Monte Carlo Methods

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	arlo methods for POMP models
	or general distributions
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Objectives	
1. To review some basic	ideas in Monte Carlo computation and simulating random variables.
-	troduction to the Monte Carlo approach, and the generation of simulated random ho haven't seen it before.

Our context: Monte Carlo methods for POMP models.

Let's consider a general POMP model. As before, let $y_{1:N}^*$ be the data, and let the model consist of a latent process $X_{0:N}$ and an observable process $Y_{1:N}$. Then the likelihood function is

$$\mathcal{L}(\theta) = f_{Y_{1:N}}(y_{1:N}^*; \theta) \tag{1}$$

$$= \int_{x_0} \cdots \int_{x_N} f_{X_0}(x_0; \theta) \prod_{n=1}^N f_{Y_n|X_n}(y_n^* | x_n; \theta) f_{X_n|X_{n-1}}(x_n | x_{n-1}; \theta) dx_0 \dots dx_N.$$
 (2)

i.e., computation of the likelihood requires integrating (or summing, for a discrete model) over all possible values of the unobserved latent process at each time point. This is very hard to do, in general.

Let's review, and/or learn, some *Monte Carlo* approaches for evaluating this and other difficult integrals. An excellent technical reference on Monte Carlo techniques is @robert04.

The fundamental theorem of Monte Carlo integration

• The basic insight of Monte Carlo methods is that we can get a numerical approximation to a challenging integral,

$$H = \int h(x) f(x) dx,$$

if we can simulate (i.e., generate random draws) from the distribution with probability density function f.

• This insight is known as the fundamental theorem of Monte Carlo integration.

Theorem. Let f(x) be the probability distribution function for a random variable X, and let $X_{1:J} = \{X_j, j = 1, ..., J\}$ be an independent and identically distributed sample of size J from f. Let H_J be the sample average of $h(X_1), ..., h(X_J)$,

$$H_J = \frac{1}{J} \sum_{j=1}^{J} h(X_j).$$

Then H_J converges to H as $J \to \infty$ with probability 1. Less formally, we write

$$H_J \approx \int h(x) f(x) dx.$$

Proof. This is the strong law of large numbers, together with the identity that

$$\mathbb{E}[h(X)] = \int h(x) f(x) dx.$$

• We can estimate the error in this approximation, because the empirical variance

$$V_J = \frac{1}{J-1} \sum_{j=1}^{J} [h(X_j) - H_J]^2$$

approximates the true variance, $\operatorname{Var}[h(X)] = \mathbb{E}[(h(X) - \mathbb{E}[h(X)])^2]$.

• The standard error on the approximation $H_J \approx \mathbb{E}[h(X)]$ is therefore

$$\sqrt{\frac{V_J}{J}}$$
.

• From the central limit theorem, the error is approximately normally distributed:

$$H_J - \mathbb{E}[h(X)] \sim \operatorname{normal}\left(0, \frac{V_J}{J}\right).$$

- The fundamental theorem of Monte Carlo inspires us to give further thought on how to simulate from a desired density function f, which may itself be a challenging problem.
- We will review simulation, but first let's consider a useful generalization of the fundamental theorem.

Importance sampling

- Sometimes it is difficult to sample directly from the distribution of X.
- In this case, we can often make use of *importance sampling*, in which we generate random samples from another distribution (easier to simulate) and make the appropriate correction.

- Specifically, suppose we wish to compute $\mathbb{E}[h(X)]$, where $X \sim f$, but it is difficult or impossible to draw random samples from f.
- Suppose g is a probability distribution from which it's relatively easy to draw samples and let $Y_{1:J}$ be i.i.d. random variables drawn from g.
- Notice that

$$\mathbb{E}[h(X)] = \int h(x) f(x) dx = \int h(x) \frac{f(x)}{g(x)} g(x) dx.$$

• So, we can generalize the Monte Carlo integration theorem to give the **Monte Carlo importance** sampling theorem,

$$\mathbb{E}[h(X)] \approx \frac{1}{J} \sum_{j=1}^{J} h(Y_j) \frac{f(Y_j)}{g(Y_j)}.$$

• We call $w_j = f(Y_j)/g(Y_j)$ the **importance weights**, and then we can write

$$\mathbb{E}[h(X)] \approx \frac{1}{J} \sum_{j=1}^{J} w_j h(Y_j).$$

• Since $\mathbb{E}[w_j] = \mathbb{E}[f(Y)/g(Y)] = 1$, we can modify this formula to give a **self-normalized importance** sampling estimate,

$$\mathbb{E}[h(X)] \approx \frac{\sum w_j h(Y_j)}{\sum w_j}.$$

- The self-normalized estimate requires computation of w_i only up to a constant of proportionality.
- The Monte Carlo variance associated with this estimate is

$$\frac{\sum w_j (h(Y_j) - \overline{h})^2}{\sum w_j}.$$

• Obtaining accurate estimates requires some thought to the importance distribution g. Specifically, if the tails of g are lighter than those of f, the Monte Carlo variance will be inflated and the estimates can be unusable.

Simulation techniques for general distributions

- Simulation refers to the generation of random variables.
- The general problem of simulation is: given a probability distribution f, find a procedure that generates random draws from f.
- This is a very important problem in scientific computing and much thought and effort has gone into producing reliable simulators for many basic random variables.
 - There are two basic ways of solving this problem:
 - 1. The transformation method,
 - 2. The rejection method.

The transformation method

- This method works for discrete or continuous scalar random variables.
- Let f be the probability distribution function we seek to draw from (known as the **target distribution**) and F be the corresponding cumulative distribution function, i.e., $F(x) = \int_{-\infty}^{x} f(v) dv$.
- Let $F^{-1}(u) = \inf\{x : F(x) \ge u\}$ be the inverse of F.
- A basic fact is that, if $X \sim f$, then $F(X) \sim \text{uniform}(0,1)$. Proof: If f(X) > 0, then

$$\mathbb{P}[F(X) \le u] = \mathbb{P}[X < F^{-1}(u)]
= F(F^{-1}(u)) = u.$$
(3)

- This suggests that, if we can compute F^{-1} , we use the following algorithm to generate $X \sim f$:
- 1. draw $U \sim \text{uniform}(0, 1)$.
- 2. let $X = F^{-1}(U)$.

The rejection method

- The transformation method is very efficient in that we are guaranteed to obtain a valid X from the density f for every $U \sim \text{uniform}(0,1)$ we generate.
- Sometimes, however, we cannot compute the inverse of the cumulative distribution function, as required by the transformation method.
- Under such circumstances, the rejection method offers a less efficient, but more flexible, alternative.
- We'll see how and why this method works.

The rejection method for uniform random variables on arbitrary sets

- Let a random variable X take values in $\mathbb{R}^{\dim(X)}$.
- Suppose that X is **uniformly distributed** over a region $D \subset \mathbb{R}^{\dim(X)}$. This means that, for any $A \subset D$,

$$\mathbb{P}[X \in A] = \frac{\operatorname{area}(A)}{\operatorname{area}(D)}.$$

We write

$$X \sim \text{uniform}(D)$$
.

- Let's suppose that we wish to simulate $X \sim \text{uniform}(D)$.
- Suppose that we don't know how to directly simulate a random draw from D, but we know D is a subset of some nicer region $U \subset \mathbb{R}^{\dim(X)}$.
- If we know how to generate $Y \sim \text{uniform}(U)$, then we can simply do so until $Y \in D$, at which point we take X = Y.

• Since for any $A \subset D$,

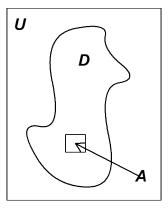
$$\mathbb{P}[X \in A] = \mathbb{P}[Y \in A | Y \in D] \tag{5}$$

$$= \frac{\operatorname{area}(A)}{\operatorname{area}(U)} / \frac{\operatorname{area}(D)}{\operatorname{area}(U)}$$
(6)

$$= \frac{\operatorname{area}(A)}{\operatorname{area}(D)},\tag{7}$$

it follows that $Y \sim \text{uniform}(D)$.

• Consider an analogy to throwing darts. If the darts are thrown in such a way as to be equally likely to land anywhere in U, then those that do land in D are equally likely to land anywhere in D.



The rejection method for dominated densities

- A useful little fact allows us to extend the rejection method from uniform distributions to arbitrary densities.
- @robert04 refer to this fact the fundamental theorem of simulation.
- Let h be an arbitary positive, integrable function.
- Define

$$D = \{(x, u) : 0 \le u \le h(x)\},\$$

i.e., D is the graph of h.

- Consider the random pair $(X, U) \sim \text{uniform}(D)$.
- What is the marginal distribution of X?

$$\int_0^{h(x)} \mathrm{d}u = h(x)$$

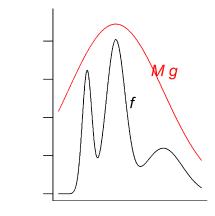
• So h is the probability distribution function for X!

• To carry out the rejection method, simulating from g to obtain a sample from f, we take our region D to be the **graph** of Mg, i.e.,

$$D = \{(x, y) : 0 \le y \le Mg(x),$$

where M is a constant chosen so that $Mg(x) \leq f(x)$ for all x.

- We propose points (X,Y) by drawing them uniformly from the area under the graph of Mg.
- We accept the point (X,Y) if it lies under the graph of f.
- Then, the X-component of the (X,Y) pair is distributed according to f.
- This suggests the following rejection method for simulating an arbitrary random variable.
- Let f be the target distribution and g be another distribution function (from which it is easy to simulate) (see Figure below).
- Let M be such that $M g(x) \ge f(x)$ for all x.
- The following procedure simulates $X \sim f$.
- 1. draw $Y \sim g$ and $U \sim \text{uniform}(0, M g(Y))$.
- 2. if $U \le f(Y)$, then let X = Y else repeat step 1.



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