

Inferential Statistics - Hypothesis Testing.

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1 Learning Objectives.

- Compute and summarize group statistics for hypothesis testing:
 - Learn to calculate sample sizes (n_i), means \bar{x}_i , and standard deviations s_i for different groups, which serve as the foundation for conducting **t - tests**, **z - tests**, **ANOVA**, and **Chi - square Tests**.
 - Organize and interpret data for decision making: Understand how to structure group-level data in tables to compare means and variances across regions, enabling proper formulation and testing of null and alternative hypotheses using both single-sample and two-sample tests.
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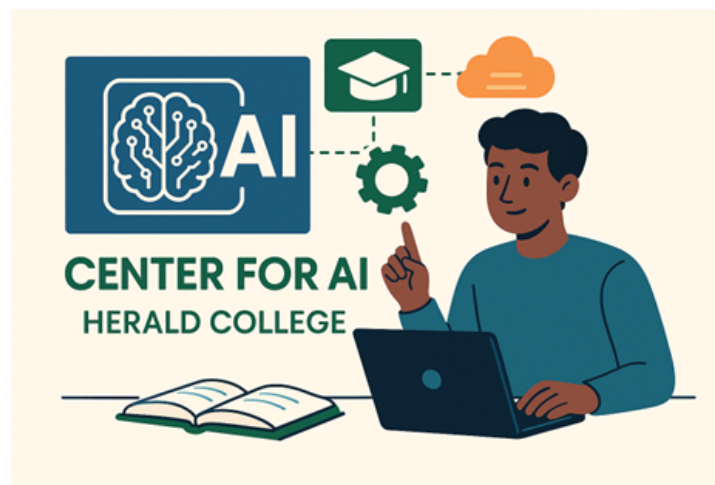


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2 Confidence Interval Approach to Hypothesis Testing:

2.1 How to ?

Confidence Interval Hypothesis Testing Procedure

1. State Hypotheses:

- $H_0 : \mu = \mu_0$ (NullHypothesis)
- $H_1 : \mu \neq \mu_0$ (AlternativeHypothesis)

2. Choose Significance Level:

- $\alpha = 0.05(95\%CI)$ or $\alpha = 0.01(99\%CI)$

3. Check Assumptions:

- Sample size, normality, data conditions for CI method

4. Find Critical Value:

- Using z^* (normal) or t^* (t-distribution) for given α

5. Compute Confidence Interval:

- $CI = \bar{x} \pm (\text{critical value} \times SE)$

6. Decision Rule:

if $\mu_0 \notin CI$ then
 Reject H_0
 \Rightarrow Evidence for H_1 ($\mu \neq \mu_0$)

else
 Fail to reject H_0
 \Rightarrow No significant difference found

7. Interpretation:

- State conclusion in context of the research question

2.2 Exercises:

Problem 1 — Email Campaign: mean purchase amount

A marketing team tests a new email layout. A random sample of $n = 40$ recipients shows an average purchase amount of $\bar{x} = \$52$ with sample standard deviation $s = \$12$. Test whether the true mean purchase amount differs from $\$50$ at the $\alpha = 0.05$ level.

Problem 1 — Sample Solution (Email Campaign)

1. Hypotheses Statement:

$$H_0 : \mu = 50 \quad \text{vs.} \quad H_a : \mu \neq 50$$

2. Choose test method, significance level and check assumptions:

- two tailed test, with significance level of $\alpha = 0.05$.
- Unknown population standard deviation, thus use **t – distributions** with $df = n - 1 = 40 - 1 = 39$.

3. Find Critical Value:

$$SE = \frac{s}{\sqrt{n}} = \frac{12}{\sqrt{40}} = \frac{12}{6.3246} \approx 1.8974.$$

Two-sided **95%** CI uses $t_{0.975,39}$. From tables or python - scipy.stats:

$$t_{0.975,39} \approx 2.0227.$$

4. Compute Confidence Interval:

- **Margin of Error:**

$$ME = t_{0.975,39} \times SE \approx 2.0227 \times 1.8974 \approx 3.838.$$

- **Confidence interval:**

$$\bar{x} \pm ME = 52 \pm 3.838 \Rightarrow (48.162, 55.838).$$

5. Decision by CI approach:

The null value **50** lies inside the **95%** CI **(48.16, 55.84)**. Therefore **Fail to reject H_0** at $\alpha = 0.05$.

- #### 6. Business Interpretation:
- Based on the sample, we do not have sufficient evidence (**at 5%**) to conclude the new email layout changes the average purchase amount from **\$50**. The observed mean **\$52** is plausible under the claim of **\$50**.

Problem 2 — Delivery time (one-sided)

A logistics manager tests whether a new routing algorithm reduces mean delivery time. From a sample of $n = 25$ shipments the mean delivery time is $\bar{x} = 3.8$ days with $s = 0.9$ days. Test $H_0 : \mu = 4$ vs $H_a : \mu < 4$ at $\alpha = 0.05$ using the CI approach.

Problem 3 — App feature adoption (proportion)

A product analyst samples $n = 500$ users; $x = 290$ have adopted a new feature ($\hat{p} = 0.58$). Test whether the true adoption proportion differs from 0.60 at $\alpha = 0.05$ (two-sided) using the CI approach.

Problem 4 — Customer satisfaction benchmark (one-sided, high confidence)

A CX team measures customer satisfaction (1–5 scale). From $n = 100$ responses the sample mean is $\bar{x} = 4.20$ with $s = 0.80$. Test whether the true mean exceeds the benchmark of 4.00 at the $\alpha = 0.01$ level using the CI approach.

Problem 5 — NFL linebacker weights (small sample)

Six randomly selected NFL linebackers have weights (lbs): **243, 238, 229, 253, 248, 225**. Given $\bar{x} = 239.3$ and $s = 10.9$, test whether the mean weight differs from **230** lb at $\alpha = 0.05$ using the CI approach.

3 Critical and p - Value Approach:

3.1 Hypothesis Testing Step for t or z Test:

Step 1 - State Hypothesis:

- Null Hypothesis:

$$H_0 : \text{No effect or no difference.} \rightarrow \mu = \mu_0$$

- Alternative Hypothesis:

$$H_a : \text{There is an effect or Difference.}$$

- Two - tailed: $\mu \neq \mu_0$
- One - Tailed (right): $\mu > \mu_0$
- One - Tailed (left) $\mu < \mu_0$

Step 2 - Choose Significance Level:

- Common choices: $\alpha = 0.05$ Or; $\alpha = 0.01$

Step 4 - Compute Test Statistics:

- Check and Validate assumptions regarding Normality, sample size requirements, independence.
- t - statistic formula: When population variance is unknown and sample sizes are ≤ 30 .

t-statistic Formula

$$t = \frac{\bar{x} - \mu_0}{s/\sqrt{n}}$$

Where:

- \bar{x} = sample mean
- μ_0 = hypothesized population mean
- s = sample standard deviation
- n = sample size
- df = n - 1 (degrees of freedom)
- z - statistic formula: When population variance is known and sample sizes are ≥ 30 .

z-statistic Formula

$$z = \frac{\bar{x} - \mu_0}{\sigma/\sqrt{n}}$$

Where σ = population standard deviation (known)

Step 4 - Determine Critical Values:

Critical Values using Python - scipy .stats.

```
from scipy import stats
if test_type.lower() == "z":
    if tails == "two-tailed":
        crit_val = stats.norm.ppf(1 - alpha/2)
    elif tails == "right-tailed":
        crit_val = stats.norm.ppf(1 - alpha)
    elif tails == "left-tailed":
        crit_val = stats.norm.ppf(alpha)
else: # T-test
    if tails == "two-tailed":
        crit_val = stats.t.ppf(1 - alpha/2, df)
    elif tails == "right-tailed":
        crit_val = stats.t.ppf(1 - alpha, df)
    elif tails == "left-tailed":
        crit_val = stats.t.ppf(alpha, df)
```

Step 5 - Decision by Critical Value:

- For Two Tailed Test:

Reject H_0 if $|z| > z_{\alpha/2}$ or $|t| > t_{\alpha/2, df}$

- For Right - Tailed Test:

Reject H_0 if $z > z_{\alpha}$ or $t > t_{\alpha, df}$

- For Left - Tailed Test:

Reject H_0 if $z < -z_{\alpha}$ or $t < -t_{\alpha, df}$

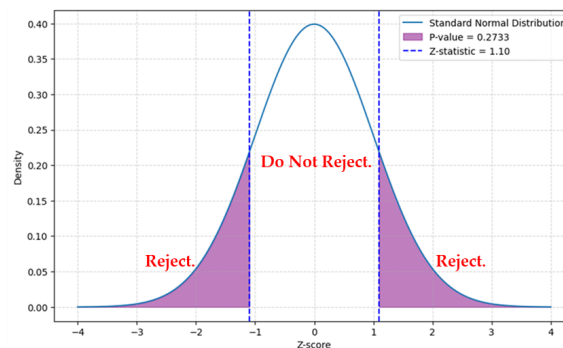


Figure 1: Example Representation of Reject or Not Reject Area on Z - test.

Step 6 - Verification by p - value:

Computing p - value using scipy stats.

```
from scipy import stats
if test_type.lower() == "z":
    if tails == "two-tailed":
        p_value = 2 * (1 - stats.norm.cdf(abs(test_stat)))
    elif tails == "right-tailed":
        p_value = 1 - stats.norm.cdf(test_stat)
    elif tails == "left-tailed":
        p_value = stats.norm.cdf(test_stat)
else: # T-test
    if tails == "two-tailed":
        p_value = 2 * (1 - stats.t.cdf(abs(test_stat), df))
    elif tails == "right-tailed":
        p_value = 1 - stats.t.cdf(test_stat, df)
    elif tails == "left-tailed":
        p_value = stats.t.cdf(test_stat, df)
```

- If $p \leq \alpha$ or equivalently, the test statistic is in the rejection region.
 - Reject Null Hypothesis H_0 .
 - Conclude the result H_a : is **Statistically Significant**.
- If $p > \alpha$ or equivalently, the test statistic is not in the rejection region.
 - Fail to Reject Null Hypothesis H_0 .
 - Conclude the result H_a : is **not Statistically Significant** at the chosen significance level α .

3.2 Algorithm:

1. State Hypotheses:

Two-tailed:

$$H_0 : \mu = \mu_0, \quad H_1 : \mu \neq \mu_0$$

Right-tailed:

$$H_0 : \mu \leq \mu_0, \quad H_1 : \mu > \mu_0$$

Left-tailed:

$$H_0 : \mu \geq \mu_0, \quad H_1 : \mu < \mu_0$$

2. Choose Significance Level:

$$\alpha = 0.05 \quad \text{or} \quad 0.01$$

3. Check Assumptions:

Normality, sample size requirements, independence

4. Calculate Test Statistic:

$$z = \frac{\bar{x} - \mu_0}{SE} \quad \text{or} \quad t = \frac{\bar{x} - \mu_0}{s/\sqrt{n}}$$

5. Find Critical Value:

z^* or t^* for given α

6. Decision Rule:

- **Two-tailed:** If $|z| > z^*$ or $|t| > t^* \Rightarrow \text{Reject } H_0 \Rightarrow \text{Significant}$
- **Right-tailed:** If $z > z^*$ or $t > t^* \Rightarrow \text{Reject } H_0 \Rightarrow \text{Significant}$
- **Left-tailed:** If $z < -z^*$ or $t < -t^* \Rightarrow \text{Reject } H_0 \Rightarrow \text{Significant}$
- **Else** $\Rightarrow \text{Fail to reject } H_0 \Rightarrow \text{Not significant}$

7. Verification by p-value:

- Two-tailed: $p = 2 \times [1 - T(|\text{statistic}|)]$
- Right-tailed: $p = 1 - T(\text{statistic})$
- Left-tailed: $p = T(\text{statistic})$

If $p < \alpha \Rightarrow \text{Reject } H_0$ else $\Rightarrow \text{Fail to reject } H_0$

8. Interpretation:

Summarize in context of problem

3.3 Exercises:

Problem A - Retail Price Test:

A retailer advertises that the average basket value is \$75. A quality analyst randomly samples $n = 12$ recent transactions and finds a sample mean of \$79.8 with sample standard deviation $s = \$9.6$. Test at $\alpha = 0.05$ whether the true mean basket value differs from \$75.

Problem B - Call Center Wait Time.

A call-center manager claims average wait time is at most **2.5 minutes**. A random sample of $n = 16$ calls yields **mean – wait = 2.8min**, $s = 0.9\text{min}$. Test at $\alpha = 0.10$ whether the mean wait time exceeds **2.5minutes**.

Problem C - Email Click - Through.

A marketing team believes the click-through rate (CTR) of a new campaign is 8%. In an **A/B trial**, the new campaign got $x = 78$ clicks out of $n = 900$ impressions. Test at $\alpha = 0.05$ whether the CTR differs from 8% (two-sided).

Part D - Delivery Reliability.

A logistics KPI target is that on-time delivery **rate = 95%**. Over one quarter, a sample of $n = 1200$ deliveries shows **1,100** on time. Use $\alpha = 0.01$ to test whether the true on-time rate differs from 95% (two-sided).

Problem E - A/B Test Revenue Lift (One - sided, unequal variances, two - sample t/welch)

A product team runs an A/B test to check a new checkout flow.

- **Group A (control):** $n_1 = 45$, mean revenue per user $\bar{x}_1 = \$24.50$, $s_1 = \$7.1$
- **Group B (treatment):** $n_2 = 50$, mean revenue per user $\bar{x}_2 = \$27.10$, $s_2 = \$8.0$

Test at $\alpha = 0.05$ whether the new flow increases average revenue **one – sided**, $\mu_2 > \mu_1$, using the critical-value (Welch) approach.

Problem E — Sample Solution (Retail Price Test)

1. Provided Data:

- **For Group - Control (A):**

$$n_1 = 45; \bar{x}_1 = 24.50; s_1 = 7.2$$

- **For Group - Treatment (B):**

$$n_2 = 50; \bar{x}_2 = 27.10; s_2 = 8.0$$

2. Hypotheses Statement:

$$H_0 : \mu_2 = \mu_1 \quad \text{vs.} \quad H_a : \mu_2 > \mu_1$$

3. Choose test method, significance level and check assumptions:

- one right tailed test, with significance level of $\alpha = 0.05$.
-

4. Compute Test Statistic (welch - statistic):

$$SE = \sqrt{\frac{s_1^2}{n_1} + \frac{s_2^2}{n_2}} = \sqrt{\frac{7.2^2}{45} + \frac{8.0^2}{50}} \approx 1.55949$$

Degrees of Freedom - Welch - Satterthwaite:

$$df \approx \frac{(s_1^2/n_1 + s_2^2/n_2)^2}{\frac{(s_1^2/n_1)^2}{n_1-1} + \frac{(s_2^2/n_2)^2}{n_2-1}} \approx 93.0$$

5. Compute Critical Value:

- Two-sided **95%** CI uses $t_{0.975,11}$.
- From tables or python - `stats.t.ppf(1 - alpha, df)`
- $t_{crit} = t_{0.975,93} = 1.6614$

6. Decision by Critical value approach:

$$|t| > t_{crit} \text{ i.e. } 1.6672 > 1.6614 \Rightarrow \text{Reject Null}(H_0)$$

7. Verified by p - value: Compute p - value:

- For two tailed test p - value is:

$$p - \text{value} = P(T \geq |t_{observed}|)$$

T follows a t - distribution with df degrees of freedom.

- Using python and `stats.t.cdf(t_statistic, df)`
- **Decision:** since $p - \text{value} \leq \alpha$, we Reject **Null**(H_0). $\rightarrow p - \text{value} = 0.05 \leq \alpha = 0.05$.

4 Test of Summary Statistics

4.1 One-Way ANOVA Testing Steps

1. State Hypotheses:

$$H_0 : \mu_1 = \mu_2 = \cdots = \mu_k \quad (\text{All group means are equal})$$

$$H_a : \text{At least one group mean differs}$$

2. Choose Significance Level: Select α (e.g., $\alpha = 0.05$).

3. Calculate Group Means and Overall Mean:

Let \bar{X}_i be the mean of group i , and \bar{X}_{GM} the grand mean.

(a) Group mean for group i :

$$\bar{X}_i = \frac{1}{n_i} \sum_{j=1}^{n_i} X_{ij}$$

where:

- X_{ij} = j -th observation in group i
- n_i = number of observations in group i

(b) Grand mean (mean across all observations in all groups):

$$\bar{X}_{GM} = \frac{1}{N} \sum_{i=1}^k \sum_{j=1}^{n_i} X_{ij}$$

where:

- k = number of groups
- $N = \sum_{i=1}^k n_i$ = total number of observations across all groups

4. Compute ANOVA Components:

• Between-group variability:

$$SS_B = \sum_{i=1}^k n_i (\bar{X}_i - \bar{X})^2$$

• Within-group variability:

$$SS_W = \sum_{i=1}^k \sum_{j=1}^{n_i} (X_{ij} - \bar{X}_i)^2$$

• Compute Mean Squares:

$$MS_B = \frac{SS_B}{k-1}, \quad MS_W = \frac{SS_W}{N-k}$$

5. Compute F-statistic:

$$F = \frac{MS_B}{MS_W}$$

6. Decision Rule (Critical Value Approach):

- If $F > F_{(k-1, N-k)}^*$ at significance level α , reject H_0 .

7. Verification by p-value:

- If $p < \alpha$, **reject** H_0 (statistically significant).
- Else, **fail to reject** H_0 (not significant).

4.2 Chi-Square Goodness-of-Fit Test

Goal: Test whether observed frequencies match expected frequencies.

1. State Hypothesis:

H_0 : Observed frequencies follow the expected distribution

H_a : Observed frequencies do not follow the expected distribution.

2. Test Statistics:

$$\chi^2 = \sum_{i=1}^k \frac{(O_i - E_i)^2}{E_i}$$

where:

- O_i = Observed frequency for category i
- E_i = Expected frequency for category i
- k = Number of categories.

3. Degrees of Freedom:

$$df = k - 1$$

where:

- k = Number of Groups.

4. Decision Rule:

- **Reject Null Hypothesis** H_0 if $\chi^2 > \chi_{\text{critical}, df, \alpha}^2$
- Otherwise, **fail to reject** H_0

5. Verification by p-value:

- If $p < \alpha$, **reject** H_0 (statistically significant).
- Else, **fail to reject** H_0 (not significant).

4.3 Chi-Square Test of Independence Steps

1. State Hypotheses:

H_0 :No association between the two categorical variables

H_a :There is an association between the two categorical variables.

2. Choose Significance Level: Select α (e.g., $\alpha = 0.05$).

3. Create a Contingency Table: Organize observed frequencies O_{ij} in a table.

4. Calculate Expected Frequencies:

$$E_{ij} = \frac{(\text{Row Total})_i \times (\text{Column Total})_j}{\text{Grand Total}}$$

5. Compute Chi-Square Statistic:

$$\chi^2 = \sum_i \sum_j \frac{(O_{ij} - E_{ij})^2}{E_{ij}}$$

6. Decision Rule (Critical Value Approach):

- If $\chi^2 > \chi^2_{(\alpha, df)}$ where $df = (r - 1)(c - 1)$, **reject** H_0 .

7. Verification by p-value:

- If $p < \alpha$, **reject** H_0 (statistically significant).
- Else, **fail to reject** H_0 (not significant).

4.4 Exercises:

Problem A — Which Creative Wins? Comparing Mean CTR Across Ads:

Fresh-mart tested four ad creatives in an email campaign. For each group, you collected CTR (%) from equal sub - samples.

- **Given (Summary Stats:**
 - Group A: $n_1 = 30, \bar{x}_1 = 5.2, s_1 = 0.9$
 - Group B: $n_2 = 30, \bar{x}_2 = 5.7, s_2 = 1.0$
 - Group C: $n_3 = 30, \bar{x}_3 = 5.1, s_3 = 0.8$
 - Group D: $n_4 = 30, \bar{x}_4 = 5.9, s_4 = 1.1$
- Test at $\alpha = 0.05$ using one - way ANOVA.

Problem A - sample solution (marketing creatives):

1. Provided Data:

Group A: $n_1 = 30, \bar{x}_1 = 5.2, s_1 = 0.9$

Group B: $n_2 = 30, \bar{x}_2 = 5.7, s_2 = 1.0$

Group C: $n_3 = 30, \bar{x}_3 = 5.1, s_3 = 0.8$

Group D: $n_4 = 30, \bar{x}_4 = 5.9, s_4 = 1.1$

2. Hypothesis Statement:

$$H_0 : \mu_A = \mu_B = \mu_C = \mu_D \quad \text{vs.} \quad H_a : \text{at least one mean differs}$$

3. Compute Grand Mean:

$$\bar{X}_G = \frac{\sum_{i=1}^4 n_i \bar{X}_i}{\sum_{i=1}^4 n_i} = \frac{30(5.2 + 5.7 + 5.1 + 5.9)}{120} = 5.475$$

4. Sum of squares between (SSB):

$$SSB = \sum_{i=1}^4 n_i (\bar{X}_i - \bar{X}_G)^2$$

Compute for each term:

Table 1: Computation of Sum of Squares Between (SSB)

Group(i)	\bar{X}_i	$\bar{X}_i - \bar{X}_G$	$(\bar{X}_i - \bar{X}_G)^2$	$n_i(\bar{X}_i - \bar{X}_G)^2$
1	5.2	-0.275	0.075625	2.26875
2	5.7	0.225	0.050625	1.51875
3	5.1	-0.375	0.140625	4.21875
4	5.9	0.425	0.180625	5.41875
SSB				13.425

5. Sum of squares within (SSW) from sample standard deviations:

$$SSW = \sum_{i=1}^4 (n_i - 1)s_i^2$$

$$\begin{aligned} SSW &= 29(0.9^2) + 29(1.0^2) + 29(0.8^2) + 29(1.1^2) \\ &= 29(0.81 + 1.00 + 0.64 + 1.21) = 29(3.66) = 106.14 \end{aligned}$$

6. Degrees of freedom:

$$df_B = k - 1 = 4 - 1 = 3, \quad df_W = N - k = 120 - 4 = 116.$$

7. Mean Squares:

$$MSB = \frac{SSB}{df_B} = \frac{13.425}{3} = 4.475, \quad MSW = \frac{SSW}{df_W} = \frac{106.14}{116} \approx 0.915$$

8. Compute F - statistic:

$$F = \frac{MSB}{MSW} = \frac{4.475}{0.915} \approx 4.891$$

9. p-value (from F-distribution): Using $df_1 = 3$, $df_2 = 116$:

$$p = P(F_{3,116} \geq 4.891) \approx 0.00308$$

using python and scipy.stats - `p_value = stats.f.sf(F_value, df_between, df_within)`

10. Decision at $\alpha = 0.05$:

$$\text{Since } \rightarrow p \approx 0.0031 < 0.05 \quad \boxed{\text{Reject } H_0}$$

11. **Business Interpretation:** There is strong evidence that at least one creative's mean CTR differs from the others. Next step: perform post-hoc pairwise comparisons (e.g., Tukey HSD) to identify which creatives differ and by how much.

Problem B - Regional Sales Performance

A retailer collected the following average order values (\$) from three regions:

North (15 stores)	South (12 stores)	West (18 stores)
82.50	76.80	88.10
85.30	74.20	90.50
79.80	78.50	85.70
83.10	75.90	92.30
81.40	77.30	87.60

Tasks:

1. Complete the summary statistics table:

Region	number of sample n_i	Mean \bar{x}_i	Standard Deviation s_i
North	15 stores		
South	12 stores		
West	18 stores		

2. Perform ANOVA at $\alpha = 0.01$

Problem C — Channel Mix Shift - Has the Sales Channel Mix Changed?

- **Scenario:** Historically, order share by channel is:
 - **Web:** 60%
 - **App:** 30%
 - **Phone:** 10%
- During a new campaign week ($N = 500$), observed orders were:
 - **Web** = 275
 - **App** = 185
 - **Phone** = 40
- Test at $\alpha = 0.05$ using a **chi-square goodness-of-fit test**.

5 Appendix - Sample Python Implementations.

5.1 T Test and Z Test:

Hypothesis Testing with Python

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

def hypothesis_test(sample_mean, mu0, s=None, n=None, sigma=None, alpha=0.05,
                    tails="two-tailed", test_type="t", plot=True):
    """
    General hypothesis testing with Z-test or T-test, including critical value, p-value, and
    visualization.
    """
    """
    Parameters:
        sample_mean : float
            Sample mean
        mu0 : float
            Hypothesized mean
        s : float, optional
            Sample standard deviation (needed for t-test)
        n : int
            Sample size
        sigma : float, optional
            Population standard deviation (needed for z-test)
        alpha : float
            Significance level
        tails : str
            'two-tailed', 'right-tailed', or 'left-tailed'
        test_type : str
            'z' for Z-test, 't' for T-test

    """

    # --- Step 1: Test Statistic ---
    if test_type.lower() == "z":
        if sigma is None:
            raise ValueError("Population standard deviation (sigma) required for Z-test")
        se = sigma / np.sqrt(n)
        test_stat = (sample_mean - mu0) / se
        df = None
    elif test_type.lower() == "t":
        if s is None or n is None:
            raise ValueError("Sample std deviation (s) and sample size (n) required for T-test")
        se = s / np.sqrt(n)
        test_stat = (sample_mean - mu0) / se
        df = n - 1
    else:
        raise ValueError("test_type must be 'z' or 't'")

    # --- Step 2: Critical Value ---
    if test_type.lower() == "z":
        if tails == "two-tailed":
            crit_val = stats.norm.ppf(1 - alpha/2)
```

```

        crit_val_lower = stats.norm.ppf(alpha/2)
        crit_val_upper = crit_val
    elif tails == "right-tailed":
        crit_val = stats.norm.ppf(1 - alpha)
        crit_val_lower = None
        crit_val_upper = crit_val
    elif tails == "left-tailed":
        crit_val = stats.norm.ppf(alpha)
        crit_val_lower = crit_val
        crit_val_upper = None
else: # T-test
    if tails == "two-tailed":
        crit_val = stats.t.ppf(1 - alpha/2, df)
        crit_val_lower = stats.t.ppf(alpha/2, df)
        crit_val_upper = crit_val
    elif tails == "right-tailed":
        crit_val = stats.t.ppf(1 - alpha, df)
        crit_val_lower = None
        crit_val_upper = crit_val
    elif tails == "left-tailed":
        crit_val = stats.t.ppf(alpha, df)
        crit_val_lower = crit_val
        crit_val_upper = None

# --- Step 3: Decision Rule (Critical Value Method) ---
if tails == "two-tailed":
    reject_cv = abs(test_stat) > crit_val
    decision_cv = "Reject H" if reject_cv else "Do not reject H"
elif tails == "right-tailed":
    reject_cv = test_stat > crit_val
    decision_cv = "Reject H" if reject_cv else "Do not reject H"
elif tails == "left-tailed":
    reject_cv = test_stat < crit_val
    decision_cv = "Reject H" if reject_cv else "Do not reject H"

# --- Step 4: P-value ---
if test_type.lower() == "z":
    if tails == "two-tailed":
        p_value = 2 * (1 - stats.norm.cdf(abs(test_stat)))
    elif tails == "right-tailed":
        p_value = 1 - stats.norm.cdf(test_stat)
    elif tails == "left-tailed":
        p_value = stats.norm.cdf(test_stat)
else: # T-test
    if tails == "two-tailed":
        p_value = 2 * (1 - stats.t.cdf(abs(test_stat), df))
    elif tails == "right-tailed":
        p_value = 1 - stats.t.cdf(test_stat, df)
    elif tails == "left-tailed":
        p_value = stats.t.cdf(test_stat, df)

reject_pv = p_value < alpha
decision_pv = "Reject H" if reject_pv else "Do not reject H"

# --- Step 5: Plot ---
if plot:

```

```

x = np.linspace(-5, 5, 1000) if test_type.lower() == "t" else np.linspace(-4,4,1000)
y = stats.t.pdf(x, df) if test_type.lower() == "t" else stats.norm.pdf(x)
plt.figure(figsize=(8,4))
plt.plot(x, y, label=f'{test_type.upper()} distribution')
# Rejection regions
if tails == "two-tailed":
    plt.fill_between(x, 0, y, where=(x <= crit_val_lower) | (x >= crit_val_upper), color='red',
        alpha=0.3, label="Rejection region")
elif tails == "right-tailed":
    plt.fill_between(x, 0, y, where=(x >= crit_val_upper), color='red', alpha=0.3, label="
        Rejection region")
elif tails == "left-tailed":
    plt.fill_between(x, 0, y, where=(x <= crit_val_lower), color='red', alpha=0.3, label="
        Rejection region")

plt.axvline(test_stat, color='blue', linestyle='--', linewidth=2, label=f"Test statistic ({
    test_stat:.2f})")
if crit_val_lower is not None:
    plt.axvline(crit_val_lower, color='red', linestyle='-', linewidth=2, label=f"Critical value
        ({crit_val_lower:.2f})")
if crit_val_upper is not None:
    plt.axvline(crit_val_upper, color='red', linestyle='-', linewidth=2)

plt.title(f"Hypothesis Test Visualization\nCritical Value Decision: {decision_cv}, P-value
    Decision: {decision_pv}")
plt.xlabel("Test Statistic")
plt.ylabel("Density")
plt.legend()
plt.grid(True)
plt.show()

return {
    "test_statistic": test_stat,
    "critical_value": crit_val,
    "reject_by_critical_value": reject_cv,
    "decision_critical_value": decision_cv,
    "p_value": p_value,
    "reject_by_p_value": reject_pv,
    "decision_p_value": decision_pv,
    "df": df
}

# ----- Example Test -----
# One-tailed right T-test - Adjusted to get test statistic between 2 and 3
result = hypothesis_test(sample_mean=46.8, mu0=45, s=8, n=100, alpha=0.05,
    tails="right-tailed", test_type="t")

print("\nHypothesis Test Results:")
print(f"Test Statistic: {result['test_statistic']:.4f}")
print(f"Critical Value: {result['critical_value']:.4f}")
print(f"P-value: {result['p_value']:.4f}")
print(f"Decision (Critical Value): {result['decision_critical_value']}")
print(f"Decision (P-value): {result['decision_p_value']}")

```

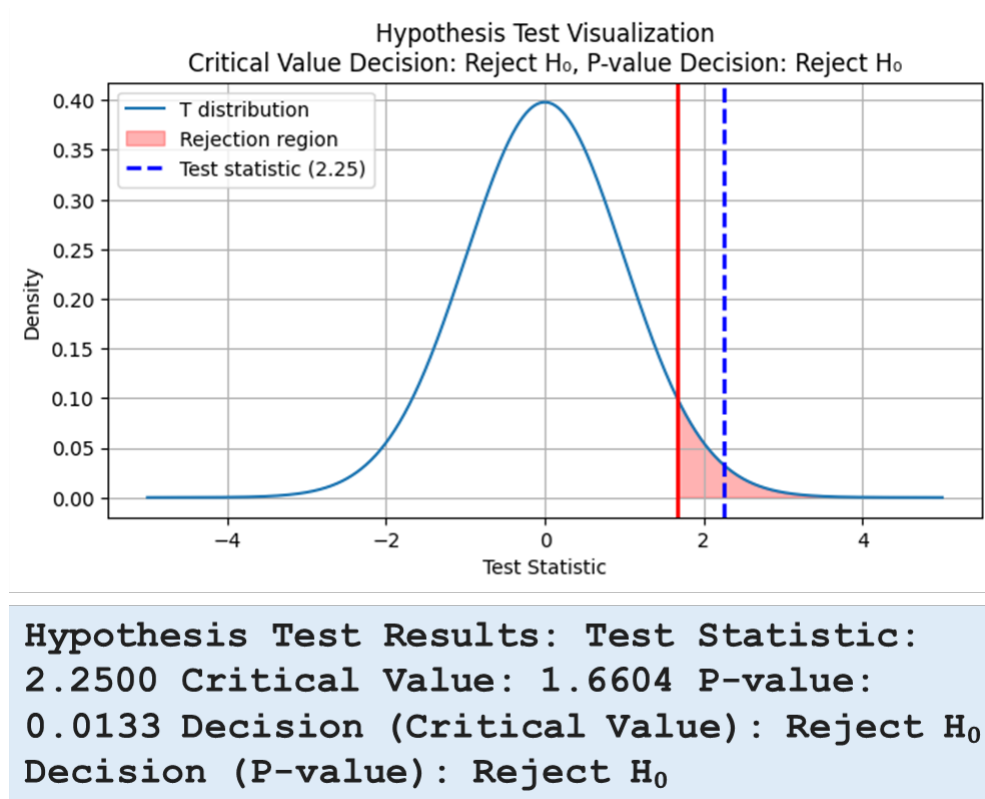


Figure 2: Sample output T - Test

5.2 One - Way ANOVA Test:

One-Way-ANOVA with Python

```
import numpy as np
import matplotlib.pyplot as plt
from scipy.stats import f, stats

# Sample data: click-through rates (%) for 4 ad creatives
# Each group represents CTRs from 10 randomly sampled customers
group_A = np.array([5.2, 5.5, 5.1, 5.4, 5.3, 5.6, 5.2, 5.5, 5.4, 5.3])
group_B = np.array([5.7, 5.8, 5.6, 5.9, 5.5, 5.6, 5.7, 5.8, 5.7, 5.6])
group_C = np.array([5.1, 5.0, 5.2, 5.1, 5.3, 5.0, 5.2, 5.1, 5.3, 5.0])
group_D = np.array([5.9, 6.0, 5.8, 5.7, 5.9, 5.8, 5.9, 6.0, 5.8, 5.9])

# Perform one-way ANOVA
F_statistic, p_value = stats.f_oneway(group_A, group_B, group_C, group_D)
print(f"F-statistic: {F_statistic:.3f}")
print(f"P-value: {p_value:.4f}")
# Decision at alpha = 0.05
alpha = 0.05
if p_value < alpha:
    print("Reject H0: There is a significant difference among the group means.")
else:
    print("Fail to reject H0: No significant difference among the group means.")

# Degrees of freedom for the F-test in one-way ANOVA
num_groups = 4
```

```

observations_per_group = 10
total_observations = num_groups * observations_per_group
df_numerator = num_groups - 1
df_denominator = total_observations - num_groups
# Calculate critical F-value
critical_f_value = f.ppf(1 - alpha, df_numerator, df_denominator)
# Plotting the F-distribution
x = np.linspace(0, F_statistic + 2, 500) # Adjust upper limit based on F-statistic
y = f.pdf(x, df_numerator, df_denominator)
plt.figure(figsize=(10, 6))
plt.plot(x, y, color='blue', lw=2, label=f'F-distribution (df={df_numerator}, {df_denominator})')
# Shade the rejection region
x_reject = np.linspace(critical_f_value, x[-1], 300) # Shade from critical value to the right
plt.fill_between(x_reject, f.pdf(x_reject, df_numerator, df_denominator), color='red', alpha=0.4,
                label=f'Rejection Region')
# Mark the F-statistic
plt.axvline(F_statistic, color='green', linestyle='--', lw=2, label=f'F-statistic = {F_statistic:.2f}')
# Mark the critical F-value
plt.axvline(critical_f_value, color='red', linestyle='--', lw=2, label=f'Critical F-value = {
    critical_f_value:.2f}')
plt.title('F-Test for One-Way ANOVA with Rejection Region')
plt.xlabel('F-value')
plt.ylabel('Density')
plt.legend()
plt.grid(True)
plt.show()

```

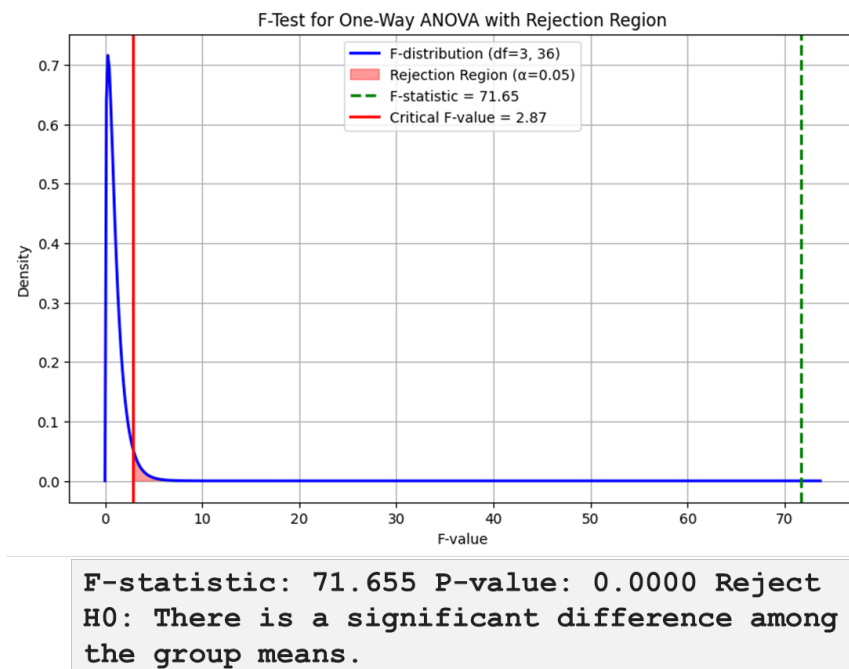


Figure 3: Sample Output One - Way - ANOVA

The - End