Kernel Interpolation for Scalable Online Gaussian Processes

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Abstract

Gaussian processes (GPs) provide a gold standard for performance in online settings, such as sample-efficient control and black box optimization, where we need to update a posterior distribution as we acquire data in a sequential fashion. However, updating a GP posterior to accommodate even a single new observation after having observed n points incurs at least $\mathcal{O}(n)$ computations in the exact setting. We show how to use structured kernel interpolation to efficiently reuse computations for constant-time $\mathcal{O}(1)$ online updates with respect to the number of points n, while retaining exact inference. We demonstrate the promise of our approach in a range of online regression and classification settings, Bayesian optimization, and active sampling to reduce error in malaria incidence forecasting. Code is available at https: //github.com/wjmaddox/online_gp.

1 INTRODUCTION

The ability to repeatedly adapt to new information is a defining feature of intelligent agents. Indeed, these online or streaming settings, where we observe data in an incremental fashion, are ubiquitous — from real-time adaptation in robotics (Nguyen-Tuong et al., 2008) to click-through rate predictions for ads (Liu et al., 2017).

Bayesian inference is naturally suited to the online setting, where after each new observation, an old posterior becomes a new prior. However, these updates can be prohibitively slow. For Gaussian processes, if we have already observed n data points, observing even a single

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new point requires introducing a new row and column into an $n \times n$ covariance matrix, which can incur $\mathcal{O}(n^2)$ operations for the predictive distribution and $\mathcal{O}(n^3)$ operations for kernel hyperparameter updates.

Since Gaussian processes are now frequently applied in online settings, such as Bayesian optimization (Yamashita et al., 2018; Letham et al., 2019), or modelbased robotics (Xu et al., 2014; Mukadam et al., 2016), this scaling is particularly problematic. Moreover, despite the growing need for scalable online inference, recent research on this topic is scarce.

Existing work has typically focused on data sparsification schemes paired with low-rank kernel updates (e.g., Nguyen-Tuong et al., 2008), or sparse variational posterior approximations (Cheng and Boots, 2016; Bui et al., 2017). Low-rank kernel updates are sensible but still costly, and data-sparsification can incur significant error. Variational approaches, while promising, can provide miscalibrated uncertainty representations compared to exact inference (Jankowiak et al., 2020; Lázaro-Gredilla and Figueiras-Vidal, 2009; Bauer et al., 2016), and often involve careful tuning of many hyperparameters. In the online setting, these limitations are especially acute. Uncertainty representation can be particularly crucial for determining the balance of exploration and exploitation in choosing new query points. Moreover, while tuning of hyperparameters and manual intervention may be feasible for a fixed dataset, it can become particularly burdensome in the online setting if it must be repeated after we observe each new point.

Intuitively, we ought to be able to recycle computations to efficiently update our predictive distribution after observing an additional point, rather than starting training anew on n+1 points. However, it is extremely challenging to realize this intuition in practice, for if we observe a new point, we must compute its interaction with every previous point. In this paper, we show it is in fact possible to perform constant-time $\mathcal{O}(1)$ updates in n, and $\mathcal{O}(m^2)$ for m inducing points, to the Gaussian process predictive distribution, marginal

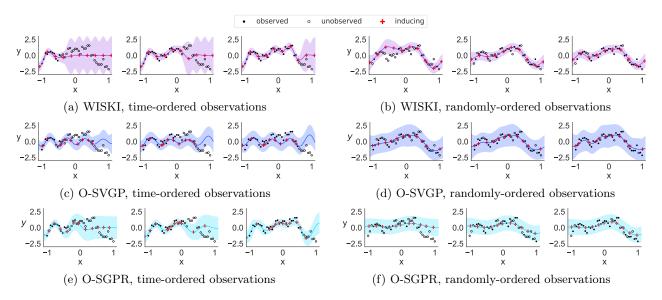


Figure 1: Online GP regression on exchange rate time series data (N = 40). The shaded regions in each panel corresponds to a 95% credible interval. In each subplot, the left subpanel shows the predictive distribution of the corresponding model after training in batch on an initial set of 10 observations. The middle and right subpanels show the evolution of the predictive distribution after 10 and 20 online updates, respectively. The **left** plots, (a,c,e), show WISKI, O-SVGP, and O-SGPR using spectral mixture kernels (Wilson and Adams, 2013) trained on observations in a time-ordered fashion. O-SVGP heavily overfits to the initial data by interpolating the first batch of data points, and struggles to recover on the next batches. WISKI and O-SGPR perform well in this situation by picking up the signal on the first batches and updating the mean as the data comes in. The **right** plots, (b,d,f) show the methods trained on observations in a randomly ordered fashion. Here, O-SVGP is still very under-confident, while O-SGPR clumps its inducing points in the middle of the data. By comparison, WISKI learns more of the high frequency trend than either variational approach.

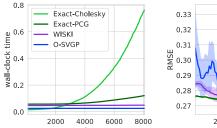
likelihood, and its gradients, while retaining exact inference. We achieve this scaling through a careful combination of caching, structured kernel interpolation (SKI) (Wilson and Nickisch, 2015), and reformulations involving the Woodbury identity. We name our approach Woodbury Inversion with SKI (WISKI). We find that WISKI achieves promising results across a range of online regression and classification problems, Bayesian optimization, and an active sampling problem for estimating malaria incidence where fast online updates, exact inference for calibrated uncertainty, and fast test-time predictions are particularly crucial.

As a motivating example, in Figure 1, we fit GPs with spectral mixture kernels (Wilson and Adams, 2013) on British pound to USD foreign exchange data. In this task, we observe points one at a time, after observing the first 10 points in batch, and update the predictive distributions for WISKI, O-SVGP and O-SGPR (Bui et al., 2017), state-of-the-art streaming sparse variational GPs. We illustrate snapshots after

having observed n=10,20, and 30 points. We see that WISKI is able to more easily capture signal in the data, whereas O-SVGP tends to underfit and O-SGPR underfits on the random data setting. In addition to the general tendency of stochastic variational GP (SVGP) models to underfit the data and overestimate noise variance (Lázaro-Gredilla and Figueiras-Vidal, 2009; Bauer et al., 2016), the variational posterior of an O-SVGP is discouraged from adapting to surprising new observations (See Appendix B). We also see that O-SVGP particularly struggles when we observe new points in a time-ordered fashion, which is a standard setup in the online setting.

The initialization heuristics used to train SVGPs in the batch setting, such as initializing the inducing points with k-means or freezing the GP hyperparameters at the beginning of training, are not effective for O-SVGPs since the full dataset is not available. In order to obtain reasonable fits with O-SVGP on even this motivating example, we carefully tuned tempering parameters using generalized variational inference (Knoblauch et al., 2019), executed 6 optimization steps for each new observation, and trained in batch on the first 10 points. WISKI, by contrast, requires no tuning, only 1 opti-

 $^{^{1}}$ https://raw.githubusercontent.com/trungngv/cogp/master/data/fx/fx2007-processed.csv, fourth column. We rescaled the inputs to [-1,1] and standardized the responses.



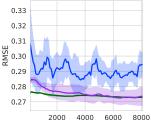


Figure 2: Left: Incorporating new observations becomes increasingly expensive for exact GPs (Exact-Cholesky), even when preconditioned conjugate gradients (Exact-PCG), as quantified in the left panel by the wall-clock time per iteration on the UCI Powerplant dataset. Variational GPs (O-SVGP) are an economical alternative by virtue of being constant time. WISKI has the constant-time profile of a variational method, but retains exact inference, is simple to train, and does not underfit. Right: RMSE on the UCI power plant dataset. Shown are mean and two standard deviations over 10 trials. O-SVGP tends to overestimate noise and converges to a sub-optimal solution, while WISKI matches the performance of the exact methods trained in an incremental fashion.

mization step for each new observation, and does not require any batch training to find reasonable solutions.

Indeed, the problems with O-SVGP² are more noticeable on more complicated datasets than just time series. In Figure 2, we plot the incremental RMSE on a held out test set on the UCI powerplant dataset, while optimizing for only a single step as we observe new data points, finding that O-SVGP underfits and sub-optimal solution, while WISKI matches the performance of an exact GP also fit incrementally. The exact GP also uses pre-conditioned conjugate gradients (Gardner et al., 2018) here. However, WISKI and O-SVGP are both constant time (shown in the left panel), while using an exact GP with Cholesky factorization is cubic time, and using CG with the GP is quadratic time. Both are much slower than WISKI and O-SVGP after t=5000.

2 RELATED WORK

2.1 Prior Approaches

Despite its timeliness, there has not been much recent work on online learning with GPs. Older work considers sparse variational approximations to GPs in the streaming setting. Csató and Opper (2002) proposed a variational sparse GP based algorithm in $\mathcal{O}(nm^2+m^3)$ time, specifically for deployment in streaming tasks; however, it assumes that the hyperparameters are fixed. Nguyen-Tuong et al. (2008) proposed local fits to GPs with weightings based on the distance of the test point to the local models. More recently, Koppel (2019) extended the types of distances used for these types of models while using an iteratively constructed coreset of data points. Evans and Nair (2018) proposed a structured eigenfunction based approach that requires one $\mathcal{O}(n)$ computation of the kernel and uses fixed kernel hyperparameters but learns interpolation weights. Cheng and Boots (2016) proposed a variational stochastic functional gradient descent method in incremental setting with the same time complexity; however, like stochastic variational GPs (Hensman et al., 2013), Cheng and Boots (2016) assumes the number of data points the model will see is known and set before training begins. Hoang et al. (2015) proposed a similar variational natural gradient ascent approach, but assumed that the hyper-parameters are fixed during the training procedure, a major limitation for flexible kernel learning.

2.2 Streaming SVGP and Streaming SGPR

The current state-of-the-art for streaming Gaussian processes is the sparse variational O-SVGP approach of Bui et al. (2017) and its "collapsed" non-stochastic variant, O-SGPR, which does not use an explicit variational distribution, like its batch equivalent, SGPR (Titsias, 2009).

O-SVGP: Unlike its predecessors, O-SVGP is fully compatible with online inference, since it has no requirements to choose the number of data points a priori, and it can update both model parameters and inducing point locations; however, it has the same time complexity as its predecessors: $\mathcal{O}(bm^2 + m^3)$, where b is the size of the batch used to update the predictive distribution and model hyper-parameters. Bui et al. (2017)'s experiments primarily focused on large batch sizes practically $b = \mathcal{O}(n)$ — rather than the pure streaming setting. A major limitation of variational methods in the streaming setting is that conditioning on new observations effectively requires the model parameters to be re-optimized to a minima after every new batch, increasing latency. In Appendix B we include a detailed discussion of the requirements of the original O-SVGP algorithm, and provide a modified generalized variational update by downweighting the prior by a factor of β < 1 better adapted to the streaming setting to have a strong baseline for comparison. We compare to the generalized O-SVGP implementation in our experiments as O-SVGP.

²O-SGPR has slightly different weaknesses, including numerical instability. We further consider O-SGPR in our larger study of incremental regression.

O-SGPR is also potentially promising O-SGPR: but like O-SVGP falls prey to several key limitations. First, O-SGPR relies on analytic marginalization and so can only be used for Gaussian likelihoods. In Figure 1, we implemented the O-SGPR bound in GPyTorch (Gardner et al., 2018) and it has fair performance for both the random ordering and time ordering settings. though not as good as WISKI. However, this performance comes with two caveats. First, we need to re-sample the inducing points to include some of the new data at each iteration, as is done in Bui et al. (2017)'s implementation. Second, we found that even in double precision we needed to add a large amount of jitter $\epsilon = 0.01$, while doing the required Cholesky decompositions (there is a matrix subtraction) to prevent numerical instability.

3 BACKGROUND

For a complete treatment on Gaussian Processes, see Williams and Rasmussen (2006). Here we briefly review the key ideas for efficient exact GPs, SKI, and the conditioning of GPs on new observations online. We note SKI provides scalable exact inference through creating an approximate kernel which admits fast computations.

3.1 Exact GP Regression

Starting with the regression setting, suppose $\mathbf{y} = f(x) + \varepsilon$, $f \sim \mathcal{GP}(0, k_{\theta}(x, x'))$, and $\varepsilon \sim \mathcal{N}(0, \sigma^2)$. Here, $k_{\theta}(x, x')$ is the kernel function with hyperparameters θ , and $K_{AB} := k_{\theta}(A, B)$ is the covariance between two sets of data inputs A and B. Given training data $\mathcal{D} = (X, \mathbf{y})$, we can train the GP hyperparameters by maximizing the marginal log-likelihood,

$$\log p(\mathbf{y}|X,\theta) = -\frac{1}{2}\mathbf{y}^{\top}(K_{XX} + \sigma^2 I)^{-1}\mathbf{y}$$
$$-\frac{1}{2}\log|K_{XX} + \sigma^2 I| - \frac{n}{2}\log 2\pi. \quad (1)$$

Conventionally, solving the linear system, $(K_{XX} + \sigma^2 I)^{-1} \mathbf{y}$, in Eq. 1 costs $\mathcal{O}(n^3)$ operations. The posterior predictive distribution of a new test point $p(f(\mathbf{x}^*)|\mathbf{x}^*, \mathcal{D}, \theta) = \mathcal{N}(\mu_{f|\mathcal{D}}, \sigma^2_{f|\mathcal{D}})$, where

$$\mu_{f|\mathcal{D}}(\mathbf{x}^*) = K_{\mathbf{x}^*X}(K_{XX} + \sigma^2 I)^{-1}\mathbf{y}, \tag{2}$$

$$\sigma_{f|\mathcal{D}}^2(\mathbf{x}^*) = K_{\mathbf{x}^*\mathbf{x}^*} - K_{\mathbf{x}^*X}(K_{XX} + \sigma^2 I)^{-1}K_{X\mathbf{x}^*}$$
(3)

We build on previous work on scaling GP training and prediction by exploiting kernel structure and efficient GPU matrix vector multiply routines to quickly compute gradients (CG) of Eq. 1 for training, and by caching terms in Eq. 2 and Eq. 3 for fast prediction (Gardner et al., 2018). Conjugate gradient methods improve the asymptotic complexity of GP regression

and to $\mathcal{O}(jn^2)$, where j is the number of CG steps used. These recent advances in GP inference have enabled exact GP regression on datasets of up to one million data points in the batch setting (Wang et al., 2019).

3.2 Structured Kernel Interpolation (SKI) and Lanczos Variance Estimates

GPs are often sparsified through the introduction of inducing points (also known as pseudo-inputs), which are small subset of fixed points (Snelson and Ghahramani, 2006). In particular, Wilson and Nickisch (2015) proposed structured kernel interpolation (SKI) to approximate the kernel matrix as $K_{XX} \approx K_{XX} = WK_{UU}W^{\top}$, where U represents the m inducing points, and $W \in$ $\mathbb{R}^{n\times m}$ is a sparse cubic interpolation matrix composed of n vectors $\mathbf{w}_i \in \mathbb{R}^m$. Each vector \mathbf{w}_i is sparse, containing 4^d non-zero entries, where d is the dimensionality of the input data. SKI places the inducing points on a multi-dimensional grid. When k_{θ} is stationary and factorizes across dimensions, K_{UU} can often be expressed as a Kronecker product of Toeplitz matrices, leading to fast multiplies. Overall multiplies with K_{XX} take $\mathcal{O}(n+q(m))$ time, where $q(m)\approx m$ (Wilson and Nickisch, 2015), compared to the $\mathcal{O}(nm^2)$ complexity associated with most inducing point methods (Quinonero-Candela and Rasmussen, 2005). In short, SKI provides scalable exact inference, through introducing an approximate kernel that admits fast computations.

Pleiss et al. (2018) propose to cache (i.e. to store in memory) all parts of the predictive mean and covariance that can be computed before prediction, enabling constant time predictive means and covariances. Directly substituting the SKI kernel matrix, \tilde{K}_{XX} , into Eq. 2, the predictive mean becomes

$$\mu_{f|\mathcal{D}}\left(\mathbf{x}^*\right) = \mathbf{w}_{\mathbf{x}^*}^{\top} \underbrace{K_{UU}W^{\top} \left(WK_{UU}W^{\top} + \sigma^2I\right)^{-1}\mathbf{y}}_{\mathbf{a}},$$

where **a** is the *predictive mean cache*.³ Similarly, Eq. 3 becomes

$$\sigma_{f|\mathcal{D}}^{2}\left(\mathbf{x}_{i}^{*}, \mathbf{x}_{j}^{*}\right) = k(\mathbf{x}_{i}, \mathbf{x}_{j}^{*}) - \mathbf{w}_{\mathbf{x}^{*}}^{\top} \underbrace{K_{UU}W^{\top} \left(\tilde{K}_{XX} + \sigma^{2}I\right)^{-1} W K_{UU}}_{C} \mathbf{w}_{\mathbf{x}^{*}},$$

where $C \approx SS^{\top}$, is the predictive covariance cache. S is formed by computing a rank-k root decomposition of $(\tilde{K}_{XX} + \sigma^2 I)^{-1} \approx RR^{\top}$ and taking $S = K_{UU}W^{\top}R$. The complexity of the root decomposition is $\mathcal{O}(km^2)$,

³We refer to entities that can be computed, stored in memory, and used in subsequent computations as *caches*. We use blue font to identify which cached expressions.

requiring $k \leq m$ iterations of the Lanczos algorithm (Lanczos, 1950) and a subsequent eigendecomposition of the resulting $k \times k$ symmetric tridiagonal matrix. Further details on Lanczos decomposition and the caching methods of (Pleiss et al., 2018) are in Appendix A.

3.3 Online Conditioning and Low-Rank Matrix Updates

GP models are conditioned on new observations through Gaussian marginalization (Williams and Rasmussen, 2006, Chapter 2). Suppose we have past observations $\mathcal{D} = \{(\mathbf{x}_i, y_i)\}_{i=1}^n$ used to make predictions via $p(y^*|\mathbf{x}^*, \mathcal{D}, \theta)$. We subsequently observe a new data point (\mathbf{x}', y') . For clarity, let $X = \mathbf{x}_{1:n}$, $X' = X \cup \{\mathbf{x}'\}$. The new kernel matrix is

$$K_{X'X'} = \begin{pmatrix} K_{XX} & k(X, \mathbf{x}') \\ k(\mathbf{x}', X) & k(\mathbf{x}', \mathbf{x}') \end{pmatrix}$$
(4)

We would like to update our posterior predictions to incorporate the new data point without recomputing our caches that are not hyper-parameter dependent from scratch. If the hyperparameters are fixed, this can be a $\mathcal{O}(n^2)$ low-rank update to the predictive covariance matrix (e.g. a Schur complement update or low rank Cholesky update to a decomposition of $(K_{XX} + \sigma^2 I)^{-1}$). If we additionally wish to update hyper-parameters, we must recompute the marginal log likelihood in Eq. 1, which costs $\mathcal{O}(n^3)$. Similarly, if we naively use the SKI approximations in Eq. 4 we additionally have an $\mathcal{O}(n)$ cost for both adding a new data point and to update the hyper-parameters afterwards. Thus, as n increases, training and prediction will slow down (Figure 2).

4 WISKI: ONLINE CONSTANT TIME SKI UPDATES

We now propose WISKI, which through a careful combination of caching, SKI, and the Woodbury identity, achieves constant time (in n) updates in the streaming setting, while retaining exact inference. To begin, we present two key identities that result from the application of the Woodbury matrix identity to the inverse of the updated SKI approximated kernel, $\tilde{K}_{X_tX_t}$, after having received t data points. First, we can rewrite the SKI kernel inverse as

$$(\tilde{K}_{XX} + \sigma^2 I)^{-1} = \frac{1}{\sigma^2} I - \frac{1}{\sigma^2} W M W^{\top}.$$
 (5)

$$M := (\sigma^2 K_{UU}^{-1} + W^{\top} W)^{-1}.$$
 (6)

Second, after observing a new data point at time t+1, the inner matrix inverse term M can be updated via a rank-one update,

$$M_{t+1}^{-1} = M_t^{-1} + \mathbf{w}_{t+1} \mathbf{w}_{t+1}^{\top}, \tag{7}$$

where \mathbf{w}_{t+1} is an interpolation vector for the t+1th data point.⁴ The exploitation of this rank-one update on a fixed size matrix by storing $W^{\top}W$ will form the basis of our work.

Computing Eq. 7 as written requires explicit computation of K_{UU}^{-1} . In general, K_{UU} will have significant structure as U is a dense grid, which yields fast matrix inversion algorithms; however, the inverse will be very ill-conditioned because many kernel matrices on gridded data have (super-)exponentially decaying eigenvalues (Bach and Jordan, 2002). We will instead focus on reformulating SKI into expressions that depend only on K_{UU} , W, and \mathbf{y} with a constant $\mathcal{O}(m^2)$ memory footprint and can be computed in $\mathcal{O}(m^2)$ time.

4.1 Computing the Marginal Log-Likelihood, Predictive Mean and Predictive Variance

Substituting Eq. (5) into Eqs. (1), (2), and (3), we obtain the following expressions for the marginal log-likelihood (MLL), predictive mean, and predictive variance⁵:

$$\log p(\mathbf{y}|X,\theta) = -\frac{1}{2\sigma^2}(\mathbf{y}^{\top}\mathbf{y} - \mathbf{y}^{\top}WMW^{\top}\mathbf{y}) -$$

$$\frac{1}{2} \left(\log |K_{UU}| - \log |M| + (n-m) \log \sigma^2 \right), \tag{8}$$

$$\mu_{f|\mathcal{D}}\left(\mathbf{x}^*\right) = \mathbf{w}_{\mathbf{x}^*}^{\top} M W^{\top} \mathbf{y},\tag{9}$$

$$\sigma_{f|\mathcal{D}}^{2}\left(\mathbf{x}_{i}^{*}, \mathbf{x}_{j}^{*}\right) = \sigma^{2}\mathbf{w}_{\mathbf{x}_{i}^{*}}^{\top} M \mathbf{w}_{\mathbf{x}_{i}^{*}}.$$
(10)

For all derivations see Appendix A. We begin by constructing a rank r root decomposition of the matrix $W^\top W \approx LL^\top$, along with the factorization of the (pseudo-)inverse, $JJ^\top \approx (W^\top W)^+$. The root decomposition LL^\top can be a full Cholesky factorization (r=m) for relatively small m (i.e. $m \leq 1000$) or an approximate Lanczos decomposition for larger m, at a one-time cost of $\mathcal{O}(m^2r)$. Applying the Woodbury matrix identity to Eq. (6) and substituting $W^\top W \approx LL^\top$, we have

$$M = \sigma^{-2} K_{UU} - \sigma^{-2} K_{UU} L Q^{-1} L^{\top} \sigma^{-2} K_{UU}, \quad (11)$$

$$Q := I + \mathbf{L}^{\top} \sigma^{-2} K_{UU} \mathbf{L}. \tag{12}$$

 $Q^{-1}L^{\top}$ is a $r \times r$ system, so directly computing Eq. (11) requires $\mathcal{O}(r^2m)$ time for the solve using conjugate gradients, $\mathcal{O}(rm\log m)$ time for the matrix multiplications with K_{UU} if it has Toeplitz structure, and $\mathcal{O}(m^2)$ for the dense matrix additions, and $\mathcal{O}(km)$ for the root decomposition of $W^{\top}W$, for a final total of $\mathcal{O}(r^2m+km\log m+m^2)$. However, we do not explicitly

 $^{^4\}mathrm{We}$ have dropped the dependence on $\mathbf x$ for simplicity of notation.

⁵A similar result holds for fixed noise heteroscedastic likelihoods as well. See Appendix A.5 for further details.

store the matrix M as doing so would require m solves of a $r \times r$ system since $L \in \mathbb{R}^{m \times r}$.

Eqs. (8) - (10) involve computations of the form

$$M\mathbf{v} = \sigma^{-2}K_{UU}\mathbf{v} - \sigma^{-2}K_{UU}LQ^{-1}L\sigma^{-2}K_{UU}\mathbf{v},$$

which can be computed using only a *single* solve against the matrix Q via first multiplying out $\mathbf{a} = L\sigma^{-2}K_{UU}\mathbf{v}$, and then computing $\mathbf{b} = Q^{-1}\mathbf{a}$. Applying the matrix determinant identity to $\log |M|$ results in a simplified expression in terms of $\log |Q|$. Taking $\mathbf{v} = W^{\top}\mathbf{y}$, we obtain a practical expression for the MLL,

$$\log p(\mathbf{y}|X,\theta) = -\frac{1}{2\sigma^2} \left(\mathbf{y}^{\top} \mathbf{y} - \mathbf{y}^{\top} W K_{UU} W^{\top} \mathbf{y} + \mathbf{a}^{\top} Q^{-1} \mathbf{a} \right) - \frac{1}{2} \left(-\log|Q| + (n-m)\log\sigma^2 \right). \tag{13}$$

Computing a costs $\mathcal{O}(m \log m + rm)$, so computing the two quadratic forms are $\mathcal{O}(m \log m + m)$ and $\mathcal{O}(jr^2)$ respectively, assuming j steps of conjugate gradients. We use stochastic Lanczos quadrature to compute the log determinant of |Q| which costs $\mathcal{O}(jr^2)$ (Gardner et al., 2018). Overall, computation of the MLL becomes $\mathcal{O}(rm + m \log m + jr^2)$.

The predictive mean is similarly computed by taking $\mathbf{v} = W^{\top} y$, resulting in the expression

$$\mu_{f|\mathcal{D}}(\mathbf{x}^*) = \mathbf{w}_{\mathbf{x}^*}^{\top} \left(\sigma^{-2} K_{UU}(\mathbf{W}^{\top} \mathbf{y} - L\mathbf{b}) \right). \tag{14}$$

The only term that remains is the predictive variance, for which we take $\mathbf{v} = \mathbf{w}_{\mathbf{x}_{i}^{*}}$ and obtain

$$\sigma_{f|\mathcal{D}}^{2}\left(\mathbf{x}_{i}^{*}, \mathbf{x}_{j}^{*}\right) = \sigma^{2}\mathbf{w}_{\mathbf{x}_{i}^{*}}^{\top}\left(K_{UU}(\mathbf{w}_{\mathbf{x}_{j}^{*}} - \mathbf{L}\mathbf{b})\right).$$
 (15)

4.2 Conditioning on New Observations

When we observe a new data point $(\mathbf{x}_{t+1}, y_{t+1})$, we need to update $(W^{\top}\mathbf{y})_t$, $(\mathbf{y}^{\top}\mathbf{y})_t$, and $L_tL_t^{\top} = (W^{\top}W)_t$. The update to the first two terms is simple:

$$(W^{\top}y)_{t+1} = (W^{\top}\mathbf{y})_t + y_{t+1}\mathbf{w}_{\mathbf{x}_{t+1}}$$
 (16)

$$(\mathbf{y}^{\top}\mathbf{y})_{t+1} = (\mathbf{y}^{\top}\mathbf{y})_t + y_{t+1}^2$$
(17)

We can update L_t in $\mathcal{O}(mr+r)$ time by exploiting the rank-one structure of the expression

$$(W^{\top}W)_{t+1} = (W^{\top}W)_t + \mathbf{w}_{\mathbf{x}_{t+1}}\mathbf{w}_{\mathbf{x}_{t+1}}^{\top}.$$

Recalling that $JJ^{\top} = (W^{\top}W)^+$, let $\mathbf{p} = J_t^{\top}\mathbf{w}_{\mathbf{x}_{t+1}}$. We compute the decomposition $BB^{\top} = I_r + \mathbf{p}\mathbf{p}^{\top}$ and obtain the expression for the updated root $L_{t+1} = L_t B$. Since BB^{\top} is a decomposition of I_r plus a rank-one correction, it can be computed in $\mathcal{O}(r)$ time. Since the updates to the first two caches are $\mathcal{O}(1)$ and $\mathcal{O}(m)$, respectively, the total complexity of conditioning on a new observation is $\mathcal{O}(mr^2 + r)$. Further details and an extended proof are given in Appendix A.

4.3 Updating Kernel Hyperparameters

Conventionally, learning the kernel hyperparameters θ of a GP online presents two major challenges. First, the basic form of the gradient of the MLL naively costs at least $\mathcal{O}(n)$, even if scalable methods are employed. Second, after a parameter update, any cached terms that depend on the kernel matrix must be recomputed (e.g. a new factorization of K_{XX}). The reformulation of the MLL in Eq. (13) addresses the first challenge, with a complexity of $\mathcal{O}(rm + m \log m + jr^2)$ (after computing the necessary caches). To address the second challenge, we observe that the combination of the SKI approximation to the kernel matrix and the Woodbury matrix identity in Section 4.1 has allowed us to reformulate GP inference entirely in terms of computations whose cost depends only on the number of inducing points and the rank of the matrix decompositions (which is at most m, and typically much less than m). As a result, we can recompute the necessary caches as needed without any increase in computational effort as n increases.

The computational efficiency of SKI is a direct result of the grid structure imposed on the inducing points. The reduced computational complexity comes at the cost of memory complexity that is exponential in the dimension of the input. In practice, if the input data has more than three or four dimensions, the inputs must be projected into a low-dimensional space. The projection may be random (Delbridge et al., 2020) or learned (Wilson et al., 2016), depending on the requirements of the task. If the projection is learned, then the parameters ϕ of the projection operator h are treated as additional kernel hyperparameters and trained through the marginal log-likelihood.

In the batch setting the interpolation weights W are updated after every optimization iteration to adapt to the new projected features $h(\mathbf{x}_{1:n};\phi)$. In the online setting, updating W for every previous observation would be $\mathcal{O}(n)$. Since we cannot update the interpolation weights for old observations, the gradient for the projection parameters at time t can be rewritten as follows:

$$\nabla_{\phi} \mathcal{L}(\phi) = \nabla_{\phi} \frac{1}{2} \left((\mathbf{y}^{\top} W)_{t} M_{t-1} (W^{\top} \mathbf{y})_{t} \right)$$
$$-\frac{1}{1 + \mathbf{v}^{\top} \mathbf{w}} (\mathbf{v}_{t}^{\top} (W^{\top} \mathbf{y})_{t})^{2} - \log(1 + \mathbf{v}_{t} \mathbf{w}_{t}), \quad (18)$$
$$\mathbf{w}_{t} = w(h(\mathbf{x}_{t}; \phi)), \quad \mathbf{v}_{t} = M_{t-1} \mathbf{w}_{t},$$
$$(W^{\top} \mathbf{y})_{t} = (W^{\top} \mathbf{y})_{t-1} + y_{t} \mathbf{w}_{t}.$$

The gradient in Eq. (18) will move the projection parameters ϕ in a direction that maximizes the marginal likelihood, assuming $\mathbf{w}_{1:t-1}$ are fixed. That is, only projections on new data are updated, while the old projections remain fixed. In contrast to the batch setting,

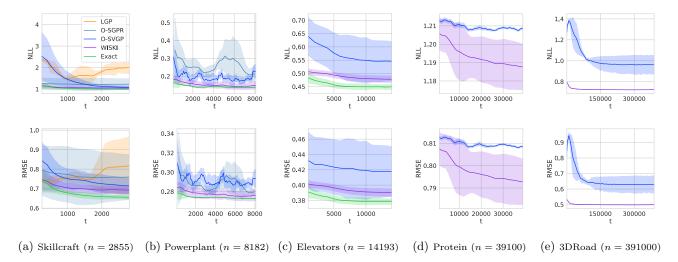


Figure 3: Online homoskedastic regression on UCI datasets. We compare to local GPs (LGP), O-SGPR, O-SVGP, and exact GPs. Due to memory constraints or numerical issues for other methods, only O-SVGP and WISKI were easily capable of running on the larger tasks (Protein and 3DRoad). WISKI has comparable accuracy to exact methods, with comparable runtime to scalable approximations like OSVGP. **Top:** Test set NLLs. **Bottom:** Test set RMSEs.

where ϕ is jointly optimized with θ , the online update is sequential; whenever a new observation is received ϕ is updated through Eq. (18), then the GP is conditioned on $(h(\mathbf{x}_t; \phi_t), y_t)$, and finally θ is updated through Eq. (1). See Appendix A.4 for the full derivation.

5 EXPERIMENTS

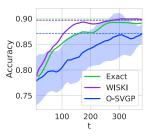
To evaluate WISKI, we first consider online regression and binary classification. We then demonstrate how WISKI can be used to accelerate Bayesian optimization, a fundamentally online algorithm often applied to experiment design and hyperparameter tuning (Frazier, 2018). Finally we consider an active learning problem for measuring malaria incidence, and show that the scalability of WISKI enables much longer horizons than a conventional GP.

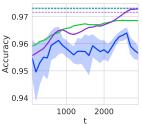
We compare against exact GPs (no kernel approximations), O-SGPR, O-SVGP (Bui et al., 2017), sparse variational methods that represents the current gold standard for scalable online Gaussian processes, and local GPs (LGPs) (Nguyen-Tuong et al., 2008). All experimental details (hyper-parameters, data preparation, etc.) are given in Appendix C, where we also include ablation studies on the β parameter for O-SVGP as well as the the number of inducing points (as we had to modify it to achieve good results in the incremental setting for O-SVGP), m, for WISKI and O-SVGP. Unless stated otherwise, shaded regions in the plots correspond to $\overline{\mu} \pm 2\overline{\sigma}$, estimated from 10 trials.

5.1 Regression

We first consider online regression on several datasets from the UCI repository (Dua and Graff, 2017). In each trial we split the dataset into a 90%/10% train/test split. We scaled the raw features to the unit hypercube $[-1,1]^d$ and normalized the targets to have zero-mean and unit variance. Each model learned a linear projection from \mathbb{R}^d to \mathbb{R}^2 that was transformed via a batchnorm operation and the non-linear tanh activation to ensure that the features were constrained to $[-1, 1]^2$. Each model learned an RBF-ARD kernel on the transformed features, except on the 3DRoad dataset, which did not require dimensionality reduction. We used the same number of inducing points for WISKI, O-SGPR, and O-SVGP and set $n_{\text{max}} = m$ for local GPs. The models were pretrained on 5% of the training examples, and then trained online for the remaining 95%. When adding a new data point, we update with a single optimization step for each method, such that the runtime is similar for the scalable methods. Since O-SVGP can be sensitive to the number of gradient updates per timestep, in Figure A.2 in the Appendix we provide results for an ablation.

We show the test NLL and RMSE for each dataset in Figure 3. For the two largest datasets we only report results for WISKI and O-SVGP. We found that O-SGPR was fairly unstable numerically, even after using an exceptionally large jitter value (0.01) and switching from single to double precision. The exact baseline and the WISKI model overfit less to the initial examples than O-SGPR or O-SVGP. Note that O-SVGP is a





- (a) Banana (n = 400)
- (b) SVM Guide 1 (n = 3000)

Figure 4: A comparison of Dirichlet-based exact and WISKI GP classifiers against an O-SVGP with a binomial likelihood. The exact and WISKI models overfit less to the initial data and ultimately match the performance of their hindsight counterparts trained on the full dataset, shown as a dotted line.

much stronger baseline in this experiment since the observations are independently observed than would typically be the case for correlated time-series data, as seen in Figure 1.

5.2 Classification

We extend WISKI to classification though the Dirichletbased GP (GPD) classification formulation of Milios et al. (2018), which reformulates classification as a regression problem with a fixed noise Gaussian likelihood. Empirically the approach has been found to be competitive with the conventional softmax likelihood formulation. In Figure 4 we compare exact and WISKI GPD classifiers to O-SVGP with binomial likelihood on two binary classification tasks, Banana⁶ and SVM Guide 1 (Chang and Lin, 2011). Banana has 2D features, and SVM Guide has 4D features, so we did not need to learn a projection. As in the UCI regression tasks, WISKI and O-SVGP both had 256 inducing point, each model used an RBF-ARD kernel, and the classifiers were pretrained on 5% of the training examples and trained online on the remaining 95%. In both cases the WISKI classifier outperformed the O-SVGP baseline, and matched the accuracy of the the exact baseline.

5.3 Bayesian Optimization

In Bayesian optimization (BO) one optimizes a black-box function by iteratively conditioning a surrogate model on observed data and choosing new observations by optimizing an acquisition function based on the model posterior (Frazier, 2018). Thus, BO re-

quires efficient posterior predictions, updates to caches as new data are observed, and hyperparameter updates. While BO has historically been applied only to expensive-to-query objective functions, we demonstrate here that large-scale Bayesian optimization is possible with WISKI. Our BO experiments are conducted as follows: we choose 5 initial observations using random sampling, then iteratively optimize a batched version of upper confidence bound (UCB) (with q=3) using BoTorch (Balandat et al., 2020) and compute an online update to each GP model, before re-fitting the model. Accurate model fits are critical to high performance; therefore, we wish to use as many inducing points for WISKI and O-SVGP as possible. For both methods, we 1000 inducing points. We show the results over four trials plotting mean and two standard deviations in Figure 5a for the Ackley benchmarks. On both problems, WISKI is significantly faster than the exact GP and O-SVGP, while achieving comparable performance on Levy. We show results over a wider range of test functions in Appendix C.2, along with the best achieved point plotted against the number of steps and the average time per step.

5.4 Active Learning

Finally, we apply WISKI to an active learning problem inspired by Balandat et al. (2020). We consider data describing the infection rate of *Plasmodium falciparum* (a parasite known to cause malaria)⁷ in 2017. We wish to choose spatial locations to query malaria incidence in order to make the best possible predictions on withheld samples. To selectively choose points, we minimize the negative integrated posterior variance (NIPV, Seo et al., 2000), defined as

$$NIPV(x) := -\int_{\mathcal{X}} \mathbb{E}(\mathbb{V}(f(x)|\mathcal{D}_{x})|\mathcal{D}) dx.$$

Optimizing this acquisition function amounts to finding the batch of data points $\mathbf{x}_{1:q}$, the fantasy points, which when added into the GP model will reduce the variance on the domain of the model the most. Here, we randomly sample 10,000 data points in Nigeria to serve as a test set that we wish to reduce variance on and select q = 6 data points at a time from a held-out training set (to act as a simulator) at a time; the inner expectation drops out because the posterior variance only depends on the fantasy points and the currently observed data, and not any fantasized responses. Stochastic variational models do not have a straightforward mechanism for fantasizing (i.e. recomputing the posterior variance after updating a new data point conditional on the fantasy points), so we instead query the test set predictive variance and then

⁶https://raw.githubusercontent.com/thangbui/ streaming_sparse_gp/master/data

⁷Downloaded from the Malaria Global Atlas.

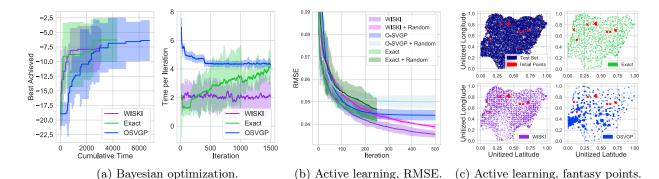


Figure 5: (a): Objective value as a function of cumulative time and time per iteration on the Levy test problem with noise standard deviation 10.0, while performing Bayesian optimization with EI acquisition for 1500 steps with a batch size of 3 so that by the end 4500 data points have been acquired. WISKI allows rapid updates of the posterior surrogate objective out to thousands of observations, while preserving the rapid convergence rate and asymptotic optimality of the exact GP. (b): RMSE on the test set after choosing new points either randomly or with qnIPV (for WISKI and exact GPs) or by the maximal posterior variance (for O-SVGP). WISKI is able to continue improving the downstream error throughout the entire experiment matching the performance of the exact GP, while O-SVGP's performance flatlines. We also compare against the RMSE of models that have data points randomly selected (shown with Random in the legend). (c): The test set (navy), as well as points chosen for all three methods with initial points (red). WISKI and the exact GP query the entire support, while O-SVGP

choose the training points closest to the six test set points with maximum predictive variance.

As both mean and variances are available for the given locations, we model the data with a fixed noise Gaussian process with scaled Matern 0.5 kernels, beginning with an initial set of 10 data points, and iterating out for 500 iterations for WISKI and O-SVGP and 250 iterations for an exact GP model (the limit of data points that a single GPU could handle due to the large amount of test points). We show the results of the experiment in Figure 5b across three trials, finding that all of the models reduce the RMSE considerably from the initial fits; however, O-SVGP and its random counterpart stagnate in RMSE after about 250 trials, while both the exact GP and WISKI continue improving throughout the entire experiment. Closer examination of the points queried by all of the three methods in Figure 5c, we find that the points queried by O-SVGP tend to clump together, locally reducing variance, while WISKI and the exact GP choose points throughout the entire support of the test set, choosing points which better reduce global variance.

6 CONCLUSION

queries clump together.

We have shown how to achieve constant-time online updates with Gaussian processes while retaining exact inference. Our approach, WISKI, achieves comparable performance to Gaussian processes with exact kernels, and comparable speed to state-of-the-art streaming Gaussian processes based on variational inference. Despite the present day need for scalable online probabilistic inference, recent research into online Gaussian processes has been relatively scarce. We hope that our work is a step towards making streaming Bayesian inference more widely applicable in cases when both speed and accuracy are crucial for online decision making.

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