STAT 801: Mathematical Statistics

Monte Carlo

Suppose you are given random variables X_1, \ldots, X_n whose joint density f (or distribution) is specified and a statistic $T(X_1, \ldots, X_n)$ whose distribution you want to know. To compute something like P(T > t) you can proceed as follows:

- 1. Generate X_1, \ldots, X_n from the density f.
- 2. Compute $T_1 = T(X_1, ..., X_n)$.
- 3. Repeat steps 1 and 2 N times getting T_1, \ldots, T_N .
- 4. Estimate p = P(T > t) as $\hat{p} = M/N$ where M is number of repetitions where $T_i > t$.
- 5. Estimate the accuracy of \hat{p} using $\sqrt{\hat{p}(1-\hat{p})/N}$.

Notice that the accuracy is inversely proportional to \sqrt{N} . There are a number of tricks to make the method more accurate (but they only change the constant of proportionality – the SE is still inversely proportional to the square root of the sample size).

In the rest of this section I begin by discussing techniques for generating random variables with a given distribution under the assumption that you have a source of independent uniformly distributed variables. Then I survey a variety of methods for improving the accuracy of Monte Carlo methods.

Generating the Sample

We will consider here several methods for converting a source of indepedent Uniform [0,1] random variables to random variables with another desired joint distribution. Most computer languages have a facility for generating pseudo uniform random numbers, that is, variables U which have (approximately of course) a Uniform [0,1] distribution. They are produced by a deterministic algorithm so they are not really random (that's why we say "pseudo") but the algorithms are tested to check that they have approximately the properties of such a sequence. Of course numbers stored in a computer are recorded only to a certain number of bits so they are really discrete uniforms not continuous uniforms. Sometimes this matters in programming.

The methods we will consider here are:

- 1. Transformation
- 2. Acceptance-Rejection sampling
- 3. Markov Chain Monte Carlo

Transformation

Other distributions are generated by transformation:

Exponential: $X = -\log U$ has an exponential distribution:

$$P(X > x) = P(-\log(U) > x)$$
$$= P(U \le e^{-x}) = e^{-x}$$

Random uniforms generated on the computer sometimes have only 6 or 7 digits or so of detail. This can make the tail of your distribution grainy. If U were actually a multiple of 10^{-6} for instance then the largest possible value of X is $6 \log(10)$. This problem can be ameliorated by the following algorithm:

- Generate U a Uniform [0,1] variable.
- Pick a small ϵ like 10^{-3} say. If $U > \epsilon$ take $Y = -\log(U)$.
- If $U \leq \epsilon$ remember that the conditional distribution of Y y given Y > y is exponential. You use this by generating a new U' and computing $Y' = -\log(U')$. Then take $Y = Y' \log(\epsilon)$. The resulting Y has an exponential distribution. You should check this by computing P(Y > y).

General technique: inverse probability integral transform.

If X is to have cdf F: Generate $U \sim Uniform[0, 1]$. Take $X = F^{-1}(U)$:

$$P(Y \le y) = P(F^{-1}(U) \le y)$$
$$= P(U \le F(y)) = F(y)$$

Example: X exponential. $F(x) = 1 - e^{-x}$ and $F^{-1}(u) = -\log(1-u)$. Compare to previous method. (Use U instead of 1 - U.) **Normal:** $F = \Phi$ (common notation for standard normal cdf). No closed form for F^{-1} . One way: use numerical algorithm to compute F^{-1} . Alternative: Box Müller Generate U_1, U_2 two independent Uniform[0,1] variables. Define $Y_1 = \sqrt{-2\log(U_1)}\cos(2\pi U_2)$

$$Y_2 = \sqrt{-2\log(U_1)}\sin(2\pi U_2)$$

Exercise: (use change of variables) Y_1 and Y_2 are independent N(0,1) variables.

Acceptance Rejection

Suppose: can't calculate F^{-1} but know f. Find density g and constant c such that

- 1. $f(x) \leq cg(x)$ for each x and
- 2. G^{-1} is computable or can generate observations W_1, W_2, \ldots independently from g.

Algorithm:

- 1. Generate W_1 .
- 2. Compute $p = f(W_1)/(cg(W_1)) \le 1$.
- 3. Generate uniform [0,1] random variable U_1 independent of all Ws.
- 4. Let $Y = W_1$ if $U_1 \le p$.
- 5. Otherwise get new W, U; repeat until you find $U_i \leq f(W_i)/(cg(W_i))$.
- 6. Make Y be last W generated.

This Y has density f.

Markov Chain Monte Carlo

Recently popular tactic, particularly for generating multivariate observations.

Theorem Suppose W_1, W_2, \ldots is an (ergodic) Markov chain with stationary transitions and the stationary initial distribution of W has density f. Then starting the chain with *any* initial distribution

$$\frac{1}{n}\sum_{i=1}^n g(W_i) \to \int g(x)f(x)dx \,.$$

Estimate things like $\int_A f(x) dx$ by computing the fraction of the W_i which land in A.

Many versions of this technique including Gibbs Sampling and Metropolis-Hastings algorithm. The technique was invented in 1950s by physicists in a paper by Metropolis et al. One of the authors was Edward Teller the so-called "father of the hydrogen bomb".

Importance Sampling

If you want to compute

$$\theta \equiv E(T(X)) = \int T(x)f(x)dx$$

you can generate observations from a different density g and then compute

$$\hat{\theta} = n^{-1} \sum T(X_i) f(X_i) / g(X_i)$$

Then

$$E(\hat{\theta}) = n^{-1} \sum E \{T(X_i)f(X_i)/g(X_i)\}$$
$$= \int \{T(x)f(x)/g(x)\}g(x)dx$$
$$= \int T(x)f(x)dx$$
$$= \theta$$

Variance reduction

Consider the problem of estimating the distribution of the sample mean for a Cauchy random variable. The Cauchy density is

$$f(x) = \frac{1}{\pi(1+x^2)}$$

We generate U_1, \ldots, U_n uniforms and then define $X_i = \tan^{-1}(\pi(U_i - 1/2))$. Then we compute $T = \overline{X}$. Now to estimate p = P(T > t) we would use

$$\hat{p} = \sum_{i=1}^N \mathbbm{1}(T_i > t) / N$$

after generating N samples of size n. This estimate is unbiased and has standard error $\sqrt{p(1-p)/N}$.

We can improve this estimate by remembering that $-X_i$ also has Cauchy distribution. Take $S_i = -T_i$. Remember that S_i has the same distribution as T_i . Then we try (for t > 0)

$$\tilde{p} = \left[\sum_{i=1}^{N} 1(T_i > t) + \sum_{i=1}^{N} 1(S_i > t)\right]/(2N)$$

which is the average of two estimates like \hat{p} . The variance of \tilde{p} is

$$(4N)^{-1}Var(1(T_i > t) + 1(S_i > t)) = (4N)^{-1}Var(1(|T| > t))$$

which is

$$\frac{2p(1-2p)}{4N} = \frac{p(1-2p)}{2N}$$

Notice that the variance has an extra 2 in the denominator and that the numerator is also smaller – particularly for p near 1/2. So this method of variance reduction has resulted in a need for only half the sample size to get the same accuracy.

Regression estimates

Suppose we want to compute

$$\theta = E(|Z|)$$

where Z is standard normal. We generate N iid N(0,1) variables Z_1, \ldots, Z_N and compute $\hat{\theta} = \sum |Z_i|/N$. But we know that $E(Z_i^2) = 1$ and can see easily that $\hat{\theta}$ is positively correlated with $\sum Z_i^2/N$. So we consider using

$$\tilde{\theta} = \hat{\theta} - c(\sum Z_i^2/N - 1)$$

Notice that $E(\tilde{\theta}) = \theta$ and

$$Var(\hat{\theta}) = Var(\hat{\theta}) - 2c \text{Cov}(\hat{\theta}, \sum Z_i^2/n) + c^2 Var(\sum Z_i^2/N)$$

The value of c which minimizes this is

$$c = \frac{\operatorname{Cov}(\hat{\theta}, \sum Z_i^2/n)}{\operatorname{Var}(\sum Z_i^2/N)}$$

and this value can be estimated by regressing the $|Z_i|$ on the Z_i^2 !