

STAT 22000 Lecture Slides

Hypothesis Testing About Population Means

Yibi Huang
Department of Statistics
University of Chicago

Outline

- Hypothesis Testing About Population Means (Section 4.3)
- Relationships Between Confidence Intervals and Hypothesis Tests (Section 4.3.2)
- Common Misunderstandings About Hypothesis Testing (Not in the textbook)

Hypothesis Tests about Population Means

Example: Number of College Applications

To know how many colleges students applied to, the dean of a certain university took a random sample of size 106 from their newly admitted students. This sample yielded an average of 9.7 college applications with a standard deviation of 7. College Board website states that counselors recommend students apply to roughly 8 colleges. Do these data provide convincing evidence that the average number of colleges all freshmen in this university apply to is higher than recommended?

<http://www.collegeboard.com/student/apply/the-application/151680.html>

Example: Number of College Applications – Hypotheses

- *Population*: all freshmen in this university
- The *parameter of interest* μ is the average number of schools applied to by all freshmen in this university
- There are two explanations why the sample mean is higher than the recommended 8 schools.
 - The true population mean is different.
 - The true population mean is 8, and the difference between the true population mean and the sample mean is simply due to natural sampling variability.
- $H_0 : \mu = 8$ (the average number of colleges freshmen in this university have applied to is 8, as recommended)
- $H_A : \mu > 8$ (the average number of colleges freshmen in this university have applied to is > 8)

Wrong Ways to State H_0 and H_A

H_0 and H_A are **ALWAYS** stated in terms of population parameters, not sample statistics

Neither

$$H_0 : \bar{x} = 8, \quad H_A : \bar{x} > 8$$

or

H_0 : average number of colleges applied **in the sample** is 8

H_A : average number of colleges applied **in the sample** is 9.7

is correct. The correct statements should be

$$H_0 : \mu = 8, \quad H_A : \mu > 8$$

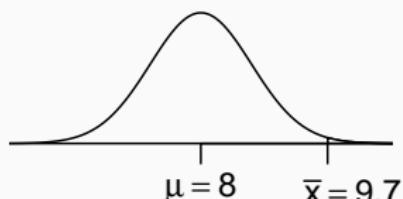
Also please **clearly specify what is μ .**

e.g., μ is the average number of colleges freshmen in this university have applied to.

Number of College Applications — Test Statistic

By CLT, under $H_0: \mu = 8$, the sampling distribution of the sample mean is

$$\bar{x} \sim N\left(\mu = 8, SE = \frac{7}{\sqrt{106}} = 0.68\right)$$

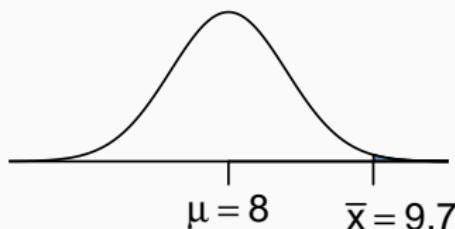


To gauge how unusual the observed sample mean $\bar{x} = 9.7$ is relative to its hypothesized sampling distribution above, the *test statistic* we used is the *z-statistic*, which is the *z-score* of the sample mean relative to the distribution above

$$z\text{-statistic} = \frac{\bar{x} - \mu_0}{SE} = \frac{\bar{x} - \mu_0}{\sigma / \sqrt{n}} = \frac{9.7 - 8}{7 / \sqrt{106}} \approx 2.5$$

Number of College Applications — P -Value

Recall *p-value* is the probability of observing data such that the evidence for the H_A is at least as strong as our current data set ($\bar{x} > 9.7$), if in fact $H_0: \mu = 8$ were true.



$$p\text{-value} = P(\bar{x} > 9.7 | \mu = 8) = P(Z > 2.50) = 0.0062$$

- Since *p-value* is *low* (lower than 5%) we *reject H_0* .
- The data provide convincing evidence that freshmen in this university have applied to more than 8 schools on average.
- The diff. between the null value of 8 schools and observed sample mean of 9.7 schools is *not due to chance* or sampling variability.

Example: Number of College Applications – Conditions

As CLT is used in the hypothesis test above, we need to check the same conditions as we construct confidence intervals for the population mean.

- Observations must be *independent*
 - Use your knowledge to judge if the data might be dependent
- The population distribution of the number of colleges students apply to should not be extremely skewed.
- In the z -statistic = $\frac{\bar{x} - \mu_0}{\sigma / \sqrt{n}}$, if the unknown population SD σ is replaced with the sample SD s , we need to further check that
 - sample size cannot be too small (at least 30)
 - no outliers & not too skewed ⇒ Check the histogram of data!

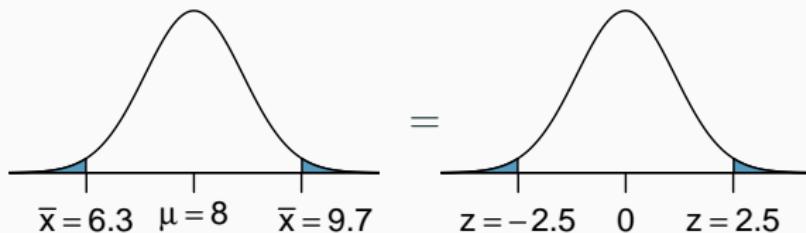
Two-Sided Hypothesis Test

If the dean wanted to know whether the data provide convincing evidence that the average number of colleges applied is *different* than the recommended 8 schools, the alternative hypothesis would be different.

$$H_0 : \mu = 8$$

$$H_A : \mu \neq 8$$

In this case, a sample mean \bar{x} far below 8 would also be evidence in favor of H_A . Hence the *p*-value would be the *two-tail* probability



$$\begin{aligned} p\text{-value} &= 0.0062 \times 2 \\ &= 0.0124 \end{aligned}$$

Recap: Hypothesis Testing for a Population Mean

1. Set the hypotheses

- $H_0 : \mu = \mu_0$
- $H_A : \mu < \text{or} > \text{or} \neq \mu_0$

2. Check assumptions and conditions

- Independence
- Normality: nearly normal population or $n \geq 30$, no extreme skew – or use the t distribution (Section 5.1)

3. Calculate a *test statistic* and a *p-value* (draw a picture!)

$$Z = \frac{\bar{x} - \mu_0}{SE}, \text{ where } SE = \frac{s}{\sqrt{n}}$$

4. (Optional) Make a decision

- If $p\text{-value} < \alpha$, reject H_0
- If $p\text{-value} > \alpha$, do not reject H_0

Relationship Between Confidence Intervals and Two-Sided Hypothesis Tests

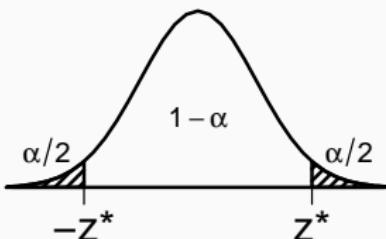
Confidence Intervals and Two-Sided Hypothesis Tests

For a two-sided test:

$$H_0 : \mu = \mu_0 \quad \text{versus} \quad H_A : \mu \neq \mu_0$$

the following are equivalent:

- $p\text{-value} > \alpha$ (and hence $H_0 : \mu = \mu_0$ is not rejected at level α)
- $|z\text{-statistic}| = |(\bar{x} - \mu_0)/SE| < z^*$, where z^* is a value such that



- μ_0 is in the $100(1 - \alpha)\%$ confidence interval for μ

$$\bar{x} - z^* SE < \mu_0 < \bar{x} + z^* SE$$

Example

Suppose in a study,

- 90% CI for μ is (4.81, 11.39);
- 95% CI for μ : (4.18, 12.02);
- 99% CI for μ : (2.95, 13.25).

Then

- $H_0 : \mu = 4$ is rejected at 5% level but not at 1% level
(2-sided p -value is between 1% and 5%)
because 4 is in the 99% CI but not in the 95% CI
- $H_0 : \mu = 4.5$ is rejected at 10% level but not at 5% level
because 4.5 is in the 95% CI but not in the 90% CI

Common Misunderstandings About Hypothesis Testing

In the lecture slide “General Framework of Hypothesis Testing”, we have introduced a number of common misunderstanding about hypothesis testing

- Rejecting H_0 doesn't mean we are 100% that H_0 is false. We might make Type 1 errors. Setting a significance level just guarantee we won't make Type 1 error too often
- P -value is not $P(H_0 \text{ is true} | \text{data})$ but it is $P(\text{data} | H_0 \text{ is true})$.

We are going to talk about more common misunderstanding about hypothesis testing here.

Failing to Reject H_0 Does Not Prove H_0 to Be True

Another mistake is to conclude from a high p -value that the H_0 is probably true

- We have said that if our p -value is low, then this is evidence that the H_0 is not true
- If our p -value is high, can we conclude that H_0 is true?
 - No, we could make a type 2 error when failing to reject H_0
 - Moreover, unlike type 1 error rate is controlled at a low level, type 2 error rate is usually quite high. It is quite often that H_0 is not true but the data fail to reject it.
- When we fail to reject H_0 , often it just means the data are not able to distinguish between H_0 and H_A (because the data are too noisy, etc)

Real Example

- As an example, the Women's Health Initiative found that low-fat diets reduce the risk of breast cancer with a *p*-value of 0.07
- The New York Times headline: “*Study finds low-fat diets won't stop cancer*”
- The lead editorial claimed that the trial represented “*strong evidence that the war against fats was mostly in vain*” and sounded “*the death knell for the belief that reducing the percentage of total fat in the diet is important for health*”
- Failing to prove the effect of low-fat diets doesn't prove that low-fat diets have no effect

<http://www.nytimes.com/2006/02/07/health/study-finds-lowfat-diet-wont-stop-cancer-or-heart-disease.html>

Don't Take the 0.05 Significance Level Too Seriously

- A p -value of 0.049 and a p -value of 0.051 give nearly the same strength of evidence against H_0
- For example, in the highly publicized 2009 study involving a vaccine that may protect against HIV infection, the two-sided p -value is 0.08, and the one-sided p -value of is 0.04
- Much debate and controversy ensued, partially because the two ways of analyzing the data produce p -values on either side of 0.05
- Much of this debate and controversy is fairly pointless; both p -values tell you essentially the same thing — that the vaccine holds promise, but that the results are not yet conclusive

Hypothesis Testing Cannot Tell Us...

Hypothesis testing cannot tell us

- whether the design of a study is flawed
- whether the data is appropriately collected

So we cannot conclude from a small P -value about whether one variable has a causal effect on another variable or whether the conclusion can be generalized to a bigger population.

Garbage In → Garbage Out

Statistical Significance Does Not Mean Practical Importance

Another mistake is reading too much into the term “statistically significant”

- Saying that results are statistically significant informs the reader that the findings are unlikely to be due to chance alone
- However, it says nothing about the practical importance of the finding.
- E.g., rejecting the $H_0: \mu_1 = \mu_2$ just tells us $\mu_1 \neq \mu_2$, but not how big and how important $\mu_1 - \mu_2$ is. It is possible that the difference is too small to be relevant even if it is significant.
- Remedy: *Attach a confidence interval* for the parameter so that people can decide whether the difference is big enough to be relevant.

Example

A 95% CI for the average number of colleges freshmen in this university have applied is

$$\bar{x} \pm 1.96 \frac{s}{\sqrt{n}} = 9.7 \pm 1.96 \frac{7}{\sqrt{106}} \approx 9.7 \pm 1.3 = (8.4, 11.0).$$

from which one can decide whether the difference from 8 is big enough to be relevant.

Recap: Common Misunderstandings about Hypothesis Testing

- Rejecting H_0 doesn't mean we are 100% sure that H_0 is false.
We might make Type 1 errors
- P -value is not the probability that the H_0 is true
- Failing to reject H_0 does not prove H_0 to be true
- Don't take the 0.05 significance level too seriously
- Hypothesis testing cannot tell us if data were collected properly or if the design of a study was flawed
- Statistical significance does not mean practical importance