Generic Inference in Latent Gaussian Process Models

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Abstract

We develop an automated variational method for inference in models with Gaussian
process (gp) priors and general likelihoods. The method supports multiple outputs and
multiple latent functions and does not require detailed knowledge of the conditional
likelihood, only needing its evaluation as a black-box function. Using a mixture of
Gaussians as the variational distribution, we show that the evidence lower bound and
its gradients can be estimated efficiently using empirical expectations over univariate
Gaussian distributions. Furthermore, the method is scalable to large datasets which
is achieved by using an augmented prior via the inducing-variable approach underpin-
ning most sparse gp approximations, along with parallel computation and stochastic
optimization. We evaluate our method with experiments on small datasets, medium-
scale datasets and a large dataset, showing its competitiveness under different likelihood
models and sparsity levels. Moreover, we analyze learning in our model under batch
and stochastic settings, and study the effect of optimizing the inducing inputs. Finally,
in the large-scale experiment, we investigate the problem of predicting airline delays
and show that our method is on par with the state-of-the-art hard-coded approach for
scalable gp regression.

1 Introduction

Developing fully automated methods for inference in complex probabilistic models has be-
come arguably one of the most exciting areas of research in machine learning, with notable
examples in the probabilistic programming community (see e.g. Hoffman and Gelman, 2014;
Wood et al., 2014; Goodman et al., 2008; Domingos et al., 2006). Indeed, while these prob-
abilistic programming systems allow for the formulation of very expressive and flexible
probabilistic models, building efficient inference methods for reasoning in such systems is a
significant challenge.

One particularly interesting setting is when the prior over the latent parameters of the
model can be well described with a Gaussian process (gp, Rasmussen and Williams, 2006).
Gp priors are a popular choice in practical non-parametric Bayesian modeling with perhaps
the most straightforward and well-known application being the standard regression model

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with Gaussian likelihood, for which the posterior can be computed in closed form (see e.g. Rasmussen and Williams, 2006, §2.2).

Interest in GP models stems from their functional non-parametric Bayesian nature and there are at least three critical benefits from adopting such a modeling approach. Firstly, by being a prior over functions, they represent a much more elegant way to address problems such as regression, where it is less natural to think of a prior over parameters such as the weights in a linear model. Secondly, by being Bayesian, they provide us with a principled way to combine our prior beliefs with the observed data in order to predict a full posterior distribution over unknown quantities, which of course is much more informative than a single point prediction. Finally, by being non-parametric, they address the common concern of parametric approaches about how well we can fit the data, since the model space is not constrained to have a parametric form.

1.1 Key Inference Challenges in Gaussian Process Models

Nevertheless, such a principled and flexible approach to machine learning comes at the cost of facing two fundamental probabilistic inference challenges, (i) scalability to large datasets and (ii) dealing with nonlinear likelihoods. With regards to the first challenge, scalability, GP models have been notorious for their poor scalability as a function of the number of training points. Despite ground-breaking work in understanding scalable GPs through the so-called sparse approximations (Quiñonero-Candela and Rasmussen, 2005) and in developing practical inference methods for these models (Titsias, 2009), recent literature on addressing this challenge gives a clear indication that the scalability problem is still a very active area of research, see e.g. Hensman et al. (2013); Gal et al. (2014); Das et al. (2015); Deisenroth and Ng (2015); Hoang et al. (2015).

Concerning the second challenge, dealing with nonlinear likelihoods, the main difficulty is that of estimating a posterior over latent functions distributed according to a GP prior, given observations assumed to be drawn from a possibly nonlinear conditional likelihood model. In general, this posterior is analytically intractable and one must resort to approximate methods. These methods can be roughly classified into stochastic approaches and deterministic approaches. The former, stochastic approaches, are based on sampling algorithms such as Markov Chain Monte Carlo (MCMC, see e.g. Neal, 1993, for an overview) and the latter, deterministic approaches, are based on optimization techniques and include variational inference (VI, Jordan et al., 1998), the Laplace approximation (see e.g. MacKay, 2003, Chapter 27) and expectation propagation (EP, Minka, 2001).

On the one hand, although stochastic techniques such as MCMC provide a flexible framework for sampling from complex posterior distributions of probabilistic models, their generality comes at the expense of a very high computational cost as well as cumbersome convergence analysis. On the other hand, deterministic methods such as variational inference build upon the main insight that optimization is generally easier than integration. Consequently, they estimate a posterior by maximizing a lower bound of the marginal likelihood, the so-called evidence lower bound (ELBO). Variational methods can be considerably faster
than MCMC but they lack MCMC’s broader applicability, usually requiring mathematical derivations on a model-by-model basis\(^1\).

\subsection*{1.2 Contributions}

In this paper we address the above challenges by developing a scalable automated variational method for inference in models with Gaussian process priors and general likelihoods. This method reduces the overhead of the tedious mathematical derivations traditionally inherent to variational algorithms, allowing their application to a wide range of problems. In particular, we consider models with multiple latent functions, multiple outputs and non-linear likelihoods that satisfy the following properties: (i) factorization across latent functions and (ii) across observations. The former assumes that, when there are more than one latent function, they are generated from independent GPs. The latter assumes that, given the latent functions, the observations are conditionally independent.

Existing GP models, such as regression (Rasmussen and Williams, 2006), binary and multi-class classification (Nickisch and Rasmussen, 2008; Williams and Barber, 1998), warped GPs (Snelson et al., 2003), log Gaussian Cox process (Møller et al., 1998), and multi-output regression (Wilson et al., 2012), all fall into this class of models. Furthermore, our approach goes well beyond standard settings for which elaborate learning machinery has been developed, as we only require access to the likelihood function in a black-box manner. As we shall see below, our inference method can scale up to very large datasets, hence we will refer to it as SAVIGP, which stands for scalable automated variational inference for Gaussian process models. The key characteristics of our method and contributions of this work are summarized below.

- **Black-box likelihoods**: As mentioned above, the main contribution of this work is to be able to carry out posterior inference with GP priors and general likelihood models, without knowing the details of the conditional likelihood (or its gradients) and only requiring its evaluation as a black-box function.

- **Scalable inference and stochastic optimization**: By building upon the inducing-variable approach underpinning most sparse approximations to GP models (Quinonero-Candela and Rasmussen, 2005; Titsias, 2009), the computational complexity of our method is dominated by \(O(M^3)\) operations in time, where \(M \ll N\) is the number of inducing points and \(N\) the number of observations. This is in contrast to naive inference in GP models which has a time complexity of \(O(N^3)\). As the resulting ELBO decomposes over the training datapoints, our model is amenable to parallel computation and stochastic optimization. In fact, we provide an implementation that can scale up to a very large number of observations, exploiting stochastic optimization, multi-core architectures and GPU computation.

- **Joint learning of model parameters**: As our approach is underpinned by variational inference principles, SAVIGP allows for learning of all model parameters, including

\(^1\)However, we note recent developments such as variational auto-encoders (Kingma and Welling, 2014) and refer the reader to the related work and the discussion (Sections 2.4 and 10, respectively).
posterior parameters, inducing inputs, covariance hyperparameters and likelihood parameters, within the same framework via maximization of the evidence lower bound (ELBO).

- **Multiple outputs and multiple latent functions:** SAVIGP is designed to support models with multiple outputs and multiple latent functions, such as in multi-class classification (Williams and Barber, 1998) and non-stationary multi-output regression (Wilson et al., 2012). It does so in a very flexible way, allowing the different GP priors on the latent functions to have different covariance functions and inducing inputs.

- **Flexible posterior:** SAVIGP uses a mixture of Gaussians as the approximating posterior distribution. This is a very general approach as it is well known that, with a sufficient number of components, almost any continuous density can be approximated to arbitrary accuracy (Maz’ya and Schmidt, 1996).

- **Statistical efficiency:** By using knowledge of the approximate posterior and the structure of the GP prior, we exploit the decomposition of the ELBO, into a KL-divergence term and an expected log likelihood term, to provide statistically efficient parameter estimates. In particular, we derive an analytical lower bound for the KL-divergence term and we show that, for general black-box likelihood models, the expected log likelihood term and its gradients can be computed efficiently using empirical expectations over univariate Gaussian distributions.

- **Efficient re-parametrization:** For the case of a single full Gaussian variational posterior, we show that it is possible to re-parametrize the model so that the optimal posterior can be represented using a parametrization that is linear in the number of observations. This parametrization becomes useful for denser models, i.e. for models that have a larger number of inducing variables.

- **Extensive experimentation:** We evaluate SAVIGP with experiments on small datasets, medium-scale datasets and a large dataset. The experiments on small datasets ($N < 1,300$) evaluate the effectiveness of the method under different likelihood models and sparsity levels (as determined by the number of inducing variables), including problems such as regression, classification, Log Gaussian Cox processes, and warped GPs (Snelson et al., 2003). We show that SAVIGP can perform as well as hard-coded inference methods under high levels of sparsity. The medium-scale experiments consider binary and multi-class classification using the MNIST dataset ($N = 60,000$) and non-stationary regression under the GPRN model (Wilson et al., 2012) using the SARCOS dataset ($N \approx 45,000$). Besides showing the competitiveness of our model for problems at this scale, we analyze the effect of learning the inducing inputs, i.e. the location of inducing variables. Finally, in the large-scale experiment, we study the problem of predicting airline delays (using $N = 700,000$), and show that our method is on par with the state-of-the-art approach for scalable GP regression (Hensman et al., 2013), which uses full knowledge of the likelihood model.

Before describing the family of GP models that we focus on, we start by relating our work to the previous literature concerning the key inference challenges mentioned above, i.e. scalability and dealing with non-linear likelihoods.
2 Related Work

As pointed out by Rasmussen and Williams (2006), there has been a long-standing interest in Gaussian processes with early work dating back at least to the 1880s when Danish astronomer and mathematician T. N. Thiel, concerned with determining the distance between Copenhagen and Lund from astronomical observations, essentially proposed the first mathematical formulation of Brownian motion (see e.g. Lauritzen, 1981, for details). In geostatistics, GP regression is known as Kriging (see e.g. Cressie, 1993) where, naturally, it has focused on 2-dimensional and 3-dimensional problems.

While the work of O’Hagan and Kingman (1978) has been quite influential in applying GPs to general regression problems, the introduction of GP regression to main-stream machine learning by Williams and Rasmussen (1996) sparked significant interest in the machine learning community. Indeed, henceforth, the community has taken GP models well beyond the standard regression setting, addressing other problems such as non-stationary and heteroscedastic regression (Paciorek and Schervish, 2004; Kersting et al., 2007; Wilson et al., 2012); nonlinear dimensionality reduction (Lawrence, 2005); classification (Williams and Barber, 1998; Nickisch and Rasmussen, 2008); multi-task learning (Bonilla et al., 2008; Yu and Chu, 2008; Alvarez and Lawrence, 2009; Wilson et al., 2012); preference learning (Bonilla et al., 2010); and ordinal regression (Chu and Ghahramani, 2005). In fact, their book (Rasmussen and Williams, 2006) is the de facto reference in any work related to Gaussian process models in machine learning. As we shall see below, despite all these significant advances, most previous work in the GP community has focused on addressing the scalability and the non-linear likelihood challenges in isolation.

2.1 Scalable Models

The cubic scaling on the number of observations of Gaussian process models, until very recently, has hindered the use of these models in a wider variety of applications. Work on approaching this problem has ranged from selecting informative (inducing) datapoints from the training data so as to facilitate sparse approximations to the posterior (Lawrence et al., 2002) to considering these inducing points as continuous parameters and optimizing them within a coherent probabilistic framework (Snelson and Ghahramani, 2006).

Although none of these methods actually scale to very large datasets as their time complexity is $O(M^2N)$, where $N$ is the number of observations and $M$ is the number of inducing points, unifying such approaches from a probabilistic perspective (Quiñonero-Candela and Rasmussen, 2005) has been extremely valuable to the community, not only to understand what those methods are actually doing but also to develop new approaches to sparse GP models. In particular, the framework of Titsias (2009) has become the underpinning machinery of modern scalable approaches to GP regression and classification. This has been taken one step further by allowing optimization of variational parameters within a stochastic optimization framework (Hensman et al., 2013), hence enabling the applicability of these inference methods to very large datasets.

Besides inducing-variable approaches to scalability in GP-models, other approaches have exploited the relationship between infinite-feature linear-in-the-parameters models and GPs. In particular, early work investigated truncated linear-in-the-parameters models as approx-
imations to GP regression (see e.g. Tipping, 2001). This idea has been developed in an alternative direction that exploits the relationship between a covariance function of a stationary process and its spectral density, hence providing random-feature approximations to the covariance function of a GP, similarly to the work of Rahimi and Recht (2008, 2009), who focused on non-probabilistic kernel machines. For example, Lázaro-Gredilla et al. (2010) and Yang et al. (2015) have followed these types of approaches, with the latter mostly concerned about having fast kernel evaluations.

Unlike our work, none of these approaches deals with the harder task of developing scalable inference methods for multi-output problems and general likelihood models.

### 2.2 Multi-output and Multi-task Gaussian Processes

Developing multi-task or multi-output learning approaches using Gaussian processes has also proved an intensive area of research. Most GP-based models for these problems assume that correlations between the outputs are induced via linear combinations of a set of independent latent processes. Such linear combinations can use fixed coefficients (see e.g. Teh et al., 2005) or input-dependent coefficients (Wilson et al., 2012; Nguyen and Bonilla, 2013). Tasks dependencies can also be defined explicitly through the covariance function as done by Bonilla et al. (2008), who assume a covariance decomposition between tasks and inputs. More complex dependencies can be modeled by using the convolution formalism as done in earlier work by Boyle and Frean (2005) and generalized by Alvarez and Lawrence (2009); Álvarez et al. (2010), whose later work also provides efficient inference algorithms for such models (Álvarez and Lawrence, 2011). Finally, the work of Nguyen and Bonilla (2014b) also assumes a linear combination of independent latent processes but it exploits the developments of Hensman et al. (2013) to scale up to very large datasets.

Besides the work of Nguyen and Bonilla (2014b), these approaches do not scale to a very large number of observations and all of them are mainly concerned with regression problems.

### 2.3 General Nonlinear Likelihoods

The problem of inference in models with GP priors and nonlinear likelihoods has been tackled from a sampling perspective with algorithms such as elliptical slice sampling (ESS, Murray et al., 2010), which are more effective at drawing samples from strongly correlated Gaussians than generic MCMC algorithms. Nevertheless, as we shall see in Section 9.3.5 the sampling cost of ESS remains a major challenge for practical usages.

From variational inference perspective, the work by (Opper and Archambeau, 2009) has been considerably under-recognized by the community even though it proposes an efficient full Gaussian posterior approximation for GP models with iid observations. Our work pushes this breakthrough further by allowing multiple latent functions, multiple outputs, and more importantly, scalability to large datasets.

Another approach to deterministic approximate inference is the integrated nested Laplace approximation (INLA, Rue et al., 2009). INLA uses numerical integration to approximate the marginal likelihood, which makes it unsuitable for GP models that contain a large number of hyperparameters.
2.4 Recent developments

As mentioned in Section 1, the scalability problem continues to attract the interest of researchers working with GP models, with recently developed distributed inference frameworks (Gal et al., 2014; Deisenroth and Ng, 2015), and the variational inference frameworks for scalable GP regression and classification by Hensman et al. (2013) and Hensman et al. (2015), respectively. As with the previous work described in Section 2.1, these approaches have been limited to classification or regression problems.

Contemporarily to our work, black-box variational inference (BBVI, Ranganath et al., 2014) has been developed for general latent variable models. Due to this generality, it under-utilizes the rich amount of information available in Gaussian process models. For example, BBVI approximates the KL-divergence term of the evidence lower bound but this is computed analytically in our method. Additionally, for practical reasons, BBVI imposes the variational distribution to fully factorize over the latent variables, while we make no such a restriction. A clear disadvantage of BBVI is that it does not provide a practical way of learning the covariance hyperparameters of GPs — in fact, these are set to fixed values. In principle, these values can be learned in BBVI using stochastic optimization, but experimentally, we have found this to be problematic, ineffectual, and time-consuming.

Very recently, Bonilla et al. (2016) have used the random-feature approach mentioned in Section 2.1 along with linearization techniques to provide scalable methods for inference in GP models with general likelihoods. Unlike our approach for estimating expectations of the conditional likelihood, such linearizations of the conditional likelihood are approximations and generally do not converge to the exact expectations even in the limit of a large number of observations.

Following the recent advances in making automatic differentiation widely available and easy to use in practical systems (see e.g. Baydin et al., 2015), developments on stochastic variational inference for fairly complex probabilistic models (Rezende et al., 2014; Kingma and Welling, 2014) can be used when the likelihood can be implemented in such systems and can be expressed using the so-called re-parametrization trick (see e.g. Kingma and Welling, 2014, for details). Nevertheless, the advantage of our method over these generic approaches is twofold. Firstly, as with BBVI, in practice such a generality comes at the cost of making assumptions about the posterior (such as factorization), which is not suitable for GP models. Secondly, and more importantly, such methods are not truly black-box as they require explicit access to the implementation of the conditional likelihood.

Finally, a related area of research is that of modeling complex data with deep belief networks based on Gaussian process mappings (Damianou and Lawrence, 2013), which have been proposed primarily as hierarchical extensions of the Gaussian process latent variable model (Lawrence, 2005). Unlike our approach, these models target the unsupervised problem of discovering structure in high-dimensional data and have focused mainly on small-data applications.

3 Latent Gaussian Process Models

Before starting our description of the types of models we are addressing in this paper, we refer the reader to Appendix K for a summary of the notation used henceforth.
We consider supervised learning problems where we are given a dataset \( D = \{x_n, y_n\}_{n=1}^N \), where \( x_n \) is a \( D \)-dimensional input vector and \( y_n \) is a \( P \)-dimensional output. Our goal is to learn the mapping from inputs to outputs, which can be established via \( Q \) underlying latent functions \( \{f_j\}_{j=1}^Q \). In many problems these latent functions have a physical interpretation, while in others they are simply nuisance parameters and we just want to integrate them out in order to make probabilistic predictions.

A sensible modeling approach to the above problem is to assume that the \( Q \) latent functions \( \{f_j\} \) are uncorrelated a priori and that they are drawn from \( Q \) zero-mean Gaussian processes (Rasmussen and Williams, 2006):

\[
p(f_j|\theta_j) \sim \text{GP}(0, \kappa_j(\cdot, \cdot; \theta_j)), \quad j = 1, \ldots, Q,
\]

\[
p(f|\theta) = \prod_{j=1}^Q p(f_j|\theta_j) = \prod_{j=1}^Q \mathcal{N}(f_j; 0, K_{xx}^j),
\]

where \( f \) is the set of all latent function values; \( f_j = \{f_j(x_n)\}_{n=1}^N \) denotes the values of latent function \( j \); \( K_{xx}^j \) is the covariance matrix induced by the covariance function \( \kappa_j(\cdot, \cdot; \theta_j) \) evaluated at every pair of inputs; and \( \theta = \{\theta_j\} \) are the parameters of the corresponding covariance functions. Along with the prior in Equation (1), we can also assume that our multi-dimensional observations \( \{y_n\} \) are iid given the corresponding set of latent functions \( \{f_n\} \):

\[
p(y|f, \phi) = \prod_{n=1}^N p(y_n|f_n, \phi),
\]

where \( y \) is the set of all output observations; \( y_n \) is the \( n \)th output observation; \( f_n = \{f_j(x_n)\}_{j=1}^Q \) is the set of latent function values which \( y_n \) depends upon; and \( \phi \) are the conditional likelihood parameters. In the sequel, we will refer to the covariance parameters \( (\theta) \) as the model hyperparameters.

In other words, we are interested in models for which the following criteria are satisfied:

(i) factorization of the prior over the latent functions, as specified by Equation (1), and

(ii) factorization of the conditional likelihood over the observations given the latent functions, as specified by Equation (2).

We refer to the models satisfying the above assumptions (when using GP priors) as latent Gaussian process models (LGPMs). Interestingly, a large class of problems can be well modeled with the above assumptions. For example binary classification (Nickisch and Rasmussen, 2008; Williams and Barber, 1998), warped GPs (Snelson et al., 2003), log Gaussian Cox processes (Møller et al., 1998), multi-class classification (Williams and Barber, 1998), and multi-output regression (Wilson et al., 2012) all belong to this family of models.

More importantly, besides the iid assumption, there are not additional constraints on the conditional likelihood which can be any linear or nonlinear model. Furthermore, as we shall see in Section 4, our proposed inference algorithm only requires evaluations of this likelihood model in a black-box manner, i.e. without requiring detailed knowledge of its implementation or its gradients.
3.1 Inference Challenges

As mentioned in Section 1, for general LGPMs, the inference problem of estimating the posterior distribution over the latent functions given the observed data \( p(f|D) \) and, ultimately, for a new observation \( x_\star \), estimating the predictive posterior distribution \( p(f_\star |x_\star, D) \), poses two important challenges (even in the case of a single latent function \( Q = 1 \)) from the computational and statistical perspectives. These challenges are (i) scalability to large datasets and (ii) dealing with nonlinear likelihoods.

We address the scalability challenge inherent to GP models (given their cubic time complexity on the number of observations) by augmenting our priors using an inducing-variable approach (see e.g. Quiñonero-Candela and Rasmussen, 2005) and embedding our model into a variational inference framework (Titsias, 2009). In short, inducing variables in sparse GP models act as latent summary statistics, avoiding the computation of large inverse covariances. Crucially, unlike other approaches (Quiñonero-Candela and Rasmussen, 2005), we keep an explicit representation of these variables (and integrate them out variationally), which facilitates the decomposition of the variational objective into a sum on the individual datapoints. This allows us to devise stochastic optimization strategies and parallel implementations in cloud computing services such as Amazon EC2. Such a strategy, also allows us to learn the location of the inducing variables (inducing inputs) in conjunction with variational parameters and hyperparameters, which in general provides better performance, especially in high-dimensional problems.

To address the nonlinear likelihood challenge, which from a variational perspective boils down to estimating expectations of a nonlinear function over the approximate posterior, we follow a stochastic estimation approach, in which we develop low-variance Monte Carlo estimates of the expected log-likelihood term in the variational objective. Crucially, we will show that the expected log-likelihood term can be estimated efficiently by using only expectations over univariate Gaussian distributions.

4 Scalable Inference

Here we describe our scalable inference method for the model class specified in Section 3. We build upon the inducing-point formalism underlying most sparse GP approximations (Quiñonero-Candela and Rasmussen, 2005; Titsias, 2009) and obtain an algorithm with time complexity \( O(M^3) \), where \( M \) is the number of inducing points per latent process. We show that, under this sparse approximation and the variational inference framework, the expected log likelihood term and its gradient can be estimated using only expectations over univariate Gaussian distributions.

4.1 Augmented Prior

In order to make inference scalable we redefine our prior in terms of some auxiliary variables \( \{u_{j}\}_{j=1}^{Q} \), which we will refer to as the inducing variables. These inducing variables lie in the same space as \( \{f_{j}\} \) and are drawn from the same zero-mean GP priors. As before, we assume factorization of the prior across the \( Q \) latent functions. Hence the resulting
The augmented prior is given by:

\[
p(u) = \prod_{j=1}^{Q} \mathcal{N}(u_j; 0, K_{zz}^j), \quad p(f|u) = \prod_{j=1}^{Q} \mathcal{N}(f_j; \tilde{\mu}_j, \tilde{K}_j),
\]

where \( \tilde{\mu}_j = K_{xz}^j (K_{zz}^j)^{-1} u_j \), and
\[
\tilde{K}_j = K_{xx}^j - A_j K_{zx}^j \text{ with } A_j = K_{xz}^j (K_{zz}^j)^{-1},
\]

where \( u_j \) are the inducing variables for latent process \( j \); \( u \) is the set of all the inducing variables; \( Z_j \) are all the inducing inputs for latent process \( j \); \( X \) is the matrix of all input locations \( \{x_i\} \); and \( K_{uv}^j \) is the covariance matrix induced by evaluating the covariance function \( \kappa_j(\cdot, \cdot) \) at all pairwise columns of matrices \( U \) and \( V \). We note that while each of the inducing variables in \( u_j \) lies in the same space as the elements in \( f_j \), each of the \( M \) inducing inputs in \( Z_j \) lies in the same space as each input data point \( x_n \). Therefore, while \( u_j \) is a \( M \)-dimensional vector, \( Z_j \) is a \( M \times D \) matrix where each of the rows corresponds to a \( D \)-dimensional inducing input. We refer the reader to Appendix K for a summary of the notation and dimensionality of the above kernel matrices.

As thoroughly investigated by Quiñonero-Candela and Rasmussen (2005), most GP approximations can be formulated using the augmented prior above and additional assumptions on the training and test conditional distributions \( p(f|u) \) and \( p(f_*|u) \), respectively. Such approaches have been traditionally referred to as sparse approximations and we will use this terminology as well. Analogously, we will refer to models with a larger number of inducing inputs as denser models.

It is important to emphasize that the joint prior \( p(f, u) \) defined in Equation (3) is an equivalent prior to that in the original model, as if we integrate out the inducing variables \( u \) from this joint prior we will obtain the prior \( p(f) \) in Equation (1) exactly. Nevertheless, as we shall see later, following a variational inference approach and having an explicit representation of the (approximate) posterior over the inducing variables will be fundamental to scaling up inference in these types of models without making the assumptions on the training or test conditionals described in Quiñonero-Candela and Rasmussen (2005).

Along with the joint prior defined above, we maintain the factorization assumption of the conditional likelihood given in Equation (2).

### 4.2 Variational Inference and the Evidence Lower Bound

Given the prior in Equation (3) and the likelihood in Equation (2), posterior inference for general (non-Gaussian) likelihoods is analytically intractable. Therefore, we resort to approximate methods such as variational inference (Jordan et al., 1998). Variational inference methods entail positing a tractable family of distributions and finding the member of the family that is “closest” to the true posterior in terms of their the Kullback-Leibler divergence. In our case, we are seeking to approximate the joint posterior \( p(f, u|y) \) with a variational distribution \( q(f, u|\lambda) \).
4.3 Approximate Posterior

Motivated by the fact that the true joint posterior is given by \( p(f, u|y) = p(f|u, y)p(u|y) \), our approximate posterior has the form:

\[
q(f, u|\lambda) = p(f|u)q(u|\lambda),
\]

where \( p(f|u) \) is the conditional prior given in Equation (3) and \( q(u|\lambda) \) is our approximate (variational) posterior. This decomposition has proved effective in regression problems with a single latent process and a single output (see e.g. Titsias, 2009).

Hence, we can define our variational distribution using a mixture of Gaussians (MoG):

\[
q(u|\lambda) = \sum_{k=1}^{K} \pi_k q_k(u|m_k, S_k) = \sum_{k=1}^{K} \pi_k \prod_{j=1}^{Q} \mathcal{N}(u_j; m_{kj}, S_{kj}),
\]

where \( \lambda = \{\pi_k, m_{kj}, S_{kj}\} \) are the variational parameters: the mixture proportions \( \{\pi_k\} \), the posterior means \( \{m_{kj}\} \) and posterior covariances \( \{S_{kj}\} \) of the inducing variables corresponding to mixture component \( k \) and latent function \( j \). We also note that each of the mixture components \( q_k(u|m_k, S_k) \) is a Gaussian with mean \( m_k \) and block-diagonal covariance \( S_k \).

4.4 Variational Lower Bound

It is easy to show that minimizing the Kullback-Leibler divergence between our approximate posterior and the true posterior, \( \text{KL}(q(f, u|\lambda)||p(f, u|y)) \) is equivalent to maximizing the log-evidence lower bound (\( \mathcal{L}_{\text{elbo}} \)), which is composed of a KL-term (\( \mathcal{L}_{\text{kl}} \)) and an expected log-likelihood term (\( \mathcal{L}_{\text{ell}} \)). In other words:

\[
\log p(y) \geq \mathcal{L}_{\text{elbo}}(\lambda) \overset{\text{def}}{=} \mathcal{L}_{\text{kl}}(\lambda) + \mathcal{L}_{\text{ell}}(\lambda),
\]

where

\[
\mathcal{L}_{\text{kl}}(\lambda) = -\text{KL}(q(f, u|\lambda)||p(f, u)),
\]

and

\[
\mathcal{L}_{\text{ell}}(\lambda) = \mathbb{E}_{q(f, u|\lambda)}[\log p(y|f)],
\]

where \( \mathbb{E}_{q(x)}[g(x)] \) denotes the expectation of function \( g(x) \) over distribution \( q(x) \). Here we note that \( \mathcal{L}_{\text{kl}} \) is a negative KL divergence between the joint approximate posterior \( q(f, u|\lambda) \) and the joint prior \( p(f, u) \). Therefore, maximization of the \( \mathcal{L}_{\text{elbo}} \) in Equation (7) entails minimization of an expected loss (given by the negative expected log-likelihood \( \mathcal{L}_{\text{ell}} \)) regularized by \( \mathcal{L}_{\text{kl}} \), which imposes the constraint of finding solutions to our approximate posterior that are close to the prior in the KL-sense.

An interesting observation of the decomposition of the \( \mathcal{L}_{\text{elbo}} \) objective is that, unlike \( \mathcal{L}_{\text{ell}} \), \( \mathcal{L}_{\text{kl}} \) in Equation (8) does not depend on the conditional likelihood \( p(y|f) \), for which we do not assume any specific parametric form (i.e. black-box likelihood). We can thus address the technical difficulties regarding each component and their gradients separately using different approaches. In particular, for \( \mathcal{L}_{\text{kl}} \) we will exploit the structure of the variational posterior in order to avoid computing KL-divergences over distributions involving all the data. Furthermore, we will obtain a lower bound for \( \mathcal{L}_{\text{kl}} \) in the general case of \( q(u|\lambda) \) being a mixture-of-Gaussians (MoG) as given in Equation (6). For the expected log-likelihood term (\( \mathcal{L}_{\text{ell}} \)) we will use a Monte Carlo approach in order to estimate the required expectation and its gradients.
4.5 Computation of the KL-divergence term ($\mathcal{L}_{kl}$)

In order to have an explicit form for $\mathcal{L}_{kl}$ and its gradients, we start by expanding Equation (8):

$$\mathcal{L}_{kl}(\lambda) = -\text{KL}(q(f,u|\lambda) \| p(f,u)) = -\mathbb{E}_{q(f,u|\lambda)} \left[ \log \frac{q(f,u|\lambda)}{p(f,u)} \right],$$ (10)

$$= -\mathbb{E}_{p(f|u)q(u|\lambda)} \left[ \log \frac{q(u|\lambda)}{p(u)} \right],$$ (11)

$$= -\text{KL}(q(u|\lambda) \| p(u)),$$ (12)

where we have applied the definition of the KL-divergence in Equation (10); used the variational joint posterior $q(f,u|\lambda)$ given in Equation (5) to go from Equation (10) to Equation (11); and integrated out $f$ to obtain Equation (12). We note that the definition of the joint posterior $q(f,u|\lambda)$ in Equation (5) has been crucial to transform a KL-divergence between the joint approximate posterior and the joint prior into a KL-divergence between the variational posterior $q(u|\lambda)$ and the prior $p(u)$ over the inducing variables. In doing that, we have avoided computing a KL-divergence between distributions over the $N$-dimensional variables $f_j$. As we shall see later, this implies a reduction in time complexity from $\mathcal{O}(N^3)$ to $\mathcal{O}(M^3)$, where $N$ is the number of datapoints and $M$ is the number of inducing variables.

We now decompose the resulting KL-divergence term in Equation (12) as follows,

$$\mathcal{L}_{kl}(\lambda) = -\text{KL}(q(u|\lambda) \| p(u)) = \mathcal{L}_{\text{ent}}(\lambda) + \mathcal{L}_{\text{cross}}(\lambda),$$

where:

$$\mathcal{L}_{\text{ent}}(\lambda) = -\mathbb{E}_{q(u|\lambda)} \left[ \log q(u|\lambda) \right],$$ and

$$\mathcal{L}_{\text{cross}}(\lambda) = \mathbb{E}_{q(u|\lambda)} \left[ \log p(u) \right],$$ (13)

where $\mathcal{L}_{\text{ent}}(\lambda)$ denotes the differential entropy of the approximating distribution $q(u|\lambda)$ and $\mathcal{L}_{\text{cross}}(\lambda)$ denotes the negative cross-entropy between the approximating distribution $q(u|\lambda)$ and the prior $p(u)$.

Computing the entropy of the variational distribution in Equation (6), which is a mixture-of-Gaussians (MoG), is analytically intractable. However, a lower bound of this entropy can be obtained using Jensen’s inequality (see e.g. Huber et al., 2008) giving:

$$\mathcal{L}_{\text{ent}}(\lambda) \geq -\sum_{k=1}^{K} \pi_k \log \sum_{\ell=1}^{K} \pi_\ell \mathcal{N}(\mathbf{m}_k; \mathbf{m}_\ell, \mathbf{S}_k + \mathbf{S}_\ell)^{\text{det}} \equiv \hat{\mathcal{L}}_{\text{ent}}.\quad (14)$$

The negative cross-entropy in Equation (13) between a Gaussian mixture $q(u|\lambda)$ and a Gaussian $p(u)$, can be obtained analytically,

$$\mathcal{L}_{\text{cross}}(\lambda) = -\frac{1}{2} \sum_{k=1}^{K} \sum_{j=1}^{Q} \pi_k \sum_{j=1}^{Q} \left[ M \log 2\pi + \log |\mathbf{K}_{zz}^j| + \mathbf{m}_{kj}^T \left( \mathbf{K}_{zz}^j \right)^{-1} \mathbf{m}_{kj} + \text{tr} \left( \mathbf{K}_{zz}^j \right)^{-1} \mathbf{S}_{kj} \right].\quad (15)$$

The derivation of the entropy term and the cross-entropy term in Equations (14) and (15) is given in Appendix A.
4.6 Estimation of the expected log likelihood term ($\mathcal{L}_{\text{ell}}$)

We now address the computation of the expected log likelihood term in Equation (9). The main difficulty of computing this term is that, unlike the $\mathcal{L}_{\text{kl}}$ where we have full knowledge of the prior and the approximate posterior, here we do not assume a specific form for the conditional likelihood $p(y|f, \phi)$. Furthermore, we only require evaluations of $\log p(y_n|f_n, \phi)$ for each datapoint $n$, hence yielding a truly black-box likelihood method.

We will show one of our main results, that of statistical efficiency of our $\mathcal{L}_{\text{ell}}$ estimator. This means that, despite having a full Gaussian approximate posterior, estimation of the $\mathcal{L}_{\text{ell}}$ and its gradients only requires expectations over univariate Gaussian distributions. We start by expanding Equation (9) using our definitions of the approximate posterior and the factorization of the conditional likelihood:

$$\mathcal{L}_{\text{ell}}(\lambda) = \mathbb{E}_{q(f,u|\lambda)}[\log p(y|f, \phi)] = \mathbb{E}_{q(f|\lambda)}[\log p(y|f, \phi)],$$

where, given the definition of the approximate joint posterior $q(f,u|\lambda)$ in Equation (5), the distribution $q(f|\lambda)$ resulting from marginalizing $u$ from this joint posterior can be obtained analytically,

$$q(f|\lambda) = \sum_{k=1}^{K} \pi_k q_k(f|\lambda_k) = \sum_{k=1}^{K} \pi_k \prod_{j=1}^{Q} \mathcal{N}(f_j; b_{kj}, \Sigma_{kj}),$$

where $\pi_k$ and $\Sigma_{kj}$ are given in Equation (4) and, as defined before, $\{m_{kj}\}$ and $\{s_{kj}\}$ are the posterior means and posterior covariances of the inducing variables corresponding to mixture component $k$ and latent function $j$. We are now ready to state our result of a statistically efficient estimator for the $\mathcal{L}_{\text{ell}}$:

**Theorem 1** For the GP model with prior defined in Equations (3) to (4), and conditional likelihood defined in Equation (2), the expected log likelihood over the variational distribution in Equation (5) and its gradients can be estimated using expectations over univariate Gaussian distributions.

The proof is constructive and can be found in Appendix B. We note that a less general result, for the case of one latent function and a single variational Gaussian posterior, was obtained in Opper and Archambeau (2009) using a different derivation. Here we state our final result on how to compute these estimates:

$$\mathcal{L}_{\text{ell}}(\lambda) = \sum_{n=1}^{N} \sum_{k=1}^{K} \pi_k \mathbb{E}_{q_k(n)}(f_n|\lambda_k)[\log p(y_n|f_n, \phi)],$$

$$\nabla_{\lambda_k} \mathcal{L}_{\text{ell}}(\lambda) = \pi_k \sum_{n=1}^{N} \mathbb{E}_{q_k(n)}(f_n|\lambda_k)[\nabla_{\lambda_k} \log q_k(n)(f_n|\lambda_k) \log p(y_n|f_n, \phi)],$$

for $\lambda_k \in \{m_k, s_k\}$.
\[ \nabla_{\pi_k} \mathcal{L}_{\text{ell}}(\lambda) = \sum_{n=1}^{N} \mathbb{E}_{q_k(n)}(f_n|\lambda_k)[\log p(y_n|f_n, \phi)], \]  
(22)

where \( q_k(n)(f_n, \lambda_k) \) is a \( Q \)-dimensional Gaussian with:
\[
q_k(n)(f_n|\lambda_k) = \mathcal{N}(f_n; b_k(n), \Sigma_k(n)),
\]
where \( \Sigma_k(n) \) is a diagonal matrix. The \( j \)th element of the mean and the \((j,j)\)th entry of the covariance of the above distribution are given by:
\[
[b_k(n)]_j = a^T_{jn}m_{kj}, \quad [\Sigma_k(n)]_{j,j} = [\tilde{K}_j]_{n,n} + a^T_{jn}S_{kj}a_{jn},
\]  
(23)

where \( a_{jn} \) denotes the \( M \)-dimensional vector corresponding to the \( n \)th column of matrix \( A_j \); \( \tilde{K}_j \) and \( A_j \) are given in Equation (4); and, as before, \( \{m_{kj}, S_{kj}\} \) are the variational parameters corresponding to the mean and covariance of the approximate posterior over the inducing variables for mixture component \( k \) and latent process \( j \).

We emphasize that when \( Q > 1 \), \( q_k(n)(f_n, \lambda_k) \) is not a univariate marginal but a \( Q \)-dimensional marginal posterior with diagonal covariance. Therefore, only expectations over univariate Gaussians are required to estimate the expressions in Equations (20) to (22).

### 4.6.1 Practical consequences and unbiased estimates

There are two immediate practical consequences of the result in Theorem 1. The first consequence is that we can use unbiased empirical estimates of the expected log likelihood term and its gradients. In our experiments we use Monte Carlo (MC) estimates, hence we can compute \( \mathcal{L}_{\text{ell}} \) as:
\[
\hat{\mathcal{L}}_{\text{ell}} = \frac{1}{S} \sum_{n=1}^{N} \sum_{k=1}^{K} \pi_k \sum_{i=1}^{S} \log p(y_n|f^{(k,i)}_n, \phi),
\]  
(24)

where \( b_k(n) \) and \( \Sigma_k(n) \) are the vector and matrix forms of the mean and covariance of the posterior over the latent functions as given in Equation (23). Analogous MC estimates of the gradients are given in Appendix C.2.

The second practical consequence is that in order to estimate the gradients of the \( \mathcal{L}_{\text{ell}} \), using Equations (21) and (22), we only require evaluations of the conditional likelihood in a black-box manner, without resorting to numerical approximations or analytical derivations of its gradients.

### 5 Parameter Optimization

In order to learn the parameters of our model we seek to maximize our (estimated) log-evidence lower bound \( \hat{\mathcal{L}}_{\text{elbo}} \) using gradient-based optimization. Let \( \eta = \{\lambda, \theta, \phi\} \) be all the model parameters for which point estimates are required. We have that:
\[
\hat{\mathcal{L}}_{\text{elbo}}(\eta) \equiv \hat{\mathcal{L}}_{\text{ent}}(\eta) + \mathcal{L}_{\text{cross}}(\eta) + \hat{\mathcal{L}}_{\text{ell}}(\eta),
\]
\[ \nabla \eta \hat{\mathcal{L}}_{\text{elbo}} = \nabla \eta \hat{\mathcal{L}}_{\text{ent}} + \nabla \eta \mathcal{L}_{\text{cross}} + \nabla \eta \hat{\mathcal{L}}_{\text{ell}}, \]

where we have made explicit the dependency of the log-evidence lower bound on any parameter of the model and \( \hat{\mathcal{L}}_{\text{ent}}, \mathcal{L}_{\text{cross}}, \) and \( \hat{\mathcal{L}}_{\text{ell}} \) are given in Equations (14), (15), (24) respectively. The gradients of \( \hat{\mathcal{L}}_{\text{elbo}} \) wrt variational parameters \( \lambda \), covariance hyperparameters \( \theta \) and likelihood parameters \( \phi \) are given in Appendices C, D.1, and D.2, respectively. As shown in these appendices, not all constituents of \( \hat{\mathcal{L}}_{\text{elbo}} \) contribute to learning all parameters, for example \( \nabla \theta \hat{\mathcal{L}}_{\text{ent}} = \nabla \phi \hat{\mathcal{L}}_{\text{ent}} = \nabla \phi \mathcal{L}_{\text{cross}} = 0. \)

Using the above objective function (\( \hat{\mathcal{L}}_{\text{elbo}} \)) and its gradients we can consider batch optimization with limited-memory BFGS for small datasets and medium-size datasets. However, under this batch setting, the computational cost can be too large to be afforded in practice even for medium-size datasets on single-core architectures.

### 5.1 Stochastic optimization

To deal with the above problem, we first note that the terms corresponding to the KL-divergence \( \hat{\mathcal{L}}_{\text{ent}} \) and \( \mathcal{L}_{\text{cross}} \) in Equations (14) and (15) do not depend on the observed data, hence their computational complexity is independent of \( N \). More importantly, we note that \( \hat{\mathcal{L}}_{\text{ell}} \) in Equation (24) decomposes as a sum of expectations over individual datapoints. This makes our inference framework amenable to parallel computation and stochastic optimization (Robbins and Monro, 1951). More precisely, we can rewrite \( \hat{\mathcal{L}}_{\text{elbo}} \) as:

\[ \hat{\mathcal{L}}_{\text{elbo}} = \sum_{n=1}^{N} \left[ \frac{1}{N} \left( \hat{\mathcal{L}}_{\text{ent}} + \mathcal{L}_{\text{cross}} \right) + \hat{\mathcal{L}}^{(n)}_{\text{ell}} \right], \]

where \( \hat{\mathcal{L}}^{(n)}_{\text{ell}} = \frac{1}{S} \sum_{k=1}^{K} \tau_k \sum_{i=1}^{S} \log p(y_n|f^{(n)}_{k,i}, \phi) \), which enables us to apply stochastic optimization techniques such as stochastic gradients descend (SGD, Robbins and Monro, 1951) or ADADELTA (Zeiler, 2012). The complexity of the resulting algorithm is independent of \( N \) and dominated by algebraic operations that are \( \mathcal{O}(M^3) \) in time, where \( M \) is the number of inducing points per latent process. This makes our automated variational inference framework practical for very large datasets.

### 5.2 Reducing the variance of the gradients with control variates

Theorem 1 is fundamental to having a statistical efficient algorithm that only requires sampling from univariate Gaussian distributions (instead of sampling from very high-dimensional Gaussians) for the estimation of the expected log likelihood term and its gradients. However, the variance of the gradient estimates may be too high for the algorithm to work in practice, and variance reduction techniques become necessary. Here we use the well-known technique of control variates (see e.g. Ross, 2006, §8.2), where a new gradient estimate is constructed so as to have the same expectation but lower variance than the original estimate. Our control variate is the so-called score function \( h(f_n) = \nabla_{\lambda_k} \log q_{(n)}(f_n|\lambda_k) \) and full details are given in Appendix F.
5.3 Optimization of the inducing inputs

So far we have discussed optimization of variational parameters ($\lambda$), i.e. the parameters of the approximate posterior ($\{\pi_k, m_k, S_k\}$); covariance hyperparameters ($\theta$); and likelihood parameters ($\phi$). However, as discussed by Titsias (2009), the inducing inputs $\{Z_j\}$ can be seen as additional variational parameters and, therefore, their optimization should be somehow robust to overfitting. As described in the Experiments (Section 9.4.2), learning of the inducing inputs can improve performance, requiring a lower number of inducing variables than when these are fixed. This, of course, comes at an additional computational cost which can be significant when considering high-dimensional input spaces. As with the variational parameters, we study learning of the inducing inputs via gradient-based optimization, for which we use the gradients provided in Appendix E.

6 Dense Posterior and Practical Distributions

In this section we consider the case when the inducing inputs are placed at the training points, i.e. $Z_j = X$ and consequently $M = N$. As mentioned in Section 4.1, we refer to this setting as dense to distinguish it from the case when $M < N$, for which the resulting models are usually called sparse approximations. It is important to realize that not all real datasets are very large and that in many cases the resulting time and memory complexity $O(N^3)$ and $O(N^2)$ can be afforded. Besides the dense posterior case, we also study some particular variational distributions that make our framework more practical, especially in large-scale applications.

6.1 Dense Approximate Posterior

When our posterior is dense the only approximation made is the assumed variational distribution in Equation (6). We will show that, in this case, we recover the objective function in Nguyen and Bonilla (2014a) and that hyper-parameter learning is easier as the terms in the resulting objective function that depend on the hyperparameters do not involve $MC$ estimates. Therefore, their analytical gradients can be used. Furthermore, in the following section, we will provide an efficient exact parametrization of the posterior covariance that reduces the $O(N^2)$ memory complexity to a linear complexity $O(N)$.

We start by looking at the components of $L_{elbo}$ when we make $Z_j = X$, which we can simply obtain by making $K_{zz}^j = K_{xx}^j$ and $M = N$, and realizing that the posterior parameters $\{m_{kj}, S_{kj}\}$ are now $N$-dimensional objects. Therefore, we leave the entropy term $L_{ent}$ in Equation (14) unchanged and we replace all the appearances of $K_{zz}^j$ with $K_{xx}^j$ and all the appearances of $M$ with $N$ for the $L_{cross}$. We refer the reader to Appendix G for details of the equations but it is easy to see that the resulting $L_{ent}$ and $L_{cross}$ are identical to those obtained by Nguyen and Bonilla (2014a, Equations 5 and 6).

For the expected log likelihood term, the generic expression in Equation (16) still applies but we need to figure out the resulting expressions for the approximate posterior parameters in Equations (18) and (19). It is easy to show that the resulting posterior means and covariances are in fact $b_{kj} = m_{kj}$ and $\Sigma_{kj} = S_{kj}$ (see Appendix G for details). This means that in the dense case we simply estimate the $\hat{L}_{ell}$ by using empirical expectations over the
unconstrained variational posterior \( q(f|\lambda) \), with ‘free’ mean and covariance parameters. In contrast, in the sparse case, although these expectations are still computed over \( q(f|\lambda) \), the parameters of the variational posterior \( q(f|\lambda) \) are constrained by Equations (18) and (19) which are functions of the prior covariance and the parameters of the variational distribution over the inducing variables \( q(u|\lambda) \). As we shall see in the following section, this distinction between the dense case and sparse case has critical consequences on hyperparameter learning.

6.1.1 Exact Hyperparameter Optimization

The above insight reveals a remarkable property of the model in the dense case. Unlike the sparse case, the expected log likelihood term does not depend on the covariance hyperparameters, as the expectation of the conditional likelihood is taken over the variational distribution \( q(f|\lambda) \) with ‘free’ parameters. Therefore, only the cross-entropy term \( L_{\text{cross}} \) depends on the hyperparameters (as we also know that \( \nabla_{\theta} L_{\text{ent}} = 0 \)). For this term, as seen in Appendix G.1 and corresponding gradients in Equation (37), we have derived the exact (analytical) expressions for the objective function and its gradients, avoiding empirical MC estimates altogether. This has a significant practical implication: despite using black-box inference, the hyperparameters are optimized wrt the true evidence lower bound (given fixed variational parameters). This is an additional and crucial advantage of our automated inference method over other generic inference techniques (see e.g. Ranganath et al., 2014), which do not exploit knowledge of the prior.

6.1.2 Exact Solution with Gaussian Likelihoods

Another interesting property of our approach arises from the fact that, as we are using MC estimates, \( \hat{L}_{\text{ell}} \) is an unbiased estimator of \( L_{\text{ell}} \). This means that, as the number of samples \( S \) increases, \( L_{\text{elbo}} \) will converge to the true value \( L_{\text{elbo}} \). In the case of a Gaussian likelihood, the posterior over the latent functions is also Gaussian and a variational approach with a full Gaussian posterior will converge to the true parameters of the posterior, see e.g. Opper and Archambeau (2009) and Appendix G.2 for details.

6.2 Practical Variational Distributions

As we have seen in Section 5, learning of all parameters in the model can be done in a scalable way through stochastic optimization for general likelihood models, providing automated variational inference for models with Gaussian process priors. However, the general MoG approximate posterior in Equation (6) requires \( O(M^2) \) variational parameters for each covariance matrix of the corresponding latent process, yielding a total requirement of \( O(QKM^2) \) parameters. This may cause difficulties for learning when these parameters are optimized simultaneously. In this section we introduce two special members of the assumed variational posterior family that improve the practical tractability of our inference framework. These members are a full Gaussian posterior and a mixture of diagonal Gaussians posterior.
6.2.1 Full Gaussian Posterior

This instance considers the case of only one component in the mixture ($K = 1$) in Equation (6), which has a Gaussian distribution with a full covariance matrix for each latent process. Therefore, following the factorization assumption in the posterior across latent processes, the posterior distribution over the inducing variables $u$, and consequently over the latent functions $f$, is a Gaussian with block diagonal covariance, where each block is a full covariance corresponding to that of a single latent function. We thus refer to this approximate posterior as the full Gaussian posterior (FG).

6.2.2 Mixture of Diagonal Gaussians Posterior

Our second practical variational posterior considers a mixture distribution as in Equation (6), constraining each of the mixture components for each latent process to be a Gaussian distribution with diagonal covariance matrix. Therefore, following the factorization assumption in the posterior across latent processes, the posterior distribution over the inducing variables $u$ is a mixture of diagonal Gaussians. However, we note that, as seen in Equations (18) and (19), the posterior over the latent functions $f$ is not a mixture of diagonal Gaussians in the general sparse case. Obviously, in the dense case (where $Z_j = X$) the posterior covariance over $f$ of each component does have a diagonal structure. Henceforth, we will refer to this approximation simply as MoG, to distinguish it from the FG case above, while avoiding the use of additional notation. One immediate benefit of using this approximate posterior is computational, as we avoid the inversion of a full covariance for each component in the mixture.

As we shall see in the following sections, there are additional benefits from the assumed practical distributions and they concern the efficient parametrization of the covariance for both distributions and the lower variance of the gradient estimates for the MoG posterior.

6.3 Efficient Re-parametrization

As mentioned above, one of the main motivations for having specific practical distributions is to reduce the computational overhead due to the large number of parameters to optimize. For the MoG approximation, it is obvious that only $O(M)$ parameters for each latent process and mixture component are required to represent the posterior covariance, hence one obtains an efficient parametrization by definition. However, for the full Gaussian (FG) approximation, naively, one would require $O(M^2)$ parameters. The following theorem states that for settings that require a large number of inducing variables, the FG variational distribution can be represented using a significantly lower number of parameters.

**Theorem 2** The optimal full Gaussian variational posterior can be represented using a parametrization that is linear in the number of observations ($N$).

Before proceeding with the analysis of this theorem, we remind the reader that the general form of our variational distribution in Equation (6) requires $O(KQM^2)$ parameters for the covariance, for a model with $K$ mixture components, $Q$ latent processes and $M$ inducing variables. Nevertheless, for simplicity and because usually $K \ll M$ and $Q \ll M$, we will omit $K$ and $Q$ in in the following discussion.
The proof the theorem can be found in Appendix H, where it is shown that in the FG case the optimal solution for the posterior covariance is given by:

$$\hat{S}_j = K_{zz}^j (K_{zz}^j + K_{zx}^j \Lambda_j K_{xz}^j)^{-1} K_{zz}^j,$$

where $\Lambda_j$ is a $N$-dimensional diagonal matrix. Since the optimal covariance can be expressed in terms of fixed kernel computations and a free set of parameters given by $\Lambda_j$, only $N$ parameters are necessary to represent the posterior covariance. As we shall see below, this parametrization becomes useful for denser models, i.e. for models that have a large number of inducing variables.

6.3.1 Sparse Posterior

In the sparse case the inducing inputs $Z_j$ are at arbitrary locations and, more importantly, $M \ll N$, the number of inducing variables is considerably smaller than the number of training points. The result in Equation (25) implies that if we parameterize $S_j$ in that way, we will need $O(N)$ parameters instead of $O(M^2)$. Of course this is useful when roughly $N < M^2/2$. A natural question arises when we define the number of inducing points as a fraction of the number of training points, i.e. $M = \epsilon N$ with $0 \leq \epsilon \leq 1$, when is such a parameterization useful? In this case, using the alternative parametrization will become beneficial when $\epsilon > \sqrt{2}/N$. To illustrate this, consider for example the MNIST dataset used in our medium-scale experiments in Section 9.4 where $N = 60,000$. This yields a beneficial regime around roughly $\epsilon > 0.006$, which is a realistic setting. In fact, in our experiments on this dataset we did consider sparsity factors of this magnitude. For example, our biggest experiment used $\epsilon = 0.04$. With a naive parametrization of the posterior covariance, we would need roughly $2 \times 10^6$ parameters. In contrast, by using the efficient parametrization we only need $50 \times 10^3$ parameters. As shown below, these gains are greater as the model becomes denser, yielding a dramatic reduction in the number of parameters when having a fully dense Gaussian posterior.

6.3.2 Dense Posterior

In the dense case we have that $Z_j = X$, $\forall j = 1, \ldots, Q$ and consequently $M = N$. Therefore we have that the optimal posterior covariance is given by:

$$\hat{S}_j = \left( (K_{xx}^j)^{-1} + \Lambda_j \right)^{-1}.$$

In principle, the parametrization of the posterior covariance would require $O(N^2)$ parameters for each latent process. However, the above result shows that we can parametrize these covariances efficiently using only $O(N)$ parameters.

6.4 Automatic Variance Reduction with a Mixture-of-Diagonals Posterior

An additional benefit of having a mixture-of-diagonals (MoG) posterior in the dense case is that optimization of the variational parameters will typically converge faster when using a mixture of diagonal Gaussians. This is an immediate consequence of the following theorem.
Theorem 3  When having a dense posterior, the estimator of the gradients wrt the vari-
ational parameters using the mixture of diagonal Gaussians has a lower variance than the full Gaussian posterior's.

The proof is in Appendix I and is simply a manifestation of the Rao-Blackwellization tech-
nique (Casella and Robert, 1996). The theorem is only made possible due to the analytical
tractability of the KL-divergence term \( \mathcal{L}_{kl} \) in the variational objective \( \mathcal{L}_{elbo} \). The practical consequence of this theorem is that optimization will typically converge faster when using
a mixture-of-diagonals Gaussians than when using a full Gaussian posterior approximation.

7 Predictions

Given the general posterior over the inducing variables \( q(u|\lambda) \) in equation (6), the predictive
distribution for a new test point \( x_* \) is given by:

\[
p(y_*|x_*) = \sum_{k=1}^{K} \pi_k \int p(y_*|f_*) \int p(f_*|u)q_k(u|\lambda_k)du df_*.
\]

\[
= \sum_{k=1}^{K} \pi_k \int p(y_*|f_*)q_k(f_*|\lambda_k)df_* , \tag{26}
\]

where \( q_k(f_*|\lambda_k) \) is the predictive distribution over the \( Q \) latent functions corresponding to
mixture component \( k \) given the learned variational parameters \( \lambda = \{ \mathbf{m}_k, \mathbf{S}_k \} \):

\[
q_k(f_*|\lambda_k) = \prod_{j=1}^{Q} \mathcal{N}(f_{*j}; \mu_{kj}^*, \sigma_{kj}^*),
\]

with

\[
\mu_{kj}^* = \kappa_j(x_*, Z_j)(K_{zz})^{-1}m_{kj},
\]

\[
\sigma_{kj}^* = \kappa_j(x_*, x_*) - \kappa_j(x_*, Z_j)((K_{zz})^{-1} - (K_{zz})^{-1}S_{kj}(K_{zz})^{-1})\kappa_j(Z_j, x_*).
\]

Thus, the probability of the test points taking values \( y_* \) (e.g. in classification) in Equation
(26) can be readily estimated via Monte Carlo sampling.

8 Complexity analysis

Throughout this paper we have only considered computational complexity with respect to
the number of inducing points and training points for simplicity. Although this has also
been common practice in previous work (see e.g. Dezfouli and Bonilla, 2015; Nguyen and
Bonilla, 2014a; Hensman et al., 2013; Titsias, 2009), we believe it is necessary to provide
an in-depth complexity analysis of the computational cost of our model. Here we analyze
the computational cost of evaluating the \( \mathcal{L}_{elbo} \) and its gradients once.

We begin by reminding the reader of the dimensionality notation used so far and by
introducing additional notation specific to this section. We analyze the more general case of
stochastic optimization using mini-batches. Let \( K \) be the number of mixture components
in our variational posterior; \( Q \) the number of latent functions; \( D \) the dimensionality of the
input data; $S$ the number of samples used to estimate the required expectations via Monte Carlo; $B$ the number of inputs used per mini-batch; $M$ the number of inducing inputs; and $N$ the number of observations. We note that in the case of batch optimization $B = N$, hence our analysis applies to both the stochastic and batch setting. We also let $T(e)$ represent the computational costs of expression $e$. Furthermore, we assume that the kernel function is simple enough such that evaluating its output between two points is $\mathcal{O}(D)$ and we denote with $T(\log p(y_n | f_n, \phi)) \in \mathcal{O}(L)$ the cost of a single evaluation of the conditional likelihood.

### 8.1 Overall Complexity

While the detailed derivations of the computational complexity are in Appendix J, here we state the main results. Assuming that $K \ll M$, the total computational cost is given by:

$$T(\mathcal{L}_{\text{elbo}}) \in \mathcal{O}(Q(M^2D + BMD + K^2M^3 + KBM^2 + KBSL)),$$

and for diagonal posterior covariances we have:

$$T(\mathcal{L}_{\text{elbo}}) \in \mathcal{O}(Q(M^2D + BMD + M^3 + KBM + KBSL)).$$

We note that it takes $\lceil N/B \rceil$ gradient updates to go over an entire pass of the data. If we assume that $K$ is a small constant, the asymptotic complexity of a single pass over the data does not improve by increasing $B$ beyond $B = M$. Hence if we assume that $B \in \mathcal{O}(M)$ the computational cost of a single gradient update is given by

$$T(\mathcal{L}_{\text{elbo}}) \in \mathcal{O}(QM(MD + M^2 + SL)),$$

for both diagonal and full covariances. If we assume that it takes a constant amount of time to compute the likelihood function between a sample and an output, we see that setting $S \in \mathcal{O}(MD + M^2)$ does not increase the computational complexity of our algorithm. As we shall see in section 9.4, even a complex likelihood function only requires 10,000 samples to approximate it. Since we expect to have more than 100 inducing points in most large scale settings, it is reasonable to assume that the overhead from generated samples will not be significant in most cases.

### 9 Experiments

In this section we analyze the behavior of our model on a wide range of experiments involving small-scale to large-scale datasets. The main aim of the experiments is to evaluate the performance of the model when considering different likelihoods and different dataset sizes. We furthermore elaborate how the performance is affected by the density and location of inducing points, and how the performance of batch and stochastic optimization compare. We first evaluate the algorithm using five small-scale datasets ($N < 1,300$) under different likelihood models and number of inducing points (section 9.3). Then, using medium-scale experiments ($N < 70,000$), we compare stochastic and batch optimization settings, and determine the effect of learning the location of inducing points on the performance (section 9.4). Finally, we apply SAVIGP to a large-scale dataset involving the prediction of airline delays, and we compare the convergence properties of our algorithm to models that leverage full knowledge of the likelihood model (section 9.5). Before proceeding with the experimental set-up, we give details of our implementation which uses GPUs.
9.1 Implementation

We have implemented our SavIGP method in Python and all the code is publicly available at https://github.com/Karl-Krauth/Sparse-GP. Most current mainstream implementations of Gaussian process models do not support GPU computation, and instead opt to offload most of the work to the CPU. For example, neither of the popular packages GPML\(^2\) or GPy\(^3\) provide support for GPUs. This is despite the fact that matrix manipulation operations, which are easily parallelizable, are at the core of any Gaussian process model. In fact, the rate of progress between subsequent GPU models has been much larger than for CPUs, thus ensuring that any GPU implementation would run at an accelerated rate as faster hardware gets released.

With these advantages in mind, we provide an implementation of SavIGP that uses Theano (Al-Rfou et al., 2016), a library that allows users to define symbolic mathematical expressions that get compiled to highly optimized GPU CUDA code. Any operation that involved the manipulation of large matrices or vectors was done in Theano. Most of our experiments were either run on g2.2 AWS instances, or on a desktop machine with an Intel core i5-4460 CPU, 8GB of RAM, and a GTX760 GPU.

Despite using a low-end outdated GPU, we found a time speed-up of 5x on average when we offloaded work to the GPU. For example, in the case of the MNIST-B dataset (used in section 9.4), we averaged the time it took to compute ten gradient evaluations of the \(L_{elbo}\) with respect to the posterior parameters over the entire training set, where we expressed the posterior as a full Gaussian and used a sparsity factor of 0.04. While it took 42.35 seconds, on average, per gradient computation when making use of the CPU only, it took a mere 8.52 seconds when work was offloaded to the GPU. We expect the difference to be even greater given a high-end current-generation GPU.

9.2 Details of the Experiments

Performance measures. We evaluated the performance of the algorithm using non-probabilistic and probabilistic measures according to the type of learning problem we are addressing. The standardized squared error (SSE) and the negative log predictive density (NLPD) were used in the case of continuous-output problems. The error rate (ER) and the negative log probability (NLP) were used in the case of discrete-output problems. In the experiments using the airline dataset, we used the root mean squared error (RMSE) instead of the SSE to be able to compare our method with previous work that used RMSE for performance evaluation.

Experimental settings. In small-scale experiments, inducing points were placed on a subset of the training data in a nested fashion, so that experiments on less sparse models contained the inducing points of the sparser models. In medium-scale and large-scale experiments the location of the inducing points was initialized using the \(k\)-means clustering method. In all experiments the squared exponential covariance function was used.

\(^2\)Available at http://www.gaussianprocess.org/gpml/code/matlab/doc/.
\(^3\)Available at https://github.com/SheffieldML/GPy.
Table 1: Details of the datasets used in the small-scale experiments and their corresponding likelihood models. $N_{\text{train}}, N_{\text{test}}, D$ are the number of training points, test points and input dimensions respectively; ‘likelihood’ is the conditional likelihood used on each problem; and ‘model’ is the name of the model associated with that likelihood. For mining $N_{\text{test}} = 0$ as only posterior inference is done (predictions on a test set are not made for this problem).

| Dataset | $N_{\text{train}}$ | $N_{\text{test}}$ | $D$ | Likelihood $p(y|f)$ | Model                  |
|---------|---------------------|-------------------|-----|---------------------|------------------------|
| MINING  | 811                 | 0                 | 1   | $\lambda^y \exp(-\lambda)/y!$ | Log Gaussian Cox process |
| BOSTON  | 300                 | 206               | 13  | $\mathcal{N}(y; f, \sigma^2)$ | Standard regression     |
| CREEP   | 800                 | 1266              | 30  | $\nabla_y t(y)\mathcal{N}(t(y); f, \sigma^2)$ | Warped Gaussian processes |
| ABALONE | 1000                | 3177              | 8   | $\nabla_y t(y)\mathcal{N}(t(y); f, \sigma^2)$ | Warped Gaussian processes |
| CANCER  | 300                 | 383               | 9   | $1/(1 + \exp(-f))$ | Binary classification    |
| USPS    | 1233                | 1232              | 256 | $\exp(f_c)/\sum_{i=1} \exp(f_i)$ | Multi-class classification |

Optimization methods. Learning the model involves optimizing variational parameters, hyperparameters, likelihood parameters, and inducing inputs. These were optimized iteratively in a global loop. In every iteration, each set of parameters was optimized separately while keeping the other sets of parameters fixed, and this process was continued until the change in the objective function between two successive iterations was less than $10^{-6}$. For optimization in the batch settings, each set of parameters was optimized using L-BFGS, with the maximum number of global iterations limited to 200. In the case of stochastic optimization, we used the ADADELTA method (Zeiler, 2012) with parameters $\epsilon = 10^{-6}$ and a decay rate of 0.95.

Reading the graphs. Results are presented using boxplots and bar/line charts. In the case of boxplots, the lower, middle, and upper hinges correspond to the 25th, 50th, and 75th percentile of the data. The upper whisker extends to the highest value that is within $1.5 \times$ IQR of the top of the box, and the lower whisker extends to the lowest value within $1.5 \times$ IQR of the bottom of the box (IQR is the distance between the first and third quantile). In the case of bar charts, the error bars represent 95% confidence intervals computed over multiple partitions of the dataset.

Model configurations. We refer to the ratio of inducing points to training points as the sparsity factor (SF = $M/N$). For each sparsity factor, three different variations of SAVIGP corresponding to (i) a full Gaussian posterior, (ii) a diagonal Gaussian posterior, and (iii) a mixture of two diagonal Gaussian posteriors were tested, which are denoted respectively by FG, MoG1, and MoG2 in the graphs.

9.3 Small-scale Experiments

We tested SAVIGP on six small-scale datasets with different likelihood models. The datasets are summarized in Table 1, and are the same as those used by Nguyen and Bonilla (2014a). For each dataset, the model was tested five times across different subsets of the data; except
Figure 1: The distributions of $\text{sse}$ and $\text{nlpd}$ for a regression problem with a univariate Gaussian likelihood model on the BOSTON housing dataset. Three approximate posteriors in SAVIGP are used: FG (full Gaussian), MoG1 (diagonal Gaussian), and MoG2 (mixture of two diagonal Gaussians), along with various sparsity factors (SF = $M/N$). The smaller the SF the sparser the model, with SF = 1 corresponding to the dense model. GP corresponds to the performance of exact inference using standard Gaussian process regression.

for the mining dataset where only training data were used for evaluation of the posterior distribution.

9.3.1 Standard regression

The model was evaluated on the BOSTON dataset (Bache and Lichman, 2013), which involves a standard regression problem with a univariate Gaussian likelihood model, i.e., $p(y_n|f_n) = \mathcal{N}(y_n|f_n, \sigma^2)$. Figure 1 shows the performance of SAVIGP for different sparsity factors, as well as the performance of exact Gaussian process inference (GP). As we can see, $\text{sse}$ increases slightly on sparser models. However, the $\text{sse}$ of all the models (FG, MoG1, MoG2) across all sparsity factors are comparable to the performance of exact inference (GP). In terms of $\text{nlpd}$, as expected, the dense (SF = 1) FG model performs exactly like the exact inference method (GP). In the sparse models, $\text{nlpd}$ shows less variation in lower sparsity factors (especially for MoG1 and MoG2), which can be attributed to the tendency of such models to make less confident predictions under high sparsity settings.
Figure 2: The distributions of SSE and NLPD for a warped Gaussian process likelihood model on the ABALONE dataset. Three approximate posteriors in SAVINGP are used: FG (full Gaussian), MoG1 (diagonal Gaussian), and MoG2 (mixture of two diagonal Gaussians), along with various sparsity factors (SF = M/N). The smaller the SF the sparser the model, with SF = 1 corresponding to the non-sparse model. WGP corresponds to the performance of the exact inference method for warped Gaussian process models (Snelson et al., 2003), and GP is the performance of exact inference on a univariate Gaussian likelihood model.

9.3.2 Warped Gaussian process

In warped Gaussian processes, the likelihood function is \( p(y_n|f_n) = \nabla_y t(y_n)N(t(y_n); f_n, \sigma^2) \), for some transformation \( t \). We used the same neural-net style transformation as Snelson et al. (2003), and evaluated the performance of our model on two datasets: CREEP (Cole et al., 2000), and ABALONE (Bache and Lichman, 2013). The results are compared with the performance of exact inference for warped Gaussian processes (WGP) described by Snelson et al. (2003), and also with the performance of exact Gaussian process inference with a univariate Gaussian likelihood model (GP). As shown in Figure 2, in the case of the ABALONE dataset, the performance is similar across all the models (FG, MoG1 and MoG2) and sparsity factors, and is comparable to the performance of the exact inference method for warped Gaussian processes (WGP). In the case of the CREEP dataset (Figure 3), SSE and NLPD are better in denser models. As expected, the performance of the dense (SF = 1) FG model is identical to the performance of WGP. We also note that NLPD in GP has less variation compared to the other models, which can be attributed to its assumption of Gaussian noise.
9.3.3 Binary classification

For binary classification we used the logistic likelihood $p(y_n = 1 | f_n) = 1/(1 + e^{-f_n})$ on the breast cancer dataset (Bache and Lichman, 2013) and compared our model against the expectation propagation (EP) and variational bounds (VBO) methods described by Nickisch and Rasmussen (2008). Results depicted in Figure 4 indicate that the error rate remains comparable across all models (FG, MoG1, MoG2) and sparsity factors, and is almost identical to the error rates obtained by inference using EP and VBO. Interestingly, the NLP shows more variation and generally degrades as the number of inducing points is increased, especially for MoG1 and MoG2 models, which can be attributed to the fact that these denser models are overly confident in their predictions.

9.3.4 Multi-class classification

For multi-class classification, we used the softmax likelihood $p(y_n = c) = e^{-f_c} / \sum_i e^{-f_i}$, and trained the model to classify the digits 4, 7, and 9 from the USPS dataset (Rasmussen and Williams, 2006). We compared our model against a variational inference method (VQ)
Figure 4: Error rates and NLP for binary classification using a logistic likelihood model on the Wisconsin breast cancer dataset. Three approximate posteriors are used: FG (full Gaussian), MoG1 (diagonal Gaussian), and MoG2 (mixture of two diagonal Gaussians), along with various sparsity factors (SF = M/N). The smaller the SF, the sparser the model, with SF = 1 corresponding to the original model without sparsity. The performance of inference using expectation propagation and variational bounds are denoted by EP and VBO respectively.

which represents the ELBO using a quadratic lower bound on the likelihood terms (Khan et al., 2012). As we see in Figure 5, the error rates are slightly lower in denser FG models. We also note that all versions of SavGp achieve comparable error rates to VQ's. Similarly to the binary classification case, NLP shows higher variation with higher sparsity factor, especially in MoG1 and MoG2 models.

9.3.5 Log Gaussian Cox Process

The log Gaussian Cox process (LGCP) is an inhomogeneous Poisson process in which the log-intensity function is a shifted draw from a Gaussian process. Following Murray et al. (2010), we used the likelihood \( p(y_n|f_n) = \frac{\lambda_n^{y_n} \exp(-\lambda_n)}{y_n!} \), where \( \lambda_n = \exp f_n + m \) is the mean of a Poisson distribution and \( m \) is the offset of the log mean. We applied SavGp with the LGCP likelihood on a coal-mining disaster dataset (Jarrett, 1979), which can be seen in Figure 6 (top).

As baseline comparisons, we use hybrid Monte Carlo (HMC) and elliptical slice sampling (ESS) described by Duane et al. (1987) and Murray et al. (2010) respectively. We collected every 100th sample for a total of 10k samples after a burn-in period of 5k samples and used
Figure 5: Classification error rates and NLP for the multi-class classification problem using a softmax likelihood model on the USPS dataset. Three approximate posteriors in SAVIGP are used: FG (full Gaussian), MoG1 (diagonal Gaussian), and MoG2 (mixture of two diagonal Gaussians), along with various sparsity factors (SF = M/N). The smaller the SF the sparser the model, with SF = 1 corresponding to the original model without sparsity. VQ corresponds to a variational inference method, which represents the ELBO as a quadratic lower bound to the likelihood terms.

the Gelman-Rubin potential scale reduction factors (Gelman and Rubin, 1992) to check for convergence.

The bottom plot of Figure 6 shows the mean and variance of the predictions made by SAVIGP, HMC and ESS. We see that the FG model provides similar results across all sparsity factors, which is also comparable to the results provided by HMC and ESS. MoG1 and MoG2 models provide the same mean as the FG models, but tend to underestimate the posterior variance. This under-estimation of posterior variance is well known for variational methods, especially under factorized posteriors. This results are more significant when comparing the running times across models. When using a slower Matlab implementation of our model, for a fair comparison across all methods, SAVIGP was at least two orders of magnitude faster than HMC and one order of magnitude faster than ESS.

9.4 Medium-scale Experiments

In this section we investigate the performance of SAVIGP on four medium-scale problems on the datasets summarized in Table 2. Our goal here is to evaluate the model on medium-scale datasets, to study the effect of optimising the location of inducing points, and to
Figure 6: Top: the coal-mining disasters data. Bottom: the posteriors for a Log Gaussian Cox process on the data when using a \textit{FG} (full Gaussian), \textit{MoG1} (diagonal Gaussian), and \textit{MoG2} (mixture of two diagonal Gaussians), along with various sparsity factors (SF = $M/N$). The smaller the SF the sparser the model, with SF = 1 corresponding to non-sparse model. The solid line is the posterior mean and the shaded area represents 95% confidence intervals. \textit{HMC} and \textit{ESS} correspond to hybrid Monte Carlo and elliptical slice sampling inference methods and are represented by orange dots and dashed green lines respectively.

compare the performance of batch and stochastic optimization of the parameters. The first problem that we consider is multi-class classification of handwriting digits on the \textit{MNIST} dataset using a softmax likelihood model. The second problem uses the same dataset, but the task involves binary classification of odd and even digits using the logistic likelihood model. Therefore we refer to this dataset as the \textit{MNIST} binary dataset (\textit{MNIST-B}). The third problem uses the \textit{SARCOS} dataset (Vijayakumar and Schaal, 2000) and concerns an inverse-dynamics problem for a seven degrees-of-freedom \textit{SARCOS} anthropomorphic robot.
Table 2: Datasets used on the medium-scale experiments. GPRN stands for Gaussian process regression networks; \( N_{\text{train}}, N_{\text{test}}, D \) are the number of training points, test points and input dimensions respectively; \( Q \) is the number of latent processes; \( P \) is the dimensionality of the output data; and ‘model’ is the model associated with the conditional likelihood used.

| Dataset      | \( N_{\text{train}} \)| \( N_{\text{test}} \)| \( D \)| \( Q \)| \( P \)| Model                                      |
|-------------|----------------|----------------|-------|-----|-----|---------------------------------|
| MNIST-B     | 60,000         | 10,000         | 784   | 1   | 1   | Binary classification           |
| MNIST       | 60,000         | 10,000         | 784   | 10  | 10  | Multi-class classification      |
| SARCOS-2    | 44,484         | 4,449          | 21    | 3   | 2   | GPRN (Wilson et al., 2012)      |
| SARCOS      | 44,484         | 4,449          | 21    | 8   | 7   | GPRN (Wilson et al., 2012)      |

Arm. The task is to map from a 21-dimensional input space (7 joint positions, 7 joint velocities, 7 joint accelerations) to the corresponding 7 joint torques. For this multi-output regression problem we use the Gaussian process regression network (GPRN) likelihood model of Wilson et al. (2012), which allows for nonlinear models where the correlation between the outputs can be spatially adaptive. This is achieved by taking a linear combination of latent Gaussian processes, where the weights are also drawn from Gaussian processes. Finally, SARCOS-2 is the same as SARCOS, but the model is learned using only data from joints 4 and 7. For both SARCOS and SARCOS-2, the mean of the SSE and NLPD across joints 4 and 7 are used for performance evaluation. We only consider joints 4 and 7 for SARCOS, despite the fact that predictions are made across all 7 joints to provide a direct comparison with SARCOS-2 and with previous literature (Nguyen and Bonilla, 2014b). We also made use of automatic relevance determination (ARD) for both the SARCOS and SARCOS-2 datasets.

9.4.1 Batch optimization

Binary and multi-class classification. Here we evaluate the performance of batch optimization of model parameters on the MNIST-B and MNIST datasets. We optimized the kernel hyperparameters and the variational parameters, but fixed the inducing point locations using \( k \)-means clustering. Unlike most previous approaches (Ranzato et al., 2006; Jarrett et al., 2009), we did not tune model parameters using the validation set. Instead, we considered the validation set to be part of the training set and used our variational framework to learn all model parameters. As such, the current setting likely provides a lower performance on test accuracy compared to the approaches that use a validation dataset, however our goal is simply to show that we were able to achieve competitive performance in a sparse setting when no knowledge of the likelihood model is used.

We see in Figure 7 that the performance improves significantly with denser models, i.e. when we use a larger number of inducing variables. Overall, our model achieves an accuracy of 98.1% and a mean NLP of 0.062 on MNIST-B with a sparsity factor of 0.04. This is slightly better than the performance obtained by Hensman et al. (2015), who report an accuracy of 97.8% and NLP of 0.069, although they used a smaller number of inducing variables. We will show later that our model also achieves similar performance when using the same sparsity factor as that used by Hensman et al. (2015) when the inducing inputs (i.e. the locations of the inducing variables) are learned.
Figure 7: Error rates and NLP for binary classification on the MNIST-B dataset. We used a full Gaussian (FG) posterior approximation across various sparsity factors (SF = M/N). The smaller the SF the sparser the model.

Figure 8: Error rates and NLP for multi-class classification on the MNIST dataset. We used a full Gaussian (FG) posterior approximation across various sparsity factors (SF = M/N). The smaller the SF the sparser the model.

Figure 8 shows the result on the MNIST dataset. As in the case of MNIST-B, the performance improves with denser models. Overall, SavIGP achieves an error rate of 2.77% at SF = 0.04. This is a significant improvement over the results reported by Gal et al. (2014), in which a separate model was trained for each digit, which achieved an error rate of 5.95%. As a reference, previous literature reports about 12% error rate by linear classifiers and less than 1% error rate by state-of-the-art deep convolutional nets. Our results show that our method reduces the gap between GPs and deep nets while solving the harder problem of full posterior estimation.

Gaussian process regression networks. Figure 9 shows that SARCOS-2 gets a significant benefit from a higher number of inducing points, which is consistent with previous work that found that the performance on this dataset improves as more data is being used to train the model (Vijayakumar and Schaal, 2000). The performance is significantly better than the results reported by Nguyen and Bonilla (2014b), who achieved a mean standardized squared error (MSSE) of 0.2631 and 0.0127 across joints 4 and 7, against our values of
0.0033 and 0.0065 for SF = 0.04. However, we note that their setting was much sparser than ours on joint 4.

Figure 10 shows the performance on the sarcos dataset. Compared to sarcos-2, learning all joints yields a slight increase in MSSE for joints 4 and 7, but also resulted in a significant decrease in NLPD. This shows that the transition from learning only 2 joints to all 7 joints (as in sarcos-2), while unhelpful for point estimates, led to a significantly better posterior density estimate. We also note that we had to make use of 10,000 samples for this dataset to get stable results over different sparsity factors. This is a much larger number than the 100 samples used in mnist. This is likely because the GPRN likelihood model involves multiplication between latent processes, which increases the variance of sampling results by a squared factor, and therefore, it is reasonable to require more samples in this complex likelihood model.

9.4.2 Inducing point learning

We now compare the effect of adaptively learning the inducing inputs, versus initializing them using k-means and leaving them fixed. We look at the performance of our model
under two settings: (i) a low number of inducing variables (SF = 0.001, 0.004) where the inducing inputs are learned, and (ii) a large number of inducing variables (SF = 0.02, 0.04) without learning of their locations.

Figures 11, 12 show the performance of the model under the two settings on MNIST and MNIST-B. We see that learning the location of the inducing variables yields a large gain in performance. In fact, the sparser models with inducing point learning performed similarly to the denser models, despite the fact that the two models differed by an order of magnitude when it came to the number of inducing variables.

Figures 13, 14 show the performance on SARCOS and SARCOS-2. We observe an improvement in performance from learning inducing inputs. However, the improvement is less significant than for MNIST, which shows that different datasets have significantly different sensitivity to the location of the inducing variables. We note that there is a trade-off between the reduction in computational complexity gained by reducing the number of inducing variables and the increased computational cost of calculating inducing input gradients. As such, the advantage of learning the inducing inputs is dataset dependent and it is affected mainly by the input dimensionality ($D$).
9.4.3 Stochastic optimization

In this section we compare the performance of our model after it has been trained in a batch setting versus a stochastic setting. We used ADADELTA as our stochastic optimization algorithm as it requires less hand-tuning than other algorithms such as vanilla stochastic gradient descend (SGD).

Table 3 shows only a slight deterioration in predictive performance on MNIST-B when using stochastic optimization instead of batch optimization. In fact, our exploratory experiments showed that the error metrics could have significantly been reduced by meticulously hand-tuning momentum stochastic gradient descent (SGD). However, our goal was simply to show that our model does not suffer from a large loss in performance when going from a batch to a stochastic setting in medium-scale experiments, without requiring extensive hand tuning of the optimization algorithm. As we shall see in the next section, when batch optimization is not feasible, stochastic optimization in our model performs well when compared to state-of-the-art approaches for GP regression on very large datasets.
Table 3: Error rate (ER) and negative log probability (NLP) obtained with SAVIGP optimized using batch optimization (with L-BFGS) and stochastic optimization (with ADADELTA) on the MNIST-B dataset. The inducing inputs are optimized in both cases.

| Method | Batch SF = 0.001 | Batch SF = 0.004 | Stochastic SF = 0.001 | Stochastic SF = 0.004 |
|--------|-----------------|-----------------|---------------------|---------------------|
| ER     | 3.17%           | 2.12%           | 3.11%               | 2.68%               |
| NLP    | 0.097           | 0.068           | 0.099               | 0.083               |

9.5 Large-scale Experiments

In this section we evaluate SAVIGP on a large-scale dataset that involves prediction of airline delays using a univariate Gaussian likelihood model (Hensman et al., 2013). We note here that this dataset has outliers that can significantly affect the performance of regression algorithms when using metrics such as the RMSE. This has also been pointed out by Das et al. (2015). Additionally, in order to match the application of the algorithms to a realistic setting, evaluation of learning algorithms on this dataset should always consider making predictions in the future. Therefore, unlike Hensman et al. (2013), we did not randomly select the training and test sets, instead we selected the first 700,000 data points starting at a given offset as the training set and the next 100,000 data points as the test set. We generated five training/test sets by setting the initial offset to 0 and increasing it by 200,000 each time.

We used the squared exponential covariance function with automatic relevance determination (ARD), and optimized all the parameters using ADADELTA (Zeiler, 2012). We compare the performance of our model with the stochastic variational inference on GPs method (SVIGP) described by Hensman et al. (2013), which assumes full knowledge of the likelihood function. Our method (SAVIGP) and SVIGP were optimized using ADADELTA with identical settings. We also report the results of Bayesian linear regression with a zero-mean unit-variance prior over the weights (LINEAR) and Gaussian process regression using subsets of the training data (GP1000 and GP2000 using 1,000 and 2,000 datapoints respectively). For each run we fit GP1000 and GP2000 ten times using a randomly selected subset of the training data. Kernel hyper-parameters of GP1000 and GP2000 were optimized using L-BFGS.

Figure 15 shows the performance of SAVIGP and the four baselines. We see that SAVIGP converges at a very similar rate to SVIGP, despite making no assumption about the likelihood model. Furthermore, SAVIGP performs better than all three simple baselines after less than 5 epochs. We note that our results are not directly comparable with those reported by Hensman et al. (2013), since we deal with the harder problem of predicting future events due to the way our dataset is selected.

An additional interesting question about the behavior of our model relates to how well the training objective function correlates to the test error metrics. Figure 16 shows how the performance metrics (RMSE and NLPD) on the test data vary along with the negative evidence lower bound (NELBO). As we can see, changes in the NELBO closely mirror changes.
Figure 15: The RMSE (left) and NLPD (right) of SVIGP (Hensman et al., 2013), and SAVIGP (our method) averaged across all 5 airline experiment runs. The x-axis represents the number of passes over the data (epochs). The figure also shows the performance of GP1000, GP2000 and Bayesian linear regression (LINEAR) after training is completed.

Figure 16: The RMSE (left) and NLPD (right) of SAVIGP on the test data alongside the negative evidence lower bound (NELBO) on the training data, averaged over five runs of the airline experiment.

in RMSE and NLPD in our model, which is an indication that the variational objective generalizes well to test performance when using these metrics.

10 Conclusions and Discussion

We have developed scalable automated variational inference for Gaussian process models (SAVIGP), an inference method for models with Gaussian process (GP) priors, multiple outputs, and nonlinear likelihoods. The method is generally applicable to black-box likelihoods, i.e. it does not need to know the details of the conditional likelihood (or its gradients), only requiring its evaluation as a black-box function.

One of the key properties of this method is that, despite using a flexible variational posterior such as a mixture-of-Gaussians distribution, it is statistically efficient in that it requires only expectations over univariate Gaussian distributions to estimate the evidence lower bound (ELBO) and its gradients.
In order to provide scalability to very large datasets, we have used an augmented prior via the so-called inducing variables, which are prevalent in most sparse GP approximations. This has allowed us to decompose the ELBO as a sum of terms over the datapoints, hence giving way for the application of stochastic optimization and parallel computation.

Our small-scale experiments have shown that SAVIGP can perform as accurately as solutions that have been hard-coded specifically for the conditional likelihood of the problem at hand, even under high sparsity levels.

Our medium-scale experiments also have shown the effectiveness of SAVIGP, when considering problems such as multi-class classification on the MNIST dataset and highly nonlinear likelihoods such as that used in GPRN. On these experiments, we also have analyzed the effect of learning the inducing inputs, i.e. the locations corresponding to the inducing variables, and concluded that doing this can yield significant improvements in performance, reducing the numbers of inducing variables required to achieve similar accuracy by an order of magnitude. Nevertheless, there is generally a trade-off between reducing the time cost gained by having a lower number of inducing inputs and the overhead of carrying out optimization over these locations.

Finally, our large-scale experiments on the problem of predicting airline delays showed that SAVIGP is on par with the state-of-the-art approach for Gaussian process regression on big data (Hensman et al., 2013). In fact, our results show that SAVIGP is slightly better but we attribute those differences to implementation specifics. The important message is that SAVIGP can attain state-of-the-art performance even without exploiting specific knowledge of the likelihood model.

Overall, we believe SAVIGP has the potential to be a powerful tool for practitioners and researchers when devising models for new or existing problems with Gaussian process priors for which variational inference is not yet available. As mentioned in Section 2, we are very much aware of recent developments in the areas of probabilistic programming, stochastic variational inference and Bayesian deep learning. Advances such as those in black-box variational inference (BBVI, Ranganath et al., 2014) and variational auto-encoders (Rezende et al., 2014; Kingma and Welling, 2014) are incredibly exciting for the machine learning community. While the former, BBVI, is somewhat too general to be useful in practice for GP models, the latter (variational auto-encoders) requires specific knowledge of the likelihood and is not a truly black-box method.

Nevertheless, we can enhance our contributions by using results such as the so-called re-parametrization trick in Kingma and Welling (2014) and build more tailored methods that may require a lower number of samples to estimate the expectations in our framework. Indeed, we are extending our framework in this direction, exploring the limits and capabilities of kernel machines such as GP models.

Furthermore, we are also working on extending our models to more complex settings such as structured prediction problems, i.e. where the conditional likelihood is a non-iid model such as a chain or a tree. Such settings provide incredible challenges from the computational and modeling perspectives. For example, how to deal with the exponential increase in the number of parameters of the model, and how to reduce the number of calls to an expensive conditional likelihood model. We believe that the benefits of being Bayesian well outweigh the effort in exploring such challenges.
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A Derivation of the KL-divergence Term

Here we derive the expressions for the terms composing the KL-divergence ($L_{\text{kl}} = L_{\text{ent}} + L_{\text{cross}}$) part of the log-evidence lower bound ($L_{\text{elbo}}$).

The entropy term is given by:

$$L_{\text{ent}}(\lambda) = -\mathbb{E}_{q(u|\lambda)}[\log q(u|\lambda)]$$

$$= -\int \sum_{k=1}^{K} \pi_k q_k(u|m_k, S_k) \log \sum_{\ell=1}^{K} \pi_\ell q_\ell(u|m_\ell, S_\ell) du$$

$$= -\sum_{k=1}^{K} \pi_k \int \mathcal{N}(u; m_k, S_k) \log \sum_{\ell=1}^{K} \pi_\ell \mathcal{N}(u; m_\ell, S_\ell) du$$

$$\geq -\sum_{k=1}^{K} \pi_k \log \int \mathcal{N}(u; m_k, S_k) \sum_{\ell=1}^{K} \pi_\ell \mathcal{N}(u; m_\ell, S_\ell) du$$

$$= -\sum_{k=1}^{K} \pi_k \log \sum_{\ell=1}^{K} \pi_\ell \int \mathcal{N}(u; m_k, S_k) \mathcal{N}(u; m_\ell, S_\ell) du$$

$$= -\sum_{k=1}^{K} \pi_k \log \sum_{\ell=1}^{K} \pi_\ell \mathcal{N}(m_k; m_\ell, S_k + S_\ell) \text{def} = \hat{L}_{\text{ent}},$$

where we have used Jensen’s inequality to bring the logarithm out of the integral from Equations (27) to (28).

The negative cross-entropy term can be computed as:

$$L_{\text{cross}}(\lambda) = \mathbb{E}_{q(u|\lambda)}[\log p(u)] = \sum_{k=1}^{K} \pi_k q_k(u|m_k, S_k) \log p(u) du$$

$$= \sum_{k=1}^{K} \sum_{j=1}^{Q} \pi_k \int \mathcal{N}(u_j; m_{kj}, S_{kj}) \log \mathcal{N}(u_j; 0, K_{zz}^j) du_j$$

$$= \sum_{k=1}^{K} \sum_{j=1}^{Q} \pi_k \left[ \log \mathcal{N}(m_{kj}; 0, K_{zz}^j) - \frac{1}{2} \text{tr} \left( K_{zz}^j \right)^{-1} S_{kj} \right]$$

$$= -\frac{1}{2} \sum_{k=1}^{K} \sum_{j=1}^{Q} \left[ M \log 2\pi + \log |K_{zz}^j| + m_{kj}^T \left( K_{zz}^j \right)^{-1} m_{kj} + \text{tr} \left( K_{zz}^j \right)^{-1} S_{kj} \right].$$
B Proof of Theorem 1

In this section we prove Theorem 1 concerning the statistical efficiency of the estimator of the expected log likelihood and its gradients, i.e. that both can be estimated using expectations over Univariate Gaussian distributions.

B.1 Estimation of $L_{\text{ell}}$

Taking the original expression in Equation (16) we have that:

$$L_{\text{ell}}(\lambda) = E_{q(f|\lambda)}[\log p(y|f, \phi)],$$

$$= \sum_{n=1}^{N} \int f q(f|\lambda) \log p(y_n|f_n, \phi) df,$$

$$= \sum_{n=1}^{N} \int f_{-n} \int f_n q(f_{-n}|f_n) q(f_n) \log p(y_n|f_n, \phi) df_{-n} df_n,$$

$$= \sum_{n=1}^{N} \mathbb{E}_{q(n)}(f_n) [\log p(y_n|f_n, \phi)],$$

$$= \sum_{n=1}^{N} \sum_{k=1}^{K} \pi_k \mathbb{E}_{q_k(n)}(f_n|\lambda_k) [\log p(y_n|f_n, \phi)],$$

where we have applied the linear property of the expectation operation in Equation (29); used $f_{-n}$ to denote all the latent functions except those corresponding to the nth observation, and integrated these out to obtain Equation (30); and used the form of the marginal posterior (MoG in Equation (17)) to get Equation (31).

As described in Section 4.6, $q_{k(n)}(f_n|\lambda_k)$ is a $Q$-dimensional Gaussian with diagonal covariance, hence computation of Equation (31) only requires expectations over univariate Gaussian distributions.

B.2 Estimation of the Gradients of $L_{\text{ell}}$

Denoting the kth term for the nth observation of the expected log likelihood with $L_{\text{ell}}^{(k,n)}$ we have that:

$$L_{\text{ell}}^{(k,n)} = \mathbb{E}_{q_{k(n)}(f_n|\lambda_k)}[\log p(y_n|f_n, \phi)],$$

$$= \int f_{n} q_{k(n)}(f_n|\lambda_k) \log p(y_n|f_n, \phi) df_n.$$

$$\nabla_{\lambda_k} L_{\text{ell}}^{(k,n)} = \int f_{n} q_{k(n)}(f_n|\lambda_k) \nabla_{\lambda_k} \log q_{k(n)}(f_n|\lambda_k) \log p(y_n|f_n, \phi) df_n,$$

$$= \mathbb{E}_{q_{k(n)}(f_n|\lambda_k)} [\nabla_{\lambda_k} \log q_{k(n)}(f_n|\lambda_k) \log p(y_n|f_n, \phi)],$$

for $\lambda_k \in \{m_k, S_k\}$, and for the mixture proportions the gradients can be estimated straight-forwardly using Equation (22). We have used in Equation (32) the fact that $\nabla_x f(x) =$
\( f(x) \nabla_x \log f(x) \) for any nonnegative function \( f(x) \). We see that our resulting gradient estimate has an analogous form to that obtained in Equation (31), hence its computation only requires expectations over univariate Gaussian distributions.

C Gradients of the Evidence Lower Bound wrt Variational Parameters

Here we specify the gradients of the log-evidence lower bound (\( \mathcal{L}_{\text{elbo}} \)) wrt variational parameters. For the covariance, we consider \( S_{kj} \) of general structure but also give the updates when \( S_{kj} \) is a diagonal matrix, denoted with \( \tilde{S}_{kj} \).

C.1 KL-divergence Term

Let \( K_{zz} \) be the block-diagonal covariance with \( Q \) blocks \( K^j_{zz} \), \( j = 1, \ldots, Q \). Additionally, let us assume the following definitions:

\[
C_{kl} \overset{\text{def}}{=} S_k + S_\ell, \quad (34)
\]

\[
N_{k\ell} \overset{\text{def}}{=} \mathcal{N}(m_k; m_\ell, C_{kl}),
\]

\[
z_k \overset{\text{def}}{=} \sum_{\ell=1}^K \pi_\ell N_{k\ell}.
\]

The gradients of \( \mathcal{L}_{kl} \) wrt the posterior mean and posterior covariance for component \( k \) are:

\[
\nabla_{m_k} \mathcal{L}_{\text{cross}} = -\pi_k K_{zz}^{-1} m_k,
\]

\[
\nabla_{S_k} \mathcal{L}_{\text{cross}} = -\frac{1}{2} \pi_k K_{zz}^{-1}, \text{ and for diagonal covariance we have:}
\]

\[
\nabla_{\pi_k} \mathcal{L}_{\text{cross}} = -\frac{1}{2} \pi_k \text{diag}(K_{zz}^{-1}),
\]

\[
\nabla_{m_k} \hat{\mathcal{L}}_{\text{ent}} = \frac{1}{2} \sum_{\ell=1}^Q [M \log 2\pi + \log |K^j_{zz}| + m^T_{kj} (K^j_{zz})^{-1} m_{kj} + \text{tr} (K^j_{zz})^{-1} S_{kj}],
\]

where we note that we compute \( K_{zz}^{-1} \) by inverting the corresponding blocks \( K^j_{zz} \) independently. The gradients of the entropy term wrt the variational parameters are:

\[
\nabla_{m_k} \hat{\mathcal{L}}_{\text{ent}} = \pi_k \sum_{\ell=1}^K \pi_\ell \left( \frac{N_{k\ell}}{z_k} + \frac{N_{\ell k}}{z_\ell} \right) C^{-1}_{kl} (m_k - m_\ell),
\]

\[
\nabla_{S_k} \hat{\mathcal{L}}_{\text{ent}} = \frac{1}{2} \pi_k \sum_{\ell=1}^K \pi_\ell \left( \frac{N_{k\ell}}{z_k} + \frac{N_{\ell k}}{z_\ell} \right) \left[ C^{-1}_{kl} - C^{-1}_{kl} (m_k - m_\ell) (m_k - m_\ell)^T C^{-1}_{kl} \right],
\]

and for diagonal covariance we have:

\[
\nabla_{S_k} \hat{\mathcal{L}}_{\text{ent}} = \frac{1}{2} \pi_k \sum_{\ell=1}^K \pi_\ell \left( \frac{N_{k\ell}}{z_k} + \frac{N_{\ell k}}{z_\ell} \right) \left[ \tilde{C}^{-1}_{kl} - \tilde{C}^{-1}_{kl} \text{diag} ((m_k - m_\ell) \odot (m_k - m_\ell)) \tilde{C}^{-1}_{kl} \right],
\]
\[ \nabla_{\pi_k} \hat{L}_{\text{ent}} = -\log z_k - \sum_{\ell=1}^{K} \pi_{\ell} \frac{N_{k\ell}}{z_{\ell}}, \]

where \( \hat{C}_{kl} \) is the diagonal matrix defined analogously to \( C_{kl} \) in Equation (34) and \( \odot \) is the Hadamard product.

### C.2 Expected Log Likelihood Term

Monte Carlo estimates of the gradients of the expected log likelihood term are:

\[
\nabla_{m_{kj}} \hat{L}_{\text{ell}} = \pi_k \sum_{n=1}^{N} k_{zn}[\Sigma_{k(n)}]^{-1}_{j,j} \sum_{i=1}^{S} \left[ f_{nj}^{(k,i)} - [b_{k(n)}]_j \right] \log p(y_n|f_n^{(k,i)}),
\]

\[
\nabla_{s_{kj}} \hat{L}_{\text{ell}} = \frac{\pi_k}{2S} \sum_{n=1}^{N} (a_{jn}a_{jn}^T) \sum_{i=1}^{S} \left[ (\Sigma_{k(n)})_{j,j}^{-2} \left( f_{nj}^{(k,i)} - [b_{k(n)}]_j \right)^2 - (\Sigma_{k(n)})_{j,j}^{-1} \right] \log p(y_n|f_n^{(k,i)}),
\]

\[
\nabla_{\pi_k} \hat{L}_{\text{ell}} = \frac{1}{S} \sum_{n=1}^{N} \sum_{i=1}^{S} \log p(y_n|f_n^{(k,i)}),
\]

and for diagonal covariance \( \hat{S}_{kj} \) we replace \( (a_{jn}a_{jn}^T) \) with \( \text{diag}(a_{jn} \odot a_{jn}) \) in Equation (35), where \( \odot \) is the Hadamard product and \( \text{diag}(v) \) takes the input vector \( v \) and outputs a matrix with \( v \) on its diagonal. We have also defined above \( k_{zn} \overset{\text{def}}{=} \kappa_j(Z_{j}, x_n) \), i.e. the vector obtained from evaluating the covariance function \( j \) between the inducing points \( Z_j \) and datapoint \( x_n \).

### D Gradients of the Evidence Lower Bound wrt Covariance Hyperparameters and Likelihood Parameters

Here we give the gradients of the variational objective (\( \mathcal{L}_{\text{elbo}} \)) wrt the covariance hyperparameters and, when required, the conditional likelihood parameters. We note here that our method does not require gradients of the conditional likelihood \( p(y|f, \phi) \) wrt the latent functions \( f \). However, if the conditional likelihood is parametrized by \( \phi \) and point-estimates of these parameters are needed, these can also be learned in our variational framework.

#### D.1 Covariance Hyperparameters

The gradients of the terms in the KL-divergence part of \( \mathcal{L}_{\text{elbo}} \) wrt a covariance hyperparameter \( \theta_j \) are:

\[
\nabla_{\theta_j} \hat{L}_{\text{ent}} = 0,
\]

\[
\nabla_{\theta_j} \mathcal{L}_{\text{cross}} = -\frac{1}{2} \sum_{k=1}^{K} \pi_k \text{tr} \left[ (K_{zz})^{-1} \nabla_{\theta_j} K_{zz}^j - (K_{zz}^j)^{-1} \nabla_{\theta_j} K_{zz}^j (K_{zz}^j)^{-1} \left( m_{kj}m_{kj}^T + S_{kj} \right) \right].
\]
For the $\widehat{L}_{\text{ell}}$ we have that:

$$\nabla_{\theta_j} \widehat{L}_{\text{ell}} = \frac{1}{2} \sum_{n=1}^{N} \sum_{k=1}^{K} \frac{\pi_k}{\sum_{i=1}^{S} \pi_{k(i)}} q_{k(n)}(f_n | A_k) \left[ \nabla_{\theta_j} \log q_{k(n)}(f_n | A_k) \log p(y_n | f_n) \right],$$

and computing the corresponding gradient we obtain:

$$\nabla_{\theta_j} \widehat{L}_{\text{ell}} = -\frac{1}{2} \sum_{n=1}^{N} \sum_{k=1}^{K} \frac{\pi_k}{\sum_{i=1}^{S} \pi_{k(i)}} q_{k(n)}(f_n | A_k) \left[ \left( \nabla_{\theta_j} \log q_{k(n)}(f_n | A_k) \right) \log p(y_n | f_n) \right]$$

for which we need:

$$\nabla_{\theta_j} \pi_k = \frac{\pi_k}{\sum_{i=1}^{S} \pi_{k(i)}},$$

$$\nabla_{\theta_j} \log q_{k(n)}(f_n | A_k) = \left( \nabla_{\theta_j} q_{k(n)}(f_n | A_k) \right) \log p(y_n | f_n),$$

$$\nabla_{\theta_j} q_{k(n)}(f_n | A_k) = \sum_{i=1}^{S} \pi_{k(i)} q_{k(i)}(f_n | A_{k(i)}) \left( \nabla_{\theta_j} \log q_{k(i)}(f_n | A_{k(i)}) \right),$$

and as in the main text we have defined $a_{j,n} \equiv [A_j]_{n,n}$, i.e. the vector corresponding to the $n$th column of $A_j$. Furthermore, as in the previous section, $k_{zn} \equiv \kappa_j(Z_j, x_n)$.

### D.2 Likelihood Parameters

Since the terms in the KL-divergence do not depend on the likelihood parameters we have that $\nabla_{\phi} \widehat{L}_{\text{ent}} = \nabla_{\phi} \widehat{L}_{\text{cross}} = 0$. For the gradients of the expected log likelihood term we have that:

$$\nabla_{\phi} \widehat{L}_{\text{ell}} = \frac{1}{S} \sum_{n=1}^{N} \sum_{k=1}^{K} \frac{\pi_k}{\sum_{i=1}^{S} \pi_{k(i)}} \nabla_{\phi} \log p(y_n | f_n^{(k,i)}),$$

where $\{f_n^{(k,i)} \sim \mathcal{N}(f_n^{(k)}, \Sigma_k(n))$, for $k = 1, \ldots, K$ and $i = 1, \ldots, S$. 

### E Gradients of the Log-Evidence Lower Bound wrt Inducing Inputs

These gradients can be obtained by using the same expression as the gradients wrt covariance hyperparameters in Appendix D.1, considering the inducing inputs as additional hyperparameters of the covariances. Therefore, we rewrite the gradients above keeping in
mind that \( \theta_j \) is an element of inducing input \( Z_j \), hence dropping those gradient terms that do not depend on \( Z_j \). As before we have that \( \nabla_{\theta_j} \tilde{\ell}_{\text{ent}} = 0 \).

\[
\nabla_{\theta_j} \mathcal{L}_{\text{cross}} = -\frac{1}{2} \sum_{k=1}^{K} \pi_k \text{tr} \left\{ \left[ (K_{zz}^j)^{-1} - (K_{zz}^j)^{-1} (m_{kj} m_{kj}^T + S_j) (K_{zz}^j)^{-1} \right] \nabla_{\theta_j} K_{zz}^j \right\} \tag{37}
\]

Similarly, for the gradients of \( \tilde{\ell}_{\text{ell}} \) in Equation (36) we have that:

\[
\nabla_{\theta_j} [b_{k(n)}]_j = \nabla_{\theta_j} (k_{zn}^j)^T (K_{zz}^j)^{-1} m_{kj} - a_{jn}^T \nabla_{\theta_j} K_{zz}^j (K_{zz}^j)^{-1} m_{kj},
\]

and

\[
\nabla_{\theta_j} [\Sigma_{k(n)}]_{jj} = \nabla_{\theta_j} (k_{zn}^j)^T \left(- (K_{zz}^j)^{-1} k_{zn}^j - a_{jn} + 2 (K_{zz}^j)^{-1} S_{kj} a_{jn} \right) + a_{jn}^T \nabla_{\theta_j} K_{zz}^j (K_{zz}^j)^{-1} k_{zn}^j - 2 a_{jn}^T \nabla_{\theta_j} K_{zz}^j (K_{zz}^j)^{-1} S_{kj} a_{jn}.
\]

From the equations above, we see that in the computation of the gradients of \( \tilde{\ell}_{\text{ell}} \) there are two types of terms. The first term is of the form:

\[
\nabla_{\theta_l} t_{pn}^{(1)} \overset{\text{def}}{=} v_n^T \nabla_{\theta_l} k_{zn}^j w_n,
\]

where we have dropped the index \( j \) on the LHS of the equation for simplicity in the notation and \( v_n, w_n \) are \( M \)-dimensional vectors. Let \( V \) and \( W \) be the \( M \times N \) matrices corresponding to the \( N \) vectors \( \{v_n\} \) and \( \{w_n\} \), respectively. Furthermore, let us assume \( \tilde{Z}_j^{(d)} \) is the \( M \times M \) matrix of all pairwise differences on dimension \( d \) of all inducing points divided by the squared length-scale of the dimension \( \ell_d^2 \),

\[
[\tilde{Z}_j^{(d)}]_{o,p} = \frac{[Z_j]_{o,d} - [Z_j]_{p,d}}{\ell_d^2}.
\]

Hence, in the case of the squared exponential covariance function the above gradient can be calculated as follows (for all data points):

\[
\nabla_{[Z_j]_{.,d}} t^{(1)} = -((\tilde{Z}_j^{(d)} \odot K_{zz}^j) W) \odot V - ((\tilde{Z}_j^{(d)} \odot K_{zz}^j) V) \odot W,
\]

where \( \nabla_{[Z_j]_{.,d}} t^{(1)} \) is the \( M \times N \) matrix of gradients corresponding to dimension \( d \) for all \( m = 1, \ldots, M \) and \( n = 1, \ldots, N \).

Similarly, the second type of term is of the form:

\[
\nabla_{\theta_l} t_{pn}^{(2)} \overset{\text{def}}{=} v_n^T \nabla_{\theta_l} k_{zn}^j,
\]

where \( v_n \) and \( V \) are defined as before. The gradients in the above expression wrt to the inducing points (for all datapoints or a mini-batch) can be calculated as follows:

\[
\nabla_{[Z_j]_{.,d}} t^{(2)} = -(K_{xz} \odot V^T \odot \tilde{X}^{(d)})^T,
\]

where in the above equation \( \tilde{X}^{(d)} \) is the \( N \times M \) matrix of all pairwise differences on dimension \( d \) between all datapoints and inducing points divided by the squared length-scale of the dimension \( \ell_d^2 \):

\[
[\tilde{X}^{(d)}]_{o,p} = \frac{[X]_{o,d} - [Z_j]_{p,d}}{\ell_d^2}.
\]
F  Control Variates

We use control variates (see e.g. Ross, 2006, §8.2) to reduce the variance of the gradient estimates. In particular, we are interested in estimating gradients of the form:

\[ \nabla \lambda_k \mathbb{E}_{q_k(n)(f_n | \lambda_k)} \left[ \log p(y_n | f_n) \right] = \mathbb{E}_{q_k(n)(f_n | \lambda_k)} \left[ g(f_n) \right], \]

with

\[ g(f_n) = \nabla \lambda_k \log q_k(n)(f_n | \lambda_k) \log p(y_n | f_n), \]

where the expectations are computed using samples from \( q_k(n)(f_n | \lambda_k) \), which depends on the variational parameter \( \lambda_k \). As suggested by Ranganath et al. (2014), a sensible control variate is the so-called score function

\[ h(f_n) = \nabla \lambda_k \log q_k(n)(f_n | \lambda_k), \]

whose expectation is zero. Hence, the function:

\[ \tilde{g}(f_n) = g(f_n) - \hat{a} h(f_n), \]

has the same expectation as \( g(f_n) \) but lower variance when \( \hat{a} \) is given by:

\[ \hat{a} = \frac{\text{Cov}[g(f_n), h(f_n)]}{\text{V}[h(f_n)]}, \]

where \( \text{Cov}[g(f_n), h(f_n)] \) is the covariance between \( g(f_n) \) and \( h(f_n) \); \( \text{V}[h(f_n)] \) is the variance of \( h(f_n) \); and both are estimated using samples from \( q_k(n)(f_n | \lambda_k) \). Therefore, our corrected gradient is given by:

\[ \tilde{\nabla} \lambda_k \mathbb{E}_{q_k(n)(f_n | \lambda_k)} \left[ \log p(y_n | f_n) \right] \overset{\text{def}}{=} \mathbb{E}_{q_k(n)(f_n | \lambda_k)} \left[ \tilde{g}(f_n) \right] = \mathbb{E}_{q_k(n)(f_n | \lambda_k)} \left[ \nabla \lambda_k \log q_k(n)(f_n | \lambda_k)(\log p(y_n | f_n) - \hat{a}) \right]. \]

G  Derivations for the Dense Posterior Case

In this section we consider the case of having a dense variational posterior, i.e. not using the so-called sparse approximations. We derive the expressions for the components of the log-evidence lower bound and show that the posterior parameters of the variational distribution \( q(f | \lambda) \) are, in fact, of ‘free’-form. Furthermore, we analyze the case of a Gaussian likelihood showing that, in the limit of a large number of samples, our estimates converge to the exact analytical solution.

G.1  Evidence Lower Bound

Here we show the derivations of all the terms in \( \mathcal{L}_{\text{elbo}} \) when considering the dense case, i.e. \( M = N \) and \( Z_j = X \). As described in the main text, the resulting changes to these terms can be obtained by replacing \( M \) with \( N \) and \( K_{zz}^j \) with \( K_{xx}^j \). Hence, for the terms in the KL-divergence part of \( \mathcal{L}_{\text{elbo}} \) we have that:

\[ \hat{\mathcal{L}}_{\text{ent}}(\lambda) = -\sum_{k=1}^{K} \pi_k \log \sum_{\ell=1}^{K} \pi_{\ell} N(m_k; m_{\ell}, S_k + S_{\ell}), \quad (38) \]
\[
\mathcal{L}_{\text{cross}}(\lambda) = -\frac{1}{2} \sum_{k=1}^{K} \sum_{j=1}^{Q} \left[ N \log 2\pi + \log |K_{xx}| + m_{kj}^T (K_{xx}^{-1}) m_{kj} + \text{tr} (K_{xx}^{-1}) S_{kj} \right]
\]

where we recall that now \( m_{kj} \) and \( S_{kj} \) are \( N \)-dimensional objects. For the \( \hat{L}_{\text{ell}} \) we still need to compute empirical expectations over the variational distribution \( q(f|\lambda) \), where the corresponding parameters are given by:

\[
\begin{align*}
b_{kj} &= A_j m_{kj} = K_{xx}^j (K_{xx}^j)^{-1} m_{kj} = m_{kj}, \\
\Sigma_{kj} &= \tilde{K}_j - A_j K_{xx}^j + A_j S_{kj} A_j \\
&= K_{xx}^j - K_{xx}^j (K_{xx}^j)^{-1} K_{xx}^j + K_{xx}^j (K_{xx}^j)^{-1} S_{kj} K_{xx}^j (K_{xx}^j)^{-1} \\
&= S_{kj}.
\end{align*}
\]

**G.2 Gaussian Likelihoods**

Consider the case of a single output and a single latent function \((Q = P = 1)\) with a Gaussian conditional likelihood and a single full Gaussian variational posterior \((K = 1)\).

\[
p(y|f) = \mathcal{N}(y; f, \sigma^2 I), \quad q(f|\lambda) = \mathcal{N}(f; m, S).
\]

The entropy and the cross entropy terms are given in Equations (38) and (39) with \( Q = K = 1 \). The expect log likelihood term can be determined analytically,

\[
\mathcal{L}_{\text{ell}}(\lambda) = \sum_{n=1}^{N} \mathbb{E}_{q(f|\lambda)}[\log p(y_n|f_n, \phi)],
\]

\[
= \log \mathcal{N}(y; m, \sigma^2 I) - \frac{1}{2\sigma^2} \text{tr} S.
\]

Hence the gradients of \( \mathcal{L}_{\text{elbo}} \) can also be determined analytically, yielding the optimal mean posterior as follows:

\[
\nabla m \mathcal{L}_{\text{elbo}} = K_{xx}^{-1} m + \frac{1}{\sigma^2} (y - m) = 0
\]

\[
\left( K_{xx}^{-1} + \frac{1}{\sigma^2} \right) m = \frac{1}{\sigma^2} y
\]

\[
\hat{m} = (\sigma^2 K_{xx}^{-1} + I)^{-1} y
\]

\[
= K_{xx} (K_{xx} + \sigma^2 I)^{-1} y.
\]

Similarly for the posterior covariances we have that:

\[
\nabla S \mathcal{L}_{\text{elbo}} = \frac{1}{2} S^{-1} - \frac{1}{2} K_{xx}^{-1} - \frac{1}{2\sigma^2} I = 0
\]

\[
\hat{S} = \left( K_{xx}^{-1} + \frac{1}{\sigma^2} I \right)^{-1}
\]

(40)
\begin{align*}
&= K_{xx} - K_{xx} \frac{1}{\sigma^2} \left( I + K_{xx} \frac{1}{\sigma^2} \right)^{-1} K_{xx} \\
&= K_{xx} - K_{xx} (K_{xx} + \sigma^2 I)^{-1} K_{xx},
\end{align*}

where we have used Woodbury’s formula to go from Equation (40) to Equation (41). The Equations above for the optimal posterior mean and posterior covariance, \( \hat{m} \) and \( \hat{S} \), are the exact analytical expressions for regressions with Gaussian process priors and isotropic noise likelihoods, see Rasmussen and Williams (2006, §2.2).

\section{Efficient Parametrization}

In this appendix we prove Theorem 2, showing that it is possible to obtain an efficient parameterization of the posterior covariances when using a full Gaussian approximation. In this case we have that:

\[
\nabla_S \mathcal{L}_{\text{elbo}} = \nabla_S \mathcal{L}_{\text{ent}} + \nabla_S \mathcal{L}_{\text{cross}} + \nabla_S \mathcal{L}_{\text{ell}}
= -\frac{1}{2} (K_{zz}^j)^{-1} + \frac{1}{2} S_j^{-1} + \sum_{n=1}^N \nabla_S \mathcal{E}_{q(\omega)}(f_n) [\log p(y_n|f_n)]
\]

Setting the gradients to zero we have that

\[
\frac{1}{2} S_j^{-1} = \frac{1}{2} (K_{zz}^j)^{-1} + \frac{1}{2} (K_{zz}^j)^{-1} \sum_{n=1}^N k_{zn}^j \lambda_{jn} (k_{zn}^j)^T (K_{zz}^j)^{-1},
\]

where

\[
\lambda_{jn} = -2 \frac{d\bar{\ell}_n}{d\Sigma_j}_{n,n}
\]

\[
\bar{\ell}_n = \mathcal{E}_{q(\omega)}(f_n) [\log p(y_n|f_n)].
\]

Therefore, the optimal solution for the posterior covariance is given by:

\[
\hat{S}_j = ((K_{zz}^j)^{-1} + (K_{zz}^j)^{-1} K_{zx}^j A_j K_{xz}^j (K_{zz}^j)^{-1})^{-1},
\]

\[
= K_{zz}^j (K_{zz}^j + K_{zx}^j A_j K_{xz}^j)^{-1} K_{zz}^j,
\]

where \( A_j \) is a \( N \times N \) diagonal matrix with \( \{\lambda_{jn}\}_{n=1}^N \) on the diagonal.

\section{Lower Variance of Mixture-of-Diagonals Posterior}

In this section we prove Theorem 3. First we review Rao-Blackwellization (Casella and Robert, 1996), which is also known as partial averaging or conditional Monte Carlo. Suppose we want to estimate \( V = \mathbb{E}_{p(X,Y)}[h(X,Y)] \) where \( (X,Y) \) is a random variable with
probability density \( p(x, y) \) and \( h(X, Y) \) is a random variable that is a function of \( X \) and \( Y \). It is easy to see that

\[
E_{p(x,y)}[h(X,Y)] = \int p(x,y)h(x,y)dxdy
\]

\[
= \int p(y)p(x|y)h(x,y)dxdy
\]

\[
= \mathbb{E}_{p(y)}[\hat{h}(Y)], \text{ with } \hat{h}(Y) = E_{p(x|y)}[h(X,Y)|Y],
\]

and, from the conditional variance formula,

\[
\nabla[\hat{h}(Y)] < \nabla[h(X,Y)].
\]

Therefore when \( \hat{h}(Y) \) is easy to compute, it can be used to estimate \( V \) with a lower variance than the original estimator. When \( p(x,y) = p(x)p(y) \), then \( \hat{h}(Y) \) is simplified to

\[
\hat{h}(Y = y) = \int p(x)h(x,y)d = E_{p(x)}[h(X,Y)|Y].
\]

We apply Rao-Blackwellization to our problem with \( f_n \) playing the role of the conditioning variable \( Y \) and \( f_{-n} \) playing the role of \( X \), where \( f_{-n} \) denotes \( f \) excluding \( f_n \). We note that this Rao-Blackwellization analysis is only applicable to the dense case with a mixture of diagonals posterior, as it satisfies the independence condition, i.e. \( p(x,y) = p(x)p(y) \).

First, we express the gradient of \( \lambda_{k(n)} \) as an expectation by interchanging the integral and gradient operators giving

\[
\nabla_{\lambda_{k(n)}} E_{q_k(f|\lambda_k)}[\log p(y|\phi, \phi)] = E_{q_k(f|\lambda_k)}[\nabla_{\lambda_{k(n)}} \log q_k(f|\lambda_k) \log p(y|f, \phi)].
\]

The Rao-Blackwellized estimator is thus

\[
\hat{h}(f_{-n}) = \int q(f_{-n}) \nabla_{\lambda_{k(n)}} \log q_k(f|\lambda_k) \log p(y|f) df_{-n}.
\]

\[
= \int q(f_{-n}) \nabla_{\lambda_{k(n)}} \log q_k(f_n|\lambda_{k(n)}) \log p(y|f) df_{-n}.
\]

\[
= \nabla_{\lambda_{k(n)}} \log q_k(f_n|\lambda_{k(n)}) \int q(f_{-n}) [\log p(y_n|f_{-n}) + \log p(y_{-n}|f_{-n})] df_{-n}.
\]

\[
= \nabla_{\lambda_{k(n)}} \log q_k(f_n|\lambda_{k(n)}) [\log p(y_n|f_{-n}) + C], \tag{42}
\]

where we have used the factorization of the conditional likelihood in Equation (42), and where \( C \) is a constant w.r.t \( f_{-n} \). This gives the Rao-Blackwellized gradient,

\[
\nabla_{\lambda_{k(n)}} E_{q_k(f|\lambda_k)}[\log p(y|f)] = E_{q_k(f_n|\lambda_{k(n)})}[\hat{h}(f_{-n})].
\]

\[
= E_{q_k(f_n|\lambda_{k(n)})}[\nabla_{\lambda_{k(n)}} \log q_k(f_n|\lambda_{k(n)}) \log p(y_n|f_n)],
\]

where we have used the fact that \( E_q[\nabla \log q] = 0 \) for any \( q \). We see then that the expression above is exactly the gradient obtained in Equation (33).
J Details of Complexity analysis

Here we give the details of the computational complexity of a single evaluation of the $L_{\text{elbo}}$ and its gradients. Let $A$ be some matrix of size $p \times q$, $B$ be some matrix of size $q \times r$, $C$ be some matrix of size $q \times p$, and $D$ be a square matrix of size $s \times s$. Then we assume that $T(AB) \in \mathcal{O}(pqr)$, $T(D^{-1}) \in \mathcal{O}(s^3)$, $T(|D|) \in \mathcal{O}(s^3)$, $T(\text{diag}(AC)) \in \mathcal{O}(pq)$, and $T(\text{tr}(AC)) \in \mathcal{O}(pq)$.

J.1 Interim values

We first provide an analysis of the cost of various interim expressions that get re-used throughout the computation of the $L_{\text{elbo}}$. The interim expressions include: various values computed from the kernel functions ($K_{zz}^j$, $(K_{zz}^j)^{-1}$, $|K_{zz}^j|$, $k_{zn}^j$), values used to determine the shape of the Gaussian distributions from which we sample ($[b_k^j(n)]$, $[\Sigma_k^j(n)]_{j,j}$), and various intermediate values ($[\widetilde{K}_j^j n,n]$, $a_{jn}$). We note that expressions with a subscript or superscript of $j$, $n$, or $k$ are dependent on $Q$, $B$, and $K$ respectively. For example, in the case of $k_{zn}^j$, we actually need to consider the cost of $Q \times B$ vectors of size $M$.

As stated in the main text, we assume that the kernel function is simple enough such that evaluating its output between two points is in $\mathcal{O}(D)$. Hence the cost of evaluating interim values is given by:

$$
T(K_{zz}^j) \in \mathcal{O}(QM^2D),
T((K_{zz}^j)^{-1}) \in \mathcal{O}(QM^3),
T(|K_{zz}^j|) \in \mathcal{O}(QM^3),
T(k_{zn}^j) \in \mathcal{O}(QBM^3),
T([b_{k(n)}^j]_j) \in \mathcal{O}(KQBM),
T([\Sigma_{k(n)}^j]_{j,j}) \in \mathcal{O}(KQBM^2),
T([\widetilde{K}_j^j]_{n,n}) \in \mathcal{O}(QBM),
T(a_{jn}) \in \mathcal{O}(QBM^2),
$$

and for diagonal covariance we have:

$$
T([\Sigma_{k(n)}^j]_{j,j}) \in \mathcal{O}(KQBM^2).
$$

The computational cost of evaluating all interim values is thus in $\mathcal{O}(QM(MD + M^2 + BD + KBM))$ for the case of full covariance and in $\mathcal{O}(QM(MD + M^2, BD, KB))$ for diagonal covariances.

J.2 Analysis of the $L_{\text{kl}}$

The computational complexity of terms in the KL-divergence and its gradients is given by:

$$
T(L_{\text{cross}}) \in \mathcal{O}(KQM^2),
T(L_{\text{ent}}) \in \mathcal{O}(K^2QM^3),
$$

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\[ T(\nabla_m \mathcal{L}_{\text{cross}}) \in \mathcal{O}(KQM^2) , \]
\[ T(\nabla_S \mathcal{L}_{\text{cross}}) \in \mathcal{O}(KQM^2) , \]
\[ T(\nabla_{\pi_k} \mathcal{L}_{\text{cross}}) \in \mathcal{O}(KQM^2) , \]
\[ T(\nabla_m \mathcal{L}_{\text{ent}}) \in \mathcal{O}(K^2QM^3) , \]
\[ T(\nabla_S \mathcal{L}_{\text{ent}}) \in \mathcal{O}(K^2QM^3) , \]
\[ T(\nabla_{\pi_k} \mathcal{L}_{\text{ent}}) \in \mathcal{O}(K^2QM^3) , \]

and for diagonal covariance we have:
\[ T(\nabla_m \mathcal{L}_{\text{cross}}) \in \mathcal{O}(KQM) , \]
\[ T(\nabla_m \mathcal{L}_{\text{ent}}) \in \mathcal{O}(K^2QM) , \]
\[ T(\nabla_S \mathcal{L}_{\text{ent}}) \in \mathcal{O}(K^2QM) , \]
\[ T(\nabla_{\pi_k} \mathcal{L}_{\text{ent}}) \in \mathcal{O}(K^2QM) . \]

Hence the computational cost of evaluating the KL-term and its gradients is in \(\mathcal{O}(K^2QM^3)\) for full covariances, and \(\mathcal{O}(KQM(K + M))\) for diagonal covariances.

### J.3 Analysis of the \(\mathcal{L}_{\text{ell}}\)

Let \(T(\log p(y_n|f_n, \phi)) \in \mathcal{O}(L)\), then the cost of evaluating the expected log likelihood and its gradients is given by:
\[ T(\mathcal{L}_{\text{ell}}) \in \mathcal{O}(KBS\langle L + Q \rangle) , \]
\[ T(\nabla_m \mathcal{L}_{\text{ell}}) \in \mathcal{O}(KQ\langle M^2 + BM + BSL \rangle) , \]
\[ T(\nabla_S \mathcal{L}_{\text{ell}}) \in \mathcal{O}(KQB\langle M^2 + SL \rangle) , \]
\[ T(\nabla_{\pi_k} \mathcal{L}_{\text{ell}}) \in \mathcal{O}(KBS\langle L + Q \rangle) , \]

and for diagonal covariance we have:
\[ T(\nabla_S \mathcal{L}_{\text{ell}}) \in \mathcal{O}(KQB\langle M + SL \rangle) . \]

Hence the computational cost of evaluating the ell term and its gradients is in \(\mathcal{O}(KQB\langle M^2 + SL \rangle)\) for full covariances, and \(\mathcal{O}(KQB\langle M + SL \rangle)\) for diagonal covariances.

### K Notation

A summary of the notation used in this paper is shown in the table below.

| Symbol | Description |
|--------|-------------|
| \(v\)  | Bold lower case denotes a vector |
| \(C\)  | Bold upper case denotes a matrix |
| \(|C|\) | Determinant of matrix \(C\) |
| \(\text{tr } C\) | Trace of matrix \(C\) |
| \(D\)  | Input dimensionality |
| \(N\)  | Number of training datapoints |
| Symbol | Description |
|--------|-------------|
| $M$    | Number of inducing variables per latent process |
| $P$    | Number of outputs |
| $Q$    | Number of latent functions |
| $K$    | Number of mixture components in variational posterior |
| $S$    | Number of samples in Monte Carlo estimates |
| $x_n$  | $n$th input datapoint |
| $X$    | $N \times D$ matrix of input data |
| $y_n$  | $P$-dimensional vector of outputs (label) for $n$th observation |
| $f_n$  | $Q$-dimensional vector of latent function values for $n$th observation |
| $f_{j}$| $N$-dimensional vector of latent function values for $j$th process |
| $u_{j}$| Inducing variables of latent process $j$ |
| $Z_j$  | $M \times D$ matrix of inducing inputs for latent process $j$ |
| $\kappa_j(x, x')$ | Covariance function of latent process $j$ evaluated at $x, x'$ |
| $K_{xx}^j$ | $N \times N$ covariance matrix obtained by evaluating $\kappa_j(X, X)$ |
| $K_{zz}^j$ | $M \times M$ covariance matrix obtained by evaluating $\kappa_j(Z_j, Z_j)$ |
| $K_{xz}^j$ | $N \times M$ covariance between $X$ and $Z_j$ |
| $k_{zn}$ | $M \times 1$ covariance vector between $Z_j$ and $x_n$ |
| $q(u|\lambda)$ | Variational distribution over inducing variables |
| $q(f|\lambda)$ | Variational distribution over latent functions |
| $m$    | Posterior mean over inducing variables |
| $S$    | Posterior covariance over inducing variables |
| $b$    | Posterior mean over latent functions |
| $\Sigma$ | Posterior covariance over latent functions |

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