8. Monte Carlo Simulations

#### Observables in classical statistical mechanics

the average value of an observable A is

$$\langle A \rangle = \int p(\mathbf{r}_1, \mathbf{r}_2, \ldots) A(\mathbf{r}_1, \mathbf{r}_2, \ldots) d\mathbf{r}_1 d\mathbf{r}_2 \ldots = \frac{\int \exp(-\beta U(\mathbf{r}_1, \mathbf{r}_2, \ldots)) A(\mathbf{r}_1, \mathbf{r}_2, \ldots) d\mathbf{r}_1 d\mathbf{r}_2 \ldots }{\int \exp(-\beta U(\mathbf{r}_1, \mathbf{r}_2, \ldots)) d\mathbf{r}_1 d\mathbf{r}_2 \ldots }$$

- $\blacktriangleright$  a force field specifies  $U({\bf r}_1,{\bf r}_2,\ldots)$  for a given system
- ▶ to calculate the average value of an observable, we need to draw samples from the Boltzmann distribution  $p(\mathbf{r}_1, \mathbf{r}_2, ...)$

## Sampling from a probability distribution

- for a probability distribution p(x), drawing samples from it means generating a sequence of random numbers  $\{x_1, x_2, \ldots\}$  such that the probability of x being in the sequence is p(x)
- in other words, the histogram of sampled numbers should match the probability distribution
- when p(x) is a standard distribution, such as the uniform or normal distribution, specialized procedures are available to draw samples from it
- however, these specialized procedures do not work for general probability distributions

## Sampling from standard distributions

 $\blacktriangleright$  assume we know how to draw samples from a uniform distribution on the interval [0,1]  $x = {\rm rand}()$ 

b how to draw samples from a uniform distribution on the interval [a, b]

### Sampling from standard distributions

▶ assume we know how to draw samples from a uniform distribution on the interval [0,1]x = rand()

• how to draw samples from a uniform distribution on the interval [a, b]

 $y = a + (b - a) \times x$ 

how to draw samples from a normal distribution with mean 0 and standard deviation 1 the Box-Muller transform

$$u = \operatorname{rand}(); v = \operatorname{rand}()$$
$$x = \sqrt{-2 \ln u} \cos(2\pi v)$$
$$y = \sqrt{-2 \ln u} \sin(2\pi v)$$

▶ many programming languages provide functions to draw samples from standard distributions
(ML ∪ MD) ∩ Biophysics

## General methods for sampling

#### Monte Carlo methods

- rejection sampling
- importance sampling
- Markov chain Monte Carlo Metropolis-Hastings algorithm
- Molecular dynamics simulations

# **Rejection sampling**

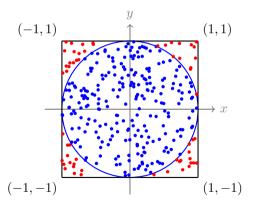
how to draw samples from a uniform distribution inside a circle of radius 1 centered at the origin in 2D

# **Rejection sampling**

how to draw samples from a uniform distribution inside a circle of radius 1 centered at the origin in 2D

repeat the following steps

- 1. draw a random point (x, y) in the square  $[-1, 1] \times [-1, 1]$
- 2. if the point is inside the circle, keep it; otherwise, discard it



### Importance sampling

 $\blacktriangleright$  the average value of an observable A(x) with respect to a probability distribution p(x) is

$$\langle A\rangle = \int p(x)A(x)\,dx$$

▶ if drawing samples from p(x) is difficult whereas drawing samples from another probability distribution q(x) is easy, the average value can be estimated as

$$\langle A \rangle = \int q(x) \frac{p(x)}{q(x)} A(x) \, dx \approx \frac{1}{N} \sum_{i=1}^{N} \frac{p(x_i)}{q(x_i)} A(x_i),$$

where  $\{x_1, x_2, \ldots, x_N\}$  are samples drawn from q(x)

•  $w(x_i) = p(x_i)/q(x_i)$  is called the importance weight

## Importance sampling

 $\blacktriangleright$  to generate samples from a probability distribution p(x) using importance sampling

- 1. draw samples  $\{x_1, x_2, \ldots, x_N\}$  from a probability distribution q(x)
- 2. calculate the importance weights  $\{w(x_1), w(x_2), \ldots, w(x_N)\}$

 $\blacktriangleright$  the average value of an observable A(x) with respect to p(x) is

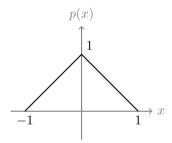
$$\langle A \rangle \approx \frac{1}{N} \sum_{i=1}^{N} w(x_i) A(x_i)$$

▶ for q(x),  $\{x_1, x_2, ..., x_N\}$  are samples with equal weights; for p(x),  $\{x_1, x_2, ..., x_N\}$  are samples with importance weights

•  $\{x_1, x_2, \dots, x_N\}$  and their importance weights can be used to generate samples of p(x) with equal weights by resampling

## Sampling from a triangular distribution

how to draw samples from the following triangular distribution



- using rejection sampling
- using importance sampling

# Rejection sampling and importance sampling

- **b** both requires sampling from a proposal distribution q(x)
- their efficiency depends on the choice of q(x)
- q(x) should be close to the target distribution p(x)
- q(x) should be easy to sample from
- $\blacktriangleright$  difficult to find a good proposal distribution for a general target distribution p(x)

### Markov chain Monte Carlo

> a general method to sample from a probability distribution p(x)

- generates a sequence of samples  $\{x_1, x_2, \ldots\}$ , where each sample is generated using the previous sample, i.e., it is a Markov chain
- ▶ assume that the sample at step i is  $x_o$ , the sample at step i + 1 is generated based on the transition probability  $T(x_n|x_o)$
- the transition probability is designed such that the samples generated by the Markov chain converge to the target distribution p(x)
- one design principle is to make the transition probability satisfy the *detailed balance* condition

$$p(x_o)T(x_n|x_o) = p(x_n)T(x_o|x_n)$$

#### **Detailed balance condition**

the detailed balance condition

$$p(x_o)T(x_n|x_o) = p(x_n)T(x_o|x_n)$$

implies that the probability flow going from  $x_o$  to  $x_n$  is equal to the probability mass going from  $x_n$  to  $x_o$ 

• along with other conditions, the detailed balance condition ensures that p(x) is invariant with respect to the transition probability

$$x_o \sim p(x) \implies x_n \sim p(x)$$

proof

$$\sum_{x_o} p(x_o) T(x_n | x_o) = \sum_{x_o} p(x_n) T(x_o | x_n) = p(x_n) \sum_{x_o} T(x_o | x_n) = p(x_n)$$

 $(ML \cup MD) \cap Biophysics$ 

Ding

## **Metropolis-Hastings algorithm**

▶ a special case of the Markov chain Monte Carlo method

#### the algorithm

- 1. given the sample  $x_o$  at step i, propose a new sample  $x_n$  using a proposal distribution  $q(x_n|x_o)$
- 2. calculate the acceptance probability

$$\alpha(x_n|x_o) = \min\left(1, \frac{p(x_n)q(x_o|x_n)}{p(x_o)q(x_n|x_o)}\right)$$

3.  $x_{i+1} = x_n$  with probability  $\alpha(x_n|x_o)$ ; otherwise,  $x_{i+1} = x_o$ 

the transition probability is

$$T(x_n|x_o) = q(x_n|x_o)\alpha(x_n|x_o)$$

#### **Metropolis-Hastings algorithm**

#### it satisfies the *detailed balance* condition

 $p(x_o)T(x_n|x_o) = p(x_o)q(x_n|x_o)\alpha(x_n|x_o) = p(x_n)q(x_o|x_n)\alpha(x_o|x_n) = p(x_n)T(x_o|x_n)$ 

▶ when  $q(x_n|x_o) = q(x_o|x_n)$ , the acceptance probability simplifies to

$$\alpha(x_n|x_o) = \min\left(1, \frac{p(x_n)}{p(x_o)}\right)$$

and the detailed balance condition simplifies to

$$p(x_o)\alpha(x_n|x_o) = p(x_n)\alpha(x_o|x_n)$$

#### Metropolis-Hastings algorithm for Boltzmann distributions

 $\blacktriangleright$  for a system with fixed N,V,T, the Boltzmann distribution is

$$p(\mathbf{r}_1, \mathbf{r}_2, \ldots) = \frac{1}{Z} \exp(-\beta U(\mathbf{r}_1, \mathbf{r}_2, \ldots))$$

the Metropolis-Hastings algorithm for the Boltzmann distribution

- 1. given the sample  $\mathbf{r}_o$  at step *i*, propose a new sample  $\mathbf{r}_n$  using a proposal distribution  $q(\mathbf{r}_n|\mathbf{r}_o)$  that satisfies  $q(\mathbf{r}_n|\mathbf{r}_o) = q(\mathbf{r}_o|\mathbf{r}_n)$
- 2. calculate the acceptance probability

$$\alpha(\mathbf{r}_{n}|\mathbf{r}_{o}) = \min\left(1, \frac{\exp(-\beta U(\mathbf{r}_{n}))}{\exp(-\beta U(\mathbf{r}_{o}))}\right) = \min\left(1, \exp(-\beta \Delta U)\right)$$

where  $\Delta U = U(\mathbf{r}_n) - U(\mathbf{r}_o)$ 3.  $\mathbf{r}_{i+1} = \mathbf{r}_n$  with probability  $\alpha(\mathbf{r}_n | \mathbf{r}_o)$ ; otherwise,  $\mathbf{r}_{i+1} = \mathbf{r}_o$