

# Monte Carlo *and* Simulation

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# Introduction

# Monte Carlo: principle

$$\mathbb{E}[\varphi(X)] \approx \frac{1}{N} \sum_{n=1}^N \varphi(X_n)$$

with  $\varphi : \mathcal{X} \rightarrow \mathbb{R}$ .

Rationale: MSE, Law of large number, central limit theorems.

Also: **confidence intervals!**

Need for *simulation* methods. Note that simulation has other uses beyond Monte Carlo.

# Bibliography

- Stats, MCMC: *Monte Carlo statistical methods*, C.P Robert, Springer
- IID simulation: *non-uniform random variate generation*, L. Devroye, Springer
- in French: *La simulation de Monte Carlo*, B. Tuffin, Lavoisier
- QMC: *Monte Carlo and quasi-Monte Carlo sampling*, C. Lemieux, Springer
- finance: *Monte Carlo methods in financial engineering*, P. Glasserman, Springer

# Pseudo-random number generators

# Famous quotes, outline

*Anyone who uses software to produce random numbers is in a “state of sin”.*

John von Neumann

*One should not use a random method to generate random numbers.* Donald

Knuth

A PRNG is a *convenient fiction*. Ideally, it should:

- be *fast*,
- be *reproducible*,
- look *random* (at least according to statistical tests, e.g. “die-hard”).

# PRNGs: a few facts

- The general structure of a PRNG:  $x_t = f(x_{t-1})$ , where  $x_t \in \{0, \dots, 2^k - 1\}$ ; by construction,  $x_t$  is **periodic**.
- LCG (linear congruential generators):

$$x_{t+1} = (ax_t + c) \pmod{m}$$

and take  $u_t = x_t/m$  so that the  $u_t$ 's are in  $[0, 1]$ .

- Take  $c = 0$  for simplicity (then seed 0 is forbidden; and 0 is never generated, provided  $m$  is prime, and  $a < m$ ).
- Assuming  $m$  is prime, the period is  $m - 1$  iff  $a^k - 1$  is a multiple of  $m$  for  $k = m - 1$ , but not  $k \leq m - 2$ .



# Lattice structure

- Vectors of dim  $d$  lie on at most  $(d!m)^{1/d}$  hyperplanes in the  $d$ -dimensional unit cube; e.g. for  $m = 2^{31} - 1$ , 108 for  $d = 3$  and 39 for  $d = 10$ .
- RANDU, the most ill-conceived random number generators ever designed. . . has  $a = 65539 = 2^{16} + 3$ ,  $c = 2^{31}$ , and is such that  $x_t = 6x_{t-1} - 9x_{t-2}$ .
- See Table 2.1 p 44 of Glasserman for better choices of  $(a,c)$ .
- note that if  $a$  is not small, then computing  $a * x$  is not easy even using floating point operations. We could take  $a = 2^k$ , but then generators typically have bad properties (see RANDU).

# More modern PRNGs

- basic LCGs (even with *good* values of  $a$  and  $c$ ) are now considered obsolete.
- Combine several generators to (a) increase period; and (b) reduce lattice structure: e.g. take the sum of  $K$  generators modulo one (Wichmann-Hill).
- Mersenne twister: very popular 32-bit PRNG (Python, R, Matlab, etc), has period  $2^{19937} - 1$ .
- Also push for 64-bit PRNG.

# Main conclusion

- **DO NOT** use C standard implementation `rand()`.
- **DO NOT** implement your own PRNG.
- **DO** resort to some **modern** implementation of a **modern** generator, such as Mersenne twister; see e.g. GSL in C.

## Non-uniform simulation

# Outline

A few general recipes:

- inversion
- rejection
- chain rule

plus several specialised ones (e.g. Box-Muller).

# inversion

## inversion algorithm

If  $X$  has CDF  $F$ , take

$$X = F^{-1}(U), \quad U \sim \mathcal{U}[0, 1].$$

Applications: exponential, Laplace, Gaussian?

# Box-Muller

## Box-Muller

$$\begin{cases} X &= \sqrt{-2 \log(U)} * \cos(2\pi V) \\ Y &= \sqrt{-2 \log(U)} * \sin(2\pi V) \end{cases}$$

Then  $X, Y \sim N(0, 1)$ , independently.

# A sneaky introduction to rejection

To understand the coming slides, note that the following algorithm

## Rejection

Repeat  $X \sim \mathcal{U}(\mathcal{A})$

Until  $X \in \mathcal{B}$ .

draws from  $\mathcal{U}(\mathcal{B})$  (provided  $\mathcal{B} \subset \mathcal{A}$ ).



# Modified Box-Muller

## Box-Muller with rejection

Repeat

$$U, V \sim \mathcal{U}[-1, 1]$$

Until  $S := U^2 + V^2 \leq 1$ .

Return

$$\begin{cases} X &= U\sqrt{-2\log(S)/S} \\ Y &= V\sqrt{-2\log(S)/S} \end{cases}$$

Then  $X, Y \sim N(0, 1)$ , independently.

Note: avoid computing sin and cos.

# Rejection

Let  $f, g$  PDFs such that  $f \leq Mg$  (with  $M \geq 1$ ).

Accept-reject

Repeat

$$X \sim g, \quad U \sim \mathcal{U}[0, 1],$$

Until  $U \leq f(X)/Mg(X)$ .

Properties:  $X \sim f$ , number of draws until acceptance is Geometric( $1/M$ ).

Justification: uniform sampling under the graph, see next slide.

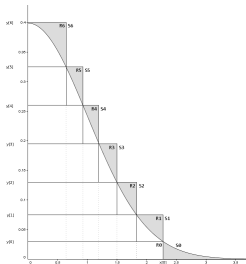
# Uniform sampling under the graph

For a function  $f$ , let  $\mathcal{G} = \{(x, y) \in \mathbb{R}^2 : 0 \leq y \leq Mf(x)\}$ , then

$$(X, Y) \sim \mathcal{U}(\mathcal{G}) \Leftrightarrow \begin{cases} X & \sim f \\ Y|X = x & \sim \mathcal{U}[0, Mf(x)] \end{cases}$$

Note: this construction is in fact not restricted to real-valued random variables.

# Ziggurat algorithm for $N(0, 1)$ (Marsaglia, 60?)



$K$  Slices  $S_k = [-x_k, x_k] \times [y_k, y_{k+1}]$  constructed to have the same area.

- 1 Choose slice  $k$  (uniformly).
- 2 Sample  $(X, Y)$  within slice  $k$ .
- 3 If  $X \leq x_{k+1}$ , return  $X$ , else, if  $Y \leq \varphi(X)$ , return  $X$ , else go to 1.

Note: If slice 0 is selected, extra steps required (truncated Gaussian distribution).

# Multivariate simulation: chain-rule decomposition

The inverse transform method is restricted to real-valued random variables, the inverse transform *is not*.

General recipe to generate jointly  $(X, Y, Z)$ , with PDF  $f(x, y, z)$ :

- 1 Generate  $X \sim f_X(x)$  (marginal). Call  $x$  the output.
- 2 Generate  $Y|X = x \sim f_{Y|X}(y|x)$  (conditional given  $X = x$ ). Call  $y$  the output.
- 3 Generate  $Z|X = x, Y = y \sim f_{Z|Y,X}(z|x, y)$  (full conditional). Call  $z$  the output.

# Gaussian vectors

The standard method to generate  $X \sim N_d(\mu, \Sigma)$  is:

- Generate  $Z_1, \dots, Z_d \sim N(0, 1)$ .
- Compute  $C = \text{Choleksy}(\Sigma)$ . (i.e.  $\Sigma = CC^T$ , and  $C$  is lower triangular)
- Return  $X = \mu + CZ$ .

The Cholesky decomposition costs  $\mathcal{O}(d^3)$ .

# Non-uniform simulation in spaces other than $\mathbb{R}^d$

# Outline

Some recipes to sample specific cases of

- distributions over constrained sets
- discrete distributions



# How to sample $N$ sorted uniforms

Naive method: sample  $U_n \sim \mathcal{U}[0, 1]$  for  $n = 1, \dots, N$ , return  $\text{sort}(U_{1:N})$ .  
Cost is  $\mathcal{O}(N \log N)$  (not bad).

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Smart  $\mathcal{O}(N)$  method:

- Sample  $E_1, \dots, E_{N+1} \sim \text{Exp}(1)$ .
- Compute  $V_{1:(N+1)} = \text{cumsum}(E_{1:(N+1)})$ .
- Return  $(V_1/V_{N+1}, \dots, V_N/V_{N+1})$ .

# How to sample uniformly on the sphere

- Sample  $X \sim N_d(0, I_d)$ .
- Return  $X/\|X\|$ .

# How to sample from a discrete distribution over $\mathbb{N}$

The inverse method extends to the discrete case. Simply define:

$$F^{-1}(u) = \inf\{x : F(x) \geq u\}$$

In practice:

- Sample  $U \sim \mathcal{U}[0, 1]$
- If  $U \leq p_0$ , return 0
- If  $p_0 < U \leq p_0 + p_1$ , return 1
- etc

# What if $N$ and $K$ are large

Suppose we want to sample  $N$  times from a distribution over  $\{0, \dots, K-1\}$ . If we run the algorithm of the previous slide  $N$  times, we do  $\mathcal{O}(NK)$  operations (on average). Can we do better?

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Solution: use as input  $N$  sorted uniforms. Then cost is  $\mathcal{O}(N + K)$ .

Application: (weighted) bootstrap.

# Inverse CDF algorithm

```
def inversecdf(su,W):
    """ Input:  su[0:N] sorted uniforms
            W[0:K] normalised weights (sum to one)
    Output: A[0:N] indexes (in {0,...,K-1})
    """
    j=0; s=W[0]; N = su.shape[0]
    A = empty(N,'int')
    for n in xrange(N):
        while su[n]>s:
            j += 1
            s += W[j]
        A[n] = j
    return A
```

# How to sample a permutation

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Smart  $\mathcal{O}(N)$  method:

- Let  $\sigma = (1, 2, \dots, N)$ .
- $I \sim \mathcal{U}(1, \dots, N)$ , swap  $\sigma(1)$  and  $\sigma(I)$ .
- $I \sim \mathcal{U}(2, \dots, N)$ , swap  $\sigma(2)$  and  $\sigma(I)$ .
- etc.

## Variance reduction

# Objectives, outline

Given a certain quantity

$$I = \mathbb{E}[\varphi(X)] = \int_{\mathcal{X}} f(x)\varphi(x) dx$$

find a Monte Carlo estimator with smaller variance than the the standard estimator

$$\frac{1}{N} \sum_{n=1}^N \varphi(X_n).$$

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Recipes:

- antithetic variables
- control variates

# Antithetic variables

In cases where  $\varphi(-X)$  has the same distribution as  $\varphi(X)$ , use:

$$\hat{l}_{\text{anti}} = \frac{1}{2N} \sum_{n=1}^N \{\varphi(X_n) + \varphi(-X_n)\}$$

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**Lemma:**

$$\text{Var}(\hat{l}_{\text{anti}}) \leq \text{Var}(\hat{l})$$

Note: we have less variance, but twice as many evaluations of  $\varphi \dots$

# Control variates (univariate case)

Let  $Z$  a real-valued r.v. such that  $\mathbb{E}(Z) = 0$ . For any  $\beta$ ,

$$\hat{l}_{\text{cv}} = \frac{1}{N} \sum_{n=1}^N \{\varphi(X_n) - \beta Z_n\}$$

is an unbiased estimator of  $I = \mathbb{E}[\varphi(X)]$ .



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The smallest variance is obtained by taking

$$\beta_{\text{opt}} = \frac{\text{Cov}(\varphi(X), Z)}{\text{Var}(Z)}.$$

# Control variates (multivariate case)

$Z^1, \dots, Z^K$  are mean-zero real-valued r.v. Take

$$\hat{I}_{\text{cv}} = \frac{1}{N} \sum_{n=1}^N \left\{ \varphi(X_n) + \sum_{k=1}^K \beta_k Z_n^k \right\}.$$

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In practice, replace  $\beta_k$  by  $\hat{\beta}_k$ , the OLS estimate for regression:

$$\varphi(X_n) = \alpha + \sum_{k=1}^K \beta_k Z_n^k + \varepsilon_n.$$

# Variance reduction and Rao-Blackwellisation

Often variance reduction techniques may be cast as particular **Rao-Blackwellisation** schemes, i.e. the idea that

$$\text{Var} [\mathbb{E}[\varphi(X)|Z]] \leq \text{Var}[\varphi(X)].$$

# Importance sampling

# A simple identity

$$\begin{aligned}\mathbb{E}[\varphi(X)] &= \int_{\mathcal{X}} \varphi(x) f(x) dx \\ &= \int_{\mathcal{X}} \varphi(x) \frac{f(x)}{g(x)} g(x) dx = \mathbb{E}_g \left[ \frac{f(X)}{g(X)} \varphi(X) \right]\end{aligned}$$

assuming  $\text{Supp}(f) \subset \text{Supp}(g)$ .

Any expectation w.r.t. PDF  $f$  may be rewritten thusly as an expectation w.r.t. PDF  $g$  (which may be easier to simulate from):

$$\hat{I}_{IS} = \frac{1}{N} \sum_{n=1}^N \frac{f(X_n)}{g(X_n)} \varphi(X_n).$$

# How to choose proposal $g$ ?

- 1 Check that variance exists,  $\Leftrightarrow \mathbb{E}_g \left[ \varphi(X)^2 \frac{f(X)^2}{g(X)^2} \right] < \infty$ . (Sufficient condition:  $f/g \leq M$ , and  $\mathbb{E}_f[\varphi^2] < \infty$ .)

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- 2 Optimal  $g$  (in terms of minimizing variance) is

$$g_{\text{opt}}(x) \propto f(x)|\varphi(x)|.$$

It is often not possible to simulate from  $g_{\text{opt}}$ , so more generally, it is recommended to take  $g \approx f$ .



# Auto-normalised IS

Sometimes either  $f$  or  $g$  are known only up to a constant:  $f = f_u/Z_f$ ,  $g = g_u/Z_g$ , and  $Z_f$ ,  $Z_g$  are intractable. In that case, we use the auto-normalised IS estimator:

$$\hat{I}_{\text{AIS}} = \frac{\sum_{n=1}^N w_n \varphi(X_n)}{\sum_{n=1}^N w_n}, \quad w_n = \frac{f_u(X_n)}{g_u(X_n)}.$$

This estimator is biased, and asymptotically Gaussian:

$$\sqrt{N} (\hat{I}_{\text{AIS}} - I) \Rightarrow N(0, v_{f/g})$$

with  $v_{f/g} = \mathbb{E}_g \left[ \left( \frac{f}{g} \right)^2 (\varphi - I)^2 \right]$  (assuming this quantity is  $< \infty$ ).

# How to choose $g$ (bis repetita)

Same points as for standard IS:

- ① Check that at least  $v_{f/g} < \infty$ ; sufficient condition is (a)  $f/g < M$  and (b)  $\text{Var}_f(\varphi) < \infty$ .
- ② Optimal  $g$  is

$$g_{\text{opt}}(x) \propto f(x)|\varphi(x) - I|$$

which depends on  $I \dots$  In practice, take  $g \approx f$ .

# Estimating the $Z$ 's, effective sample size

Note that AIS also provides an estimate of  $Z_f/Z_g$ :

$$\mathbb{E} \left[ \frac{1}{N} \sum_{n=1}^N w_n \right] = \frac{Z_f}{Z_g}$$

and of  $v_{f/g}$ :

$$\frac{N \sum_{n=1}^N w_n^2 \left\{ \varphi(X_n) - \hat{l} \right\}^2}{\left( \sum_{n=1}^N w_n \right)^2}.$$

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Similarly, the **effective sample size**

$$\frac{\left( \sum_{n=1}^N w_n \right)^2}{\sum_{n=1}^N (w_n)^2} \in [1, N]$$

is a good indicator of AIS efficiency.

# Curse of dimensionality

For  $\mathcal{X} = \mathbb{R}^d$ ,  $f(x) = \prod_{i=1}^d f_1(x_i)$ ,  $g(x) = \prod_{i=1}^d g_1(x_i)$ , one has:

$$\mathbb{E}_g[f^2/g^2] = C^d, \quad C \geq 1.$$

We expect the variance of IS to grow exponentially with the dimension.

# Resampling

How to transform weighted sample  $(w_n, X_n)$  into an **unweighted** sample?

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Simply draw randomly  $\tilde{X}_n$  from

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(as in the bootstrap).

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(as in the bootstrap).

See previous chapter on multinomial sampling.



# Quasi-Monte Carlo

# Principle

Often one may rewrite quantity of interest as:

$$I = \mathbb{E}[\varphi(U)], \quad U \sim \mathcal{U}[0, 1]^d$$

and then use

$$\hat{I} = \frac{1}{N} \sum_{n=1}^N \varphi(U_n).$$

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Can we construct (deterministic or random) points  $U_1, \dots, U_N$  in  $[0, 1]^d$  so that the approximation error is smaller than with standard Monte Carlo (i.e.  $U_n$  are IID uniforms)?

# Stratification ( $d = 1$ )

- Generate  $N/K$  uniforms in each interval  $[(k-1)/K, k/K]$ ,  $k = 1, \dots, K$ . (Note the connection with antithetic variables.)

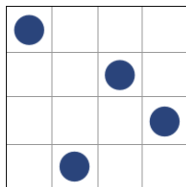
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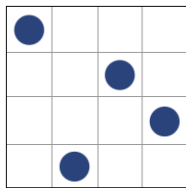
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- Or even take  $K = N$ , i.e. generate  $U_n \sim \mathcal{U}[(n-1)/N, n/N]$ .
- or even take  $u_n = (2n-1)/2N$ , the (deterministic) centre of interval  $[(n-1)/N, n/N]$ .

# Stratification for $d > 1$ : Latin hypercube sampling



Generate the  $U_n$ 's so that exactly one point falls in each horizontal or vertical strip (of area  $1/N$ ).

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**Tip:** use random permutations.



# Koksma–Hlawka inequality

$$|\hat{I} - I| \leq V(\varphi) D^*(u_{1:N})$$

where  $V(\varphi)$  is the variation of  $\varphi$  (in the sense of Hardy and Krause), and

$$D^*(u_{1:N}) = \sup_{[0,b] \subset [0,1]^d} \left| N^{-1} \sum_{n=1}^N \mathbb{I}_{[0,b]}(u_n) - \prod_{i=1}^d b_i \right|$$

is the **star discrepancy**.

# Proof for $d = 1$

$$N^{-1} \sum_{n=1}^N \varphi(u_n) - \int_0^1 \varphi(u) du = \int_0^1 \delta(u) \varphi'(u) du$$

where  $\delta(u) = u - N^{-1} \sum_{n=1}^N \mathbf{1}(u_n \leq u)$ .

# Why is the previous inequality so important?

Because we know how to construct:

- point-sets such that  $D^*(u_{1:N}) = \mathcal{O}\left(\frac{(\log N)^{d-1}}{N}\right)$
- sequences such that  $D^*(u_{1:N}) = \mathcal{O}\left(\frac{(\log N)^d}{N}\right)$

hence we can do **better** than Monte Carlo, i.e.  $\mathcal{O}_P\left(\frac{1}{\sqrt{N}}\right)$ .

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hence we can do **better** than Monte Carlo, i.e.  $\mathcal{O}_P\left(\frac{1}{\sqrt{N}}\right)$ .

Side note: there are good reasons to believe that these rates are optimal.

$$d = 1$$

Take  $u_n = (2n - 1)/2N$ ,  $n = 1, \dots, N$ . Then

$$D^*(u_{1:N}) = \frac{1}{2N}.$$

# Van der Corput (sequence for $d = 1$ )

In base  $b$ , for  $n = \sum_{j=0}^k a_j(n)b^j$ , take

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e.g. for  $b = 2$ :  $1/2, 1/4, 3/4, 1/8, \dots$

For  $b = 3$ :  $1/3, 2/3, 1/9, \dots$

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For  $b = 3$ :  $1/3, 2/3, 1/9, \dots$

Then  $D^*(u_{1:N}) = \mathcal{O}(\log N/N)$ .



## $d > 1$ : Halton & Hammersley

**Halton sequence:** component  $j$  is a van der Corput sequence in base  $b_j$ , where the  $b_j$  are the first  $d$  prime numbers. Discrepancy is  $\mathcal{O}((\log N)^d / N)$ .

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**Hammersley point set** (of size  $N$ ): take  $N$  first elements of Halton sequence of dimension  $d$ , replace last component by  $n/N$ . Discrepancy is  $\mathcal{O}((\log N)^{d-1}/N)$ .

## $d > 1$ : Halton & Hammersley

**Halton sequence:** component  $j$  is a van der Corput sequence in base  $b_j$ , where the  $b_j$  are the first  $d$  prime numbers. Discrepancy is  $\mathcal{O}((\log N)^d/N)$ .

**Hammersley point set** (of size  $N$ ): take  $N$  first elements of Halton sequence of dimension  $d$ , replace last component by  $n/N$ .

Discrepancy is  $\mathcal{O}((\log N)^{d-1}/N)$ .

Note however that for large  $d$ , both Halton and Hammersley require many points to cover the space. . .

# Other low-discrepancy sequences and point sets

- Niederreiter
- Faure
- Sobol'
- ...

# RQMC (randomised QMC)

QMC is purely deterministic. It lacks a simple way to evaluate the numerical error. Imagine we are able to randomise  $U_{1:N}$  so that

- ①  $U_n \sim \mathcal{U}[0, 1]^d$  (marginally).
- ②  $U_{1:N}$  is still a low-discrepancy point-set (or sequence).

Then

$$\mathbb{E} \left[ \frac{1}{N} \sum_{n=1}^N \varphi(U_n) \right] = \mathbb{E}[\varphi(U)]$$

and we can evaluate the numerical error through the empirical variance (over repeated runs).

# RQMC: random shift

The simplest RQMC strategy is to generate a low-discrepancy point set  $v_{1:N}$ ,  $W \sim \mathcal{U}[0, 1]^d$ , then take:

$$U_n = v_n + W \pmod{1} \quad (\text{componentwise})$$

# RQMC: a surprising result

Owen (1998) showed that for smooth functions  $\varphi$

$$\text{Var}[\hat{I}] = \mathcal{O}\left(\frac{(\log N)^{(d-1)/2}}{N^3}\right)$$

when scrambling (a particular RQMC technique) is used.

## conclusion: QMC vs MC

- QMC has a better convergence rate.
- But for large  $d$ , QMC might need a very large  $N$  to beat MC.
- With MC, the (square) error is simple to estimate, whereas for QMC, we have only a deterministic bound, which is hard to evaluate, and is often pessimistic. See RQMC however.
- Variance reduction: may be used in conjunction with (R)QMC. (Recommendation is to do variance reduction, *then* replace MC with QMC).
- Practical recommendation: scrambled Sobol' seems like a good default choice (or Latin Hypercube sampling for very high dimensions).



# Markov chain Monte Carlo

# Outline

In some settings, simulating *independently*  $X \sim \pi(dx)$  is difficult, but it is possible to simulate a Markov chain  $(X_n)$  that leaves  $\pi(dx)$  **invariant**. Then, we still have

$$\frac{1}{N} \sum_{n=1}^N \varphi(X_n) \approx \mathbb{E}_{\pi}[\varphi(X)]$$

in some sense.

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in some sense.

This is the case in particular when density  $\pi$  is known only up to a constant.

# Definitions

- A **Markov kernel**  $K(x, dy)$  is an application  $\mathcal{X} \rightarrow \mathcal{P}(\mathcal{X})$ .

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- A Markov kernel  $K$  leaves distribution  $\pi$  **invariant** iff

$$\int_{\mathcal{X}} \pi(dx) K(x, dy) = \pi(dy).$$

- A Markov kernel is **reversible** w.r.t.  $\pi$  iff

$$\pi(dx) K(x, dy) = \pi(dy) K(y, dx).$$

This implies that  $\pi$  is invariant.

# Metropolis-Hastings

Let  $Q(x, dy)$  a Markov kernel, such that  $Q(x, dy) = q(x, y)dy$ .

## Hastings-Metropolis step

Input:  $X_{n-1}$

- ① Generate  $Y \sim Q(X_{n-1}, dy)$
- ② With probability  $1 \wedge r(X_{n-1}, Y)$ , where

$$r(x, y) = \frac{\pi(y)q(y, x)}{\pi(x)q(x, y)}$$

accept  $Y$ , i.e.  $X_n = Y$ ; otherwise  $X_n = X_{n-1}$ .

**Property:** This kernel is reversible (w.r.t.  $\pi$ ).

# An important practical point

Note that Hastings-Metropolis may be implemented even if  $\pi$  is known only up to a constant:  $\pi(x) = \pi_u(x)/Z$ ,  $Z$  is intractable. Then

$$r(x, y) = \frac{\pi(y)q(y, x)}{\pi(x)q(x, y)} = \frac{\pi_u(y)q(y, x)}{\pi_u(x)q(x, y)}$$



# Examples of MH algorithms

- ①  $q(x, y) = q(y, x)$ , for instance  $q(x, y) = N(x; y, \Sigma)$  (Gaussian **random walk**); then

$$r = \frac{\pi(y)}{\pi(x)}$$

- ②  $q(x, y) = q(y)$ : **independent Metropolis**; then

$$r = \frac{\pi(y)q(x)}{\pi(x)q(y)}$$

- ③ Langevin proposal:

$$Y \sim N\left(x + \frac{1}{2}\Sigma\nabla \log \pi(X_n), \Sigma\right)$$

## (two-block) Gibbs sampling

Assume  $X = (X_1, X_2)$ ,  $\pi(x) = \pi(x_1, x_2)$ , with conditional distributions  $\pi_{1|2}(x_1|x_2)$ ,  $\pi_{2|1}(x_2|x_1)$ .

### Gibbs sampling step

Input:  $X_{n-1} = (X_{n-1,1}, X_{n-1,2})$

- 1 Generate  $X_{n,1} \sim f_{1|2}(\bullet|X_{n-1,2})$ .
- 2 Generate  $X_{n,2} \sim f_{2|1}(\bullet|X_{n,1})$ .

Again, this leaves invariant  $\pi$ . Gibbs can be generalised to  $k > 2$  blocks.

# Combining chains

- 1 If  $K_1$ ,  $K_2$  leave  $\pi$  invariant, then so does  $K_1 K_2$ .
- 2 **Within** Gibbs, we can replace the exact simulation of  $X_1|X_2$  (say) by a Metropolis step that leaves invariant  $\pi_{1|2}$ .

# MCMC in practice

- ① Assess how long it takes for the chain to reach stationarity;
- ② When chain seems stationary, check for intra-correlations, i.e. look at ACF (Auto-Correlation Function).

Then we compute averages

$$\frac{1}{N - N_0} \sum_{n=N_0+1}^N \varphi(X_n)$$

where  $N_0$  is burn-in time, and  $N - N_0$  is sufficiently large relative to the **auto-correlation time** (i.e. time  $k$  so that  $X_n$  and  $X_{n+k}$  are nearly uncorelated).

# scaling random walk Metropolis

One big hurdle with random walk Metropolis is the choice of  $\Sigma$ , in the proposal  $N(x, \Sigma)$ . If too small, chain moves slowly, if too large, proposals always get rejected.

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One big hurdle with random walk Metropolis is the choice of  $\Sigma$ , in the proposal  $N(x, \Sigma)$ . If too small, chain moves slowly, if too large, proposals always get rejected.

Theory (e.g. Roberts and Rosenthal, 2004) indicates that one should take

$$\Sigma = c\Sigma_{\pi}$$

where  $\Sigma_{\pi}$  is the covariance matrix of target  $\pi$ , and  $c$  calibrated so that acceptance rate is  $\approx 0.25$ .

# a tiny bit of MCMC theory

- ① From an arbitrary starting point  $X_0 = x_0$ , and any  $\varepsilon > 0$ , we have

$$\|K^n(x_0, dx_n) - \pi(dx_n)\|_{\text{TV}} \leq \varepsilon$$

for  $n$  large enough.

- ② CLT:

$$\sqrt{N} \left( \frac{1}{N} \sum_{n=1}^N \varphi(X_n) - I \right) \Rightarrow N(0, V(\varphi))$$

with

$$V(\varphi) = \text{Var}_{\pi}(\varphi) + 2 \sum_{k=1}^{\infty} \gamma_k(\varphi)$$

and  $\gamma_k(\varphi) = \text{Cov}[\varphi(X_n), \varphi(X_{n+k})]$ .

# Adaptive MCMC?

Can we use past samples to automatically calibrate Metropolis-Hastings?  
e.g. at time  $t$ , do a random walk Metropolis step, of size  $\Sigma = c\hat{\Sigma}_t$ , where  $\hat{\Sigma}_t$  is the empirical covariance matrix computed from  $X_0, \dots, X_{t-1}$ .



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Big theoretical problem: we are not simulating a Markov chain any more ( $X_t$  depends on the whole past). Convergence is more difficult to establish.

## Interlude: Bayesian classification

# Outline

Consider model with responses  $y_i \in \{-1, 1\}$ , covariates  $x_i \in \mathbb{R}^p$ , likelihood

$$L(x, y; \beta) = \prod_{i=1}^{n_d} F(y_i \beta^T x_i)$$

with  $F = \Phi$  (probit), or  $F = L$  (logit), and prior

$$\pi(\beta) = 1$$

# Outline

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with  $F = \Phi$  (probit), or  $F = L$  (logit), and prior

$$\pi(\beta) = 1$$

The posterior is

$$\pi(\beta|x, y) \propto \prod_{i=1}^{n_d} F(y_i \beta^T x_i)$$

We will use this example to discuss many of the approaches seen so far.

# Monte Carlo optimisation

# Objectives

Numerical maximisation:

$$\max_{\theta \in \Theta} \psi(\theta)$$

when

- ①  $\psi$  can be evaluated point-wise, but is difficult to maximise by standard methods: **Exploration**
- ②  $\psi$  is an (intractable) expectation:

$$\psi(\theta) = \mathbb{E}_{\theta}[h(X, \theta)]$$

## Stochastic approximation

Statistical applications: MLE

# Exploration

When  $\psi$  can be evaluated point-wise, one may sample  $N$  times from some distribution  $\pi(d\theta)$ , and return  $\max_{n=1,\dots,N} \psi(\theta_n)$ ; for instance if  $\Theta$  is compact, take  $\pi(d\theta)$  to be the Uniform dist. over  $\Theta$ .

# Exploration

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In particular, consider

$$\pi_\lambda(\theta) \propto \exp\{\lambda\psi(\theta)\}.$$

When  $\lambda$  (inverse temperature) increases, support of  $\pi_\lambda$  gets more concentrated around modal regions, but in return it may be more difficult to sample from  $\pi_\lambda$ .



# Simulated annealing

Simulate a (inhomogeneous) Markov chain as follows: at iteration  $t$ , do a Metropolis step w.r.t.  $\pi_{\lambda_t}$ , and make  $\lambda_t$  increase at a logarithmic rate.

# The Cross-Entropy method

For some parametric family  $\{f_\xi, \xi \in \Xi\}$ , choose initial  $\xi_0$ , then iteratively:

- 1 Sample  $\theta_1, \dots, \theta_n \sim f_{\xi_t}$ .
- 2 Estimate (using e.g. MLE)  $\xi_{t+1}$  from the 10% best of the  $\theta_i$  (in terms of  $\psi(\theta_i)$ ).

# Other heuristic optimisation procedures

- genetic algorithms
- tabu search
- ant colony algorithm

and also more specialised ones.

# Stochastic approximation

One has:  $\psi(\theta) = \mathbb{E}_\theta[h(X, \theta)]$  (double dependence on  $\theta$ ). Possible approaches:

- 1 If Expectation is w.r.t. a fixed dist'  $f$ ,  $\psi(\theta) = \mathbb{E}[h(X, \theta)]$ , generate  $X_1, \dots, X_n \sim f$ , maximise  $\theta \rightarrow N^{-1} \sum_{n=1}^N h(X_n, \theta)$ .
- 2 Gradient-based approach, e.g.

$$\theta_{t+1} = \theta_t + \alpha_t \hat{\nabla} \psi(\theta_t)$$

where  $\hat{\nabla} \psi(\theta_t)$  is some MC estimate of the gradient of  $\psi$ .

# Robins-Monroe

Take  $\alpha_t$  such that  $\alpha_t \rightarrow 0$ , and  $\sum_t \alpha_t = \infty$ ; e.g.  $\alpha_t = Ct^{-b}$ ,  $1/2 < b \leq 1$ .

# Robins-Monroe

Take  $\alpha_t$  such that  $\alpha_t \rightarrow 0$ , and  $\sum_t \alpha_t = \infty$ ; e.g.  $\alpha_t = Ct^{-b}$ ,  $1/2 < b \leq 1$ .  
To estimate the gradient, if  $\psi(\theta) = \mathbb{E}_\theta[h(X)]$ , one has

$$\nabla \psi(\theta) = \mathbb{E}_\theta[h(X)s_\theta(X)], \quad s_\theta(x) = \frac{\partial}{\partial \theta} \log f_\theta(x)$$

and thus a possible choice is:

$$\hat{\nabla} \psi(\theta) = \frac{1}{N} \sum_{n=1}^N h(X_n) s_\theta(X_n)$$

## Selected applications of Monte Carlo

# Outline

- 1 Derivative pricing
- 2 Statistical applications: MCEM, Bayesian inference, ABC
- 3 Enumeration
- 4 Go playing. . .



# Derivative pricing: statement

There,  $X$  is continuous-time process on  $[0, T]$ , and  $\varphi$  could be:

- $\varphi(X) = (K - X_T)^+$
- $\varphi(X) = (K - \frac{1}{k} \sum_i X_{t_i})^+$
- $\varphi(X) = (K - \int X_t dt)^+$
- $\varphi(X) = \mathbb{I}\{\tau_b > T\}(X_T - K)^+$ , with  $\tau_b = \inf\{t : X_t \leq b\}$
- etc.

# Simulating Brownian paths

For a Brownian motion  $\{W_t\}$ , several ways to simulate *exactly*  $(W_{t_1}, \dots, W_{t_k})$ :

- random walk:  $W_{t_i} | W_{t_{i-1}} \sim N(W_{t_{i-1}}, t_i - t_{i-1})$
- Brownian bridge:  $W_{t_i} | W_{t_{i-1}}, W_{t_{i+1}} \sim$

$$N\left(\frac{(t_{i+1} - t_i)W_{t_{i-1}} + (t_i - t_{i-1})W_{t_{i+1}}}{t_{i+1} - t_{i-1}}, \frac{(t_{i+1} - t_i)(t_i - t_{i-1})}{t_{i+1} - t_{i-1}}\right)$$

- principal components

Try to think about the implications for QMC...

# QMC and Brownian paths

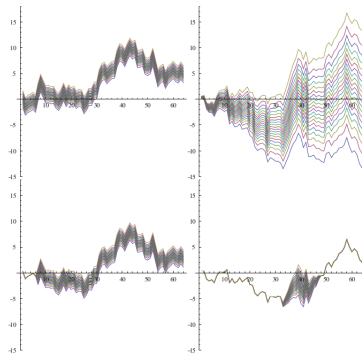


Fig. 8.6 Paths of Brownian motion obtained by the forward construction (*left*) and the Brownian bridge construction (*right*). All but one parameters are fixed

Top: all but first component fixed; bottom, all but seventh component fixed

# Discretization

In general, diffusion processes need to be discretized:

$$dX_t = \mu(X_t)dt + \sigma(X_t)dW_t$$

becomes

$$X_{t+1} - X_t = \delta\mu(X_t) + \sigma(X_t)\epsilon_t, \quad \epsilon_t \sim N(0, \delta)$$

where  $\delta$  is the discretization step.

Choice of  $\delta$ : trade-off between discretization bias and CPU time.

# Multi-level Monte Carlo

Consider a sequence of decreasing steps:  $\delta_0 > \dots > \delta_L$ ; say  $\delta_l = 2^{-l}$ .

$$\mathbb{E}_{\delta_L}(\varphi) = \mathbb{E}_{\delta_0}(\varphi) + \sum_{l=1}^L \left\{ \mathbb{E}_{\delta_l}(\varphi) - \mathbb{E}_{\delta_{l-1}}(\varphi) \right\}$$

To get a low-variance estimate for each level, use **coupling**: e.g. use Brownian bridge construction to obtain the finer level from the coarser level.

To minimise variance, choose  $N_l$  (number of samples for level  $l$ ) to be:

$$N_l \propto \sqrt{V_l / C_l}$$

where  $V_l$  (resp.  $C_l$ ) is variance (resp. CPU cost per sample) of estimate for level  $l$ .

# Other worthy points

- control variates: simulation involves many Gaussian variables, with known mean and variance
- antithetic variables (Gaussian variables are symmetric)
- QMC very popular nowadays in option pricing

# Statistical applications

- Bayesian estimation: already covered, see MCMC
- Frequentist estimation: MC for the E part of EM: MCEM
- Likelihood-free inference: see next slide

# ABC (likelihood-free inference)

Data  $y^*$ , model  $p(y|\theta)$  such that (a) one can simulate from  $p(y|\theta)$ ; (b) one cannot compute the likelihood  $p(y|\theta)$ . (Many scientific models fall in this category.)



# ABC (likelihood-free inference)

Data  $y^*$ , model  $p(y|\theta)$  such that (a) one can simulate from  $p(y|\theta)$ ; (b) one cannot compute the likelihood  $p(y|\theta)$ . (Many scientific models fall in this category.)

ABC (Approximate Bayesian inference) samples from:

$$p_\varepsilon(\theta, y|y^*) \propto p(\theta)p(y|\theta)\mathbb{I}(\|s(y) - s(y^*)\| \leq \varepsilon).$$