

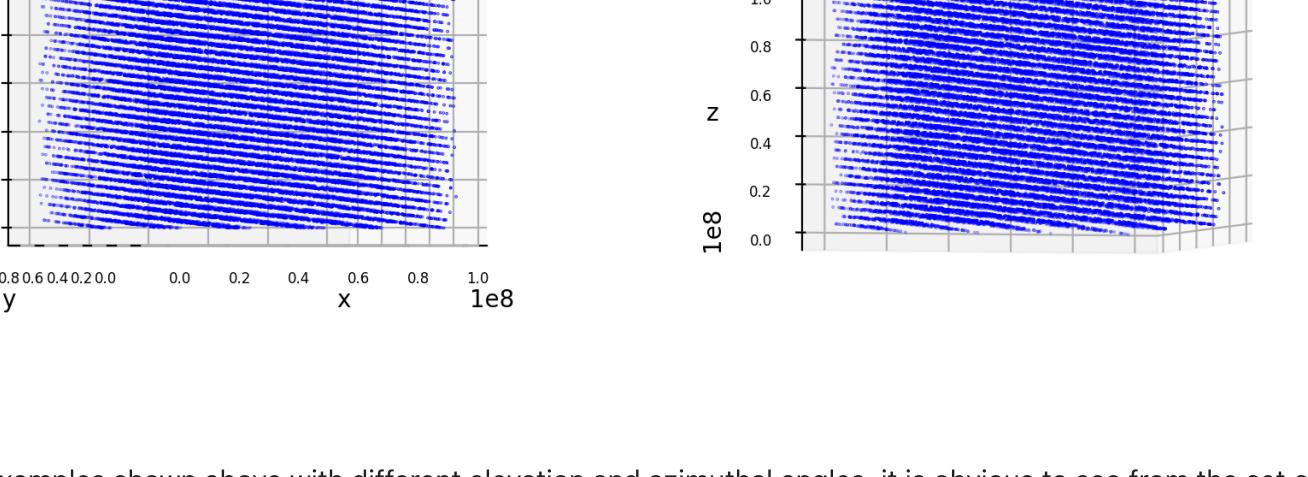
```
In [1]: import numpy as np
from matplotlib import pyplot as plt
from mpl_toolkits import tk3dm
import time
from IPython.display import display, Latex
from datetime import datetime
import os
from scipy import signal
%matplotlib notebook
```

PROBLEM 1

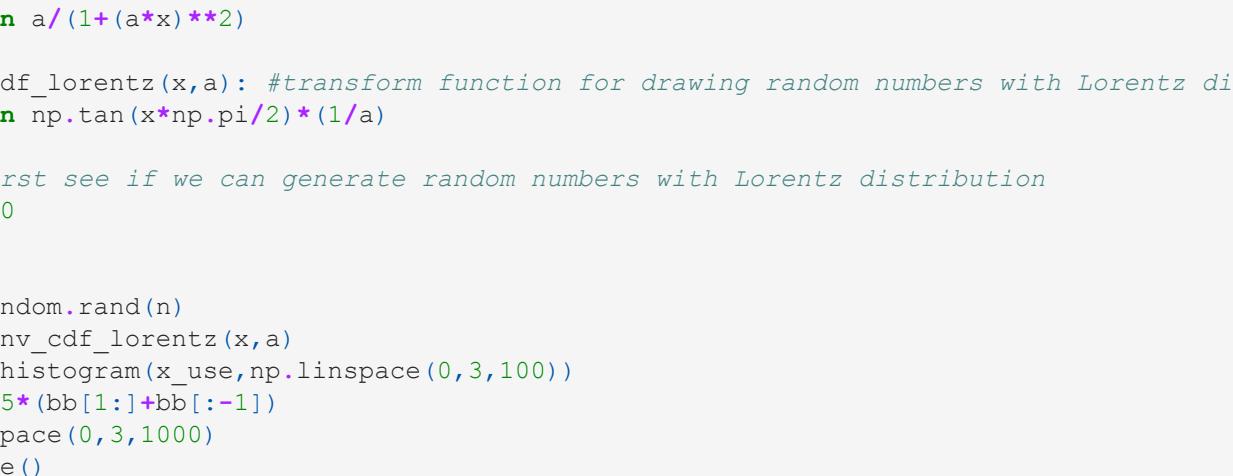
We want to see the broken default random number generator in the C standard library. If we generate (x,y,z) coordinates, for "true" random numbers, we should see uniformly dispersed points with no underlying pattern. Hence, if we generate (x,y,z) random numbers from the C library and observe a pattern as we plot them, it is evident that the default C library's PRNG is flawed!

```
In [2]: rand_points = np.loadtxt('rand_points.txt') #download random points from C library
```

```
fig = plt.figure()
ax = plt.axes(projection='3d',proj_type = 'ortho')
ax.scatter3D(rand_points[:,0], rand_points[:,1], rand_points[:,2],marker='.',color='blue',s=5)
ax.set_xlabel('x')
ax.set_ylabel('y')
ax.set_zlabel('z')
ax.tick_params(axis='both', which='major', labelsize=6)
ax.view_init(-179,129)
fig.set_size_inches(8,8)
```

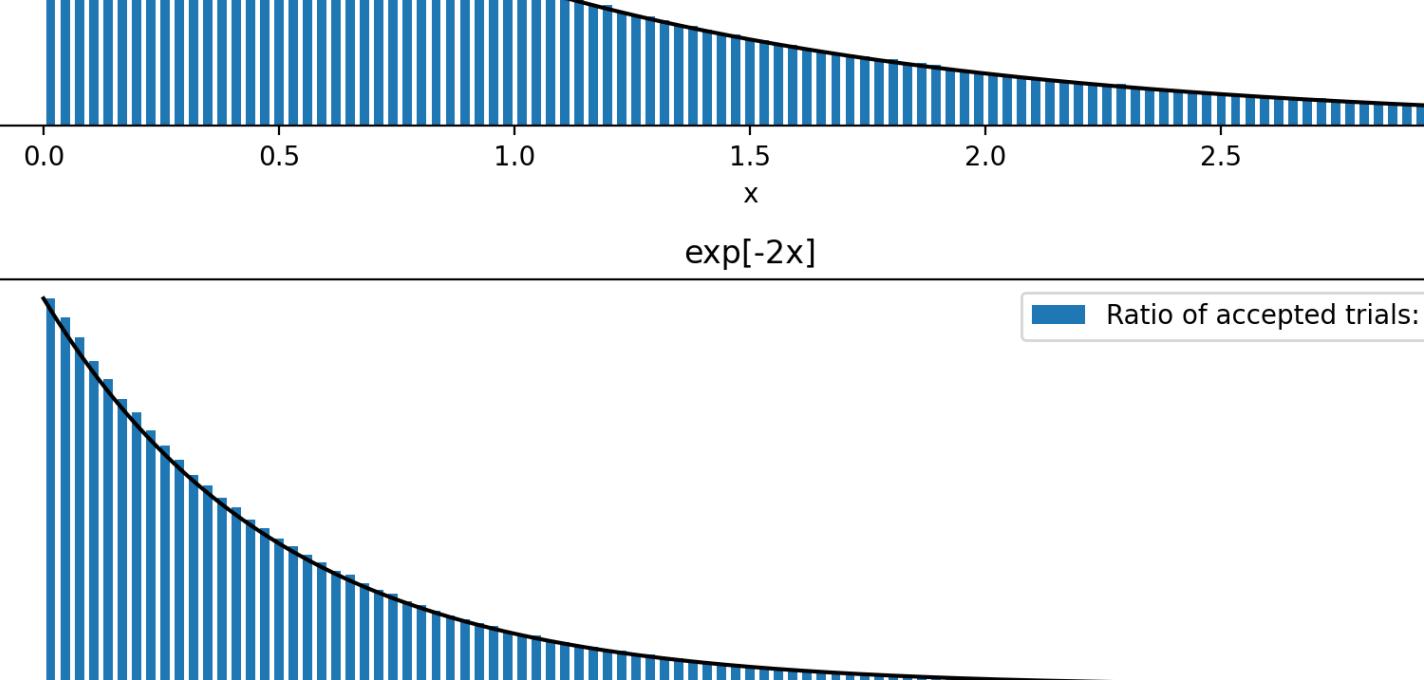
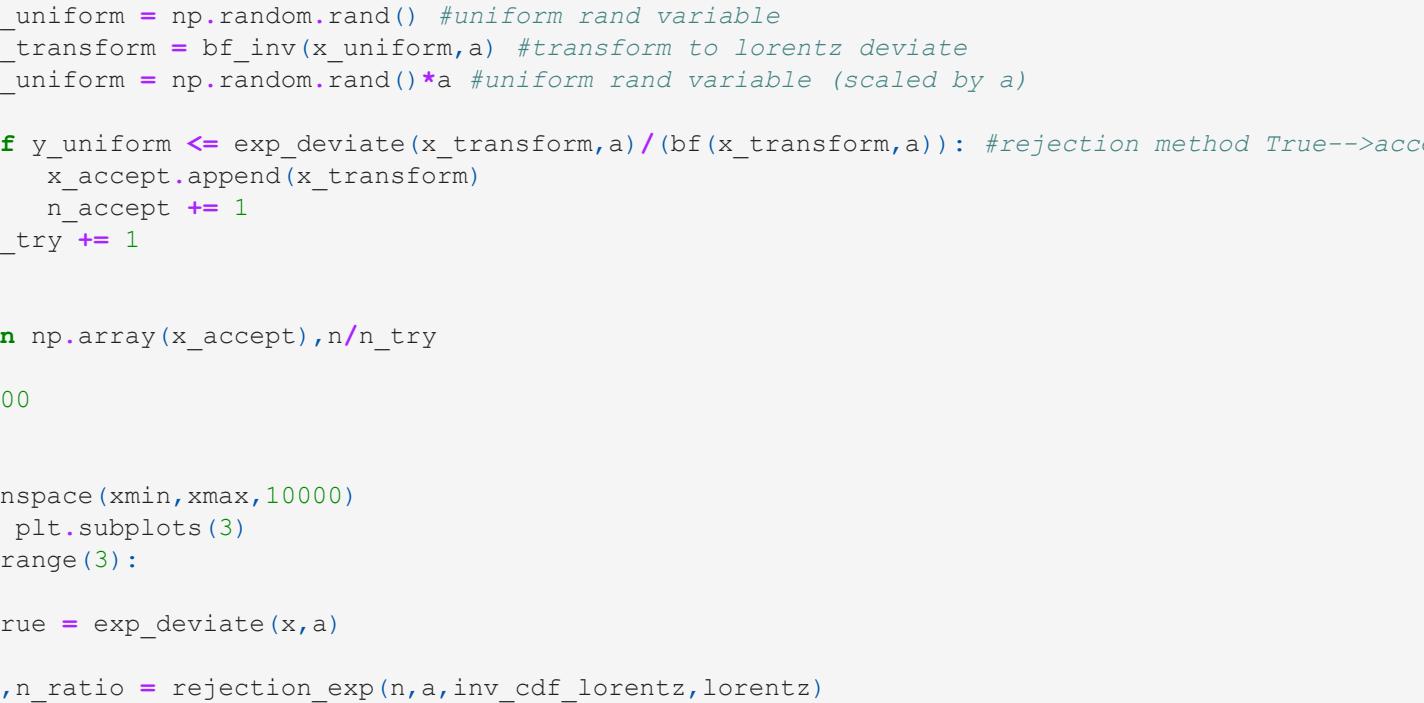


```
In [3]: fig = plt.figure()
ax = plt.axes(projection='3d',proj_type = 'ortho')
ax.scatter3D(rand_points[:,0], rand_points[:,1], rand_points[:,2],marker='.',color='blue',s=5)
ax.set_xlabel('x')
ax.set_ylabel('y')
ax.set_zlabel('z')
ax.tick_params(axis='both', which='major', labelsize=6)
ax.view_init(0,-12)
fig.set_size_inches(8,8)
```



```
In [4]: angles = [(1,-179),[0,-112],[-2,-109]]
fig = plt.figure()
pos = [221,222,223,224]
for i in range(4):
    ax=plt.subplot(pos[i],projection='3d',proj_type = 'ortho')
    ax.scatter3D(rand_points[:,0], rand_points[:,1], rand_points[:,2],marker='.',color='blue',s=0.5)
    ax.set_xlabel('x')
    ax.set_ylabel('y')
    ax.set_zlabel('z')
    ax.tick_params(axis='both', which='major', labelsize=6)
    if i==0:
        ax.view_init(angles[i-1][0],angles[i-1][1])
    else:
        ax.view_init(angles[i-1][0],angles[i-1][1])
    ax.set_title('Elevation:{}\nAzimuth:{}'.format(angles[i-1][0],angles[i-1][1]))
```

Elevation:-179
Azimuth:-129



From the three examples shown above with different elevation and azimuthal angles, it is obvious to see from the set of planes (about 30) that the triples are very not much randomly distributed in 3D space!

We can also view this by plotting plane projections i.e. $(ax + by, z)$ by choosing suitable a, b . Below shows the projection for $a = 1$ and $b = 0.5$, which again shows that the triples are very far from being randomly distributed viewed like this.

```
In [5]: #We can also view this by plotting a projection with suitable a,b
a=1
b=0.5
fig=plt.figure()
plt.plot(rand_points[:,0]+b*rand_points[:,1],rand_points[:,2],'.',markersize=2)
```

The centered at 0 Lorentz is given by
 $L(x,a) = \frac{a}{1+(ax)^2}$. We first normalize it, where $\int_0^\infty \frac{a}{1+(ax)^2} dx = \frac{\pi}{2}$ such that

$pdf_L(x) = \frac{2a}{\pi(1+(ax)^2)}$.

The transform method requires to simply take the cdf of the pdf and then inverse it to get x deviates:

$$\int_0^x \frac{2}{\pi(1+(ax)^2)} dx = \frac{2}{\pi} \arctan(ax) \Rightarrow \left[\frac{2}{\pi} \arctan(ax) \right]^{-1} = \frac{1}{a} \tan\left(\frac{\pi x}{2}\right)$$

So our function to draw random numbers from a Lorentz distribution is given by $\frac{1}{a} \tan\left(\frac{\pi x}{2}\right)$.

From the rejection method, the "ideal" ratio of accepted deviates is given the ratio of the area of $f(x)$ to the area of $p(x)$ (not normalized). In other words, accrate = $\frac{\int_0^\infty f(x) dx}{\int_0^\infty p(x) dx}$

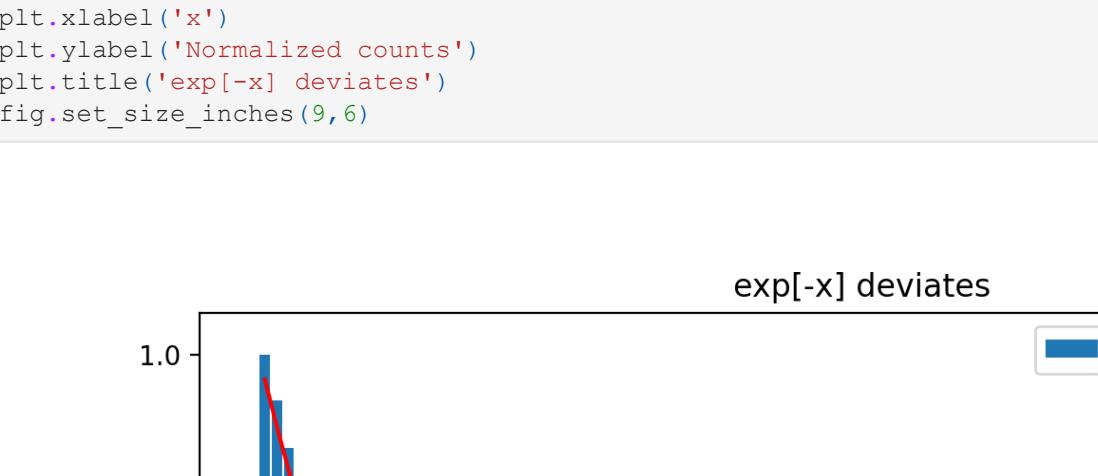
```
In [7]: def lorentz(x,a): #Lorentz centered at 0, scaled by 1/a
    return a/(1+(a*x)**2)

def inv_cdf_lorentz(x,a): #transform function for drawing random numbers with Lorentz distribution
    return np.tan((np.pi*x)/2)/(1/a)
```

#Let's first see if we can generate random numbers with Lorentz distribution
n=1000000
a=1

x = np.random.rand()
u=inv_cdf_lorentz(x,a)
aa,b=histogram(x,u,np.linspace(0,3,100))
b.cent=0.5*(bb[1]+bb[-1])

x=np.linspace(0,3,1000)
plt.figure()
plt.bar(b.cent,aa/n,0.02)
plt.plot(x,aa/n,'r--')
plt.xlabel('x')
plt.ylabel('Normalized Counts')
plt.title('Normalized Counts')



Indeed, we see that the lorentz deviates match very well with the lorentz distribution function! We can now use the rejection method to draw exponential deviates.

```
In [10]: def exp_deviates(x,a):
    return a*np.exp(-x*a) #normalized exponential pdf
```

```
def rejection_exp(n,a,bf_inv_ratio):
    n_accept=0
    n_try=0 #variable to then calculate the ratio of accepted steps
    x_accept = []
    while n_accept < n:
        x_uniform = np.random.rand() #uniform rand variable
        y_uniform = np.random.rand()*a #uniform rand variable (scaled by a)
        if y_uniform < exp_deviates(x_uniform,a): #rejection method True-->accept
            x_accept.append(x_uniform)
            y_accept.append(y_uniform)
            n_accept += 1
        n_try += 1
```

```
return np.array(x_accept),n/n_try
```

```
n=100000
xmin=0
xmax=3
x = np.linspace(xmin,xmax,1000)
fig,axs = plt.subplots(3)
for i in range(3):
    a=i+1
    exp_true = exp_deviates(x,a)

    x_exp_n_ratio = rejection_exp(n,a,inv_cdf_lorentz)
    aa,b=histogram(x_exp_n_ratio,np.linspace(xmin,xmax,100))
    b.cent=0.5*(bb[1]+bb[-1])

    axs[i].bar(b.cent,aa/n,0.02,label="Ratio of accepted trials: {}".format(f'{n_ratio:.3}'))
```

```
axs[i].set_xlabel('exp(-{}x)'.format(a))
axs[i].set_title('exp(-{}x)'.format(a))
axs[0].set_ylabel('Normalized Counts')
fig.set_size_inches(9,9)
fig.tight_layout()
```


From the plots we can see how everything went as planned! We have a matching exponential distribution and the ratio of accepted trials is approximately $\frac{1}{a}$.

Problem 3: RATIO OF UNIFORMS

Now we want to use the method of the ratio of uniforms. The general idea is to map the number line to a box. We know that this works since the Jacobian is a constant!

We take the (u,v) plane where $0 < u < \sqrt{v/b}$, sample u, v uniformly in the given region (where 0 is the lower bound). So, we use this and apply the ratio of uniforms method.

```
In [11]: #First we can plot our region
u=np.linspace(0,1,2001)
uu[1:]=u**0.5
v=u**0.5-np.log(u)
v=2*u**0.5-np.log(u)
w=2*u**0.5-np.log(u)

fig = plt.figure()
plt.plot(u,v,'k')
plt.fill_between(u,v,color='darkgreen',alpha=0.6)
plt.xlabel('u')
plt.ylabel('v')
plt.title('(u,v) Region')
fig.set_size_inches(9,6)
```


Above is the (u, v) region from which we draw u, v uniformly.

```
In [12]: def ratio_uniforms(N):
    u=np.linspace(0,1,2001)
    u[1:]=u**0.5
    v=u**0.5-np.log(u) #upper bound
    v_max = np.max(v) #approx max value that v can take

    n_accept = 0
    n_try = 0
    deviate = []
    while n_accept < N:
        u=np.random.rand() #loop until we have N deviates
        v = np.random.rand() * v_max
        r = v/u #ratio
        if r < np.exp(-b[1]): #if true --> accept deviate
            deviate.append(r)
            n_accept += 1
        n_try += 1

    return np.array(deviate),N/n_try
```

```
n=1000000
bb=0.5*(b[1]+b[-1])
bb=0.5*(bb[1]+bb[-1])

x=np.linspace(0,1,2001)
y=np.exp(-x)
y=y[1:]*bb**0.5
```

```
fig,axs = plt.subplots(3)
for i in range(3):
    a=i+1
    exp_deviates = ratio_uniforms(n)

    x_exp_n_ratio = exp_deviates[0]/np.max(exp_deviates)
    aa,b=histogram(x_exp_n_ratio,np.linspace(0,1,200))
    b.cent=0.5*(bb[1]+bb[-1])

    axs[i].bar(b.cent,aa/n,0.02,label="Ratio of accepted trials: {}".format(f'{n_ratio:.3}'))
```

```
axs[i].set_xlabel('exp(-{}x) deviates'.format(a))
axs[i].set_title('exp(-{}x) deviates'.format(a))
fig.set_size_inches(9,9)
```


So we see the the exponential deviates with the ratio of uniforms match the exponential distribution! The method is slightly more efficient than the rejection method (by about 5%) so this is the most efficient method between the two so far!

