

Numerical Methods I

Monte Carlo Methods

Aleksandar Donev

*Courant Institute, NYU*¹

donev@courant.nyu.edu

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Outline

1 Background

2 Pseudo-Random Numbers

- Inversion Method
- Rejection

3 Monte Carlo Integration

4 Variance Reduction

5 Histogramming

6 Conclusions

What is Monte Carlo?

- Monte Carlo is any numerical algorithm that uses random numbers to compute a deterministic (non-random) answer: **stochastic** or **randomized algorithm**.
- An important example is **numerical integration in higher dimensions**:

$$J = \int_{\Omega \subseteq \mathbb{R}^n} f(\mathbf{x}) d\mathbf{x}$$

- Recall that using a **deterministic method** is very accurate and fast for low dimensions.
- But for large dimensions we have to deal with the **curse of dimensionality**:

The number of quadrature nodes scales like at least 2^n (exponentially). E.g., $2^{20} = 10^6$, but $2^{40} = 10^{12}!$

Probability Theory

- First define a set Ω of possible **outcomes** $\omega \in \Omega$ of an “experiment”:
 - A coin toss can end in heads or tails, so two outcomes.
 - A sequence of four coin tosses can end in one of $4^2 = 16$ outcomes, e.g., HHTT or THTH.
- The set Ω can be finite (heads or tails), countably infinite (the number of atoms inside a box), or uncountable (the weight of a person).
- An **event** $A \subseteq \Omega$ is a **set of possible outcomes**: e.g., more tails than heads occur in a sequence of four coin tosses,

$$A = \{HHHH, THHH, HTHH, HHTH, HHHT\}.$$

- Each event has an associated **probability**

$$0 \leq P(A) \leq 1,$$

with $P(\Omega) = 1$ and $P(\emptyset) = 0$.

Conditional Probability

- A basic axiom is that probability is **additive** for disjoint events:

$$P(A \cup B) = P(A \text{ or } B) = P(A) + P(B) \text{ if } A \cap B = \emptyset$$

- Bayes formula gives the **conditional probability** that an outcome belongs to set B if it belongs to set C :

$$P(B|C) = \frac{P(B \cap C)}{P(C)} = \frac{P(B \text{ and } C)}{P(C)}$$

- Two events are said to be **independent** if their probabilities are multiplicative:

$$P(A \cap B) = P(A \text{ and } B) = P(A)P(B)$$

Probability Distribution

- If Ω is uncountable, think of outcomes as **random variables**, that is, variables whose value is determined by a random outcome:

$$X = X(\omega) \in \mathbb{R}.$$

- The **probability density function** $f(x) \geq 0$ determines the probability for the outcome to be close to x , in one dimension

$$P(x \leq X \leq x + dx) = f(x)dx,$$

$$P(A) = P(X \in A) = \int_{x \in A} f(x)dx$$

- The concept of a **measure** and the **Lebesgue integral** generalizes the traditional Riemann integral in probability theory.

Mean and Variance

- We call the **probability density** or the **probability measure** the **law** or the **distribution** of a random variable X , and write:

$$X \sim f.$$

- The **cummulative distribution function** is

$$F(x) = P(X \leq x) = \int_{-\infty}^x f(x')dx',$$

and we will assume that this function is continuous.

- The **mean** or **expectation value** of a random variable X is

$$\mu = \bar{X} = E[X] = \int_{-\infty}^{\infty} xf(x)dx.$$

- The **variance** σ^2 and the **standard deviation** σ measure the **uncertainty** in a random variable

$$\sigma^2 = \text{var}(X) = E[(X - \mu)^2] = \int_{-\infty}^{\infty} (x - \mu)^2 f(x)dx.$$

Multiple Random Variables

- Consider a set of two random variables $Z = (X, Y)$ and the **joint probability distribution** $Z \sim f(x, y)$.
- The **marginal density** for X is the distribution of just X , without regard to Y :

$$g(x) = \int_y f(x, y) dy, \text{ similarly } h(y) = \int_x f(x, y) dx$$

- The **conditional probability distribution** is the distribution of X for a known Y :

$$f(x|y) = \frac{f(x, y)}{h(y)}$$

- Two random variables X and Y are **independent** if

$$f(x, y) = g(x)h(y) \Rightarrow f(x|y) = g(x).$$

Covariance

- The term **i.i.d. \equiv independent identically-distributed** random variables is used to describe independent **samples** $X_k \sim f$, $k = 1, \dots$
- The generalization of variance for two variables is the **covariance**:

$$C_{XY} = \text{cov}(X, Y) = E[(X - \bar{X})(Y - \bar{Y})] = E(XY) - E(X)E(Y).$$

- For independent variables

$$E(XY) = \int xy f(x, y) dx dy = \int xg(x) dx \int yh(y) dy = E(X)E(Y)$$

and so $C_{XY} = 0$.

- Define the **correlation coefficient** between X and Y as a measure of how correlated two variables are:

$$r_{XY} = \frac{\text{cov}(X, Y)}{\sqrt{\text{var}(X)\text{var}(Y)}} = \frac{C_{XY}}{\sigma_X \sigma_Y}.$$

Law of Large Numbers

- The average of N i.i.d. samples of a random variable $X \sim f$ is itself a random variable:

$$A = \frac{1}{N} \sum_{k=1}^N X_k.$$

- A is an **unbiased estimator** of the mean of X , $E(A) = \bar{X}$.
- Numerically we often use a **biased estimate** of the variance:

$$\sigma_X^2 = \lim_{N \rightarrow \infty} \frac{1}{N} \sum_{k=1}^N (X_k - \bar{X})^2 \approx \frac{1}{N} \sum_{k=1}^N (X_k - A)^2 = \left(\frac{1}{N} \sum_{k=1}^N X_k^2 \right) - A^2$$

- The weak **law of large numbers** states that the estimator is also **consistent**:

$$\lim_{N \rightarrow \infty} A = \bar{X} = E(X) \text{ (almost surely).}$$

Central Limit Theorem

- The central value theorem says that if σ_X is finite, in the limit $N \rightarrow \infty$ the random variable A is **normally-distributed**:

$$A \sim f(a) = (2\pi\sigma_A^2)^{-1/2} \exp\left[-\frac{(a - \bar{X})^2}{2\sigma_A^2}\right]$$

- The **error of the estimator** A decreases as N^{-1} , more specifically,

$$E[(A - \bar{X})^2] = E\left\{\left[\frac{1}{N} \sum_{k=1}^N (X_k - \bar{X})\right]^2\right\} = \frac{1}{N^2} E\left[\sum_{k=1}^N (X_k - \bar{X})^2\right]$$

$$\text{var}(A) = \sigma_A^2 = \frac{\sigma_X^2}{N}.$$

- The slow convergence of the error, $\sigma \sim N^{-1/2}$, is a fundamental characteristic of Monte Carlo.

Monte Carlo on a Computer

- In order to compute integrals using Monte Carlo on a computer, we need to be able to generate samples from a distribution, e.g., uniformly distributed inside an interval $I = [a, b]$.
- Almost all randomized software is based on having a **pseudo-random number generator** (PRNG), which is a routine that returns a pseudo-random number $0 \leq u \leq 1$ from the **standard uniform distribution**:

$$f(u) = \begin{cases} 1 & \text{if } 0 \leq u \leq 1 \\ 0 & \text{otherwise} \end{cases}$$

- Since computers (Turing machines) are deterministic, it is not possible to generate truly random samples (outcomes): Pseudo-random means **as close to random as we can get it**.
- There are well-known good PRNGs that are also efficient: One should **use other-people's PRNGs**, e.g., the **Marsenne Twister**.

PRNGs

- The PRNG is a procedure (function) that takes a collection of m integers called the **state of the generator** $\mathbf{s} = \{i_1, \dots, i_m\}$, and updates it:

$$\mathbf{s} \leftarrow \Phi(\mathbf{s}),$$

and produces (returns) a number $u = \Psi(\mathbf{s})$ that is a pseudo-random sample from the standard uniform distribution.

- So in pseudo-MATLAB notation, $[u, \mathbf{s}] = \text{rng}(\mathbf{s})$, often called a **random stream**.
- Simple built-in generator such as the MATLAB/C function *rand* or the Fortran function *RANDOM_NUMBER* hide the state from the user (but the state is stored somewhere in some global variable).
- All PRNGs provide a routine to **seed the generator**, that is, to set the seed \mathbf{s} to some particular value.

This way one can generate the same sequence of “random” numbers over and over again (e.g., when debugging a program).

Generating Non-Uniform Variates

- Using a uniform (pseudo-)random number generator (**URNG**), it is easy to generate an outcome drawn uniformly in $I = [a, b]$:

$$X = a + (b - a)U,$$

where $U = \text{rng}()$ is a standard uniform variate.

- We often need to generate **(pseudo)random samples** or **variates** drawn from a distribution $f(x)$ other than a uniform distribution, where $f(x) \geq 0$ and $f(x)$ is normalized, $\int f(x)dx = 1$.
- Almost all **non-uniform samplers are based on a URNG**.
- Sometimes it may be more efficient to replace the URNG with a **random bitstream**, that is, a sequence of random bits, if only a few random bits are needed (e.g., for discrete variables).
- We need a method to convert a uniform variate into a non-uniform variate.

Generating Non-Uniform Variates

- Task: We want to sample a random number with **probability distribution** $f(x)$. For now assume $f(x)$ is a **probability density**:

$$P(x \leq X \leq x + dx) = f(x)dx,$$

- Tool: We can generate samples from some special distributions, e.g., a sample U from the standard uniform distribution.
- Consider applying a non-linear **differentiable one-to-one** function $g(x)$ to U :

$$X \equiv X(U) = g(U) \quad \Rightarrow \quad dx = g'(U)du$$

- We can find the probability density of X by using the informal differential notation

$$P(u \leq U \leq u + du) = du = \frac{dx}{g'(u)} = P(x \leq X \leq x + dx) = f(x)dx$$

$$f[x(u)] = [g'(u)]^{-1}$$

Inverting the CDF

$$f[x(u)] = [g'(u)]^{-1}$$

- Can we find $g(u)$ given the target $f(x)$? It is simpler to see this if we invert $x(u)$:

$$u = g^{-1}(x) = F(x).$$

- Repeating the same calculation

$$P(u \leq U \leq u + dx) = du = F'(x)dx = f(x)dx$$

$$F'(x) = f(x)$$

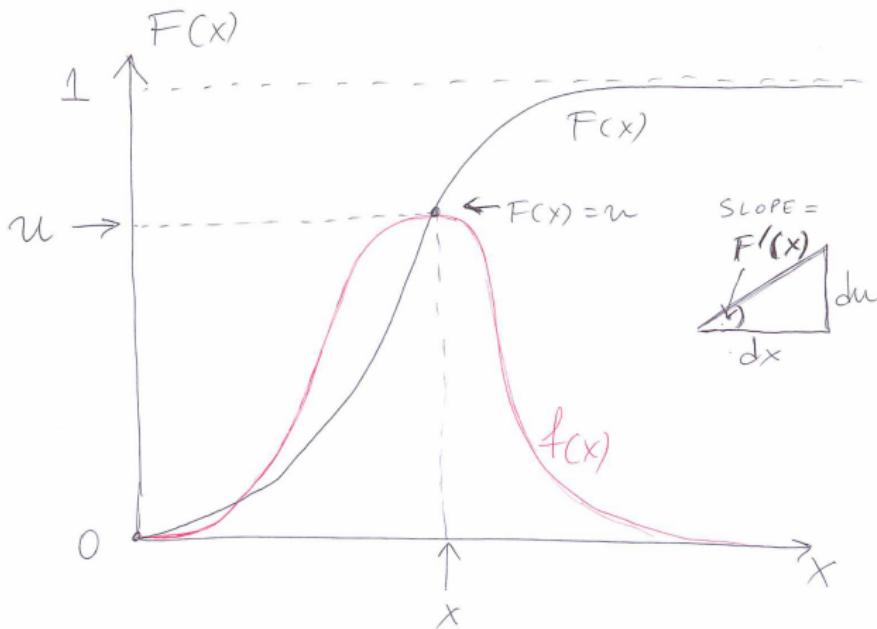
- This shows that $F(x) = g^{-1}(x)$ is the **cummulative probability distribution**:

$$F(x) = P(X \leq x) = \int_{-\infty}^x f(x')dx'.$$

- Note that $F(x)$ is monotonically non-decreasing because $f(x) \geq 0$.

Sampling by Inversion

Inversion algorithm: Generate a standard uniform variate u and then solve the **non-linear equation** $F(x) = u$ to get x .



Exponentially-Distributed Number

- As an example, consider generating a sample from the **exponential distribution with rate λ** :

$$f_{\lambda}(t) = \begin{cases} \lambda e^{-\lambda t} & \text{if } t \geq 0 \\ 0 & \text{otherwise} \end{cases}$$

- Related to the **Poisson process** of events whose rate of occurrence is λ and whose occurrence does not depend on the past (history):

$$P(t \leq T \leq t + dt \mid T \geq t) = P(T < dt) = \lambda dt.$$

- Using the **inversion technique** we get

$$F(t) = P(T \leq t) = \int_{t'=0}^t \lambda e^{-\lambda t'} dt' = 1 - e^{-\lambda t} = u' \equiv 1 - u$$

$$T = -\lambda^{-1} \ln(U),$$

where numerical care must be taken to ensure the log does not overflow or underflow.

Rejection Sampling

- An alternative method is to use **rejection sampling**:
Generate a sample X from some other distribution $g(x)$ and accept them with **acceptance probability** $p(X)$, otherwise **reject** and try again.
- The rejection requires sampling a standard uniform variate U :
Accept if $U \leq p(X)$ and return X , otherwise reject and try a new X .
- What should $p(X)$ be in order for the distribution of X to be $f(x)$?
- Since X and U are **independent** the marginal distribution of x is

$$f(x) \sim g(x)p(x) \quad \Rightarrow \quad p(x) = Z \frac{f(x)}{g(x)},$$

where Z is determined from the **normalization condition**:

$$\int f(x)dx = 1 \quad \Rightarrow \quad \int p(x)g(x)dx = Z$$

Envelope Function

$$p(x) = \frac{f(x)}{Z^{-1}g(x)} = \frac{f(x)}{\tilde{g}(x)}$$

- Since $0 \leq p(x) \leq 1$, we see that $\tilde{g}(x) = Z^{-1}g(x)$ must be a **bounding** or **envelope** function:

$$\tilde{g}(x) = Z^{-1}g(x) \geq f(x), \text{ which implies } Z \leq \frac{g(x)}{f(x)}.$$

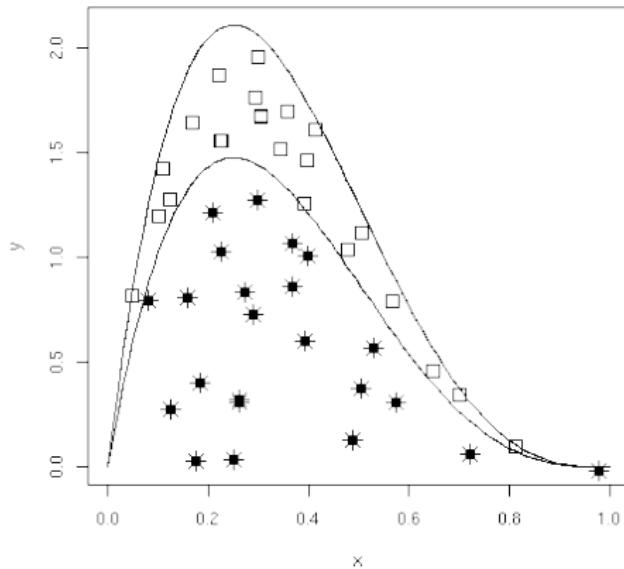
- For efficiency, we want to have the **highest possible acceptance probability**, that is, **maximize**

$$P_{acc} = \frac{\int f(x)dx}{\int \tilde{g}(x)dx} = Z \frac{\int f(x)dx}{\int g(x)dx} = Z \leq 1.$$

- We know $f(x)$ and so once we choose $g(x)$ we can determine $Z = \min \frac{g(x)}{f(x)}$ and thus $\tilde{g}(x) = Z^{-1}g(x)$.

Rejection Sampling Illustrated

Rejection algorithm: Generate a sample X from $g(x)$ and an independent standard uniform variate U . If $U \leq \frac{f(x)}{\tilde{g}(x)}$ accept and return X , otherwise reject and try again.



Normally-Distributed Numbers

- The **standard normal distribution** is a Gaussian “bell-curve”:

$$f(x) = (2\pi\sigma^2)^{-1/2} \exp\left(-\frac{(x-\mu)^2}{2\sigma^2}\right),$$

where μ is the **mean** and σ is the **standard deviation**.

- The **standard normal distribution** has $\sigma = 1$ and $\mu = 0$.
- If we have a sample X_s from the standard distribution we can generate a sample X from $f(x)$ using:

$$X = \mu + \sigma X_s$$

- Consider sampling the positive half of the standard normal, that is, sampling:

$$f(x) = \sqrt{\frac{2}{\pi}} e^{-x^2/2} \text{ for } x \geq 0.$$

If we can do that then we can choose a random \pm sign and sample from the Gaussian distribution.

Optimizing Rejection Sampling

- We want the **tighest possible** (especially where $f(x)$ is large) **easy-to-sample** $g(x) \approx f(x)$.
- Since we want $f(x) < Z^{-1}g(x)$ we must have that $g(x)$ does **not decay faster** than $f(x)$ for large x .
- We already know how to sample an exponential:

$$g(x) = e^{-x}$$

- We want the tightest possible $\tilde{g}(x)$:

$$\min [\tilde{g}(x) - f(x)] = \min \left[Z^{-1}e^{-x} - \sqrt{\frac{2}{\pi}}e^{-x^{2/2}} \right] = 0$$

$$\tilde{g}'(x^*) = f'(x^*) \text{ and } \tilde{g}(x^*) = f(x^*)$$

- Solving this system of two equations gives $x^* = 1$ and

$$Z = P_{acc} = \sqrt{\frac{\pi}{2}}e^{-1/2} \approx 76\%$$

Integration via Monte Carlo

- Define the random variable $Y = f(\mathbf{X})$, and generate a sequence of N **independent uniform samples** $\mathbf{X}_k \in \Omega$, i.e., N random variables distributed uniformly inside Ω :

$$\mathbf{X} \sim g(\mathbf{x}) = \begin{cases} |\Omega|^{-1} & \text{for } \mathbf{x} \in \Omega \\ 0 & \text{otherwise} \end{cases}$$

and calculate the mean

$$\hat{Y} = \frac{1}{N} \sum_{k=1}^N Y_k = \frac{1}{N} \sum_{k=1}^N f(\mathbf{X}_k)$$

- According to the weak law of large numbers,

$$\lim_{N \rightarrow \infty} \hat{Y} = E(Y) = \bar{Y} = \int f(\mathbf{x}) g(\mathbf{x}) d\mathbf{x} = |\Omega|^{-1} \int_{\Omega} f(\mathbf{x}) d\mathbf{x}$$

Accuracy of Monte Carlo Integration

- This gives a Monte Carlo approximation to the integral:

$$J = \int_{\Omega \in \mathbb{R}^n} f(\mathbf{x}) d\mathbf{x} = |\Omega| \bar{Y} \approx |\Omega| \hat{Y} = |\Omega| \frac{1}{N} \sum_{k=1}^N f(\mathbf{x}_k).$$

- Recalling the central limit theorem, for large N we get an **error estimate** by evaluating the standard deviation of the estimate \hat{Y} :

$$\sigma^2(\hat{Y}) \approx \frac{\sigma_Y^2}{N} = N^{-1} \int_{\Omega} \left[f(\mathbf{x}) - |\Omega|^{-1} J \right]^2 d\mathbf{x}$$

$$\sigma(\hat{Y}) \approx \frac{1}{\sqrt{N}} \left[\int_{\Omega} \left[f(\mathbf{x}) - \bar{f}(\mathbf{x}) \right]^2 d\mathbf{x} \right]^{1/2}$$

- Note that this error goes like $N^{-1/2}$, which is order of convergence 1/2: **Worse than any deterministic quadrature.**
- But, the same number of points are needed to get a certain accuracy **independent of the dimension**.

Monte Carlo Error Bars

- Monte Carlo (MC) answers should always be reported with **error bars**, or equivalently, with confidence intervals!
- Since the answer is approximately normally-distributed, we have the well-known **confidence intervals**:

$$P\left(\frac{J}{|\Omega|} \in [\hat{Y} - \sigma, \hat{Y} + \sigma]\right) \approx 66\%$$

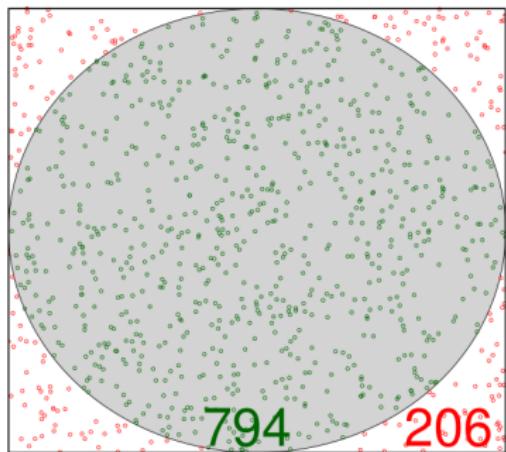
$$P\left(\frac{J}{|\Omega|} \in [\hat{Y} - 2\sigma, \hat{Y} + 2\sigma]\right) \approx 95\%$$

- In practice we estimate the uncertainty **empirically** as

$$\sigma^2(\hat{Y}) \approx \frac{1}{N^2} \sum (Y_i - \bar{Y})^2 = \frac{1}{N} \left[\left(\frac{1}{N} \sum Y_i^2 \right) - \left(\frac{1}{N} \sum Y_i \right)^2 \right].$$

- This is done in a **single MC loop**: Average the Y 's to get the answer but also average the squares Y^2 to get the uncertainty in the answer.

Employing Rejection



Note how this becomes **less efficient as dimension grows** (most points are outside the sphere).

- Integration requires $|\Omega|$, which is hard to compute for complicated domains,

$$\int_{\Omega \in \mathbb{R}^n} f(\mathbf{x}) d\mathbf{x} \approx |\Omega| \frac{1}{N} \sum_{k=1}^N f(\mathbf{X}_k)$$

- Consider Ω being the unit circle of radius 1.
- Rejection: Integrate by sampling points inside an **enclosing region**, e.g, a square of area $|\Omega_{encl}| = 4$, and rejecting any points outside of Ω :

$$\int_{\Omega \in \mathbb{R}^n} f(\mathbf{x}) d\mathbf{x} \approx |\Omega_{encl}| \frac{1}{N} \sum_{\mathbf{X}_k \in \Omega} f(\mathbf{X}_k)$$

Sampling Function

- In the basic MC algorithm described above, the samples \mathbf{X} have a uniform distribution over the integration domain. This does not work if the domain of integration is infinite.
- Instead, we can sample our points from some probability distribution function $g(\mathbf{X}) \geq 0$, $\int g(\mathbf{x}) d\mathbf{x} = 1$, and rewrite:

$$\int f(\mathbf{x}) d\mathbf{x} = \int \frac{f(\mathbf{x})}{g(\mathbf{x})} g(\mathbf{x}) d\mathbf{x} = E \left[\frac{f(\mathbf{X})}{g(\mathbf{X})} \right] \text{ where } \mathbf{X} \sim g.$$

- This now corresponds to taking **samples from the sampling function** $g(\mathbf{x})$:

$$\int f(\mathbf{x}) d\mathbf{x} \approx \frac{1}{N} \sum_{k=1}^N \frac{f(\mathbf{X}_k)}{g(\mathbf{X}_k)} \text{ where } \mathbf{X} \sim g$$

- Note that $|\Omega|$ does not appear since it is implicitly included in the normalization of $g(\mathbf{x})$. The previous uniform sampling algorithm corresponds to $g(\mathbf{x}) = |\Omega|^{-1}$ for $\mathbf{x} \in \Omega$.

Example: Option Pricing

- The fair price of a simple financial option is the expectation of the discounted future value,

$$V = E [e^{-rt} \cdot h(S(t))] ,$$

where $S(t)$ is the stochastic (fluctuating) price of the underlying asset, and r is the risk-free interest rate, and $h(S)$ is the payoff function, for example, $h = \max(S_{\text{strike}} - S, 0)$.

- The price of $S(t)$ is a random process (a random variable that is a function of time), and can be generated either based on a theoretical model.
- For example, a simple theoretical model with volatility ν and average return μ would give

$$S(t) = S(0)e^{\mu t + \nu \sqrt{t} X} ,$$

where X is a normally-distributed random variable.

- So here the natural choice for $g(X)$ is the unit normal distribution, $g(x) = (2\pi)^{-\frac{1}{2}} e^{-x^2/2}$.

Option pricing example

- In the end, combining the pieces together we define

$$\tilde{f}(x) = \frac{f(x)}{g(x)} = e^{-rt} \max \left[S_{\text{strike}} - S(0)e^{\mu t + \nu \sqrt{t}x}, 0 \right]$$

- The option value is simply the expectation value of $\tilde{f}(X)$ when $X \sim g$, i.e., an integral that can be computed using Monte Carlo:

$$V = \int f(x)dx \approx \frac{1}{N} \sum_{k=1}^N \frac{f(\mathbf{X}_k)}{g(\mathbf{X}_k)} = \frac{1}{N} \sum_{k=1}^N \tilde{f}(X_k).$$

- Of course, for a one-dimensional example like this we can probably do this better with a deterministic algorithm.
- But if there are many assets or the payoff function is complicated or if the dynamics of the assets is complex, Monte Carlo may be unavoidable.

Variance Reduction

- The order of convergence (accuracy) of Monte Carlo is always $1/2$ and cannot be improved. Instead, all of the focus is on improving the error constant, i.e., the **variance** for a constant number of samples N .
- The most important thing in Monte Carlo is **variance reduction**, i.e., finding methods that give the same answers in the limit $N \rightarrow \infty$ but have a much smaller σ .
- There are several methods for variance reduction, the most general and powerful of which is **importance sampling**.
- Importance sampling simply means choosing the sampling function $g(x)$ to give more importance to those points that dominate the value of the integral. We call $g(x)$ an **importance sampling function**.

Importance Sampling

- Repeating the variance calculation for

$$Y(\mathbf{X}) = \frac{f(\mathbf{X})}{g(\mathbf{X})}, \text{ with mean } \bar{Y} = \int f(\mathbf{x}) g(\mathbf{x}) d\mathbf{x}$$

- The variance of the empirical mean $\hat{Y} = N^{-1} \sum Y_i$ is

$$\sigma^2(\hat{Y}) \approx \frac{\sigma_Y^2}{N} = N^{-1} \int [Y(\mathbf{x}) - \bar{Y}]^2 g(\mathbf{x}) d\mathbf{x}$$

$$\sigma(\hat{Y}) \approx \sqrt{\frac{1}{N}} \left[\int \left[\frac{f(\mathbf{x})}{g(\mathbf{x})} - \bar{Y} \right]^2 g(\mathbf{x}) d\mathbf{x} \right]^{1/2}.$$

The Importance Function

- We therefore want $f(\mathbf{x})/g(\mathbf{x}) = \bar{Y}$ to be as close as possible to a constant, **ideally**

$$g_{ideal}(\mathbf{x}) = \frac{f(\mathbf{x})}{\int f(\mathbf{x}) d\mathbf{x}}$$

but this requires being able to create independent samples from $f(\mathbf{x})$.

- Instead, what we try to do is to find an importance function that is focused on regions where f is large (the peaks).
- The importance sampling function $g(\mathbf{x})$ must be a probability distribution function that we know how to sample from, such that

$$h(\mathbf{x}) = \frac{f(\mathbf{x})}{g(\mathbf{x})}$$

is **as close to constant as possible**, and in particular, it must be **bounded** from above (i.e., finite for all \mathbf{x} in the relevant domain).

Antithetic Variables

- Consider sampling a random variable $X \sim f$, and assume that f has a **symmetry**, $S(X) \sim f$.
- Examples: For uniform variates, X and $1 - X$, and for normal variates, X and $-X$ both sample the same distribution.
- Variance reduction is achieved if one combines the **antithetic variables** X and $S(X)$ when averaging:

$$\frac{1}{N} \sum_{k=1}^N h(\mathbf{X}_k) \leftarrow \frac{1}{2N} \sum_{k=1}^N [h(\mathbf{X}_k) + h(S(\mathbf{X}_k))].$$

- If there is a **negative correlation** between $h(X)$ and $h(S(X))$, then the variance of the right hand side can be much smaller than the left hand side. This is similar to the use of **control variables** to reduce variance (not covered).
- One example is integrating stochastic differential equations in option pricing: change the sign of the noise term and average the two results.

Quasi-Random Numbers

- If you generate a lot of random samples inside a domain Ω it is easy to observe that there are some regions where there are many points, and others where there are very few points.
- If instead, one uses **quasi-random** or **sub-random** or **low-discrepancy** samples, then one can more uniformly fill the space and get a lower variance.

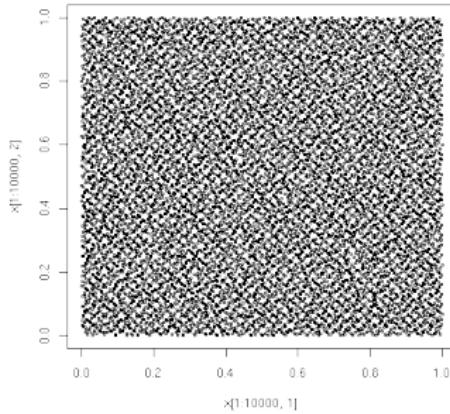
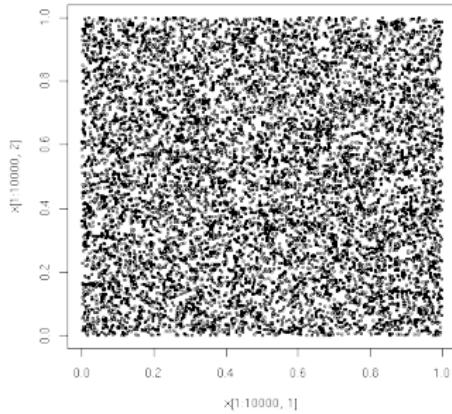
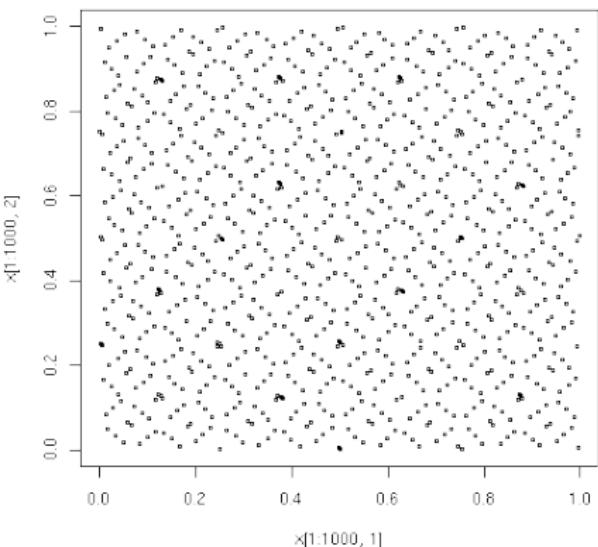
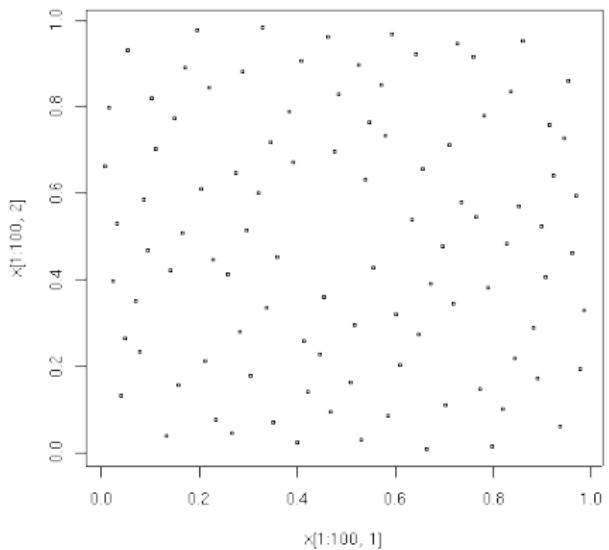


Illustration of the Sobol Sequence



Histogram Validation

- We need some way to test that a sampler is correct, that is, that the generated sequence of random numbers really comes from the specified distribution $f(x)$. One easy way to do that is by computing the **histogram** of the samples.
- Count how many N_x samples of the N samples are inside a **bin of width h** centered at x :

$$f(x) \approx P_x = \frac{1}{h} P(x - h/2 \leq X \leq x + h/2) \approx \frac{N_x}{hN}.$$

- If we make the bins smaller, the **truncation error** will be reduced:

$$P_x - f(x) = \frac{1}{h} \int_{x-h/2}^{x+h/2} f(x') dx' - f(x) = \alpha h^2 + O(h^4)$$

- But, this means there will be fewer points per bin, i.e., **statistical errors** will grow. As usual, we want to find the optimal tradeoff between the the two types of error.

Statistical Error in Histogramming

- For every sample point X , define the **indicator** random variable Y :

$$Y = \mathbb{I}_x(X) = \begin{cases} 1 & \text{if } x - h/2 \leq X \leq x + h/2 \\ 0 & \text{otherwise} \end{cases}$$

- The mean and variance of this **Bernoulli random variable** are:

$$E(Y) = \bar{Y} = hP_x \approx hf(x)$$

$$\sigma_Y^2 = \int (y - \bar{Y})^2 f(y) dy = \bar{Y} \cdot (1 - \bar{Y}) \approx \bar{Y} \approx hf(x)$$

- The number N_x out of N trials inside the bin is a sum of N random Bernoulli variables Y_i :

$$f(x) \approx \frac{1}{h} \frac{N_x}{N} = h^{-1} \left(\frac{1}{N} \sum_{i=1}^N Y_i \right) = \hat{P}_x$$

Optimal Bin Width

- The central limit theorem gives us the uncertainty in our estimate of $f(x)$

$$\sigma(\hat{P}_x) \approx h^{-1} \frac{\sigma_Y}{\sqrt{N}} = \sqrt{\frac{f(x)}{hN}} = \frac{\sqrt{N_x}}{hN}.$$

- This means that the **empirical distribution** $f(x)$ should be reported with a 95% **confidence interval**,

$$P \left\{ f(x) \in \left[\frac{N_x - 2\sqrt{N_x}}{hN}, \frac{N_x + 2\sqrt{N_x}}{hN} \right] \right\} \approx 95\%.$$

- The optimal bin width is when the truncation and statistical errors are equal:

$$\alpha h^2 \approx \sqrt{\frac{f(x)}{hN}} \Rightarrow h \sim N^{-1/5},$$

with total error $\varepsilon \sim (hN)^{-1/2} \sim N^{-2/5}$.

- Typically we choose h based on how well we want to resolve $f(x)$, and accept the fact that **statistical errors dominate**.

Markov Chain Monte Carlo

- Getting **independent samples** from complicated multi-dimensional distributions $f(x)$ is often difficult to do. This is called **static sampling**.
- Instead, it is often easier to construct **dynamic samplers**, which generate a sequence of samples, X_1, X_2, \dots, X_m , which, if scrambled randomly, are essentially independent samples [formally, the marginal distribution of each X_i is $f(x)$].
- But successive samples, for example, X_1 and X_2 , are not independent, and are usually (strongly) correlated. The number of samples N_{corr} over which correlations persist is called the **correlation length**.
- To generate N independent samples, and thus get variance reduction of $N^{-1/2}$, will require generating $N \times N_{corr}$ dynamic samples.

Conclusions/Summary

- Monte Carlo is an umbrella term for **stochastic computation** of deterministic answers.
- Monte Carlo answers are random, and their accuracy is measured by the **variance** or uncertainty of the estimate, which typically scales like $\sigma \sim N^{-1/2}$, where N is the number of **samples**.
- Implementing Monte Carlo algorithms on a computer requires a PRNG, almost always a **uniform pseudo-random number generator** (URNG).
- One often needs to convert a sample from a URNG to a sample from an arbitrary distribution $f(x)$, including inverting the cumulative distribution and rejection sampling.

contd.

- Sampling complex distributions in high-dimensional spaces is often done using **Markov Chain MC**, also called Dynamic or Kinetic MC, which may be based on a model of the actual **dynamics** of a real system (e.g., the market), or simply a computational tool (e.g., Gibbs sampler for Gaussians).
- Monte Carlo can be used to perform **integration in high dimensions** by simply evaluating the function at random points.
- **Variance reduction** is the search for algorithms that give the same answer but with less statistical error. General-purpose variance-reduction techniques include **quasi-random numbers**, **antithetic variables**, **control variates**, and **importance sampling**. They are usually based on some specific knowledge about the model.