Statistical Analysis in Physics

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Problem 1: Distribution function and CLT theorem

- a) Generate a random sample space of size N using following probability distribution functions :
 - I. Binomial
 - II. Poisson
 - III. Normal
 - IV. Cauchy-Lorentz

and make histogram plot for at least three different values of their parameters .

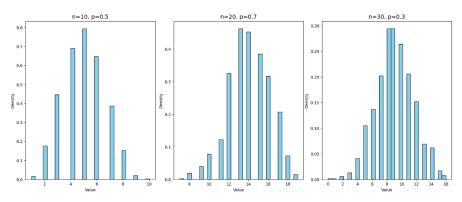
- b) Verify CLT theorem for the functions mentioned in part (a) for fixed sampling (M: take large value) and vary sample size (N). Show it graphically.
- c) With sample size (N) as large as the CLT theorem is verified in part (b), check for minimum sampling (M) required to achieve the normal distribution of sample means. You may take M = 100, 500, 1000, 5000, 10000. Show it graphically.

• Binomial Distribution Function

a) Histogram Plot

```
#GARIMA SINGH
import numpy as
                  np
import matplotlib.pyplot as plt
from scipy.stats import binom
np.random.seed(42)
N = 1000
params = [(10, 0.5), (20, 0.7), (30, 0.3)]
fig, axs = plt.subplots(1, 3, figsize=(18, 5))
fig.suptitle('Binomial Distribution - Different Parameters', fontsize=20)
for idx, (n, p) in enumerate(params):
    data = binom.rvs(n=n, p=p, size=N)
    axs[idx].hist(data, bins=30, density=True, edgecolor='black', color='skyblue')
    axs[idx].set_title(f'n={n}, p={p}', fontsize=14)
axs[idx].set_xlabel('Value')
axs[idx].set_ylabel('Density')
plt.tight layout()
plt.show()
```



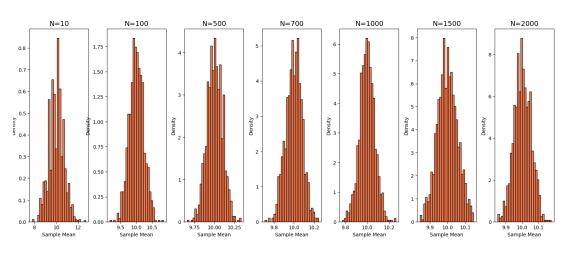


b) CLT Theorem verification by varying sample size N

```
import numpy as np
 import matplotlib.pyplot as plt
 from scipy.stats import binom
M = 1000
N values = [ 10, 100, 500,700,1000,1500,2000]
n, p = 20, 0.5
fig, axs = plt.subplots(1, len(N_values), figsize=(22, 5))
fig.suptitle('CLT Verification - Binomial Distribution', fontsize=20)
for idx, N in enumerate(N_values):
     sample_means = []
     for _ in range(M):
          sample = binom.rvs(n=n, p=p, size=N)
          sample_means.append(np.mean(sample))
     axs[idx].hist(sample_means, bins=30, density=True, edgecolor='black', color='coral')
axs[idx].set_title(f'N={N}', fontsize=14)
axs[idx].set_xlabel('Sample Mean')
axs[idx].set_ylabel('Density')
plt.tight layout()
plt.show()
```

Output

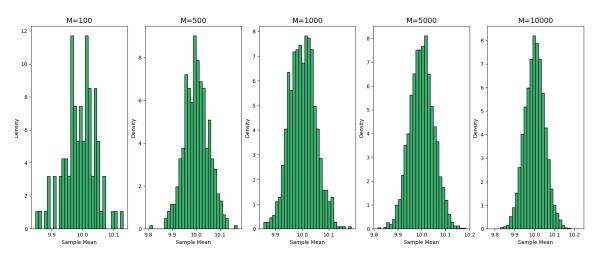
CLT Verification - Binomial Distribution



c)Minimum sampling (M) required to achieve normal distribution

```
import numpy as np
import matplotlib.pyplot as plt
from scipy.stats import binom
N = 2000
M values = [100, 500, 1000, 5000, 10000]
n, p = 20, 0.5
fig, axs = plt.subplots(1, len(M_values), figsize=(22, 5))
fig.suptitle('Finding Minimum M for CLT - Binomial Distribution', fontsize=20)
for idx, M in enumerate(M_values):
    sample_means = []
    for _ in range(M):
    sample = binom.rvs(n=n, p=p, size=N)
         sample_means.append(np.mean(sample))
     axs[idx].hist(sample\_means, bins=30, density=True, edgecolor='black', color='mediumseagreen') \\ axs[idx].set\_title(f'M=\{M\}', fontsize=14) 
    axs[idx].set_xlabel('Sample Mean')
    axs[idx].set_ylabel('Density')
plt.tight_layout()
plt.show()
```

Finding Minimum M for CLT - Binomial Distribution



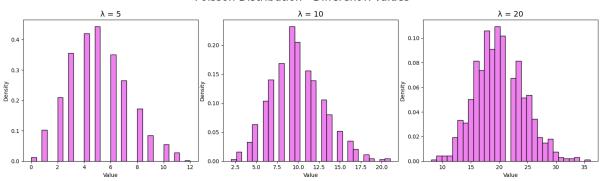
• Poisson Distribution Function

a)Histogram Plot

```
import numpy as np
import matplotlib.pyplot as plt
from scipy.stats import poisson
np.random.seed(42)
N = 1000
lambdas = [ 5, 10,20]
fig, axs = plt.subplots(1, 3, figsize=(18, 5))
fig.suptitle('Poisson Distribution - Different \( \lambda \) Values', fontsize=20)
for idx, lam in enumerate(lambdas):
    data = poisson.rvs(mu=lam, size=N)
    axs[idx].hist(data, bins=30, density=True, edgecolor='black', color='violet')
    axs[idx].set_title(f'\( \lamble \) = (lam\( \lamble \)', fontsize=14)
    axs[idx].set_xlabel('Value')
    axs[idx].set_ylabel('Density')
plt.tight_layout()
plt.show()
```

Output

Poisson Distribution - Different λ Values

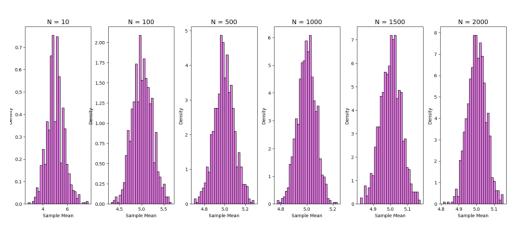


b)CLT Theorem verification by varying sample size N

```
import numpy as np
import matplotlib.pyplot as plt
from scipy.stats import poisson
M = 1000
N values = [10,100,500,1000,1500,2000]
lam = 5
fig, axs = plt.subplots(1, len(N_values), figsize=(22, 5)) fig.suptitle('CLT Verification - Poisson Distribution', fontsize=20)
for idx, N in enumerate(N_values):
    sample_means = []
     for _ in range(M):
    sample = poisson.rvs(mu=lam, size=N)
          sample means.append(np.mean(sample))
     axs[idx].hist(sample_means, bins=30, density=True, edgecolor='black', color='violet')
     axs[idx].set_title(f'N = {N}', fontsize=14)
axs[idx].set_xlabel('Sample Mean')
     axs[idx].set_ylabel('Density')
plt.tight_layout()
plt.show()
```

Output

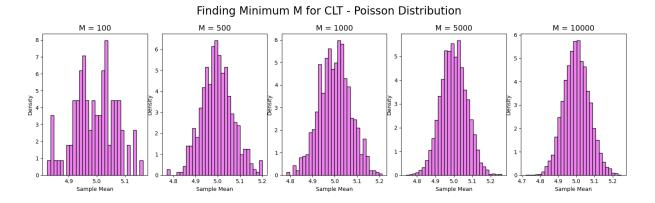
CLT Verification - Poisson Distribution



c)Minimum sampling (M) required to achieve normal distribution

```
import numpy as np
import matplotlib.pyplot as plt
from scipy.stats import poisson
N = 1000
M_values = [100, 500, 1000, 5000, 10000]
lam = 5
fig, axs = plt.subplots(1, len(M_values), figsize=(22, 5))
fig.suptitle('Finding Minimum M_for CLT - Poisson Distribution', fontsize=20)
for idx, M in enumerate(M_values):
    sample means = []
    for _ in range(M):
        sample = poisson.rvs(mu=lam, size=N)
        sample_means.append(np.mean(sample))

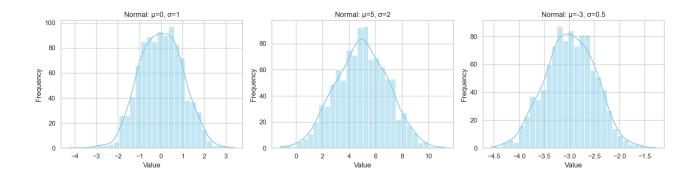
axs[idx].hist(sample_means, bins=30, density=True, edgecolor='black', color='violet')
    axs[idx].set_title(f'M = {M}', fontsize=14)
    axs[idx].set_vlabel('Sample Mean')
    axs[idx].set_ylabel('Density')
plt.tight_layout()
plt.show()
```



Normal Distribution Function

a)Histogram Plot

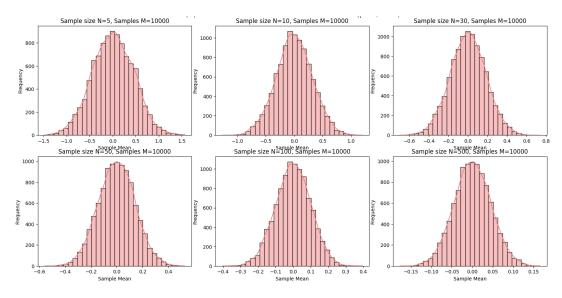
```
import numpy as np
import matplotlib.pyplot as plt
import seaborn as sns
sns.set(style="whitegrid")
def part_a normal_distribution(N, param_sets):
    plt.figure(figsize=(15, 4))
    for i, (mu, sigma) in enumerate(param_sets):
        samples = np.random.normal(loc=mu, scale=sigma, size=N)
        plt.subplot(1, 3, i + 1)
        sns.histplot(samples, bins=30, kde=True, color='skyblue')
        plt.title(f'Normal: \mu=\{mu\}, \sigma=\{sigma\}')
        plt.xlabel('Value')
        plt.ylabel('Frequency')
    plt.tight layout()
    plt.suptitle(f'Part (a): Normal Distributions with N={N}', y=1.05, fontsize=16)
    plt.show()
N = 1000
param_sets_normal = [(0, 1), (5, 2), (-3, 0.5)]
part a normal distribution(N, param sets normal)
```



b) CLT Theorem verification by varying sample size N

```
import numpy as np
import matplotlib.pyplot as plt
import seaborn as sns
def part_b_verify_clt_normal(M, N_values, mu=0, sigma=1):
    plt.figure(figsize=(18, 10))
    for i, N in enumerate(N_values):
        sample_means = []
        for _ in range(M):
            sample_means.append(np.mean(sample))
        plt.subplot(2, len(N_values) // 2, i + 1)
        sns.histplot(sample_means, bins=30, kde=True, color='lightcoral')
        plt.title(f'Sample_means, bins=30, kde=True, color='lightcoral')
        plt.tylabel('Sample_Mean')
        plt.ylabel('Sample_Mean')
        plt.ylabel('Frequency')
    plt.tight_layout()
    plt.suptitle(f'Part (b): CLT Verification for Normal Distribution (µ={mu}, o={sigma})', y=1.02, fontsize=16)
    plt.show()
M = 10000
N_values = [5, 10, 30, 50, 100, 500]
part_b_verify_clt_normal(M, N_values)
```

Output

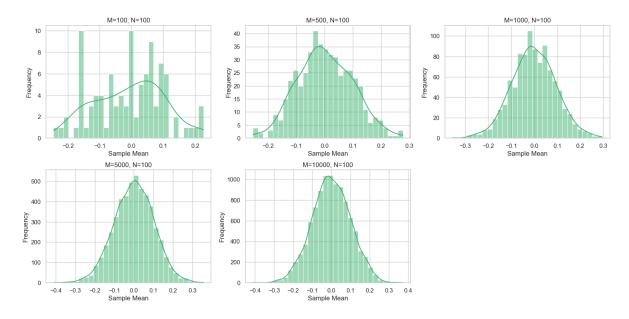


c)Minimum sampling (M) required to achieve normal distribution

```
import numpy as np
import matplotlib.pyplot as plt
import seaborn as sns
sns.set(style="withtegrid")
def part_c_minimum_m for_clt(N, M_values, mu=0, sigma=1):
    rows = 2
    cols = int(np.ceil(len(M_values) / rows))
    plt.figure(figsize=(cols * 5, rows * 4))
    for i, M in enumerate(M_values):
        sample means = []
        for _ in range(M):
            sample means = []
            for _ in range(M):
                sample means.append(np.mean(sample))

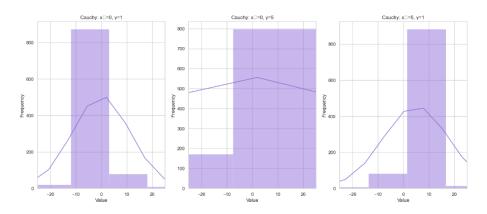
        plt.subplot(rows, cols, i + 1)
        sns.histplot(sample_means, bins=30, kde=True, color='mediumseagreen')
        plt.title(f'M=(M), N=(N)')
        plt.ylabel('Sample Mean')
        plt.ylabel('Sample Mean')
        plt.ylabel('Frequency')

plt.tight_layout()
    plt.supritle(f'Part (c): Minimum M Needed for Normality (Fixed N={N})', y=1.05, fontsize=16)
        plt.show()
        N_fixed = 100
        M_values = [100, 500, 1000, 5000, 10000]
        part_c_minimum_m_for_clt(N_fixed, M_values)
```



Cauchy-Lorentz Distribution Function

a)Histogram Plot



b) CLT Theorem verification by varying sample size N

```
import numpy as np
import matplotlib.pyplot as plt
import seaborn as sns

import subject (18, 10)

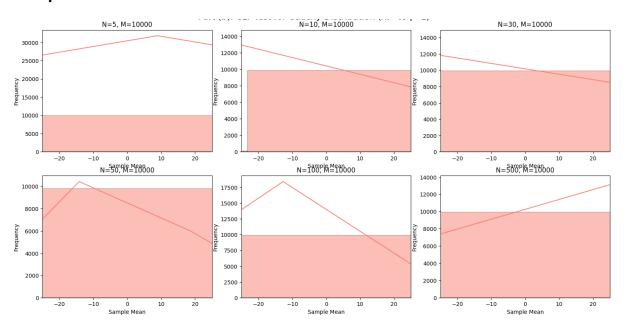
for in range (M, N_values):

    sample means = []

    for in range (M):

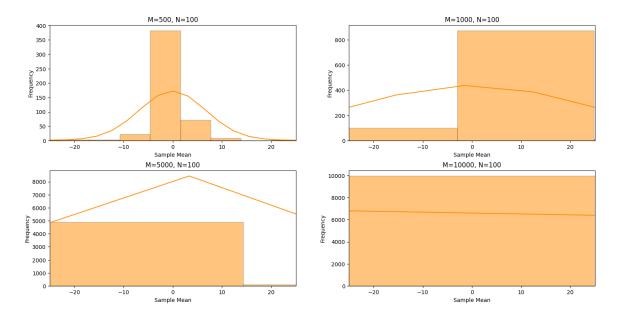
        sample = np.random.standard_cauchy(size=N) * gamma + x0

        samp
```



c) Minimum sampling (M) required to achieve normal distribution

```
import numpy as np
import matplotlib.pyplot as plt
 import seaborn as sns
 def part_c_minimum_m_for_cauchy(N, M_values, x0=0, gamma=1):
     cols = int(np.ceil(len(M_values) / rows))
plt.figure(figsize=(cols * 5, rows * 4))
     for i, M in enumerate(M values):
          sample_means = []
          for _ in range(M):
    sample = np.random.standard_cauchy(size=N) * gamma + x0
                sample_means.append(np.mean(sample))
          plt.subplot(rows, cols, i + 1)
           sns.histplot(sample_means, bins=100, kde=True, color='darkorange')
          plt.xlim(-25, 25)
plt.title(f'M={M}, N={N}')
plt.xlabel('Sample Mean')
          plt.ylabel('Frequency')
     plt.tight_layout()
plt.suptitle(f'Part (c): Varying M for CLT in Cauchy (Fixed N={N})', y=1.05, fontsize=16)
     plt.show()
N fixed cauchy = 100
M_values_cauchy = [ 500, 1000, 5000, 10000]
part_c_minimum_m_for_cauchy(N_fixed_cauchy, M_values_cauchy)
```



Cauchy distribution does **NOT** satisfy the Central Limit Theorem (CLT) well — because it has infinite variance.

So **even if we increase N or M**, the sample mean won't stabilize nicely into a normal distribution, unlike Normal or Binomial.

In graphs, we can see "wild behaviour" (many outliers, fat tails).

Problem 2: Joint distribution

a) Discrete case: Use random number generator (integer random number, Binomial, Poisson) to generate two random variables (X and Y) with sample size more than 50 and perform following:-

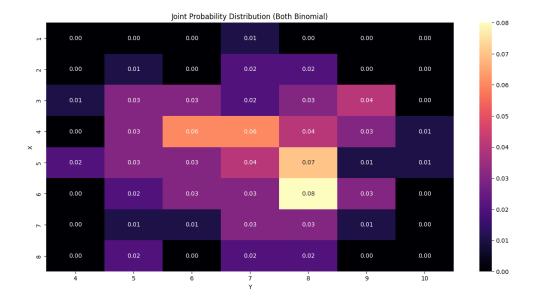
- For the two random variables, make a joint table and show the joint probability graphically. You may use 'heatmap()' or 'imshow()' function for graphical display.
- ii. Calculate and display the marginal distribution of the random variables.
- iii. Check if two variables are independent or not. [Optional]
- b) Continuous case: Generate two random variables (X and Y) of sample size more than 200 using normal distribution and perform following:
 - Make a 3dplot or surface plot of joint probability density of X and Y (probability surface).
 - ii. On the plot obtained in part (i) mark/show the region representing joint probability for X and Y in the range [a,b] and [c, d] respectively. (a > min(X), b < max(X) and c > min(Y), d < max(Y)).

BINOMIAL RANDOM NUMBER GENERATOR

Graphical Display of Joint Probability

```
import numpy as np
import pandas as pd
import seaborn as sns
import matplotlib.pyplot as plt
np.random.seed(42)
N = 100
X = np.random.binomial(n=10, p=0.5, size=N)
Y = np.random.binomial(n=10, p=0.7, size=N)
joint_table = pd.crosstab(X, Y, normalize='all')
print("\nJoint Probability Table (X vs Y):\n")
print(joint_table)
# Plot joint probability heatmap
plt.figure(figsize=(10, 8))
sns.heatmap(joint_table, cmap='magma', annot=True, fmt=".2f")
plt.title('Joint Probability Distribution (Both Binomial)')
plt.xlabel('Y')
plt.ylabel('X')
plt.show()
```

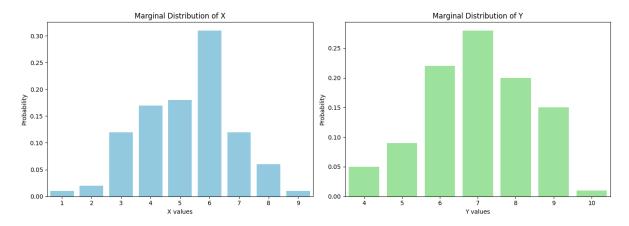
```
Joint Probability Table (X vs Y):
                    7
                        8
col 0
               6
                                   10
row 0
     0.00 0.00 0.00 0.01 0.00 0.00 0.00
     0.00
         0.01
              0.00
                   0.02
                        0.02
                            0.00
3
              0.03
                   0.02
     0.01
         0.03
                        0.03
                            0.04
                                 0.00
     0.00
         0.03 0.06 0.06
                       0.04
                            0.03
                                 0.01
5
     0.02 0.03 0.03 0.04 0.07 0.01 0.01
6
     0.00 0.02 0.03 0.03 0.08 0.03 0.00
     0.00 0.01 0.01 0.03 0.03 0.01 0.00
     0.00 0.02 0.00 0.02 0.02 0.00 0.00
```



Marginal Distribution of Random Variables

```
import numpy as np
import pandas as pd
import seaborn as sns
import matplotlib.pyplot as plt
N = 100
X = np.random.binomial(n=10, p=0.5, size=N)
Y = np.random.binomial(n=10, p=0.7, size=N)
joint_table = pd.crosstab(X, Y, normalize='all')
marginal_X = joint_table.sum(axis=1)
marginal_Y = joint_table.sum(axis=0)
print("\nMarginal Distribution of X:\n")
print(marginal X)
print("\nMarginal Distribution of Y:\n")
print(marginal Y)
fig, axs = plt.subplots(1, 2, figsize=(14, 5))
# Plot Marginal X
sns.barplot(x=marginal_X.index, y=marginal_X.values, ax=axs[0], color='skyblue')
axs[0].set title('Marginal Distribution of X')
axs[0].set_xlabel('X values')
axs[0].set_ylabel('Probability')
# Plot Marginal Y
sns.barplot(x=marginal Y.index, y=marginal Y.values, ax=axs[1], color='lightgreen')
axs[1].set_title('Marginal Distribution of Y')
axs[1].set xlabel('Y values')
axs[1].set ylabel('Probability')
plt.tight layout()
plt.show()
```

```
Marginal Distribution of X:
row_0
    0.01
    0.02
     0.12
    0.17
    0.18
    0.31
    0.12
    0.06
    0.01
dtype: float64
Marginal Distribution of Y:
col_0
     0.05
      0.09
      0.22
      0.28
      0.20
      0.15
10
     0.01
dtype: float64
```



Independence of Random Variables

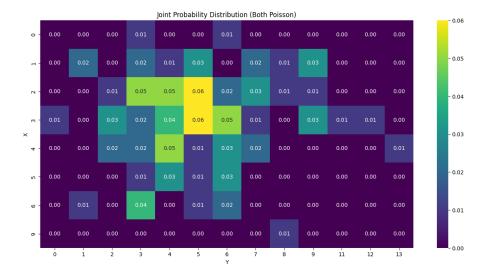
```
import numpy as np
import pandas as pd
import seaborn as sns
import matplotlib.pyplot as plt
N = 100
X = np.random.binomial(n=10, p=0.5, size=N)
Y = np.random.binomial(n=10, p=0.7, size=N)
joint_table = pd.crosstab(X, Y, normalize='all')
marginal_X = joint_table.sum(axis=1)
marginal_Y = joint_table.sum(axis=0)
product_marginals = np.outer(marginal_X, marginal_Y)
product_table = pd.DataFrame(product_marginals, index=joint_table.index, columns=joint_table.columns)
difference = np.abs(joint_table - product_table)
print("\nMaximum absolute difference between Joint and Product of Marginals:", difference.values.max())

# Decision based on threshold
threshold = 0.05  # you can set stricter value
if difference.values.max() < threshold:
    print("\n\ X and Y can be considered approximately independent.")
else:
    print("\n\ X and Y are NOT independent.")</pre>
```

POISSON RANDOM NUMBER GENERATOR

Graphical Display of Joint Probability

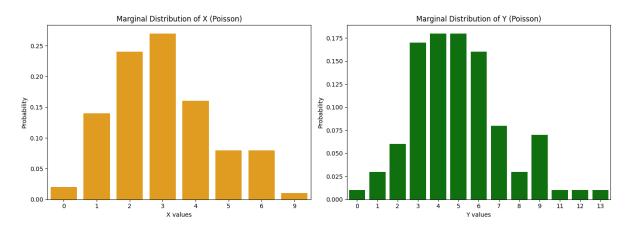
```
import numpy as np
import pandas as pd
import seaborn as sns
import matplotlib.pyplot as plt
np.random.seed(123)
N = 100
X = np.random.poisson(lam=3, size=N)
Y = np.random.poisson(lam=5, size=N)
joint table = pd.crosstab(X, Y, normalize='all')
print("\nJoint Probability Table (X vs Y) for Poisson:\n")
print(joint table)
# Plot joint probability heatmap
plt.figure(figsize=(10, 8))
sns.heatmap(joint table, cmap='viridis', annot=True, fmt=".2f")
plt.title('Joint Probability Distribution (Both Poisson)')
plt.xlabel('Y')
plt.ylabel('X')
plt.show()
```



Marginal Distribution of Random Variables

```
import numpy as np
import pandas as pd
import seaborn as sns
import matplotlib.pyplot as plt
np.random.seed(123)
N = 100
X = np.random.poisson(lam=3, size=N)
Y = np.random.poisson(lam=5, size=N)
joint table = pd.crosstab(X, Y, normalize='all')
marginal X = joint table.sum(axis=1)
marginal_Y = joint_table.sum(axis=0)
print("\nMarginal Distribution of X:\n")
print(marginal X)
print("\nMarginal Distribution of Y:\n")
print(marginal Y)
# Plot Marginal Distributions
fig, axs = plt.subplots(1, 2, figsize=(14, 5))
# Marginal X
sns.barplot(x=marginal X.index, y=marginal X.values, ax=axs[0], color='orange')
axs[0].set title('Marginal Distribution of X (Poisson)')
axs[0].set xlabel('X values')
axs[0].set_ylabel('Probability')
# Marginal Y
sns.barplot(x=marginal_Y.index, y=marginal_Y.values, ax=axs[1], color='green')
axs[1].set title('Marginal Distribution of Y (Poisson)')
axs[1].set xlabel('Y values')
axs[1].set ylabel('Probability')
plt.tight layout()
plt.show()
```

```
Marginal Distribution of X:
row_0
0 0.02
1 0.14
 1
3
4
5
6
      0.24
      0.27
       0.08
       0.08
      0.01
dtype: float64
Marginal Distribution of Y:
       0.01
1
2
3
4
5
6
7
8
9
11
12
        0.03
        0.06
        0.17
        0.18
        0.08
        0.07
        0.01
        0.01
13 0.01
dtype: float64
```



Independence of Random Variables

```
File Edit Format Run Options Window Help
import numpy as np
 import pandas as pd
import seaborn as sns
import matplotlib.pyplot as plt
np.random.seed (123)
N = 100
X = np.random.poisson(lam=3, size=N)
Y = np.random.poisson(lam=5, size=N)
joint table = pd.crosstab(X, Y, normalize='all')
marginal X = joint table.sum(axis=1)
marginal Y = joint table.sum(axis=0)
product_marginals = np.outer(marginal_X, marginal_Y)
product table = pd.DataFrame(product marginals, index=joint table.index, columns=joint table.columns)
difference = np.abs(joint table - product table)
print("\nMaximum absolute difference between Joint and Product of Marginals:", difference.values.max())
 threshold = 0.05
 if difference.values.max() < threshold:</pre>
    print("\n☑ X and Y can be considered approximately independent (Poisson).")
   print("\n X X and Y are NOT independent (Poisson).")
joint_flat = joint_table.stack()
product_flat = product_table.stack()
difference_flat = difference.stack()
comparison_df = pd.DataFrame({
     'Joint P(X,Y)': joint_flat,
'Product P(X)P(Y)': product_flat,
     'Absolute Difference': difference flat
})
comparison df = comparison df.reset index()
comparison df.rename(columns={'index': 'X', 'level 1': 'Y'}, inplace=True)
print("\n  Clean Comparison Table (Poisson case):\n")
print(comparison df)
 comparison df sorted = comparison df.sort values('Absolute Difference', ascending=False)
print("\nTop 10 biggest deviations (Poisson case):\n")
print(comparison_df_sorted.head(10))
```

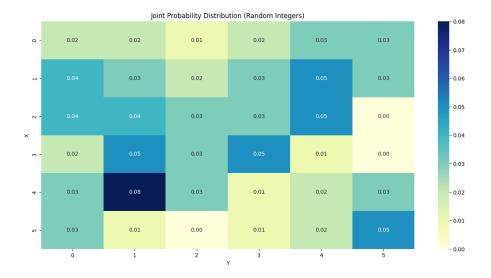
```
Maximum absolute difference between Joint and Product of Marginals: 0.0264
igspace X and Y can be considered approximately independent (Poisson).
Clean Comparison Table (Poisson case):
    row_0 col_0 Joint P(X,Y) Product P(X)P(Y) Absolute Difference
                  0.00
0.00
                              0.0002
0.0006
                                                0.0002
                                                          0.0006
                        0.00
                                       0.0012
                                                          0.0012
                      0.01
                                     0.0036
4
       0
                                                         0.0036
99
                      0.01
                                     0.0003
                                                         0.0097
                      0.00
0.00
0.00
100
                                      0.0007
                                                         0.0007
101
                                      0.0001
102
             12
                                      0.0001
                                                          0.0001
                                      0.0001
                                                          0.0001
103
             13
                       0.00
[104 rows x 5 columns]
Top 10 biggest deviations (Poisson case):
   row_0 col_0 Joint P(X,Y) Product P(X)P(Y) Absolute Difference
81
                       0.04
                                      0.0136
0.0459
                                                        0.0264
42
19
                       0.00
                                      0.0224
                                                         0.0224
56
                       0.05
                                      0.0288
                                                         0.0212
22
                       0.03
                                      0.0098
                                                         0.0202
57
                       0.01
                                      0.0288
                                                         0.0188
32
                       0.02
                                      0.0384
                                                         0.0184
                       0.03
                                      0.0128
31
                       0.06
                                      0.0432
                                                         0.0168
14
                       0.02
                                     0.0042
                                                         0.0158
```

INTEGER RANDOM NUMBER GENERATOR

Graphical Display of Joint Probability

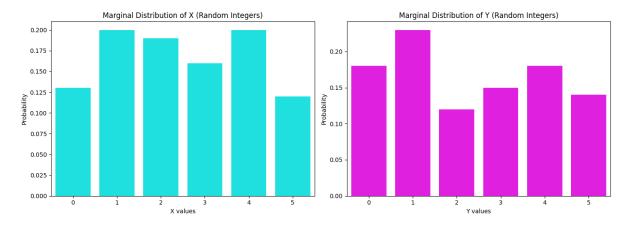
```
import numpy as np
import pandas as pd
import seaborn as sns
import matplotlib.pyplot as plt
np.random.seed(1234)
N = 100
X = np.random.randint(low=0, high=6, size=N)
Y = np.random.randint(low=0, high=6, size=N)
joint table = pd.crosstab(X, Y, normalize='all')
print("\nJoint Probability Table (X vs Y) for Random Integers:\n")
print(joint_table)
# Plot joint probability heatmap
plt.figure(figsize=(10, 8))
sns.heatmap(joint_table, cmap='YlGnBu', annot=True, fmt=".2f")
plt.title('Joint Probability Distribution (Random Integers)')
plt.xlabel('Y')
plt.ylabel('X')
plt.show()
```

```
Joint Probability Table (X vs Y) for Random Integers:
               1
                           3
                                       5
col 0
         0
                     2
                                 4
row 0
       0.02
            0.02
                  0.01
                        0.02
                              0.03
       0.04
            0.03
                  0.02
                        0.03
                              0.05
                                   0.03
2
       0.04
            0.04
                  0.03
                        0.03
                              0.05
                                   0.00
3
       0.02
            0.05
                  0.03
                        0.05
                              0.01
                                   0.00
4
      0.03 0.08
                  0.03
                        0.01 0.02
                                   0.03
5
      0.03 0.01 0.00 0.01 0.02
                                   0.05
```



Marginal Distribution of Random Variables

```
import numpy as np
import pandas as pd
import seaborn as sns
import matplotlib.pyplot as plt
np.random.seed(1234)
N = 100
X = np.random.randint(low=0, high=6, size=N)
Y = np.random.randint(low=0, high=6, size=N)
joint table = pd.crosstab(X, Y, normalize='all')
marginal_X = joint_table.sum(axis=1)
marginal_Y = joint_table.sum(axis=0)
print("\nMarginal Distribution of X:\n")
print(marginal X)
print("\nMarginal Distribution of Y:\n")
print(marginal Y)
# Plot Marginal Distributions
fig, axs = plt.subplots(1, 2, figsize=(14, 5))
# Marginal X
sns.barplot(x=marginal X.index, y=marginal X.values, ax=axs[0], color='cyan')
axs[0].set title('Marginal Distribution of X (Random Integers)')
axs[0].set xlabel('X values')
axs[0].set_ylabel('Probability')
# Marginal Y
sns.barplot(x=marginal_Y.index, y=marginal_Y.values, ax=axs[1], color='magenta')
axs[1].set_title('Marginal Distribution of Y (Random Integers)')
axs[1].set xlabel('Y values')
axs[1].set ylabel('Probability')
plt.tight layout()
plt.show()
```



Independence of Random Variables

```
import numpy as np
 import pandas as pd
 import seaborn as sns
 import matplotlib.pyplot as plt
np.random.seed(1234)
N = 100
X = np.random.randint(low=0, high=6, size=N)
 Y = np.random.randint(low=0, high=6, size=N)
joint_table = pd.crosstab(X, Y, normalize='all')
marginal X = joint_table.sum(axis=1)
marginal Y = joint_table.sum(axis=0)
product_marginals = np.outer(marginal_X, marginal_Y)
product_marginals = np.outer(marginal_x, marginal_x, m
threshold = 0.05 # set threshold
 if difference.values.max() < threshold:</pre>
           print("\n☑ X and Y can be considered approximately independent (Random Integers).")
print("\n X X and Y are NOT independent (Random Integers).")
joint_flat = joint_table.stack()
product_flat = product_table.stack()
difference flat = difference.stack()
comparison_df = pd.DataFrame({
            'Joint P(X,Y)': joint_flat,
             'Product P(X)P(Y)': product_flat,
            'Absolute Difference': difference_flat
comparison_df = comparison_df.reset_index()
comparison_df.rename(columns={'index': 'X', 'level_1': 'Y'}, inplace=True)
print("\n Tolean Comparison Table (Random Integer case):\n")
print(comparison df)
 comparison_df_sorted = comparison_df.sort_values('Absolute Difference', ascending=False)
print("\nTop 10 biggest deviations (Random Integer case):\n")
print(comparison_df_sorted.head(10))
```

```
\ensuremath{ullet} X and Y can be considered approximately independent (Random Integers).
🗒 Clean Comparison Table (Random Integer case):
   row 0 col 0 Joint P(X,Y) Product P(X)P(Y) Absolute Difference
                                                      3.400000e-03
                        0.02
                                        0.0299
                                                      9.900000e-03
                                                      5.600000e-03
                        0.02
                                        0.0195
                                                      5.000000e-04
                                        0.0234
                                                      6.600000e-03
                        0.03
                                        0.0182
0.0360
                        0.03
                                                      1.180000e-02
                                                      4.000000e-03
                        0.04
                                        0.0460
                                                      1.600000e-02
                                                      4.000000e-03
                        0.02
                                        0.0240
10
                        0.05
                                        0.0360
                                                      1.400000e-02
11
                                        0.0280
                                                      2.000000e-03
12
13
              0
                        0.04
                                        0.0342
                                                      5.800000e-03
                                        0.0437
                                                      3.700000e-03
                        0.04
                                        0.0228
                                                      7.200000e-03
15
16
                        0.03
                                        0.0285
                                                      1.500000e-03
                                        0.0342
17
18
                                        0.0266
0.0288
              5
0
                        0.00
                                                      2.660000e-02
                                                      8.800000e-03
                        0.02
19
20
                        0.05
                                        0.0368
                                                      1.320000e-02
                                                      1.080000e-02
                        0.03
                                        0.0192
22
23
                        0.01
                                        0.0288
                                                      1.880000e-02
                        0.00
                                        0.0224
                                                      2.240000e-02
24
25
              0
                        0.03
                                        0.0360
                                                      6.000000e-03
                                                      3.400000e-02
                                        0.0460
                        0.08
                                                      6.000000e-03
                        0.01
                                        0.0300
                                                      2.000000e-02
                                        0.0360
29
                        0.03
                                        0.0280
                                                      2.000000e-03
                                                      8.400000e-03
30
                                        0.0216
                         0.03
31
                        0.01
                                        0.0276
                                                      1.760000e-02
32
                                        0.0144
                                                      1.440000e-02
                        0.00
                                                      8.000000e-03
34
35
                        0.02
                                        0.0216
                                                      1.600000e-03
                                        0.0168
                                                      3.320000e-02
                                                0.0300
                                                                 z.000000e-02
28
                                                0.0360
                             0.02
                                                                 1.600000e-02
29
                             0.03
                                                0.0280
                                                                 2.000000e-03
30
                 0
                                                0.0216
                                                                 8.400000e-03
                             0.03
31
                                                0.0276
                             0.01
                                                                 1.760000e-02
32
                                                                 1.440000e-02
                                                0.0144
                             0.00
33
                             0.01
                                               0.0180
                                                                 8.000000e-03
                                                                1.600000e-03
34
                             0.02
                                                0.0216
35
                                                0.0168
                                                                3.320000e-02
                             0.05
Top 10 biggest deviations (Random Integer case):
             col_0 Joint P(X,Y) Product P(X)P(Y) Absolute Difference
25
                             0.08
                                                0.0460
                                                                       0.0340
35
                             0.05
                                                0.0168
                                                                       0.0332
17
                             0.00
                                                0.0266
                                                                       0.0266
21
                 3
                             0.05
                                                0.0240
                                                                       0.0260
23
                             0.00
                                                0.0224
                                                                       0.0224
27
                             0.01
                                                0.0300
                                                                       0.0200
22
                             0.01
                                                0.0288
                                                                       0.0188
31
                             0.01
                                                0.0276
                                                                       0.0176
                             0.03
                                                0.0460
                                                                       0.0160
28
                             0.02
                                                0.0360
                                                                       0.0160
```

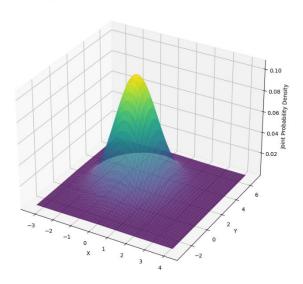
NORMAL RANDOM VARIABLE GENERATOR

3D PLOT

```
import numpy as np
import matplotlib.pyplot as plt
from scipy.stats import norm
from mpl_toolkits.mplot3d import Axes3D
np.random.seed(5678)
# Generate two continuous random variables X and Y (Normal Distribution)
N = 300 # sample size > 200
# Parameters for normal distribution
mu_X, sigma_X = 0, 1
mu_Y, sigma_Y = 2, 1.5
# Generate samples
X = np.random.normal(mu X, sigma X, N)
Y = np.random.normal(mu Y, sigma Y, N)
# Create grid for evaluating joint PDF
x = np.linspace(min(X)-1, max(X)+1, 100)
y = np.linspace(min(Y)-1, max(Y)+1, 100)
X_grid, Y_grid = np.meshgrid(x, y)
# Calculate joint PDF assuming independence
pdf_X = norm.pdf(X_grid, mu_X, sigma_X)
pdf_Y = norm.pdf(Y_grid, mu_Y, sigma_Y)
joint_pdf = pdf_X * pdf_Y
fig = plt.figure(figsize=(12, 9))
ax = fig.add_subplot(111, projection='3d')
ax.plot_surface(X_grid, Y_grid, joint_pdf, cmap='viridis', edgecolor='none', alpha=0.8)
ax.set_xlabel('X')
ax.set_ylabel('Y')
ax.set zlabel('Joint Probability Density')
ax.set_title('Joint Probability Density Surface (Normal Distribution)')
```

OUTPUT

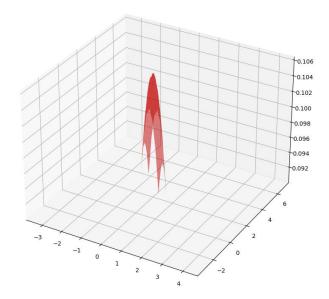




Highlight Region [a, b] and [c, d]

OUTPUT

```
Selected Region:
X in [-0.50, 0.50], Y in [1.50, 2.50]
```



Problem 3: Hypothesis testing

From an experiment of tossing of a coin n times we observe x number of heads (success) with probability of head being p_0 . Perform a hypothesis test with

Null hypothesis $H_0\colon p=p_0$, and Alternate hypothesis (i) $H_1\colon p\neq p_0$ Two-tailed test (ii) $H_1\colon p>p_0$ one-tailed test (right tail) (ii) $H_1\colon p< p_0$ one-tailed test (left tail)

Find P-value and make a decision with level of significance $\alpha = 0.05$ and 0.01.

Also calculate z-value for the experiment where $z=(\hat{p}-p_0)/\sqrt{p_0(1-p_0)/n}$ with statistical proportion $\hat{p}=x/n$. Compare it with critical value of z (i.e. $z_{\alpha/2}$ or z_{α}) to make your decision.

critical z	$\alpha = 0.05$	$\alpha = 0.01$
$Z_{\alpha/2}$	1.96	2.575
Z_{α}	1.645	2.325

Reference: Probability & Statistics for Engineers &; Scientists; Ronald E. Walpole, R.H. Myers ...

Solution

```
from scipy.stats import norm
def hypothesis_test_coin_toss(x, n, p0):
    # Step 1: Calculate sample proportion
      p_hat = x / n
# Step 2: Calculate z-value
      # Step 2: Calculate z-value
standard_error = math.sqrt(p0 * (1 - p0) / n)
z = (p_hat - p0) / standard_error
# Step 3: Calculate P-values
p_value_two_tailed = 2 * (1 - norm.cdf(abs(z)))
p_value_right_tailed = 1 - norm.cdf(z)
      p_value_left_tailed = norm.cdf(z)
       critical_values = {
    0.05: {"two_tailed": 1.96, "one_tailed": 1.645},
    0.01: {"two_tailed": 2.575, "one_tailed": 2.325}
       # Step 5: Display clean output
print("\n" + "="*50)
print(" HYPOTHESIS TESTING REPORT ".center(50, "="))
       print("="*50)
       print(f"Sample size (n): {n}")
       print(f"Number of successes (x): {x}")
       print(f"Hypothesized proportion (po): {p0}")
      print("-"*50)
print(f"Sample proportion (p^): {p_hat:.4f}")
print(f"Standard Error (SE): {standard_error:.4f}")
print(f"z-value: {z:.4f}")
print("-"*50)
       print("-"*50)
      print("-"*50)
print(f"P-value (Two-tailed): {p_value_two_tailed:.4f}")
print(f"P-value (Right-tailed): {p_value_right_tailed:.4f}")
print(f"P-value (Left-tailed): {p_value_left_tailed:.4f}")
print("="*50)
       for alpha in [0.05, 0.01]: 
 print(f"\n{'*'*50}") 
 print(f"DECISION AT SIGNIFICANCE LEVEL \alpha = {alpha}".center(50))
              print('*'*50)
               # Two-tailed Test
              decision_two_p = "Reject Ho" if p_value_two_tailed <= alpha else "Do not reject Ho" decision_two_z = "Reject Ho" if abs(z) > critical_values[alpha]["two_tailed"] else "Do not reject Ho"
              \sharp Right-tailed Test decision right p = "Reject Ho" if p value right tailed <= alpha else "Do not reject Ho"
```

```
# Two-tailed Test
decision_two_p = "Reject Ho" if p_value_two_tailed <= alpha else "Do not reject Ho"
decision_two_z = "Reject Ho" if abs(z) > critical_values[alpha]["two_tailed"] else "Do not reject Ho"
          # Right-tailed Test
decision_right_p = "Reject Ho" if p_value_right_tailed <= alpha else "Do not reject Ho"
decision_right_z = "Reject Ho" if z > critical_values[alpha]["one_tailed"] else "Do not reject Ho"
          # Left-tailed Test
decision_left_p = "Reject Ho" if p_value_left_tailed <= alpha else "Do not reject Ho"
decision_left_z = "Reject Ho" if z < -critical_values[alpha]["one_tailed"] else "Do not reject Ho"</pre>
           # Print all decisions
           print(f"Two-tailed Test (P-value method): {decision two p}")
           print(f"Two-tailed Test (Critical value method): {decision two z}")
           print("-"*50)
           print(f"Right-tailed Test (P-value method): {decision_right_p}")
           print(f"Right-tailed Test (Critical value method): {decision_right_z}")
           print("-"*50)
           print(f"Left-tailed Test (P-value method): {decision_left_p}")
           print(f"Left-tailed Test (Critical value method): {decision_left_z}")
# Example usage
# x = number of heads observed
# n = number of tosses
# p0 = hypothesized probability of head (under H0)
n = 50
hypothesis_test_coin_toss(x, n, p0)
```

```
====== HYPOTHESIS TESTING REPORT ========
Sample size (n): 50
Number of successes (x): 32
Hypothesized proportion (p<sub>0</sub>): 0.5
Sample proportion (p^): 0.6400
Standard Error (SE): 0.0707 z-value: 1.9799
P-value (Two-tailed): 0.0477
P-value (Right-tailed): 0.0239
P-value (Left-tailed): 0.9761
***********
DECISION AT SIGNIFICANCE LEVEL \alpha = 0.05
Two-tailed Test (P-value method): Reject Ho
Two-tailed Test (Critical value method): Reject Ho
Right-tailed Test (P-value method): Reject Ho
Right-tailed Test (Critical value method): Reject Ho
Left-tailed Test (P-value method): Do not reject Ho
Left-tailed Test (Critical value method): Do not reject Ho
************
DECISION AT SIGNIFICANCE LEVEL \alpha = 0.01
Two-tailed Test (P-value method): Do not reject H\circ Two-tailed Test (Critical value method): Do not reject H\circ
Right-tailed Test (P-value method): Do not reject Ho
Right-tailed Test (Critical value method): Do not reject Ho
Left-tailed Test (P-value method): Do not reject H₀
Left-tailed Test (Critical value method): Do not reject Ho
```

Problem 4: Bayesian Inference

From an experiment of flipping of a coin N times M heads showed up with statistical proportion $\hat{\theta} = M/N$. With the prior distribution $(\pi(\theta))$ given below

- (a) Beta distribution $B(\theta; \alpha, \beta)$ with given values of α, β
- (b) Gaussian distribution $n(\theta;\mu,\sigma)$ with a given mean μ) and stantdard deviation σ .

perform following

- i) Plot the prior distributions $(\pi(\theta))$ with θ .
- ii) Plot the likelihood $l(\theta|x)$ with θ and determine the value of θ that maximizes the probability of the data.
- iii) Plot the posterior distribution $\pi(\theta|x)=\pi(\theta)l(\theta|x)/g(x)$, where $g(x)=\sum_{\theta}\pi\left(\theta\right)l(\theta|x)$ is marginal distribution.
- iv) Make a single plot for posterior distribution for both the priors with the following values of (M,N)=(0,0);(1,1);(2,2);(2,3);(2,4);(3,8);(5,16);(10,32);20,64);(40,128);(80,256);(160,512);(320,1024);(640,2048);(1280,4096), discuss the behaviour of posterior distribution by increasing number of trials and number of successes keeping statistical proportion almost constant.

Solution

```
import numpy as np
import matplotlib.pyplot as plt
 Import matprovide import beta, norm theta = np.linspace(0, 1, 500) # 0 varies between 0 and 1 # Prior parameters alpha_prior = 2
 alpha_prior = 2
beta_prior = 2
mu_prior = 0.5
sigma_prior = 0.2
# Function: Likelihood for Binomial trials
def likelihood(theta, M, N):
    return theta**M * (1 - theta)**(N - M)
# Function: Posterior Calculation
def posterior(prior_func, theta, M, N):
    lik = likelihood(theta, M, N)
    numerator = prior_func(theta) * lik
           lik = likelinood(theta, M, N)
numerator = prior_func(theta) * lik
denominator = np.trapz(numerator, theta)
return numerator / denominator
 # Prior Functions
prior_beta = lambda t: beta.pdf(t, alpha_prior, beta_prior)
prior_gauss = lambda t: norm.pdf(t, mu_prior, sigma_prior)
# Example: Pick M and N
M = 5
N = 10
 theta_MLE = M / N
# ------
# (i) Plot Priors
  plt.figure(figsize=(10.5))
 plt.plot(theta, prior_beta(theta), label=f'Beta Prior (\alpha={alpha prior}, \beta={beta prior})') plt.plot(theta, prior_gauss(theta), label=f'Gaussian Prior (\mu={mu_prior}, \sigma={sigma_prior})')
  plt.title("Prior Distributions")
 plt.xlabel('0')
plt.ylabel('Prior Probability Density')
 plt.legend()
 plt.grid()
plt.show()
  # (ii) Plot Likelihood
  plt.figure(figsize=(10,5))
plt.ligure(figsize=[10,5))
plt.plot(theta, likelihood(theta, M, N), color='purple', label=f'Likelihood (M={M}, N={N})')
plt.axvline(theta_MLE, color='red', linestyle='--', label=f'MLE \theta_MLE:.2f)')
plt.title("Likelihood Function")
plt.xlabel('0')
plt.ylabel('Likelihood')
A 20°C
```

```
File Edit Format Run Options Window Help

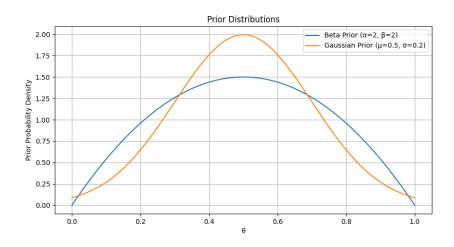
plt.plot(theta, prior_gauss(theta), label=f'Gaussian Prior (µ=(mu_prior), σ={sigma_prior})')

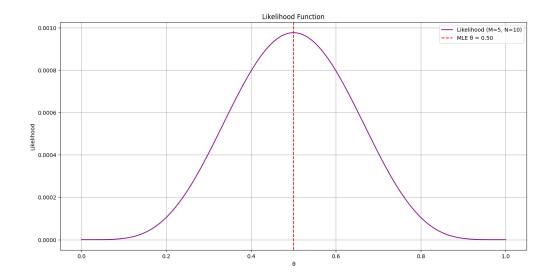
plt.title("Prior Distributions")

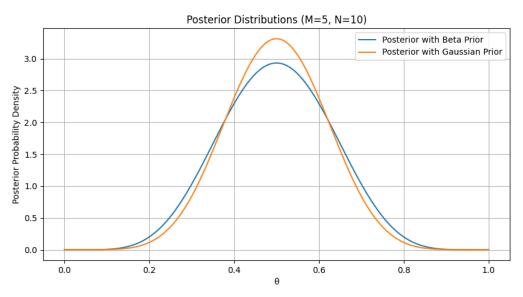
plt.xlabel('0')

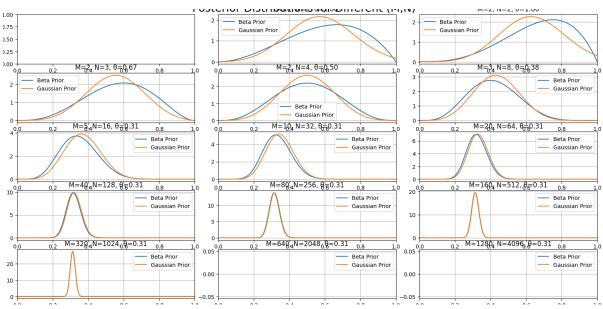
plt.ylabel('Prior Probability Density')

plt.legend()
 plt.grid()
plt.show()
 # ------
# (ii) Plot Likelihood
# -----
# ------
plt.figure(figsize=(10,5))
plt.pjot(theta, likelihood(theta, M, N), color='purple', label=f'Likelihood (M={M}, N={N})')
plt.axvline(theta_MLB, color='red', linestyle='--', label=f'MLE 0 = {theta_MLE:.2f}')
plt.title("Likelihood Function")
plt.ylabel('Ulikelihood')
plt.ylabel('Likelihood')
plt.legend()
plt.grid()
plt.show()
 posterior_beta = posterior(prior_beta, theta, M, N)
posterior_gauss = posterior(prior_gauss, theta, M, N)
plt.figure(figsize=(10,5))
plt.plot(theta, posterior_beta, label='Posterior with Beta Prior')
plt.plot(theta, posterior_gauss, label='Posterior with Gaussian Prior')
plt.title(f"Posterior Distributions (M={M}), N={N})")
plt.xlabel('0')
plt.ylabel('Posterior Probability Density')
plt.legend()
plt.grid()
 plt.grid()
plt.show()
 # ------
# (iv) Single plot for multiple (M,N)
# ------
 # Plotting
 fig, axs = plt.subplots(5, 3, figsize=(20, 20))
axs = axs.ravel()
 for i, (M, N) in enumerate(MN_pairs):
    if N == 0:
          post_beta = posterior(prior_beta, theta, M, N)
post_gauss = posterior(prior_gauss, theta, M, N)
          axs[i].plot(theta, post_beta, label='Beta Prior')
axs[i].plot(theta, post_gauss, label='Gaussian Prior')
axs[i].set_title(f"M={M}, N={N}, θ={M/N:.2f}")
axs[i].set_xlim(0,1)
axs[i].legend()
axs[i].prid()
           axs[i].grid()
 plt.tight layout()
 plt.suptitle("Posterior Distributions for Different (M,N)", y=1.02, fontsize=20)
 plt.show()
```









- As **N** increases, the posterior becomes sharper and narrower around the true $\vartheta = M/N$.
- **Prior** effect **decreases** as N becomes large (data dominates).
- When N is **small**, prior shape **significantly affects** the posterior.
- The statistical proportion (M/N) is kept **roughly constant**, so posterior centres remain near the same value, but become much **tighter**.