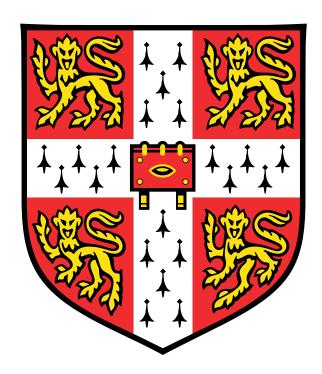
Parameter Optimization using high-dimensional Bayesian Optimization

Bachelor Thesis



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This dissertation is submitted for the degree of

0.1 Project Proposal for Bachelor Thesis

0.1.1 Motivation

Tuning hyperparameters is considered a computationally intensive and tedious task, be it for neural networks, or complex physical instruments such as free electron lasers. Users for such applications could benefit from a 'one-click-search' feature, which would find optimal parameters in as few function evaluations as possible. This project aims to find such an algorithm which is both efficient and holds certain convergence guarantees. We focus our efforts on Bayesian Optimization (BO) and revise techniques for high-dimensional BO.

0.1.2 Background

In Bayesian Optimization, we want to use a Gaussian Process to find an optimal parameter setting \mathbf{x}^* that maximizes a given utility function f. We assume the response surface to be Lipschitz-continuous.

Assume we have observations $\mathscr{Y} = \{y^{(1)}, \dots, y^{(N)}\}$, each evaluated at a point $\mathscr{X} = \{\mathbf{x}^{(1)}, \dots, \mathbf{x}^{(N)}\}$. The relationship between the observations y and individual parameter settings \mathbf{x} is $y = f(\mathbf{x}) + \varepsilon$ where $\varepsilon \sim \mathscr{N}(0, \sigma_n^2)$. Any quantity to be predicted has a subscript-star (e.g. y_* is the function evaluation we want to predict).

In it's simplest form, a Gaussian Process is described by the following equation:

$$\begin{pmatrix} y \\ y_* \end{pmatrix} \sim N \left(\mu, \begin{pmatrix} K & K_*^T \\ K_* & K_{**} \end{pmatrix} \right), \tag{1}$$

Where μ is a mean function, $K = \text{kernel}(\mathbf{X}, \mathbf{X})$, $K_* = \text{kernel}(\mathbf{x}_*, \mathbf{X})$ and $K_{**} = \text{kernel}(\mathbf{x}_*, \mathbf{x}_*)$. We predict any new point y_* , (given all previously sampled points y) by estimating the probability $p(y_*|y) \sim N(K_*K^{-1}y, K_{**} - K_*K^{-1}K_*')$

This, in turn, can be used to build an acquisition function. This acquisition function describes where to best sample points next. Some popular acquisition functions include GP-UCB, Most probable improvement (MPI) and Expected Improvement (EI). The choice of the acquisition function has great influence on the performance of the optimization procedure.

We will talk about the problems and possible solutions for the task at hand in the next section.

0.1.3 Scope of the Project

Bayesian optimization suffers from the curse of dimensionality. The goal of this project is to arrive at a solution that resolves the curse of dimensionality for the specific task with regards to Bayesian optimization. This project includes, but is not limited to the following methods.

1. [7] Assume $f(x) \approx g(\mathbf{W}^T x)$ where $\mathbf{W} \in \mathbb{R}^{D \times d}$ and D >> d. We assume that \mathbf{W} is orthogonal.

This algorithm does not require gradient-information (thus, easier to implement, and robust to noise). The standard-deviation, kernel parameters and W can be found iteratively. First we fix W, and optimize over the standard-deviation, kernel parameters. Then we fix the standard-deviation, kernel parameters. and optimize over W. We repeat this procedure until the change of the log-likelihood between iterations is below some ε_l .

2. [6] Assume $f(x) = f^{(1)}(x^{(1)}) + f^{(2)}(x^{(2)}) + \ldots + f^{(M)}(x^{(M)})$ where $x^{(i)} \in \mathcal{X}^{(i)} \subseteq \mathcal{X}$, i.e. each function component $f^{(i)}$ takes some lower-dimensional subspace as the input. The lower-dimensional subspaces may overlap. The mean and covariance of f(x) is then the sum of the individual component's means and covariances.

An additive decomposition (as described above) can be represented by a dependency graph. The dependency graph is built by joining variables i and j with an edge whenever they appear together within some set x(k).

The goal is to maximize an acquisition function $\phi_t(x) = \sum_{i=1}^M \phi_t^{(i)}(x^{(i)})$. This maximization is achieved by maximizing the probability of Markov Random Fields within the graph. A junction tree is created from the graph, which is then used to find the global maximum of the acquisition function.

The dependencies between the variable-subsets are represented through a graph, which can be learned through Gibbs sampling. This, in turn, is used to create a kernel for the GP.

3. [?] A function $f: \mathbf{R}^D \to \mathbf{R}$ is said to have effective dimensionality d_e (where $d_e < D$), if there exists a linear subspace \mathscr{T} of dimension d_e such that for all $x_{\top} \in \mathscr{T} \subset \mathbf{R}^D$ and $x_{\perp} \in \mathscr{T}_{\perp} \subset \mathbf{R}^D$, we have $f(x) = f(x_{\top} + x_{\perp}) = f(x_{\top})$. \mathscr{T}^{\perp} is the orthogonal complement of \mathscr{T} .

Assume $f: \mathbf{R}^D \to \mathbf{R}$ has effective dimensionality d_e . Given a random matrix $\mathbf{A} \in \mathbf{R}^{D \times d}$ (where $d \ge d_e$) with independent entries sampled from $\mathcal{N}(0,1)$. For any $x \in \mathbf{R}^D$, there

exists a $y \in \mathbf{R}^d$ such that $f(x) = f(\mathbf{A}y)$. We now only need to optimize over all possible $y \in \mathbf{R}^d$, instead of all possible $x \in \mathbf{R}^D$.



Fig. 1 This function in D=2 dimesions only has d=1 effective dimension. Hence, the 1-dimensional embedding includes the 2-dimensional function's optimizer. It is more efficient to search for the optimum along the 1-dimensional random embedding than in the original 2-dimensional space

If, for some reason, finding an active subspace or an effective lower dimension is not possible, we are open to adapt the procedure of optimization.

Abstract

This is where you write your abstract ...

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Background

1.1

Lorem Ipsum is simply dummy text of the printing and typesetting industry (see

1.2 Solvers

BFGS: https://stats.stackexchange.com/questions/284712/how-does-the-l-bfgs-work

Existing Approaches

This section will cover approaches taken so far to solve the problem. We will present algorithms that solve the approach, and the dataset that the algorithm was evaluated on, as well as their effectiveness.

2.1 Using a projection matrix

2.1.1 Active learning of linear subspace

[3] Assume f depends only on $u := xR^T$ with $R \in \mathbf{R}^{d \times D}$ where d << D. Learn an algorithm that learns $\hat{f}(u) = f(x)$ and R.

Steps include: 1.) Create a probability distribution over possible embeddings to learn R (Laplace approximation). 2.) Use this to create a probability distribution over f. 3.) Perform active selection over all possible points.

3.) is done using using Bayesian Active Learning by disagreement, where the utility function is the expected reduction in entropy (equal to the mutual information), as opposed to uncertainty sampling, which simply minimizes the entropy.

Tests are conducted on a real, and synthetic dataset with up to D=318 and selecting N=100 observations.

2.1.2 High dimensional Gaussian bandits

[1] Assume there exists a function $g : \mathbf{R}^k \Longrightarrow [0,1]$ and a matrix $A \in \mathbf{R}d \times D$ with orthogonal rows, such that f(x) = g(Ax). Assume $g \in \mathcal{C}^2$. Assume that $B = \mathbf{B}^D(1 + \varepsilon)$. We want to

maximize $f: B \Longrightarrow [0,1]$.

The SI-BO algorithm has a two-step approach: 1.) subspace identification. 2.) Bayesian Optimization on the learned subspace.

2.1.3 Random embeddings (REMBO)

[8] Let $x \in \mathbb{R}^D$ and $y \in \mathbb{R}^d$. Assume, that f(x) = f(Ax). We can generate $A \in \mathbb{R}^{D \times d}$ by randomly generating this matrix.

2.1.4 Applications to high-dimensional uncertainty propogation

[7] Assume $f(x) \approx g(\mathbf{W}^T y)$ where $\mathbf{W} \in \mathbb{R}^{D \times d}$ and D >> d. We assume that \mathbf{W} is orthogonal. This algorithm does not require gradient-information (thus, easier to implement, and robust to noise). The standard-deviation, kernel parameters and \mathbf{W} can be found iteratively. First we fix \mathbf{W} , and optimize over the standard-deviation, kernel parameters. Then we fix the standard-deviation, kernel parameters and optimize over \mathbf{W} . We repeat this procedure until the change of the log-likelihood between iterations is below some ε_l .

A more detailed description:

The quantities of interest are:

$$\mu_f = \int f(x)p(x)dx \tag{2.1}$$

$$\sigma_f^2 = \int (f(x) - \mu_f)^2 p(x) dx$$
 (2.2)

$$f \sim p(f) = \int \delta(f - f(x))p(x)dx \tag{2.3}$$

We further assume that the response surface has the form of

$$f(x) \approx g(W^T x)$$

Because all matrices yield identical approximations, one can focus on orthogonal approximations.

For the family of orthogonal matrices of dimension $d \times D$, we write $\mathbf{W} \in V_d(\mathbb{R}^D)$. This quantity is also known as the Stiefel manifold.

.

The paper focuses on identifying the low dimensional map $g(\dot{)}$, the effective dimensionality d and the orthogonal matrix W.

We determine the hyperparameters by optimizing over the following loss function, where Q are the input samples, t are the corresponding output samples, θ are the ..., and t are the ...

$$\mathcal{L}(\theta, s_n; Q, t) = \log p(t|Q, \theta, s_n)$$

where we find the individual variables by maximizing this log-likelihood.

The authors introduce a new kernel function

$$k_{AS}: \mathbb{R}^D \times \mathbb{R}^D \times V_d(\mathbb{R}^D) \times \phi - > \mathbb{R}$$
 (2.4)

where the kernel has the form

$$k_{AS}(x, x'; W, \phi) = k_d(W^T x, W^T x'; \phi)$$
 (2.5)

Within the paper, we use the following kernel function

$$1 (2.6)$$

Optimize $W \in V_d(\mathbb{R}^D)$ and keep ϕ and s_n^2 fixed

For this subsection, redefine the loss function as (ϕ are the hyperparameters of the covariance function):

(We instantiate θ with W)

$$F(W) := \mathcal{L}(W, s_n; X, y) \tag{2.7}$$

$$= \log p(y|X, W, s_n) \tag{2.8}$$

$$= -\frac{1}{2}(y-m)^{T}(K+s_{n}^{2}I_{N})^{-1}(y-m) - \frac{1}{2}\log|K+s_{n}^{2}I_{N}| - \frac{N}{2}\log 2\pi$$
 (2.9)

(2.10)

where ϕ , s_n ; X, y are fixed and m is the prior mean function.

The derivative of this function F with respect to the weights-matrix is:

$$\nabla_{W}F(W) := \nabla_{W}\mathcal{L}(W, s_{n}; X, y)$$

$$= \frac{1}{2} \text{tr} \left[\left\{ (K + s_{n}^{2}I_{N})^{-1} (y - m) \left((K + s_{n}^{2}I_{N})^{-1} (y - m) \right)^{T} - (K + s_{n}^{2}I_{N})^{-1} \right\} \nabla_{W} (K + s_{n}^{2}I_{N}) \right]$$
(2.11)

both these functions depend on the kernel K, and it's derivative $\nabla_W K$.

We use the 32-Matern kernel function (for two vectors a and b) with derivative:

$$K(a,b,\theta) = s^2 \left(1 + \sqrt{3} \sum_{i=1}^{l} \frac{(a_i - b_i)^2}{l_i} \right) exp \left(-\sqrt{3} \sum_{i=1}^{l} \frac{(a_i - b_i)^2}{l_i} \right)$$
(2.13)

$$\nabla_W K(a, b, \theta) = s^2 \left(1 + \sqrt{3} \sum_{i=1}^{l} \frac{(a_i - b_i)^2}{l_i} \right) exp \left(-\sqrt{3} \sum_{i=1}^{l} \frac{(a_i - b_i)^2}{l_i} \right)$$
(2.14)

where s, l_1, \dots, l_l are hyper-parameters included within θ

We optimize over this non-convex optimization problem by using a gradient-descent technique that optimizes over the Stiefel manifold (class of orthogonal matrices).

Identification of active subspace dimension

2.2 Exploiting additive structures within the function

2.2.1 Independent additive structures within the target function

[2] Assume that $f(x) = \sum_{i=1}^{|P|} f_i(x[P_i])$, i.e. f is fully additive, and can be represented as a sum of smaller-dimensional functions f_i , each of which accepts a subset of the input-variables. The kernel also results in an additive structure: $f(x) = \sum_{i=1}^{|P|} k_i(x[P_i], x[P_i])$. The posterior is calculated usin ghte Metropolis Hastings algorithm. The two actions for the sampling algorithm are 'Merge two subsets', and 'Split one set into two subsets'. k models are sampled, and we respectively approximate $p(f_*|D,x^*) = \frac{1}{k} \sum_{j=1}^k p(f(x^*|D,x,M_j))$, where M_j denotes the partition amongst all input-variables of the original function f.

2.2.2 Ideas

We could have a function sth like: $f(x) = g(Ax_1) + g(Bx_2) + ...$ as this does not assume that the dimensions are directly correlated. We could also use PCA or anything similar to choose the axis (i.e. A, B, C...)

2.3 Other approaches

2.3.1 Elastic Gaussian Processes

[5] Use a process where the space is iteratively explored. The key insight here is that with low length-scales, the acquisition function is extremely flat, but with higher length-scales, the acquisition function starts to have significant gradients. The two key-steps is to 1.) additively increase the length-scale for the gaussian process if the length-scale is not maximal and if $||x_{init} - x^*|| = 0$. And 2.) exponentially decrease the length-scale for the gaussian process if the length-scale is below the optimum length-scale and if $||x_{init} - x^*|| = 0$.

2.3.2 Bayesian Optimization using Dropout

[4] propose that the assumption of an active subspace is restrictive and often not fulfilled in real-world applications. They propose three algorithms, to iteratively optimize amongst certain dimensions that are not within the d 'most influential' dimensions: 1.) Dropout Random, which picks dimensions to be optimized at random, 2.) Dropout copy, which continuous optimizing the function values from the found local optimum configuration, and 3.) which does method 1. with probability p, and else method 2. The d 'most influential' dimensions are picked at random at each iteration.

2.3.3 Ideas

I think optimizing the hyper-parameters of Gaussian processes, and also the kernel is a valueable insight not much explored.

2.4 Datasets for benchmarking

1. [3] Synthetic in-model data matching the proposed model, with d = 2, 3, and D = 10, 20.

- 2. [3] (Synthetic) Braning function, d = 2, hidden in a higher dimensional space D = 10,20.
- 3. [3] Temperature data D = 106 and d = 2.
- 4. [3] Communities and Crime dataset d = 2, and D = 96.
- 5. [3] Relative location of CT slices on axial axis with d = 2 and D = 318.
- 6. [1] (Synthetic) Random GP samples from 2-dimensional Matern-Kernel-output, embedded within 100 dimensions
- 7. [1] Gabor Filters: Determine visual stimuli that maximally excite some neurons which reacts to edges in the image. We have $f(x) = \exp(-(\theta^T x 1)^2)$. θ is of size 17x17, and the set of admissible signals is d.
- 8. [8] (Synthethic) d = 2 and $D = 1 * 10^9$.
- 9. [8] D = 47 where each dimension is a parameter of a mixed integer linear programming solver.
- 10. [8] D = 14 with d for a random forest body part classifier.
- 11. [7] (Synthetic) Use d = 1, 10 and D = 10.
- 12. [7] (Half-synthetic) Stochastic elliptic partial differential equation, where D = 100, and an assume value for d of 1 or 2.
- 13. [7] Granular crystals $X \in \mathbb{R}^{1000 \times 2n_p + 1}$, and $y \in \mathbb{R}^{1000}$.
- 14. [2] (Synthetic) Styblinski–Tang function where *D* is freely choosable.
- 15. [2] (Synthetic) Michalewicz function where *D* is freely choosable.
- 16. [2] (Simulated) NASA cosmological constant data where D = 9.
- 17. [2] Simple matrix completion with D = 3.
- 18. [5] (Synthetic) Hertmann6d in [0, 1].
- 19. [5] (Synthetic) Unnormalized Gaussian PDF with a maximum of 1 in $[-1,1]^d$ for D = 20 and $[-0.5, 0.5]^d$ for D = 50
- 20. [5] (Synthetic) Generalized Rosenbrock function $[-5, 10]^d$

- 21. [5] Training cascade classifiers, with D = 10 per group.
- 22. [5] Optimizing alloys D = 13.
- 23. [4] (Synthetic) Gaussian mixture function
- 24. [4] (Synthetic) Schwefel's 1.2 function.
- 25. [?] A list of optimization test functions can be found here.
- 26. [?] A more comprehensive list of general functions can be found here.

Enumeration

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Experiments

3.1 Biggest competitors

Some main competitors are:

3.1.1 REMBO

3.1.2 Linear embeddings for Gaussian processes

And now I begin my third chapter here ...

And now to cite some more people??]

3.1.3 Synthetic Datasets

5 dimensional function with 2 dimensional linear embedding It is easy to test snythetic datasets, as we can evaluate these functions at any point, and immediately get a regret value.

2D to 1D: For this function, we use a simple Parabola which is embedded in a 2D space.

5D to 1D: For this function, we use a simple Parabola which is embedded in a 5D space.

5D to 2D: For this function, we use a simple Camelback function which is embedded in a 5D space.

10D to 5D: For this function, we use the 5D Rosenbrock function which is embedded in a 10D space.

12 Experiments

3.1.4 Real Datasets

It is more difficult to test algorithms on real datasets, as we cannot quite test a metric such as regret, but rather have to use log likelihood or something similar to test how well our algorithm adapts on a test set, given a training set.

SwissFEL dataset ... and some more

Implementing methods

4.1 Difficulties when implementing models

When implementing models, the following difficulties may arise:

- Overflows and underflow are often occuring as a reusult of high dimesionality
- Making sure the numerical examples are correct
- Making sure the differentiations are correct (use finite differences for matrices in this example)
- Write theoretical tests of the entire results
- test individual parts
- Test if the expectation

$$E[f(Ax) - f_hat(A_hatx)]$$

approaches zero

- Visualize the individual functions
- write a very simple test case of a parabola, and check where the points project to
- Performance of the implementation might not be optimal, as the design is only clear after implementation

4.2 Family of matrices

It turns out that there is a family of matrices that all project to the same space. As such, we need better measures to

• choose the matrix that we will take amongst all initialized tries (should be in the paper)

•

4.3 Methods from

And now I begin my third chapter here ...

And now to cite some more people??

4.3.1 First subsection in the first section

... and some more

Extensions

5.1 Finding a matrix when we apply a polynomial kernel onto the input first

We want to analyse if the gpregression can identify the random matrix, if the input is first put through a polynomial kernel of degree 2.

More specifically, the problem looks as follows. We want to approximate the real function f through a gaussian process g, with the following condition:

$$f\left(W\begin{bmatrix} (x-x_0)^2\\ (x-x_1)^2 \end{bmatrix}\right) \approx g(x) \tag{5.1}$$

where x_0, x_1 are constants.

5.2 Our proposed improvement to existing methods

We propose a novel method which is based on additive GPs, where we use different kinds of kernels.

We model the function f, which we approximate through functions g_i as follows:

$$f(x) \sim g_1(A_1x_1) + \sum_i g_i(A_ix_i)$$
 (5.2)

where x_1 is the subvector of the input x that includes the active subspace. And each x_i is a subvector of the input x which is part of the nullspace.

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Appendix A

How to install LATEX

Mac OS X

MacTeX - TeX distribution

- Download the file from https://www.tug.org/mactex/
- 2. Extract and double click to run the installer. It does the entire configuration, sit back and relax.