
Trajectory Kernel for Bayesian Optimization

Bachelor-Thesis von Sebastian Rinder aus Sindelfingen
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Vorgelegte Bachelor-Thesis von Sebastian Rinder aus Sindelfingen

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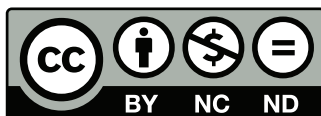
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Abstract

Reinforcement learning relies on policy gradient but the gradient is known only in expectation and most of the time stochastic policies. This leaves some room for zero order methods and BO can combine solving the problem and the exploration strategy from deterministic policies. We investigate in this paper how to integrate efficient exploration strategies stemming from Bayesian optimization for solving high dimensional reinforcement learning problems. We propose a novel optimization algorithm that is able to scale Bayesian optimization to such high dimensional tasks by restricting the search to the local vicinity of a search distribution and by proposing kernels capturing similarity in behavior rather than parameter. We show in the experiments that our approach can be very useful for applications such as robotics.

Zusammenfassung

Das Ziel im bestärkten Lernen ist das Finden einer Strategie, welche die erhaltene Belohnung eines Agenten maximiert. Da der Suchraum für mögliche Strategien sehr groß sein kann, verwenden wir Bayesian optimization, um die Anzahl der Evaluierungen durch den Agenten zu minimieren. Das hat den Vorteil, dass zeit- und kostenaufwändige Abläufe, wie beispielsweise das Bewegen eines Roboterarms, reduziert werden. Die Effektivität der Suche wird maßgeblich von der Wahl des Kernels beeinflusst. Standardkernel in der Bayesian optimization vergleichen die Parameter von Strategien um eine Vorhersage über bisher nicht evaluierte Strategien zu treffen.

Der Trajectorykernel vergleicht statt der Parameter, die aus den jeweiligen Strategien resultierenden Verhaltensweisen. Dadurch werden unterschiedliche Strategien mit ähnlichem Resultat von der Suche weniger priorisiert.

Wir zeigen die Überlegenheit des verhaltensbasierten Kernels gegenüber dem parameterbasierten anhand von Roboters-terierungssimulationen.

As a consequence the trajectory kernel makes the search process more efficient, but also has the disadvantage that the effort for kernel calculations is greatly increased.

The trajectory kernel relates policies derives a behaviour measure from the state and action parameter sets, to accordingly. This relation has the advantage that different policies with similar results are recognized as redundant and are therefore less prioritized by the search. As a consequence the trajectory kernel makes the search process more efficient, but also has the disadvantage that the effort for kernel calculations is greatly increased.

When it comes to higher dimensional problems Bayesian optimization tends to over explore because of its focus on global optimization. This can lead to not finding the optimum. We therefore also use Bayesian optimization with a locally restricted search area [1]. This area is adjusted throughout the search process to cover the most promising parts of the entire search space. As a result, we only have to optimize locally. This method is more robust against high-dimensional problems and also computationally less demanding than global optimization.

So we will compare the trajectory kernel to standard kernels in the Bayesian optimization. We also do test runs with global Bayesian optimization and Bayesian optimization in a local context. The reinforcement learning environment will be given by the classic control tasks cart pole, mountain car and acrobot [2].

Acknowledgments

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Symbols and Notation

Matrices and constants are noted as capital letters, and vectors and scalars as lower case letters. Vectors are assumed as column vectors. The first dimension of a matrix indicates its row number, the second dimension its column number.

<u>Symbol</u>	<u>Meaning</u>
S	Matrix of states
A	Matrix of actions
p	state transitioning probabilitiy
p_0	probabilitiy of starting in state s_0
r	immediate reward
\bar{r}	cumulative reward, sum of all rewards from one trajectory
M	number of initial sample points before starting the Bayesian optimization
N	number of total Bayesian optimization steps
n	number of training points present
n_*	number of sample test points
d	number of dimensions in the problem
x	training point vector of length d
x_*	test point vector of length d
X	$n \times d$ matrix of n points x^\top
X_*	$n_* \times d$ matrix of n_* test points x_*^\top
y	vector of n evaluated objective function values
$K(X, X) = K$	$n \times n$ covariance matrix
$K(X, X_*) = K_*$	$n \times n_*$ covariance between training and test points
$K(X_*, X) = K_*^\top$	same as $K(X, X_*)^\top$
$K(X_*, X_*) = K_{**}$	$n_* \times n_*$ covariance matrix
\circ	Hadamard product, element-wise product
$\text{diag}(V)$	returns the diagonal elements of the square matrix V as a vector
$\mathcal{O}(n)$	Big O notation - linear algorithm performance
$\mathcal{O}(n^2)$	Big O notation - quadratic algorithm performance

1 Introduction

The science of machine learning aims at creating well-functioning systems that are programmed to learn from data. This enables such systems to continuously improve their performance autonomously in order to accomplish a certain task. Machine learning is particularly important for tasks where manual design shows poor performance, such as image or speech recognition, effective web search or realization of self-driving cars. Considering the huge amount of data to be processed, machine learning algorithms easily surpass deterministic design. The advantage stems from generalizing learned examples. Accordingly, patterns in the learning data can be recognized and transferred to future situations without human supervision.

The goal of reinforcement learning, a sub-area of machine learning, is to teach an agent how to improve its behaviour [2]. In order to achieve a certain result, behaviour that contributes to the desired result is reinforced and behaviour that is not needed to achieve this result will be discouraged. The following example illustrates this. As an agent let us assume a robot arm with a table tennis racket and a ball. If our goal is to balance the ball for as long as possible, the agent will be penalized if it drops the ball, and rewarded for each successful time step of balancing. At the end of an evaluation episode all rewards received are summarized to get the cumulative reward as a measurement of performance.

The agent's behaviour can be defined by a set of policy parameters. These parameters tell the agent which action to perform depending on its current state. State parameters of the table tennis agent, for example, would contain the angle of the racket and the position of the ball. A well-tuned policy could tell the robot arm when and to what extent it should rotate in order to prevent the ball from falling. Finding such a well performing policy is the goal of our reinforcement learning task.

One approach of maximizing the performance of an agent could try random policies until finding a sufficient one. Unfortunately, this would take a very long time because it is highly unlikely to find suitable policy parameters by accident. Especially in more complex environments with higher dimensional policies. Besides the huge number of evaluations necessary, we would have to deal with wear and tear, and the limited movement speed of our robotic equipment [3].

For this reason, we need a more efficient search process to find properly functioning policies. A search process, which can learn from collected data, and extrapolate future behaviour. In recent years the Bayesian optimization approach has been shown to be very efficient in this field [4, 3, 5]. It tries to find an optimum of a black box function in as few steps as possible. In our reinforcement learning task the black box function would be the cumulative reward of an episode generated by a specific policy. In this case the optimum would be the policy with the highest cumulative reward achievable.

The modelling of the black box function and thus the prediction of future behaviour is based on measuring the distance between already evaluated policies with known results. The outcome of a new policy is then estimated by applying the learned distance metric. Out of many new policies Bayesian optimization selects one with the most promising enhancement. A good enhancement measure will include policies of uncharted areas in the search space and policies with high expected cumulative rewards as well. This balancing act is known as the exploration exploitation trade-off [4]. Only exploiting policies with a high expectation value can lead to a local optimum and thus to a result not as good as the best possible one. Whereas excessive exploration will cover the entire search space, but maybe will not find an optimum at all.

When using standard kernels, the distance between policies is usually measured by calculating the Euclidean distance [6] between corresponding policy parameters. A more significant measure would compare the resulting behaviour patterns stemming from certain policies rather than comparing the parameters of the policies. Such a behaviour measure is utilised by the trajectory kernel [7]. It uses the trajectory data generated by our agent during an evaluation process. This data, consisting of state and action values, relate the respective policies for modelling the black box function.

2 Foundations

2.1 Bayesian optimization

To find the maximum point of a costly black box target function, we use Bayesian optimization [4, 3, 5]. It makes the search process more efficient by incorporating a Gaussian process model to anticipate the objective function's behaviour. This model, containing estimates of the function's returns depending on points, is used by a so called acquisition function to guide the exploration for promising new points. Each selected point is evaluated by the objective function and the results are included in the Gaussian process model. Optimally these steps are repeated until found function values converge at the maximum possible function value. In reality (Algorithm 1) we iterate over N steps and analyse the results. The change of the Gaussian process model affecting the acquisition function during Bayesian optimization iterations is illustrated in Figure 2.1.

```
input :  $X = M$  uniformly random samples from the search space  
         $N$  = number of Bayesian optimization iterations  
        set hyper parameters to 1  
output:  $x_{M+N}$   
 $y$  = evaluations of the objective at the training points  $X$   
for  $n = M$  to  $M + N$  do  
    optimize hyper parameters (optional)  
    prepare a Gaussian process model depending on  $X$ ,  $y$  and hyper parameters  
     $x_{n+1}$  = point at the optimum of the acquisition function  
     $y_{n+1}$  = evaluation of the objective at the point  $x_{n+1}$   
     $X = \{X, x_{n+1}\}$   
     $y = \{y, y_{n+1}\}$ 
```

Algorithm 1: Global Bayesian optimization

The optimization of hyper parameters is optional and depends on the Gaussian process kernel as mentioned in chapter 6. Also the noise level parameter σ_n (2.1) is not considered as hyper parameter in this context. It will be tuned separately (see chapter 5).

2.1.1 Gaussian Process Regression

The Gaussian process finds a prior distribution over the possible functions that are consistent with the training data. From the regression we then get a posterior mean and variance, which describe our model of the objective function [6]. The mean represents a prediction of the true objective at a given point and the variance represents the uncertainty at that point. The more training points our model incorporates the smaller the variance, and the preciser the predictions, in the proximity around training points.

In real world applications we always have some noise in the objective observations. Therefore a Gaussian distributed error term,

$$\epsilon \sim \mathcal{N}(0, \sigma_n^2),$$

with zero mean and σ_n^2 variance is added to the function value. Ergo the observed target

$$y = f(x) + \epsilon$$

regards this noise [4, 6, 5, 3]. The knowledge our training data provides is represented by the covariance matrix $K(X, X)$. With the matrix of test points X_* we get the joint distribution of the target values and the estimated function values at the test locations [6]:

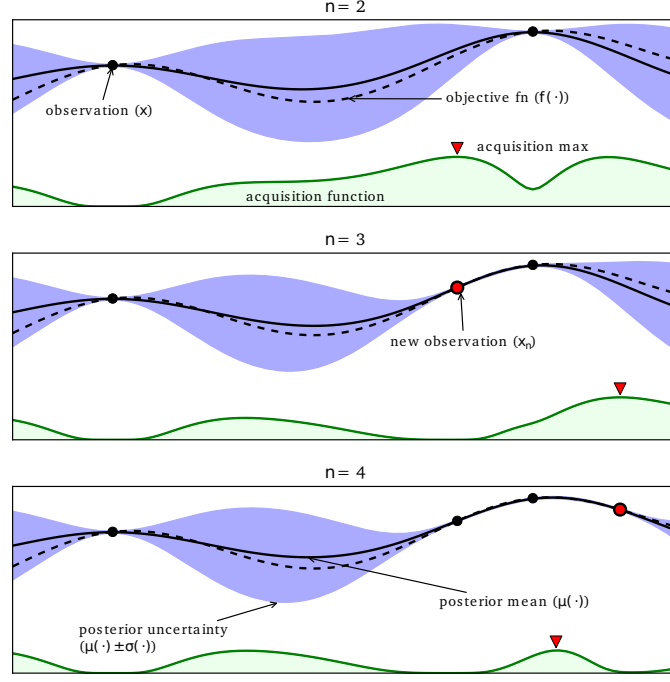


Figure 2.1.: Visualization of the change in the Gaussian process model and the resulting acquisition function [4]. The x-axis contains the search space over a one-dimensional example problem. With every Bayesian optimization iteration step a new observation point at the maximum of the acquisition function is added to the training data. This new observation and its evaluation update the Gaussian process model. Thus the model becomes increasingly accurate in the vicinity of each observation. The number of observations equals the number of iterations and is denoted by n .

$$\begin{bmatrix} y \\ f_* \end{bmatrix} \sim \mathcal{N} \left(0, \begin{bmatrix} K(X, X) + \sigma_n^2 I & K(X, X_*) \\ K(X_*, X) & K(X_*, X_*) \end{bmatrix} \right).$$

For further simplification we define $K = K(X, X)$, $K_* = K(X, X_*)$, $K_*^\top = K(X, X_*)^\top = K(X_*, X)$, and $K_{**} = K(X_*, X_*)$. Now we can calculate the posterior mean and variance at given test points X_* :

$$K_n = K + \sigma_n^2 I \quad (2.1)$$

$$\mu = K_* K_n^{-1} y \quad (2.2)$$

$$V(x_*, x_*) = K_{**} - K_* K_n^{-1} K_*^\top \quad (2.3)$$

$$v(x_*) = \text{diag}(V) \quad (2.4)$$

$$\sigma(x_*) = \sqrt{v(x_*)}. \quad (2.5)$$

The resulting vectors μ and v contain the means and variances for all corresponding test points. Also we get the whole covariance matrix V .

2.1.1.1 Standard kernels

The similarity between points is measured by the covariance function. The better this covariance function is suited for our objective function the more precise is the resulting model. We use two standard kernels [6] to compare them to the trajectory kernel (section 2.2.2). In those standard kernels the distance metric for two points is given by the Euclidean distance:

$$D(x_i, x_j) = (x_i - x_j)^\top (x_i - x_j).$$

Which we then use in the **squared exponential kernel**,

$$K(x_i, x_j, \sigma) = \sigma_f^2 \exp\left(-\frac{D(x_i, x_j)}{2\sigma_l^2}\right),$$

and the **Matérn 5/2 kernel**,

$$K(x_i, x_j, \sigma) = \sigma_f^2 \left(1 + \frac{\sqrt{5D}}{\sigma_l} + \frac{5D}{3\sigma_l^2}\right) \exp\left(-\frac{\sqrt{5D}}{\sigma_l}\right),$$

where σ_f denote the signal standard deviation and σ_l the characteristic length scale. These so called hyper parameters can be tuned by hand or set with hyper parameter optimization.

2.1.2 Hyper parameter optimization

Selecting proper hyper parameters for the Gaussian process regression can reduce the number of objective function evaluations necessary.

The likelihood function of the Gaussian process describes the probability of the Gaussian process model being suited for the black box function. To find an optimum for the signal deviation hyperparameter σ_f and the length scale hyperparameter σ_l we maximize the log likelihood function depending on those parameters [5]:

$$\log p(y = f|X, \sigma_f, \sigma_l) = -\frac{1}{2}y^\top K_n^{-1}y - \frac{1}{2}\log |K_n| - \frac{n}{2}\log 2\pi, \quad (2.6)$$

The number of observations is n , X is the $d \times n$ dataset of input points and K_n is the covariance matrix for the noisy target y .

2.1.3 Acquisition function

The basis of the Bayesian optimization consists of selecting the next evaluation point in our search space. This point is at the optimum of the acquisition function, which depends on the current Gaussian process model. We choose expected improvement [4] during the global Bayesian optimization and in the local optimization (section 3.1) we use Thompson sampling [1]. Both acquisition functions take the mean and the variance generated by the Gaussian process model into account to guide the exploration process. The difficulty lies in avoiding excessive exploration or exploitation. Exploration seeks points with a high variance and exploitation selects points with a high mean instead. The latter one would result in a local optimum whereas too much exploration may not improve at all. An exemplary comparison of acquisition function is shown in Figure 2.2.

2.1.3.1 Expected improvement

To get an expected improvement function value at a test point x_* , we need the mean value $\mu(x_*)$, and the standard deviation value $\sigma(x_*) = \sqrt{v(x_*)}$ from the Gaussian process model. Also we need the maximum of all observations y_{max} and a trade-off parameter τ . For the cumulative distribution function we write $\Phi(\cdot)$ and for the probability density function we write $\phi(\cdot)$. They are both Gaussian with zero mean and unit variance. We adopt the expected improvement function

$$EI(x_*) = \begin{cases} (\mu(x_*) - y_{max} - \tau)\Phi(z(x_*)) + \sigma(x_*)\phi(z(x_*)) & \text{if } \sigma(x_*) > 0 \\ 0 & \text{if } \sigma(x_*) = 0 \end{cases}$$

where

$$z(x_*) = \begin{cases} \frac{(\mu(x_*) - y_{max} - \tau)}{\sigma(x_*)} & \text{if } \sigma(x_*) > 0 \\ 0 & \text{if } \sigma(x_*) = 0 \end{cases}$$

as suggested in [4]. The trade-off parameter τ can be set to zero or above to gain more exploration. To get the next evaluation point,

$$x_{n+1} = \arg \max_{x_*} EI(x_*),$$

we optimize the expected improvement function over the whole search space.

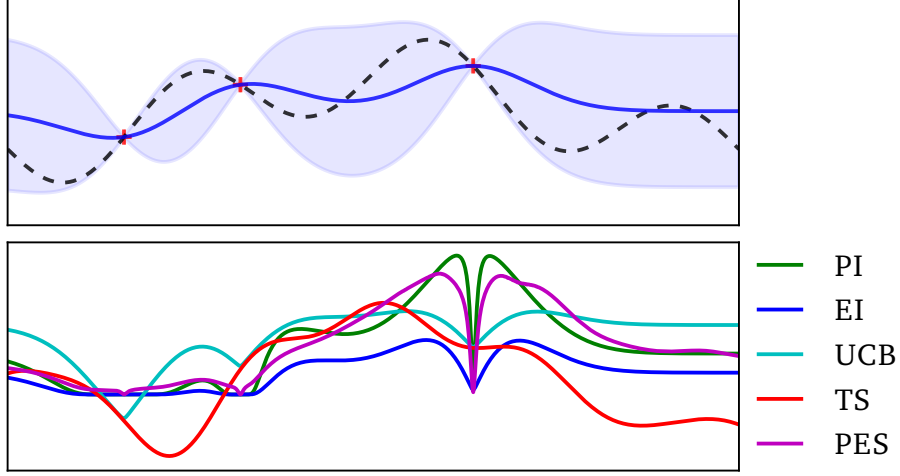


Figure 2.2.: Visualization of Gaussian process posterior values corresponding to different acquisition functions [3]. The x-axis covers the search space over a one-dimensional example problem. In the upper frame the posterior mean μ and double standard deviation 2σ is marked as the blue line with the shaded area. The black dashed line shows the true objective and the red crosses mark already evaluated points. The lower frame compares the plots of the resulting acquisition functions probability of improvement, expected improvement, upper confidence bounds, Thompson sampling and predictive entropy search. The peaks of probability of improvement next to values with high mean illustrates its exploitive behaviour compared to expected improvement, which takes the variance more into account. Thompson sampling is the one with the most explorative peak due to its random nature.

2.1.3.2 Thompson sampling

For Thompson sampling acquisition values we sample one function from the Gaussian process posterior [3],

$$\text{TS} \sim \text{GP}(0, K(X, X_*)),$$

where X is the dataset of already evaluated points, and X_* is a randomly Gaussian distributed set of points with mean and variance given by the local optimizer. These mean and variance represent our current search space. To draw function values we need the mean vector μ and the full covariance matrix V from the Gaussian process model. First we take the lower Cholesky decomposite of V such that $L_V L_V^\top = V$. Then we generate a vector g , which consists of independent Gaussian distributed values with zero mean and unit variance. Finally we get a vector TS of sampled values:

$$\text{TS}(x_*) = \mu + L_V g.$$

We take the one with the highest value such that,

$$x_{n+1} = \arg \max_{x_*} \text{TS}(x_*),$$

to get the next evaluation point.

2.2 Reinforcement learning

In our reinforcement learning task we want to maximize the cumulative reward stemming from a policy with as few evaluations as feasible. We achieve this maximization by applying Bayesian optimization, treating the cumulative reward as the objective function and policies as points. The objective function evaluation consists of the simulation of a robotic agent.

2.2.1 Markov decision process

We use the Markov decision process model to describe the agent's decisions making [2]. This model consists of a tuple (S, A, P, R, γ) , holding all states $s \in S$, all actions $a \in A$, all state transitioning probabilities $p \in P$, and all corresponding rewards $r \in R$. The discount factor γ is set to 1 because we have a finite horizon Markov decision process where the timing of the reward is irrelevant. Since the transitioning probabilities and the rewards are unknown, they are accessed through simulation runs of our environment. Assume the agent executing a policy x for T time steps, producing a trajectory ξ , and therefore receiving the final reward

$$\bar{r}(\xi) = \sum_{t=1}^T \gamma^t r(s_{t-1}, a_{t-1}, s_t),$$

as the sum of all immediate rewards given by a rewarding function $r(s_{t-1}, a_{t-1}, s_t)$. The explicit probabilities corresponding to actions and state transitionings are also obtained (see 4.1.5) throughout the simulation run.

2.2.2 Trajectory kernel

Standard kernels like the squared exponential kernel, relate policies by measuring the difference between policy parameter values. Therefore policies with similar behavior but different parameters are not compared adequately. The behavior based trajectory kernel fixes this, by relating policies to their resulting behavior [7]. This makes our policy search more efficient, since we avoid redundant search of different policies with similar behaviour.

2.2.2.1 Behaviour based measurement

For relating policies to their resulting behavior we use the Markov decision process transitioning probabilities. We formulate the conditional probability of observing trajectory ξ given policy x as proposed in [7]:

$$p(\xi|x) = p_0(s_0) \prod_{t=1}^T p(s_t|s_{t-1}, a_{t-1}) p_\pi(a_{t-1}|s_{t-1}, x).$$

Trajectory $\xi = (s_0, a_0, \dots, s_{T-1}, a_{T-1}, s_T)$ contains the sequence of state, action tuples and policy x a set of d parameters. $p_0(s_0)$ is the probability of starting in the initial state s_0 . $p(s_t|s_{t-1}, a_{t-1})$ is the probability of transitioning from state s_{t-1} to s_t when action a_{t-1} is executed. The stochastic mapping $p_\pi(a_{t-1}|s_{t-1}, x)$ is the probability for selecting the action a_{t-1} when in state s_{t-1} and executing the parametric policy x .

2.2.2.2 Distance metric

To examine the difference between two policies x_i and x_j the discrete Kullback Leibler divergence

$$\text{KL}(P(\xi|x_i)||P(\xi|x_j)) = \sum_i P(\xi|x_i) \log \frac{P(\xi|x_i)}{P(\xi|x_j)}.$$

is applied to the policy-trajectory mapping probabilities $P(\xi|x_i)$ and $P(\xi|x_j)$ [7]. The Kullback Leibler divergence measures how the two distributions diverge from another.

In general $\text{KL}(P(\xi|x_i)||P(\xi|x_j))$ is not equal to $\text{KL}(P(\xi|x_j)||P(\xi|x_i))$. But we need a symmetric distance measure. So we sum up the two divergences

$$D(x_i, x_j) = \text{KL}(P(\xi|x_i)||P(\xi|x_j)) + \text{KL}(P(\xi|x_j)||P(\xi|x_i)),$$

to achieve that $D(x_i, x_j) = D(x_j, x_i)$. An additional requirement for the kernel is the resulting matrix to be positive semi-definite and scalable[7]. Therefore we exponentiate the negative of our distance matrix D . We also apply the hyper parameters σ_f^2 to compensate for the signal variance and σ_l^2 to adjust for signal length scale. This gives us the covariance function

$$K(x_i, x_j, \sigma_f, \sigma_l) = \sigma_f^2 \exp\left(-\frac{D(x_i, x_j)}{2\sigma_l^2}\right). \quad (2.7)$$

In addition to the distance metric proposed by [7] we included the signal variance hyper parameter σ_f^2 .

2.2.2.3 Estimation of Trajectory Kernel Values

To reduce the computational effort an estimation of kernel values is applied. The approximation with the Monte-Carlo estimate

$$\hat{D}(x_i, x_j) = \sum_{\xi \in \xi_i} \log \left(\frac{P(\xi|x_i)}{P(\xi|x_j)} \right) + \sum_{\xi \in \xi_j} \log \left(\frac{P(\xi|x_j)}{P(\xi|x_i)} \right) \quad (2.8)$$

is sufficient to measure the divergences between policies with already sampled trajectories [7]. Here ξ_i is the set of trajectories generated by policy x_i . For our Gaussian process regression we also need a distance measure between a policy with known trajectories and new policies with unknown trajectories. Since there is no closed form solution to this we use the importance sampled divergence

$$\hat{D}(x_{new}, x_j) = \sum_{\xi \in \xi_j} \left[\frac{P(\xi|x_{new})}{P(\xi|x_j)} \log \left(\frac{P(\xi|x_{new})}{P(\xi|x_j)} \right) + \log \left(\frac{P(\xi|x_j)}{P(\xi|x_{new})} \right) \right]$$

to estimate the divergence between the new policy x_{new} and the policy x_j with already sampled trajectories ξ_j [7]. Since we only have a ratio of transitioning probabilities present in our trajectory kernel we can reduce the logarithmic term to:

$$\begin{aligned} \log \left(\frac{P(\xi|x_i)}{P(\xi|x_j)} \right) &= \log \left(\frac{P_0(s_0) \prod_{t=1}^T P_s(s_t|s_{t-1}, a_{t-1}) P_\pi(a_{t-1}|s_{t-1}, x_i)}{P_0(s_0) \prod_{t=1}^T P_s(s_t|s_{t-1}, a_{t-1}) P_\pi(a_{t-1}|s_{t-1}, x_j)} \right) \\ &= \log \left(\prod_{t=1}^T \frac{P_\pi(a_{t-1}|s_{t-1}, x_i)}{P_\pi(a_{t-1}|s_{t-1}, x_j)} \right) \\ &= \sum_{t=1}^T \log \left(\frac{P_\pi(a_{t-1}|s_{t-1}, x_i)}{P_\pi(a_{t-1}|s_{t-1}, x_j)} \right). \end{aligned}$$

Summing up the logarithms of the probability ratios in the end is also numerically more stable than computing the logarithm of the products.

3 Contributions

3.1 Local Bayesian optimization

- formula for search dist update

Modelling the objective function for a higher dimensional search space is challenging. Also global Bayesian optimization tends to over-explore. To perform a more robust optimization we use local Bayesian optimization as stated in [1]. It restricts the search space of the acquisition function to a local area which is moved, resized, and rotated between iterations. This local area is defined by a Gaussian distribution in which the mean and variance represent the center and the exploration reach respectively. To update that mean and variance properly we minimize the Kullback-Leibler divergence between the current search distribution π_n and the probability $p_n^* = p(\mathbf{x} = \mathbf{x}^* | \mathcal{D}_n)$ of \mathbf{x}^* being optimal. This results in a search area which neglects poorly performing regions.

To prevent the mean from moving too fast from the initial point and to avoid the variance becoming too small quickly the minimization is constrained with the hyper parameters α and β . Therefore our optimization problem is given by

$$\begin{aligned} \arg \min_{\pi} \quad & \text{KL}(\pi || p_n^*), \\ \text{subject to} \quad & \text{KL}(\pi || \pi_n) \leq \alpha, \end{aligned} \quad (3.1)$$

$$\mathcal{H}(\pi_n) - \mathcal{H}(\pi) \leq \beta, \quad (3.2)$$

where $\text{KL}(\cdot || \cdot)$ denotes the Kullback Leibler divergence $\mathcal{H}(p) = - \int p(\mathbf{x}) \log(p(\mathbf{x})) d\mathbf{x}$ is the entropy of p . We implement the local Bayesian optimization (Algorithm 2) as suggested by [1]

3.1.1 Thompson sampling

To perform Thompson sampling on the variable search distribution we first sample M points from the current search distribution (Algorithm 3). Then we compute the mahalanobis distance for each test point to discard samples, which are at the outer edge of the search space. The mahalanobis distance is used to regard the covariance of the current search distribution.

3.1.2 Trajectory kernel

If using Thompson sampling as the acquisition function, the whole covariance matrix V of test points is needed. To compute this covariance matrix we need a distance measure between unknown trajectories. The kernel proposed by [7] does not support a measurement between unevaluated policies. We therefore implement our own distance metric based on states, which are already present in the training data. First we filter a random subset from the training data and save it to S_{sub} . For the continuous action space we then calculate each mean

$$\mu_i = f_s(S_{\text{sub}})^\top x_i \quad \text{and} \quad \mu_j = f_s(S_{\text{sub}})^\top x_j$$

for the given policies x_i and x_j . The difference between the two policies is measured by the squared Euclidean distance between the means:

$$D(x_i, x_j) = (\mu_i - \mu_j)^\top (\mu_i - \mu_j).$$

For the discrete action space we sample the actions a from $P(a | S_{\text{sub}}, x)$ for the given policies x_i and x_j from (4.3). The sampled actions then select the associated probabilities. We compare the discrete probability distributions for each policy by applying the Kullback-Leibler twice to get a symmetric distance measure:

$$D(x_i, x_j) = \sum_i P(a | S_{\text{sub}}, x_i) \log \frac{P(a | S_{\text{sub}}, x_i)}{P(a | S_{\text{sub}}, x_j)} + \sum_j P(a | S_{\text{sub}}, x_j) \log \frac{P(a | S_{\text{sub}}, x_j)}{P(a | S_{\text{sub}}, x_i)}.$$

```

input : Initial search policy  $\pi_0 = \mathcal{N}(\boldsymbol{\mu}_0, \sigma_0^2 I)$ 
         step-size  $\epsilon$ 
         entropy reduction rate  $\beta$ 
output: Search policy  $\pi_N$ 

for  $n = 1$  to  $N$  do
    Fit: Gaussian  $\hat{p}_n$  from local samples of  $p_n^*$ 
    Optimize:  $(\eta^*, \omega^*) = \arg \min g_n(\eta, \omega)$ 
    Bayesian Update:  $(\pi_{n+1})^{\eta^* + \omega^*} \propto \pi_n^{\eta^*} p_n^*$ 
    Evaluate:  $\mathbf{x}_n$  from local samples of  $p_n^*$ 
     $\mathcal{D}_n \leftarrow \mathcal{D}_{n-1} \cup \{(\mathbf{x}_n, y_n)\}$ 

```

Algorithm 2: Local Bayesian Optimization

```

input :  $X = N$  already evaluated points
          $X_* = M$  random samples from the search distribution
          $d_t =$  distance threshold
output:  $x_{n+1}$ : next evaluation point

compute the mahalanobis distance from test points to current search distribution
keep samples, which are inside  $d_t$ % of the distribution density
get mean and covariance matrix from the Gaussian process
get values from Thompson sampling
 $x_{n+1} =$  sample at the highest Thompson sampled value

```

Algorithm 3: Thompson sampling acquisition for local Bayesian optimization

3.2 Numerical stability and efficiency

Due to numerical instabilities, errors may occur during calculations. These errors are caused by the discrete implementation of continuous mathematical formulas. Sometimes numerical instabilities lead to negative values for covariances. To avoid getting complex numbers we assume negative values as zero before applying the square root on covariances when computing the standard deviation from the Gaussian process model (2.5). Negative values can also occur for the distance between two trajectories, and are thresholded to zero too.

When computing the symmetrical distance matrix $\hat{D}(X, X)$ (2.8) only the upper triangle matrix \hat{D}_u is calculated. That halves the computational effort. We obtain $\hat{D} = \hat{D}_u + \hat{D}_u^\top$, since the diagonal elements of \hat{D} are zero and $\hat{D}(x_i, x_j) = \hat{D}(x_j, x_i)$.

To gain robustness we use the lower Cholesky decomposition instead doing matrix inverse calculation. Therefore the matrix to decompose must be positive definite. We achieve this by doubling the noise variance added to the diagonal elements of the original matrix as shown in algorithm 4.

```

input :  $K, \sigma_n$ 
output:  $L$ 

 $K_n = K + \sigma_n^2 I$ 
while  $K_n$  not positive definite do
    double  $\sigma_n^2$ 
     $K_n = K + \sigma_n^2 I$ 
 $L =$  lower Cholesky of  $K_n$ 

```

Algorithm 4: Lower Cholesky with variance doubling

When doing hyper parameter optimization we do not double the noise variance. Instead our log marginal likelihood function (2.6) returns $-\infty$ for hyper parameters, which produce a non positive definit matrix K_n . Therefore, such hyper parameters do not come into consideration when maximizing.

3.2.1 Gaussian Process Regression

3.2.1.1 Inverse of prior covariance matrix

Instead of calculating the inverse of K_n in (2.2) we use the lower Cholesky decomposed matrix:

$$LL^\top = K_n$$

This is considered faster and numerically more stable [6]. The mean vector μ is then computed as follows:

$$\mu = K_n^{-1} y = (L L^\top)^{-1} y = (L^{-\top} L^{-1}) y = L^{-\top} (L^{-1} y) = L^\top \setminus (L \setminus y). \quad (3.3)$$

The backslash operator denotes the matrix left division, so the solution $x = A \setminus b$ satisfies the system of linear equations $Ax = b$. Matrix K_n must be positive definite for the cholesky decomposition. So we double the noise variance hyperparameter σ_n^2 as shown in Algorithm 4.

3.2.1.2 Variances

For the expected improvement function we only need the vector of variances. Instead of calculating the whole covariance matrix V and taking the diagonal elements (2.4) we can take a shortcut. All elements on the diagonal of $K(X_*, X_*)$ equal σ_f^2 because the difference between one x_* and the same x_* is zero. Therefore we can write:

$$L_k = L \setminus K(X_*, X) \quad (3.4)$$

$$v = \sigma_f^2 - \sum_{\text{rows}} (L_k \circ L_k). \quad (3.5)$$

This adaptation is inspired by [8] and reduces the computational effort drastically from $\mathcal{O}(n^2)$ to $\mathcal{O}(n)$. For the whole covariance matrix for Thompson sampling we also avoid calculating the matrix inverse:

$$V = K(X_*, X_*) - (L_k^\top L_k)^\top$$

TODO WRITE WHOLE MODIFIED GP ALGO BOX

3.2.2 Action selection

In continuous action space our stochastic policy is Gaussian distributed (4.1). Therefore the resulting probability density of the action selection,

$$P_\pi(a|s, x) = \frac{1}{\sqrt{2\pi\sigma_a^2}} \exp\left(-\frac{(a - f_s(s)x)^2}{2\sigma_a^2}\right),$$

allows us to do the computations of the logarithm of the probability ratios in the trajectory kernel more efficient:

$$\begin{aligned} \sum_{t=0}^{T-1} \log\left(\frac{P_\pi(a_t|s_t, x_i)}{P_\pi(a_t|s_t, x_j)}\right) &= \sum_{t=0}^{T-1} \log\left(\frac{\frac{1}{\sqrt{2\pi\sigma_a^2}} \exp\left(-\frac{(a_t - f_s(s_t)x_i)^2}{2\sigma_a^2}\right)}{\frac{1}{\sqrt{2\pi\sigma_a^2}} \exp\left(-\frac{(a_t - f_s(s_t)x_j)^2}{2\sigma_a^2}\right)}\right) \\ &= \sum_{t=0}^{T-1} \log\left(\exp\left(-\frac{(a_t - f_s(s_t)x_i)^2}{2\sigma_a^2}\right) - \left(-\frac{(a_t - f_s(s_t)x_j)^2}{2\sigma_a^2}\right)\right) \\ &= \frac{1}{2\sigma_a^2} \sum_{t=0}^{T-1} ((a_t - f_s(s_t)x_j)^2 - (a_t - f_s(s_t)x_i)^2). \end{aligned}$$

3.2.3 Hyper parameter optimization

Especially for the trajectory kernel we want a hyper parameter optimization, because all the values of the distance matrix D may get very big. When dividing by a well tuned hyper parameter σ_l before applying the exponential function (2.7), we avoid getting a diagonal matrix K or a zero matrix K_* .

When calculating $\log(|K_n|)$ for the hyperparameter optimization (2.6), again we use the Cholesky decomposition of K . Thus the determinant transforms to

$$|K_n| = |L L^T| = |L| |L^T| = |L| |L| = |L|^2.$$

Since the determinant of the Cholesky decomposed matrix,

$$|L| = \prod_i L_{ii},$$

is the product of its diagonal elements, we can transform this into a numerically more stable version:

$$\log(|K_n|) = \log(|L|^2) = 2 \log(|L|) = 2 \log(\prod_i L_{ii}) = 2 \sum_i \log(L_{ii}).$$

The computation of $K_y^{-1}y$ in (2.6) is done by the same method we already use in the Gaussian process (3.3).

3.2.3.1 Independent log-normal prior

The log marginal likelihood maximization will sometimes succeed at the borders of our search space resulting in very high high or very low hyper parameters. We prevent this by adding an independent log-normal prior term as introduced in [5]:

$$\sum_{i=f,l} \left(\frac{-\log(\sigma_i)^2}{2 \cdot 10^2} - \log 10\sqrt{2\pi} \right).$$

Since the suggested prior is centered at the point of origin with standard deviation 10, but we want our search space to adapt between iterations, we make the mean and the standard deviation adjustable. We write:

$$\sum_{i=1,2} \left(\frac{-(h_i - c_i)^2}{2(b_{u_i} - b_{l_i})^2} - \log(b_{u_i} - b_{l_i})\sqrt{2\pi} \right),$$

where b_l and b_u denote the lower and upper bounds of the search space and c its center. And since we optimize over the logarithmic space $h_1 = \log \sigma_f$ and $h_2 = \log \sigma_l$.

TODO write whole hyper opt with L and stuff

4 Experiments

4.1 Implementation

4.1.1 Efficiency

To cut the time for experiment runs we vectorized every suited operation to speed up calculations in Matlab. Additionally we tuned the code to run on a parallel pool.

4.1.2 Global optimization

The expected improvement function and the log marginal likelihood function for hyper parameter optimization need to be maximized. Since we use Matlab's GlobalSearch Toolbox, which is a global minimizer, we multiply the results of the functions by -1 to be able to minimize them appropriately.

4.1.3 Optimization starting point

When optimizing the log marginal likelihood of the gaussian process we are confronted with undefined areas in the hyper parameter space because K is not positive semi definit in those areas. To start the optimization properly we randomly select points until finding a defined one.

4.1.4 Problem environment

We use Matlab implementations of Cart Pole, Acrobot, and Mountain Car that were adapted from [9]. We also implement the OpenAI Gym [10] to provide a wide range of problem environments. To use the simulations provided by the OpenAI Gym we prepare a python module which is imported to MATLAB. After loading the python module with `py.importlib.import_module(moduleName)` we can call every method it contains via the `py.moduleName.` prefix. It is vital to convert our data correctly before calling the python subroutine with it. The Matlab variables containing whole numbers are converted from Double to Int. Also all the vectors received by the python module have to be converted to an array through `numpy.asarray()`. Unfortunately, the OpenAI Gym environments do not work if we use the parallelization of MATLAB. So we used these environments only with the less computationally demanding local Bayesian optimization.

4.1.5 Action selection

In continuous action space we use a linear policy to action mapping

$$a = f_s(s)^\top x + \sigma_a, \quad (4.1)$$

with a small gaussian noise σ_a needed for stochastic policies. So the Gaussian distributed actions,

$$a \sim \mathcal{N}(f_s(s)x, \sigma_a^2),$$

result in the probability measure by the Gaussian density:

$$P_\pi(a|s, x) = \frac{1}{\sqrt{2\pi\sigma_a^2}} \exp\left(-\frac{(a - f_s(s)x)^2}{2\sigma_a^2}\right). \quad (4.2)$$

In discrete action space environments we use a parametric soft-max action selection policy:

$$P(a|s, x) = \frac{\exp(f_s(s)^\top x_a)}{\sum_{a \in A} \exp(f_s(s)^\top x_a)}. \quad (4.3)$$

Again it consists of the linear mapping $f_s(s)^\top x$, and A holds all possible actions. The resulting action is then sampled from the probability of action a given state s .

The state feature function $f_s(s)$ depends on the environment as listed in Table 4.1.

4.2 Start conditions

For the trajectory kernel, we proposed in section 3.1.2, a maximum of 500 random states as the subset S_{sub}

For the continuous action selection the standard error deviation σ_a is set to 10^{-3} . We adapted the action selection strategies from [7]. Unfortunately they do not provide any parameters. Therefore this error deviation value was guessed after a few test runs.

4.2.1 Global context

In the global search context the policy parameter boundaries are set to -10 and 10 for each dimension resulting in a d dimensional hypercubic search space.

For the Bayesian optimization we select the initial samples set X_n from a bunch of sample sets such that the Euclidean distance between points of the selected set is maximized. This grants us a well covered search space as a starting condition. We also set the number of initial samples to $M = 10$ and the iteration count to $N = 200$.

In the expected improvement function, the trade-off parameter τ is set to 0.01 as proposed by [4].

The hyper parameter optimization runs every fifth Bayesian optimization step or if the maximum of the expected improvement function drops below 10^{-6} .

4.2.2 Local context

The Bayesian optimization in the local context starts with a hypersphere centered at the origin with radius 10 as the policy search space boundary. The starting point for the search is set to the origin accordingly.

We run the hyper parameter optimization every time the search space is adjusted. We carry out a total of 400 Bayesian optimization steps, in which the search space is adapted every four steps.

Before doing Gaussian process regression we transform our observations to zero mean and uniform variance:

$$y = \frac{y_n - \text{mean}(y_n)}{\text{std}(y_n)}.$$

This standardization proposed by [1] affects the Thompson sampling exploration exploitation trade-off. The distance threshold d_t (Algorithm 3) for the samples during Thompson sampling is set to 80%.

4.3 Environments

4.3.1 Cart Pole

The Cart Pole environment consists of a cart with a pole attached to its top. The cart is accelerated to the left or the right to balance the pole for as long as possible. An episode starts with small random state values and it ends when the maximum of time steps or a specific state is reached. The state values contain the position and the velocity of the cart, and the angle and the angular velocity of the pole. Each of these four state values are set uniformly random between -0.05 and 0.05 at the beginning. The episode ends after 200 time steps or if the pole falls below 12 degrees relative to the vertical. It also ends when the cart is more than 2.4 meters away from the center. For each completed time step a reward of 1 is returned. The cart pole implementation from Open AI Gym (Figure 5.2) accepts a discrete action value, -1 or 1, for either applying force to the left or to the right. With the four state values and an additional bias value we have five dimensions for each discrete action resulting in a total of ten dimensions. In our own cart pole simulation we only use the four state parameters for calculating a continuous action value between -1 and 1. It turned out that adding a bias value would make the policy search slightly more difficult here. Therefore this policy has only four dimensions.

Environment	Description	Action	State feature	Policy dimensions d
Cart Pole	p : cart position θ : pole angle	continuous: $[-1,1]$	$(p, \dot{p}, \theta, \dot{\theta})$	4
		discrete: $\{-1,1\}$	$(p, \dot{p}, \theta, \dot{\theta}, 1)$	10
Acrobot	θ_1 : angle of pendulum 1 θ_2 : angle between pendulum 1 and 2	discrete: $\{-1,0,1\}$	$(\cos(\theta_1), \sin(\theta_1), \cos(\theta_2), \sin(\theta_2), \dot{\theta}_1, \dot{\theta}_2, 1)$	21
Mountain Car	p : horizontal car position u : car velocity	continuous: $[-1,1]$	$(p, u, p^2, u^2, pu, p^2u, pu^2, p^3, u^3, 1)$	10
		discrete: $\{-1,0,1\}$	$(p, u, p^2, u^2, pu, p^2u, pu^2, p^3, u^3, 1)$	30

Table 4.1.: Environment parameters of the experiments

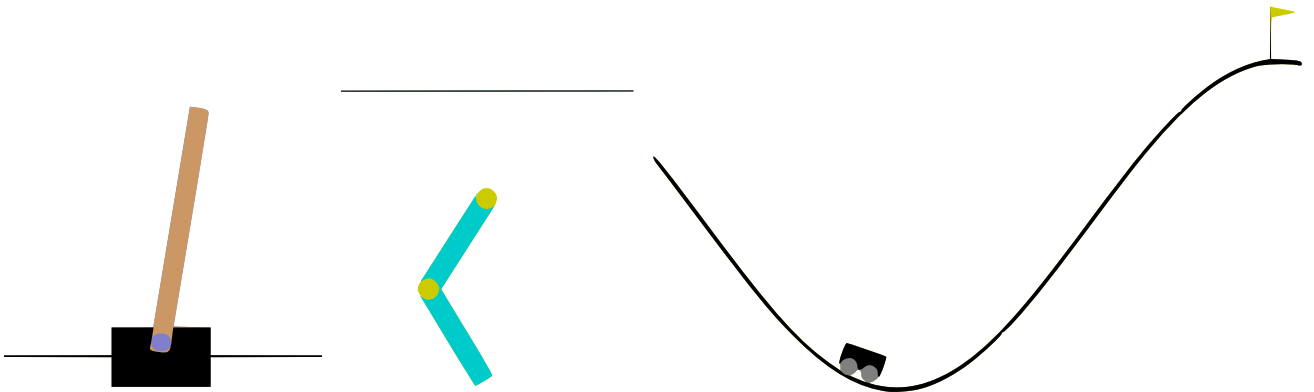


Figure 4.1.: BLABLABLAH

4.3.2 Acrobot

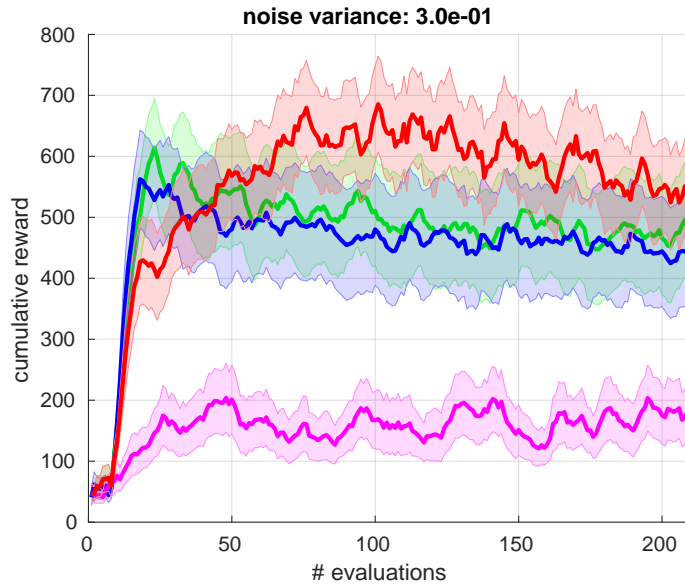
The acrobatic robot has two joints and two pendulums. The upper joint has a fixed position. It connects to the first pendulum, which is attached to the second joint. A torque can be applied to that second joint, which controls the second pendulum. The pendulums start in equilibrium position and the goal is to gain enough momentum and swing the end of the second pendulum above a certain mark. The angle between vertical and the first pendulum and the angle between the two pendulums form the six state parameters. Four parameters are the sine and the cosine of these two angles each, whereas the last two parameters are the angular velocities 4.1. The discrete action can have the values -1, 0 or 1. Each value stands for the amount of torque applied to the joint between the two pendulum links. With the six state values, an additional bias value, and three discrete action possibilities we get 21 dimensions for the Acrobot policy.

4.3.3 Mountain car

During the Mountain Car task, an underpowered car tries to drive uphill. It can only reach the goal on the right side if it uses both hills to gain momentum. The end of the left hill is an inelastic wall, so if the car hits it, the velocity is set to zero. Since the hills are formed by a sine curve, it is sufficient that the position of the car is only one-dimensional. The vertical position of the vehicle can be easily derived from the horizontal position with the sine function. The horizontal position and the car's velocity are cubically expanded (Table 4.1) to get nine state values. Also a bias parameter is added as a tenth state value. In the continuous setting, we therefore receive ten dimensions. In the discrete setting we have three action values for either applying acceleration to the left or right (-1,1) or doing nothing (0). In this case we receive 30 dimensions.

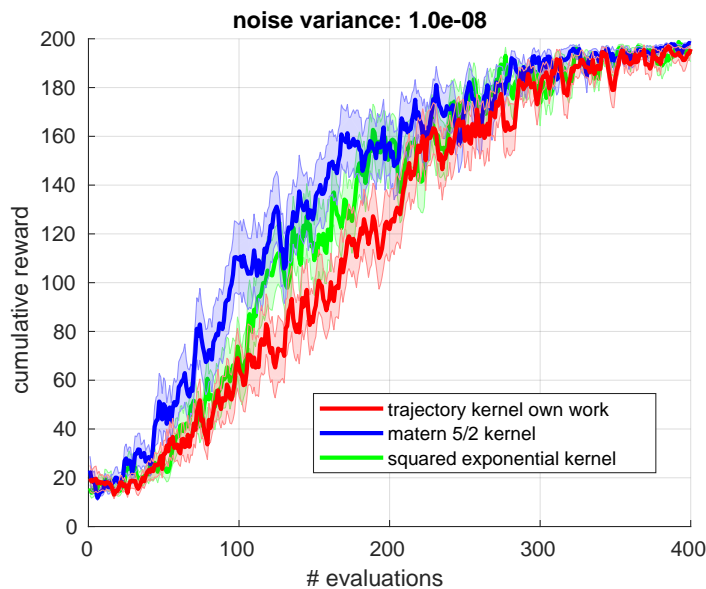
5 Results

We plot the mean and the standard deviation for all kernel runs. One run consists of 10 or 32 trials depending on the experiment. 32 trials are used on cluster runs, because we can use 32 cores to parallelize the run. Due to high variations between iteration steps, all plots are smoothed by a moving mean of five. Also, the standard deviation values are divided by five to maintain readability when three or four different kernel graphs are present in one plot.



kernel		mean(time)	std(time)
trajectory	(own work)	874.0	261.0
trajectory	(Wilson 2014)	201.1	80.0
matérn 5/2		20.7	4.8
squared exponential		19.1	4.9

Figure 5.1.: Results of the MATLAB Cart Pole implementation with a continuous action selection policy with the local optimization. Each kernel graph contains the average over 10 trials.



kernel		mean(time)	std(time)
trajectory	(own work)	61.6	3.5
matérn 5/2		8.8	1.4
squared exponential		6.4	0.6

Figure 5.2.: Results of the Open AI Cart Pole implementation with a discrete action selection policy with the local optimization. Each kernel graph contains the average over 10 trials.

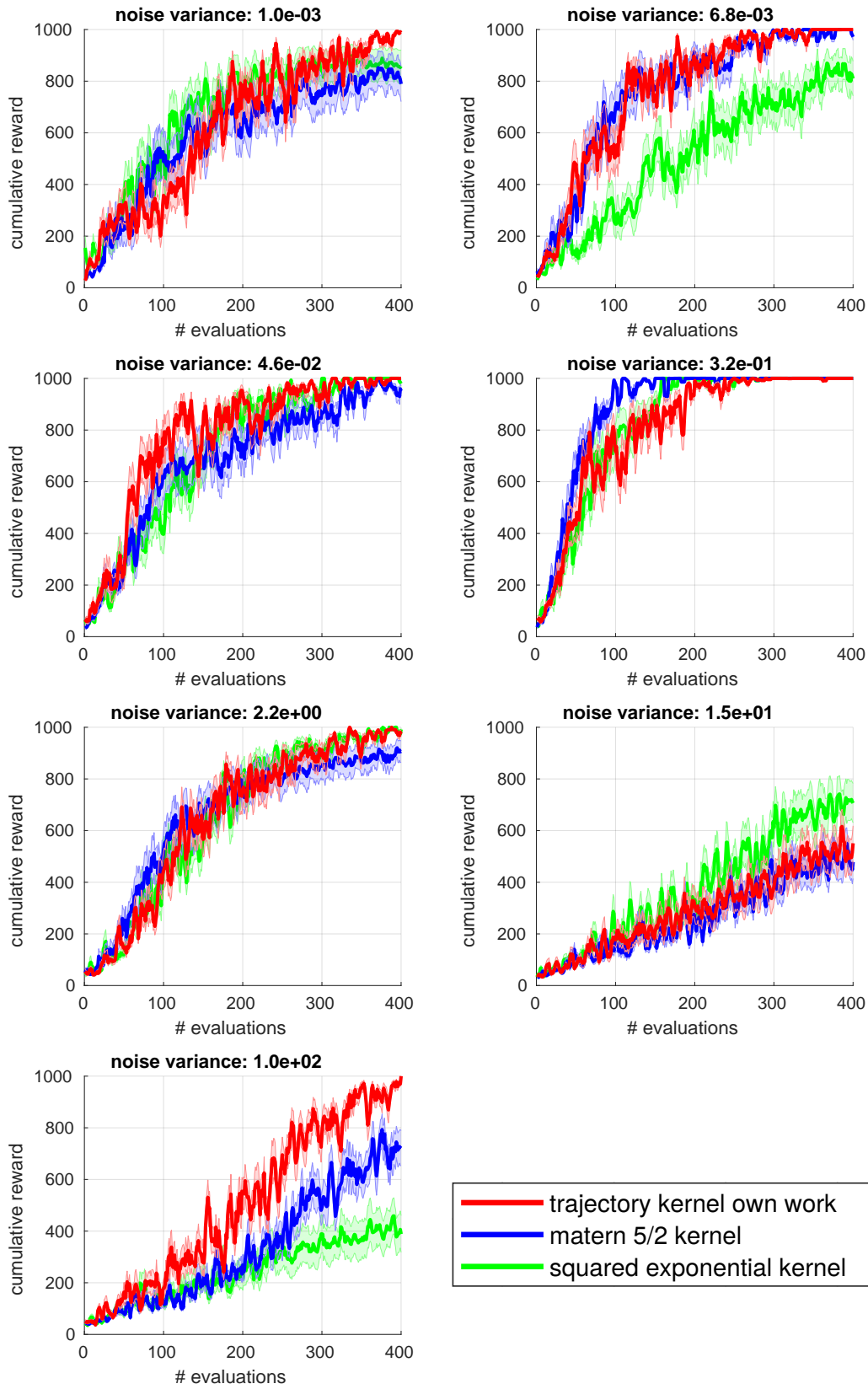


Figure 5.3.: Noise level comparison on the MATLAB Cart Pole implementation with 1000 time steps on the local Bayesian optimizer. The noise levels are logarithmically spaced and sorted in ascending order from 10^{-3} to 10^2 . Each kernel graph contains the average over 5 trials.

6 Discussion

When computing the continuous action space trajectory distance metric, proposed by [7], hyper parameter optimization is mandatory to compensate for high distance values. These values occur because of the Gaussian distribution density used as the probability measure for actions 4.2. Since we exponentiate the negative of the distance values, very high distance values will result in covariance values that are zero. The optimization of the scale length hyper parameter σ_l will prevent all-zero covariance matrices. Experiments have shown that tuning the signal deviation parameter σ_f is also helpful, because we simulate a policy only once for a result. Since a policy will not produce the exact same result after repeated evaluations, we have to assume a high variation on each result. This variation is regarded by σ_f .

Unfortunately we were not able to reproduce the outstanding results with the trajectory kernel from ???. Maybe because they do not provide any parameters or environment conditions. As shown exemplary in Figure 5.3 tuning of the noise level parameter can have a huge impact on the results. After a few test runs, we decided noise 1 2 3

The results show that trajectory kernels may have some advantage if computation time is not expensive. However, the standard kernels seem to require a similar number of black box function evaluations on the experiments shown.

7 Outlook

Implementing the OpenAI Gym gives the classic control problems like cart pole, mountain car, and acrobot. Furthermore a lot of more complex environments like a humanoid walker or some Atari games are provided [10]. This could facilitate future research if working with higher dimensional problems.

Dealing with more complex environments, can also lead to non-linear action mappings and more than one-dimensional actions. The trajectory kernels would have to be modified accordingly.

Sometimes the hyper parameter optimization fails to maximize of the log marginal likelihood. To gain more robustness the maximization with partial derivatives could be implemented as proposed by [6, 5].

Computing the Gaussian distribution density for the continuous action selection probability measure has lead to difficulties. Instead, one could implement a classification on the basis of a sigmoid function as it is used in [6]. The Gaussian cumulative distribution function could be suitable.

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A Some Appendix

Use letters instead of numbers for the chapters.