

Chapter 6: Introduction to hypothesis testing

(Ott & Longnecker Sections: 5.1, 5.4, 5.6)

<https://dzwang91.github.io/stat324/>



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- null hypothesis
- alternative hypothesis
- test statistic
- rejection region
- type I error
- type II error
- power
- p-value
- significance level



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- **Reasoning:** If you love me (if H_0 is true), you would take the trash out every week and put your socks away.
- **Data:** Some weeks the girl's boyfriend doesn't take the trash out or leave the socks where they fall.
- **Conclusion:** I don't believe you love me (reject the H_0).
- Hypothesis testing concerns on how to use a **random sample** to judge if it is evidence that supports or not the hypothesis.

- In hypothesis testing, there are two competing hypotheses:
 - H_0 : the null hypothesis;
 - H_A : the alternative hypothesis.
- The hypothesis we want to test is if H_A is “likely” true.
- One-tailed tests:

$$H_0 : \theta = \theta_0 \text{ v.s. } H_A : \theta > \theta_0.$$

or

$$H_0 : \theta = \theta_0 \text{ v.s. } H_A : \theta < \theta_0.$$

- Two-tailed test:

$$H_0 : \theta = \theta_0 \text{ v.s. } H_A : \theta \neq \theta_0.$$



- There are two possible outcomes:
 - Reject H_0 because of sufficient evidence in the sample in favor of H_A .
 - Do not reject H_0 because of insufficient evidence to support H_A .
- Note that failure to reject H_0 does not mean the null hypothesis is true. It only means that we do not have sufficient evidence to support H_A .

- 1 State the H_0 and H_A hypotheses
- 2 Devise a test statistic: collect data (X_1, \dots, X_n) and choose a test statistic $T_n = T_n(X_1, \dots, X_n)$. The test statistic is an RV. Based on data, we can calculate the realization of the test statistic.
- 3 Decide the rejection region: specify a set of values of the test statistic such that, if it realizes to one of these values, we reject H_0 . This region is called the rejection region. (how do we specify the rejection region?)
- 4 Conclusion: if the test statistic falls outside of the rejection region, then there is insufficient evidence against H_0 , and we say we fail to reject the null. Otherwise we say reject the null.

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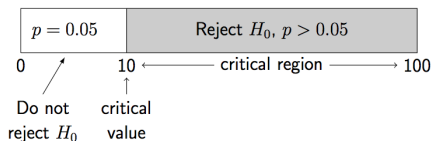
Example






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We are going to use a sample of 100 chips from the production to test. Let X denote the number of defective in the sample of 100. Reject H_0 if $X \geq 10$. Then X is a **test statistic**.



		Decision (based on sample)	
		Reject H_0	Not Reject H_0
Truth (for population studied)	H_0 True	Type I Error α 	
	H_0 False		Type II Error β

- The acceptance of H_A when H_0 is true is called a Type I error. The probability of committing a type I error is called the **level of significance** and is denoted by α :

$$\alpha = P(\text{reject } H_0 \mid H_0 \text{ is true}) = P(T_n \in \text{rejection region} \mid H_0 \text{ is true}).$$

- Smaller α is better. Typically, 0.05 or smaller.



- Use the distribution of the test statistic at H_0 to determine a rejection region: control the type I error at significance level α .
- Failure to reject H_0 when H_A is true is called a Type II error. The probability of committing a type II error is denoted by β .

$$\beta = P(\text{not reject } H_0 \mid H_A \text{ is true}).$$

- Smaller β is better.
- It is usually impossible to compute β unless we have a specific alternative hypothesis.

- Type I error:

$$\begin{aligned}\alpha = P(X \geq 10 \text{ when } p = 0.05) &= \sum_{n=10}^{100} \binom{100}{n} 0.05^n (1 - 0.05)^{100-n} \\ &= 0.0282\end{aligned}$$

- Suppose we have $H_A : p = 0.1$, then

$$\begin{aligned}\beta = P(X < 10 \text{ when } p = 0.1) &= \sum_{n=0}^9 \binom{100}{n} 0.1^n (1 - 0.1)^{100-n} \\ &= 0.4513\end{aligned}$$



$$H_0 : p = 0.05 \text{ v.s. } H_A : p = 0.01.$$

- When rejection region is $X \geq 10$:

$$\alpha = 0.0282, \beta = 0.4513$$

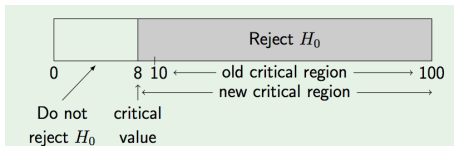
Trade-off between α and β for fixed sample size



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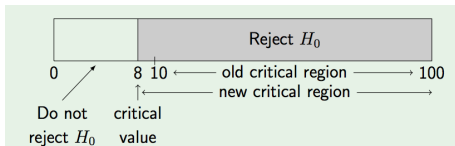
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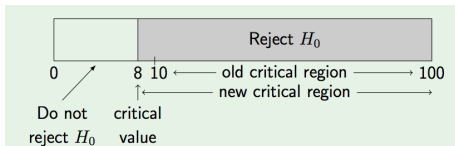
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- When the rejection region is $X \geq 8$:

$$\alpha = 0.128, \beta = 0.206.$$

How can we reduce α and β simultaneously?

- Both α and β can be reduced simultaneously by increasing the sample size.
- Consider that the sample size is $n = 150$ and the critical value is 12. Then, rejection region is $X \geq 12$, where X is the number of defectives in the sample of 150 chips.
 - The type I error is

$$\alpha = \sum_{n=12}^{150} \binom{150}{n} 0.05^n 0.95^{150-n} = 0.074,$$

lower than 0.128 for $n=100$ and critical value of 8.

- The type II error is

$$\beta = \sum_{n=0}^{11} 0.1^n 0.9^{150-n} = 0.171,$$

lower than 0.206 for $n=100$ and critical value of 8.



- The **power** of a test is the probability of rejecting H_0 given that the alternative hypothesis is true. That is,

$$\text{Power} = 1 - \beta = P(\text{reject } H_0 \text{ when } H_0 \text{ is false}).$$



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- Different significance levels require recomputing the rejection region.
- The p-value is defined to be the probability of a test statistic realizing to a value that is as or more extreme than the one actually observed given the null hypothesis is true.
- Smaller p-values indicate relatively more evidence against the null hypothesis.
- If the p-value is smaller than the given significance level α , we would reject the null, otherwise we would not reject the null.

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- Depends! At a significance level of 0.05, we cannot reject H_0 because $p = 0.09 > 0.05$. However, for significance levels greater than or equal to 0.09, we can reject H_0 .

What's the next?



We'll introduce some specific tests based on samples from one population in the next lecture.