

DA-AG-006: Applied Statistics and Inference - Assignment

Question 1: What are Type I and Type II errors in hypothesis testing, and how do they impact decision-making?

Answer:

- Type I error (α): Rejecting the null hypothesis H_0 when it is actually true. This is a false positive. The probability of a Type I error is controlled by the significance level α (commonly 0.05).
- Type II error (β): Failing to reject H_0 when the alternative hypothesis H_1 is actually true. This is a false negative. The probability of correctly rejecting a false null is the power ($1-\beta$).

Impact on decision-making:

- Trade-off: Decreasing α (making tests more conservative) typically increases β (lower power), and vice versa. Choice of α depends on consequences: if false positives are costly (e.g., claiming a drug works when it does not), choose smaller α ; if false negatives are more harmful (missing a real effect), prioritize power and increase sample size to reduce β .
- Practical response: Compute required sample size to achieve desired power for a chosen effect size and α ; consider consequences and costs of both error types when setting α and designing experiments.

Question 2: What is the P-value in hypothesis testing, and how should it be interpreted in the context of the null hypothesis?

Answer:

- Definition: The p-value is the probability of observing data at least as extreme as what was observed, assuming the null hypothesis H_0 is true.
- Interpretation: A small p-value (below the chosen α) indicates that the observed data are unlikely under H_0 , and provides evidence to reject H_0 . It is not the probability that H_0 is true.
- Cautions: Do not equate p-value with effect size—report confidence intervals and effect sizes alongside p-values. Multiple testing inflates false positive rate; adjust p-values or use corrections (e.g., Bonferroni) when performing many tests.

Question 3: Explain the difference between a Z-test and a T-test, including when to use each.

Answer:

- Z-test: Used when the population standard deviation σ is known and/or sample size is large (commonly $n \geq 30$ so CLT applies). The test statistic uses the normal (Z) distribution.
- T-test: Used when σ is unknown and estimated by the sample standard deviation s , especially for small samples ($n < 30$). The test statistic follows Student's t-distribution with $n-1$ degrees of freedom.
- Practical guidance: For means, use a t-test by default when σ is unknown. For large samples the t-distribution approaches normal, so z- and t-tests give similar results. Also consider one-sample vs two-sample and paired designs when choosing the specific t-test variant.

Question 4: What is a confidence interval, and how does the margin of error influence its width and interpretation?

Answer:

- Confidence interval (CI): A range computed from sample data that—under repeated sampling—would contain the true population parameter with a specified confidence level (e.g., 95%). For a sample mean: $CI = \text{sample_mean} \pm \text{margin_of_error}$.
- Margin of error (ME): $ME = \text{critical_value} \times \text{standard_error}$. It controls the half-width of the CI. Larger ME → wider CI (less precision); smaller ME → narrower CI (more precision).
- Factors affecting ME and width: sample size (larger $n \rightarrow$ smaller SE \rightarrow smaller ME), variability in data (larger $\sigma \rightarrow$ larger ME), and confidence level (higher confidence level \rightarrow larger critical value \rightarrow larger ME).

- Interpretation: A 95% CI means that the procedure used to compute the interval will produce intervals that include the true parameter 95% of the time across repeated samples. It does not say there's a 95% probability the specific interval contains the parameter (frequentist interpretation).

Question 5: Describe the purpose and assumptions of an ANOVA test. How does it extend hypothesis testing to more than two groups?

Answer:

- Purpose: ANOVA (Analysis of Variance) tests whether the means of three or more groups are equal (H_0 : all group means are the same) by comparing between-group variability to within-group variability.

- Key assumptions:

1. Independence of observations.

2. Normality: residuals within each group are approximately normally distributed.

3. Homogeneity of variances: group variances are approximately equal.

- How it extends hypothesis testing: Instead of multiple pairwise t-tests (which inflate Type I error), ANOVA provides a single global test. If ANOVA rejects H_0 , follow-up post-hoc tests (e.g., Tukey HSD) identify which pairs of groups differ while controlling family-wise error.

Question 6: Write a Python program to perform a one-sample Z-test and interpret the result for a given dataset.

Answer:

Below is a one-sample Z-test implementation and an example. The test compares the sample mean to a known population mean μ_0 . We assume population standard deviation σ is known.

Example result: sample mean = 56.4286, z-statistic = 3.5211, two-sided p-value = 0.0004.

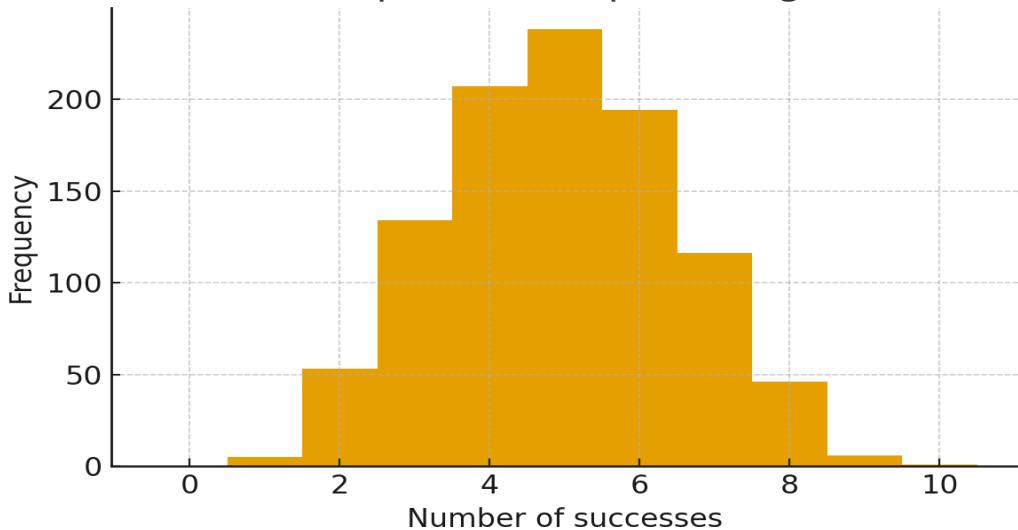
```
import numpy as np
from math import sqrt
# Example data
np.random.seed(0)
sample = np.random.normal(loc=52, scale=10, size=30) # sample drawn from population
n = len(sample)
sample_mean = sample.mean()
sigma = 10.0 # assumed known population std dev
mu0 = 50.0
z_stat = (sample_mean - mu0) / (sigma / sqrt(n))
# two-sided p-value
p_value = 2 * (1 - stats.norm.cdf(abs(z_stat)))
sample_mean, z_stat, p_value
```

Question 7: Simulate a dataset from a binomial distribution ($n = 10$, $p = 0.5$) using NumPy and plot the histogram.

Answer: Code and histogram (1000 draws) are provided below.

```
import numpy as np
samples = np.random.binomial(10, 0.5, size=1000)
# plt.hist(samples) # histogram saved as q7_binomial_hist.png
```

Q7: Binomial($n=10, p=0.5$) sample histogram (1000 d)



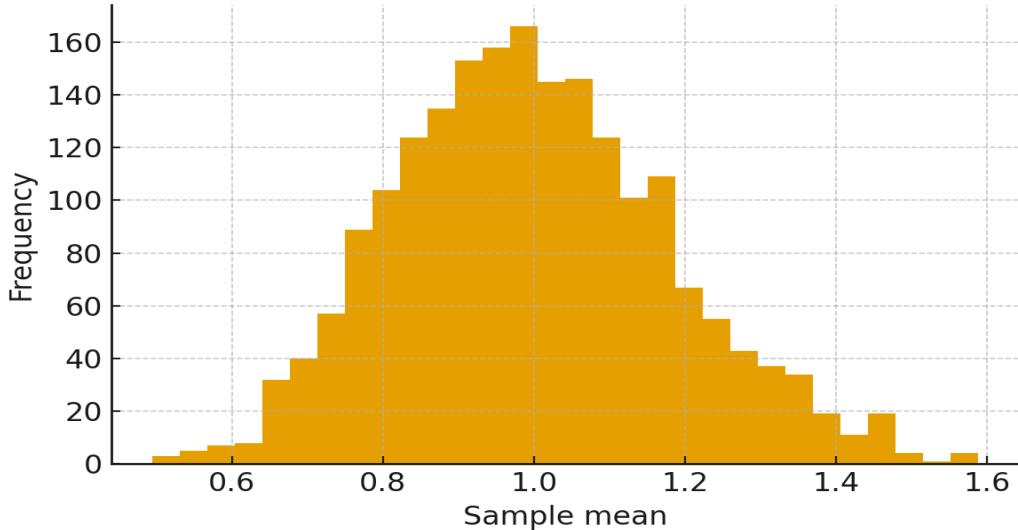
Question 8: Generate multiple samples from a non-normal distribution and implement the Central Limit Theorem using Python.

Answer: We used an exponential population, drew 2000 samples of size 30 and plotted the sampling distribution of the mean. The histogram shows approximate normality of sample means.

```
import numpy as np
```

```
sample_means = [np.mean(np.random.exponential(scale=1.0, size=30)) for _ in range(2000)]  
# histogram saved as q8_clt_hist.png
```

Sampling distribution of mean ($n=30$) from Exponential



Question 9: Write a Python function to calculate and visualize the confidence interval for a sample mean.

Answer: Example with a sample of $n=50$ drawn from $N(5, 2^2)$.

Sample mean = 5.1423; 95% CI $\approx (4.6121, 5.6725)$.

Code (computes t-based 95% CI):

```
import numpy as np  
from scipy import stats
```

```
def mean_confidence_interval(data, confidence=0.95):
```

```

n = len(data)
mean = np.mean(data)
se = np.std(data, ddof=1)/np.sqrt(n)
h = se * stats.t.ppf((1+confidence)/2., n-1)
return mean, mean-h, mean+h

```

Q9: Sample mean and 95% CI (example sample n=!



Question 10: Perform a Chi-square goodness-of-fit test using Python to compare observed and expected distributions, and explain the outcome.

Answer:

Observed counts = [50, 30, 10, 10]

Expected counts (uniform) = [25.0, 25.0, 25.0, 25.0]

Chi-square statistic = 44.0000; p-value = 0.0000.

Interpretation: With p-value = 0.0000, at $\alpha = 0.05$ we reject the null hypothesis that observed frequencies come from the expected (uniform) distribution.

```

from scipy import stats
observed = np.array([50,30,10,10])
expected = np.array([0.25,0.25,0.25,0.25]) * observed.sum()
chi2_stat, p_val = stats.chisquare(f_obs=observed, f_exp=expected)
# chi2_stat, p_val

```

Q10: Observed vs Expected counts (Chi-square tes

