

Multilevel and Multi-Index Monte Carlo: Theory, Practice, and `mimclib`

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Outline

1 Theory: Monte Carlo methods

- Problem
- Monte Carlo
- Multilevel Monte Carlo (MLMC)
- Multi-Index Monte Carlo (MIMC)

2 Implementation: `mimclib`

- Library Overview
- Installation
- Example output from MIMC example

The central question

Given an $\mathbb{R}^n$ -valued random variable, X , a function, $g : \mathbb{R}^n \rightarrow \mathbb{R}$, assume that we are interested in computing the quantity

$$\mathbb{E}[g(X)] = \int_{\mathbb{R}^n} g(x) \, dF(x).$$

Challenges:

- Only approximate samples of X can be generated, e.g., X is the solution of an SDE.
- g cannot be computed exactly or cheaply, e.g., involves FEM approximation, or a nested expectation.

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Setup

Our objective is to build an estimator $\mathcal{A} \approx \mathbb{E}[g(X)]$ with **minimal work** where

$$P(|\mathcal{A} - \mathbb{E}[g(X)]| \leq \varepsilon) \geq \eta$$

for a given accuracy ε and a given confidence level determined by $0 < \eta < 1$.

Instead, we impose the following, more restrictive, two constraints:

Bias constraint: $|\mathbb{E}[\mathcal{A} - g(X)]| \leq (1 - \theta)\varepsilon,$

Statistical constraint: $\text{Var}[\mathcal{A}] \leq \theta^2 \varepsilon^2$

For some tolerance splitting, $\theta \in (0, 1)$.

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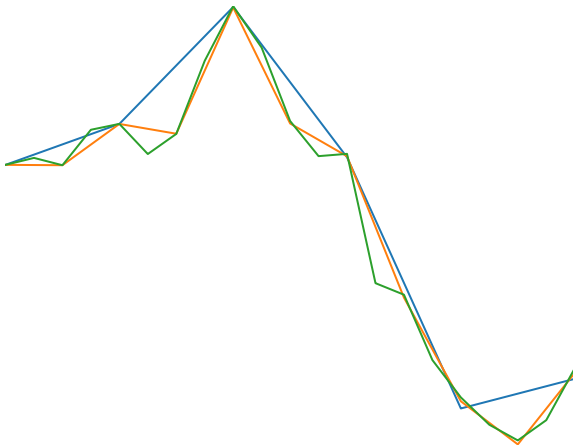
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Numerical approximation

Notation: g_ℓ for $\ell \in \mathbb{N}$ is the approximation of g calculated based on some discretization. As ℓ increases, both the accuracy and cost per sample increase.



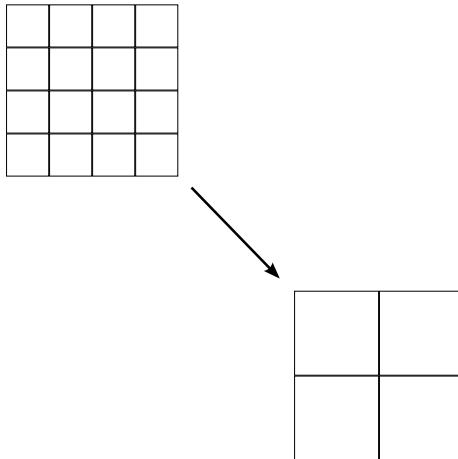
Monte Carlo

The simplest (and most popular) estimator is the Monte Carlo estimator

$$\mathcal{A}_{\text{MC}}[g_L; M] = \frac{1}{M} \sum_{m=1}^M g_L(\mathbf{x}^{(m)}),$$

for a given level, L , and number of samples, M , that we can choose to satisfy the error constraints and minimize the work.

MLMC main idea: Variance reduction



Multilevel Monte Carlo (MLMC)

(Heinrich, 1998) and (Giles, 2008)

Observe the telescopic identity

$$\mathbb{E}[g] \approx \mathbb{E}[g_L] = \mathbb{E}[g_0] + \sum_{\ell=1}^L \mathbb{E}[g_\ell - g_{\ell-1}] = \sum_{\ell=0}^L \mathbb{E}[\Delta_\ell g],$$

where

$$\Delta_\ell g = \begin{cases} g_0 & \text{if } \ell = 0, \\ g_\ell - g_{\ell-1} & \text{if } \ell > 0. \end{cases}$$

Then, using Monte Carlo to approximate each difference independently, the MLMC estimator can be written as

$$\mathcal{A}_{\text{MLMC}}[g; L] = \sum_{\ell=0}^L \mathcal{A}_{\text{MC}}[\Delta_\ell g; M_\ell].$$

Main idea: variance reduction using cheaper approximations.

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Assumptions

Assumption MC1 (Bias):

Assumption MC2 (Work):

Assumption MLMC3 (Variance):

$$\begin{aligned} E_\ell &:= |\mathbb{E}[g - g_\ell]| \lesssim 2^{-\alpha\ell}, \\ \text{Work}[\Delta_\ell g] &\lesssim 2^{d\gamma\ell}, \\ V_\ell &:= \text{Var}[\Delta_\ell g] \lesssim 2^{-\beta\ell} \end{aligned}$$

for all $\ell \in \mathbb{N}$ and positive constants α, β, γ .

Recall: Optimal cost of Monte Carlo is $\mathcal{O}(\varepsilon^{-2 - \frac{d\gamma}{\alpha}})$

The optimal cost of MLMC is

$$\begin{cases} \mathcal{O}(\varepsilon^{-2}) & \beta > d\gamma \\ \mathcal{O}(\varepsilon^{-2}) (\log(\varepsilon^{-1}))^2 & \beta = d\gamma \\ \mathcal{O}(\varepsilon^{-2 - \frac{d\gamma - \beta}{\alpha}}) & \beta < d\gamma \end{cases}$$

Notice: the cost increases with increasing work (exponentially in d).

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In practice: Continuation MLMC

- To compute M_ℓ , we need to find estimates of $V_\ell := \text{Var}[\Delta_\ell \text{func}]$.
- To determine L , we need reliable estimate of E_ℓ .
- Instead of running for the small required ε , CMLMC runs a sequence of MLMC realisations, for decreasing tolerances, ending with the required ε .
- In each step, estimates of V_ℓ are generated using a Bayesian setting which uses **Assumption MLMC3** coupled with the generated samples to produce good estimates even with a small number of samples.
- This works as long as the Kurtosis does not increase too much with ℓ .
- The value of L is also chosen in each step to minimize the work. This allows choosing a better splitting parameter, θ .
- CMLMC does not have to reuse samples between iterations, ensuring an unbiased estimator for level L approximation.
- All implemented in `mimc`lib, see later.

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A simple problem

- Consider this problem

$$\mathbb{E}[g(\mathbb{E}[f(X_T) | X_H])]$$

for an SDE solution,

$$dX_t = \mu_t dt + \sigma_t dW_t$$

- How do we approximate it?
 - Requires approximation of outer process X_H
 - Requires approximation of inner expectation $\mathbb{E}[X_T | X_H]$
 - Requires approximation of inner process X_T , given X_H .
- Using $2^{-\ell}$ for the step-size of the process and 2^ℓ inner samples the cost of approximating a single sample is $\mathcal{O}(2^{2\ell})$.
- The variance of $\Delta_\ell g$, given these approximations, converges slower than $\mathcal{O}(2^{-2\ell})$, so complexity is not optimal.
- Similar problem if particle system is used.

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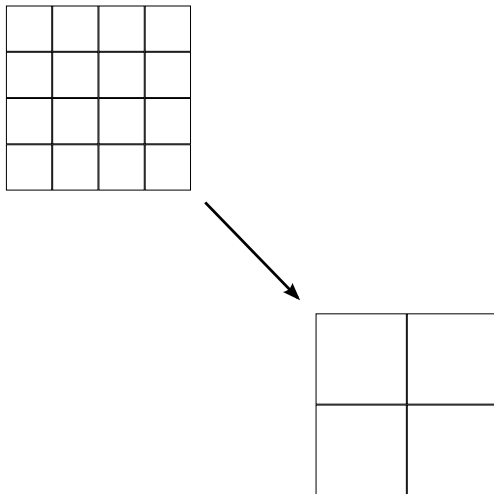
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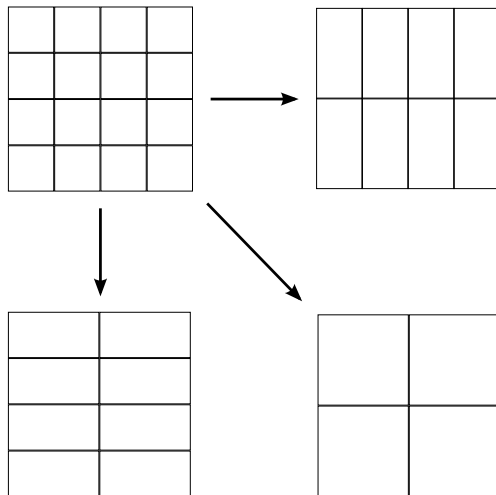
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Variance reduction: MLMC



Variance reduction: Further potential



MIMC Estimator (Haji-Ali et al., 2017)

Consider discretization parameter, $\ell = (\ell_1, \ell_2, \dots, \ell_d) \in \mathbb{N}^d$, and use the same notation g_ℓ the approximation of g calculated using a discretization defined by ℓ , e.g., using 2^{ℓ_1} time-steps and 2^{ℓ_2} inner samples.

For $i = 1, \dots, d$, define the first order difference operators

$$\Delta_i g_\ell = \begin{cases} g_\ell & \text{if } \ell_i = 0, \\ g_\ell - g_{\ell - \mathbf{e}_i} & \text{if } \ell_i > 0, \end{cases}$$

Construct the first order mixed difference

$$\begin{aligned} \Delta_\ell g &= \left(\bigotimes_{i=1}^d \Delta_i \right) g_\ell \\ &= g_{\ell_1, \ell_2} - g_{\ell_1-1, \ell_2} - g_{\ell_1, \ell_2-1} + g_{\ell_1-1, \ell_2-1} \end{aligned}$$

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MIMC Estimator

Then, assuming that

$$\mathbb{E}[g_\ell] \rightarrow \mathbb{E}[g] \quad \text{as} \quad \ell_i \rightarrow \infty \quad \text{for all } i = 1, \dots, d,$$

it is not difficult to see that

$$\mathbb{E}[g] = \sum_{\ell \in \mathbb{N}^d} \mathbb{E}[\Delta_\ell g] \approx \sum_{\ell \in \mathcal{I}} \mathbb{E}[\Delta_\ell g]$$

where $\mathcal{I} \subset \mathbb{N}^d$ is a *properly chosen* index set.

As in MLMC, approximating each term by independent MC samplers, the MIMC estimator can be written as

$$\mathcal{A}_{\text{MIMC}}[g; \mathcal{I}] = \sum_{\ell \in \mathcal{I}} \frac{1}{M_\ell} \sum_{m=1}^{M_\ell} \Delta_\ell g(\omega_{\ell,m})$$

with *properly chosen* sample sizes $(M_\ell)_{\ell \in \mathcal{I}}$.

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Assumptions for MIMC

For every ℓ , we assume the following

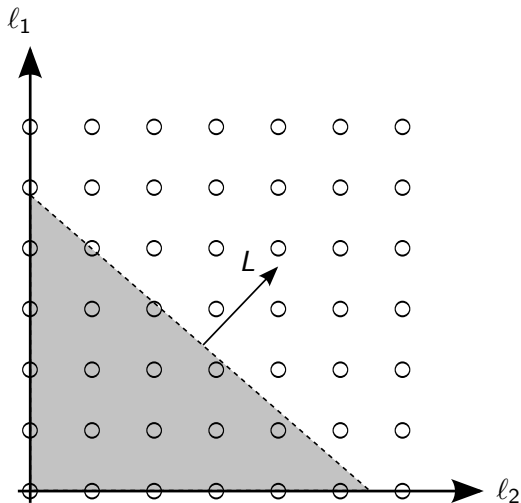
Assumption MIMC1 (Bias) : $E_\ell = |\mathbb{E}[\Delta_\ell g]| \propto \prod_{i=1}^d 2^{-\alpha_i \ell_i}$

Assumption MIMC2 (Work) : $W_\ell = \text{Work}(\Delta_\ell g) \propto \prod_{i=1}^d 2^{\gamma_i \ell_i},$

Assumption MIMC3 (Variance) : $V_\ell = \text{Var}[\Delta_\ell g] \propto \prod_{i=1}^d 2^{-\beta_i \ell_i},$

- The MIMC assumptions are stronger than the corresponding assumptions in MLMC.
They imply existence of **mixed derivatives** of the underlying map (and possibly the solution of the adjoint problem associated to the functional g), as opposed to first-order derivatives for MLMC.

Optimal index set, $\mathcal{I}$



Or find it adaptively!

Fully Isotropic case

Assume $\alpha_i = \alpha$, $\beta_i = \beta < 2\alpha$ and $\gamma_i = \gamma$ for all $i \in \mathcal{I}$. Then the optimal work is

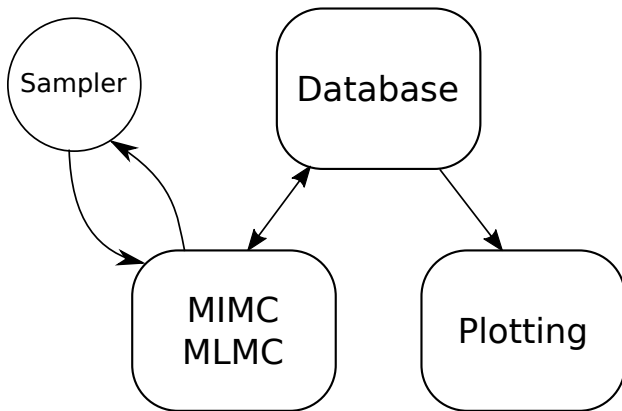
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$$\text{Work}(\text{MLMC}) = \begin{cases} \mathcal{O}(\varepsilon^{-2}), & \beta > d\gamma, \\ \mathcal{O}(\varepsilon^{-2} (\log(\varepsilon^{-1}))^2), & \beta = d\gamma, \\ \mathcal{O}(\varepsilon^{-(2 + \frac{(d\gamma - \beta)}{\alpha})}), & \beta < d\gamma. \end{cases}$$

$$\text{Work}(\text{MIMC}) = \begin{cases} \mathcal{O}(\varepsilon^{-2}), & \beta > \gamma, \\ \mathcal{O}(\varepsilon^{-2} (\log(\varepsilon^{-1}))^{2d}), & \beta = \gamma, \\ \mathcal{O}(\varepsilon^{-(2 + \frac{\gamma - \beta}{\alpha})} \log(\varepsilon^{-1})^{(d-1)\frac{\gamma - \beta}{\alpha}}), & \beta < \gamma. \end{cases}$$

mimclib

- These methods and others are common in UQ applications.
- UQ research and applications require **systematic** studies of random runs.
- mimclib is a library that was developed with that objective in mind.
 - ▶ Provide an **easy to use**, **customizable** and **extendable** open source library for UQ problems, both forward and inverse.
 - ▶ Multilevel and Multi-Index versions of Monte Carlo, Stochastic collocation, Least square projection.
 - ▶ Parallel computation-friendly.
 - ▶ Provide easy to use storage facility (SQLite and MySQL, allowing asynchronous, remote access).
 - ▶ Provide easy to customize plotting facility (for common plots) – using matplotlib.
 - ▶ Provide some simple (and not so simple) test cases.
 - ▶ Use Python for easier implementation of most parts of code and use object code (C++) for computationally expensive parts.
- mimclib can also be used for non-academic purpose too.



Installation

- Prerequisites: gcc (supporting c++11), Python 3.4, pip.
- ```
> git clone \
 https://github.com/haji-ali/mimclib.git
> cd mimclib
> make pip
```

Or just google: “mimclib github”
- Done!.
- Uses sqlite3 by default. A bit more complicated to install MySQL but very convenient if you are planning to run thousands of simulations on a cluster.

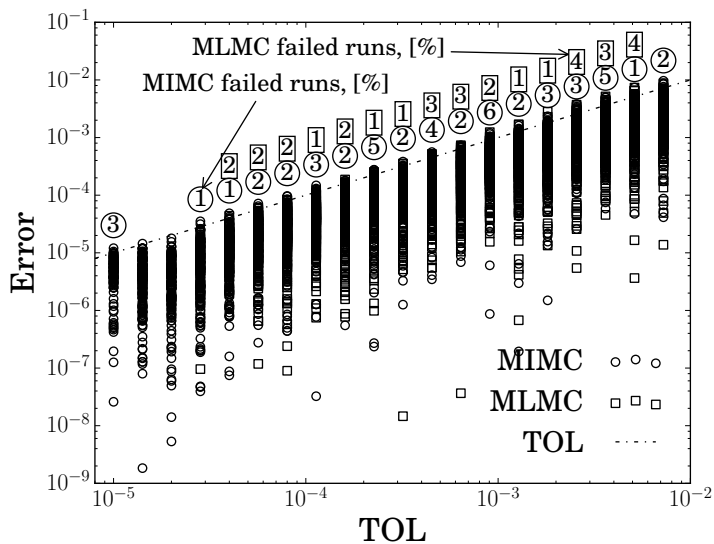


# Minimal Sampler

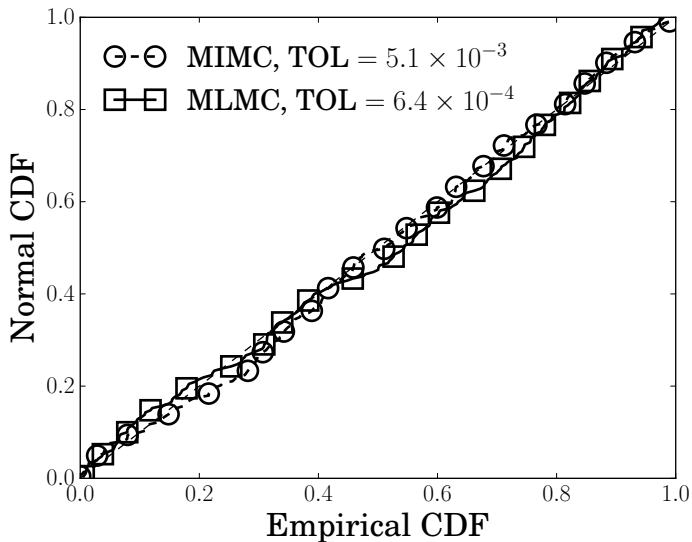
```
def mySampleQoI(run, inds, M):
 # run: is a reference to current run
 # inds: is a dxN array that specifies the
 # discretization-level along each dimension
 # M: number of samples

 return solves, time
solves: NxM array that contains the correlated
samples for each discretization
time: time taken to generate these samples.
```

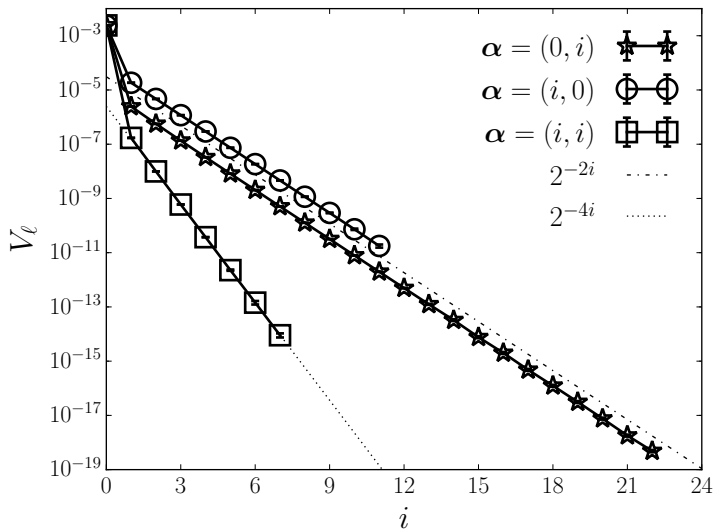
# Errors, customised



# PP-plot



# Variance convergence



# Running time

