Monte Carlo Methods for Uncertainty Quantification

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ERCOFTAC course on Mathematical Methods and Tools in Uncertainty Management and Quantification

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Lecture outline

Lecture 1: Introduction and Monte Carlo basics

- some model applications
- random number generation
- Monte Carlo estimation
- Central Limit Theorem and confidence interval

Lecture 2: Variance reduction

- basic manipulations
- control variate
- importance sampling
- stratified sampling
- Latin Hypercube
- randomised quasi-Monte Carlo

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Lecture outline

Application 1

Consider a bridge with 7 elements and pinned joints:



- weak and strong convergence
- mean square error decomposition
- multilevel Monte Carlo
- PDEs with uncertainty



Design will compute a force balance, work out compression / extension of each element, and therefore determine the natural length to be cast.

However, the manufactured elements will vary from design in both length and extensibility -14 uncertain inputs.

If two supporting joints have fixed position, then analysis has 6 unknowns (coordinates of free joints) and 6 equations (force balance at free joints).

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Application 1

Given manufacturing data on the variability of the natural length and extensibility, what might we want to know?

- RMS deviation of joint position from design
- RMS deviation of forces from design
- probability of maximum compression / extension force being outside some specified range

Note: if we turn this into a full finite element analysis, then the computational cost becomes much larger.

Application 2

Consider a square trampoline, with vertical position given by

$$T\left(\frac{\partial^2 Z}{\partial x^2} + \frac{\partial^2 Z}{\partial y^2}\right) = L(x, y), \quad 0 < x < 1, \ 0 < y < 1$$

where T is the tension and L(x, y) is the applied load.

Here the uncertainty could be in the boundary conditions:

- simplest case would be uncertainty in the 4 corner values of Z(x, y) with straight line interpolation along each edge
- a more complicated case might add a Fourier decomposition of the perturbation from the straight line interpolation

$$Z(x,0) = (1-x) Z_{0,0} + x Z_{1,0} + \sum_{n=1}^{\infty} a_n \sin(n\pi x)$$

Could also have uncertainty in the tension and the loading.

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Application 2

Again there are various outputs we might be interested in:

- average values for min Z(x, y) and max Z(x, y)
- RMS varation in these due to uncertainty

Note: biggest displacements likely to occur in the middle, not significantly affected by high order Fourier perturbations on the boundary.

Application 3

In modelling groundwater flow in nuclear waste repositories, or oil flow in oil reservoirs, we use Darcy's Law:

$$abla \cdot \left(\kappa
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ight) = 0$$

where p is the pressure and κ is the permeability of the rock.

The uncertainty here is in $\kappa(x)$ – typically might know the values at a few bore holes, but there is a large amount of uncertainty elsewhere.

We do know that if two points x_1, x_2 are close, then $\kappa(x_1) \approx \kappa(x_2)$, but if they are far apart then they can be quite different. Hence, often model log κ as having a Normal distribution, with a spatial covariance of the form

$$\operatorname{cov}(\log \kappa(x_1), \log \kappa(x_2)) = \sigma^2 \exp(-\|x_1 - x_2\|/\lambda)$$

where λ is the correlation length.

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Application 3

What does this application require?

- generation of samples of the stochastic field $\log \kappa(x)$ with correct distribution
- computation of p(x) by solving Darcy PDE
- evaluation of outputs of interest (e.g. water or oil flux across some boundary)
- Monte Carlo simulation to obtain average value, RMS variation, probability of exceeding some threshold, etc.

Random Number Generation

Monte Carlo simulation starts with random number generation, usually split into 2 stages:

- generation of independent uniform (0, 1) random variables
- conversion into random variables with a particular distribution (e.g. Normal)

Very important: never write your own generator, always use a well validated generator from a reputable source

- Matlab
- NAG
- Intel MKI
- AMD ACML



Uniform Random Variables

Pseudo-random number generators use a deterministic (i.e. repeatable) algorithm to generate a sequence of (apparently) random numbers on (0,1) interval.

What defines a good generator?

- a long period how long it takes before the sequence repeats itself 2^{32} is not enough – need at least 2^{40}
- various statistical tests to measure "randomness" well validated software will have gone through these checks
- trivially-parallel Monte Carlo simulation on a compute cluster requires the ability to "skip-ahead" to an arbitrary starting point in the sequence

first computer gets first 10⁶ numbers second computer gets second 10⁶ numbers, etc

Uniform Random Variables

For information see

- Intel MKL information www.intel.com/cd/software/products/asmo-na/eng/266864.htm
- NAG library information www.nag.co.uk/numeric/CL/nagdoc_cl08/pdf/G05/g05_conts.pdf
- Matlab information www.mathworks.com/moler/random.pdf
- Wikipedia information

en.wikipedia.org/wiki/Random_number_generation en.wikipedia.org/wiki/List_of_random_number_generators en.wikipedia.org/wiki/Mersenne_Twister

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Normal Random Variables

N(0,1) Normal random variables (mean 0, variance 1) have the probability distribution

$$p(x) = \phi(x) \equiv \frac{1}{\sqrt{2\pi}} \exp(-\frac{1}{2}x^2)$$

The Box-Muller method takes two independent uniform (0, 1) random numbers y_1, y_2 , and defines

$$\begin{array}{rcl} x_1 & = & \sqrt{-2\log(y_1)} \, \cos(2\pi y_2) \\ x_2 & = & \sqrt{-2\log(y_1)} \, \sin(2\pi y_2) \end{array}$$

It can be proved that x_1 and x_2 are N(0,1) random variables, and independent:

$$p_{\text{joint}}(x_1, x_2) = p(x_1) p(x_2)$$

Inverse CDF

A more flexible alternative uses the cumulative distribution function CDF(x) for a random variable X, defined as

$$CDF(x) = \mathbb{P}(X < x)$$

If Y is a uniform (0,1) random variable, then can define X by

$$X = CDF^{-1}(Y).$$

For N(0,1) Normal random variables,

$$CDF(x) = \Phi(x) \equiv \int_{-\infty}^{x} \phi(s) ds = \frac{1}{\sqrt{2\pi}} \int_{-\infty}^{x} \exp\left(-\frac{1}{2}s^{2}\right) ds$$

 $\Phi^{-1}(y)$ is approximated in software in a very similar way to the implementation of cos, sin, log.



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My GPU CUDA code for inverse error function erfinv

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Normal Random Variables

The Normal CDF $\Phi(x)$ is related to the error function $\operatorname{erf}(x)$ through

$$\Phi(x) = \frac{1}{2} + \frac{1}{2} \operatorname{erf}(x/\sqrt{2}) \implies \Phi^{-1}(y) = \sqrt{2} \operatorname{erf}^{-1}(2y-1)$$

This is the function I use in Matlab:

```
% x = ncfinv(y)
%
% inverse Normal CDF
function x = ncfinv(y)
x = sqrt(2)*erfinv(2*y-1);
```

Correlated Normal Random Variables

We often need a vector y of Normally distributed variables with a prescribed covariance matrix, so that $\mathbb{E}[y y^T] = \Sigma$.

Suppose x is a vector of independent N(0, 1) variables, and define y = Lx. Each element of y is Normally distributed, $\mathbb{E}[y] = L\mathbb{E}[x] = 0$, and

$$\mathbb{E}[y \, y^{\,T}] = \mathbb{E}[L \times x^{\,T} L^{\,T}] = L \, \mathbb{E}[x \, x^{\,T}] \, L^{\,T} = L \, L^{\,T}$$

since $\mathbb{E}[x x^T] = I$ because

- elements of x are independent $\Longrightarrow \mathbb{E}[x_i x_j] = 0$ for $i \neq j$
- elements of x have unit variance $\Longrightarrow \mathbb{E}[x_i^2] = 1$

Hence choose *L* so that $LL^T = \Sigma$. One choice is a Cholesky factorisation in which *L* is lower-triangular. Another is PCA in which $L = U \Lambda^{1/2}$ with *U* the matrix of eigenvectors of Σ , and Λ the diagonal eigenvalue matrix.



Expectation and Integration

If X is a random variable uniformly distributed on [0, 1] then the expectation of a function f(X) is equal to its integral:

$$\overline{f} = \mathbb{E}[f(X)] = I[f] = \int_0^1 f(x) \,\mathrm{d}x.$$

The generalisation to a *d*-dimensional "cube" $I^d = [0, 1]^d$, is

$$\overline{f} = \mathbb{E}[f(X)] = I[f] = \int_{I^d} f(x) \,\mathrm{d}x.$$

Thus the problem of finding expectations is directly connected to the problem of numerical quadrature (integration), often in very large dimensions.

Expectation and Integration

Suppose we have a sequence $X^{(n)}$ of independent samples from the uniform distribution.

An approximation to the expectation/integral is given by

$$I_N[f] = N^{-1} \sum_{n=1}^N f(x^{(n)}).$$

Two key features:

• Unbiased: $\mathbb{E}\left[I_N[f]\right] = I[f]$ • Convergent: $\lim_{N \to \infty} I_N[f] = I[f]$

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Central Limit Theorem

In general, define

- error $\varepsilon_N(f) = I[f] I_N[f]$
- bias = $\mathbb{E}[\varepsilon_N(f)]$
- RMSE, "root-mean-square-error" = $\sqrt{\mathbb{E}[(\varepsilon_N(f))^2]}$

The Central Limit Theorem proves (roughly speaking) that for large N

$$\varepsilon_N(f) \sim \sigma N^{-1/2} Z$$

with Z a N(0,1) random variable and σ^2 the variance of f:

$$\sigma^2 = \mathbb{E}[(f - \overline{f})^2] = \int_{I^d} (f(x) - \overline{f})^2 \, \mathrm{d}x.$$

Central Limit Theorem

More precisely, provided σ is finite, then as $N \longrightarrow \infty$,

$$\mathsf{CDF}(N^{1/2}\sigma^{-1}\varepsilon_N)\longrightarrow \mathsf{CDF}(Z)$$

 $\mathbb{P}\left[N^{1/2}\sigma^{-1}\varepsilon_N < s\right] \longrightarrow \mathbb{P}\left[Z < s\right] = \Phi(s)$

so that

and

$$\mathbb{P}\left[\left|N^{1/2}\sigma^{-1}\varepsilon_{N}\right| > s\right] \longrightarrow \mathbb{P}\left[|Z| > s\right] = 2 \Phi(-s)$$
$$\mathbb{P}\left[\left|N^{1/2}\sigma^{-1}\varepsilon_{N}\right| < s\right] \longrightarrow \mathbb{P}\left[|Z| < s\right] = 1 - 2 \Phi(-s)$$

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Central Limit Theorem

Given N samples, the empirical variance is

$$\widetilde{\sigma}^2 = N^{-1} \sum_{n=1}^{N} \left(f(x^{(n)}) - I_N \right)^2 = I_N^{(2)} - (I_N)^2$$

where

$$I_N = N^{-1} \sum_{n=1}^N f(x^{(n)}), \qquad I_N^{(2)} = N^{-1} \sum_{n=1}^N \left(f(x^{(n)}) \right)^2$$

 $\tilde{\sigma}^2$ is a slightly biased estimator for σ^2 ; an unbiased estimator is

$$\widehat{\sigma}^2 = (N-1)^{-1} \sum_{n=1}^{N} \left(f(x^{(n)}) - I_N \right)^2 = \frac{N}{N-1} \left(I_N^{(2)} - (I_N)^2 \right)$$

Central Limit Theorem

How many samples do we need for an accuracy of $\overline{\varepsilon}$ with probability *c*?

Since

$$\mathbb{P}\left[N^{1/2}\sigma^{-1}|\varepsilon| < s
ight] pprox 1-2 \Phi(-s),$$

define s so that $1 - 2 \Phi(-s) = c$

С	0.683	0.9545	0.9973	0.99994
S	1.0	2.0	3.0	4.0

Then $|\varepsilon| < N^{-1/2} \sigma s$ with probability c, so to get $|\varepsilon| < \overline{\varepsilon}$ we can put

$$N^{-1/2} \widehat{\sigma} s(c) = \overline{\varepsilon} \implies N = \left(\frac{\widehat{\sigma} s(c)}{\overline{\varepsilon}}\right)^2.$$

Note: twice as much accuracy requires 4 times as many samples.

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Expectation and Integration

How does Monte Carlo integration compare to grid based methods for d-dimensional integration?

MC error is proportional to $N^{-1/2}$ independent of the dimension.

If the integrand is sufficiently smooth, trapezoidal integration with $M = N^{1/d}$ points in each direction has

Error
$$\propto M^{-2} = N^{-2/d}$$

This scales better than MC for d < 4, but worse for d > 4. i.e. MC is better at handling high dimensional problems.

Other outputs

Using Monte Carlo we can compute more than just simple averages.

First of all, can compute quantities like the standard deviation:

$$\sigma_f^2 = \mathbb{E}\left[(f - \mathbb{E}[f])^2\right]$$

and other higher moments.

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Other outputs

We can also approximate the cumulative distribution function

$$C(s) = \mathbb{P}[f(X) < s] = \mathbb{E}[\mathbf{1}_{f(X) < s}]$$

or the probability density function, $P(s) = \frac{\mathrm{d}C}{\mathrm{d}s}$, in various ways:

- Maximum Entropy reconstruction uses E[f^m] for m = 1, 2, ..., M to construct PDF approximation P(s) ≈ exp(p(s)), where p(s) is a polynomial and P(s) has the same moments
- Alternatively, can evaluate C(s_j) for a set of values s_j, and then interpolate to approximate the full C(s)

Summary so far

- Monte Carlo quadrature is straightforward and robust
- confidence bounds can be obtained as part of the calculation
- ${\ensuremath{\, \circ }}$ can calculate the number of samples N needed for chosen accuracy
- much more efficient than grid-based methods for high dimensions

• accuracy =
$$O(N^{-1/2})$$
, CPU time = $O(N)$

$$\implies$$
 accuracy = $O(CPU time^{-1/2})$

$$\implies$$
 CPU time = $O(\text{accuracy}^{-2})$

• the key now is to reduce number of samples required by reducing the variance

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