

## Random Numbers and Simulation

- **Generating random numbers:** Typically impossible/unfeasible to obtain truly random numbers
- Programs have been developed to generate pseudo-random numbers:
- Values generated from a complicated deterministic algorithm, which can pass any *statistical test* for randomness
- They *appear* to be independent and identically distributed.
- Random number generators for common distributions are built into R.
- For less common distributions, more complicated methods have been developed (e.g., Accept-Reject Sampling, Metropolis-Hastings Algorithm)
- STAT 740 covers these.

## **(Monte Carlo) Simulation**

### *Some Common Uses of Simulation*

1. Optimization (Example: Finding MLEs)
2. Calculating Definite Integrals (Ex: Finding Posterior Distributions)
3. Approximating the Sampling Distribution of a Statistic (Ex: Constructing CIs)

- 1. Finding the  $x$  that maximizes (or minimizes) a complicated function  $h(x)$  can be difficult analytically
- Situation even tougher if  $\mathbf{x}$  is multidimensional
- Find  $\mathbf{x}$  to maximize  $h(x_1, x_2, \dots, x_p)$

## OTHER OPTIONS:

- Simple Stochastic Search: If the maximum is to take place over a bounded region, say  $[0, 1]^p$ , then:  
Generate many uniform random observations in that region, plug each into  $h(\cdot)$ , and pick the one that gives the largest  $h(\mathbf{x})$ .
- *Advantage*: Easy to program.
- *Disadvantage*: Very slow, especially for multidimensional problems. Requires much computation.

*Example*: Maximize  $h(x_1, x_2) = (x_1^2 + 4x_2^2)e^{1-x_1^2-x_2^2}$  over  $[-3, 3]^2$ .

*More advanced: Gradient Methods*, which use derivative information to determine which area of the region to search next.

- Rule: “go up the slope”
- Disadvantage: Can get stuck on *local* maxima

*Simulated Annealing*: Tries a sequence of  $\mathbf{x}$  values:  $\mathbf{x}_0, \mathbf{x}_1, \dots$

- If  $h(\mathbf{x}_{i+1}) \geq h(\mathbf{x}_i)$ , “move” to  $\mathbf{x}_{i+1}$ .
- If  $h(\mathbf{x}_{i+1}) < h(\mathbf{x}_i)$ , “move” to  $\mathbf{x}_{i+1}$  with a certain probability which depends on  $h(\mathbf{x}_{i+1}) - h(\mathbf{x}_i)$ .

## R functions that perform optimization

```
optim()
```

```
optimize() ← one-dimensional optimization
```

```
Example: optim(par = c(0,0), fn=my.fcn,  
control=list(fnscale=-1), maxit=100000)
```

```
# Nelder-Mead optimization
```

```
Other choices: method="CG", method="BFGS", method="SANN"
```

## Calculating Definite Integrals

In statistics, we often have to calculate difficult definite integrals (Posterior distributions, expected values)

$$I = \int_a^b h(x) dx$$

(here,  $\mathbf{x}$  could be multidimensional)

**Example 1:** Find:

$$\int_0^1 \frac{4}{1+x^2} dx$$

**Example 2:** Find:

$$\int_0^1 \int_0^1 (4 - x_1^2 - 2x_2^2) dx_2 dx_1$$

## Hit-or-Miss Method

### Example 1:

$$h(x) = \frac{4}{1 + x^2}$$

- Determine  $c$  such that  $c \geq h(x)$  across entire region of interest. (Here,  $c = 4$ )
- Generate  $n$  random uniform  $(X_i, Y_i)$  pairs,  $X_i$ 's from  $U[a, b]$  (here,  $U[0, 1]$ ) and  $Y_i$ 's from  $U[0, c]$  (here,  $U[0, 4]$ )
- Count the number of times (call this  $m$ ) that the  $Y_i$  is less than the  $h(X_i)$
- Then  $I \approx c(b - a) \frac{m}{n}$

[ This is (height)(width)(proportion in shaded region) ]



## Classical Monte Carlo Integration

$$I = \int_a^b h(x) dx$$

- Take  $n$  random uniform values  $U_1, \dots, U_n$  (could be vectors) over  $[a, b]$

Then

$$I \approx \frac{b-a}{n} \sum_{i=1}^n h(U_i)$$

## Expected Value of a Function of a Random Variable

Suppose  $X$  is a random variable with density  $f$ .

Find  $E[h(X)]$  for some function  $h$ , e.g.,

$$E[X^2]$$

$$E[\sqrt{X}]$$

$$E[\sin(X)]$$

- Note  $E[h(X)] = \int_{\mathcal{X}} h(x) f(x) dx$  over whatever the support of  $f$  is.
- Take  $n$  random values  $X_1, \dots, X_n$  from the distribution of  $X$  (i.e., with density  $f$ )
- Then

$$E[h(X)] \approx \frac{1}{n} \sum_{i=1}^n h(X_i)$$

## Examples

**Example 3:** If  $X$  is a random variable with a  $N(10, 1)$  distribution, find  $E(X^2)$ .

**Example 4:** If  $Y$  is a beta random variable with parameters  $a = 5$  and  $b = 1$ , find  $E(-\log_e Y)$ .

- Some more advanced methods of integration using simulation (Importance Sampling)
- Note: R function `integrate()` does numerical integration for functions of a *single* variable (*not* using simulation techniques)
- `adapt()` in the “adapt” package does multivariate numerical integration

## Approximating the Sampling Distribution of a Statistic

To perform inference based on sample statistics, we typically need to know the sampling distribution of the statistics.

**Example:**  $X_1, \dots, X_n \sim iid N(\mu, \sigma^2)$ .

$$T = \frac{\bar{X} - \mu}{s/\sqrt{n}}$$

has a  $t(n - 1)$  distribution.

If  $\sigma^2$  known,

$$Z = \frac{\bar{X} - \mu}{\sigma/\sqrt{n}}$$

has a  $N(0, 1)$  distribution.

Then we can use these sampling distributions for inference (CIs, hypothesis tests).

What if the data's distribution is not normal?

1. Large sample: Central Limit Theorem
2. Small sample: Nonparametric procedures based on permutation distribution

- If population distribution is known, can approximate sampling distribution with simulation.
- Repeatedly ( $m$  times) generate random sample of size  $n$  from population distribution.
- Calculate statistic (say,  $S$ ) each time.
- The empirical distribution of  $S$ -values approximates its true distribution.

**Example 1:**  $X_1, \dots, X_4 \sim \text{Expon}(1)$

- What is the sampling distribution of  $\bar{X}$ ?
- What is the sampling distribution of sample midrange?

- What if we don't know the exact population distribution (more likely)?
- Can use *bootstrap methods*: Resample (randomly select  $n$  values from the original sample, with replacement). These “bootstrap samples” together mimic the population.
- For each of the, say,  $m$  bootstrap samples, calculate the statistic of interest.
- These  $m$  values will approximate the sampling distribution.

**Example 2:** Observe 7, 9, 13, 12, 4, 6, 8, 10, 10, 7 from an unknown population type.



- Bootstrap sampling built into R in the “boot” package.  
Try `library(boot); help(boot)` for details.
- If you know the *form* of the population distribution, but not the parameters, a *parametric* bootstrap can be used.
- Simple bootstrap CIs have some drawbacks
- More complicated “bias-corrected” bootstrap methods have been developed