

# Lecture 7: Hypothesis testing part I

## Statistical Methods for Data Science

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# Today

## 1 Terminology

- Experiment and parameter of interest
- Null hypothesis and alternative hypothesis
- Test statistic
- Null distribution  $f(s | H_0)$
- Significance level  $\alpha$ , power and  $p$ -value

## 2 Example

## 3 Summary

## Learning outcome

- Be able to explain the following terminology
  - Null hypothesis  $H_0$  and alternative hypothesis  $H_A$
  - Test statistic  $s$
  - Null distribution  $f(s | H_0)$
  - Significance level  $\alpha$  and power
  - $p$ -value
- Be able to design and interpret the one-sample z-test

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- What if company A tested drug D on 100 ducks and the average weight loss after one month is 2.3 kg?
- Now company B tested drug E on 30 ducks and the average weight loss after one month is 2.5 kg, while drug D results in 2.3 kg weight loss with the same setup, would you buy drug E instead of drug D?

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How would you make your decision?

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  - Hypothesis + data  $\rightarrow$  decision on rejecting/not rejecting the hypothesis

# Hypothesis testing: a list to go through

- A “boring” statement
- Experiment
- Data  $x$ , random variable  $X$
- Parameter of interest  $\theta$
- Parameter estimate  $\hat{\theta}$
- Null hypothesis  $H_0$
- Alternative hypothesis  $H_A$
- Test statistic  $s$
- Null distribution  $f(s | H_0)$
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## Experiment and parameter of interest

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Then we can use  $\bar{x}$  to approximate  $\mu_D$  and check if it is greater than diet control (2.1 kg)

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Then we use  $\bar{x}$  and  $\bar{y}$  to approximate  $\mu_D$  and  $\mu_E$  to see if they are the same

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$$H_A : \mu_D > 2.1$$

- Example 2: drug E and drug D do not work the same on average (5 sec)

# Hypotheses $H_0$ and $H_A$

- **Statistical hypothesis**: a proposed distribution - a statement about the **parameter of interest**
- **Null hypothesis  $H_0$** : the “boring” statement translated into a mathematical expression
  - Example 1: drug D is not more effective than regular diet on average

$$H_0 : \mu_D = 2.1$$

- Example 2: drug E and drug D work the same on average (5 sec)

$$H_0 : \mu_D = \mu_E$$

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Answer: no



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- Question 2: can  $H_0$  and  $H_A$  be ANYTHING I want? Like a magic mirror!?  
Answer: no
- Follow up question: what are the choices for  $H_0$  and  $H_A$ ?

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- We have seen one-sample test and two-sample test in the Q-Q plot lecture
- In practice, you can narrow down your choice of hypotheses by making a Q-Q plot



# Choices for $H_A$

## Given

$$H_0 : \theta = \beta$$

where  $\beta$  can be either a constant (one-sample test) or a parameter from another data distribution (two-sample test)

- **Alternative hypothesis  $H_A$ :**  $H_A$  can be **one-tailed** or **two-tailed**
  - **One-tailed:**

$$H_A : \theta > \beta$$

or

$$H_A : \theta < \beta$$

- **Two-tailed:**

$$H_A : \theta \neq \beta \iff \theta < \beta \text{ or } \theta > \beta$$

## Summary: choices for $H_0$ and $H_A$

Putting everything together,

	One-sample test	Two-sample test
Two-tailed	$H_0 : \theta = c, H_A : \theta \neq c$	$H_0 : \theta_1 = \theta_2, H_A : \theta_1 \neq \theta_2$
One-tailed	$H_0 : \theta = c, H_A : \theta > c$	$H_0 : \theta_1 = \theta_2, H_A : \theta_1 > \theta_2$
	$H_0 : \theta = c, H_A : \theta < c$	$H_0 : \theta_1 = \theta_2, H_A : \theta_1 < \theta_2$

where  $\theta, \theta_1, \theta_2$  are the parameters of interest and  $c$  is a constant

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Note: in this course, we only deal with null hypothesis where we are able to express the PDF/PMF  $f(s | H_0)$ , i.e.  $H_0$  with an equal sign in them

## Test statistic (cont.)

### Example 1. one-sample test

- **Data:**  $x_1, \dots, x_N$
- **Random variable:**  $X_1, \dots, X_N$  i.i.d. **Gaussian with known  $\sigma$**
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**Standardize  $\bar{x}$  (15 sec)**

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**Standardize  $\bar{x}$  (15 sec)**

$$z = \frac{\bar{x} - 2.1}{\sigma / \sqrt{N}}$$

## Test statistic (cont.)

### Example 2. two-sample test

- **Data:**  $x_1, \dots, x_{N_D}$  and  $y_1, \dots, y_{N_E}$
- **Random variable:**  $X_1, \dots, X_{N_D}$  i.i.d. **Gaussian with known  $\sigma_D$** ;  
 $Y_1, \dots, Y_{N_E}$  i.i.d. **Gaussian with known  $\sigma_E$** ;  $X_i$  and  $Y_j$  independent
- **Parameter of interest:**  $\mu_D, \mu_E$
- **Parameter estimate:**  $\bar{x}, \bar{y}$
- **Null hypothesis:**  $H_0 : \mu_D = \mu_E \iff H_0 : \mu_D - \mu_E = 0$
- **Test statistic:** standardized  $\bar{x} - \bar{y}$  assuming the null hypothesis

$$z = \frac{\bar{x} - \bar{y}}{\sqrt{\sigma_D^2/N_D + \sigma_E^2/N_E}} \quad (\text{explained later})$$

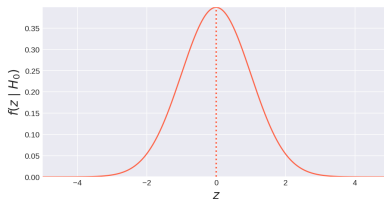
## Null distribution $f(s | H_0)$

# Null distribution

- **Null distribution  $f(s | H_0)$ :** the distribution of the test statistic given the null hypothesis
- **Example:**
  - **Data:**  $x_1, \dots, x_N$
  - **Random variable:**  $X_1, \dots, X_N$  i.i.d. Gaussian with known  $\sigma$
  - **Parameter of interest:**  $\mu$
  - **Parameter estimate:**  $\bar{x}$
  - **Null hypothesis:**  $H_0 : \mu = \mu_0$
  - **Test statistic:**

$$z = \frac{\bar{x} - \mu_0}{\sigma/\sqrt{N}}$$

- **Null distribution:** standard Gaussian distribution





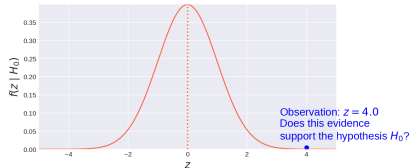
## Significance level $\alpha$ , power and $p$ -value

# Significance level

Given a null hypothesis  $H_0 : \mu = 2.1$  and the null distribution  $f(s | H_0)$ , we decide if we reject the hypothesis or not by observing data

- Run some experiments and collect data  $x_1, \dots, x_N$
- Estimate the parameter of interest  $\hat{\theta}$ , e.g.  $\hat{\mu} = \bar{x} = \frac{1}{N} \sum_{i=1}^N x_i$
- Standardize  $\hat{\theta}$  assuming  $H_0$  to compute the test statistic, e.g.

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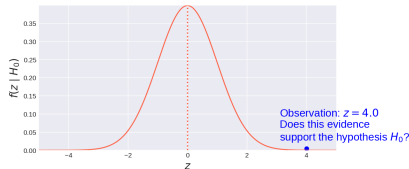
- Does this evidence support the hypothesis  $H_0$ ?

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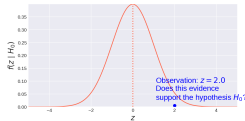
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- Does this evidence support the hypothesis  $H_0$ ? Probably not since it's so far away from the center?

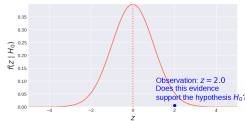
## Significance level (cont.)

- What about this observation?



## Significance level (cont.)

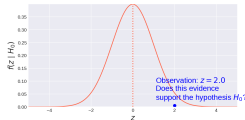
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- To be able to answer the question, you need to decide where you draw the line

## Significance level (cont.)

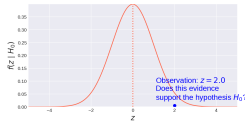
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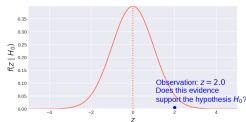
- To be able to answer the question, you need to decide where you draw the line - define a **rejection region** by choosing a significance level
- Significance level  $\alpha$** : red area under the curve



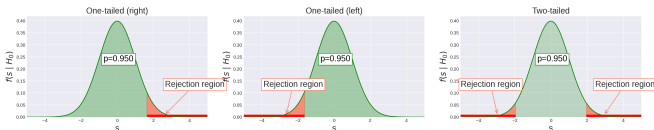
In these three images,  $\alpha = 0.05$

## Significance level (cont.)

- What about this observation?



- To be able to answer the question, you need to decide where you draw the line - define a **rejection region** by choosing a significance level
- Significance level  $\alpha$** : red area under the curve



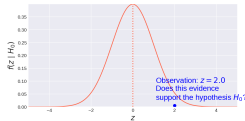
In these three images,  $\alpha = 0.05$

More conservative  $\Rightarrow$  less probable to reject  $H_0$ , which indicates a smaller rejection region



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Two-tailed  $H_A$  is more conservative

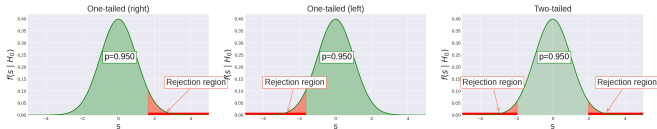
## Significance level (cont.)

What is needed for choosing a meaningful  $\alpha$ ?

- Null distribution
- $H_A$  one-tailed or two-tailed

# Interpretation of $\alpha$

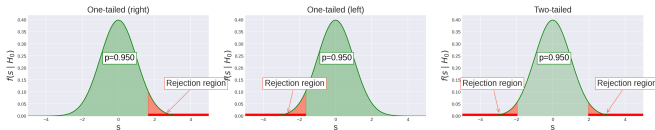
- $\alpha = P(\text{reject } H_0 \mid H_0 \text{ is true})$  - the probability of making such a mistake



- The rejection region indicates that  $H_0$  is **unlikely**, but the probability is not zero
- It is possible that  $H_0$  is true, but our observation happens to fall in the rejection region
- If  $H_0$  is true and our observation falls in the rejection region, we will **mistakenly** reject  $H_0$
- The probability of making this type of mistakes is  $\alpha$
- Similar to the confidence interval,  $1 - \alpha$  is called the **confidence level**

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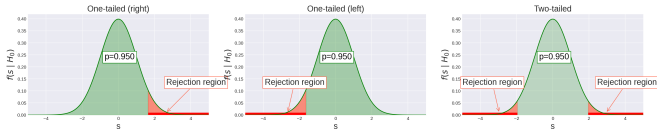
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- Similar to the confidence interval,  $1 - \alpha$  is called the **confidence level** - "with 95% confidence, rejecting  $H_0$  is the right thing to do"
- Define the significance level **before you run the experiments** so that you can't cheat!

# Significance level and power

- Contingency table:

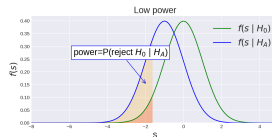
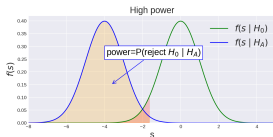
	$y = H_A$	$y = H_0$
$\hat{y} = \text{reject } H_0$	TP	FP (Type I error)
$\hat{y} = \text{do not reject } H_0$	FN (Type II error)	TN

- Significance level  $\alpha$ : incorrectly rejecting  $H_0$

$$\alpha = P(\text{type I error})$$

- Power: correctly rejecting  $H_0$

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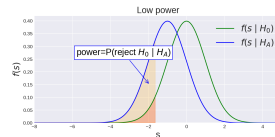
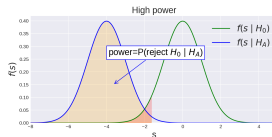
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- What is needed for computing the power?  $f(s | H_0)$ ,  $f(s | H_A)$

# $p$ -value

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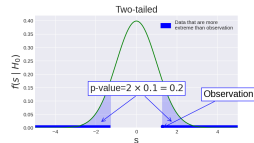
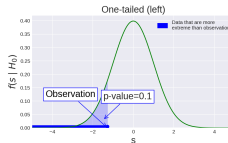
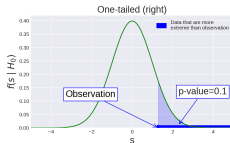
- One-tailed:

- Right tail:  $p = P(S \geq s | H_0)$
- Left tail:  $p = P(S \leq s | H_0)$

- Two-tailed:  $p = 2 \min(P(S \leq s | H_0), P(S \geq s | H_0))$

Note: for example, if  $f(s | H_0)$  is symmetric around zero and  $s < 0$ ,

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- What is needed for computing the  $p$ -value? (10 sec)



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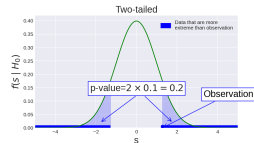
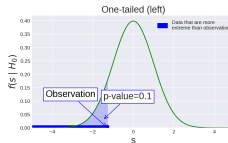
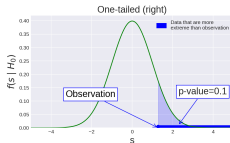
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- Null distribution
- Alternative hypothesis  $H_A$  to know one-tailed or two-tailed
- Observation - test statistic computed from data

## Summary: steps for hypothesis testing

- Step 1 Make a “boring” statement
- Step 2 Design an **experiment**
- Step 3 Describe the **data** generated from the experiment and the corresponding random variables
- Step 4 Describe the parameter of interest and their estimates
- Step 5 Translate the “boring” statement into a statistical hypothesis and call it the **null hypothesis  $H_0$**
- Step 6 Find the expression for the **test statistic  $s$**
- Step 7 Find the expression for the **null distribution**
- Step 8 Define **an alternative hypothesis  $H_A$** : one-tailed or two-tailed
- Step 9 Choose a **significance level  $\alpha$**  (the tail), which defines the **rejection region**
- Step 10 Collect **data**
- Step 11 Compute the test statistic from data
- Step 12 Compute the  $p$ -value
- Step 13 If  $p\text{-value} < \alpha$ , i.e. the test statistic falls in the rejection region of the null distribution, then we reject the hypothesis  $H_0$ ; otherwise, we fail to reject  $H_0$ .

# Today

- 1 Terminology
- 2 Example
- 3 Summary

# Example

Recall example: if you control the diet of your ducks, they lose 2.1 kg after one month on average. Company A has developed a drug D to help ducks lose weight. They claim that on average the drug works better than diet control. Here is the set up for the experiment.

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Step 10 Collect 30 ducks in 20 secs and feed them drugs

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Step 9 Choose a **significance level**  $\alpha$  (the tail), which defines the rejection region (5 secs): e.g.  $\alpha = 0.05$

Step 10 Collect 30 ducks in 20 secs and feed them drugs - great job! Weights measured after one month  $x_1, \dots, x_{30}$

## Example (cont.)

Step 8 Define **an alternative hypothesis**  $H_A$  (10 secs):

$$H_A : \mu_D \neq 2.1 \text{ or } H_A : \mu_D > 2.1$$

One-tailed or two-tailed

- **Two-tailed (5 secs):**  $H_A : \mu_D \neq 2.1$
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$$\text{Say } \frac{1}{30} \sum_{i=1}^{30} x_i = 2.2$$

Step 11 Compute the test statistic from data (5 secs):

## Example (cont.)

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Step 11 Compute the test statistic from data (5 secs):

$$z_0 = \frac{2.2 - 2.1}{0.6/\sqrt{30}} = 0.91$$



## Example (cont.)

Step 12 Compute the  $p$ -value (20 secs):

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- For  $H_A : \mu_D > 2.1$  (one-tailed):  $p = P(Z \geq z_0 \mid H_0) = 0.1807 > \alpha$

## Example (cont.)

Step 12 Compute the  $p$ -value (20 secs):

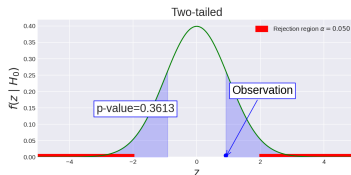
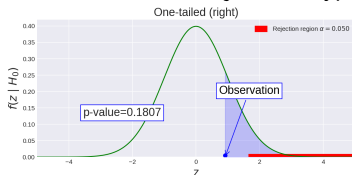
- **For  $H_A : \mu_D > 2.1$  (one-tailed):**  $p = P(Z \geq z_0 \mid H_0) = 0.1807 > \alpha$
- **For  $H_A : \mu_D \neq 2.1$  (two-tailed):**  $p = 2P(Z \geq z_0 \mid H_0) = 0.3613 > \alpha$

## Example (cont.)

Step 12 Compute the  $p$ -value (20 secs):

- For  $H_A : \mu_D > 2.1$  (**one-tailed**):  $p = P(Z \geq z_0 | H_0) = 0.1807 > \alpha$
- For  $H_A : \mu_D \neq 2.1$  (**two-tailed**):  $p = 2P(Z \geq z_0 | H_0) = 0.3613 > \alpha$

Step 13 If  $p\text{-value} < \alpha$ , i.e. the test statistic falls in the rejection region of the null distribution, then we reject the hypothesis  $H_0$



Do not reject  $H_0$  for both one-tailed and two-tailed  $H_A$

## Example (cont.)

What if  $\bar{x} = 2.3$ ?

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$$z_0 = \frac{2.3 - 2.1}{0.6/\sqrt{30}} = 1.826$$

## Example (cont.)

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## Example (cont.)

What if  $\bar{x} = 2.3$ ?

Step 11 Compute the test statistic from data (5 secs):

$$z_0 = \frac{2.3 - 2.1}{0.6/\sqrt{30}} = 1.826$$

Step 12 Compute the  $p$ -value (20 secs):

- **For  $H_A : \mu_D > 2.1$  (one-tailed):**  $p = P(Z \geq z_0 \mid H_0) = 0.0339 < \alpha$
- **For  $H_A : \mu_D \neq 2.1$  (two-tailed):**  $p = 2P(Z \geq z_0 \mid H_0) = 0.0679 > \alpha$

## Example (cont.)

What if  $\bar{x} = 2.3$ ?

Step 11 Compute the test statistic from data (5 secs):

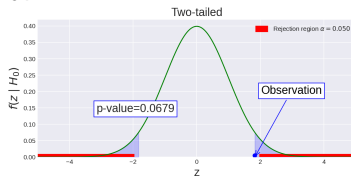
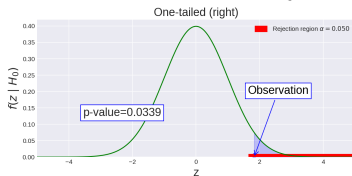
$$z_0 = \frac{2.3 - 2.1}{0.6/\sqrt{30}} = 1.826$$

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## Example (cont.)

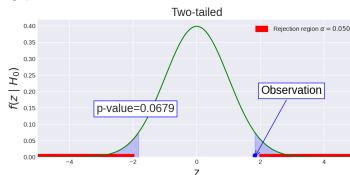
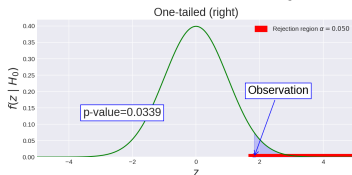
**Step 13** If  $p\text{-value} < \alpha$ , i.e. the test statistic falls in the rejection region of the null distribution, then we reject the hypothesis  $H_0$



Reject  $H_0$  for one-tailed  $H_A$ ; do not reject  $H_0$  for two-tailed  $H_A$   
for the same confidence level  $1 - \alpha = 5\%$

## Example (cont.)

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**Reject  $H_0$  for one-tailed  $H_A$ ; do not reject  $H_0$  for two-tailed  $H_A$  for the same confidence level  $1 - \alpha = 5\%$**

Note: the two-tailed test is more conservative - if the data passes a two-tailed test, it is more conclusive than one-tailed test for the same confidence level

## Example (cont.)

What if  $\bar{x} = 2.3$  with  $N = 100$ ?

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## Example (cont.)

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Step 12 Compute the  $p$ -value (20 secs):



## Example (cont.)

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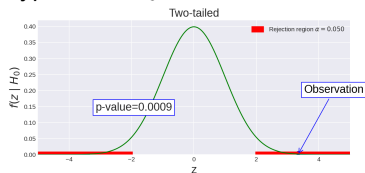
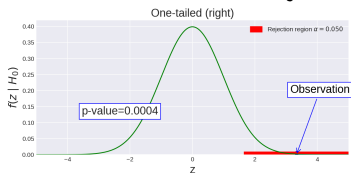
$$z_0 = \frac{2.3 - 2.1}{0.6/\sqrt{100}} = 3.33$$

Step 12 Compute the  $p$ -value (20 secs):

- **For  $H_A : \mu_D > 2.1$  (one-tailed):**  $p = P(Z \geq z_0 \mid H_0) = 0.0004 < \alpha$
- **For  $H_A : \mu_D \neq 2.1$  (two-tailed):**  $p = 2P(Z \geq z_0 \mid H_0) = 0.0009 < \alpha$

## Example (cont.)

**Step 13** If  $p\text{-value} < \alpha$ , i.e. the test statistic falls in the rejection region of the null distribution, then we reject the hypothesis  $H_0$



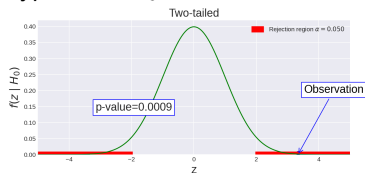
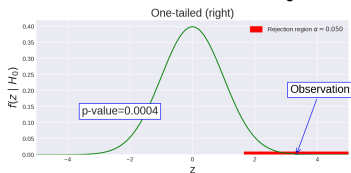
**Reject  $H_0$  for both one-tailed and two-tailed  $H_A$**

Note:

- With more data, it becomes more certain that we should reject  $H_0$  in favor of  $H_A$  given the observation  $\bar{x} = 2.3$

## Example (cont.)

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Note:

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This test is called **one-sample z-test**

# Today

- 1 Terminology
- 2 Example
- 3 Summary

# Summary

So far:

- Data types and data containers
- Descriptive data analysis: descriptive statistics, visualization
- Probability distributions, events, random variables, PMF, PDF, parameters
- CDF, Q-Q plot, how to compare two distributions (data vs theoretical, data vs data)
- Modeling
- Parameter estimation: maximum likelihood estimation (MLE) and maximum a posteriori estimation (MAP)
- Classification, multinomial naive Bayes classifier, Gaussian naive Bayes classifier
- Central limit theorem, interval estimation
- Hypothesis test

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Next:

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Before next lecture:

- Steps for hypothesis testing

