Claim: 
$$P(X>t+s|X>t) = P(X>s)$$
  $t = 0$ 

$$P(X>t+s|X>t) = P(X>t)$$

$$P(X>t+s|X>t) = P(X>t)$$

$$P(X>t+s|X>t) = P(X>t+s) \cap (X>t)$$

$$P(X>t+s) = P(X>t+s) = P(X>t+s)$$

$$P(X>t+s) = P(X>t+s)$$

$$P(X>t+s) = P(X>t+s)$$

$$P(X>t+s) = P(X>t+s)$$

$$P(X > t + s \mid X > t) = P(X > t + s)$$

$$P(X > t + s)$$

$$P(X > t + s)$$

$$A e^{-\lambda X} dx$$

$$+ e^{-\lambda X} dx$$

$$+ e^{-\lambda X} dx$$

$$+ e^{-\lambda X} dx$$

$$|P(X75) = \int_{S} |e^{-\lambda x} dx = -e^{-\lambda x}| = -(0 - e^{-\lambda 5})$$

e)
$$EX = \int x f_{\chi}(x) dx = \int x dx = \frac{1}{2} x^{2} / \frac{1}{2} = \frac{1}{2}$$

$$EY = \int y f_{\chi}(y) dy = -\int y \ln(y) dy \qquad u = \ln y \quad du = -\int y dy$$

$$V = \frac{1}{2} y^{2} \quad dv = y dy$$

$$\frac{E}{y} = \int y f_y(y) dy = -\int y \ln y dy$$

$$= \frac{1}{2}y^2 \ln y - \int \frac{1}{2}y^2 dy$$

$$= \frac{1}{2}y^2 \ln y - \int \frac{1}{2}y^2 dy$$

$$= \frac{1}{2}y^2 \ln y - \int \frac{1}{2}y^2 dy$$

$$= \frac{1}{2}y^{2}\ln y - \frac{1}{2}\int y dy = \frac{1}{2}y^{2}\ln y - \frac{1}{4}y^{2}$$

$$= \left[\frac{1}{2}\left(-\frac{1}{2}\ln y^{2}\ln y\right) - \frac{1}{4}\right] = -\frac{1}{2}\lim_{y \to 0} \frac{\ln y}{y^{2}} - \frac{1}{4}$$

$$= -\frac{1}{2} \lim_{y \to 0} \frac{y^{-1}}{-2y^{-3}} - \frac{1}{4} = -\frac{1}{2} \lim_{y \to 0} -2y^{2} - \frac{1}{4} = \frac{1}{4}$$

```
import numpy as np
import matplotlib.pyplot as plt
import math
import bisect
from collections import defaultdict
from typing import List
```

# Rejection sampling

In this worksheet you'll be doing the following tasks:

- 1. Define a function to sample from a discrete distribution.
- 2. Define classes for continuous and discrete random variables (provided).
- 3. Define a function to sample from a mixture distribution.
- 4. Define a rejection sampling function.
- 5. Test your functions on some examples.

After this assignment, we'll extract this code into a separate module and keep importing it in the future assignments.

## Discrete distribution

- 1. To start with, copy your RNG class from the previous assignment below. Make sure to delete the unimplemented mixture method if one still exists there.
- 2. Also, the method <code>np.random.random()</code> returns a single random number (and not a list with a single element). Make sure this is the same behavior in your <code>RNG</code> class for all the sampling methods. If you simply called the method <code>self.uniform</code> and used list comprehension you should be fine. If you wrote your own loop then make sure to return a single number and not a list with a single element if <code>size=None</code> (not when <code>size=1</code>). I ran into this issue when testing my code.
- 3. Then define a method discrete in the RNG class that takes a list of (unnormalized) probabilities and a list of values and returns a random sample from the corresponding discrete distribution. For example, if the input is weights= [2, 8] and list=[3, 5] then the output should be 3 with probability 0.2 and 5 with probability 0.8.

```
In [102... class RNG:
    def __init__(self, seed=None):
        self._rng = np.random.default_rng(seed)
        self.random = self._rng.random

def seed(self, seed=None):
        self._rng = np.random.default_rng(seed)
```

```
def uniform(self, low=0.0, high=1.0, size=None):
    return low + (high - low) * self.random(size)
# choose a random object from a list
def choice(self, list_, size=None):
    u = self.random(size)
    return list_[np.floor(u * len(list_)).astype(int)]
def discrete(self, weights, list =None, size=None):
    total weight = sum(weights)
    intervals = np.cumsum(weights) / total_weight
    u samples = self.random(size)
    return [list_[bisect.bisect_left(intervals, s)] for s in u_samples]
def bernoulli(self, p=0.5, size=None):
    assert 0 <= p <= 1, "p must be between 0 and 1"</pre>
    u = self.random(size)
    return (u < p).astype(int)</pre>
def binomial(self, n=1, p=0.5, size=None):
    assert 0 <= p <= 1, "p must be between 0 and 1"</pre>
    sample = lambda x, y: np.sum(self.bernoulli(x, y))
    return np.array([sample(p, n) for _ in range(size)])
def exponential(self, lambda_=1.0, size=None):
    assert lambda > 0, "lambda must be positive"
    return -1/lambda_ * np.log(1 - self.uniform(size=size))
def normal(self, mean=0.0, std=1.0, size=None):
    R = np.sqrt(-2 * np.log(1 - self.uniform(size=size))) # u1
    Theta = 2 * np.pi * self.uniform(size=size) # u2
    return mean + std * R * np.cos(Theta) # element-wise multiplication
def poisson(self, lambda =1.0, size=None):
    assert lambda_ > 0, "lambda must be positive"
    samples = np.zeros(size)
    for i in range(size):
        N, sum = 0, 0
        while True:
            X = self.exponential(lambda_, size=1)
            sum += X
            if sum > 1:
                break
            N += 1
        samples[i] = N
    return samples
def beta(self, a=1.0, b=1.0, size=None):
    assert a > 0 and b > 0, "a and b must be positive"
    assert isinstance(a, int) and isinstance(b, int), "a and b must be i
    k, n = a, b + a - 1
    samples = np.array([self.uniform(size=n) for _ in range(size)])
    samples.sort(axis=1)
    return np.array([row[k-1] for row in samples])
```

```
def triangular(self, low=0.0, high=1.0, mode=None, size=None):
    assert low <= mode <= high, "low <= mode <= high"
    samples = self.uniform(size=size)
    samples.sort()
    crit = (mode - low) / (high - low)
    return np.where(
        samples < crit,
        low + np.sqrt(samples * (high - low) * (mode - low)),
        high - np.sqrt((1 - samples) * (high - low) * (high - mode))
    )

def weibull(self, shape=1.0, scale=1.0, size=None):
    assert shape > 0, "shape must be positive"
    assert scale > 0, "scale must be positive"
    return scale * (-np.log(1 - self.uniform(size=size)))**(1/shape)
```

# Test your discrete sampling function

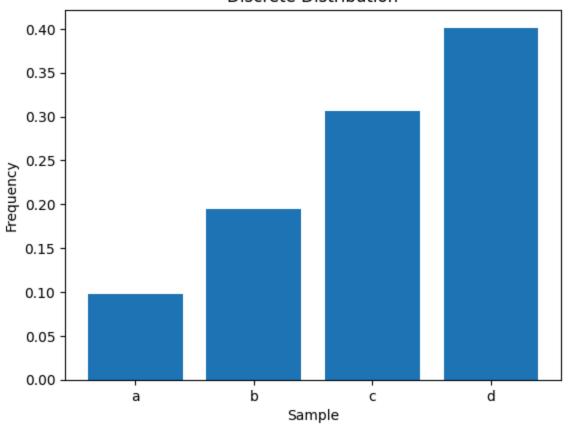
```
In [102... # test discrete distribution
    rng.seed(0)
    size = 10000

samples = rng.discrete(weights=[0.1, 0.2, 0.3, 0.4], list_=['a', 'b', 'c', 'unique, counts = np.unique(samples, return_counts=True)
    frequencies = counts / len(samples)

epsilon = 0.001 # allowance for floating point errors
    assert abs(np.sum(frequencies) - 1) < epsilon, "Probabilities do not sum to

# check that the samples are generated correctly
    plt.bar(unique, frequencies)
    plt.xlabel('Sample')
    plt.ylabel('Frequency')
    plt.title('Discrete Distribution')
    plt.show()</pre>
```

#### Discrete Distribution



## Continuous random variables

Below we define classes for continuous random variables that bundle the sampling methods together with the density functions. For now we only define the pdf method but for a full implementation we would need many more methods such as cdf, ppf, mean, var, etc. See the documentation of scipy.stats.norm class for an example. These classes are provided for you and you don't need to do anything with them. You can move on to the Mixture class.

```
In [102... class RandomVariable():
    def __init__(self, rng):
        self.rng = rng

    def pdf(self, x):
        raise NotImplementedError

    def rsv(self, size=None):
        raise NotImplementedError

class Uniform(RandomVariable):
    def __init__(self, rng, low=0.0, high=1.0):
        super().__init__(rng)
        self.low = low
        self.high = high
```

```
def rsv(self, size=None):
        return np.array(self.rng.uniform(low=self.low, high=self.high, size=
   def pdf(self, x):
        return np.array(np.where((x >= self.low) & (x <= self.high), 1 / (se
class Beta(RandomVariable):
   def __init__(self, rng, a=1.0, b=1.0):
        if not isinstance(a, int) or not isinstance(b, int):
            raise ValueError("a and b must be integers")
        super().__init__(rng)
        self.a = a
        self.b = b
   def rsv(self, size=None):
        return np.array(self.rng.beta(a=self.a, b=self.b, size=size))
   def pdf(self, x):
        return np.array(x**(self.a - 1) * (1 - x)**(self.b - 1) / self.beta(
   def beta(self, a, b):
        return math.factorial(a - 1) * math.factorial(b - 1) / math.factoria
class Exponential(RandomVariable):
   def __init__(self, rng, lambda_=1.0):
        super(). init (rng)
        self.lambda_ = lambda_
   def rsv(self, size=None):
        return np.array(self.rng.exponential(lambda_=self.lambda_, size=size
   def pdf(self, x):
        return np.array(self.lambda_ * np.exp(-self.lambda_ * x))
class Normal(RandomVariable):
   def __init__(self, rng, mean=0.0, std=1.0):
        super().__init__(rng)
        self.mean = mean
        self.std = std
   def rsv(self, size=None):
        return np.array(self.rng.normal(mean=self.mean, std=self.std, size=s
   def pdf(self, x):
        return np.array(np.exp(-0.5 * ((x - self.mean) / self.std)**2) / np.
class Bernoulli(RandomVariable):
   def __init__(self, rng, p=0.5):
        super().__init__(rng)
        self.p = p
   def rsv(self, size=None):
        return np.array(self.rng.bernoulli(p=self.p, size=size))
   def pdf(self, x):
        return np.array(np.where(x == 1, self.p, 1 - self.p))
```

```
class Poisson(RandomVariable):
   def __init__(self, rng, lambda_=1.0):
        super().__init__(rng)
        self.lambda_ = lambda_
   def rsv(self, size=None):
        return np.array(self.rng.poisson(lambda_=self.lambda_, size=size))
   def pdf(self, x):
        return np.array(self.lambda_**x * np.exp(-self.lambda_) / math.facto
class Binomial(RandomVariable):
   def __init__(self, rng, n=1, p=0.5):
        super().__init__(rng)
        self.n = n
        self.p = p
   def rsv(self, size=None):
        return np.array(self.rng.binomial(n=self.n, p=self.p, size=size))
   def pdf(self, x):
        return np.array(math.factorial(self.n) / (math.factorial(x) * math.f
class Weibull(RandomVariable):
   def init (self, rnq, shape=1.0, scale=1.0):
        super().__init__(rng)
        self.shape = shape
        self.scale = scale
   def rsv(self, size=None):
        return np.array(self.rng.weibull(shape=self.shape, scale=self.scale,
   def pdf(self, x):
        return np.array((self.shape / self.scale) * (x / self.scale)**(self.
class Triangular(RandomVariable):
   def __init__(self, rng, low=0.0, high=1.0, mode=None):
        super().__init__(rng)
        self.low = low
        self.high = high
        self.mode = mode if mode else (low + high) / 2
   def rsv(self, size=None):
        return np.array(self.rng.triangular(low=self.low, high=self.high, mc
   def pdf(self, x):
        return np.array(np.where((x >= self.low) & (x < self.mode),</pre>
                                 2 * (x - self.low) / ((self.high - self.low))
                                 2 * (self.high - x) / ((self.high - self.ld)
class Discrete(RandomVariable):
   def __init__(self, rng, weights, list_=None):
        super().__init__(rng)
        self.weights = weights
        self.list = list
```

```
def rsv(self, size=None):
    return np.array(self.rng.discrete(weights=self.weights, list_=self.l)

def pdf(self, x):
    return np.array(self.weights[self.list_.index(x)])
```

### Mixture Distribution

Next, define a class Mixture that takes a list of weights and a list of component random variables. The class should have a method rsv that samples a random variable from the mixture distribution and a method pdf that evaluates the density function of the mixture distribution. Wrap your rsv output in a np.array (see above) to make sure it is a numpy array as numpy functions do not work well with lists.

```
class Mixture(RandomVariable):
    def __init__(self, rng, weights, components: List[RandomVariable]):
        assert len(weights) == len(components), "weights and components must super().__init__(rng)
        self.weights = weights
        self.components = components

def rsv(self, size=None):
        sampled_components = Discrete(self.rng, self.weights, self.component return np.array([comp.rsv() for comp in sampled_components])

def pdf(self, x):
        return np.array(sum([w * c.pdf(x) for w, c in zip(self.weights, self.components))
```

# Test your mixture distribution

```
In [102... # test mixture of normals
    rng.seed(0)
    size = 10000

    normal1 = Normal(rng, mean=-2, std=1)
    normal2 = Normal(rng, mean=3, std=1)

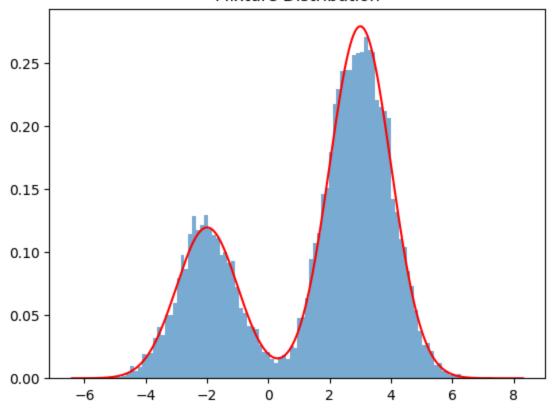
    mixture = Mixture(rng, weights=[0.3, 0.7], components=[normal1, normal2])

    samples = mixture.rsv(size=size)

# histogram of mixture distribution
    plt.hist(samples, bins=math.floor(np.sqrt(size)), density=True, alpha=0.6)
    plt.title('Mixture Distribution')

# plot scaled pdf of mixture distribution
    x = np.linspace(samples.min() - 1, samples.max() + 1, 1000)
    plt.plot(x, mixture.pdf(x), "red")
    plt.show()
```

#### Mixture Distribution



# Rejection sampler

In this section, you will implement and test a rejection sampling algorithm to generate samples from a target distribution using a proposal distribution. The rejection sampling algorithm will be implemented in the function rejection rsv.

- Inputs:
  - target\_pdf: The probability density function (pdf) of the target distribution (can be unnormalized).
  - proposal\_rv : An instance of a continuous random variable class used as the proposal distribution. You will need to use both the pdf and rsv methods of this class.
  - majorizing\_constant : A constant M such that M \*
    proposal\_rv.pdf(x) majorizes target\_pdf(x) for all x.
  - size : The number of samples to generate.
- Outputs:
  - samples : An ndarray of samples from the target distribution.
  - efficiency : The proportion of samples that were accepted.

Note that size is the number of samples to accept, not the number of samples to generate. You will need to generate more samples than size to ensure that you accept size samples.

```
In [102...

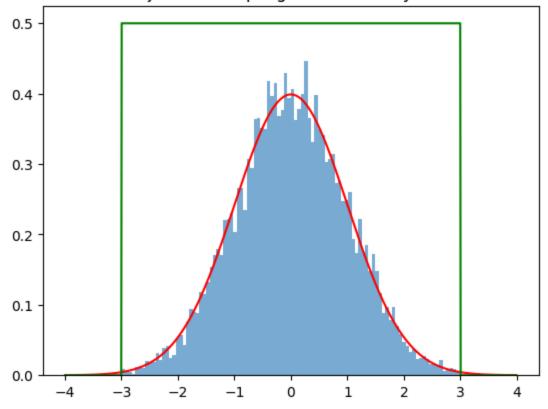
def rejection_rsv(target_pdf, proposal_rv: RandomVariable, majorizing_consta
    rng = RNG()
    accept = []
    trials = 0
    U = Uniform(rng, 0, 1)
    while len(accept) < size:
        x = proposal_rv.rsv(1)
        u = U.rsv(None)
        if u <= target_pdf(x) / (majorizing_constant * proposal_rv.pdf(x)):
        accept.append(x)
        trials += 1
    efficiency = size / trials
    return np.array(accept), efficiency</pre>
```

### Test 1

Test your algorithm on a truncated standard normal distribution. Use a uniform distribution to majorize the target distribution. A sample code is provided below. Fill in the appropriate constants. Feel free to find the majorizing constant by trial and error.

```
In [102... # test rejection rsv for normal distribution
         # find appropriate values of low, high, and M
         rng.seed(0)
         size = 10000
         target pdf = Normal(rng, mean=0, std=1).pdf
         proposal = Uniform(rng, low=-3, high=3)
         M = 3
         samples, efficiency = rejection_rsv(target_pdf, proposal, M, size)
         assert len(samples) == size, "Incorrect number of samples"
         # histogram of samples
         plt.hist(samples, math.floor(math.sqrt(size)), density=True, alpha=0.6)
         plt.title('Rejection Sampling with efficiency {:.2f}'.format(efficiency))
         # plot scaled pdf of target distribution
         x = np.linspace(samples.min() - 1, samples.max() + 1, 1000)
         plt.plot(x, target_pdf(x), "red")
         plt.plot(x, M*proposal.pdf(x), "green")
         plt.show()
```

### Rejection Sampling with efficiency 0.33

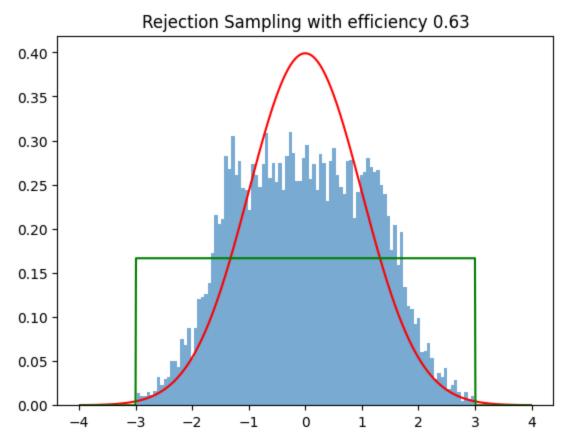


### Test 2

Next see what happens if you choose a value of M that is too small, say less than half the height of peak of the target distribution.

```
In [102... # test rejection_rsv for normal distribution
         # find appropriate values of low, high, and M
         rng.seed(0)
         size = 10000
         target_pdf = Normal(rng, mean=0, std=1).pdf
         proposal = Uniform(rng, low=-3, high=3)
         M = 1
         samples, efficiency = rejection_rsv(target_pdf, proposal, M, size)
         assert len(samples) == size, "Incorrect number of samples"
         # histogram of samples
         plt.hist(samples, math.floor(math.sqrt(size)), density=True, alpha=0.6)
         plt.title('Rejection Sampling with efficiency {:.2f}'.format(efficiency))
         # plot scaled pdf of target distribution
         x = np.linspace(samples.min() - 1, samples.max() + 1, 1000)
         plt.plot(x, target_pdf(x), "red")
         plt.plot(x, M*proposal.pdf(x), "green")
```





# Test 3

Test your algorithm on a (truncated) mixture of a normal distributions. Use a uniform distribution to majorize the target distribution. This time write your own code. The final graph should show the target distribution, the scaled proposal distribution, and a histogram of the samples.

```
In [103... # test rejection rsv
rng.seed(0)
size = 10000

normal1 = Normal(rng, mean=-2, std=1)
normal2 = Normal(rng, mean=3, std=1)

mixture = Mixture(rng, weights=[0.3, 0.7], components=[normal1, normal2])

target_pdf = mixture.pdf
proposal = Uniform(rng, low=-6, high=6)

M = 3.5

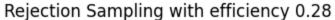
samples, efficiency = rejection_rsv(target_pdf, proposal, M, size)
```

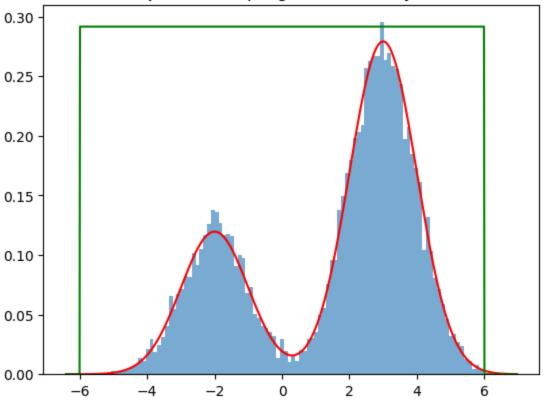
```
assert len(samples) == size, "Incorrect number of samples"

# histogram of samples
plt.hist(samples, math.floor(math.sqrt(size)), density=True, alpha=0.6)
plt.title('Rejection Sampling with efficiency {:.2f}'.format(efficiency))

# plot scaled pdf of target distribution
x = np.linspace(samples.min() - 1, samples.max() + 1, 1000)
plt.plot(x, target_pdf(x), "red")
plt.plot(x, M*proposal.pdf(x), "green")

plt.show()
```





# Test 4

Finally, test your algorithm on a (truncated) mixture of a normal distributions. Use a mixture of two or more uniform distributions to majorize the target distribution. Compare the efficiency of the algorithm to the one above.

```
mixture = Mixture(rng, weights=[0.3, 0.7], components=[normal1, normal2])
target_pdf = mixture.pdf
uniform1 = Uniform(rng, low=-4.5, high=1)
uniform2 = Uniform(rng, low=1, high=5)
proposal = Mixture(rng, weights=[1.2, 2], components=[uniform1, uniform2])
M = 0.6
samples, efficiency = rejection_rsv(target_pdf, proposal, M, size)
assert len(samples) == size, "Incorrect number of samples"
# histogram of samples
plt.hist(samples, math.floor(math.sqrt(size)), density=True, alpha=0.6)
plt.title('Rejection Sampling with efficiency {:.2f}'.format(efficiency))
# plot scaled pdf of target distribution
x = np.linspace(samples.min() - 1, samples.max() + 1, 1000)
plt.plot(x, target_pdf(x), "red")
plt.plot(x, M*proposal.pdf(x), "green")
plt.show()
# we have a higher efficiency here
```

