Gaussian Process in Practice: Scalability and Uncertainty

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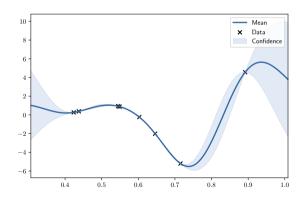
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Gaussian process

$$\mathbf{y} = (y_1, \dots, y_N), \quad \mathbf{X} = (\mathbf{x}_1, \dots, \mathbf{x}_N)^{\top}$$
$$p(\mathbf{y}|\mathbf{f}) = \mathcal{N}(\mathbf{y}|\mathbf{f}, \sigma^2 \mathbf{I}), \quad p(\mathbf{f}|\mathbf{X}) = \mathcal{N}(\mathbf{f}|0, \mathbf{K}(\mathbf{X}, \mathbf{X}))$$

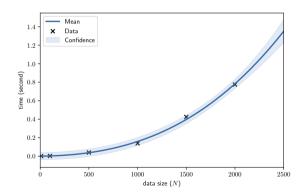


The computational time of GP

- The time complexity of Gaussian process is $O(N^3)$.
- Take 1D regression problem as an example:
 - ▶ The input and output dimensionality are both one.
 - We generate generate synthetic data.
 - ▶ Measure the time that it takes for computing the log-likelihood.

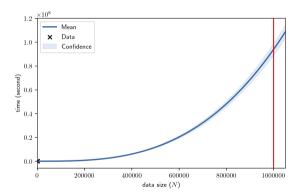
Empirical analysis of computational time

- I collect the run time for $N = \{10, 100, 500, 1000, 1500, 2000\}.$
- They take 1.3ms, 8.5ms, 28ms, 0.12s, 0.29s, 0.76s.



What if we have 1 million data points?

• The mean of predicted computational time is 9.4×10^7 seconds ≈ 2.98 years.



What about waiting for faster computers?

- Computational time $= \frac{\text{amount of work}}{\text{computer speed}}$
- If the computer speed increase at the pace of 20% year over year:
 - ▶ After 10 years, it will take about 176 days.
 - ▶ After 50 years, it will take about 2.9 hours.
- If we double the size of data, it takes 11.4 years to catch up.

What about parallel computing / GPU?

- Ongoing works about speeding up Cholesky decomposition with multi-core CPU or GPU.
- Main limitation: heavy communication and shared memory.

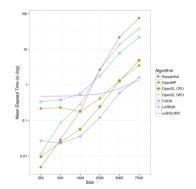


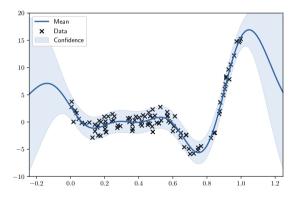
Figure 1: (RUSCHEL 2016)

Other approaches

- Apart from speeding up the exact computation, there have been a lot of works on approximation of GP inference.
- These methods often target at some specific scenario and provide good approximation for the targeted scenarios.
- Provide an overview about common approximations.

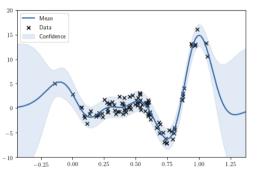
Big data (?)

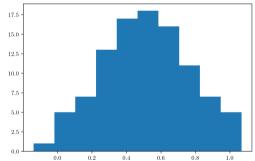
- lots of data \neq complex function
- In real world problems, we often collect a lot of data for modeling relatively simple relations.



Data subsampling?

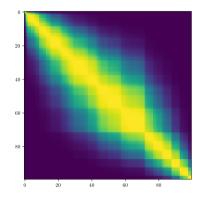
- Real data often do not evenly distributed.
- We tend to get a lot of data on common cases and very few data on rare cases.

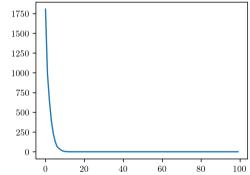




Covariance matrix of redundant data

- With redundant data, the covariance matrix becomes low rank.
- What about low rank approximation?





Low-rank approximation

• Let's recall the log-likelihood of GP:

$$\log p(\mathbf{y}|\mathbf{X}) = \log \mathcal{N}\left(\mathbf{y}|0, \mathbf{K} + \sigma^2 \mathbf{I}\right),$$

where K is the covariance matrix computed from X according to the kernel function $k(\cdot,\cdot)$ and σ^2 is the variance of the Gaussian noise distribution.

- \bullet Assume K to be low rank.
- This leads to Nyström approximation by Williams and Seeger (2001).

Nyström approximation (Williams and Seeger 2001)

- Let's randomly pick a subset from the training data: $\mathbf{Z} \in \mathbb{R}^{M \times Q}$.
- \bullet Approximate the covariance matrix K by $\tilde{K}\colon$

$$\tilde{\mathbf{K}} = \mathbf{K}_z \mathbf{K}_{zz}^{-1} \mathbf{K}_z^{\top},$$

where $\mathbf{K}_z = \mathbf{K}(\mathbf{X}, \mathbf{Z})$ and $\mathbf{K}_{zz} = \mathbf{K}(\mathbf{Z}, \mathbf{Z})$.

- Note that $\tilde{\mathbf{K}} \in \mathbb{R}^{N \times N}$, $\mathbf{K}_z \in \mathbb{R}^{N \times M}$ and $\mathbf{K}_{zz} \in \mathbb{R}^{M \times M}$.
- The log-likelihood is approximated by

$$\log p(\mathbf{y}|\mathbf{X}, \theta) \approx \log \mathcal{N}\left(\mathbf{y}|0, \mathbf{K}_z \mathbf{K}_{zz}^{-1} \mathbf{K}_z^{\top} + \sigma^2 \mathbf{I}\right).$$

Efficient computation using Woodbury formula

• The naive formulation does not bring any computational benefits.

$$\tilde{\mathcal{L}} = -\frac{1}{2}\log|2\pi(\tilde{\mathbf{K}} + \sigma^2\mathbf{I})| - \frac{1}{2}\mathbf{y}^{\top}(\tilde{\mathbf{K}} + \sigma^2\mathbf{I})^{-1}\mathbf{y}$$

Apply the Woodbury formula:

$$(\mathbf{K}_z \mathbf{K}_{zz}^{-1} \mathbf{K}_z^\top + \sigma^2 \mathbf{I})^{-1} = \sigma^{-2} \mathbf{I} - \sigma^{-4} \mathbf{K}_z (\mathbf{K}_{zz} + \sigma^{-2} \mathbf{K}_z^\top \mathbf{K}_z)^{-1} \mathbf{K}_z^\top$$

- Note that $(\mathbf{K}_{zz} + \sigma^{-2}\mathbf{K}_z^{\top}\mathbf{K}_z) \in \mathbb{R}^{M \times M}$.
- The computational complexity reduces to $O(NM^2)$.

Gaussian process with pseudo data (1)

- Snelson and Ghahramani (2006) propose the idea of having pseudo data. This
 approach is later referred to as Fully Independent Training Conditional (FITC).
- ullet Augment the training data (X, y) with pseudo data u at location Z.

$$p\left(\begin{bmatrix}\mathbf{y}\\\mathbf{u}\end{bmatrix} \mid \begin{bmatrix}\mathbf{X}\\\mathbf{Z}\end{bmatrix}\right) = \mathcal{N}\left(\begin{bmatrix}\mathbf{y}\\\mathbf{u}\end{bmatrix} \mid 0, \begin{bmatrix}\mathbf{K}_{ff} + \sigma^2 \mathbf{I} & \mathbf{K}_{fu}\\\mathbf{K}_{fu}^\top & \mathbf{K}_{uu}\end{bmatrix}\right)$$

where $K_{ff} = K(X, X)$, $K_{fu} = K(X, Z)$ and $K_{uu} = K(Z, Z)$.

Gaussian process with pseudo data (2)

• Thanks to the marginalization property of Gaussian distribution,

$$p(\mathbf{y}|\mathbf{X}) = \int_{\mathbf{u}} p(\mathbf{y}, \mathbf{u}|\mathbf{X}, \mathbf{Z}).$$

• Further re-arrange the notation:

$$p(\mathbf{y}, \mathbf{u} | \mathbf{X}, \mathbf{Z}) = p(\mathbf{y} | \mathbf{u}, \mathbf{X}, \mathbf{Z}) p(\mathbf{u} | \mathbf{Z})$$

where
$$p(\mathbf{u}|\mathbf{Z}) = \mathcal{N}\left(\mathbf{u}|0, \mathbf{K}_{uu}\right)$$
, $p(\mathbf{y}|\mathbf{u}, \mathbf{X}, \mathbf{Z}) = \mathcal{N}\left(\mathbf{y}|\mathbf{K}_{fu}\mathbf{K}_{uu}^{-1}\mathbf{u}, \mathbf{K}_{ff} - \mathbf{K}_{fu}\mathbf{K}_{uu}^{-1}\mathbf{K}_{fu}^{\top} + \sigma^2\mathbf{I}\right)$.

FITC approximation (1)

- So far, $p(\mathbf{y}|\mathbf{X})$ has not been changed, but there is no speed-up, $\mathbf{K}_{ff} \in \mathbb{R}^{N \times N}$ in $\mathbf{K}_{ff} \mathbf{K}_{fu} \mathbf{K}_{uu}^{-1} \mathbf{K}_{fu}^{\top} + \sigma^2 \mathbf{I}$.
- The FITC approximation assumes

$$\tilde{p}(\mathbf{y}|\mathbf{u}, \mathbf{X}, \mathbf{Z}) = \mathcal{N}\left(\mathbf{y}|\mathbf{K}_{fu}\mathbf{K}_{uu}^{-1}\mathbf{u}, \mathbf{\Lambda} + \sigma^2\mathbf{I}\right),$$

where
$$\mathbf{\Lambda} = (\mathbf{K}_{ff} - \mathbf{K}_{fu} \mathbf{K}_{uu}^{-1} \mathbf{K}_{fu}^{\top}) \circ \mathbf{I}$$
.

FITC approximation (2)

• Marginalize **u** from the model definition:

$$\tilde{p}(\mathbf{y}|\mathbf{X}, \mathbf{Z}) = \mathcal{N}\left(\mathbf{y}|0, \mathbf{K}_{fu}\mathbf{K}_{uu}^{-1}\mathbf{K}_{fu}^{\top} + \mathbf{\Lambda} + \sigma^{2}\mathbf{I}\right)$$

• Woodbury formula can be applied in the sam way as in Nyström approximation:

$$(\mathbf{K}_z\mathbf{K}_{zz}^{-1}\mathbf{K}_z^\top + \mathbf{\Lambda} + \sigma^2\mathbf{I})^{-1} = \mathbf{A} - \mathbf{A}\mathbf{K}_z(\mathbf{K}_{zz} + \mathbf{K}_z^\top\mathbf{A}\mathbf{K}_z)^{-1}\mathbf{K}_z^\top\mathbf{A},$$

where $\mathbf{A} = (\mathbf{\Lambda} + \sigma^2 \mathbf{I})^{-1}$.

FITC approximation (3)

- FITC allows the pseudo data not being a subset of training data.
- ullet The inducing inputs ${f Z}$ can be optimized via gradient optimization.
- Like Nyström approximation, when taking all the training data as inducing inputs, the FITC approximation is equivalent to the original GP:

$$\tilde{p}(\mathbf{y}|\mathbf{X}, \mathbf{Z} = \mathbf{X}) = \mathcal{N}(\mathbf{y}|0, \mathbf{K}_{ff} + \sigma^2 \mathbf{I})$$

- FITC can be combined easily with expectation propagation (EP).
- (Bui, Yan, and Turner 2017) provides an overview and a nice connection with variational sparse GP.

Model Approximation vs. Approximate Inference

When the exact model/inference is intractable, typically there are two types of approaches:

- Approximate the original model with a simpler one such that inference becomes tractable, like Nyström approximation, FITC.
- Keep the original model but derive an approximate inference method which is often *not* able to return the true answer, like variational inference.

Model Approximation vs. Approximate Inference

A problem with model approximation is that

- when an approximated model requires some tuning, e.g., for hyper-parameters, it is unclear how to improve it based on training data.
- In the case of FITC, we know the model is correct if $\mathbf{Z} = \mathbf{X}$, however, optimizing \mathbf{Z} will not necessarily lead to a better location.
- ullet In fact, optimizing ${f Z}$ can lead to overfitting. (Quiñonero-Candela and Rasmussen 2005)

Variational Sparse Gaussian Process (1)

- (Titsias 2009) introduces a variational approach for sparse GP.
- It follows the same concept of pseudo data:

$$p(\mathbf{y}|\mathbf{X}) = \int_{\mathbf{f}, \mathbf{u}} p(\mathbf{y}|\mathbf{f}) p(\mathbf{f}|\mathbf{u}, \mathbf{X}, \mathbf{Z}) p(\mathbf{u}|\mathbf{Z})$$

where
$$p(\mathbf{u}|\mathbf{Z}) = \mathcal{N}\left(\mathbf{u}|0, \mathbf{K}_{uu}\right)$$
,
$$p(\mathbf{y}|\mathbf{u}, \mathbf{X}, \mathbf{Z}) = \mathcal{N}\left(\mathbf{y}|\mathbf{K}_{fu}\mathbf{K}_{uu}^{-1}\mathbf{u}, \mathbf{K}_{ff} - \mathbf{K}_{fu}\mathbf{K}_{uu}^{-1}\mathbf{K}_{fu}^{\top} + \sigma^2\mathbf{I}\right)$$
.

Variational Sparse Gaussian Process (2)

- Instead of approximate the model, (Titsias 2009) derives a variational lower bound.
- Normally, a variational lower bound of a marginal likelihood, also known as evidence lower bound (ELBO), looks like

$$\begin{split} \log p(\mathbf{y}|\mathbf{X}) &= \log \int_{\mathbf{f}, \mathbf{u}} p(\mathbf{y}|\mathbf{f}) p(\mathbf{f}|\mathbf{u}, \mathbf{X}, \mathbf{Z}) p(\mathbf{u}|\mathbf{Z}) \\ &\geq \int_{\mathbf{f}, \mathbf{u}} q(\mathbf{f}, \mathbf{u}) \log \frac{p(\mathbf{y}|\mathbf{f}) p(\mathbf{f}|\mathbf{u}, \mathbf{X}, \mathbf{Z}) p(\mathbf{u}|\mathbf{Z})}{q(\mathbf{f}, \mathbf{u})}. \end{split}$$

Special Variational Posterior

• (Titsias 2009) defines an unusual variational posterior:

$$q(\mathbf{f}, \mathbf{u}) = p(\mathbf{f} | \mathbf{u}, \mathbf{X}, \mathbf{Z}) q(\mathbf{u}), \quad \text{where } q(\mathbf{u}) = \mathcal{N}\left(\mathbf{u} | \mu, \Sigma\right).$$

• Plug it into the lower bound:

$$\mathcal{L} = \int_{\mathbf{f}, \mathbf{u}} p(\mathbf{f} | \mathbf{u}, \mathbf{X}, \mathbf{Z}) q(\mathbf{u}) \log \frac{p(\mathbf{y} | \mathbf{f}) p(\mathbf{f} | \mathbf{u}, \mathbf{X}, \mathbf{Z}) p(\mathbf{u} | \mathbf{Z})}{p(\mathbf{f} | \mathbf{u}, \mathbf{X}, \mathbf{Z}) q(\mathbf{u})}$$

$$= \langle \log p(\mathbf{y} | \mathbf{f}) \rangle_{p(\mathbf{f} | \mathbf{u}, \mathbf{X}, \mathbf{Z}) q(\mathbf{u})} - \mathsf{KL} (q(\mathbf{u}) \parallel p(\mathbf{u} | \mathbf{Z}))$$

$$= \langle \log \mathcal{N} (\mathbf{y} | \mathbf{K}_{fu} \mathbf{K}_{uu}^{-1} \mathbf{u}, \sigma^{2} \mathbf{I}) \rangle_{q(\mathbf{u})} - \mathsf{KL} (q(\mathbf{u}) \parallel p(\mathbf{u} | \mathbf{Z}))$$

Tighten the Bound

• Find the optimal parameters of $q(\mathbf{u})$:

$$\mu^*, \Sigma^* = \arg \max_{\mu, \Sigma} \mathcal{L}(\mu, \Sigma).$$

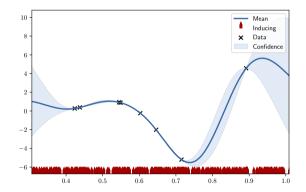
• Make the bound as tight as possible by plugging in μ^* and Σ^* :

$$\mathcal{L} = \log \mathcal{N} \left(\mathbf{y} | 0, \mathbf{K}_{fu} \mathbf{K}_{uu}^{-1} \mathbf{K}_{fu}^{\top} + \sigma^2 \mathbf{I} \right) - \frac{1}{2\sigma^2} \mathsf{tr} \left(\mathbf{K}_{ff} - \mathbf{K}_{fu} \mathbf{K}_{uu}^{-1} \mathbf{K}_{fu}^{\top} \right).$$

• The overall complexity of the lower bound remains $O(NM^2)$.

Variational sparse GP

- Note that \mathcal{L} is not a valid log-pdf, $\int_{\mathbf{v}} \exp(\mathcal{L}(\mathbf{y})) \leq 1$, due to the trace term.
- As inducing points are variational parameters, optimizing the inducing inputs **Z** always leads to a better bound.
- The model does not "overfit" with too many inducing points.



Are big covariance matrices always (almost) low-rank?

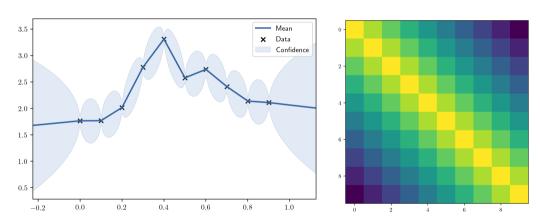
- Of course, not.
- A time series example

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

The data are collected with even time interval continuously.

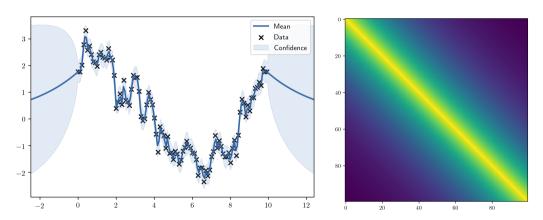
A time series example: 10 data points

• When we observe until t = 1.0:



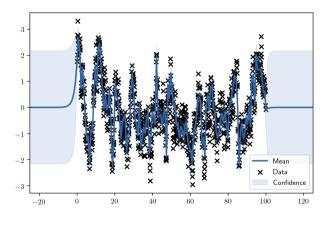
A time series example: 100 data points

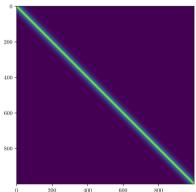
• When we observe until t = 10.0:



A time series example: 1000 data points

• When we observe until t = 100.0:





Banded precision matrix

- For the kernels like the Matern family, the precision matrix is banded.
- For example, given a Matern $\frac{1}{2}$ or known as exponential kernel: $k(x,x') = \sigma^2 \exp(-\frac{|x-x'|}{l^2})$.

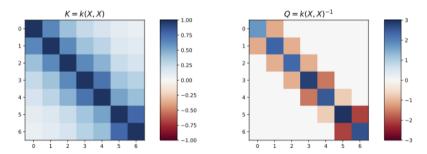


Figure 2: This slide is taken from Nicolas Durrande (2019)

Closed form precision matrix

- The precision matrix of Matern kernels can be computed in closed form.
- The lower triangular matrix from the Cholesky decomposition of the precision matrix is banded as well.

$$\log(\mathbf{y}|\mathbf{X}) = -\frac{1}{2}\log|2\pi(LL^{\top})^{-1}| - \frac{1}{2}\operatorname{tr}\left(\mathbf{y}\mathbf{y}^{\top}LL^{\top}\right)$$

where L is the lower triangular matrix from the Cholesky decomposition of the precision matrix Q, $Q=LL^{\top}$.

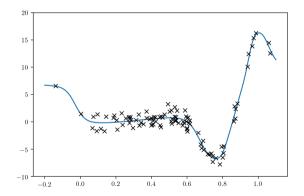
• The computational complexity becomes O(N).

Other approximations

- deterministic/stochastic frequency approximation
- distributed approximation
- conjugate gradient methods for covariance matrix inversion

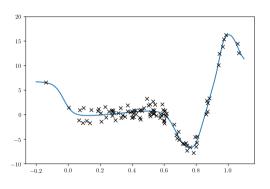
Why GP?

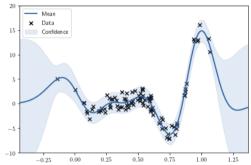
- Why do we spend all these energy on speeding up GP?
- Only for a non-parametric regressor?
- What about fitting a neural network?



What is the difference?

• The error bar!

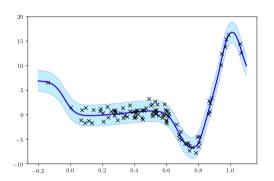




Can we learn an error bar with NN?

• Of course, we can. Let's add a likelihood to the neural network:

$$p(y|x) = \mathcal{N}\left(y|f_{\theta}(x), \sigma^2\right)$$



• Now, we have an error bar for our neural network. Are they the same?

Two types of uncertainty

• In our GP regression model, we have two "layers" of distributions:

$$p(\mathbf{y}|\mathbf{f}) = \mathcal{N}(\mathbf{y}|\mathbf{f}, \sigma^2 \mathbf{I}), \quad p(\mathbf{f}|\mathbf{X}) = \mathcal{N}(\mathbf{f}|0, \mathbf{K}(\mathbf{X}, \mathbf{X}))$$

- Aleatoric uncertainty: the uncertainty about the noise in individual data points
- Epistemic uncertainty: the uncertainty in the model

$$p(\theta|\mathcal{D}) = \frac{p(\mathcal{D}|\theta)p(\theta)}{p(\mathcal{D})}$$

Automated decision making

- Under the model assumption, the epistemic uncertainty allows us to know what the model *does not* know.
- This enables us to trade-off among different choices with limited information.
- Example: Bayesian optimization

$$x^* = \arg\min_{x} f(x)$$

Bayesian optimization

A surrogate model guided search

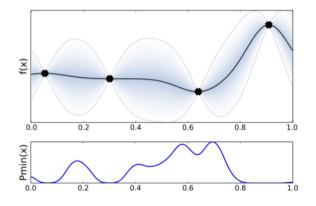


Figure 3: A 1D example

Possibility represented by uncertainty

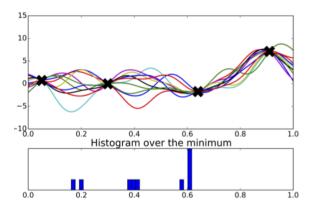


Figure 4: A 1D example

Balance exploitation and exploration

- **Exploitation**: Evaluate at the known best location will return the minimal value so far, but we learn nothing new.
- **Exploration**: Improve the understanding of the objective function, but may not be better than the current minimum.

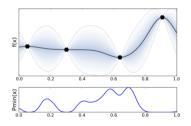


Figure 5: A 1D example

Acquisition function

- Formulate the policy of the exploitation and exploration tradeoff.
- The utility function about improvement:

$$u(f) = \max(0, f' - f)$$

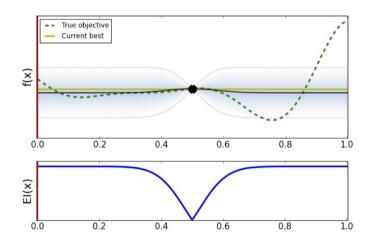
• The expected improvement under our surrogate model:

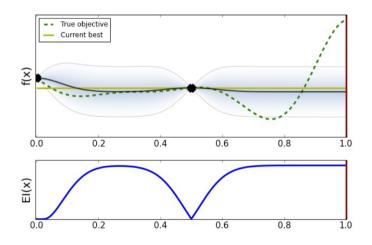
$$a_{\mathsf{EI}}(x) = \int u(f)p(f|x,\mathcal{D})\mathsf{d}f$$
$$= \int \max(0, f' - f)\mathcal{N}\left(f|m(x), c(x)\right)\mathsf{d}f$$

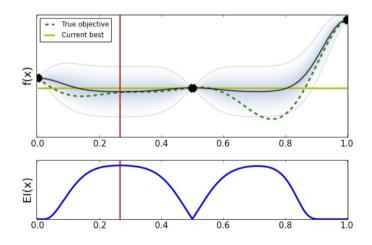
A BO algorithm

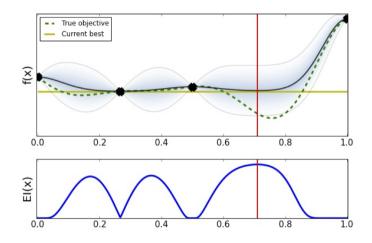
Loop of Bayesian optimization:

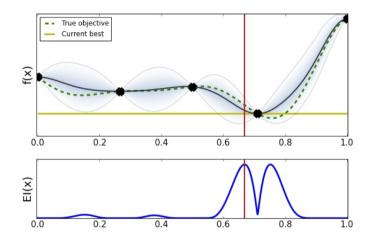
- Evaluate the objective function.
- Update the surrogate model.
- Select to the next location according to the acquisition function.

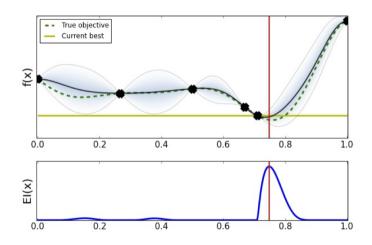


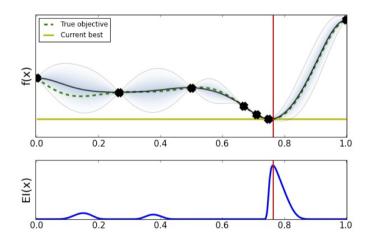


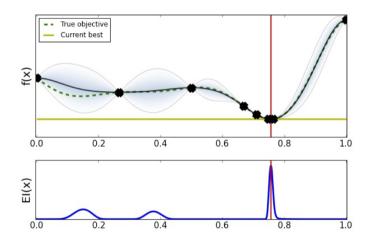












Challenges in Bayesian Optimization

- Optimizing the acquisition may be hard.
- With a high dimensional search problem, surrogate modeling becomes hard.
- Structured inputs can be hard to handle.
- Non-stationarity of an objective function
- Model mismatch
- Unknown safety constraints
- Warm-starting

• Thank you!

Lab session

• Please download the Jupyter notebook for the lab session from the following link:

http://gpss.cc/gpss18/labs/GPSS_Lab3_2018.ipynb

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