7.2 Transformation Method: Exponential and Normal Deviates

In the previous section, we learned how to generate random deviates with a uniform probability distribution, so that the probability of generating a number between x and x + dx, denoted p(x)dx, is given by

$$p(x)dx = \begin{cases} dx & 0 < x < 1\\ 0 & \text{otherwise} \end{cases}$$
 (7.2.1)

The probability distribution p(x) is of course normalized, so that

$$\int_{-\infty}^{\infty} p(x)dx = 1 \tag{7.2.2}$$

Now suppose that we generate a uniform deviate x and then take some prescribed function of it, y(x). The probability distribution of y, denoted p(y)dy, is determined by the fundamental transformation law of probabilities, which is simply

$$|p(y)dy| = |p(x)dx| \tag{7.2.3}$$

or

$$p(y) = p(x) \left| \frac{dx}{dy} \right| \tag{7.2.4}$$

Exponential Deviates

As an example, suppose that $y(x) \equiv -\ln(x)$, and that p(x) is as given by equation (7.2.1) for a uniform deviate. Then

$$p(y)dy = \left| \frac{dx}{dy} \right| dy = e^{-y} dy \tag{7.2.5}$$

which is distributed exponentially. This exponential distribution occurs frequently in real problems, usually as the distribution of waiting times between independent Poisson-random events, for example the radioactive decay of nuclei. You can also easily see (from 7.2.4) that the quantity y/λ has the probability distribution $\lambda e^{-\lambda y}$.

So we have

```
#include <math.h>
float expdev(long *idum)
Returns an exponentially distributed, positive, random deviate of unit mean, using
ran1(idum) as the source of uniform deviates.
{
    float ran1(long *idum);
    float dum;

    do
        dum=ran1(idum);
    while (dum == 0.0);
    return -log(dum);
```

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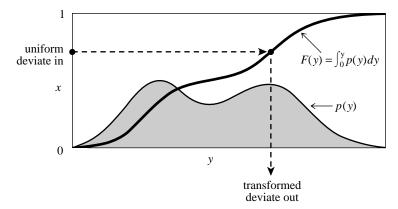


Figure 7.2.1. Transformation method for generating a random deviate y from a known probability distribution p(y). The indefinite integral of p(y) must be known and invertible. A uniform deviate x is chosen between 0 and 1. Its corresponding y on the definite-integral curve is the desired deviate.

Let's see what is involved in using the above *transformation method* to generate some arbitrary desired distribution of y's, say one with p(y) = f(y) for some positive function f whose integral is 1. (See Figure 7.2.1.) According to (7.2.4), we need to solve the differential equation

$$\frac{dx}{dy} = f(y) \tag{7.2.6}$$

But the solution of this is just x = F(y), where F(y) is the indefinite integral of f(y). The desired transformation which takes a uniform deviate into one distributed as f(y) is therefore

$$y(x) = F^{-1}(x) (7.2.7)$$

where F^{-1} is the inverse function to F. Whether (7.2.7) is feasible to implement depends on whether the *inverse function of the integral of* f(y) is itself feasible to compute, either analytically or numerically. Sometimes it is, and sometimes it isn't.

Incidentally, (7.2.7) has an immediate geometric interpretation: Since F(y) is the area under the probability curve to the left of y, (7.2.7) is just the prescription: choose a uniform random x, then find the value y that has that fraction x of probability area to its left, and return the value y.

Normal (Gaussian) Deviates

Transformation methods generalize to more than one dimension. If x_1, x_2, \ldots are random deviates with a *joint* probability distribution $p(x_1, x_2, \ldots)$ $dx_1 dx_2 \ldots$, and if y_1, y_2, \ldots are each functions of all the x's (same number of y's as x's), then the joint probability distribution of the y's is

$$p(y_1, y_2, ...) dy_1 dy_2 ... = p(x_1, x_2, ...) \left| \frac{\partial (x_1, x_2, ...)}{\partial (y_1, y_2, ...)} \right| dy_1 dy_2 ...$$
 (7.2.8)

where $|\partial(-)/\partial(-)|$ is the Jacobian determinant of the x's with respect to the y's (or reciprocal of the Jacobian determinant of the y's with respect to the x's).

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An important example of the use of (7.2.8) is the *Box-Muller* method for generating random deviates with a normal (Gaussian) distribution,

$$p(y)dy = \frac{1}{\sqrt{2\pi}}e^{-y^2/2}dy$$
 (7.2.9)

Consider the transformation between two uniform deviates on (0,1), x_1, x_2 , and two quantities y_1, y_2 ,

$$y_1 = \sqrt{-2 \ln x_1} \cos 2\pi x_2$$

$$y_2 = \sqrt{-2 \ln x_1} \sin 2\pi x_2$$
(7.2.10)

Equivalently we can write

$$x_{1} = \exp\left[-\frac{1}{2}(y_{1}^{2} + y_{2}^{2})\right]$$

$$x_{2} = \frac{1}{2\pi}\arctan\frac{y_{2}}{y_{1}}$$
(7.2.11)

Now the Jacobian determinant can readily be calculated (try it!):

$$\frac{\partial(x_1, x_2)}{\partial(y_1, y_2)} = \begin{vmatrix} \frac{\partial x_1}{\partial y_1} & \frac{\partial x_1}{\partial y_2} \\ \frac{\partial x_2}{\partial y_1} & \frac{\partial x_2}{\partial y_2} \end{vmatrix} = -\left[\frac{1}{\sqrt{2\pi}}e^{-y_1^2/2}\right] \left[\frac{1}{\sqrt{2\pi}}e^{-y_2^2/2}\right]$$
(7.2.12)

Since this is the product of a function of y_2 alone and a function of y_1 alone, we see that each y is independently distributed according to the normal distribution (7.2.9).

One further trick is useful in applying (7.2.10). Suppose that, instead of picking uniform deviates x_1 and x_2 in the unit square, we instead pick v_1 and v_2 as the ordinate and abscissa of a random point inside the unit circle around the origin. Then the sum of their squares, $R^2 \equiv v_1^2 + v_2^2$ is a uniform deviate, which can be used for x_1 , while the angle that (v_1, v_2) defines with respect to the v_1 axis can serve as the random angle $2\pi x_2$. What's the advantage? It's that the cosine and sine in (7.2.10) can now be written as $v_1/\sqrt{R^2}$ and $v_2/\sqrt{R^2}$, obviating the trigonometric function calls!

We thus have

rsq=v1*v1+v2*v2;

```
#include <math.h>
```

see if they are in the unit circle,

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```
} while (rsq >= 1.0 \mid \mid rsq == 0.0);
                                                        and if they are not, try again.
        fac=sqrt(-2.0*log(rsq)/rsq);
        Now make the Box-Muller transformation to get two normal deviates. Return one and
        save the other for next time.
        gset=v1*fac;
         iset=1;
                                                Set flag.
        return v2*fac;
    } else {
                                                We have an extra deviate handy,
        iset=0;
                                                so unset the flag,
        return gset;
                                                and return it
}
```

See Devroye [1] and Bratley [2] for many additional algorithms.

CITED REFERENCES AND FURTHER READING:

Devroye, L. 1986, *Non-Uniform Random Variate Generation* (New York: Springer-Verlag), §9.1. [1]

Bratley, P., Fox, B.L., and Schrage, E.L. 1983, *A Guide to Simulation* (New York: Springer-Verlag). [2]

Knuth, D.E. 1981, Seminumerical Algorithms, 2nd ed., vol. 2 of The Art of Computer Programming (Reading, MA: Addison-Wesley), pp. 116ff.

7.3 Rejection Method: Gamma, Poisson, Binomial Deviates

The rejection method is a powerful, general technique for generating random deviates whose distribution function p(x)dx (probability of a value occurring between x and x+dx) is known and computable. The rejection method does not require that the cumulative distribution function [indefinite integral of p(x)] be readily computable, much less the inverse of that function — which was required for the transformation method in the previous section.

The rejection method is based on a simple geometrical argument:

Draw a graph of the probability distribution p(x) that you wish to generate, so that the area under the curve in any range of x corresponds to the desired probability of generating an x in that range. If we had some way of choosing a random point *in two dimensions*, with uniform probability in the *area* under your curve, then the x value of that random point would have the desired distribution.

Now, on the same graph, draw any other curve f(x) which has finite (not infinite) area and lies everywhere *above* your original probability distribution. (This is always possible, because your original curve encloses only unit area, by definition of probability.) We will call this f(x) the *comparison function*. Imagine now that you have some way of choosing a random point in two dimensions that is uniform in the area under the comparison function. Whenever that point lies outside the area under the original probability distribution, we will *reject* it and choose another random point. Whenever it lies inside the area under the original probability distribution, we will *accept* it. It should be obvious that the accepted points are uniform in the accepted area, so that their x values have the desired distribution. It

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