



Lecture 7: Composite Hypotheses

STA261 – Probability & Statistics II

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Outline

Composite Hypotheses

- Power Functions and Power Calculations

- Uniformly Most Powerful Tests

- The p-value

The Connection to Confidence Intervals

- Two-Tailed Tests and Confidence Intervals



Composite Hypotheses

- When the alternative hypothesis is of the form $\mathcal{H}_1 : \theta \in \Theta_1$, where Θ_1 consists of more than a single possible value, we call \mathcal{H}_1 a *composite* alternative.
- Consider for example $X_1, \dots, X_n \stackrel{\text{i.i.d.}}{\sim} \mathcal{N}(\mu, \sigma^2)$ (with σ^2 known for now). One may choose to test –

- $\mathcal{H}_0 : \mu = \mu_0$ vs. $\mathcal{H}_1 : \mu > \mu_0$ (a *right-tailed test*),
- $\mathcal{H}_0 : \mu = \mu_0$ vs. $\mathcal{H}_1 : \mu < \mu_0$ (a *left-tailed test*), or
- $\mathcal{H}_0 : \mu = \mu_0$ vs. $\mathcal{H}_1 : \mu \neq \mu_0$ (a *two-tailed test*),

- Recall that when we tested $\mathcal{H}_0 : \theta = \theta_0$ vs. $\mathcal{H}_1 : \theta = \theta_1$, the power of the test was defined to be **rejecting when we should reject**

$$\pi = 1 - \beta = \mathbb{P} \left(\begin{array}{c} \text{reject} \\ \mathcal{H}_0 \end{array} \middle| \theta = \theta_1 \right).$$

- Now there is no θ_1 anymore, but rather a set of alternatives $\theta \in \Theta_1$.

cannot talk about power anymore, dunno which alternative we are talking about..



Power Functions

Definition the graph of a function

The *power function* of a statistical test is

$$\pi(\theta^*) = \mathbb{P} \left(\begin{array}{c} \text{reject} \\ \mathcal{H}_0 \end{array} \mid \theta = \theta^* \right).$$

- Consider for example the LRT test we developed for the simple hypotheses $\mathcal{H}_0 : \mu = \mu_0$ vs. $\mathcal{H}_1 : \mu = \mu_1$ ($\mu_1 > \mu_0$) for Normal data with known variance, based on the rejection region

$$\mathcal{C} = \left\{ \underline{x} \in \mathbb{R}^n : \bar{x} \geq \mu_0 + \frac{\sigma}{\sqrt{n}} z_{1-\alpha} \right\}.$$

- Note that \mathcal{C} does not depend on μ_1 : the test is the same for any $\mu_1 > \mu_0$.
- We will use it then for the right-tailed test – $\mathcal{H}_0 : \mu = \mu_0$ vs. $\mathcal{H}_1 : \mu > \mu_0$.



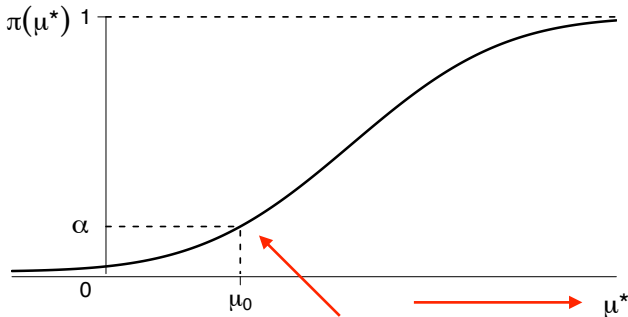
Power Functions (cont.)

power: reject null when alternative is true

$$\pi(\mu^*) = \mathbb{P}(\underline{X} \in \mathcal{C} | \mu = \mu^*) = \mathbb{P}\left(\bar{X} \geq \mu_0 + \frac{\sigma}{\sqrt{n}} z_{1-\alpha} \mid \mu = \mu^*\right)$$

$$= 1 - \mathbb{P}\left(\frac{\bar{X} - \mu^*}{\sigma/\sqrt{n}} \leq \frac{\mu_0 - \mu^*}{\sigma/\sqrt{n}} + z_{1-\alpha} \mid \mu = \mu^*\right) = 1 - \Phi\left(\frac{\sqrt{n}(\mu_0 - \mu^*)}{\sigma} + z_{1-\alpha}\right).$$

standardize w.r.t. μ^* , not μ_0



when $\mu = \mu_0$; then π is probability of rejecting when null is true, equivalent to type 1 error



Power calculations

$$\pi(\mu^*) = 1 - \Phi\left(\frac{-\sqrt{n}(\mu^* - \mu_0)}{\sigma} + z_{1-\alpha}\right)$$

- A quick look at $\pi(\mu^*)$ reveals that –
 1. The larger n is – the more powerful the test is (explain)
 2. The further apart the null hypothesis and the alternative are – the greater the power is (explain) **expect to reject null easier**
 3. The larger σ is – the less powerful the test is (explain)
 4. The smaller α is – the smaller the power is (has been discussed already)
- With that said, for any given α and we can make β arbitrarily small by increasing n – so long as we know the kind of differences $\mu^* - \mu$ that we wish to uncover.

since H_1 and alpha are set before the test. and sigma is based on the data.
only thing that can change is the size of samples.



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note that we only need to ensure that power at μ_1 is greater than $1 - \beta$, because for $\mu > \mu_1$, power is larger further apart null and alternative hypothesis are

Power calculations (cont.)

- Suppose that we wish to test $\mathcal{H}_0 : \mu = \mu_0$ vs. $\mathcal{H}_1 : \mu > \mu_0$ with σ^2 assumed to be known, as before, at significance level α .
- In addition, differences smaller than $\delta = \mu_1 - \mu_0$ are deemed non-consequential, but beyond that difference we wish to keep the probability of a type II error at less than β .

note power function is higher for higher μ_1

$$1 - \beta \leq \pi(\mu_1) = 1 - \Phi\left(-\frac{\sqrt{n}(\mu_1 - \mu_0)}{\sigma} + z_{1-\alpha}\right)$$

calculate n required to have higher output to power function than specified beta

$$\Rightarrow \Phi\left(-\frac{\sqrt{n}(\mu_1 - \mu_0)}{\sigma} + z_{1-\alpha}\right) \leq \beta$$

$$\Rightarrow -\frac{\sqrt{n}(\mu_1 - \mu_0)}{\sigma} + z_{1-\alpha} \leq z_\beta = -z_{1-\beta}$$

$$\Rightarrow \frac{\sqrt{n}(\mu_1 - \mu_0)}{\sigma} \geq z_{1-\alpha} + z_{1-\beta} \Rightarrow n \geq \left\{ \frac{\sigma(z_{1-\alpha} + z_{1-\beta})}{\mu_1 - \mu_0} \right\}^2$$

calculate minimum sample size to achieve the specified alpha, beta beyond the μ_1



Power calculations (cont.)

- For example, Suppose we know that the serum cholesterol levels for all 20-24 year-old males in Canada is normally distributed with $\mu = 180$ mg/100ml and the $\sigma = 46$ mg/100ml.
- We would expect that the mean cholesterol level of a special diet group in this population to be higher than 180 mg/100ml. (Assuming the same $\sigma = 46$ mg/100ml.)
 μ_1
 beyond what μ_1 do we care, i.e. ensure power is at least 90% beyond $180 + 20 = 200$
- If the mean cholesterol level increases by at least 20 mg/100ml, we would like to be able to show that the new diet is effective (i.e. reject $\mathcal{H}_0 : \mu = 180$ in favor of $\mathcal{H}_1 : \mu > 180$) at a significance level of 5% and with a power of 90%.
- To do that we will need a sample of size

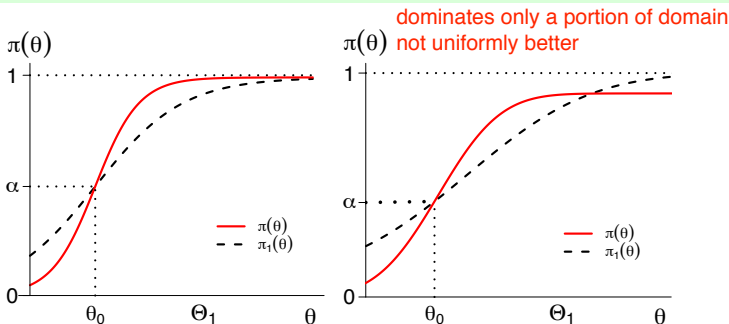
$$n \geq \left\{ \frac{\sigma(z_{1-\alpha} + z_{1-\beta})}{\mu_1 - \mu_0} \right\}^2 = \left\{ \frac{46(\overbrace{1.645}^{z_{0.95}} + \overbrace{1.282}^{z_{0.9}})}{200 - 180} \right\}^2 = 45.3 \implies n \geq 46.$$



Uniformly Most Powerful Tests

Definition

Consider testing $\mathcal{H}_0 : \theta = \theta_0$ vs. $\mathcal{H}_1 : \theta \in \Theta_1$ (a composite alternative). We say that a test at level α with power function $\pi(\theta)$ is a *uniformly most powerful* (UMP) test, if for any other test at level α with power function $\pi_1(\theta)$, we have $\pi_1(\theta) \leq \pi(\theta)$ for all $\theta \in \Theta_1$.



UMP (left panel) and not UMP (right panel).



An example of a UMP test

- For $X_1, \dots, X_n \stackrel{\text{i.i.d.}}{\sim} \mathcal{N}(\mu, \sigma^2)$ (with a known σ^2), we have proposed the rejection region

$$\mathcal{C} = \left\{ \underline{x} \in \mathbb{R}^n : \bar{x} \geq \mu_0 + \frac{\sigma}{\sqrt{n}} z_{1-\alpha} \right\}$$

to test $\mathcal{H}_0 : \mu = \mu_0$ vs. the right-tailed alternative $\mathcal{H}_1 : \mu > \mu_0$.

- Here $\Theta_1 = (\mu_0, \infty]$ (the set of alternatives)
- This has been shown to be the most powerful (MP) test at level α for testing $\mathcal{H}_0 : \mu = \mu_0$ vs. $\mathcal{H}_1 : \mu = \mu_1$ ($\mu_1 > \mu_0$) **for simple hypotheses**
- It does not depend on the value of μ_1 , hence it is the most powerful test for any $\mu \in \Theta_1$ **test is MP for all $\mu_1 > \mu_0 \implies$ test is UMP**
- But this is the definition of a UMP test!



The p-value

- We have just shown that

$$C_{\text{right}} = \left\{ \bar{X} \geq \mu_0 + \frac{\sigma}{\sqrt{n}} z_{1-\alpha} \right\}$$

forms a rejection for a UMP test at level α for testing $\mathcal{H}_0 : \mu = \mu_0$ vs. the right-tailed alternative $\mathcal{H}_1 : \mu > \mu_0$.

- It is equally simple to show that

$$C_{\text{left}} = \left\{ \bar{X} \leq \mu_0 - \frac{\sigma}{\sqrt{n}} z_{1-\alpha} \right\}$$

forms a rejection region for a UMP test at level α for testing $\mathcal{H}_0 : \mu = \mu_0$ vs. the left-tailed alternative $\mathcal{H}_1 : \mu < \mu_0$.

- If $\mathcal{H}_0 : \mu = \mu_0$ is true, $\bar{X} \sim \mathcal{N}\left(\mu_0, \frac{\sigma^2}{n}\right)$

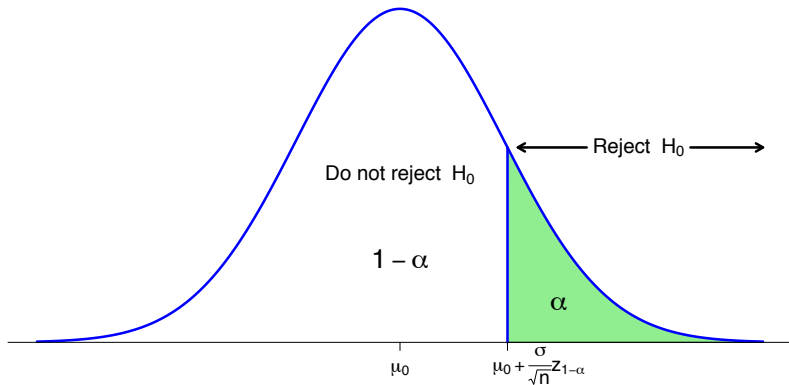


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The p-value (cont.)

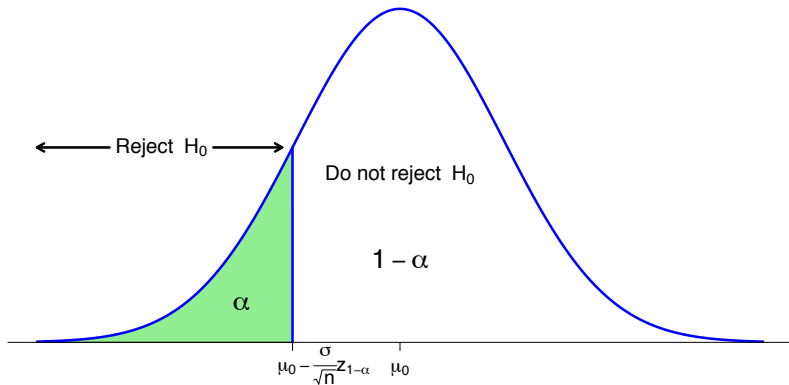
The distribution of \bar{X} under \mathcal{H}_0 and the rejection region for the right-tailed test:





The p-value (cont.)

The distribution of \bar{X} under \mathcal{H}_0 and the rejection region for the left-tailed test:





Example

- Suppose that we have a sample of size 25 from a $\mathcal{N}(\mu, 5^2)$ distribution, and we wish to test $\mathcal{H}_0 : \mu = 175$ vs. $\mathcal{H}_1 : \mu > 175$ at 5% level. The observed sample mean was 177.
- The rejection region is

$$\mathcal{C} = \left\{ \bar{X} \geq \mu_0 + \frac{\sigma}{\sqrt{n}} z_{1-\alpha} \right\} = \left\{ \bar{X} \geq 175 + \frac{5}{5} \underbrace{z_{0.95}}_{1.645} \right\} = \{ \bar{X} \geq 176.6 \}.$$

- In this example we are told that $\bar{x} = 177$ was observed – inside the rejection region.
- at 5% significance level then, we reject \mathcal{H}_0 .

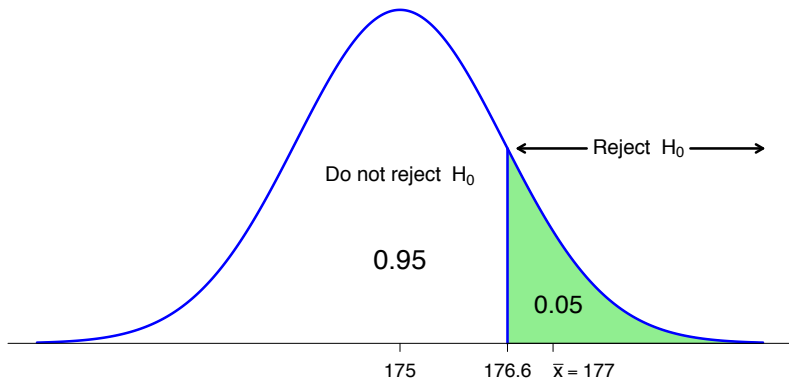


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Example (cont.)

The distribution of \bar{X} under \mathcal{H}_0 and the rejection region:



if alpha is set lower, i.e decreases, then its harder to reject,



Example (cont.)

- What if changed our mind and would now like to test the hypotheses at 1% level?
- The new rejection region is

$$\mathcal{C} = \left\{ \bar{X} \geq \mu_0 + \frac{\sigma}{\sqrt{n}} z_{1-\alpha} \right\} = \left\{ \bar{X} \geq 175 + \frac{5}{5} \underbrace{z_{0.99}}_{2.326} \right\} = \{ \bar{X} \geq 177.3 \}.$$

- Now the observed sample mean $\bar{x} = 177$ is outside of the rejection region.
- At a 1% significance level then, we do not reject \mathcal{H}_0 .

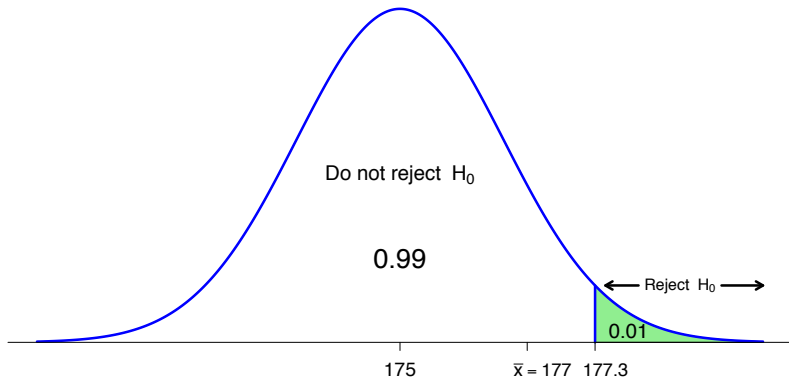


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Example (cont.)

The distribution of \bar{X} under \mathcal{H}_0 and the rejection region:



decrease alpha, line move from left to right, area of rejection gets smaller



The p-value

- We rejected \mathcal{H}_0 at 5% level – when \bar{X} was outside of the rejection region
- We did not reject \mathcal{H}_0 at 1% level – when \bar{X} was inside the rejection region
- What if \bar{X} was right at the boundary of the rejection region?
 - i.e. the rejection region was $\mathcal{C} = \{\bar{X} \geq \bar{x} = 177\}$
 gives smallest alpha that we will still reject H_0
- That would give us the minimum significance level for which \mathcal{H}_0 would be rejected. We call that number the *p-value*.
- Let us calculate it:

$$\begin{aligned} \text{p-value} &= \mathbb{P} \left(\begin{array}{c} \text{Type I} \\ \text{error} \end{array} \right) = \mathbb{P}(\underline{X} \in \mathcal{C} | \mu = 175) = \mathbb{P}(\bar{X} \geq 177 | \mu = 175) \\ &= \mathbb{P} \left(\frac{\bar{X} - 175}{5/\sqrt{25}} \geq \frac{177 - 175}{5/\sqrt{25}} \middle| \mu = 175 \right) = 1 - \Phi(2) = 0.023. \end{aligned}$$



Understanding p-values

- We calculated that the p-value for the last example was 0.023
smaller alpha => harder to reject
- It is the minimum α for which \mathcal{H}_0 would be rejected – hence we rejected \mathcal{H}_0 at 5% but did not reject it at 1% level.
- We no longer need to calculate a new rejection region for every α like we did before, because the rule is

$$\text{Reject } \mathcal{H}_0 \text{ at level } \alpha \iff \text{p-value} \leq \alpha$$

- At 3% level we would reject \mathcal{H}_0
- At 2% level we would not reject \mathcal{H}_0 because $0.02 < 0.023$
- And so on...
no longer need to calculate different rejection region for different alpha
- Recall that we calculated

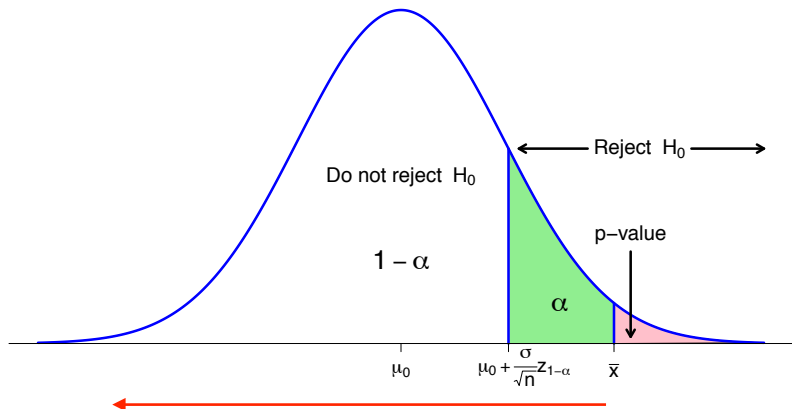
$$\text{p-value} = \mathbb{P}(\overline{X} \geq \overline{x} \mid \mu = 175)$$



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Understanding p-values (cont.)

The distribution of \bar{X} under \mathcal{H}_0 :



p-value = $\mathbb{P}(\bar{X} \geq \bar{x} | \mu = \mu_0)$ p-value = minimum alpha to reject



Understanding p-values (cont.)

$$\text{p-value} = \mathbb{P}(\bar{X} \geq \bar{x} \mid \mu = \mu_0)$$

- The p-value is therefore the probability of observing an effect “at least as extreme” as the one in your data, assuming the truth of \mathcal{H}_0 .
- Remember that we reject $\mathcal{H}_0 \iff \text{p-value} \leq \alpha$ (for a predetermined α)
- The philosophy of hypothesis testing is then, in a nutshell –

i.e. P(Type I error) is less than pre-determined alpha

If the probability of observing an affect at least as extreme as the one in your data, assuming the truth of \mathcal{H}_0 , is very low, then \mathcal{H}_0 is most likely false. so we reject null



Notes on composite null hypotheses

- Clearly, if we reject $\mathcal{H}_0 : \mu = 175$ in favor of $\mathcal{H}_1 : \mu > 175$, we would also reject $\mathcal{H}_0 : \mu = 170$, $\mathcal{H}_0 : \mu = 150$ or in general $\mathcal{H}_0 : \mu = \mu_0$ for any $\mu_0 \leq 175$.
- This allows us to write the hypotheses in a more “complete” way:
writing null and alternative hypotheses complementing each other

$$\begin{cases} \mathcal{H}_0 : \underline{\mu \leq \mu_0} \\ \mathcal{H}_1 : \mu > \mu_0 \end{cases} \quad (\text{right-tailed}), \quad \text{or} \quad \begin{cases} \mathcal{H}_0 : \underline{\mu \geq \mu_0} \\ \mathcal{H}_1 : \mu < \mu_0 \end{cases} \quad (\text{left-tailed})$$

- One could also write down the hypotheses as appearing above, and define the significance level of the test to be

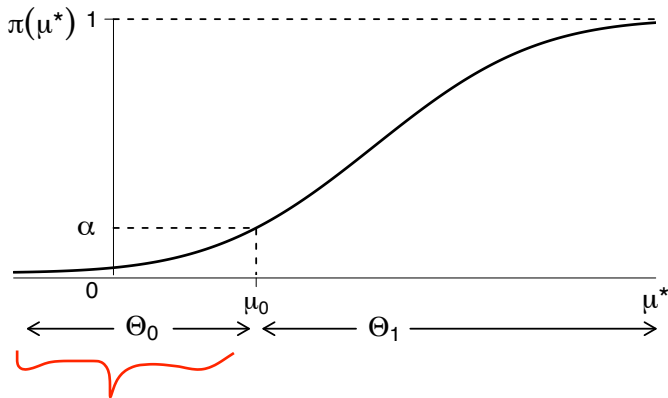
$$\alpha = \sup_{\theta \in \Theta_0} \pi(\theta),$$

where $\pi(\theta)$ is the power function of the test, as discussed at the beginning of this lecture.



Composite null-hypotheses (cont.)

The power function of the right-tailed test and the significance level:





Two-tailed test for the Normal mean (known σ^2)

- Consider now the problem of testing $\mathcal{H}_0 : \mu = \mu_0$ vs. the two-tailed alternative $\mathcal{H}_1 : \mu \neq \mu_0$.
- When testing \mathcal{H}_0 vs. the right-tailed alternative, very positive values of $\bar{X} - \mu_0$ support the alternative, hence the rejection region is of the form

$$\mathcal{C}_{\text{right}} = \left\{ \bar{X} - \mu_0 \geq \frac{\sigma}{\sqrt{n}} z_{1-\alpha} \right\}.$$

- Likewise, when testing \mathcal{H}_0 vs. the left-tailed alternative, very negative values of $\bar{X} - \mu_0$ are in support of the alternative, hence the rejection region is of the form

$$\mathcal{C}_{\text{left}} = \left\{ \bar{X} - \mu_0 \leq -\frac{\sigma}{\sqrt{n}} z_{1-\alpha} \right\}.$$

- However, when testing \mathcal{H}_0 vs. the two-tailed alternative, a strong evidence against \mathcal{H}_0 would be large values of $|\bar{X} - \mu_0|$.

want absolute value to be larger -> evidence against null



Two-tailed test for μ with known σ^2 (cont.)

- Trivially, we look for a rejection region of the form $\mathcal{C} = \{ |\bar{X} - \mu_0| \geq c \}$.
- For a test at significance level α , c must satisfy

$$\begin{aligned}
 \alpha &= \mathbb{P}(\underline{X} \in \mathcal{C} | \mu = \mu_0) = \mathbb{P}(|\bar{X} - \mu_0| \geq c | \mu = \mu_0) \\
 &= \mathbb{P}(\bar{X} - \mu_0 \geq c | \mu = \mu_0) + \mathbb{P}(\bar{X} - \mu_0 \leq -c | \mu = \mu_0) \\
 &= \mathbb{P}\left(\frac{\bar{X} - \mu_0}{\sigma/\sqrt{n}} \geq \frac{c}{\sigma/\sqrt{n}} \middle| \mu = \mu_0\right) + \mathbb{P}\left(\frac{\bar{X} - \mu_0}{\sigma/\sqrt{n}} \leq -\frac{c}{\sigma/\sqrt{n}} \middle| \mu = \mu_0\right) \\
 &= 1 - \Phi\left(\frac{c\sqrt{n}}{\sigma}\right) + \Phi\left(-\frac{c\sqrt{n}}{\sigma}\right) = 2 \left\{ 1 - \Phi\left(\frac{c\sqrt{n}}{\sigma}\right) \right\}
 \end{aligned}$$

$$\Rightarrow \Phi\left(\frac{c\sqrt{n}}{\sigma}\right) = 1 - \alpha/2 \Rightarrow \frac{c\sqrt{n}}{\sigma} = z_{1-\alpha/2} \Rightarrow c = \frac{\sigma}{\sqrt{n}} z_{1-\alpha/2}.$$

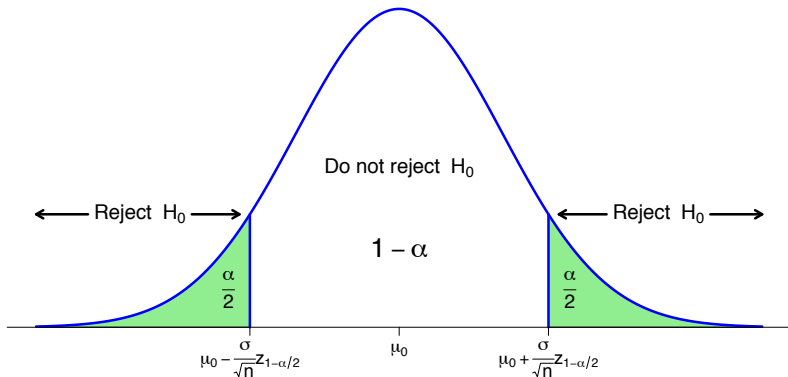
note its $1 - \alpha/2$ for two tailed test; $1 - \alpha$ for one tailed test



Two-tailed test for μ with known σ^2 (cont.)

$$\mathcal{C} = \left\{ |\bar{X} - \mu_0| \geq \frac{\sigma}{\sqrt{n}} z_{1-\alpha/2} \right\} = \left\{ \bar{X} \leq \mu_0 - \frac{\sigma}{\sqrt{n}} z_{1-\alpha/2} \right\} \cup \left\{ \bar{X} \geq \mu_0 + \frac{\sigma}{\sqrt{n}} z_{1-\alpha/2} \right\}$$

- The distribution of \bar{X} under \mathcal{H}_0 and the rejection region:





Two-tailed test for μ with known σ^2 (cont.)

- Let us now test $\mathcal{H}_0 : \mu = 175$ vs. $\mathcal{H}_1 : \mu \neq 175$ at 4% level, using the same data as before: $n = 25$, $\bar{x} = 177$, $\sigma = 5$.
- The same data yielded p-value = 0.023 for the right-tailed test, hence we would reject \mathcal{H}_0 at any level $\alpha \geq 0.023$ – in particular we would reject \mathcal{H}_0 at level 4%.
- The rejection region is

$$\begin{aligned}
 \mathcal{C} &= \left\{ \bar{X} \leq \mu_0 - \frac{\sigma}{\sqrt{n}} z_{1-\alpha/2} \right\} \cup \left\{ \bar{X} \geq \mu_0 + \frac{\sigma}{\sqrt{n}} z_{1-\alpha/2} \right\} \\
 &= \left\{ \bar{X} \leq 175 - \frac{5}{\sqrt{25}} \underbrace{z_{0.98}}_{2.05} \right\} \cup \left\{ \bar{X} \geq 175 + \frac{5}{\sqrt{25}} \underbrace{z_{0.98}}_{2.05} \right\} \\
 &= \left\{ \bar{X} \leq 172.95 \right\} \cup \left\{ \bar{X} \geq 177.05 \right\} \implies \text{we do not reject } \mathcal{H}_0!
 \end{aligned}$$

because $x = 177$



p-value for the two-tailed test

one tail -> hints to where alternative hypothesis lies... so needs more evidence

two tail -> first time, no prior knowledge of where H_1 lies

- Well, that's confusing – the same data that lead us to conclude at 4% level that $\mu > 175$, does not provide enough evidence to conclude that $\mu \neq 175$...?
- The supposed paradox is resolved by re-examining our initial standpoint: testing \mathcal{H}_0 vs. a one-tailed alternative suggests that we have an idea as to where to expect the parameter to be with respect to \mathcal{H}_0 , ergo, we need less convincing.

easier to reject null for one tail alternative than two tail alternative

- Let us reaffirm the result by calculating the p-value.
- Here, observing an effect that is “at least as extreme” as that in your data (assuming the truth of \mathcal{H}_0) means observing $|\bar{X} - \mu_0| \geq |\bar{x} - \mu_0|$, hence

$$\text{p-value} = \mathbb{P} \left(\underline{|\bar{X} - \mu_0| \geq |\bar{x} - \mu_0|} \mid \mu = \mu_0 \right).$$



p-value for the two-tailed test (cont.)

$$\begin{aligned}
 \text{p-value} &= \mathbb{P} \left(|\bar{X} - \mu_0| \geq |\bar{x} - \mu_0| \mid \mu = \mu_0 \right) \\
 &= \mathbb{P} \left(\frac{|\bar{X} - \mu_0|}{\sigma/\sqrt{n}} \geq \frac{|\bar{x} - \mu_0|}{\sigma/\sqrt{n}} \mid \mu = \mu_0 \right) \\
 &= 2\mathbb{P} \left(\frac{\bar{X} - \mu_0}{\sigma/\sqrt{n}} \geq \frac{|\bar{x} - \mu_0|}{\sigma/\sqrt{n}} \mid \mu = \mu_0 \right) = 2 \left\{ 1 - \Phi \left(\frac{|\bar{x} - \mu_0|}{\sigma/\sqrt{n}} \right) \right\} \\
 &= 2 \left\{ 1 - \Phi \left(\frac{|177 - 175|}{5/\sqrt{25}} \right) \right\} = 2 \{ 1 - \Phi(2) \} = 0.046.
 \end{aligned}$$

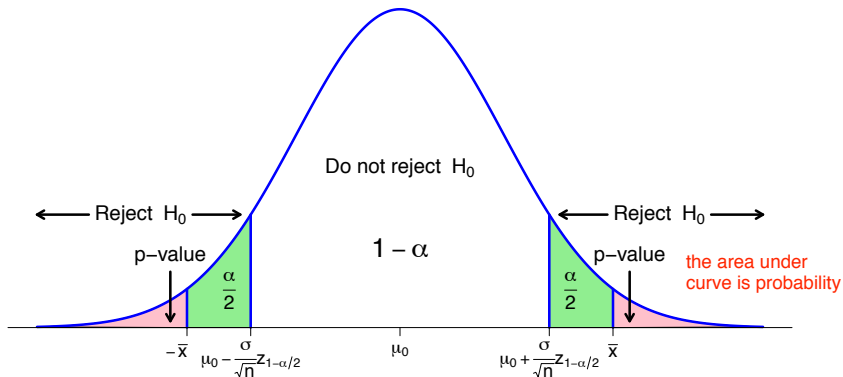
- Note that this is exactly double the p-value of the right-tailed test.
- We did not reject \mathcal{H}_0 at 4% level, but we would have rejected it at 5% level.

reject as long as p value is smaller than alpha



p-value for the two-tailed test (cont.)

The distribution of \bar{X} under \mathcal{H}_0 and the rejection region:



rejecting two tail tests is harder because p-value is doubled in this case



Power function for the two-tailed test

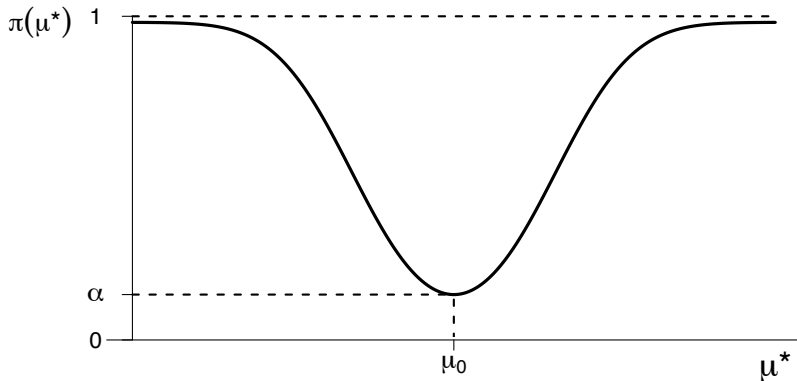
We may now proceed to calculate the power function for the two-tailed test –

$$\begin{aligned}
 \pi(\mu^*) &= \mathbb{P}\left(\underline{X} \in \mathcal{C} \mid \mu = \mu^*\right) \\
 &= \mathbb{P}\left(\bar{X} \geq \mu_0 + \frac{\sigma}{\sqrt{n}} z_{1-\alpha/2} \mid \mu = \mu^*\right) + \mathbb{P}\left(\bar{X} \leq \mu_0 - \frac{\sigma}{\sqrt{n}} z_{1-\alpha/2} \mid \mu = \mu^*\right) \\
 &= \mathbb{P}\left(\frac{\bar{X} - \mu^*}{\sigma/\sqrt{n}} \geq \frac{\mu_0 - \mu^* + \frac{\sigma}{\sqrt{n}} z_{1-\alpha/2}}{\sigma/\sqrt{n}} \mid \mu = \mu^*\right) \text{ standardize with respect to } \mu^* \\
 &\quad + \mathbb{P}\left(\frac{\bar{X} - \mu^*}{\sigma/\sqrt{n}} \leq \frac{\mu_0 - \mu^* - \frac{\sigma}{\sqrt{n}} z_{1-\alpha/2}}{\sigma/\sqrt{n}} \mid \mu = \mu^*\right) \\
 &= 1 - \Phi\left(-\frac{\sqrt{n}(\mu^* - \mu_0)}{\sigma} + z_{1-\alpha/2}\right) + \Phi\left(-\frac{\sqrt{n}(\mu^* - \mu_0)}{\sigma} - z_{1-\alpha/2}\right).
 \end{aligned}$$



Power function for the two-tailed test (cont.)

$$\pi(\mu^*) = 1 - \Phi\left(-\frac{\sqrt{n}(\mu^* - \mu_0)}{\sigma} + z_{1-\alpha/2}\right) + \Phi\left(-\frac{\sqrt{n}(\mu^* - \mu_0)}{\sigma} - z_{1-\alpha/2}\right)$$





Comments on the two-tailed tests

two tail test not more powerful than one tail test on their respective domain

- Is the two-tailed test for the Normal mean (known variance) a UMP test? No!
 - The right-tailed test is more powerful for any alternative $\mu^* > \mu_0$
 - The left-tailed test is more powerful for any $\mu^* < \mu_0$
- For large n and small α we have $\Phi\left(-\frac{\sqrt{n}(\mu^* - \mu_0)}{\sigma} - z_{1-\alpha/2}\right) \approx 0$, hence

$$\pi(\mu^*) \approx 1 - \Phi\left(-\frac{\sqrt{n}(\mu^* - \mu_0)}{\sigma} + z_{1-\alpha/2}\right),$$

allowing us to calculate the required n for any prespecified α , β and $\delta = \mu_1 - \mu_0$ (see practice problems).



Two-tailed tests and confidence intervals equivalent

- Recall the two-tailed test for the normal mean with known variance at level α , based on the rejection region

$$\mathcal{C} = \left\{ \bar{X} \leq \mu_0 - \frac{\sigma}{\sqrt{n}} z_{1-\alpha/2} \right\} \cup \left\{ \bar{X} \geq \mu_0 + \frac{\sigma}{\sqrt{n}} z_{1-\alpha/2} \right\}.$$

- Note that

We do not reject

$$\mathcal{H}_0 : \mu = \mu_0$$

$$\iff \bar{X} \notin \mathcal{C}$$

1 - \alpha Confidence interval for normal mean

$$\iff \mu_0 - \frac{\sigma}{\sqrt{n}} z_{1-\alpha/2} \leq \bar{X} \leq \mu_0 + \frac{\sigma}{\sqrt{n}} z_{1-\alpha/2}$$

$$\iff \bar{X} - \frac{\sigma}{\sqrt{n}} z_{1-\alpha/2} \leq \mu_0 \leq \bar{X} + \frac{\sigma}{\sqrt{n}} z_{1-\alpha/2}$$

$$\iff \mu_0 \text{ is contained in the } (1 - \alpha)100\% \text{ confidence interval for } \mu.$$



Two-tailed tests and confidence intervals (cont.)

We do not reject $\mathcal{H}_0 : \mu = \mu_0$ at level α $\iff \mu_0$ is contained in the $(1 - \alpha)100\%$ confidence interval for μ .

- This is hardly surprising if we think of a confidence interval as a set of “plausible” values for μ : we reject $\mathcal{H}_0 : \mu = \mu_0 \iff \mu_0$ is not a plausible value for μ .
isnt theta_0 a single number not a set even for composite hypothesis
- In general, the set of all θ_0 for which $\mathcal{H}_0 : \theta = \theta_0$ would not get rejected in a two-tailed set at level α , forms a $(1 - \alpha)100\%$ confidence set for θ .
- Conversely, the set of all \underline{X} 's for which the $(1 - \alpha)100\%$ confidence interval for θ based on \underline{X} would not contain θ_0 , forms a rejection region of size α for $\mathcal{H}_0 : \theta = \theta_0$ vs. $\mathcal{H}_1 : \theta \neq \theta_0$.
- In other words: every confidence set has a corresponding two-tailed test and vice versa.