

# **Computational Physics (PHYS6350)**

Lecture 18: Random numbers: Part II

- Non-uniformly distributed random numbers
- Importance sampling

Reference: Chapter 10 of Computational Physics by Mark Newman

#### **Instructor:** Volodymyr Vovchenko (vvovchenko@uh.edu)

**Course materials:** <u>https://github.com/vlvovch/PHYS6350-ComputationalPhysics</u> **Online textbook:** <u>https://vovchenko.net/computational-physics/</u>

### Nonuniformly distributed random numbers

In many cases we deal with random numbers  $\xi$  that are distributed non-uniformly.

Common examples are:

- Exponential distribution  $\rho(x) = e^{-x}$ .
- Gaussian distribution  $\rho(x) \propto e^{-\frac{x^2}{2\sigma^2}}$ .
- Power-law distribution  $\rho(x) \propto x^{\alpha}$ .
- Arbitrary peaked distributions.

There are two common methods for generating nonuniform random variates. They both make use of uniformly distributed variates.

- Inverse transform sampling
- Rejection sampling

The basic idea is that if  $\eta$  is a uniformly distibuted random variable, some function of it,  $\xi = f(\eta)$ , is not. The idea is to sample  $\eta$  and calculate  $\xi$  via this function such that  $\xi$  corresponds to a desired probability density  $\rho(\xi)$ . How to find the function  $f(\eta)$ ?

Without the loss of generality assume that  $\xi \in (-\infty, \infty)$  and that  $f(\eta)$  maps  $\eta$  to  $\xi$  such that  $f(0) \to -\infty$ . Consider now the cumulative distribution function

$$G(x) = Pr(\xi < x) = \int_{-\infty}^{x} \rho(\xi) d\xi.$$

It corresponds to the probability that  $\eta < y$  where y is such that x = f(y). Since  $\eta$  is uniformly distributed, this probability equals to y. Therefore,

$$G[x = f(y)] = y,$$

thus

$$f(y) = G^{-1}(y).$$

If we can calculate the inverse of  $G^{-1}(y)$  of the cumulative distribution function for  $\xi$ , we are good.



#### **Inverse transform sampling**

The algorithm follows these steps:

1. Calculate the cumulative distribution function (CDF)

$$G(x) = \int_{-\infty}^{x} \rho(\xi) d\xi$$

2. Find the inverse function  $G^{-1}(y)$  as the solution to the equation

G(x) = y

3. Sample uniformly distributed random variables  $\eta$  and compute  $\xi$  using the inverse function:

**Challenges:** Sometimes, evaluating G(x) and/or  $G^{-1}(y)$  explicitly is difficult. In such cases, numerical integration and/or non-linear equation solvers may be required.



#### **Example: Exponential Distribution**

1. Recall the radioactive decay process. The time of decay is distributed according to the probability density function:  $1 - \frac{1}{1 - t}$ 

$$\rho(t) = \frac{1}{\tau} e^{-\frac{t}{\tau}}.$$

2. The cumulative distribution function is given by:

$$F(x) = \int_0^x \frac{1}{\tau} e^{-\frac{t}{\tau}} dt = 1 - e^{-\frac{x}{\tau}}.$$

3. To apply inverse transform sampling, we need to invert F(x) by solving:

$$1-e^{-\frac{t}{\tau}}=\eta$$

4. Solving for *t*, we obtain:

$$t(\eta) = -\tau \ln(1-\eta).$$



One way to sample points inside a unit circle is by switching to **polar coordinates**:

 $x = r \cos(\phi), \qquad y = r \sin(\phi),$ 

```
where we sample r \in [0,1) and \phi \in [0, 2\pi).
```

Naively, one could sample r and  $\phi$  independently from two uniform distributions. Let's see what happens!

```
def sample_xy_naive():
    r = np.random.rand()
    phi = 2 * np.pi * np.random.rand()
    return r*np.cos(phi), r*np.sin(phi)

xplot = []
yplot = []
N = 1000
for i in range(N):
    x, y = sample_xy_naive()
    xplot.append(x)
    yplot.append(y)

plt.plot(xplot,yplot,'o',color='r')
plt.show()
```

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The points clump more in the center!

- The points clump more in the center because r is not uniformly distributed.
- Recall the **differential area element** in polar coordinates:  $dxdy = rdrd\phi$
- This leads to the probability density functions:  $\rho_r(r) = 2r$ ,  $\rho_\phi(\phi) = \frac{1}{2\pi}$ .
- Cumulative Distribution Function (CDF)

$$F_r(r) = \int_0^r \rho_r(r')dr' = r^2.$$

• To obtain a properly distributed r, we solve  $F_r(r) = \eta$  which gives:

$$r = \sqrt{\eta}$$

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 $r = \sqrt{\eta}$ 

### Sampling of an Isotropic Direction in 3D

One common problem in **Monte Carlo simulations** is the **random sampling of an isotropic direction in 3D space**. This issue arises in various contexts, such as:

- Sampling a random orientation of an axially symmetric object (e.g., a rod).
- Sampling the **momentum direction** of a particle.

This problem is equivalent to **choosing a random point on a unit sphere**. The coordinates x, y, z on the sphere can be parametrized using **azimuthal** and **polar angles**:

- $\phi \in [0, 2\pi)$  (azimuthal angle)
- $\theta \in [0, \pi)$  (polar angle)

Using these angles, the Cartesian coordinates are given by:

 $x = \sin \theta \cos \phi$  $y = \sin \theta \sin \phi$  $z = \cos \theta$ 

### Sampling of an Isotropic Direction in 3D

 $x = \sin \theta \cos \phi$  $y = \sin \theta \sin \phi$  $z = \cos \theta$ 

• The solid angle element is:

 $d\Omega = \sin(\theta) d\theta d\phi,$ 

- The random variables  $\phi$  and  $\theta$  are independent.
- $\phi$  is uniformly distributed in [0,  $2\pi$ ], making its sampling straightforward.
- However,  $\theta$  has a **weighted probability density function**:

The cumulative distribution function is:

$$F_{\theta}(\theta) = \int_0^{\theta} \frac{1}{2} \sin(\theta') d\theta' = \frac{1 - \cos(\theta)}{2}$$

Solving  $F_{\theta}(\theta) = \eta$ , we obtain:

$$\theta = \arccos(2\eta - 1).$$

In practice, work directly with  $\cos \theta$  and  $\sin \theta$ :

$$\cos(\theta) = 2\eta - 1, \qquad \sin(\theta) = \sqrt{1 - [\cos(\theta)]^2}$$

$$\rho_{\theta}(\theta) = \frac{1}{2}\sin(\theta),$$

### Sampling an isotropic direction

```
def sample xyz isotropic():
    phi = 2 * np.pi * np.random.rand()
    costh = 2 * np.random.rand() - 1
    sinth = np.sqrt(1-costh*costh)
    return sinth * np.cos(phi), sinth * np.sin(phi), costh
xplot = []
yplot = []
zplot = []
N = 500
for i in range(N):
    x, y, z = sample_xyz_isotropic()
   xplot.append(x)
   yplot.append(y)
    zplot.append(z)
fig = plt.figure()
ax = fig.add_subplot(projection='3d')
ax.scatter(xplot,yplot,zplot)
ax.set_xlabel('x')
ax.set_ylabel('y')
ax.set_zlabel('z')
plt.show()
```



## Sampling normally distributed variables



One of the most common probability distributions is the **normal (or Gaussian) distribution**, given by:

$$\rho(x) = \frac{1}{\sqrt{2\pi\sigma}} e^{-\frac{(x-\mu)^2}{2\sigma^2}}$$

There are many standard methods for sampling from this distribution. One common approach is to **standardize the variable** by making the transformation  $x \rightarrow \mu + \sigma x$ 

After this transformation, the new variable follows a **standard normal distribution** with **zero mean** and **unit standard deviation**:  $1 - \frac{1}{2}$ 

$$\rho(x)=\frac{1}{\sqrt{2\pi}}e^{-\frac{x^2}{2}}.$$

Calculating the cumulative distribution function  $F(x) = \int_{-\infty}^{x} \rho(x') dx'$  is **not straightforward**, use numerical methods.

#### Sampling normally distributed variables

Instead of one variable, we can consider a pair of independent normally distributed variables x, y:

$$\rho(x, y) = \frac{1}{2\pi} e^{-\frac{x^2}{2}} e^{-\frac{y^2}{2}},$$

Making a change of variables to polar coordinates

$$x = r \cos(\phi), \qquad y = r \sin(\phi),$$

and taking into account

$$dxdy = rdrd\phi$$

we get

$$\rho(r,\phi) = \frac{1}{2\pi} r e^{-r^2/2}.$$

Therefore, we can sample *x* and *y* by sampling two independent random variables *r* and  $\phi$ .  $\phi$  is uniformly distributed in  $[0, 2\pi)$ . For *r* we have the following probability density

$$\rho_r(r)=re^{-r^2/2},$$

and the cumulative distribution function

$$F_r(r) = \int_0^r r' e^{-r'^2/2} dr' = 1 - e^{-r^2/2},$$

therefore

$$r=\sqrt{-2\ln(1-\eta)}.$$

```
def sample_xy_normal():
    phi = 2 * np.pi * np.random.rand()
    eta = np.random.rand()
    r = np.sqrt(-2*np.log(1-eta))
    return r * np.cos(phi), r * np.sin(phi)

N = 10000
samples = []
for i in np.arange(0,N,2):
    x, y = sample_xy_normal()
```





### **Rejection sampling**

The **rejection sampling method** allows one to sample a variable  $\xi$  from an **envelope distribution** and accept or reject it with a certain probability.

Consider again the probability density function for the polar angle:

$$\rho_{\theta}(\theta) = \frac{\sin(\theta)}{2}.$$

Since  $\rho_{\theta}$  is bounded from above, we define:  $\rho_{\theta}^{\max} = 1/2$ .

1. Sample a candidate value  $\theta_{cand}$  from a uniform distribution over (0,  $\pi$ ).

2. Accept  $\theta_{cand}$  with probability:  $p = \rho_{\theta}(\theta_{cand})/\rho_{\theta}^{max}$ .

3. This step can be performed by sampling y from a uniform distribution over  $(0, \rho_{\theta}^{max})$  and accepting  $\theta_{cand}$  if  $y < \rho_{\theta}(\theta_{cand})$ 



**Geometric Interpretation:** If we consider  $\theta_{cand} = x$  and y as the **coordinates of a point** in a plane, we accept  $\theta_{cand}$  if it lies below the curve defined by  $\rho_{\theta}(\theta)$ . This ensures that  $\theta_{cand}$  values are accepted at a rate proportional to  $\rho_{\theta}(\theta)$ , as desired.

#### Advantages of Rejection Sampling:

 $\rho_{\theta}(\theta)$  does not need to be a normalized distribution for the method to work.

```
def sample_rejection(rho, a, b, rhomax):
    while True:
        x_cand = a + (b-a)*np.random.rand()
        y = rhomax * np.random.rand()
        if (y < rho(x_cand)):</pre>
            return x_cand
    return 0.
def rho_theta(theta):
    return np.sin(theta) / 2.
N = 100000
samples = []
for i in np.arange(0,N,1):
    theta = sample_rejection(rho_theta, 0., np.pi, 0.5)
    samples.append(theta)
```



## **Pros and Cons of Rejection Sampling**

#### **Pros:**

- Does not require the distribution to be normalized.
- Works even if  $y_{max}$  is larger than the true maximum of  $\rho(x)$
- Applicable to **generic distributions** and does not require the evaluation of the cumulative distribution function.

#### Cons:

- Can be **inefficient** if the rejection rate is high (e.g., for highly peaked distributions).
- Not directly applicable to distributions over infinite ranges.

#### **Generalizations of Rejection Sampling**

To address some of its limitations, several generalizations of rejection sampling can be used, including:

- Adaptive rejection sampling by considering multiple enveloping rectangles.
- Variable transformation to map an infinite interval into a finite one.
- Sampling from a non-uniform enveloping distribution for better efficiency.

#### **Importance** sampling

Recall the calculation of an integral as statistical average

$$I = \int_{a}^{b} f(x)dx = (b-a)\langle f \rangle, \qquad \text{where}$$

$$\langle f \rangle = \frac{1}{N} \sum_{i=1}^{N} f(x_i), \qquad x_i \in U(a, b)$$

Some issues with the method:

- Sample unimportant regions (e.g. f is highly peaked)
- Integrable singularities

#### Importance sampling:

Sample  $x_i$  from a *non-uniform* distribution w(x) that resembles f(x).

The integrand is then calculated as

$$I = \int_{a}^{b} \frac{f(x)}{w(x)} w(x) dx = \left\langle \frac{f(x)}{w(x)} \right\rangle_{w}$$

#### **Normalization:**

$$\int_{a}^{b} w(x)dx = 1.$$

Error: 
$$\delta I = \frac{\sqrt{\left\langle \left[\frac{f(x)}{w(x)}\right]^2 \right\rangle_w - \left\langle \frac{f(x)}{w(x)} \right\rangle_w^2}}{\sqrt{N}}$$

#### **Importance** sampling

$$I = \int_{a}^{b} \frac{f(x)}{w(x)} w(x) dx = \left\langle \frac{f(x)}{w(x)} \right\rangle_{w} \qquad \qquad \delta I = \frac{\sqrt{\left\langle \left[\frac{f(x)}{w(x)}\right]^{2} \right\rangle_{w} - \left\langle \frac{f(x)}{w(x)} \right\rangle_{w}^{2}}}{\sqrt{N}}.$$

• For w(x)=1/(b-a) we recover the mean value method

$$I = \int_{a}^{b} f(x)dx = (b-a)\langle f \rangle$$

• For w(x) 
$$\propto$$
 f(x) one has  $\left\langle \frac{f(x)}{w(x)} \right\rangle = \text{const} = 1$  and  $\delta I = 0$ 

```
# Calculate integral \int_a^b f(x) dx using importance sampling
\# f = f(x) is the integrand
# N is the number of random samples
\# wx = w(x) is the normalized probability density from which
# the sampling takes place
# sampler is a function which samples a random number from w(x)
def intMC_weighted(f, N, wx, sampler):
    total = 0
    total sq = 0
    for i in range(N):
        x = sampler()
        fval = f(x)
        total += fval / wx(x)
        total_sq += (fval / wx(x)) **2
    fw_av = total / N
    fwsq_av = total_sq / N
    return fw_av, np.sqrt((fwsq_av - fw_av*fw_av)/N)
```

### **Importance sampling: Example**



Integrable singularity at x=0

### **Importance sampling: Example**



$$w(x) = \frac{1}{b-a}$$

def uniform\_sample(): eta = np.random.rand() return eta def uniform\_w(x): return 1. np.random.seed(1) N = 1000000 I, err = intMC\_weighted(f, N, uniform\_w, uniform\_sample) print("I = ",I," +- ",err) I = 0.8374063441946126 +- 0.0017772180714415427

#### Importance sampling

 $w(x) = \frac{1}{2\sqrt{x}}$   $I = \left\langle \frac{2}{e^x + 1} \right\rangle_w$   $x = \eta^2$ 

def rsqrt\_sample():
 eta = np.random.rand()
 return eta \* eta

N = 1000000

 $\int_{0}^{1} \frac{x^{-1/2}}{e^{x}+1} dx$ 

def rsqrt\_w(x):
 return 1. / (2. \* np.sqrt(x))



I, err = intMC\_weighted(f, N, rsqrt\_w, rsqrt\_sample)
print("I = ",I," +- ",err)

I = 0.839014917136739 + - 0.0001409071521618816

Statistical error is more than  $\times 10$  smaller than in the mean value method. We would need more than  $\times 100$  samples in the mean value method to reach the same accuracy as importance sampling in this case.