

Statistical Methods

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Topics

- 1 CI for a Population Proportion p
- 2 Large Sample Test for a Population Proportion p

Objectives

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- Calculate and interpret a CI for a population proportion p when n is large.
- Carry out a hypothesis test for a population proportion p when the sample size is large.

CI for a Population Proportion p

- Consider a population of **successes** and **failures**, and let p denote the ***population proportion*** of successes.

CI for a Population Proportion p

- Consider a population of **successes** and **failures**, and let p denote the ***population proportion*** of successes.
- Suppose our goal is to **estimate** p using a random sample from the population.

- The **point estimator** of p is the ***sample proportion***, denoted \hat{P} .

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Sample Proportion:

$$\hat{P} = \frac{\text{\# of Successes in the Sample}}{\text{Sample Size}} = \frac{X}{n}$$

where

X = The number of successes in the sample.

Example

A random sample of $n = 10$ patrons at a restaurant were asked whether they smoke cigarettes (Yes or No). Here are the data.

Yes No Yes No Yes No No Yes No No

The **sample proportion** of smokers is

$$\hat{p} = \frac{4}{10} = 0.4.$$

We'd **estimate** that the true (unknown) proportion p that smokes in the population is 0.4, or 40%.

- The **sampling error** of the sample proportion is

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- It's preferable to estimate p using a **confidence interval** because its **width** will reflect how big the **sampling error** might be.
- To derive the CI, we'll need the **sampling distribution** of \hat{P} .

Sampling Distribution of the Sample Proportion \hat{P}

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$$E(\hat{P}) = \frac{1}{n}E(X) = p$$

and

$$V(\hat{P}) = \frac{1}{n^2}V(X) = \frac{p(1 - p)}{n}.$$

Mean and Variance of \hat{P} : For a **random sample** from a population of **successes** and **failures** whose proportion of successes is p , the **sampling distribution of \hat{P}** has mean

$$E(\hat{P}) = p,$$

and variance and standard deviation

$$V(\hat{P}) = \frac{p(1-p)}{n} \quad \text{and} \quad SD(\hat{P}) = \sqrt{\frac{p(1-p)}{n}}.$$

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- The **standard error** will be **small** if either:
 1. The population proportion p is close to 0 or 1, or
 2. The sample size n is **large**

- The following is a consequence of the **Normal Approximation to the Binomial**:

Proposition

Normality of \hat{P} : For a **random sample** from a population of **successes** and **failures** whose proportion of successes is p , if n is large,

$$\hat{P} \sim N \left(p, \sqrt{\frac{p(1-p)}{n}} \right) \quad (\text{approximately}).$$

Thus the **standardized** version of \hat{P} follows a **standard normal** distribution, i.e.

$$Z = \frac{\hat{P} - p}{\sqrt{\frac{p(1-p)}{n}}} \sim \text{N}(0, 1) \quad (\text{approximately}).$$

One-Sample z CI for p (7.2)

- To derive **95% CI for p** , note that

$$0.95 \approx P \left(-1.96 < \frac{\hat{P} - p}{\sqrt{\frac{p(1-p)}{n}}} < 1.96 \right)$$

$$\vdots$$

$$= P \left(\hat{P} - 1.96\sqrt{\frac{p(1-p)}{n}} < p < \hat{P} + 1.96\sqrt{\frac{p(1-p)}{n}} \right)$$

- Thus with probability 0.95, the (unknown) population proportion p will lie in the interval

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This leads to the following CI.

Large-Sample $100(1 - \alpha)\%$ Confidence Interval for p :

$$\hat{P} \pm z_{\alpha/2} \sqrt{\frac{\hat{P}(1 - \hat{P})}{n}}.$$

This is called the ***one-sample z confidence interval for p*** .

It's valid when the sample is from a population of successes and failures and n is large.

- In practice, n **is large** enough for the **one-sample z CI** for p to be valid as long as

$$n\hat{P} \geq 10 \quad \text{and} \quad n(1 - \hat{P}) \geq 10.$$

i.e. as long as there are at least **10 successes** and at least **10 failures** in the sample.

Example

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The **sample proportion** is

$$\hat{p} = \frac{284}{516} = 0.55,$$

and this is the **point estimate** of p , the true (unknown) proportion of all Americans that oppose legalizing the sale of organs.

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This gives a range of estimates of p , and we can be **95% confident** that p is in the interval somewhere.

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- Suppose we have a random sample of size n from a population of **successes** and **failures**.
- We'll see how to use the sample to decide if the **population proportion of successes** p is different from some **hypothesized value** p_0 .

The appropriate test is called the ***one-sample z test for p*** .

- The **null hypothesis** is that the population proportion p is equal to p_0 :

Null Hypothesis:

$$H_0 : p = p_0$$

- The **alternative hypothesis** will depend on what we're trying to "prove":

Alternative Hypothesis: The alternative hypothesis will be one of

1. $H_a : p > p_0$ (one-sided, upper-tailed)
2. $H_a : p < p_0$ (one-sided, lower-tailed)
3. $H_a : p \neq p_0$ (two-sided, two-tailed)

depending on what we're trying to verify using the data.

One-Sample Z Test Statistic for p :

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- Z measures how many standard errors \hat{P} is away from p_0 .
- \hat{P} is an estimator of the unknown population proportion p , so ...
 1. Z will be approximately **zero** (most likely) if $p = p_0$.
 2. It will be **positive** (most likely) if $p > p_0$.
 3. It will be **negative** (most likely) if $p < p_0$.

1. **Large positive** values of Z provide **evidence against H_0 in favor of $H_a : p > p_0$.**
2. **Large negative** values of Z provide **evidence against H_0 in favor of $H_a : p < p_0$.**
3. **Large positive and large negative** values of Z provide **evidence against H_0 in favor of $H_a : p \neq p_0$.**

- Recall that **if n is large**,

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Sampling Distribution of the Test Statistic Under H_0 :

If Z is the one-sample Z test statistic, then when

$$H_0 : p = p_0$$

is true,

$$Z \sim N(0, 1).$$

- The $N(0, 1)$ curve gives us:

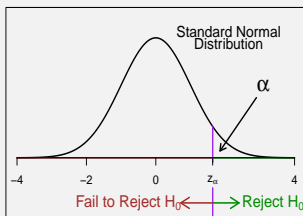
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 - The **rejection region** as the **extreme $100\alpha\%$ of z values** (in the direction(s) specified by H_a).
 - The **p -value** as the **tail area(s) beyond the observed z value** (in the direction(s) specified by H_a).

Rejection Region: The **rejection region** is the **set of z values** in the tail of the $N(0, 1)$ curve:

1. To the **right of z_α** if the alternative hypothesis is $H_a : p > p_0$:

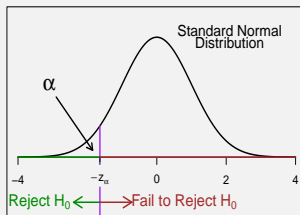
Rejection Region for Upper-Tailed Z Test



Values of Z

2. To the **left of** z_α if the alternative hypothesis is $H_a : p < p_0$:

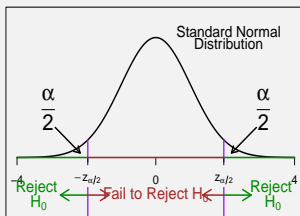
Rejection Region for Lower-Tailed Z Test



Values of Z

3. To the **left of** $-z_{\alpha/2}$ **and right of** $z_{\alpha/2}$ if the alternative hypothesis is $H_a : p \neq p_0$:

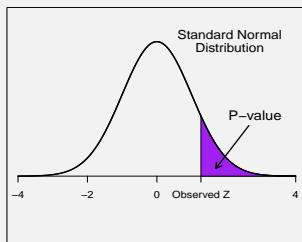
Rejection Region for Two-Tailed Z Test



Values of Z

1. **P-value** = Area to the **right** of the observed z if the alternative hypothesis is $H_a : p > p_0$.

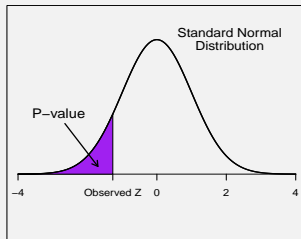
P-Value for Upper-Tailed Z Test



Values of Z

1. **P-value** = Area to the **left** of the observed z if the alternative hypothesis is $H_a : p < p_0$.

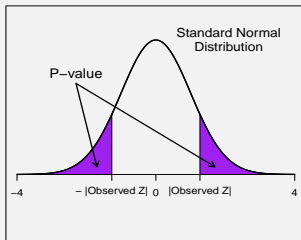
P-Value for Lower-Tailed Z Test



Values of Z

1. **P-value** = Area to the **left** of $-|z|$ **and right** of $|z|$ if the alternative hypothesis is $H_a : p \neq p_0$.

P-Value for Two-Tailed Z Test



Values of Z

- In practice, n **is large** enough for the **one-sample z test for p** to be valid as long as

$$np_0 \geq 10 \quad \text{and} \quad n(1 - p_0) \geq 10.$$

Example

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A new vaccine is meant to prevent meningitis in infants.

In a clinical trial, the vaccine was administered to **710** infants. Of these, **121** experienced appetite loss (a side effect).

Is there statistically significant evidence that the **true proportion** of infants experiencing appetite loss from the vaccine is **greater than 0.135**, the proportion that experience this side effect from competing medications?

We'll test the **hypotheses**

$$H_0 : p = 0.135$$

$$H_a : p > 0.135$$

where p is the true (unknown) population proportion that experiences appetite loss from the new vaccine.

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The **sample proportion** is

$$\hat{P} = \frac{121}{710} = \mathbf{0.170}.$$

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Thus the observed **test statistic** is

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Thus the observed **test statistic** is

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Thus the **sample proportion** that experiences appetite loss, $\hat{P} = 0.170$, is **2.73 standard errors above 0.135**.

Thus the observed **test statistic** is

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Thus the **sample proportion** that experiences appetite loss, $\hat{P} = 0.170$, is **2.73 standard errors above 0.135**.

The **p-value** is the **probability** that we'd get a z value this far above zero by chance **if the population proportion p was 0.135**.

From the **upper tail** area of the $N(0, 1)$ distribution, to the **right** of **2.73**,

$$\mathbf{p\text{-value} = 0.0032.}$$

Using a **level of significance** $\alpha = 0.05$, the **decision rule** is

Reject H_0 if p-value < 0.05 .

Fail to reject H_0 if p-value ≥ 0.05 .

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Fail to reject H_0 if p-value ≥ 0.05 .

Because $0.0032 < 0.05$, we **reject** H_0 .

There's **statistically significant evidence** that the population proportion of infants p that experience appetite loss from the vaccine is greater than 0.135.

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Fail to reject H_0 if p-value ≥ 0.05 .

Because $0.0032 < 0.05$, we **reject** H_0 .

There's **statistically significant evidence** that the population proportion of infants p that experience appetite loss from the vaccine is greater than 0.135.

The observed result cannot easily be explained by chance variation (sampling error).