# Chapter 9 - Lecture 4 P-values

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#### Introduction

In the 5-step procedure, we are given a significant level  $\alpha$  and calculate the rejection region. If the value of test statistic falls in the rejection region, we reject  $H_0$ ; otherwise, fail to reject  $H_0$ .

Today, we introduce an alternative way of reaching a conclusion in hypothesis testing. In this approach, we will calculate a certain probability called *p-value*.

#### Definition

**P-value** is the probability, of obtaining a test statistic at least as contradictory to the null hypothesis as the one we have calculated from the available sample, assuming the null hypothesis is true.

- P-values are important for the following reason:
  - If p-value  $\leq \alpha$  we reject  $H_0$
  - If p-value  $> \alpha$  we do not reject  $H_0$
- So in the 5-step procedure of testing hypotheses, we can replace "Step 4: Determine the rejection/critical region *C*" with the following step:
  - Step 4: Calculate the p-value

#### **Proposition**

The P-value is the smallest significance level  $\alpha$  at which the null hypothesis can be rejected. Because of this, the P-value is alternatively referred to as the **observed significance level** for the data.

## Calculating p- values

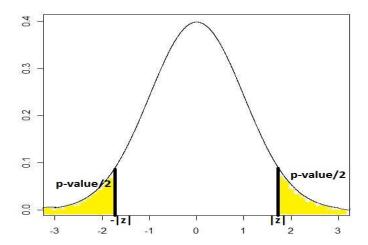
How do we calculate p-values?

- p-values depend on the tests conducted. Hence, the calculation will be done most of the time by definition.
- 1 z test / t test;
- upper-tailed / lower-tailed / two-tailed;

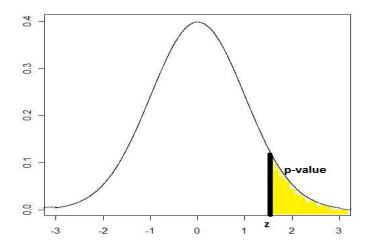
- **1** Two-sided test  $H_1: \mu \neq \mu_0: P(|Z| > z), P(|T| > t)$ ;
- 2 Upper-tailed test  $H_1: \mu > \mu_0: P(Z > z)$ , P(T > t);
- **3** Lower-tailed test  $H_1: \mu < \mu_0$ : P(Z < z), P(T < t).

Next 3 slides will use the z test as an example to illustrate the p-value. (z test has z curve and t test will have  $t_{n-1}$  curve.)

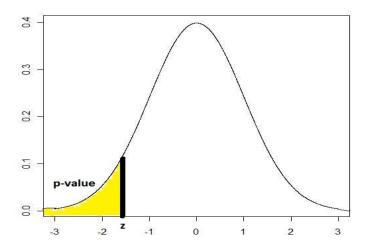
## Two-sided test $H_1: \mu \neq \mu_0$



#### Upper-tailed test $H_1: \mu > \mu_0$



#### Lower-tailed test $H_1$ : $\mu < \mu_0$



### Example 9.17

Example 9.17: The target thickness for silicon wafers used in a type of integrated circuit is 245  $\mu m$ . A sample of 50 wafers is obtained and the thickness of each one is determined, resulting in a sample mean thickness of 246.18  $\mu m$  and a sample standard deviation of 3.60  $\mu m$ . Does this data suggest that true average wafer thickness is something other than the target value?

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- step2: Test statistic value:  $z = \frac{\bar{x}-245}{s/\sqrt{n}}$
- step3: Null distribution (Distribution of Z under  $H_0$ ) is N(0,1).
- step4: Determine the P-value. Since it's a two-sided test,  $P-vlaue=P(|Z|>z)=P(Z>z)+P(Z<-z)=2[1-\Phi(z)]$

- step1:  $H_0: \mu = 245$  vs  $H_1: \mu \neq 245$ , where  $\mu =$  true average wafer thickness
- step2: Test statistic value:  $z = \frac{\bar{x}-245}{s/\sqrt{n}}$
- step3: Null distribution (Distribution of Z under  $H_0$ ) is N(0,1).
- step4: Determine the P-value. Since it's a two-sided test,  $P-vlaue = P(|Z|>z) = P(Z>z) + P(Z<-z) = 2[1-\Phi(z)]$
- step5: Based on the data, plug in  $\bar{x} = 246.18$ , s = 3.60  $\rightarrow z = 2.32 \rightarrow P value = 2[1 \Phi(2.32)] = 0.0204$ .
  - If  $\alpha = 0.01$ , fail to reject  $H_0$ .
  - If  $\alpha = 0.05$ , reject  $H_0$ .