BRSM Take Home Assignment (21-01-2025)

Submitted by - Ashish Chokhani (2021102016)

```
import numpy as np
import matplotlib.pyplot as plt
import seaborn as sns
from scipy.stats import norm
```

Question 1

a. Assume that your population distribution is N(100,15)

```
mean = 100
std_dev = 15
```

b. Sample 10 random numbers (i.e., your sample size) from N(100,15) and calculate the mean and standard deviation of those numbers

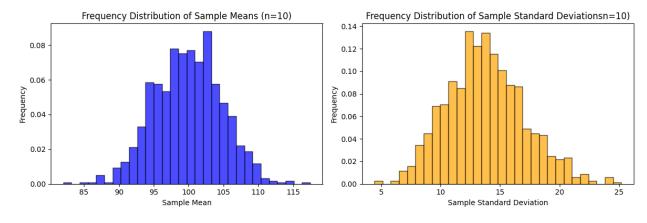
```
sample_size = 10
sample = np.random.normal(mean, std_dev, sample_size)
mean_sample = np.mean(sample)
std_dev_sample = np.std(sample)
print("Sample Mean:",mean_sample)
print("Sample Standard Deviation:",std_dev_sample)

Sample Mean: 101.17520214856617
Sample Standard Deviation: 10.865099743121363
```

c. Repeat this for 1000 trials, and plot the frequency distribution of the obtained means and standard deviations separately.

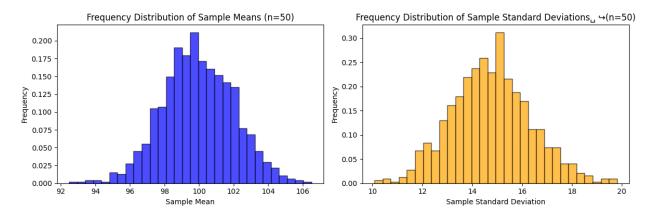
```
num_trials = 1000
means = np.zeros(num_trials)
std_devs = np.zeros(num_trials)
for i in range(num_trials):
    sample = np.random.normal(mean, std_dev, sample_size)
    means[i] = np.mean(sample)
    std_devs[i] = np.std(sample)
plt.figure(figsize=(12, 4))
plt.subplot(1, 2, 1)
plt.hist(means, bins=30, density=True, color='blue', alpha=0.7, edgecolor='black')
plt.title(f'Frequency Distribution of Sample Means (n={sample_size})')
plt.xlabel('Sample Mean')
plt.ylabel('Frequency')
```

```
plt.subplot(1, 2, 2)
plt.hist(std_devs, bins=30, density=True, color='orange', alpha=0.7,
edgecolor='black')
plt.title(f'Frequency Distribution of Sample Standard
Deviationsn={sample_size})')
plt.xlabel('Sample Standard Deviation')
plt.ylabel('Frequency')
plt.tight_layout()
plt.show()
```



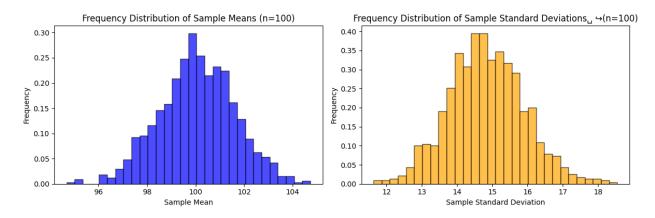
d. Repeat steps b and c for 50, 100, 500, and 1500 numbers.

```
sample sizes = [50, 100, 500, 1500]
for sample size in sample sizes:
    print("Sample Size =",sample_size)
    sample = np.random.normal(mean, std dev, sample size)
    mean sample = np.mean(sample)
    std dev sample = np.std(sample)
    print("Sample Mean:",mean_sample)
    print("Sample Standard Deviation:",std dev sample)
    num trials = 1000
    means = np.zeros(num trials)
    std devs = np.zeros(num_trials)
    for i in range(num trials):
        sample = np.random.normal(mean, std dev, sample size)
        means[i] = np.mean(sample)
        std devs[i] = np.std(sample)
    plt.figure(figsize=(12, 4))
    plt.subplot(1, 2, 1)
    plt.hist(means, bins=30, density=True, color='blue',
alpha=0.7,edgecolor='black')
    plt.title(f'Frequency Distribution of Sample Means
(n={sample size})')
    plt.xlabel('Sample Mean')
    plt.ylabel('Frequency')
```



Sample Mean: 100.94235976048219

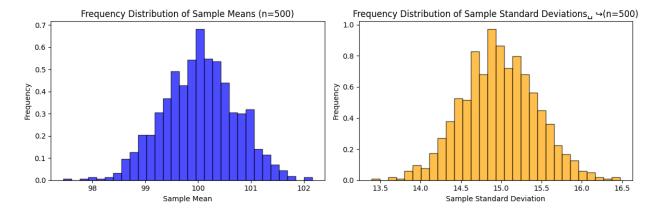
Sample Standard Deviation: 14.343823101952092



Sample Size = 500

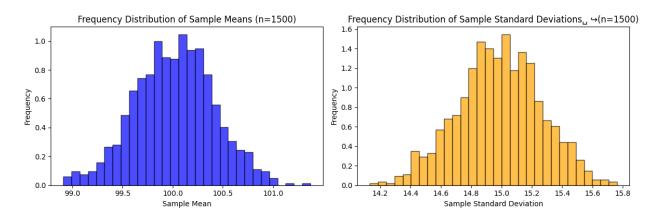
Sample Mean: 99.8472477782398

Sample Standard Deviation: 15.680709057453392



Sample Mean: 100.79486221261719

Sample Standard Deviation: 15.361824875812749



e. State and explain the observations and inferences you make regarding the above-produced histograms w.r.t to central limit theorem.

In the histograms, we observe that as the sample size increases, the distribution of sample means becomes more bell-shaped and symmetric, resembling a normal distribution. The standard deviation of the sample means decreases as the sample size increases, indicating that larger samples provide more precise estimates of the population mean. These observations align with the central limit theorem.

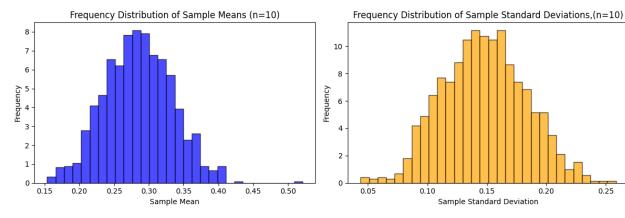
QUESTION 2

Repeat the above for a population distribution that is a Beta distribution with shape parameters 2 and 5.

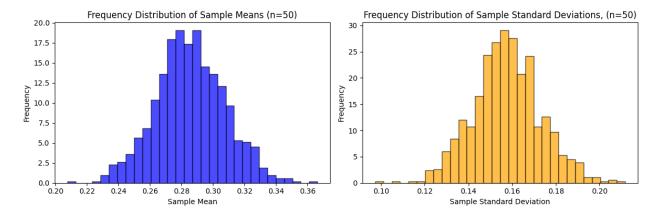
```
alpha = 2
beta = 5

mean_beta = alpha / (alpha + beta)
std_dev_beta = np.sqrt((alpha * beta) / ((alpha + beta)**2 * (alpha + beta + 1)))
```

```
print("Distribution Mean:",mean_beta)
print("Distribution Standard Deviation:",std dev beta)
sample size = 10
sample beta = np.random.beta(alpha, beta, sample size)
mean sample = np.mean(sample beta)
std \overline{dev} sample = np.std(sample beta)
print("Sample Mean:", mean sample)
print("Sample Standard Deviation:",std dev sample)
Distribution Mean: 0.2857142857142857
Distribution Standard Deviation: 0.15971914124998499
Sample Mean: 0.27298599530316275
Sample Standard Deviation: 0.16746596703231217
num trials = 1000
means = np.zeros(num trials)
std devs = np.zeros(num trials)
for i in range(num trials):
    sample beta = np.random.beta(alpha, beta, sample size)
    means[i] = np.mean(sample beta)
    std devs[i] = np.std(sample beta)
plt.figure(figsize=(12, 4))
plt.subplot(1, 2, 1)
plt.hist(means, bins=30, density=True, color='blue',
alpha=0.7,edgecolor='black')
plt.title(f'Frequency Distribution of Sample Means (n={sample size})')
plt.xlabel('Sample Mean')
plt.ylabel('Frequency')
plt.subplot(1, 2, 2)
plt.hist(std devs, bins=30, density=True, color='orange',
alpha=0.7,edgecolor='black')
plt.title(f'Frequency Distribution of Sample Standard Deviations,
(n={sample size})')
plt.xlabel('Sample Standard Deviation')
plt.ylabel('Frequency')
plt.tight_layout()
plt.show()
```

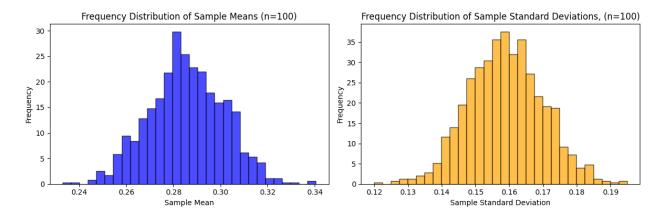


```
sample sizes = [50, 100, 500, 1500]
for sample size in sample sizes:
    print("Sample Size =",sample size)
    sample beta = np.random.beta(alpha, beta, sample size)
    mean sample = np.mean(sample beta)
    std dev sample = np.std(sample beta)
    print("Sample Mean:", mean_sample)
    print("Sample Standard Deviation:",std dev sample)
    num trials = 1000
    means = np.zeros(num trials)
    std devs = np.zeros(num trials)
    for i in range(num trials):
        sample beta = np.random.beta(alpha, beta, sample size)
        means[i] = np.mean(sample beta)
        std devs[i] = np.std(sample beta)
    plt.figure(figsize=(12, 4))
    plt.subplot(1, 2, 1)
    plt.hist(means, bins=30, density=True, color='blue',
alpha=0.7,edgecolor='black')
    plt.title(f'Frequency Distribution of Sample Means
(n={sample size})')
    plt.xlabel('Sample Mean')
    plt.ylabel('Frequency')
    plt.subplot(1, 2, 2)
    plt.hist(std devs, bins=30, density=True, color='orange',
alpha=0.7,edgecolor='black')
    plt.title(f'Frequency Distribution of Sample Standard Deviations,
(n={sample size})')
    plt.xlabel('Sample Standard Deviation')
    plt.ylabel('Frequency')
    plt.tight layout()
    plt.show()
Sample Size = 50
Sample Mean: 0.2651051496271604
Sample Standard Deviation: 0.1546243968175475
```



Sample Mean: 0.29741064538154854

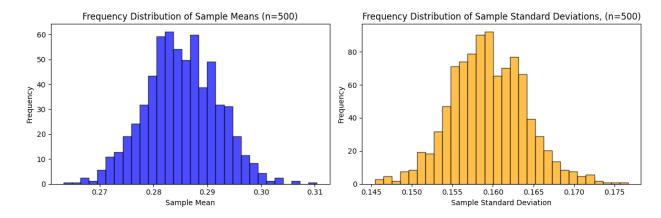
Sample Standard Deviation: 0.17651741382290384



Sample Size = 500

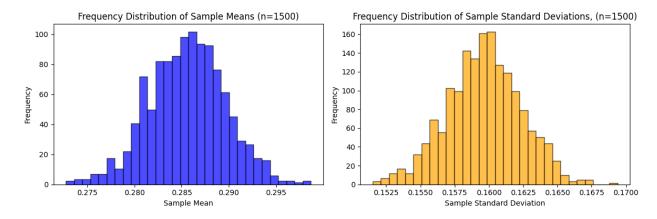
Sample Mean: 0.27573161785947153

Sample Standard Deviation: 0.15067284539580048



Sample Mean: 0.284882953222717

Sample Standard Deviation: 0.15987801928898857



Repeat the above for a population distribution that is a Beta distribution with shape parameters 2 and 5. Do you need a larger or smaller sample size now so that your sample estimate of the population mean is accurate?

- We observe that, as sample size increases the distribution tends to become more normal but it is still influenced by the original shape of beta distribution which is skewed and not symmetric.
- The distribution of sample means take longer to converge to a normal distribution compared to the normal distribution in Question 1.

Question 3

a. Sample 30 points from each of the normal distributions of mean 0 and SD 1, mean 2 and SD 0.5, and mean 3 and SD 2.

b. Plot the means and distributions of the three groups using the appropriate visualization tools. Include an error bar for each group indicating the condence interval.

c. Repeat the above two steps by sampling 70 and 100 points

```
def calculate_confidence_interval(data, confidence=0.95):
    Calculate the confidence interval for the mean of the data.
    mean = np.mean(data)
    std_error = np.std(data, ddof=1) / np.sqrt(len(data))
    margin_of_error = norm.ppf((1 + confidence) / 2) * std_error
    return mean, mean - margin_of_error, mean + margin_of_error

def plot_groups_with_confidence_intervals(group_data, group_labels, sample_sizes):
```

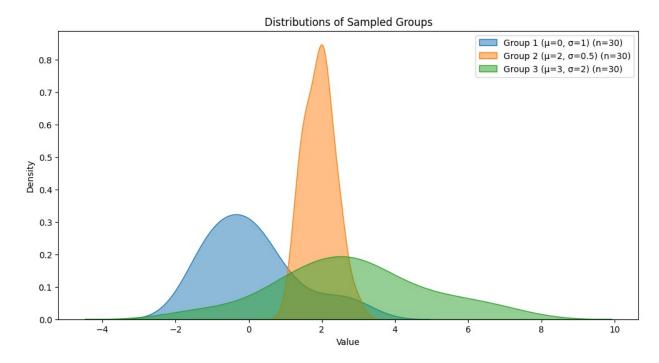
```
0.00
    Plot the means, distributions, and confidence intervals for each
group.
    num groups = len(group data)
    plt.figure(figsize=(12, 6))
    # Plot the distributions
    for i, (data, label) in enumerate(zip(group data, group labels)):
        sns.kdeplot(data, fill=True, alpha=0.5, label=f"{label}
(n={sample sizes[i]})")
    plt.title("Distributions of Sampled Groups")
    plt.xlabel("Value")
    plt.ylabel("Density")
    plt.legend()
    plt.show()
    # Plot the means with error bars
    means = []
    lower bounds = []
    upper bounds = []
    for data in group data:
        mean, lower, upper = calculate confidence interval(data)
        means.append(mean)
        lower bounds.append(lower)
        upper bounds.append(upper)
    plt.figure(figsize=(8, 5))
    x = np.arange(num groups)
    plt.bar(x, means, yerr=[np.array(means) - np.array(lower bounds),
                            np.array(upper bounds) - np.array(means)],
            capsize=5, alpha=0.6, color=['blue', 'orange', 'green'],
tick label=group labels)
    plt.title("Means with Confidence Intervals")
    plt.ylabel("Mean")
    plt.xlabel("Group")
    plt.show()
# Parameters for the distributions
params = [
    (0, 1), # mean=0, std dev=1
    (2, 0.5), # mean=2, std_dev=0.5
    (3, 2) # mean=3, std dev=2
1
sample_sizes = [30, 70, 100]
for sample size in sample sizes:
```

```
print(f"Sample Size: {sample_size}")

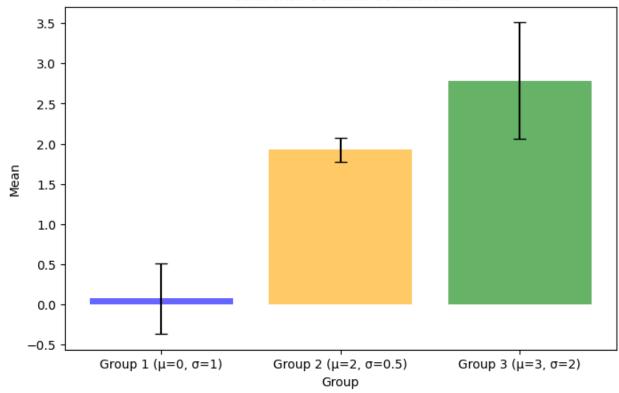
# Sampling data from the distributions
group_data = [np.random.normal(mean, std_dev, sample_size) for
mean, std_dev in params]
group_labels = [f"Group {i+1} (μ={mean}, σ={std_dev})" for i,
(mean, std_dev) in enumerate(params)]

# Plot distributions and confidence intervals
plot_groups_with_confidence_intervals(group_data, group_labels,
[sample_size] * len(params))

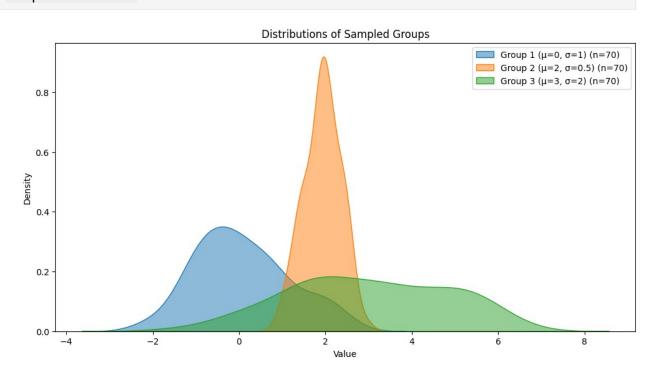
Sample Size: 30
```



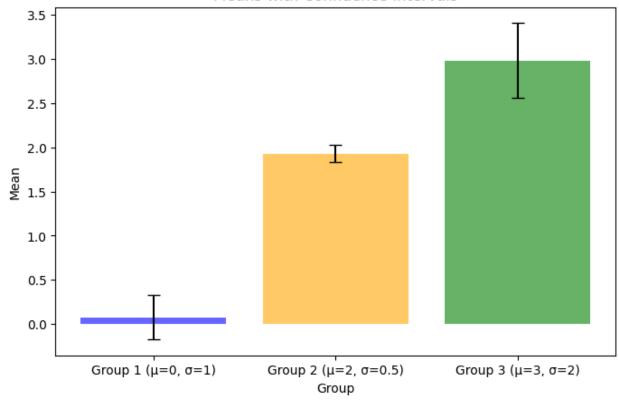
Means with Confidence Intervals



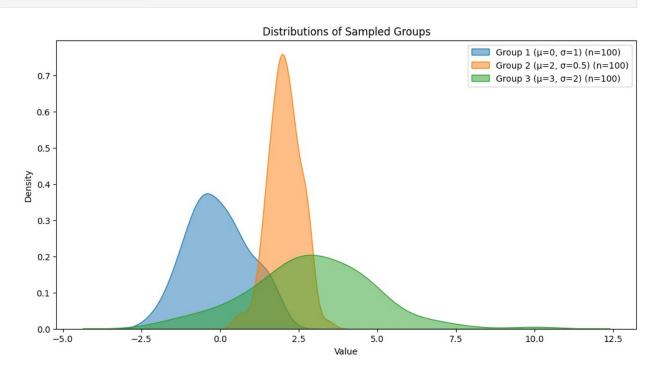
Sample Size: 70



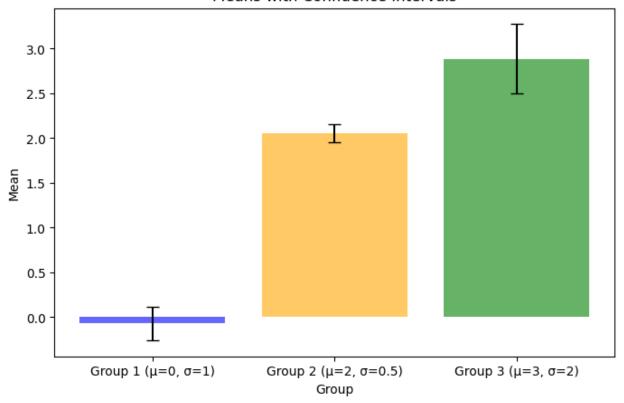
Means with Confidence Intervals



Sample Size: 100



Means with Confidence Intervals



d. How does the condence interval change along with the increase in the number of sampled points?

Part d: Effect of sample size on confidence intervals As the sample size increases:

- The confidence interval becomes narrower.
- The margin of error decreases because the standard error (sigma/sqrt(n)) is inversely proportional to the square root of the sample size.
- e. Write down your interpretation of the obtained condence intervals.
 - Confidence intervals provide a range within which the population mean is likely to fall with a certain confidence level (e.g., 95%).
 - Smaller sample sizes result in wider confidence intervals, reflecting greater uncertainty in estimating the population mean.
 - Larger sample sizes reduce uncertainty, leading to narrower confidence intervals, providing more precise estimates of the population mean.