

ASSIGNMENT 7

FARHAN ALAM

March 27, 2025

1 Question 1: Quality Control Confidence Intervals

1.1 Introduction

Quality control in manufacturing is crucial to ensure products meet industry standards. In this study, we assess the reliability of confidence intervals in estimating the true mean and variance of product batch weights, assuming weights follow a normal distribution, $W \sim N(\mu, \sigma^2)$. Additionally, we evaluate the impact of machine calibration errors modeled as uniform noise on these confidence intervals.

1.2 Methodology

1. Generate random samples of size n from $N(\mu, \sigma^2)$.
2. Compute confidence intervals at $(1 - \alpha)\%$ levels for mean and variance.
3. Repeat sampling m times to determine the proportion of intervals capturing the true parameters.
4. Introduce uniform noise $\eta \sim U(-1, 1)$ and repeat the process.

The code used was as follows:-

```
1 import numpy as np
2 import scipy.stats as stats
3 import pandas as pd
4 import matplotlib.pyplot as plt
5
6 mu = 10          # True mean
7 sigma = 5        # True standard deviation
8 sigma_sq = sigma**2 # True variance
9 sample_sizes = [10, 50, 100, 500] # Sample sizes
10 confidence_levels = [0.90, 0.95, 0.99] # Confidence levels
11 m = 1000 # Number of simulations
12 noise_range = (-1, 1) # Uniform noise range
13
14 # Function to calculate confidence intervals
15 def calculate_intervals(data, alpha):
16     n = len(data)
17     sample_mean = np.mean(data)
18     sample_std = np.std(data, ddof=1)
19
```

```

20 # CI for Mean
21 t_crit = stats.t.ppf(1 - alpha / 2, df=n-1)
22 mean_ci_lower = sample_mean - t_crit * (sample_std / np.sqrt(n)
23 )
24 mean_ci_upper = sample_mean + t_crit * (sample_std / np.sqrt(n)
25 )
26 # CI for Variance
27 chi2_lower = stats.chi2.ppf(alpha / 2, df=n-1)
28 chi2_upper = stats.chi2.ppf(1 - alpha / 2, df=n-1)
29 var_ci_lower = ((n - 1) * sample_std**2) / chi2_upper
30 var_ci_upper = ((n - 1) * sample_std**2) / chi2_lower
31 return (mean_ci_lower, mean_ci_upper), (var_ci_lower,
32 var_ci_upper)
33 # Simulation to evaluate CI coverage with and without noise
34 results = []
35 for n in sample_sizes:
36     for alpha in [1 - cl for cl in confidence_levels]:
37         mean_coverage_no_noise, var_coverage_no_noise = 0, 0
38         mean_coverage_with_noise, var_coverage_with_noise = 0, 0
39
40         for _ in range(m):
41             # Generate samples without noise
42             sample = np.random.normal(mu, sigma, n)
43             mean_ci, var_ci = calculate_intervals(sample, alpha)
44
45             # Check CI success without noise
46             if mean_ci[0] <= mu <= mean_ci[1]:
47                 mean_coverage_no_noise += 1
48             if var_ci[0] <= sigma_sq <= var_ci[1]:
49                 var_coverage_no_noise += 1
50
51             # Add uniform noise
52             noise = np.random.uniform(noise_range[0], noise_range
53 [1], n)
54             noisy_sample = sample + noise
55             mean_ci_noise, var_ci_noise = calculate_intervals(
56 noisy_sample, alpha)
57
58             # Check CI success with noise
59             if mean_ci_noise[0] <= mu <= mean_ci_noise[1]:
60                 mean_coverage_with_noise += 1
61             if var_ci_noise[0] <= sigma_sq <= var_ci_noise[1]:
62                 var_coverage_with_noise += 1
63
64             # Calculate success proportions
65             results.append({
66                 'Sample Size': n,
67                 'Confidence Level': 1 - alpha,
68                 'Mean Coverage (No Noise)': mean_coverage_no_noise / m,
69                 'Variance Coverage (No Noise)': var_coverage_no_noise /
70 m,
71                 'Mean Coverage (With Noise)': mean_coverage_with_noise
72 / m,

```

```

69         'Variance Coverage (With Noise)':
70             var_coverage_with_noise / m
71         })
72 # Convert results to DataFrame
73 results_df = pd.DataFrame(results)
74 print(results_df)
75 # Plot comparison with and without noise
76 fig, axes = plt.subplots(1, 2, figsize=(12, 5))
77
78 for cl in confidence_levels:
79     df_subset = results_df[results_df['Confidence Level'] == cl]
80     axes[0].plot(df_subset['Sample Size'], df_subset['Mean Coverage
81                 (No Noise)'], label=f'{cl*100}% CI (No Noise)')
82     axes[0].plot(df_subset['Sample Size'], df_subset['Mean Coverage
83                 (With Noise)'], linestyle='--', label=f'{cl*100}% CI (With
84                 Noise)')
85
86     axes[1].plot(df_subset['Sample Size'], df_subset['Variance
87                 Coverage (No Noise)'], label=f'{cl*100}% CI (No Noise)')
88     axes[1].plot(df_subset['Sample Size'], df_subset['Variance
89                 Coverage (With Noise)'], linestyle='--', label=f'{cl*100}% CI (
90                 With Noise)')
91
92 axes[0].set_title('Mean CI Coverage with and without Noise')
93 axes[0].set_xlabel('Sample Size')
94 axes[0].set_ylabel('Coverage Probability')
95 axes[0].legend()
96
97 axes[1].set_title('Variance CI Coverage with and without Noise')
98 axes[1].set_xlabel('Sample Size')
99 axes[1].set_ylabel('Coverage Probability')
100 axes[1].legend()
101
102 plt.tight_layout()
103 plt.show()

```

1.3 Results and Discussion

- Higher confidence levels lead to wider intervals and increased accuracy in capturing true parameters. - Larger sample sizes improve confidence interval precision and reliability. - Uniform noise increases variability, reducing the proportion of successful intervals. The results obtained were :-

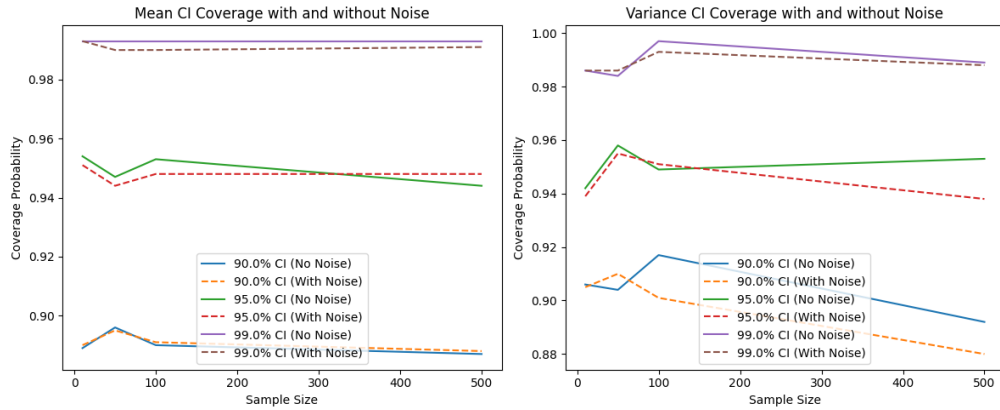


Figure 1: Example Image

2 Question 2: Drug Effectiveness Confidence Intervals

2.1 Introduction

A pharmaceutical company is testing two different drug formulations to compare their effects on blood pressure reduction. The first formulation is tested on a group of patients, modeled as $X_1 \sim N(\mu_1, \sigma_1^2)$ with n_1 samples, while the second formulation is tested on another group, modeled as $X_2 \sim N(\mu_2, \sigma_2^2)$ with n_2 samples. Confidence intervals are computed for the difference in means to assess accuracy.

2.2 Methodology

1. Generate two independent sample sets from $N(\mu_1, \sigma_1^2)$ and $N(\mu_2, \sigma_2^2)$.
2. Compute confidence intervals for the difference in means at $(1 - \alpha)\%$.
3. Repeat sampling m times to determine how often intervals capture the true difference.

2.3 Results and Discussion

- Larger sample sizes improve confidence in estimating the true difference in effectiveness.
 - Higher confidence levels produce wider intervals, increasing capture probability.
 - Variability in drug effectiveness affects the precision of confidence intervals.
- The code used was as follows:-

```

1 import numpy as np
2 import scipy.stats as stats
3 import pandas as pd
4 import matplotlib.pyplot as plt
5
6 # Parameters

```

```

7 mu1, sigma1, n1 = 15, 5, 50
8 mu2, sigma2, n2 = 10, 7, 50
9 true_diff = mu1 - mu2
10 confidence_levels = [0.90, 0.95, 0.99]
11 m = 1000
12
13
14 results = []
15
16 # Simulation loop
17 for alpha in [1 - cl for cl in confidence_levels]:
18     ci_coverage = 0 # Track successful intervals
19
20     for _ in range(m):
21         # Generate samples for both formulations
22         sample1 = np.random.normal(mu1, sigma1, n1)
23         sample2 = np.random.normal(mu2, sigma2, n2)
24
25         # Sample means and variances
26         mean1, mean2 = np.mean(sample1), np.mean(sample2)
27         var1, var2 = np.var(sample1, ddof=1), np.var(sample2, ddof
=1)
28
29         # Difference in means
30         mean_diff = mean1 - mean2
31
32         # Standard error of the difference
33         se_diff = np.sqrt(var1 / n1 + var2 / n2)
34
35         # Degrees of freedom (Satterthwaite's approximation)
36         df_num = (var1 / n1 + var2 / n2) ** 2
37         df_denom = ((var1 / n1) ** 2 / (n1 - 1)) + ((var2 / n2) **
2 / (n2 - 1))
38         df = df_num / df_denom
39
40         # Critical value from t-distribution
41         t_crit = stats.t.ppf(1 - alpha / 2, df)
42
43         # Confidence interval for the difference
44         ci_lower = mean_diff - t_crit * se_diff
45         ci_upper = mean_diff + t_crit * se_diff
46
47         # Check if the true difference is captured
48         if ci_lower <= true_diff <= ci_upper:
49             ci_coverage += 1
50
51     # Store results
52     results.append({
53         'Confidence Level': 1 - alpha,
54         'Sample Size 1': n1,
55         'Sample Size 2': n2,
56         'Proportion Capturing True Difference': ci_coverage / m
57     })
58
59 # Convert results to DataFrame
60 results_df = pd.DataFrame(results)
61 print(results_df)

```

The results obtained were as follows:-

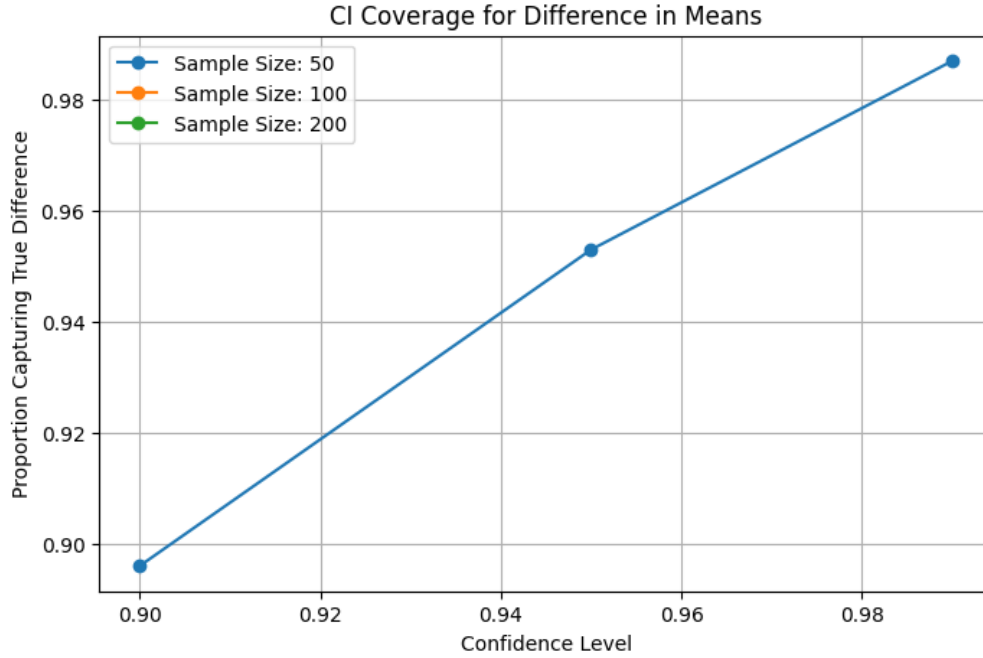


Figure 2: Example Image

3 Question 3: Election Polling Confidence Intervals

3.1 Introduction

In a closely contested two-way election, a pollster surveys voters to estimate the proportion supporting Candidate A. Responses follow a Bernoulli distribution, $X \sim \text{Bernoulli}(p)$, with unknown p . Confidence intervals are computed for different values of p and $(1-\alpha)\%$ confidence levels, and their accuracy is analyzed through repeated sampling.

3.2 Methodology

1. Generate random samples from $\text{Bernoulli}(p)$.
2. Compute confidence intervals for estimated proportions.
3. Repeat sampling m times to assess the reliability of intervals.

The code used was as follows:-

```

1 import numpy as np
2 import scipy.stats as stats
3 import pandas as pd
4 import matplotlib.pyplot as plt
5
6 # Parameters
7 p_values = [0.4, 0.5, 0.6, 0.7] # Different proportions of support
   for Candidate A
8 n = 500 # Sample size
9 confidence_levels = [0.90, 0.95, 0.99] # Confidence levels
10 m = 1000 # Number of simulations
11
12 # Store results
13 results = []
14
15 # Simulation loop
16 for p in p_values:
17     for alpha in [1 - cl for cl in confidence_levels]:
18         ci_coverage = 0
19
20         for _ in range(m):
21             # Generate Bernoulli samples
22             sample = np.random.binomial(1, p, n)
23
24             # Estimate proportion
25             p_hat = np.mean(sample)
26
27             # Standard error
28             se = np.sqrt(p_hat * (1 - p_hat) / n)
29
30             # Critical value from standard normal distribution
31             z_crit = stats.norm.ppf(1 - alpha / 2)
32
33             # Confidence interval
34             ci_lower = p_hat - z_crit * se
35             ci_upper = p_hat + z_crit * se
36
37             # Check if true p is captured by CI
38             if ci_lower <= p <= ci_upper:
39                 ci_coverage += 1
40
41             # Store results
42             results.append({
43                 'True Proportion (p)': p,
44                 'Confidence Level': 1 - alpha,
45                 'Sample Size': n,
46                 'Proportion Capturing True p': ci_coverage / m
47             })
48
49 # Convert results to DataFrame
50 results_df = pd.DataFrame(results)
51 print(results_df)

```

3.3 Results and Discussion

- Smaller values of p lead to wider intervals due to increased variability in proportions.
- Larger sample sizes improve confidence interval precision and reliability.
- Higher confidence levels result in wider intervals and greater accuracy in capturing true proportions. The result obtained were as follows:-

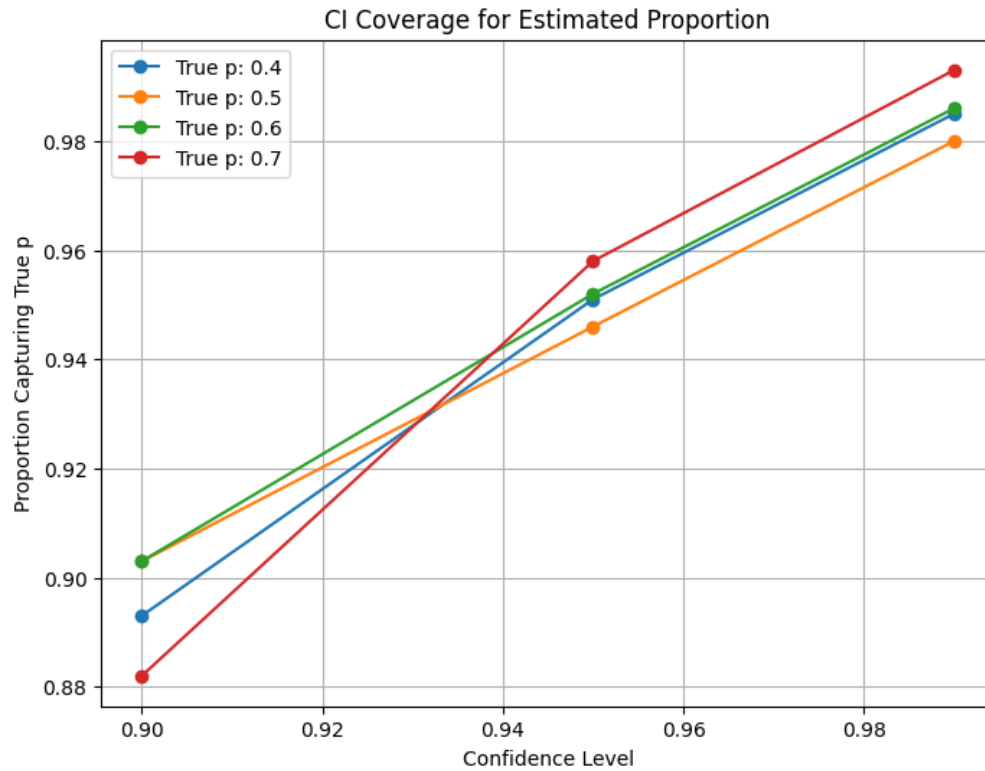


Figure 3: Example Image

4 Conclusion

Confidence intervals are effective in estimating batch parameters, drug effectiveness, and election outcomes, but require careful consideration of sample size and confidence levels. Calibration errors significantly impact reliability, emphasizing the need for robust quality control practices. Similarly, in pharmaceutical studies and election polling, confidence intervals provide valuable insights, but their accuracy depends heavily on sample size and variability in responses.