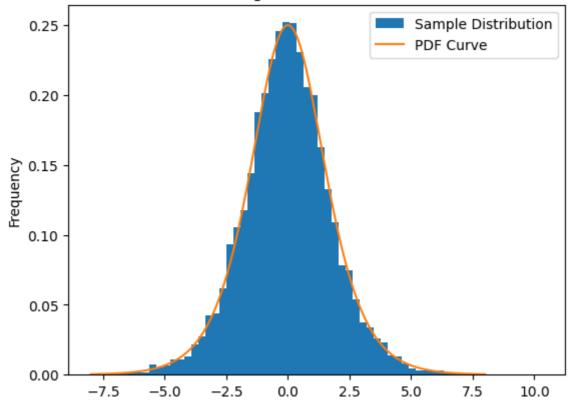
```
In [64]: import numpy as np
   import matplotlib.pyplot as plt
   import math
   import random
```

# **Problem 1. Inverse Transform Sampling**

## (a)Logistic distribution

```
In [62]: uniform_samples = np.random.uniform(0, 1, 10000)
    transformed_samples = -np.log((1 / uniform_samples) - 1)
    x_values = np.linspace(-8, 8, 1000)
    pdf_values = np.exp(-x_values) / ((1 + np.exp(-x_values))**2)
    plt.hist(transformed_samples, bins=64, density=True, label="Sample Distribution"
    plt.plot(x_values, pdf_values,label="PDF Curve")
    plt.ylabel('Frequency')
    plt.title('Logistic Distribution')
    plt.legend()
    plt.show()
    print("1")
```

#### Logistic Distribution



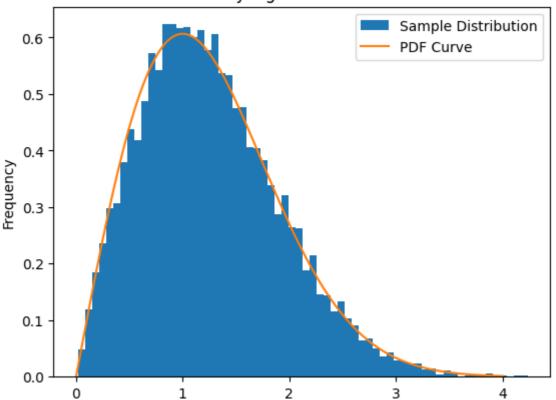
# (b) Rayleigh Distribution

1

```
In [46]: uniform_samples = np.random.uniform(0, 1, 10000)
    transformed_samples = np.sqrt(-2 * np.log(1 - uniform_samples))
    x_values = np.linspace(0, 4, 1000)
    pdf_values = x_values * np.exp(-1/2 * (x_values**2))
```

```
plt.hist(transformed_samples, bins=64, density=True, label="Sample Distribution"
plt.plot(x_values, pdf_values, label="PDF Curve")
plt.ylabel('Frequency')
plt.title('Rayleigh Distribution')
plt.legend()
plt.show()
```

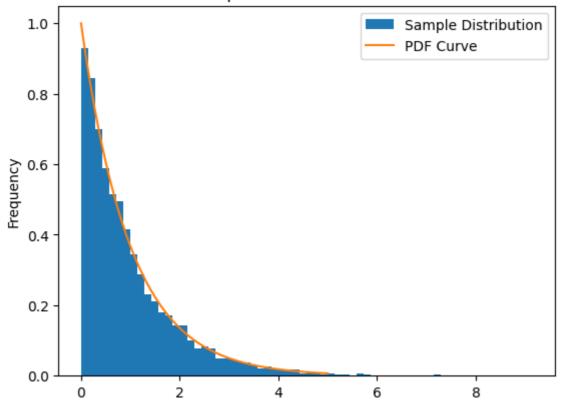
### Rayleigh Distribution



## (c) Exponential Distribution

```
In [47]: uniform_samples = np.random.uniform(0, 1, 10000)
    transformed_samples = -np.log(1 - uniform_samples)
    x_values = np.linspace(0, 5, 1000)
    pdf_values = np.exp(-x_values)
    plt.hist(transformed_samples, bins=64, density=True, label="Sample Distribution"
    plt.plot(x_values, pdf_values, label="PDF Curve")
    plt.ylabel('Frequency')
    plt.title('Exponential Distribution')
    plt.legend()
    plt.show()
```

#### **Exponential Distribution**



# **Problem 2. Samples Discrete Distributions**

## (a) Bern(0.5)

```
In [80]: N = 10000
p = 0.5

# Generate Bernoulli distributed samples directly
bern_samples = np.random.choice([0, 1], size=N, p=[1-p, p])
```

## (b) Bin(20, 0.5)

```
In [81]: N = 10000
n = 20
p = 0.5

# Generate samples using Bernoulli trials
bernoulli_trials = np.random.uniform(0, 1, (n, N)) > p
binomial_samples = bernoulli_trials.sum(axis=0)
```

## (c) Geom(0.5)

```
In [82]: N, p = 100000, 0.5
geom = np.zeros(N)

for i in range(N):
    u = np.random.random() # Generate a uniform random number
    k = 0
    while u > 1 - (1 - p) ** (k + 1):
```

```
k += 1
geom[i] = k
```

### (d) Nbin(10, 0.5)

```
In [83]: N, n, p = 10000, 10, 0.5
Ngeom = np.zeros(N)

for i in range(N):
    for _ in range(n):
        failures = 0
        while np.random.random() >= p:
            failures += 1
            Ngeom[i] += failures
```

#### (e)Pois(1)

```
In [84]: N = 10000
lam = 1

poisson_samples = np.zeros(N)

for i in range(N):
    L = np.exp(-lam)
    k = 0
    p = 1
    while p > L:
        k += 1
        p *= np.random.uniform(0, 1)
    poisson_samples[i] = k - 1
```

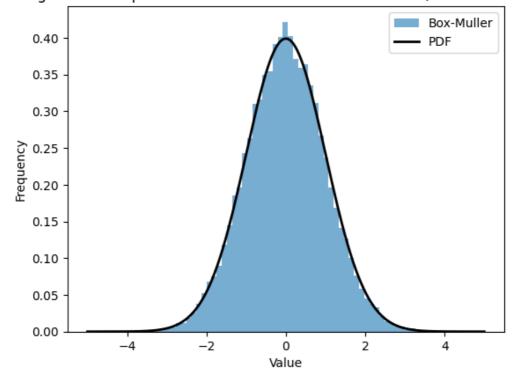
# Problem 3

## (a) Box-Muller

```
In [52]:
         import numpy as np
         import matplotlib.pyplot as plt
         from scipy.stats import norm
In [53]: def box_muller(n):
             u1, u2 = np.random.uniform(size=n), 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 np.concatenate([z1, z2])
         samples_box_muller = box_muller(10000)
         plt.hist(samples box muller, bins=60, density=True, alpha=0.6, label="Box-Muller
         x_{values} = np.linspace(-5, 5, 1000)
         pdf_values = norm.pdf(x_values, 0, 1)
         plt.plot(x_values, pdf_values, 'k', linewidth=2, label="PDF")
         plt.title("Histogram of Samples from Standard Normal Distribution (Box-Muller Me
         plt.xlabel("Value")
         plt.ylabel("Frequency")
```

```
plt.legend()
plt.show()
```

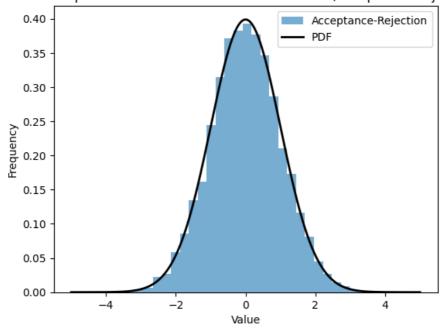
#### Histogram of Samples from Standard Normal Distribution (Box-Muller Method)



## (b) Acceptance-Rejection

```
In [54]: def acceptance_rejection(n):
              samples = []
              while len(samples) < n:</pre>
                  x = np.random.uniform(-4, 4)
                  y = np.random.uniform(0, 0.4)
                  pdf = (1/np.sqrt(2*np.pi)) * np.exp(-0.5*x**2)
                  if y < pdf:</pre>
                      samples.append(x)
              return np.array(samples)
          samples_acc_rej = acceptance_rejection(10000)
          plt.hist(samples_acc_rej, bins=30, density=True, alpha=0.6, label="Acceptance-Re
          {\tt plt.plot(x\_values, pdf\_values, 'k', linewidth=2, label="PDF")}
          plt.title("Histogram of Samples from Standard Normal Distribution (Acceptance-Re
          plt.xlabel("Value")
          plt.ylabel("Frequency")
          plt.legend()
          plt.show()
```

Histogram of Samples from Standard Normal Distribution (Acceptance-Rejection Method)



### (c) Compare

The Box-Muller method is favored for its straightforward approach to creating standard normal random variables from a uniform distribution, which is especially useful when dealing with standard normal distributions. However, this feature is also a drawback because it means that Box-Muller can't provide the variety of distributions some more complex scenarios might call for. Howeverly, the Acceptance-Rejection method ifavoreded for its adaptability. It allows us to draw samples from a broarangeum of probability distributions, including those that might be challenging to sample from directly due to their complexity. The key to its success lies in the relationship between the proposal and target distributions. For the method to be efficient, the proposal distribution must be a good fit over the entire range of the target distribution. If the fit is too tight, with the optimal constant near one, sampling becomes a tough task, and if the fit is too loose, indicated by a constant near zero, the acceptance rate plummets, which means a lot of generated samples end up being discaed.h.

### (d) bivariate

```
In [86]: def generate_bivariate_normal_samples(rho, n_samples):
    Z = np.random.normal(0, 1, n_samples)
    W = np.random.normal(0, 1, n_samples)
    X = Z
    Y = rho * Z + np.sqrt(1 - rho**2) * W
    return X, Y
    rhos = [-0.8, -0.4, 0, 0.4, 0.8]
    n_samples = 1000
    plt.figure(figsize=(20, 4))

for i, rho in enumerate(rhos, 1):
    plt.subplot(1, 5, i)
    X_samples, Y_samples = generate_bivariate_normal_samples(rho, n_samples)
```

```
plt.scatter(X_samples, Y_samples, alpha=0.5)
plt.title("p = {}".format(rho))
plt.xlabel("X")
plt.ylabel("Y")
plt.grid(True)
plt.axis('equal')

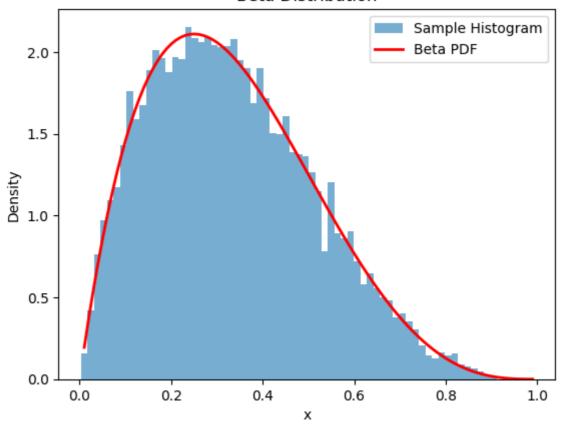
plt.tight_layout()
plt.show()
```

### Problem 4

## (a) Acceptance-Rejection

```
In [56]: import matplotlib.pyplot as plt
                                import numpy as np
                                from scipy.stats import beta
                                import random
                                import math
                                def beta_acceptance_rejection(a, b, n):
                                             i = 0
                                             result = []
                                             while i < n:
                                                           x = random.uniform(0, 1)
                                                           u = random.uniform(0, 1)
                                                           beta_value = math.factorial(a-1) * math.factorial(b-1) / math.factorial(
                                                           c = 1 / (beta_value * pow(((a-1)/(a+b-2)), a-1) * pow(((b-1)/(a+b-2)), b)
                                                           if u \leftarrow pow(x, a-1) * pow(1-x, b-1) / (c * pow(((a-1)/(a+b-2)), a-1) * pow((a-1)/(a+b-2)) * pow((a-1)/(a+b
                                                                        result.append(x)
                                                                        i += 1
                                             return result
                                a = 2
                                b = 4
                                n_reduced = 10000
                                samples_reduced = beta_acceptance_rejection(a, b, n_reduced)
                                plt.hist(samples_reduced, bins=64, density=True, alpha=0.6, label="Sample Histog
                                x = np.arange(0.01, 1, 0.01)
                                y = beta.pdf(x, a, b)
                                plt.plot(x, y, 'r-', lw=2, label='Beta PDF')
                                plt.title('Beta Distribution')
                                plt.xlabel('x')
                                plt.ylabel('Density')
                                plt.legend()
                                plt.show()
```

#### Beta Distribution



### (b) Monte Carlo

```
In [57]: n_points = 1000000
x_random = np.random.uniform(0, 1, n_points)
integrand_values = 4 / (1 + x_random**2)
integral_estimate = np.mean(integrand_values)
integral_estimate
```

Out[57]: 3.1411597720834674

#### (c) Monte Carlo

```
In [58]: def integrand(x):
    return np.sqrt(x + np.sqrt(x + np.sqrt(x + np.sqrt(x))))
    n_points = 1000000
    x_random = np.random.uniform(0, 4, n_points)
    integrand_values = integrand(x_random)
    integral_estimate = 4 * np.mean(integrand_values)
    integral_estimate
```

Out[58]: 7.676783736266756

## (d) Monte Carlo

```
In [59]: n_points = 10000000
x_random = np.random.uniform(-1, 1, n_points)
y_random = np.random.uniform(-1, 1, n_points)
inside_circle = (x_random ** 2 + y_random ** 2) <= 1</pre>
```

```
pi_estimate = 4 * np.mean(inside_circle)
pi_estimate
```

Out[59]: 3.1417264

## (e) importance sampling

```
import numpy as np
import math

def importance_sampling(n):
    sample = np.random.normal(8, 1, n)
    total_sum = np.sum(np.exp(8*(4-sample[sample > 8])))
    return total_sum / n

n = 10000000
prob_with_importance_sampling = importance_sampling(n)
print(f" importance sampling: {prob_with_importance_sampling}")
```

importance sampling: 6.217982714249921e-16