Introduction to Statistics - Young Researchers Fellowship Program

Lecture 6 - Foundations of Hypothesis Testing

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Introduction to Hypothesis Testing

Null and Alternative Hypotheses

- We use a sample to make inferences about the population
- A hypothesis test helps us determine if there's enough evidence in the sample to support a statement about the population
- Null hypothesis: the baseline assumption
- Alternative hypothesis: the opposite of the null
 - Typically what we "want", the phenomenon being studied

Defining the null

- We define the null by setting our baseline scenario: what we believe might be true in "normal circumstances"
- For instance, we may think that typically the sample mean is of a certain historical value:

$$H_0: \mu = \mu_0$$

- lacksquare μ_0 is typically called "mu naught" or hypothesized mean
 - The hypothetical baseline value

Defining the alternative

- We define the alternative as the contrary to the null
- Usually what we want to look out for (i.e. is the treatment effective?)
- For example, that the historical value is no longer the baseline and the population mean changed.

$$H_0: \mu \neq \mu_0$$

Errors

Errors 0 • 0

Decision	${\cal H}_0$ is ${\sf True}$	H_0 is False	
Reject H_0	Type I Error	Correct Decision	
Fail to reject ${\cal H}_0$	Correct Decision	Type II Error	

Type I and Type II Errors

- **Type I Error** (α): Rejecting the null hypothesis when it's true
 - Example: Null: $\mu \ge 3.0$, we conclude $\mu < 3.0$
 - \blacksquare The probability of making this error is the significance level α
- **Type II Error** (β): Failing to reject the null hypothesis when it's false

Hypothesis testing

General idea of a hypothesis test

- We are asking: "if the sampling distribution is truly distributed according to the null, what is the most extreme value of a sampling exercise I can get to not doubt the null?"
- Means computing a number of standard errors that the sample statistic is located relative to the hypothesized mean.
 - The number of standard errors is called the "test statistic"
- \blacksquare If the null is true, this shouldn't be too far from μ_0
 - If it is far, we have to reject the null
 - Otherwise, "fail to reject" H_0 .
- How far will we tolerate? Based on α , our significance level (related to confidence we want).

p-values

- The *p*-value is the probability associated to the test statistic.
- Answers the question: how likely is it to get this test statistic if the null were true?
- The general rejection rule is:

$$p < \alpha$$

because it means that the p-value shows that it is more likely that we have gotten an extreme value, which does not come from the H_0 scenario, than the tolerance we've set to commit a type-I error.

General type of tests

Case 1: Left-tailed Test (One-side)

- Null hypothesis: population parameter $\mu \ge \mu_0$
- Alternative hypothesis: $\mu < \mu_0$
- Example: Average GPA of students is less than 3.0

Case 2: Right-tailed Test (One-side)

- \blacksquare Null hypothesis: population parameter $\mu \leq \mu_0$
- lacksquare Alternative hypothesis: $\mu>\mu_0$
- Example: Average GPA of students is greater than 3.0

Case 3: Two-tailed Test (Two side)

- \blacksquare Null hypothesis: population parameter $\mu=\mu_0$
- Alternative hypothesis: $\mu \neq \mu_0$
- Example: Average GPA is different from 3.0

Introduction to Hypothesis Testing

One Sample Tests?

- This means we're only working with one sample, not several.
- Comparing a mean against a numerical value.
- Later we will work with many samples.

The z-test (one sample)

- lacktriangle Tests whether the population mean μ is equal to a given value
- Used when population standard deviation is known
- We can work with a normal distribution for computing probabilities

Z-Test Types

- Left-tailed test: $H_0: \mu \ge \mu_0$, $H_1: \mu < \mu_0$
- \blacksquare Right-tailed test: $H_0: \mu \leq \mu_0$, $H_1: \mu > \mu_0$
- \blacksquare Two-tailed test: $H_0: \mu = \mu_0, \, H_1: \mu \neq \mu_0$

- 1 Compute sample mean to be used or use the given one.
- Compute the test statistic. In the case of a Z-test, the test statistic is Z:

$$Z = \frac{x - \mu_0}{\frac{\sigma}{\sqrt{n}}}$$

- f 3 Compute the p-value associated with the test statistic and given lpha
- 4 If the p-value is smaller than α , reject H_0 .

Calculating p-values

■ We need to know how to calculate *p*-values based on each type of test, to accurately reject based on available information.

Test Type	Null Hypothesis (H_0)	Alternative Hypothesis (H_1)	Formula for p-value	R Code Example
	$H_0: \mu =$	$H_1: \mu \neq$	$2{\times}P(Z \geq z_{score})$	2 * (1 -
tailed	$H_0: \mu \geq$	$H_1 : \mu <$	$P(Z \le z_{score})$	<pre>pnorm(abs(z_score)) pnorm(z_score)</pre>
tailed		μ_0	$1 \ (Z \le z_{\text{score}})$	phorm(z_score)
	$H_0: \mu \leq$. 0	$P(Z \geq z_{\rm score})$	1 -
tailed	μ_0	μ_0		pnorm(z_score)

- Test whether population mean is **greater than or equal** to 3.0
 - $H_0: \mu \ge 3.0$
 - $H_1: \mu < 3.0$
 - \blacksquare Sample size = 100, sample mean = 2.8, $\sigma=0.3$
 - $z = \frac{2.8 3.0}{0.03} = -6.67$
- lacktriangle Use p-value approach or critical value approach for rejecting

- Base R does not offer a built-in package for this test, however, we may easily calculate probabilities using pnorm()
- For this, we should know how to calculate the *p*-value as per the table above.

R Implementation for the Example

```
# Given values
mu 0 <- 3.0 # Hypothesized population mean
x bar <- 2.8 # Sample mean
sigma <- 0.3 # Population standard deviation</pre>
n <- 100
        # Sample size
# Calculate the standard error
se <- sigma / sqrt(n)
# Calculate the Z-score
z score <- (x bar - mu 0) / se
# Calculate the p-value for a left-tailed test
p value <- pnorm(z score)</pre>
```

Proportions

- Proportions work just like means.
- \blacksquare Need only to redefine the Z test statistic to follow the modified standard error for \hat{p}

$$Z = \frac{\hat{p} - p_0}{\sqrt{\frac{p_0(1 - p_0)}{n}}}$$

where p_0 is the hypothesized population proportion.

lacksquare Apply the same rules you would for Z-tests.

R Implementation for Proportion Z-tests

```
# Given values
p_hat <- 0.55  # Sample proportion
p_0 <- 0.50  # Hypothesized population proportion (null hypoth
n <- 100  # Sample size

# Calculate the Z-score
z_score <- (p_hat - p_0) / sqrt((p_0 * (1 - p_0)) / n)

# P-value
p_value <- 2 * (1 - pnorm(abs(z_score)))</pre>
```

Errors and Significance Levels, different rejection rules

- **Significance level (** α **)**: Probability of making a Type I error
 - **Example:** $\alpha = 0.05$ means 95% confidence level
- Compare p-value to α or use critical value
- $lue{\alpha}$ Critical values are the number of standard errors associated with the lpha probability under a specific type of test.
 - Rejection rules with these vary by type of test

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α	Left-Tailed	Right-Tailed	Two-Tailed
0.05	-1.645	1.645	$\pm~1.96$
0.01	-2.33	2.33	$\pm~2.58$
0.10	-1.28	1.28	$\pm~1.64$

Test		Critical Value	
Type	Hypothesis (H_0,H_1)	(Z)	Rejection Rule
Left-	$H_0: \mu \geq \mu_0$ vs	z_{α} (negative)	Reject H_0 if
tailed	$H_1 : \mu < \mu_0$		$z < z_{\alpha}$
Right-	$H_0: \mu \leq \mu_0$ vs	z_{lpha} (positive)	Reject H_0 if
tailed	$H_1: \mu > \mu_0$		$z > z_{\alpha}$
Two-	$H_0:\mu=\mu_0$ vs	$z_{lpha/2}$ (positive	Reject H_0 if
tailed	$H_1: \mu \neq \mu_0$	and negative)	$z<-z_{lpha/2}$ or
			$z > z_{\alpha/2}$

Introduction to Hypothesis Testing

T-tests (when we don't know the population standard deviation)

As you might remember, when we cannot obtain enough information for a reliable estimate of the population standard deviation, we use the sample standard deviation as an estimate. This is common in practice when we have small samples or lack population standard deviation.

T-statistic:

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

- $\blacksquare \mu_0$: hypothesized population mean
- Denominator: standard error of the sample mean.

The t distribution is used, which has heavier tails than the normal distribution and depends on degrees of freedom (n-1).

Types of T-Tests

- **Left-tailed test**: Test for means less than the hypothesized value.
- **Right-tailed test**: Test for means greater than the hypothesized value.
- **Two-tailed test**: Test for means different from the hypothesized value.

Example: Left-tailed Test

■ Sample size: 25

■ Sample mean: 9.5

 \blacksquare Sample standard deviation: 2.5

Hypothesized mean: 10Significance level: 5%

Example: Left-tailed test

Null and Alternative Hypotheses:

$$H_0: \mu \ge 10$$

$$H_1: \mu < 10$$

T-statistic:

$$t = \frac{9.5 - 10}{2.5/\sqrt{25}} = -2$$

Critical value at 24 degrees of freedom (5% significance):

$$t_{critical} = -1.711$$

Since t=-2 is less than the critical value, we reject the null hypothesis.

Example: Right-tailed Test

Sample size: 25Sample mean: 9.5

■ Sample standard deviation: 2.5

■ Hypothesized mean: 10

Example: Right-tailed Test

Null and Alternative Hypotheses:

$$H_0: \mu \le 10$$

$$H_1: \mu > 10$$

T-statistic:

$$t = \frac{9.5 - 10}{2.5 / \sqrt{25}} = -2$$

For a right-tailed test, the critical value is $t_{critical}=1.711$. Since t=-2, we fail to reject the null hypothesis.

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■ Sample size: 25 ■ Sample mean: 12

■ Sample standard deviation: 2.5

■ Hypothesized mean: 10

Two-tailed Test Example

Null and Alternative Hypotheses:

$$H_0: \mu = 10$$

$$H_1: \mu \neq 10$$

T-statistic:

$$t = \frac{12 - 10}{2.5 / \sqrt{25}} = 4$$

Critical values are $t_{critical}=\pm 2.064$. Since t=4 exceeds the critical values, we reject the null hypothesis.

Proportions and Sampling Distribution

For categorical data, we compute proportions instead of means. Proportions follow a normal distribution with large enough samples. The sample proportion is calculated as:

$$\hat{p} = \frac{x}{n}$$

Where x is the number of successes and n is the sample size.

Standard error of proportion:

$$SE(\hat{p}) = \sqrt{\frac{p(1-p)}{n}}$$

The Interval Approach for Hypothesis Testing

We can also use confidence intervals for hypothesis testing. If the null hypothesis value falls outside the confidence interval, we reject the null hypothesis.

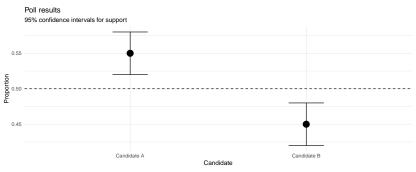


Figure 1: Confidence interval for the proportion of people who support each candidate