

Deterministic Sampling Methods



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Computational and Biological Learning Lab

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Outline

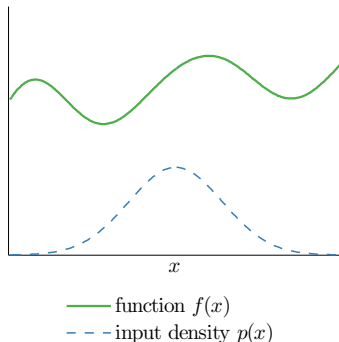
- Kernel Herding
- Bayesian Quadrature
- Unifying Results
- Demos

The Quadrature Problem

- We want to estimate an integral

$$Z = \int f(x)p(x)dx$$

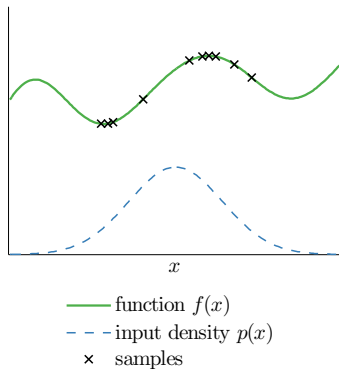
- Most computational problems in Bayesian inference correspond to integrals:
 - Expectations
 - Marginal distributions
 - Integrating out nuisance parameters
 - Normalization constants



Sampling Methods

- Monte Carlo methods:
Sample from $p(x)$, take empirical mean:

$$\hat{Z} = \frac{1}{N} \sum_{i=1}^N f(x_i)$$

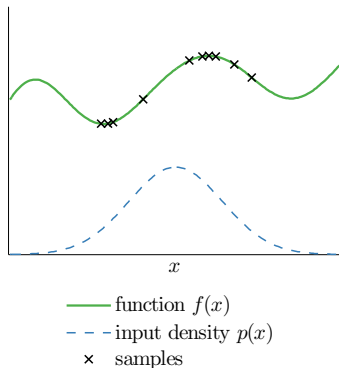


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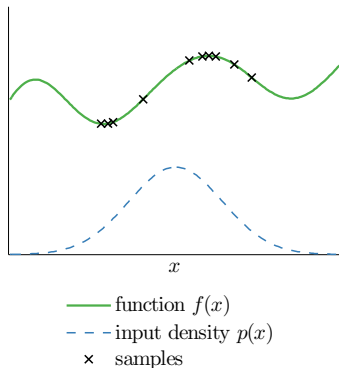


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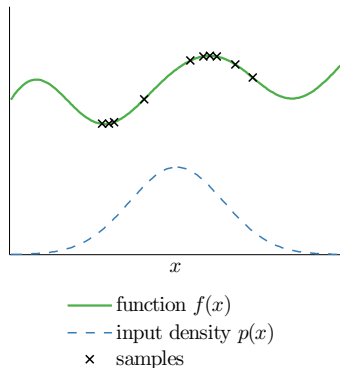


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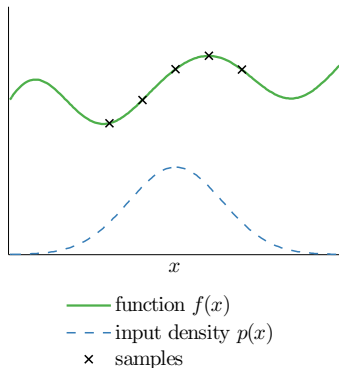


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 - Random bunching up
 - Often, nearby function values will be similar
- Quasi-Monte Carlo methods spread out samples to achieve faster convergence.



Kernel Herding [Welling et. al., 2009, Chen et. al., 2010]

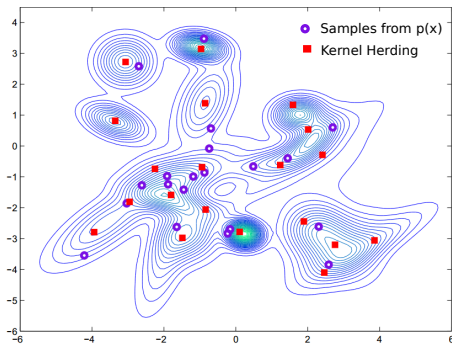
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- A sequential procedure for choosing sample locations, depending on previous locations.
- Keeps estimate rule $\hat{Z} = \frac{1}{N} \sum_{i=1}^N f(x_i)$
- Almost $\mathcal{O}(1/N)$ convergence instead of $\mathcal{O}(1/\sqrt{N})$ typical of random sampling, by spreading out samples.



Kernel Herding Objective

KH was found to minimize Maximum Mean Discrepancy:

$$\text{MMD}_{\mathcal{H}}(p, q) = \sup_{\substack{f \in \mathcal{H} \\ \|f\|_{\mathcal{H}}=1}} \left| \int f(x)p(x)dx - \int f(x)q(x)dx \right|$$

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In KH, $p(x)$ is true distribution, and $q(x)$ is a set of point masses at sample locations $\{x_1, \dots, x_N\}$:

$$\epsilon_{KH}(\{x_1, \dots, x_N\}) = \text{MMD}_{\mathcal{H}} \left(p, \underbrace{\frac{1}{N} \sum_{n=1}^N \delta_{x_n}}_{q(x)} \right)$$

Kernel Herding

- Assuming function is in a Reproducing Kernel Hilbert Space defined by $k(\cdot, \cdot)$, MMD has closed form.

Kernel Herding

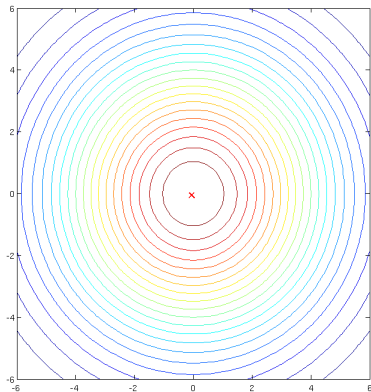
- Assuming function is in a Reproducing Kernel Hilbert Space defined by $k(\cdot, \cdot)$, MMD has closed form.
- When sequentially minimizing MMD, new point is added at:

$$x_{N+1} = \operatorname{argmax}_{x \in \mathcal{X}} \left[2 \int k(x, x') p(x') dx' - \frac{1}{N+1} \sum_{m=1}^N k(x, x_m) \right]$$

Kernel Herding in Action

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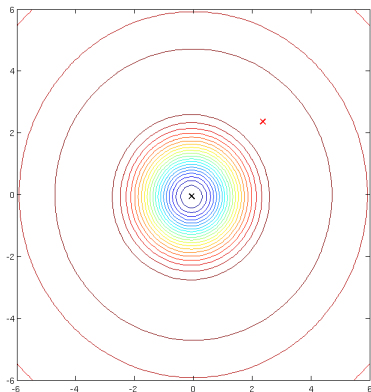
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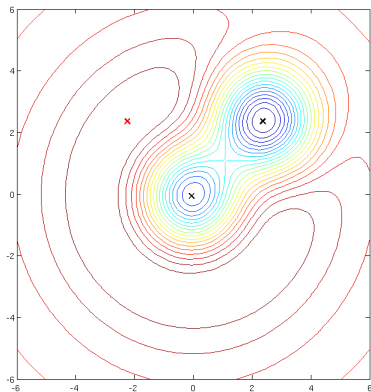


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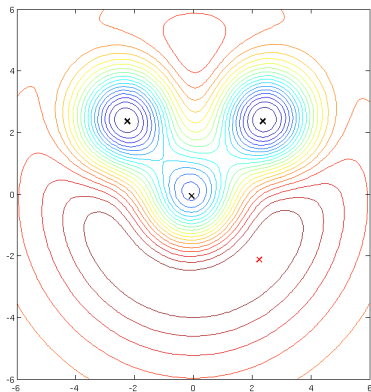
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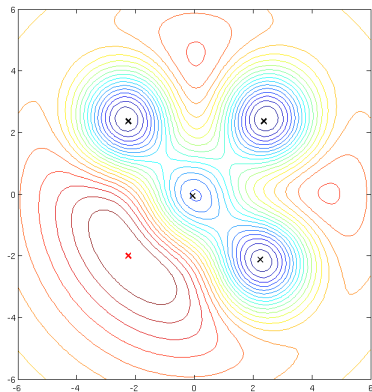
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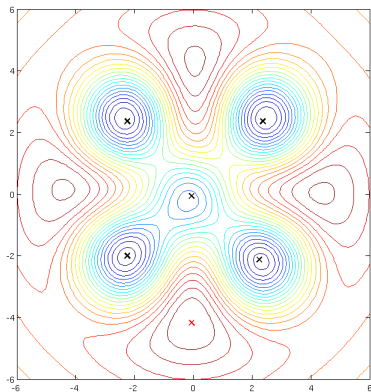
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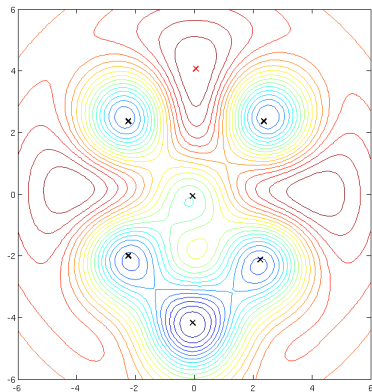
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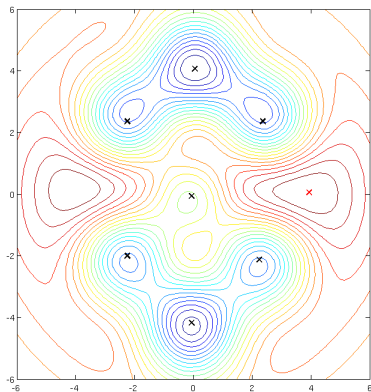
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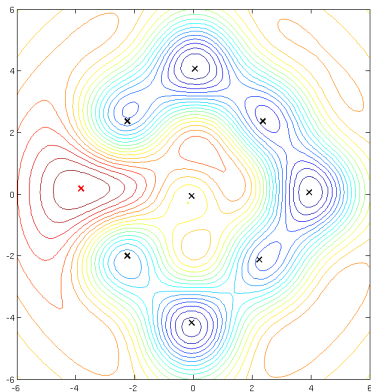
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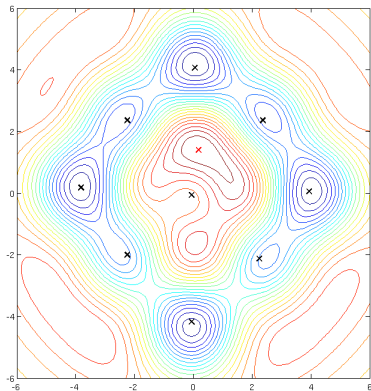
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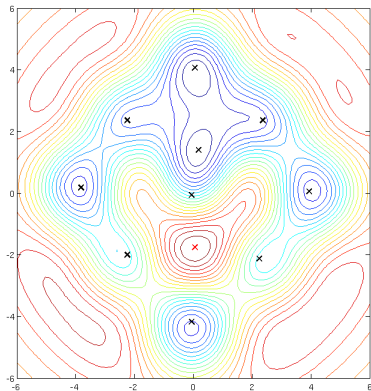
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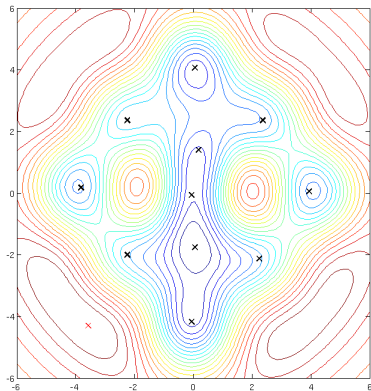


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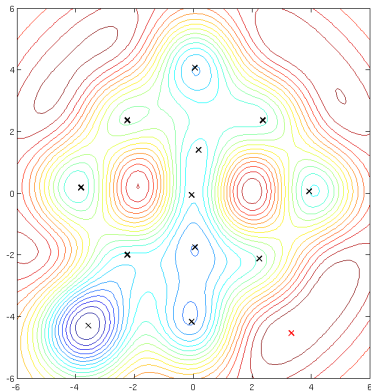
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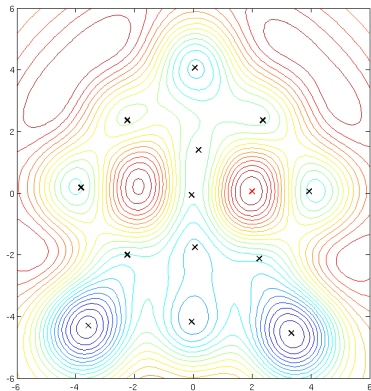
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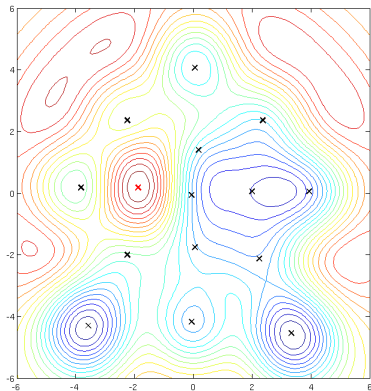
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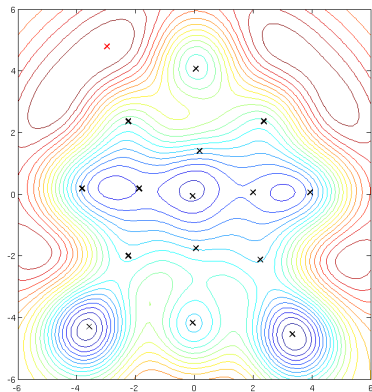
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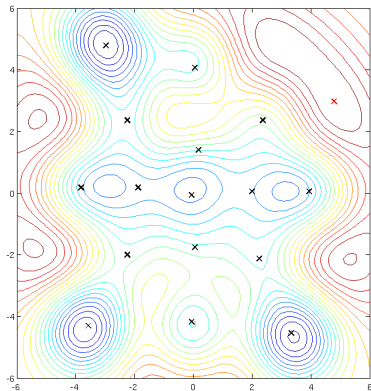
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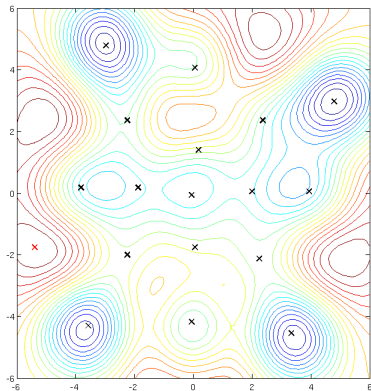


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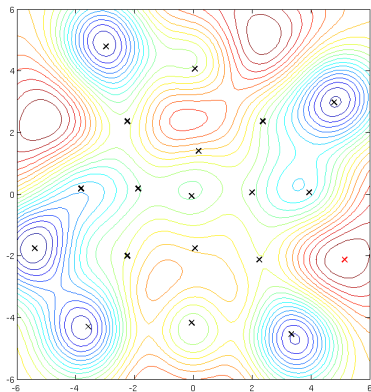


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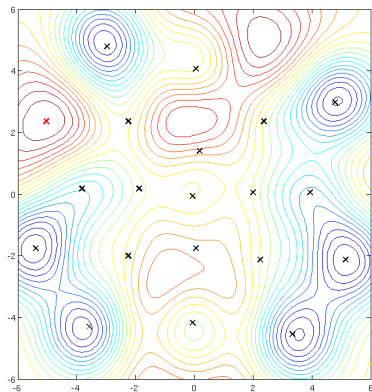


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Kernel Herding Summary

- A sequential sampling method which minimizes a worst-case divergence, given that $f(x)$ belongs to a given RKHS.
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Can we reason about the optimal weighting strategy?

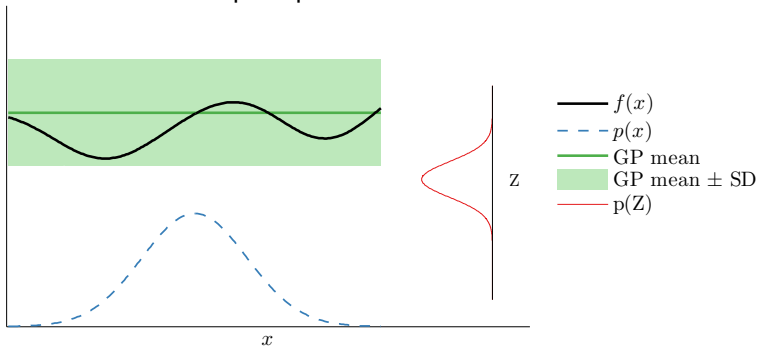
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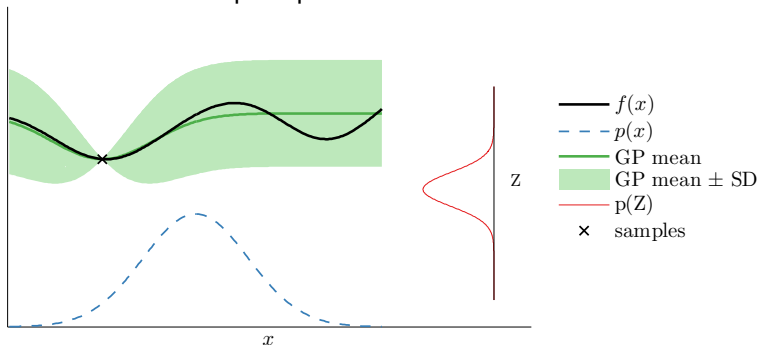
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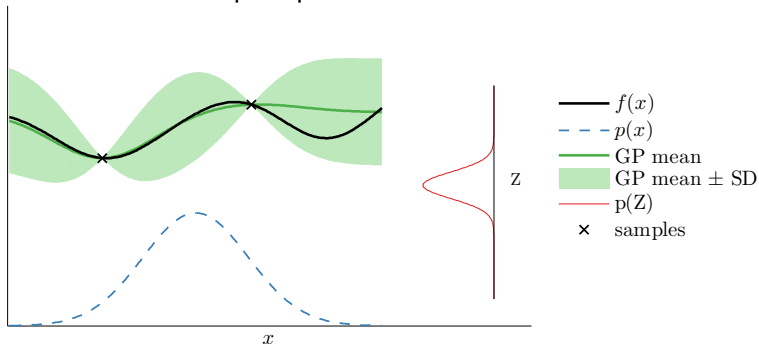
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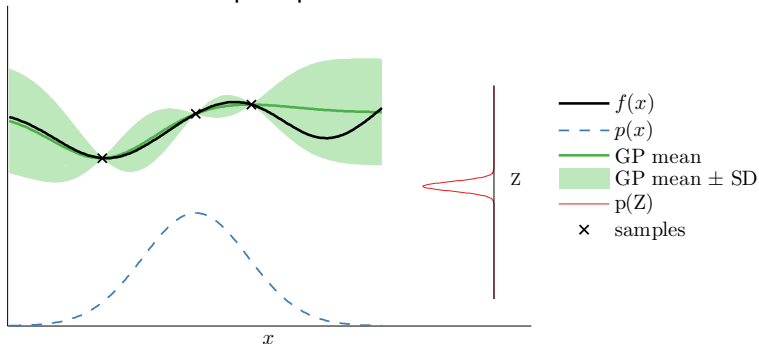
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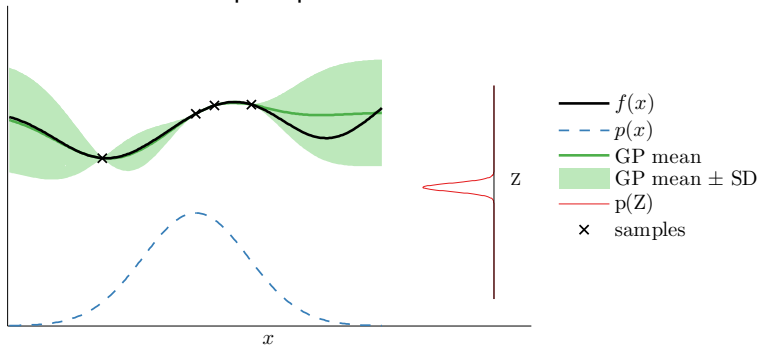
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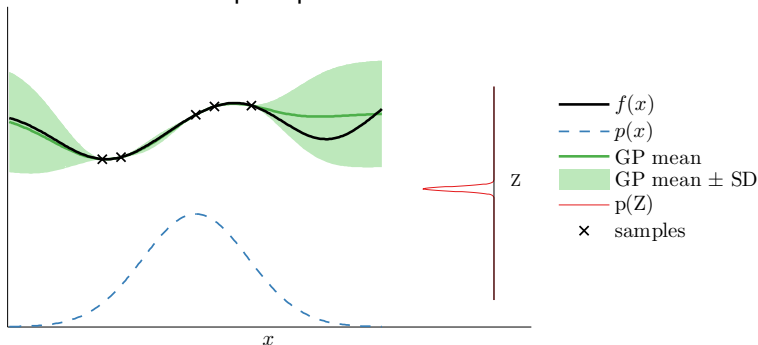
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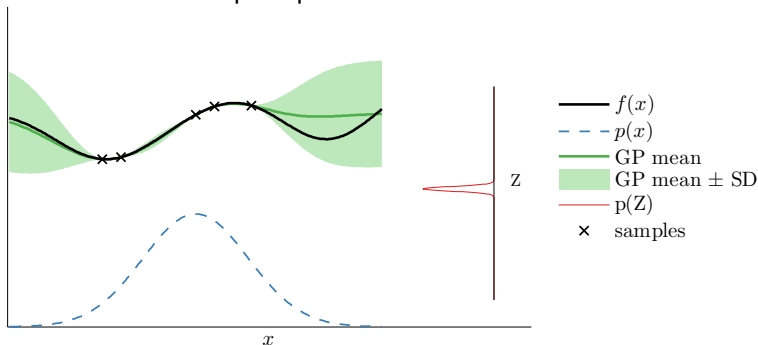
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- Can choose samples however we want.

Bayesian Quadrature Estimator

Posterior over Z has mean linear in $f(x_s)$:

$$\mathbb{E}_{\text{GP}} [Z | f(x_s)] = \sum_{i=1}^N w_{BQ}^{(i)} f(x_i)$$

where

$$w_{BQ} = z^T K^{-1} \quad \text{and} \quad z_n = \int k(x, x_n) p(x) dx$$

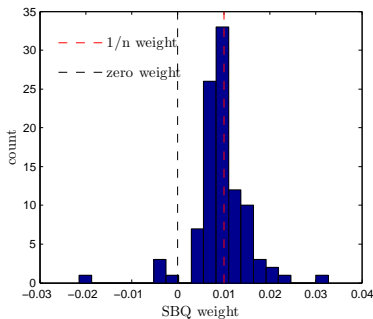
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- Does not depend on function values
- Can choose samples sequentially: Sequential Bayesian Quadrature.

Relating Objectives

KH and BQ have completely different motivations:

- KH minimizes a worst-case bound
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Is there any correspondence?

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First Main Result

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BQ is minimizing KH objective

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Second Main Result

BQ estimator is the optimal weighting strategy:

$$\mathbb{V}[Z|f(x_s)] = \inf_{w \in \mathbb{R}^N} \sup_{\substack{f \in \mathcal{H} \\ \|f\|_{\mathcal{H}}=1}} \left| \int f(x)p(x)dx - \sum_{n=1}^N w_n f(x_n) \right|^2$$

Performance

- KH and BQ are minimizing the same objective, but BQ has freedom to choose weights.
- How does this affect performance?

Second Main Result

BQ estimator is the optimal weighting strategy:

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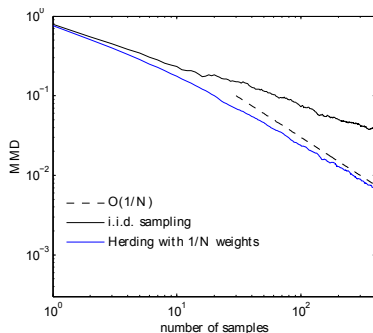
$\mathbb{V}[Z|f(x_s)]$ has two interpretations:

- Bayesian: posterior variance of Z under a GP prior.
- Frequentist: tight bound on estimation error of Z .

Rates of Convergence

What is rate of convergence of BQ?

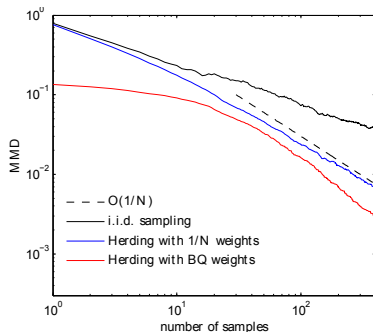
Expected Variance / MMD



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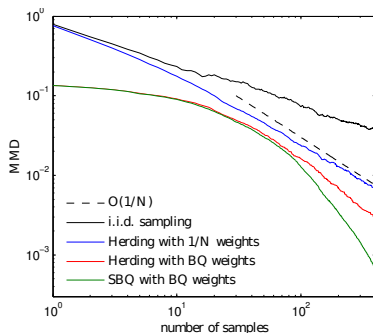
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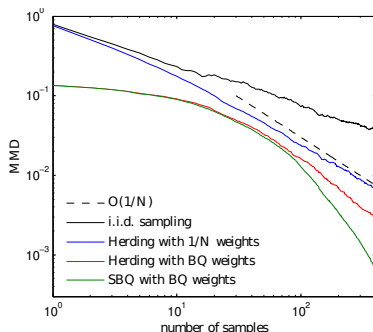
Expected Variance / MMD



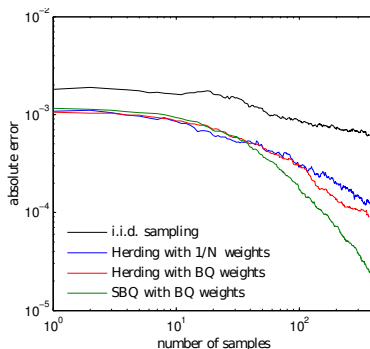
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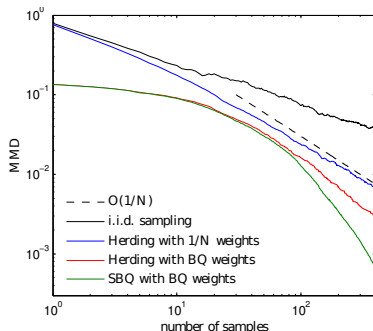
Empirical Rates in RKHS



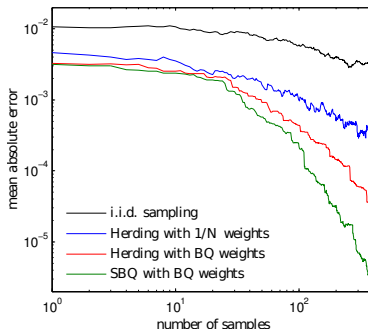
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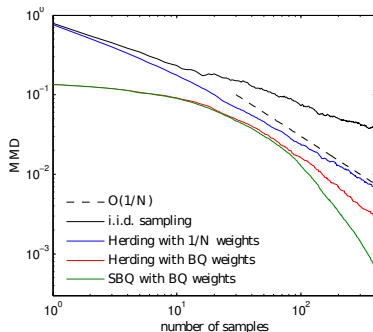
Empirical Rates out of RKHS



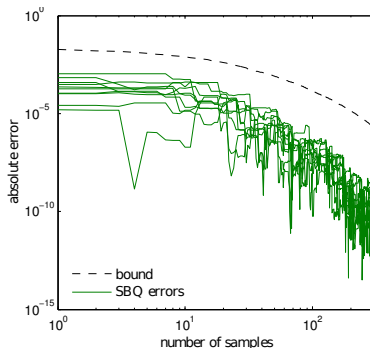
Rates of Convergence

What is rate of convergence of BQ?

Expected Variance / MMD



Bound on Bayesian Error



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- Joint work with Ferenc Huzsar