The Python programming language is one of the most popular languages in both academia and industry. It is heavily used in data science for simple data analysis and complex machine learning. By most accounts, in the last few years, Python has eclipsed the R programming language in popularity for scientific/statistical computation. Its popularity is due to intuitive and readable syntax that can be implemented in a powerful object oriented programming paradigm, if so desired, as well as being open source. It is for these reasons that I decided to transcribe the Introduction to Simulation chapter in Pishro-Nik's Introduction to Probability, Statistics and Random Processes book into Python.

This entire chapter was written in a Jupyter notebook, an interactive programming environment, primarily for Python, that can be run locally in a web browser. Jupyter notebooks are ideal for quick and interactive data analysis, incorporating markdown functionality for clean presentations and code sharing. If you are a fan of RStudio, you will most likely be fond of Jupyter notebooks. This entire notebook is available freely at https://github.com/dsrub/solutions_to_probability_statistics.

Additionally, much of this code was written using the Numpy/SciPy library, Python's main library for scientific computation and numerical methods. Numpy has a relatively clear and well documented API (https://docs.scipy.org/doc/numpy/reference/index.html), a reference which I utilize almost daily.

I start with a few basic imports, and define several functions I will use throughout the rest of this chapter.

```
#define html style element for notebook formatting
from IPython.core.display import HTML
with open('style.txt', 'r') as myfile:
    notebook_style = myfile.read().replace('\n', '')
HTML(notebook_style)
#import some relevant packages and plot inline
import matplotlib.pyplot as plt
import numpy as np
%matplotlib inline
#define a few functions I will be using throughout the rest of the notebook
#function to print several of the RGNs to the screen
def print_vals(RNG_function, *args):
    for i in range(5):
        print('X_' + str(i)+' = ', RNG_function(*args))
#plotting function
def plot_results(x, y, xlim=None, ylim=None, xlabel=None, ylabel=None, \
                 title=None, labels=None):
    plt.figure(1, figsize = (6, 4))
    plt.rc('text', usetex=True)
    plt.rc('font', family = 'serif')
```

```
if labels:
    plt.plot(x[0], y[0], label=labels[0], linewidth = 2)
    plt.plot(x[1], y[1], label=labels[1], linewidth = 2)
    plt.legend(loc='upper right')
else:
    plt.plot(x, y, linewidth = 2)
if xlim:
    plt.xlim(xlim)
if ylim:
    plt.ylim(ylim)
if xlabel:
    plt.xlabel(xlabel, size = 15)
if ylabel:
    plt.ylabel(ylabel, size = 15)
if title:
    plt.title(title, size=15)
plt.xticks(fontsize = 15)
plt.yticks(fontsize = 15);
```

Example 1. (Bernoulli) Simulate tossing a coin with probability of heads p.

Solution: We can utilize the algorithm presented in the book, which uses random variables drawn from a Unif(0,1) distribution. The following function implements this algorithm in Python to generate a Bern(p) (pseudo) random variable.

```
def draw_bern(p, N):
    """
    A Bern(p) pseudo-RNG
    """
    U = np.random.uniform(size = N)
    if N == 1: U = U[0]
    X = (U < p) + 0

    return X

#print a few examples of the RGNs to the screen
p = 0.5
print_vals(draw_bern, p, 1)

X_0 = 0
X_1 = 0
X_2 = 0
X_3 = 0</pre>
```

Note that we can directly sample from a Bern(p) distribution with Numpy's binomial random number generator (RNG) by setting n = 1 with: np.random.binomial(1, p).

 $X_{4} = 1$

Example 2. (Coin Toss Simulation) Write code to simulate tossing a fair coin to see how the law of large numbers works.

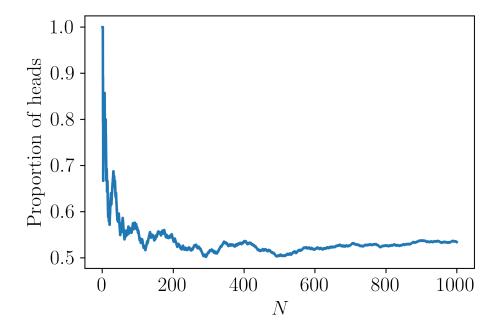
Solution: I draw 1000 Bern(0.5) random variables and compute the cumulative average.

```
#generate data, compute proportion of heads and plot

#set a seed for reproducibility
np.random.seed(2)

X = draw_bern(0.5, 1000)
avg = np.cumsum(X)/(np.arange(1000) + 1)
plot_results(np.arange(1000) + 1, avg, xlabel='$N$', ylabel='Proportion of heads')

#reset seed
np.random.seed(0)
```



Example 3. (Binomial) Generate a Bin(50, 0.2) random variable.

Solution: If $X_1, X_2, ..., X_n$ are drawn iid from a Bern(p) distribution, then we can express a Bin(n,p) random variable as $X = X_1 + X_2 + ... + X_n$. Therefore we can utilize the code we have already written for drawing a Bern(p) random variable to draw a Bin(n,p) random variable.

```
def draw_bin(n, p, N):
    """
    A Bin(n, p) pseudo-RNG
    """
    if N > 1:
        U = np.random.uniform(0, 1, (N, n))
        X = np.sum(U < p, axis = 1)

else:
        U = np.random.uniform(0, 1, n)
        X = np.sum(U < p)</pre>
```

```
return X

#print a few examples of the RGNs to the screen
n = 50
p = 0.2
print_vals(draw_bin, n, p, 1)

X_0 = 8
X_1 = 17
X_2 = 3
X_3 = 13
X_4 = 10
```

Note that we can directly sample from a Bin(n, p) distribution with Numpy's binomial RNG with: np.random.binomial(n, p).

Example 4. Write an algorithm to simulate the value of a random variable X such that:

$$P_X(x) = \begin{cases} 0.35 & \text{for } x = 1\\ 0.15 & \text{for } x = 2\\ 0.4 & \text{for } x = 3\\ 0.1 & \text{for } x = 4. \end{cases}$$

Solution: We can utilize the algorithm presented in the book which divides the unit interval into 4 partitioned sets and uses a uniformly drawn random variable.

```
def draw_general_discrete(P, R_X, N):
    """
    A pseudo-RNG for any arbitrary discrete PMF specified by R_X and
    corresponding probabilities P
    """
    F_X = np.cumsum([0] + P)

    X_arr = []
    U_arr = np.random.uniform(0, 1, size = N)
    for U in U_arr:
        X = R_X[np.sum(U > F_X)-1]

    #take care of edge case where U = 0
    if U == 0:
        X = R_X[0]
        X_arr.append(X)
    if N == 1: X_arr = X_arr[0]

    return X_arr
```

```
#print a few examples of the RGNs to the screen
P = [0.35, 0.15, 0.4, 0.1]
R_X = [1, 2, 3, 4]
print_vals(draw_general_discrete, P, R_X, 1)
```

```
X_0 = 2

X_1 = 4

X_2 = 3

X_3 = 3

X_4 = 4
```

Note that we can directly sample from a discrete PMF using Numpy's multinomial RNG. A multinomial distribution is the k dimensional analogue of a binomial distribution, where k > 2. The multinomial distribution is a distribution over random vectors, \boldsymbol{X} (of size k), where the entries in the vectors can take on values from $0, 1, \ldots, n$, subject to $X_1 + X_2 + \ldots + X_k = n$, where X_i represents the i^{th} component of \boldsymbol{X} .

If a binomial random variable represents the number of heads we flip out of n coin tosses (where the probability of heads is p), then a multinomial random variable represents the number of times we roll a 1, the number of times we roll a 2, ..., the number of times we roll a k, when rolling a k sided die n times. For each roll, the probability of rolling the i^{th} face of the die is p_i (where $\sum_{i=1}^{k} p_i = 1$). We store the value for the number times we roll the i^{th} face of the die in X_i . To denote a random vector drawn from a multinomial distribution, the notation, $X \sim Mult(n, p)$, is typical, where p denotes the k dimensional vector with the i^{th} component of p given by p_i .

To directly sample from a discrete PMF with (ordered) range array R_X and associated probability array P we can use Numpy's multinomial RNG function by setting n=1 (one roll). To sample one time we can use the code: $X = R_X[np.argmax(np.random.multinomial(1, pvals=P))]$, and to sample N times, we can use the code: $X = [R_X[np.argmax(x)]]$ for x in np.random.multinomial(1, pvals=P, size=N)].

Additionally, to sample from an arbitrary discrete PMF, we can also use Numpy's choice function, which samples randomly from a specified list, where each entry in the list is sampled according to a specified probability. To sample N values from an array R_X , with corresponding probability array P, we can use the code: $X = np.random.choice(R_X, size=N, replace=True, p=P)$. Make sure to specify replace=True to sample with replacement.

Example 5. (Exponential) Generate an Exp(1) random variable.

Solution: Using the method of inverse transformation, as shown in the book, for a strictly increasing CDF, F, the random variable $X = F^{-1}(U)$, where $U \sim Unif(0,1)$, has distribution $X \sim F$. Therefore, it is not difficult to show that,

$$-\frac{1}{\lambda}\ln(U) \sim Exp(\lambda),$$

where the fact that $1 - U \sim Unif(0,1)$ has been used.

```
def draw_exp(lam, N):
    """

An Exp(lambda) pseudo-RNG using the method of inverse transformation
    """

U = np.random.uniform(0, 1, size = N)
    if N == 1:
        U = U[0]
    X = (-1/lam)*np.log(U)

return X
```

```
#print a few examples of the RGNs to the screen
lam = 1
print_vals(draw_exp, lam, 1)

X_0 = 2.4838379957
X_1 = 0.593858616083
X_2 = 0.53703944167
X_3 = 0.0388069650697
X_4 = 1.23049637556
```

Note that we can directly sample from an $Exp(\lambda)$ distribution with Numpy's exponential RNG with: np.random.exponential(lam).

Example 6. (Gamma) Generate a Gamma(20,1) random variable.

Solution: If X_1, X_2, \ldots, X_n are drawn iid from an $Exp(\lambda)$ distribution, then $Y = X_1 + X_2 + \ldots + X_n \sim Gamma(n, \lambda)$. Therefore, to generate a $Gamma(n, \lambda)$ random variable, we need only to generate n independent $Exp(\lambda)$ random variables and add them.

```
def draw_gamma(alpha, lam, N):
    """

A Gamma(n, lambda) pseudo-RNG using the method of inverse transformation
    """

n = alpha
if N > 1:
    U = np.random.uniform(0, 1, size = (N, n))
    X = np.sum((-1/lam)*np.log(U), axis = 1)

else:
    U = np.random.uniform(0, 1, size = n)
    X = np.sum((-1/lam)*np.log(U))
return X
```

```
#print a few examples of the RGNs to the screen
alpha = 20
lam = 1
print_vals(draw_gamma, alpha, lam, 1)
```

```
X_0 = 17.4925086879

X_1 = 20.6155480241

X_2 = 26.9115218192

X_3 = 22.3654600391

X_4 = 22.331744631
```

Note that we can directly sample from a $Gamma(n, \lambda)$ distribution with Numpy's gamma RNG with: np.random.gamma(shape, scale).

Example 7. (Poisson) Generate a Poisson random variable. Hint: In this example, use the fact that the number of events in the interval [0,t] has Poisson distribution when the elapsed times between the events are Exponential.

Solution: As shown in the book, we need only to continuously generate $Exp(\lambda)$ variables

and count the number of draws it takes for the sum to be greater than 1. The Poisson random variable is then the count minus 1.

```
def draw_poiss(lam, N):
    A Poiss(lambda) pseudo-RNG
    X_list = []
    for _ in range(N):
        summ = 0
        count = 0
        while summ <= 1:
            summ += draw_exp(lam, 1)
            count += 1
        X_list.append(count-1)
    if N == 1:
        return X_list[0]
    else:
        return X_list
#print a few examples of the RGNs to the screen
print_vals(draw_poiss, lam, 1)
X O = O
X 1 = 2
X_2 = 2
X_3 = 1
X_4 = 2
```

Note that we can directly sample from a $Poiss(\lambda)$ distributions with Numpy's: np.random.poisson(lam) function.

Example 8. (Box-Muller) Generate 5000 pairs of normal random variables and plot both histograms.

Solution: Using the Box-Muller transformation as described in the book:

```
def draw_gaus_pairs(N):
    """

An N(O, 1) pseudo-RNG to draw N pairs of indepedent using the Box-Muller
    transformation
    """

U1 = np.random.uniform(size = N)
    U2 = np.random.uniform(size = N)

Z1 = np.sqrt(-2*np.log(U1))*np.cos(2*np.pi*U2)
    Z2 = np.sqrt(-2*np.log(U1))*np.sin(2*np.pi*U2)
```

```
return (Z1, Z2)

#print a few examples of the RGNs to the screen
Z1_arr, Z2_arr = draw_gaus_pairs(5)

for i, (Z1, Z2) in enumerate(zip(Z1_arr, Z2_arr)):
    print('(Z_1, Z_2)_' + str(i)+' = (', Z1, Z2, ')')

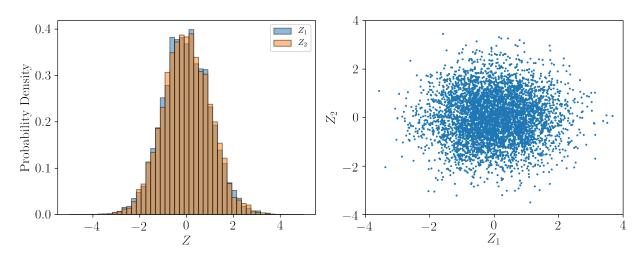
(Z_1, Z_2)_0 = (0.722134435205 -0.189448731182)
(Z_1, Z_2)_1 = (-0.918558147113 0.247330492682)
(Z_1, Z_2)_2 = (-1.42078058592 -0.914027516141)
(Z_1, Z_2)_3 = (1.19799155228 -1.49105841693)
(Z_1, Z_2)_4 = (-0.65055423687 0.179187077215)
```

In addition to plotting the histograms (plot in the first panel below) I also make a scatter plot of the 2 Gaussian random variables. The Box-Muller method produces pairs of independent random variables, and indeed, in the plot we see a bivariate Normal distribution with no correlation, i.e., it is axis-aligned (recall that independence $\implies \rho = 0$). I further compute the correlation coefficient between Z_1 and Z_2 and it is indeed very close to 0.

```
#plot the histograms and scatter plot
#set seed for reproducibility
np.random.seed(8)
#generate data
Z1_arr, Z2_arr = draw_gaus_pairs(5000)
#plot histograms
f, (ax1, ax2) = plt.subplots(1, 2, figsize=(10, 4))
bins = np.linspace(-5, 5, 50)
ax1.hist(Z1_arr, bins, alpha=0.5, normed=1, label='$Z_1$', edgecolor = 'black')
ax1.hist(Z2_arr, bins, alpha=0.5, normed=1, label='$Z_2$', edgecolor = 'black')
ax1.legend(loc='upper right')
ax1.set_xlabel('$Z$', size = 15)
ax1.set_ylabel('Probability Density', size = 15)
ax1.tick_params(labelsize=15)
#plot scatter plot
ax2.scatter(Z1_arr, Z2_arr, s=2)
ax2.set_xlabel('$Z_1$', size = 15)
ax2.set_ylabel('$Z_2$', size = 15)
ax2.set_ylim((-4, 4))
ax2.set_xlim((-4, 4))
ax2.tick_params(labelsize=15)
print('correlation coefficient = ', np.corrcoef(Z1_arr, Z2_arr)[0, 1])
#reset seed
```

np.random.seed(0)

correlation coefficient = 0.0177349514518



Note that we can directly sample from a $\mathcal{N}(0,1)$ distribution with Numpy's normal RNG with: np.random.randn(d0, d1, ..., dn), where d0, d1, ..., dn are the dimensions of the desired output array.

Exercise 1. Write Python programs to generate Geom(p) and Pascal(m, p) random variables.

Solution: As in the book, I generate Bern(p) random variables until the first success and count the number of draws to generate a Geom(p) random variable. To generate a Pascal(m,p) random variable, I generate Bern(p) random variables until I obtain m successes and count the number of draws.

```
def draw_geom(p, N):
    """
    A Geom(p) pseudo-RNG
    """
    X_list = []
    for _ in range(N):
        count = 0
        X = 0
        while X == 0:
            X = draw_bern(p, 1)
            count += 1
        X_list.append(count)

if N == 1:
    return X_list[0]
else:
    return X_list
```

```
#print a few examples of the RGNs to the screen
p = 0.2
print_vals(draw_geom, p, 1)
```

```
X_0 = 15
X_1 = 1
X_2 = 1
X_3 = 8
X_4 = 2
def draw_pascal(m, p, N):
    A Pascal(m, p) pseudo-RNG
    X_list = []
    for _ in range(N):
        count_succ = 0
        count = 0
        while count_succ < m:</pre>
            X = draw_bern(p, 1)
            count_succ += X
            count += 1
        X_list.append(count)
    if N == 1:
        return X_list[0]
    else:
        return X_list
#print a few examples of the RGNs to the screen
p = 0.2
m = 2
print_vals(draw_pascal, m, p, 1)
X_0 = 17
X_1 = 10
X_2 = 7
X_3 = 3
X_4 = 4
```

Note that we can directly sample from Geom(p) and Pascal(m,p) distributions with Numpy's np.random.geometric(p) and np.random.negative_binomial(n, p) functions respectively.

Exercise 2. (Poisson) Use the algorithm for generating discrete random variables to obtain a Poisson random variable with parameter $\lambda = 2$.

Solution:

```
from scipy.misc import factorial

def draw_poiss2(lam, N):
    """
    A Poiss(lambda) pseudo-RNG using the method to generate an
    arbitrary discrete random variable
    """
    X_list = []
```

```
for _ in range(N):
    P = np.exp(-lam)
    i = 0
    U = np.random.uniform()
    while U >= P:
        i += 1
        P += np.exp(-lam)*lam**i/(factorial(i)+0)

    X_list.append(i)

if N == 1:
    return X_list[0]

else:
    return X_list
```

```
#print a few examples of the RGNs to the screen
lam = 2
print_vals(draw_poiss2, 2, 1)
```

 $X_0 = 3$ $X_1 = 0$ $X_2 = 5$ $X_3 = 2$ $X_4 = 5$

Exercise 3. Explain how to generate a random variable with the density

$$f(x) = 2.5x\sqrt{x}$$

for 0 < x < 1.

Solution: The CDF is given by $F_X(x) = 2.5 \int_0^x x'^{3/2} dx' = x^{5/2}$, and therefore $F_X^{-1}(x) = x^{2/5}$. Using the method of inverse transformation, if $U \sim Unif(0,1)$, then $F_X^{-1}(U)$ is distributed according to the desired distribution.

```
def draw_dist3():
    """

A pseudo-RNG for the distribution in Exercise 3
    """

U = np.random.uniform()
    return U**(0.4)
```

```
#print a few examples of the RGNs to the screen
print_vals(draw_dist3)
```

 $X_0 = 0.8178201131579468$ $X_1 = 0.8861754700680049$ $X_2 = 0.27369087549414306$ $X_3 = 0.6033871249144047$ $X_4 = 0.4285059109745954$ Exercise 4. Use the inverse transformation method to generate a random variable having distribution function

$$F_X(x) = \frac{x^2 + x}{2},$$

for $0 \le x \le 1$.

Solution: By inverting the CDF, we have that.

$$F_X^{-1}(x) = -\frac{1}{2} + \sqrt{\frac{1}{4} + 2x},$$

for $0 \le x \le 1$.

```
def draw_dist4():
    """

A pseudo-RNG for the distribution in Exercise 4
    """

U = np.random.uniform()
    return -0.5 + np.sqrt(0.25 + 2*U)
```

#print a few examples of the RGNs to the screen
print_vals(draw_dist4)

 $X_0 = 0.417758353296$ $X_1 = 0.198180089883$

 $X_2 = 0.441257859881$ $X_3 = 0.538521058539$

 $X_4 = 0.115056902$

Exercise 5. Let X have a standard Cauchy distribution. function

$$F_X(x) = \frac{1}{\pi} \arctan x + \frac{1}{2}.$$

Assuming you have $U \sim Unif(0,1)$, explain how to generate X. Then, use this result to produce 1000 samples of X and compute the sample mean. Repeat the experiment 100 times. What do you observe and why?

Solution: The inverse CDF is given by $F_X^{-1}(x) = \tan[\pi(x - 1/2)]$.

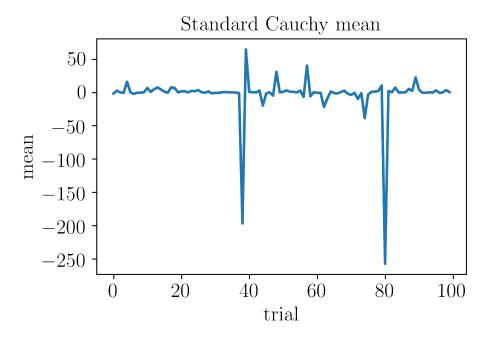
```
def draw_stand_cauchy(N):
    """

A standard Cauchy pseudo-RNG using the method of inverse transformation
    """

U = np.random.uniform(size = N)
    X = np.tan(np.pi*(U - 1/2))

if N == 1: return X[0]
    else: return X
```

```
#print a few examples of the RGNs to the screen
print_vals(draw_stand_cauchy, 1)
X 0 = 0.691013110859
X_1 = 0.212342443875
X_2 = -0.907695727473
X_3 = 0.0731660554841
X_4 = -3.28946953204
#plot means for 100 trials
#set seed for reproducibility
np.random.seed(5)
#compute means and plot
means = [np.mean(np.array(draw_stand_cauchy(1000))) for _ in range(100)]
plot_results(range(100), means, xlabel='trial', ylabel='mean', \
             title='Standard Cauchy mean')
#reset seed
np.random.seed(0)
```



We see that the means for each trial vary wildly. This is because the Cauchy distribution actually has no mean.

Exercise 6. (The Rejection Method) When we use the Inverse Transformation Method, we need a simple form of the CDF, F(x), that allows direct computation of $X = F^{-1}(U)$. When F(x) doesn't have a simple form but the PDF, f(x), is available, random variables with density f(x) can be generated by the rejection method. Suppose you have a method for generating a random variable having density function g(x). Now, assume you want to generate a random variable having

density function f(x). Let c be a constant such that $f(y)/g(y) \leq c$ (for all y). Show that the following method generates a random variable, X, with density function f(x).

1) initialize U and Y such that $U > \frac{f(Y)}{cg(Y)}$

repeat until $U \leq \frac{f(Y)}{cg(Y)}$ {

- 2) Generate Y having density g
- 3) Generate a random number U from Unif(0,1)

4) Set X = Y

Solution:

}

Firstly, as a technical matter, note that $c \ge 1$, which can be shown by integrating both sides of $f(y) \le cg(y)$.

We see that this algorithm keeps iterating until it outputs a random variable Y, given that we know that $U \leq \frac{f(Y)}{cg(Y)}$. Therefore, the goal is to show that the random variable $Y|U \leq \frac{f(Y)}{cg(Y)}$ has PDF f(y) (or equivalently CDF F(y)). In other words, we must show that $P\left(Y \leq y \middle| U \leq \frac{f(Y)}{cg(Y)}\right) = F(y)$. I show this with Baye's rule:

$$\begin{split} P\left(Y \leq y \middle| U \leq \frac{f(Y)}{cg(Y)}\right) &= \frac{P\left(U \leq \frac{f(Y)}{cg(Y)}\middle| Y \leq y\right) P(Y \leq y)}{P\left(U \leq \frac{f(Y)}{cg(Y)}\right)} \\ &= \frac{P\left(U \leq \frac{f(Y)}{cg(Y)}\middle| Y \leq y\right) G(y)}{P\left(U \leq \frac{f(Y)}{cg(Y)}\right)}. \end{split}$$

Thus, we must calculate the quantities: $P\left(U \leq \frac{f(Y)}{cg(Y)} | Y \leq y\right)$ and $P\left(U \leq \frac{f(Y)}{cg(Y)}\right)$. As an intermediate step, note that

$$\begin{split} P\left(U \leq \frac{f(Y)}{cg(Y)} \middle| Y = y\right) &= P\left(U \leq \frac{f(y)}{cg(y)} \middle| Y = y\right) \\ &= P\left(U \leq \frac{f(y)}{cg(y)}\right) \\ &= F_U\left(\frac{f(y)}{cg(y)}\right) \\ &= \frac{f(y)}{cg(y)}, \end{split}$$

where in the second line I have used that U and Y are independent and in the fourth I have used the fact that for a uniform distribution $F_U(u) = u$. Notice that the requirement that $f(y)/g(y) \le c$ (for all y) is crucial at this step. This is because $f(y)/g(y) \le c \implies c > 0$ (since f(y) and g(y) are positive), so that $0 < f(y)/cg(y) \le 1$. If this condition did not hold, then the above expression

would be $\min\{1, \frac{f(y)}{cg(y)}\}\$, for positive c and 0 for negative c, which would interfere with the rest of the derivation.

I may now calculate $P\left(U \leq \frac{f(Y)}{cg(Y)}\right)$:

$$\begin{split} P\left(U \leq \frac{f(Y)}{cg(Y)}\right) &= \int_{-\infty}^{\infty} P\left(U \leq \frac{f(Y)}{cg(Y)} \middle| Y = y\right) g(y) dy \\ &= \int_{-\infty}^{\infty} \frac{f(y)}{cg(y)} g(y) dy \\ &= \frac{1}{c} \int_{-\infty}^{\infty} f(y) dy \\ &= \frac{1}{c}. \end{split}$$

I now calculate the remaining quantity:

$$\begin{split} P\left(U \leq \frac{f(Y)}{cg(Y)}|Y \leq y\right) &= \frac{P\left(U \leq \frac{f(Y)}{cg(Y)}, Y \leq y\right)}{G(y)} \\ &= \frac{\int_{-\infty}^{\infty} P\left(U \leq \frac{f(Y)}{cg(Y)}, Y \leq y|Y = v\right)g(v)dv}{G(y)} \\ &= \frac{\int_{-\infty}^{\infty} P\left(U \leq \frac{f(Y)}{cg(Y)}|Y \leq y, Y = v\right)P(Y \leq y|Y = v)g(v)dv}{G(y)}, \end{split}$$

where in the second line I have used the law of total probability, and in the third line I have used the definition of conditional probability. Note that:

$$P(Y \le y | Y = v) = \begin{cases} 1 & \text{for } v \le y \\ 0 & \text{for } v > y, \end{cases}$$

and thus

$$\begin{split} P\left(U \leq \frac{f(Y)}{cg(Y)}|Y \leq y\right) &= \frac{\int_{-\infty}^{y} P\left(U \leq \frac{f(Y)}{cg(Y)}|Y \leq y, Y = v\right)g(v)dv}{G(y)} \\ &= \frac{\int_{-\infty}^{y} P\left(U \leq \frac{f(Y)}{cg(Y)}|Y = v\right)g(v)dv}{G(y)} \\ &= \frac{\int_{-\infty}^{y} \frac{f(v)}{cg(v)}g(v)dv}{G(y)} \\ &= \frac{\frac{1}{c}F(y)}{G(y)}, \end{split}$$

where in the second line I have used the fact that conditioning on Y=v already implies that $Y \leq y$ since we only consider values of v less than or equal to y in the integration. In the third line I have used the expression for $P\left(U \leq \frac{f(Y)}{cg(Y)} \middle| Y=y\right)$ that we derived above.

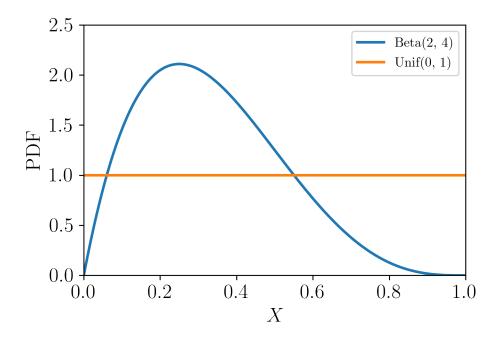
Inserting these quantities into Baye's rule:

$$\begin{split} P\left(Y \leq y \middle| U \leq \frac{f(Y)}{cg(Y)}\right) &= \frac{P\left(U \leq \frac{f(Y)}{cg(Y)}\middle| Y \leq y\right)G(y)}{P\left(U \leq \frac{f(Y)}{cg(Y)}\right)} \\ &= \frac{\frac{\frac{1}{c}F(y)}{G(y)}G(y)}{\frac{1}{c}} \\ &= F(y), \end{split}$$

which is what we set out to prove.

Exercise 7. Use the rejection method to generate a random variable having density function Beta(2,4). Hint: Assume g(x) = 1 for 0 < x < 1.

Solution: I first visualize these distributions so we can get a handle on what we are dealing with.



Since f(x)/g(x) (where f(x) is the PDF of the Beta and g(x) is the PDF of the uniform) needs to be smaller than c for all x in the support of these distributions, a fine value of c to use would be 2.5 since it is evident from the plot that this value satisfies the requirement. The book uses the

smallest possible value of c, i.e., the max of the Beta(2,4) distribution, which it derives analytically and finds to be $135/64 \approx 2.11$. It is not necessary to use the smallest value of c, but will certainly help the speed of the algorithm since the algorithm only stops when $U \leq f(Y)/cg(Y)$. I will stick with the value of 2.5 just to illustrate that the algorithm works for this value as well.

```
def draw beta 2 4(N):
    11 11 11
    A Beta(2, 4) pseudo-RNG using the rejection method
    c = 2.5
    X_list = []
    for _ in range(N):
        U = 1
        f_Y = 0
        g_Y = 1
        while U > f_Y/(c*g_Y):
            Y = np.random.uniform()
            U = np.random.uniform()
            f_Y = 20*Y*(1-Y)**3
            g_Y = 1
        X_list.append(Y)
    if N == 1:
        return X_list[0]
    else:
        return X_list
```

#print a few examples of the RGNs to the screen
print_vals(draw_beta_2_4, 1)

```
X_0 = 0.4236547993389047

X_1 = 0.07103605819788694

X_2 = 0.11827442586893322

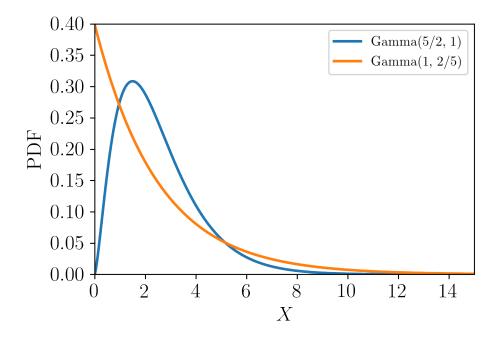
X_3 = 0.5218483217500717

X_4 = 0.26455561210462697
```

Note that we can directly sample from a $Beta(\alpha, \beta)$ distribution with Numpy's beta RNG with: np.random.beta(a, b).

Exercise 8. Use the rejection method to generate a random variable having the Gamma(5/2, 1) density function. Hint: Assume g(x) is the PDF of the Gamma(1, 2/5).

Solution: Note that there is a mistake in the phrasing of the question in the book. The PDF for g(x) should be Gamma(1,2/5), not Gamma(5/2,1). Also note that we cannot use the method that we used in **Example**, 6. since in this case α is not an integer (however, we can use that method to draw from g(x)). I first visualize these distributions so we can get a handle on what we are dealing with.



The $\max\{f(x)/g(x)\}\$ for x>0 is approximately given by:

np.max(f/g)

1.6587150033103788

As a sanity check, this value is very close to the analytically derived value in the book, which is $\frac{10}{3\sqrt{\pi}}\left(\frac{5}{2}\right)^{3/2}e^{-3/2}\approx 1.6587162$. Therefore, I set the value of c to be 1.7, and use the function I wrote in **Example**. 6, draw_gamma(alpha, lam, N), to draw from g(x).

```
def draw_gamma_2(alpha, lam, N):
    """

A Gamma(5/2, 1) pseudo-RNG using the rejection method
    """

c = 1.7

X_list = []
  for _ in range(N):

U = 1
  f_Y = 0
```

```
g_Y = 1
        while U > f_Y/(c*g_Y):
            Y = draw_gamma(1, 0.4, 1)
            U = np.random.uniform()
            f_Y = (4/(3*np.sqrt(np.pi)))*(Y**1.5)*np.exp(-Y)
            g_Y = 0.4*np.exp(-0.4*Y)
        X_list.append(Y)
    if N == 1:
        return X_list[0]
    else:
        return X_list
#print a few examples of the RGNs to the screen
print_vals(draw_gamma_2, 5/2, 1, 1)
X_0 = 1.96233211971
X_1 = 1.22716649756
X_2 = 2.55754781375
X_3 = 0.900161721137
X_4 = 3.89706921546
```

Exercise 9. Use the rejection method to generate a standard normal random variable. Hint: Assume g(x) is the PDF of the exponential distribution with $\lambda = 1$.

Solution As in the book, to solve this problem, I use the rejection method to sample from a half Gaussian:

$$f(x) = \frac{2}{\sqrt{2\pi}} e^{-\frac{x^2}{2}},$$

with range $(0, \infty)$, with an Exp(1) distribution for g(x). The book analytically computes $\max\{f(x)/g(x)\}$ to be $\sqrt{2e/\pi}\approx 1.32$, and I thus use c=1.4. Once the algorithm is able to sample from the half Gaussian, to turn this distribution into a full Gaussian with range \mathbb{R} , one need only to randomly multiply by -1. I therefore sample $Q \in \{0,1\}$ from a Bern(0.5) distribution and multiply by 1-2Q $\in \{-1,1\}$ in order to sample from the full Gaussian.

```
def draw_standard_normal(N):
    """
    A standard normal pseudo-RNG using the rejection method
    """
    c = 1.4

    X_list = []
    for _ in range(N):
        U = 1
```

```
f_Y = 0
g_Y =1

while U > f_Y/(c*g_Y):
    Y = draw_exp(1, 1)
    U = np.random.uniform()

    f_Y = (2/np.sqrt(2*np.pi))*np.exp(-(Y**2)/2)
    g_Y = np.exp(-Y)

# draw Bern(0.5) random variable for the sign
Q = draw_bern(0.5, 1)

X_list.append(Y*(1-2*Q))

if N == 1:
    return X_list[0]
else:
    return X_list
```

#print a few examples of the RGNs to the screen
print_vals(draw_standard_normal, 1)

```
X_0 = 1.1538237197

X_1 = -2.28234324111

X_2 = -0.426012274543

X_3 = -1.40884434358

X_4 = -0.421092193245
```

Exercise 10. Use the rejection method to generate a Gamma(2,1) random variable conditional on its value being greater than 5. Hint: Assume g(x) is the density function of exponential distribution.

Solution As in the book, I use an Exp(0.5) conditioned on X > 5 as the distribution for g(x). It is not difficult to show by integrating the PDF of this distribution that $G^{-1}(x) = 5 - 2\ln(1-x)$ (where G is the CDF). I therefore use the method of inverse transformation to first draw a random variable from this distribution (Y). Note that for $U \sim Unif(0,1)$, $1-U \sim Unif(0,1)$, and therefore the formula for $G^{-1}(U)$ can be simplified to $5-2\ln(U)$. I then use the rejection method to sample from the desired distribution. By maximizing f(x)/g(x), the book shows that c must be greater than 5/3, and I therefore use c = 1.7.

```
def draw_gamma_2_1_cond_5(N):
    """

A Gamma(2, 1) conditional on X>5 pseudo-RNG using the rejection method
    """

c = 1.7

X_list = []
for _ in range(N):
    U = 1
```

```
f_Y = 0
g_Y =1

while U > f_Y/(c*g_Y):
    Y = 5 - 2*np.log(np.random.uniform())
    U = np.random.uniform()

    f_Y = Y*np.exp(5-Y)/6
    g_Y = np.exp((5-Y)/2)/2

X_list.append(Y)

if N == 1:
    return X_list[0]
else:
    return X_list
```

```
#print a few examples of the RGNs to the screen
print_vals(draw_gamma_2_1_cond_5, 1)
```

```
X_0 = 6.76250850879

X_1 = 5.73497460514

X_2 = 5.14665551227

X_3 = 5.8087003199

X_4 = 5.66723645483
```

Notice that, as required, the random variables are all > 5.

As a final check to close this chapter, I draw samples from most of the RNG functions that I implemented above, compute the corresponding PMFs/PDFs, and compare to the theoretical distributions. I first check the discrete distributions, and I start by writing a function that will compute the empirical PMFs. Note that the phrase, "empirical PMF", (and "empirical PDF") is standard terminology to refer to the probability distribution associated with a sample of data. Formally, for a collection of data, $\{x_i\}_{i=1}^N$, they are given by

$$P_X(x) = \frac{1}{N} \sum_{i=1}^{N} \mathbf{I}(x = x_i)$$

for the empirical PMF, and by

$$f_X(x) = \frac{1}{N} \sum_{i=1}^{N} \delta(x = x_i)$$

for the empirical PDF (where $I(\cdot)$ is the indicator function, and $\delta(\cdot)$ is the delta function).

```
def compute_PMFs(counts, xrange):
    """
    Compute empirical PMFs from a specified array of random variables,
    and a specified range
    """
```

```
count_arr = []
xrange2 = range(np.max([np.max(xrange), np.max(counts)])+1)
for i in xrange2:
    count_arr.append(np.sum(counts==i))
pmf = np.array(count_arr)/np.sum(np.array(count_arr))
return pmf[np.min(xrange):np.max(xrange)+1]
```

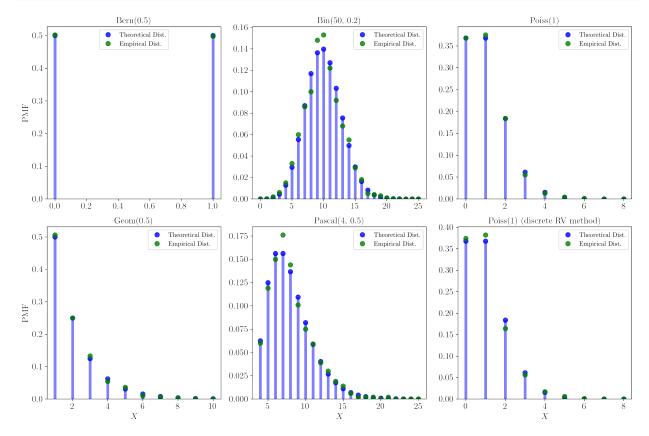
I now compute the theoretical distributions, generate the data and compute the empirical distributions.

```
from scipy.stats import bernoulli, binom, poisson, geom, nbinom
#set seed for reproducibility
np.random.seed(1984)
x_ranges = [range(2), range(26), range(9), range(1, 11), range(4, 26), range(9)]
#compute PMF arrays for the theoretical distributions
numpy dists = [bernoulli, binom, poisson, geom, nbinom, poisson]
numpy_args = [[0.5], [50, 0.2], [1], [0.5], [4, 0.5, 4], [1]]
numpy_y = [np_dist.pmf(xrange, *np_args) for np_dist, xrange, np_args in \
           zip(numpy_dists, x_ranges, numpy_args)]
N = 1000 #number of points to sample
# draw random variables from my functions and compute corresponding PMFs
my_rngs = [draw_bern, draw_bin, draw_poiss, draw_geom, draw_pascal, draw_poiss2]
my_{args} = [[0.5, N], [50, 0.2, N], [1, N], [0.5, N], [4, 0.5, N], [1, N]]
my_counts = [rng(*args) for rng, args in zip(my_rngs, my_args)]
my_y = [compute_PMFs(np.array(counts), xrange) for counts, xrange in \
        zip(my_counts, x_ranges)]
```

Finally, I plot the results.

```
ax.set_title(names[i], size = 15)
ax.set_ylim(ymin=0)
ax.tick_params(labelsize=15)

if i in [3, 4, 5]:
    ax.set_xlabel('$X$', size = 15)
if i in [0, 3]:
    ax.set_ylabel('PMF', size = 15)
```



We see that the empirical distributions match almost perfectly with the theoretical distributions, with even better correspondence for larger N.

I now check some of the continuous RNG functions that I implemented in this chapter. I first start by computing the theoretical distributions and generating the data.

I now plot normalized histograms of the data and compare to the theoretical distributions. Again, we see almost perfect correspondence between the empirical and theoretical distributions. The correspondence becomes even better with larger values of N.

```
#plot theoretical and empirical PDFs
names = ['Exp(1) (inverse trans.)', 'Gamma(20, 1) (inverse trans.)', \
         'Cauchy(0, 1) (inverse trans.)', 'Beta(2, 4) (rejection)', \
         'Gamma(5/2, 1) (rejection)', 'N(0, 1) (rejection)']
bin_arr = [50, 35, 60, 45, 45, 35]
xlims=[(0, 8), (0, 50), (-20, 20), (0, 1), (0, 15), (-5, 5)]
range_arr = [None] *6
range_arr[2] = (-20, 20)
f, [[ax1, ax2, ax3], [ax4, ax5, ax6]] = plt.subplots(2, 3, figsize=(15, 10))
ax_arr = [ax1, ax2, ax3, ax4, ax5, ax6]
for i, ax in enumerate(ax_arr):
    ax.plot(x_ranges[i], numpy_y[i], label='Theoretical Dist.', color='black', \
            linewidth=3, alpha=.7)
    ax.hist(my_rvs[i], bins=bin_arr[i], alpha=.5, edgecolor='black',normed=True, \
            label='Empirical Dist.', range=range_arr[i])
    ax.set_title(names[i], size = 15)
    ax.legend(loc='upper right', fontsize=10)
    ax.set_xlim(xlims[i])
    ax.tick_params(labelsize=15)
    if i in [3, 4, 5]:
        ax.set_xlabel('$X$', size = 15)
    if i in [0, 3]:
        ax.set_ylabel('PDF', size = 15)
```

