

# Error analysis for parallel Monte Carlo estimation from many short Markov chains

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# Motivation

# Goal in MCMC

The goal is to estimate integrals with respect to a probability measure via simulation with modern computation. This is crucial for modern Bayesian inference.

For some real-valued function  $\varphi : \mathbf{X} \rightarrow \mathbb{R}$ , estimate

$$\int_{\mathbf{X}} \varphi d\Pi$$

## MCMC in the single-chain regime

Simulate realizations of a *single* Markov chain  $X_0, \dots, X_t, \dots$  with unique invariant distribution  $\Pi$  until convergence (approximate).

The marginal realizations stabilize to realizations from  $\Pi$ . Use the empirical average to estimate an expectation:

$$\frac{1}{T} \sum_{s=1}^T \varphi(X_{t+s}) \approx \int_{\mathbf{x}} \varphi d\Pi.$$

## Single-chain regime theoretical properties

- Theoretically solid foundation
- The dependence in the Markov chain restricts many advantages of modern parallel computation
- Knowing when the Markov chain converged is an extremely challenging problem
- Obtaining explicit and useful convergence rates is challenging

**A generally long serial Markov chain path simulation**

# The many-short-chains regime

Simulate realizations from *multiple* independent *short* Markov chain paths  $X_0^m, \dots, X_t^m$  for  $m = 1, \dots, M$  and ensemble them.

Ensemble examples:

- Take the average of  $X_t^1, \dots, X_t^m$
- Take the average of averages

## Many-short-chains regime properties

- Short Markov chain length = fast simulation
- Parallel simulations able to utilize modern parallel compute

### A possibly fast, parallel simulation

- The bias can be large or even unknown due to the short simulation length = estimate incorrectly
- Many-short-chains has been used [Gelman and Rubin, 1992] and debated, and criticized due to its weaknesses [Geyer, 1992].

## Contribution of this work

Construct an ensemble MCMC estimator with an **error guarantee** in the **many-short-chains** regime.

i.e. **short** Markov chain simulations in **parallel**.



# Many-short-chains (MSC) estimator

## MSC estimator requirements

- Let  $Y_1, \dots, Y_N$  independently from an importance sampling proposal
- Construct an initial distribution for the Markov chains using the self-normalized importance weights
- Use independent Markov chains  $X_0^m, X_1^m, \dots, X_t^m, \dots$  for  $m = 1, \dots, M$  initialized with  $X_0^m$  from this initial distribution.
- An explicit set  $C$  (we will use a drift condition for this later).

**Note:** This initialization does not require normalizing constant of  $\Pi$ .

## MSC estimator construction

$\tau_C$  is the first return time to the set  $C$  and

$$S_{\tau_C}(\varphi) = \begin{cases} \sum_{k=1}^{\tau_C} \varphi(X_k) & X_0 \in C \\ 0 & X_0 \notin C. \end{cases}$$

The *MSC estimator* is the average over these independent sums of Markov chains, that is,

$$\overline{S_{\tau_C}(\varphi)} = \frac{1}{M} \sum_{m=1}^M S_{\tau_C}^m(\varphi).$$

# Mean squared error analysis

## Geometric drift condition

There is a function  $V \geq 1$  and constants  $\gamma \in (0, 1)$  and  $K > 0$  such that

$$\mathbb{E} [V(X_t) \mid X_{t-1}] \leq \gamma V(X_{t-1}) + K.$$

A geometric drift condition ensures the Markov chain visits

$$C = \{x \in \mathbf{X} : V(x) \leq R\} \tag{1}$$

for any  $R > K/(1 - \gamma)$ .

# Mean squared error analysis

## Theorem

*Under the geometric drift condition,*

$$\sup_{|\varphi| \leq \sqrt{V}} \mathbb{E} \left[ \overline{S_{\tau_C}(\varphi)} - \int \varphi d\Pi \right]^2 \lesssim \frac{R + K}{M(2 - \gamma_R)^2} + \frac{(R + K)^2 \int w d\Pi}{N(1 - \gamma_R)^2}$$

*where  $\gamma_R > \gamma$ .*

- The actual constants are explicit and non-asymptotic
- If  $V(\cdot) \geq \|\cdot\|_2^2$  can estimate the posterior mean
- Does not require convergence analysis of the Markov chain

# MSC estimator properties

- The drift condition ensures the Markov chain simulation is short in length. Not always available e.g. Metropolis-Hastings.
- Requires importance sampling proposal and drift condition, but does not require convergence analysis

## MSC estimator concentration

Under uniformly bounded importance weights and a multiplicative drift condition, there is a stronger sub-Gaussian concentration inequality.



# Applications

## Toy example

Consider the autoregressive process

$$X_t = \rho X_{t-1} + \sqrt{1 - \rho^2} \xi_t \quad \xi_t \sim N(0, I) \text{ independent}$$

has invariant distribution  $N(0, I)$ .

# MSC construction

- Use the autoregressive process as the Markov chain
- Use importance sampling proposal  $N(0, (1/2 + h)I_d)$ .
- For any  $r > 1$ , define the set

$$C_r = \{\|x\|_2^2 \leq rd\}$$

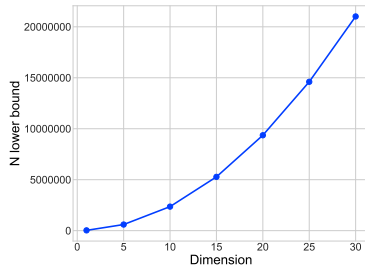
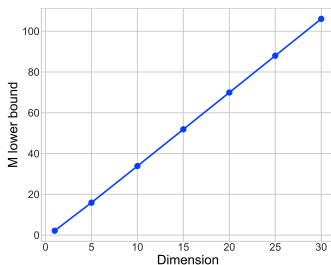
# MSE error bound

If  $N, M$  sufficiently large, then for  $|\varphi| \leq \|\cdot\|_2$ ,

$$\overline{S_{\tau_{C_r}}(\varphi)} \approx \mathbb{E}[\varphi(Z)]$$

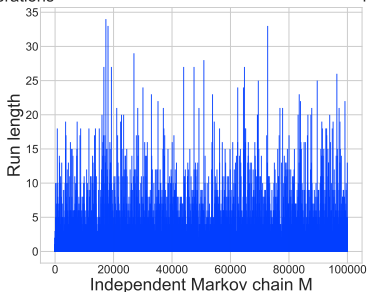
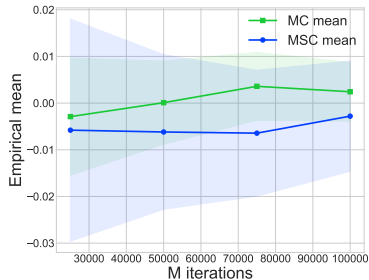
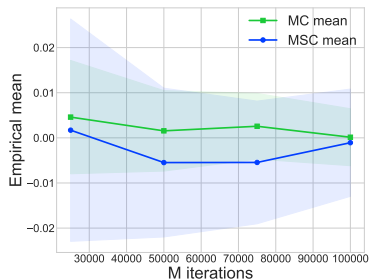
$$Z \sim N(0, I)$$

with high probability.



# Simulation result

Use  $N = 10^6$  and  $M = 10^5$  in dimension 2 to estimate the mean.



# Predicting cardiovascular disease

The aim is to predict cardiovascular disease from a data set provided by the Cleveland Clinic [[Detrano et al., 1989](#)]. The data consists of 303 patients with binary responses determining if cardiovascular disease is present and 21 covariates on patient characteristics.

# Bayesian logistic regression

Consider Bayesian logistic regression with a Gaussian prior

$$\beta \sim N(0, \Sigma)$$

The Pólya-Gamma Gibbs sampler [[Nicholas G. Polson and Windle, 2013](#)] is popular but the convergence rate [[Choi and Hobert, 2013](#)] can be problematic in moderate dimensions.

# MSC construction

- Importance sampling proposal  $N(\beta_n^*, (1/2 + h)\Sigma)$  with  $h \in (0, 1/2]$ .
- Marginal Pólya-Gamma Gibbs Markov chain  $(\beta_t)_t$ .
- For any  $r > 1$ ,

$$C_r = \{\|\beta\|_2^2 \leq rL\}. \quad (2)$$

where  $L = \|\Sigma\|_2^2 \|X^T(Y - 1/2\mathbf{1}_d)\|_2^2$ .



# MSC error analysis

Using the MSC estimator with this Pólya-Gamma Gibbs sampler:

## Proposition

Then for  $|\varphi| \leq \|\cdot\|_2$ ,

$$MSE[S_{\tau_{Cr}}(\varphi)] \lesssim \frac{rL}{M(2 - \gamma_r)^2} + \frac{[rL]^2 \left( \frac{1}{2\sqrt{2h}} + \sqrt{\frac{h}{2}} \right)^d}{N(1 - \gamma_r)^2}$$

where  $\gamma_r \in (0, 1)$ .

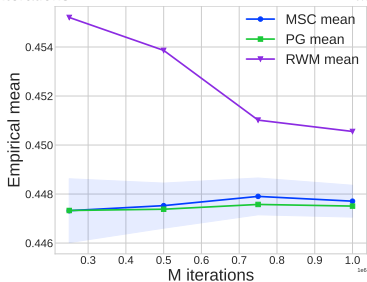
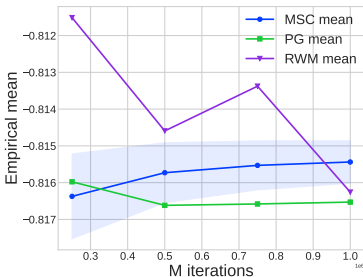
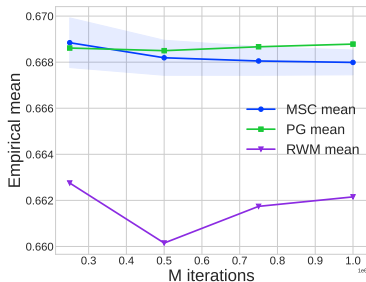
- The bound is computable
- Since the drift condition could be improved this will result 1 step runs of the Markov chain.

## Simulation setup

Use  $N = 10^7$  and  $M = 10^6$  to estimate the posterior mean for the cardiovascular data. Use some reasonable tuning parameters and prior choice.

Coefficients  $\beta_1$  indicating male versus female patients,  $\beta_2$  for the number of major blood vessels colored by flourosocopy, and  $\beta_3$  for resting blood pressure in mm/Hg on admission to the hospital.

# Simulation result



# Summary

- Developed an estimator guaranteed in the *many-short-chains* regime and error analysis that does not depend on convergence of the Markov chain.
- The bounds can have issues scaling to high dimensional problems
- The bounds appear useful compared to importance sampling bounds for unbounded functions [[Agapiou et al., 2017](#), Theorem 2.3].

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