### 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 Cls)

# **Optimization**

- ullet 1. Finding the x that maximizes (or minimizes) a complicated function h(x) can be difficult analytically
- ullet Situation even tougher if  ${\bf x}$  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 x values:  $x_0, x_1, \dots$ 

- 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, 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:

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

- ullet Determine c such that  $c \geq h(x)$  across entire region of interest. (Here, c=4)
- ullet 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])
- ullet Count the number of times (call this m) that the  $Y_i$  is less than the  $h(X_i)$
- ullet Then  $I pprox c(b-a) rac{m}{n}$

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

### **Classical Monte Carlo Integration**

$$I = \int_{a}^{b} h(x) \, dx$$

ullet Take n random uniform values  $U_1,\ldots,U_n$  (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.
- ullet Take n random values  $X_1,\ldots,X_n$  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:  $X_1, \ldots, X_n \sim iid \ N(\mu, \sigma^2)$ .

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

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).

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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.
- ullet The empirical distribution of S-values approximates its true distribution.

Example 1:  $X_1, \ldots, X_4 \sim Expon(1)$ 

- $\bullet$  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.
- ullet For each of the, say, m bootstrap samples, calculate the statistic of interest.
- ullet 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