP inference has motivated a voluminous literature on faster approximate methods over the last decades; see [30] for a recent survey. Most of these methods rely on the notion of inducing inputs either on the actual Gaussian process domain [18, 33, 38, 39], or the spectral domain [14, 17, 27]. There is also a plethora of works pursuing kernel matrix approximations, either by using the Nyström method [13, 15, 21, 26, 29, 37, 41, 47, 48], or by approximating the kernel function [8, 16, 23, 28, 31, 35]. More flexible GPs that mimic Bayesian hierarchical formulations have been introduced by [8] and further combined with random Fourier features in [7] to obtain a scalable inferential framework.
A similar theoretical investigation to ours has been given by [5] for the sparse variational GP regression framework [19, 39]. They have showed that under certain conditions, such as normality of the input vectors $\mathbf{x}$ and use of a squared exponential kernel, the total number of inducing points needed to approach the posterior distribution arbitrarily close with respect to KL divergence is $O(\log^D N)$. The number of inducing points can be seen as the equivalent of the rank $r$ of our kernel approximation since sparse variational GP regression makes use of the Nyström approximation to achieve inference in linear time over $N$.

There have also been attempts of combining kernels and neural networks, see [46], where activation functions and weight connections of a Bayesian neural network (BNN) are replaced by GPs, and [8], where multiple layers of stacked GPs are jointly used. Nonetheless, both of those methods suffer from scalability issues, too. A more recent method, proposed in [7], combines the Deep GP framework of [8] with random Fourier features [34], to obtain a highly-scalable model based on Cholesky-free optimization which leads to improved performance over standard approximation GP methods.

The idea of combining neural networks with GPs to extract more meaningful representations from high-dimensional data has been used by [6, 20, 22] but all approaches cannot scale to more than a few thousand training points. A more recent work [40] exploits convolutional architectures and GPs based on random Fourier features to tackle image classification problems. The work that is closest to ours is [45] in which scalability issues have been dealt with by exploiting Kronecker/Toeplitz algebra combined with the inducing inputs framework and simultaneous estimation of all parameters. While we propose a low-rank kernel approximation, [45] make use of the KISS-GP framework [43]. We perform extensive evaluations of our method against this and several other state-of-the-art scalable GP methods in the next section.

4 Experiments

4.1 Curve learning via low-rank kernel approximations

Supplement B.2 describes an illustrative small data example in which random Fourier features GP (FGP) and Mercer GP (MGP), which can be seen as DFGP of Section 2.1.2 and DMGP of Section 2.1.3 equipped with a trivial neural network $g_w(\mathbf{x}) = \mathbf{x}$, are compared against exact Gaussian process regression. Mercer GP achieves identical results; random Fourier features GP tends to provide better point estimates with tighter posterior regions. Robustness to the number of eigenfunctions/spectral frequencies is also illustrated.
We provide implementation details on how we implement DMGP and DFGP where \( \xi \) has to be pre-fixed or learnt from the data. We choose to keep it fixed with its spectral density. Regarding DFGP, we follow the implementation based in algorithm 1 of [34], input to the GP. We measure and we standardize the outputs of DNN, matrix using a Gaussian kernel. In both cases, the crux is to compute the low-rank standard deviations in parentheses. No results are reported for DMGP for \( d = 3, \sqrt{r} = 32 \) since computational tractability breaks for these values.

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<td>( \xi )</td>
<td>Negative log-predictive density DMGP</td>
</tr>
<tr>
<td>( d = 1 )</td>
<td>( d = 2 )</td>
</tr>
<tr>
<td>( \sqrt{r} )</td>
<td>Training Time DMGP</td>
</tr>
<tr>
<td>( 2 )</td>
<td>115(4)</td>
</tr>
<tr>
<td>( 4 )</td>
<td>112(1)</td>
</tr>
<tr>
<td>( 8 )</td>
<td>114(1)</td>
</tr>
<tr>
<td>( 16 )</td>
<td>116(5)</td>
</tr>
<tr>
<td>( 32 )</td>
<td>117(1)</td>
</tr>
<tr>
<td>( 64 )</td>
<td>122(1)</td>
</tr>
</tbody>
</table>

### 4.2 Implementation details for DMGP and DFGP

We provide implementation details on how we implement DMGP and DFGP using a Gaussian kernel. In both cases, the crux is to compute the low-rank matrix \( \Sigma \) for a fixed rank \( r \). For DMGP, we compute \( \Sigma \) by using \( \sqrt{r} \in \mathbb{N} \) eigenfunctions/eigenvalues per dimension for the Mercer expansion in [12]:

\[
\Sigma = \sum_{n \in \mathbb{N}^d, n \leq (\sqrt{r}, \ldots, \sqrt{r})} \lambda_n \xi_n \xi_n^T,
\]

where \( \xi_n = [\epsilon_n(z_1), \ldots, \epsilon_n(z_N)]^T \in \mathbb{R}^N \). Note that the parameter \( \xi_j \) in [11] has to be pre-fixed or learnt from the data. We choose to keep it fixed with its value being set \( 1/\sqrt{2} \) which corresponds to a standard \( d \)-dimensional Gaussian measure and we standardize the outputs of DNN, \( Z \), before we feed it as an input to the GP.

Regarding DFGP, we follow the implementation based in algorithm 1 of [34], where we first sample, for even number \( r, \sqrt{r} \) spectral frequencies \( \eta_1, \ldots, \eta_{\frac{r}{2}} \) from the spectral density \( p(\eta) \) of the stationary kernel \( k_0(\cdot, \cdot) \) and then create the
feature map \( \phi(z) : \mathbb{R}^d \to \mathbb{R}^r \), defined by the vector

\[
\sqrt{\frac{2}{r}} [\cos(\eta_1^\top z), \ldots, \cos(\eta_r^\top z), \sin(\eta_1^\top z), \ldots, \sin(\eta_r^\top z)]^\top.
\]

Hence, the rank of \( \Sigma \) is always an even number. The spectral frequencies are only sampled once before training and are then kept fixed throughout optimization of the log-marginal likelihood. Finally, the spectral density in the case of Gaussian kernel in [10] is given by

\[
p(\eta) = \sqrt{2\pi \Delta^{-1}} \sigma_f^{-2} \exp(-2\pi^2 \eta^\top \Delta^{-1} \eta).
\]

4.3 Real data experiments

We compare the following methods: (i) DMGP of Section 2.1.3 with \( d = 1 \) and \( r = 15 \); (ii) DFGP of Section 2.1.2 with \( d = 4 \), \( r = 40 \), and random Fourier features; (iii) Stochastic Variational Inference GP with 250 (SVIGP) and 500 (SVIGP+) inducing points [18] (code used from GPflow [31]); (iv) Sparse GP Regression [39] with 250 (SGPR) and 500 (SGPR+) inducing points (code used from GPflow); (v) Deep Kernel Learning with 5000 (DKL) and 10000 (DKL+) inducing points and \( d = 1 \) since we found that larger values of \( d \) did not improve performance (code used from [https://gpytorch.ai](https://gpytorch.ai) [45]); (vi) Deep GPs with random Fourier features (RFEDGP), see [7], with two hidden layers, three GPs per layer, and spectral frequencies being optimized variationally with fixed randomness—we used 20 Monte Carlo samples throughout training since we found it is much faster and as accurate as the training procedure followed by [7] and 100 Monte Carlo samples for prediction as in [7] (code used from [https://github.com/mauriziofilippone/deep_gp_random_features](https://github.com/mauriziofilippone/deep_gp_random_features)).

All data have been retrieved from UCI repository [9] or the official site of [36].

DMGP and DFGP require joint estimation of the parameters \( w \) and \( \theta \) through maximization of the log marginal likelihood which is a non-decomposable loss function, see [24], so we used the semi-stochastic asynchronous gradient descent suggested in [11]. More details about the practical implementation of DMGP and DFGP are discussed in Supplement 4.2. We emphasise that for maintaining fairness among comparisons, we kept hyperparameter tuning to the minimum for the DNN-based methods, by using, across all datasets, the same \([D - 512 - 256 - 64 - d]\) architecture with hyperbolic tangent activation functions, while the DNN weights of these methods were initialized by pre-training the DNN as suggested by [44, 45]. We ran all methods for 100 epochs using Adam optimizer [25] and mini-batch optimization with mini-batches of size 1000. All GPs used Gaussian kernels with separate length-scale per dimension. All results have been averaged over five random splits (90% train, 10% test).

Table 2 presents comparisons of all methods in terms of NLPD and training time, whereas Supplement 3.1 presents comparisons in terms of RMSE, which carry the same message. Both DFGP and DMGP clearly outperform all other methods in speed and NLPD performance. The last three rows of the two sub-tables of
Table 2: Negative log-predictive density and training time comparison (standard deviations reported in parentheses) on seven standard benchmark real-world datasets; \( N, N^* \) and \( D \) represent training data size, test data size, and feature dimension, respectively.

<table>
<thead>
<tr>
<th></th>
<th>Elevators</th>
<th>Protein</th>
<th>SARCOMA</th>
<th>IDRoad</th>
<th>Song</th>
<th>Movie</th>
<th>Electric</th>
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<td>14939</td>
<td>41157</td>
<td>44039</td>
<td>39136</td>
<td>463810</td>
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<td>1844352</td>
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<td>( N^* )</td>
<td>1600</td>
<td>4573</td>
<td>4894</td>
<td>43488</td>
<td>51535</td>
<td>58325</td>
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<tr>
<td>( D )</td>
<td>18</td>
<td>9</td>
<td>21</td>
<td>3</td>
<td>90</td>
<td>77</td>
<td>19</td>
</tr>
<tr>
<td>SVGP</td>
<td>0.44(0.021)</td>
<td>1.041(0.007)</td>
<td>-0.422(0.006)</td>
<td>0.652(0.008)</td>
<td>1.208(0.005)</td>
<td>0.087(0.006)</td>
<td>0.804(0.003)</td>
</tr>
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<td>SVGP+</td>
<td>0.435(0.018)</td>
<td>1.091(0.006)</td>
<td>-0.479(0.004)</td>
<td>0.541(0.008)</td>
<td>1.205(0.005)</td>
<td>0.078(0.005)</td>
<td>0.769(0.002)</td>
</tr>
<tr>
<td>SVGP+</td>
<td>0.433(0.017)</td>
<td>0.997(0.007)</td>
<td>-0.370(0.007)</td>
<td>0.799(0.007)</td>
<td>1.202(0.006)</td>
<td>0.216(0.005)</td>
<td>0.871(0.002)</td>
</tr>
<tr>
<td>SVGP+</td>
<td>0.420(0.017)</td>
<td>0.944(0.005)</td>
<td>-0.488(0.009)</td>
<td>0.737(0.011)</td>
<td>1.138(0.006)</td>
<td>0.186(0.004)</td>
<td>0.810(0.001)</td>
</tr>
<tr>
<td>SKL</td>
<td>0.527(0.011)</td>
<td>0.595(0.020)</td>
<td>0.395(0.040)</td>
<td>0.744(0.129)</td>
<td>1.261(0.057)</td>
<td>0.460(0.033)</td>
<td>0.447(0.013)</td>
</tr>
<tr>
<td>SKL+</td>
<td>0.536(0.011)</td>
<td>0.691(0.037)</td>
<td>0.430(0.034)</td>
<td>0.687(0.047)</td>
<td>1.315(0.158)</td>
<td>0.438(0.037)</td>
<td>0.448(0.012)</td>
</tr>
<tr>
<td>RFEDGP</td>
<td>0.434(0.021)</td>
<td>1.028(0.006)</td>
<td>-0.303(0.001)</td>
<td>0.583(0.009)</td>
<td>1.207(0.006)</td>
<td>0.238(0.032)</td>
<td>0.616(0.004)</td>
</tr>
<tr>
<td>DFGP</td>
<td>0.371(0.036)</td>
<td>0.875(0.015)</td>
<td>-0.777(0.015)</td>
<td>1.468(0.010)</td>
<td>1.185(0.004)</td>
<td>-0.008(0.022)</td>
<td>0.078(0.002)</td>
</tr>
<tr>
<td>DFGP</td>
<td>0.350(0.029)</td>
<td>0.853(0.018)</td>
<td>-0.777(0.020)</td>
<td>0.139(0.012)</td>
<td>1.189(0.005)</td>
<td>-0.016(0.002)</td>
<td>0.057(0.004)</td>
</tr>
</tbody>
</table>

Table [2] describe results of extra experiments in which a DNN regression model with RMSE as loss function was first trained on the data, then its fitted outputs \( Z \) were independently used as input to fit a Mercer GP (\( DNN+M \)), random Fourier features GP (\( DNN+F \))\(^2\) or simply an isotropic model \( Y \sim \mathcal{N}(Z, \sigma I_N) \) (\( DNN+S \)). These methods do not perform as well in terms of NLPD, emphasizing the necessity of our suggested joint parameter optimization. However, notice the improvement of the non-parametric \( DNN+M \) and \( DNN+F \) over the naive \( DNN+S \). We also applied an exact GP regression model (using GPflow) to the smallest dataset Elevators. The average NLPD (± one std.) was 0.377±0.024 with total average running time 53550 ± 2099 seconds. Comparing with the results of Table [2] we see that both DFGP and NLPD exhibited superior NLPD performance confirming the effectiveness of DNN feature engineering.

Figure [1] depicts how NLPD and training time over 100 epochs depend on the number of training points in the ELECTRIC dataset, illustrating that our methods can achieve equally good precision with less training points and less time. In particular, notice that DFGP scales better than DMPG. Table [3] presents the performance of DMGP and DFGP for a series of values of \( d_r \) for the smaller size datasets PROTEIN and SARCOMA. Similar results for ELEVATORS dataset

\(^2\)As discussed in Section [3] Mercer GP and random Fourier features GP are respectively DMGP and DFGP without the neural net.
can be found in Supplement B. There is evidence that large values of $d$ and $r$ offer only marginally better performance for both DMGP and DFGP, while severely affecting the training time for DMGP. This suggests using relatively small $d$ and $r$ for DMGP and slightly increase these values for DFGP. For all our data experiments we used $d = 1, r = 15$ for DMGP and $d = 4, r = 40$ for DFGP.

4.4 Summary of results

The extensive experiments of this section were designed to answer specific performance questions, the answers to which are summarized here. There is strong evidence that both instantiations of our framework, DFGP and DMGP described in Sections 2.1.2 and 2.1.3 respectively, (i) outperform all state-of-the-art baselines in both time efficiency and prediction accuracy measured in NLPD and RMSE (ii) outperform simple DNN regression without the use of a GP verifying the need for incorporating both our proposed ingredients (iii) achieve competitive performance and are much faster against the competitors with quite fewer training points (iv) outperform exact GP regression inference confirming the importance of the DNN feature extraction (v) illustrate the importance of our proposed joint parameter estimation framework since they clearly outperform consecutive estimation of the DNN first and the kernel parameters after. We also illustrate robustness with respect to $r$ and $d$ and provide practical guidelines.

Acknowledgements

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References


Appendix

A Omitted proofs

Proof of Theorem 1: We first show (4). Recall that the KL divergence between two Gaussians with non-singular covariances has a closed form expression:

$$\text{KL} \left[ \mathcal{N}(0, \Sigma_1) \mid \mid \mathcal{N}(0, \Sigma_2) \right] = \frac{1}{2} \left( \text{Tr}(\Sigma_2^{-1}\Sigma_1) - N + \ln \left| \Sigma_2 \right| \left| \Sigma_1 \right| \right). \quad (19)$$

Because $\Sigma_2$ is positive definite, $\Sigma_2^{-1}$ is too and it has a square root. Thus, by using properties of the trace we can write:

$$\text{Tr}(\Sigma_2^{-1}\Sigma_1) = \text{Tr}(\Sigma_2^{-1/2}\Sigma_1\Sigma_2^{-1/2})$$

$$= \text{Tr}(\Sigma_2^{-1/2}(\Sigma_1 - (1 - \gamma)\Sigma_2 + (1 - \gamma)\Sigma_2^{-1/2}))$$

$$= \text{Tr}(\Sigma_2^{-1/2}(\Sigma_1 - (1 - \gamma)\Sigma_2^{-1/2})) + \text{Tr}(\Sigma_2^{-1/2}(1 - \gamma)\Sigma_2^{-1/2})$$

$$= \text{Tr}(\Sigma_2^{-1/2}(\Sigma_1 - (1 - \gamma)\Sigma_2^{-1/2})) + (1 - \gamma)\text{Tr}(I_N)$$

$$= \text{Tr}(\Sigma_2^{-1/2}(\Sigma_1 - (1 - \gamma)\Sigma_2^{-1/2})) + (1 - \gamma)N$$

Plugging this into (19) yields:

$$\text{KL} \left[ \mathcal{N}(0, \Sigma_1) \mid \mid \mathcal{N}(0, \Sigma_2) \right] = \frac{1}{2} \left( \text{Tr}(\Sigma_2^{-1/2}(\Sigma_1 - (1 - \gamma)\Sigma_2^{-1/2})) - \gamma N + \ln \left| \Sigma_2 \right| \left| \Sigma_1 \right| \right). \quad (20)$$

Next we argue the following:

Lemma 6. If $A, B$ are positive definite, and $B - A$ is positive semidefinite, then

$$\ln \left( \frac{|A|}{|B|} \right) \leq 0.$$ 

Proof of Lemma 6: Let $\ell_1 \geq \ell_2 \geq \ldots \geq \ell_N > 0$ be the eigenvalues of $A$, and $\ell'_1 \geq \ell'_2 \geq \ldots \geq \ell'_N > 0$ be the eigenvalues of $B$, in non-increasing order. Because $B \succeq A$, by the min-max theorem we have $\ell_i \leq \ell'_i, \forall i$. Thus,

$$\frac{|A|}{|B|} = \prod_{i=1}^{N} \frac{\ell_i}{\ell'_i} \leq 1 \Rightarrow \ln \left( \frac{|A|}{|B|} \right) \leq 0.$$

Because $(1 + \gamma)\Sigma_1 \succeq \Sigma_2$, it follows from Lemma 6 that

$$0 \geq \ln \left( \left| \frac{\Sigma_2}{(1 + \gamma)\Sigma_1} \right| \right) = \ln \left( \frac{\left| \Sigma_2 \right|}{(1 + \gamma)^N \left| \Sigma_1 \right|} \right) = \ln \left( \frac{\left| \Sigma_2 \right|}{\left| \Sigma_1 \right|} \right) - N \ln(1 + \gamma) \geq \ln \left( \frac{\left| \Sigma_2 \right|}{\left| \Sigma_1 \right|} \right) - N \gamma.$$
Combining the last inequality with (20) yields Bound (4).

To prove (6), we note that if additionally \((1 + \gamma)\Sigma_2 \succeq \Sigma_1\) then:

\[
\frac{1}{2} \text{Tr}(\Sigma_2^{-1/2}((1 + \gamma)\Sigma_2 - \Sigma_1)\Sigma_2^{-1/2}) \geq 0. \tag{21}
\]

This follows by noticing that matrix \(\Sigma_2^{-1/2}((1 + \gamma)\Sigma_2 - \Sigma_1)\Sigma_2^{-1/2} \succeq 0\). Indeed, for all \(x \in \mathbb{R}^N\) and using that \((\Sigma_2^{-1/2})^T = \Sigma_2^{-1/2}\):

\[
x^T\Sigma_2^{-1/2}((1 + \gamma)\Sigma_2 - \Sigma_1)\Sigma_2^{-1/2}x = (\Sigma_2^{-1/2}x)^T((1 + \gamma)\Sigma_2 - \Sigma_1)(\Sigma_2^{-1/2}x) \geq 0,
\]

where the last inequality follows from the positive semidefiniteness of \((1 + \gamma)\Sigma_2 - \Sigma_1\).

Now combining (21) with (4) and using properties of the trace we get:

\[
\text{KL} [\mathcal{N}(0, \Sigma_2) \mid \mathcal{N}(0, \Sigma_2)] \\
\leq \frac{1}{2} \left( \text{Tr}(\Sigma_2^{-1/2}((1 - (1 - \gamma)\Sigma_2)\Sigma_2^{-1/2}) + \text{Tr}(\Sigma_2^{-1/2}((1 + \gamma)\Sigma_2 - \Sigma_1)\Sigma_2^{-1/2})) \right) \\
\leq \frac{1}{2} \left( \text{Tr}(\Sigma_2^{-1/2}(2\gamma\Sigma_2)\Sigma_2^{-1/2})) \right) \\
\leq \gamma \text{Tr}(I_N) = \gamma N.
\]

Let us now move to the proof of (6). We plug \(\Sigma_4 = \sigma^2 I_N + K_1\) and \(\Sigma_2 = \sigma^2 I_N + K_2\) into (6) to get:

\[
\text{KL} [\mathcal{N}(0, \Sigma_1) \mid \mathcal{N}(0, \Sigma_2)] \leq \frac{1}{2} \text{Tr}(\Sigma_2^{-1/2}(K_1 - (1 - \gamma)K_2 + \gamma\sigma^2 I_N)\Sigma_2^{-1/2}) \\
\leq \frac{1}{2} \text{Tr}(\Sigma_2^{-1}(K_1 - (1 - \gamma)K_2 + \gamma\sigma^2 I_N)) \tag{22}
\]

where we used properties of the trace. Because \(K_2\) is positive semidefinite, it has eigenvalues \(\ell_1 \geq \ell_2 \geq \ldots \geq \ell_N \geq 0\), which implies that \(\Sigma_2 = \sigma^2 I + K_2\) has eigenvalues \(\sigma^2 + \ell_1 \geq \sigma^2 + \ell_2 \geq \ldots \geq \sigma^2 + \ell_N > 0\), which in turn implies that \(\Sigma_2^{-1}\) has eigenvalues \((\sigma^2 + \ell N)^{-1} \geq (\sigma^2 + \ell_{N-1})^{-1} \geq \ldots \geq (\sigma^2 + \ell_1)^{-1} > 0\). Now using (22) and properties of the trace we have that:

\[
\text{KL} [\mathcal{N}(0, \Sigma_1) \mid \mathcal{N}(0, \Sigma_2)] \leq \frac{1}{2} \text{Tr}(\Sigma_2^{-1}(K_1 - (1 - \gamma)K_2 + \gamma\sigma^2 I_N)) \\
\leq \frac{1}{2} \lambda_{\text{max}}(\Sigma_2^{-1}) \text{Tr}(K_1 - (1 - \gamma)K_2 + \gamma\sigma^2 I_N) \\
= \frac{1}{2} \cdot \frac{1}{\sigma^2 + \ell_N} \cdot \text{Tr}(K_1 - (1 - \gamma)K_2 + \gamma\sigma^2 I_N) \\
\leq \frac{1}{2\sigma^2} \text{Tr}(K_1 - (1 - \gamma)K_2 + \gamma\sigma^2 I_N),
\]
where in the above derivation $\lambda_{\text{max}}(\Sigma_2^{-1})$ is the maximum eigenvalue of matrix $\Sigma_2^{-1}$.

**Proof of Theorem 3**: Set $\Sigma_1 = \sigma^2 I_N + K$ and $\Sigma_2 = \sigma^2 I_N + \Sigma$. Notice that $s_{\sigma^2}(K) = \text{Tr}((\sigma^2 I + K)^{-1} K) \leq \text{Tr}(I_N) \leq N$. Thus, given our choice of $r$, Theorem 2 implies that, with probability at least $1 - \delta$, $\Sigma_1$ and $\Sigma_2$ satisfy:

$$(1 - \varepsilon)\Sigma_1 \preceq \Sigma_2 \preceq (1 + \varepsilon)\Sigma_1.$$  

Given that for $\varepsilon \in (0, \frac{1}{2}]$, we get that $1 - \varepsilon \geq \frac{1}{1 + 2\varepsilon}$, the above implies that:

$$(1 + 2\varepsilon)^{-1}\Sigma_1 \preceq \Sigma_2 \preceq (1 + 2\varepsilon)\Sigma_1.$$  

Now we use (5) of Theorem 1 to get that the KL divergence from distribution (3) to distribution (1) is bounded by $2\varepsilon N$. □

**Proof of Theorem 5**: First, notice that, because $\Sigma$ is a truncation of $K$, $K - \Sigma$ is positive semidefinite. To prove (16), we set $K_1 = K$, $K_2 = \Sigma$, and use (6) from Theorem 2 with $\gamma = 0$ to get that the KL divergence from distribution (3) to distribution (1) is bounded by:

$$\frac{1}{2\sigma^2} \text{Tr}(K - \Sigma) \leq \frac{N}{2\sigma^2} \left( \Lambda_{>r} + \sqrt{B\Lambda_{>r} N \delta} \right).$$  

To prove the second part of the theorem, we use properties of the spectrum of Gaussian kernels, as discussed in Section 2.1.3. As per Equations (12), (13), (14), the eigenfunctions and eigenvalues of the Gaussian kernel can be indexed by vectors $n \in \mathbb{N}^d$. Moreover, the eigenvalues take the form $\lambda_n = c^D \lambda^T n$, for some absolute constants $c > 0$ and $\lambda \in (0, 1)$, with $I$ being a vector of all ones. In particular, the eigenvalues are ordered in terms of the “level sets” of $\lambda^T n$; namely the larger $\lambda^T n$ is, the smaller the eigenvalue is, while every $n$ with the same value of $\lambda^T n$ has the same eigenvalue, $\lambda_n \equiv c^D \lambda^T n$. For $m = \Omega(D \log D + \log \frac{1}{\varepsilon \delta})$,  

\[ 21 \]
let us take \( r = |\{ \mathbf{n} \in \mathbb{N}^D \mid \mathbf{1}^T \mathbf{n} < m \}| \). We have that

\[
\Lambda_{>r} = \sum_{\mathbf{n} : \mathbf{1}^T \mathbf{n} \geq m} \lambda_{\mathbf{n}} \\
= \sum_{\mathbf{n} : \mathbf{1}^T \mathbf{n} \geq m} c^D \lambda^{1^T \mathbf{n}} \\
\leq \sum_{\ell = m}^{\infty} \ell^D c^D \lambda^\ell \\
= c^D \sum_{\ell = m}^{\infty} (\ell^D \lambda^{\ell/2}) \lambda^{\ell/2} \\
\leq c^D \sum_{\ell = m}^{\infty} \lambda^{\ell/2} \\
\leq c^D \lambda^{m/2} \cdot \frac{1}{1 - \lambda},
\]

where the second to last inequality follows from the fact that \( \ell^D \lambda^{\ell/2} \leq 1 \) for \( m = \Omega(D \log D) \). To conclude the proof notice that the Gaussian kernel \( k(\mathbf{z}, \mathbf{z}') = \exp(-2\pi^2 ||\mathbf{z} - \mathbf{z}'||_2^2) \) satisfies \( k(\mathbf{z}, \mathbf{z}) = 1 \), hence we can use (16) with \( B = 1 \) to bound the KL divergence from distribution (3) to distribution (1) by

\[
\frac{N}{2\sigma^2} \cdot \left( \Lambda_{>r} + \sqrt{\frac{\Lambda_{>r}}{N\delta}} \right) \leq \varepsilon N,
\]

where the last inequality uses (23) and that \( m = \Omega(D \log D + \log \frac{1}{\varepsilon \sigma \delta}) \). Given that \( r = |\{ \mathbf{n} \in \mathbb{N}^D \mid \mathbf{1}^T \mathbf{n} < m \}| \), we get that to attain (24) it suffices to choose the rank to be \( r = (D \log D + \log \frac{1}{\varepsilon \sigma \delta}) \Omega(D) \).

\( \Box \)

## B Additional experimental results

### B.1 Extra results on real data

Table 3 demonstrates the RMSE values of all methods. As in Table 1 of the main paper, RMSE values follow similar trends as the corresponding NLPD values, with DMGP and DFGP outperforming all the baselines across all datasets.

Table 4 presents how rank \( r \) affects performance and training time over the Elevators dataset where results show similar patterns as in Table 1 of the main paper. Increasing embedding’s dimension \( d \) does not reduce any further NLPD for both DMGP and DFGP while computational time for DMGP increases fast with \( d \). Similarly, \( \sqrt{r} \) and \( d \) does not seem to provide any performance boost for values larger than 4 and 2, respectively.
Table 3: RMSE comparison between state-of-the-art baselines and our methods DMGP and DFGP. The experimental set-ups are the same as in Table 2 of the main paper.

<table>
<thead>
<tr>
<th>N</th>
<th>Elevators</th>
<th>Protein</th>
<th>Sarcom</th>
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<td>0.810(0.005)</td>
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<td>0.540(0.002)</td>
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<td>0.375(0.007)</td>
<td>0.649(0.005)</td>
<td>0.151(0.001)</td>
<td>0.413(0.004)</td>
<td>0.807(0.004)</td>
<td>0.270(0.003)</td>
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<td></td>
<td>SVGP</td>
<td>0.375(0.007)</td>
<td>0.653(0.005)</td>
<td>0.168(0.002)</td>
<td>0.537(0.004)</td>
<td>0.806(0.005)</td>
<td>0.315(0.003)</td>
</tr>
<tr>
<td></td>
<td>DGP</td>
<td>0.370(0.007)</td>
<td>0.620(0.004)</td>
<td>0.153(0.002)</td>
<td>0.596(0.006)</td>
<td>0.802(0.005)</td>
<td>0.308(0.003)</td>
</tr>
<tr>
<td></td>
<td>DKL</td>
<td>0.352(0.010)</td>
<td>0.630(0.012)</td>
<td>0.206(0.047)</td>
<td>0.499(0.074)</td>
<td>0.815(0.006)</td>
<td>0.274(0.014)</td>
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<td>DKL+</td>
<td>0.361(0.009)</td>
<td>0.632(0.022)</td>
<td>0.276(0.035)</td>
<td>0.474(0.024)</td>
<td>0.813(0.004)</td>
<td>0.268(0.014)</td>
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<tr>
<td></td>
<td>DFGP</td>
<td>0.346(0.010)</td>
<td>0.564(0.007)</td>
<td>0.111(0.002)</td>
<td>0.277(0.003)</td>
<td>0.791(0.003)</td>
<td>0.237(0.000)</td>
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</table>

Table 4: Comparative NLPD performance and training time (in seconds) of DMGP and DFGP on Elevators dataset for several values of rank r. No results are reported for DMGP for d = 3, √r = 32 since computational tractability breaks for these values. Experimental set-ups are the same as in Table 1 of the main paper.

<table>
<thead>
<tr>
<th>d</th>
<th>DMGP</th>
<th>DFGP</th>
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<tbody>
<tr>
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<td>NLPD</td>
<td></td>
</tr>
<tr>
<td>2</td>
<td>0.381(0.037)</td>
<td>0.361(0.032)</td>
</tr>
<tr>
<td>4</td>
<td>0.371(0.036)</td>
<td>0.351(0.032)</td>
</tr>
<tr>
<td>8</td>
<td>0.371(0.036)</td>
<td>0.351(0.032)</td>
</tr>
<tr>
<td>10</td>
<td>0.371(0.037)</td>
<td>0.351(0.032)</td>
</tr>
<tr>
<td>16</td>
<td>0.371(0.036)</td>
<td>0.351(0.032)</td>
</tr>
<tr>
<td>32</td>
<td>0.371(0.036)</td>
<td>0.351(0.032)</td>
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</tbody>
</table>

<table>
<thead>
<tr>
<th>d</th>
<th>Training Time</th>
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<tbody>
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<td>40(1)</td>
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<tr>
<td>4</td>
<td>41(1)</td>
</tr>
<tr>
<td>8</td>
<td>41(2)</td>
</tr>
<tr>
<td>10</td>
<td>41(2)</td>
</tr>
<tr>
<td>16</td>
<td>42(1)</td>
</tr>
<tr>
<td>32</td>
<td>44(2)</td>
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<table>
<thead>
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<th></th>
<th>Elevators</th>
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<tr>
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<td>NLPD</td>
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<tr>
<td>2</td>
<td>0.381(0.037)</td>
<td>0.381(0.040)</td>
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<td>4</td>
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<tr>
<td>32</td>
<td>0.371(0.036)</td>
<td>0.357(0.032)</td>
</tr>
</tbody>
</table>

23
B.2 Curve learning via low-rank kernel approximations

We examine the flexibility of our models by comparing them to exact Gaussian process regression models via the following simple example. We generate an artificial dataset based on the function $f(x) = \frac{1}{2} (3\sin(2x) + \cos(10x) + \frac{x}{4})$; exact Gaussian process models can easily recover such a smooth function and, therefore, they provide a sound baseline for comparison with our methods. Our simulated dataset has one-dimensional training points $\{x_i, f(x_i)\}_{i=1}^{25}$ where $x_i \sim \mathcal{N}(0, 1)$. We omit to include any DNN for our two methods, i.e. no embedding is being learnt, facilitating thus comparisons with exact Gaussian processes. We call those methods MGP and FGP since they only depend on Mercer (see Sections 4.2 from supplement and 2.1.3 from main paper) and random Fourier features frameworks (see 2.1.2 from main paper), respectively. For all three methods, a Gaussian kernel is used. The exact Gaussian process model has been trained using GPflow.

Figure 2 illustrates how MGP and FGP compare to exact Gaussian process. MGP presents identical behavior, leading to same posterior mean and predictive intervals. The posterior mean of FGP approximates better the underlying curve with more ‘confidence’ to unseen function values. We use $r = 34$ and $r = 68$ for MGP and FGP respectively.

Figure 3 depicts how MGP and FGP inference is affected by considering different values for $r$ (i.e. eigenfunctions or spectral frequencies) for approximating $\Sigma$ on the simulated dataset. For MGP, as $r$ increases, the uncertainty decreases and posterior mean estimates tend to approximate very well those of the exact GP model. FGP performs well with high confidence even with $r = 4$ and after $r = 24$ learns the true function impressively well.

C Code

All experiments were carried out on a Linux machine with 32 2.20GHz CPU cores and 64GB RAM. The implementation of our code is provided at https://github.com/aresPanos/dmgp_dfgp_regression.
Figure 2: Recovering the function $f(x) = \frac{1}{2} (3\sin(2x) + \cos(10x) + \frac{x}{2})$. From top to bottom: Predictive mean and 95% of the predictive probability mass of exact Gaussian process, MGP and FGP, respectively. We make use of 34 eigenfunctions for MGP and 34 spectral frequencies for FGP, i.e. $r = 34$ and $r = 68$, respectively. Black crosses depict the training data, solid red line shows $f(x)$, and the dashed yellow line shows the approximate methods MGP and FGP.
Figure 3: Recovering the function $f(x) = \frac{1}{2} (3 \sin(2x) + \cos(10x) + \frac{x}{2})$ by using different ranks $r$ for $\Sigma$ to approximate the true kernel $K$. Training points are denoted by black crosses, $f(x)$ by solid red line, MGP and FGP with dashed yellow lines.