

# Introduction to Hypothesis Testing

1. One-sample Student t-test
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3. Two-sample Student t-test

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# One-sample Student t-test

## One-sample Student t-test (shortly, one sample t-test)

- The one-sample Student t-test is one of the most frequently applied tests in statistics. It is used to **test a certain hypothesis about the unknown population mean  $\mu$**

## Background

- The t-test was devised by William Sealy Gosset, working for **Guinness brewery** in Dublin, to cheaply monitor the quality of stout
- Published in Biometrika in 1908 under the pen name **“Student”** as Guinness regarded the fact that they used statistics a trade secret



# P-value

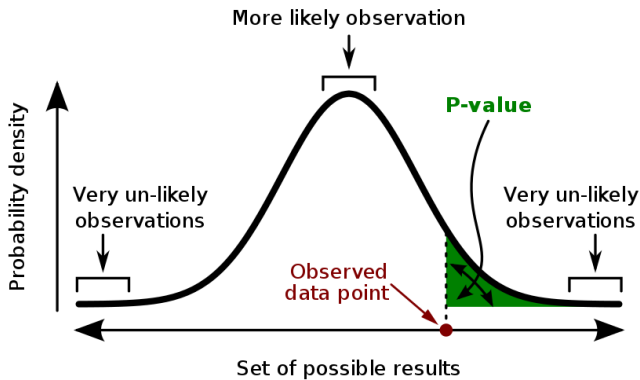
## Definition

The probability that the observed result, or a result more extreme, is true, given  $H_0$  is true.

Then:

- The p-value is a measure of how likely our observed result is, under the  $H_0$  assumption.
- If the p-value is small, then what we have observed is rare under  $H_0$ , which means we have *evidence against it*.
- p-values are used to *evaluate* the hypothesis test result, in terms of the *strength of the evidence* that the test provides.

# P-value



## The one sample t-test: an example

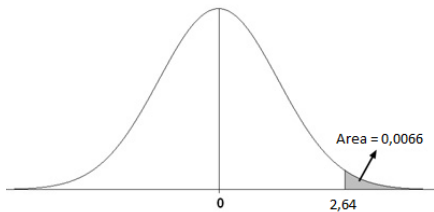
- 30 measures of lactate dehydrogenase (LD)
- **Question:**  $\mu = 105$ ?
- **Test:**  $H_0: \mu = 105$ ,  $H_a: \mu > 105$
- We know that if  $H_0$  is true, then

$$T_0 = \frac{\bar{X} - \mu_0}{s/\sqrt{n}} \rightarrow t(n-1),$$

$T_0$  is called **test statistic**

- For our example:  $T_0 = \frac{108.8-105}{7.88/\sqrt{30}} = 2.64$
- When would you reject the null hypothesis? Two options:
  - ▶ when  $T_0$  is large, meaning when  $T_0 > t_{n-1,\alpha}$ , **OR**
  - ▶ when  $p < \alpha$
- In our example:  $T_0 = 2.64 > t_{29,0.05} = 1.699 \rightarrow$  Rejection
- When the test is **two-sided** ( $H_a: \mu \neq 105$ ), use  $t_{29,0.025} = 2.04$

## How to get the P-value



- If two-sided test:  $p = 2P_{H_0}(t > |T_0|)$
- R or other statistical softwares produce the p-value automatically

# Paired data

## Paired measurements

- In medical settings we often deal with **paired measurements**, which is two outcomes measured on
  - ▶ the *same* individual under different exposure (or treatment) circumstances
  - ▶ *two* individuals matched by certain key characteristics
- The pairing in the data is taking into account by considering the *differences* between each pair of outcome observations. In that way the **data are turned into a single sample** of differences

## Paired measurements

2 measures of each individual (for example before/after treatment)

Individual	Measure 1	Measure 2
1	$X_{11}$	$X_{12}$
2	$X_{21}$	$X_{22}$
3	$X_{31}$	$X_{32}$
...	...	...



## Example: 7.3 in Kirkwood & Sterne

We consider the results of a **clinical trial** to test the effectiveness of a sleeping drug. The sleep of ten patients was observed during one night with the **drug** and one night with **placebo**. For each patient a *pair* of sleep times, was recorded and the *difference* between these calculated

Patient	Hours of sleep		Difference
	Drug	Placebo	
1	6.1	5.2	0.9
2	6.0	7.9	-1.9
3	8.2	3.9	4.3
4	7.6	4.7	2.9
5	6.5	5.3	1.2
6	5.4	7.4	-2.0
7	6.9	4.2	2.7
8	6.7	6.1	0.6
9	7.4	3.8	3.6
10	5.8	7.3	-1.5
Mean	$\bar{X}_1 = 6.66$	$\bar{X}_0 = 5.58$	$\bar{X} = 1.08$

The observed **mean difference** in sleep time was  $\bar{X} = 1.08$  hours, and the empirical **standard deviation** of the differences was  $s = 2.31$ . The estimated **standard error** of the differences is  $s/\sqrt{n} = 2.31/\sqrt{10} = 0.73$  hours

A 95% **confidence interval** for the mean difference in sleep time in the population is given by

$$(1.08 - 2.26 \times 0.73, 1.08 + 2.26 \times 0.73) = (-0.57, 2.73),$$

where 2.26 is the two-sided **5% point** of the  $t$  distribution with  $(n - 1) = 9$  degrees of freedom

The **mean difference** in sleep time was  $\bar{X} = 1.08$  hours, and the estimated **standard error** was  $s/\sqrt{n} = 0.73$  hours. The **test statistic** is given by

$$t = 1.08/0.73 = 1.48,$$

which is  $t$  distributed with  $(n - 1) = 9$  degrees of freedom when the null hypothesis of no effect is true. The corresponding  **$P$ -value**, which is the probability of getting a  $t$  value with a size as large as this or larger in a  $t$  distribution with 9 degrees of freedom, is

$$p = 0.17$$

*So, there is no evidence against the null hypothesis that the drug does not affect sleep time*

# Two sample t-test

So far...

- Tests and confidence intervals for
  - ▶ Single sample
  - ▶ Paired samples
- We know how to test (the procedure)

Now:

- Test for the difference in the mean of two independent samples

## The data: two different settings. Now focus on situation 2

- 1 Paired data: 2 measures of each individual (for example before/after treatment)

Individual	Measure 1	Measure 2
1	$X_{11}$	$X_{12}$
2	$X_{21}$	$X_{22}$
3	$X_{31}$	$X_{32}$
...	...	...

- 2 2 groups: 1 measure of each individual, each which corresponds to a group (for example sick/healthy people)

Group 1		Group 2	
Ind.	Measure	Ind.	Measure
1	$X_{11}$	1	$X_{12}$
2	$X_{21}$	2	$X_{22}$
...		...	
		14	$X_{14,2}$
15	$X_{15,1}$		

## The two sample t-test

- The **null hypothesis** is given by

$$H_0 : \mu_1 = \mu_0 \quad \text{or} \quad H_0 : \mu_1 - \mu_0 = 0,$$

i.e. there is *no difference between the population means* in the two groups

- The **test statistic** is given by

$$t = \frac{\bar{X}_1 - \bar{X}_0}{s\sqrt{(1/n_1 + 1/n_0)}},$$

which follow a  $t$  distribution with  $(n_1 + n_0 - 2)$  degrees of freedom. Here,  $s$  is the common estimate of the population standard deviation:

$$s = \sqrt{\left[ \frac{(n_1 - 1)s_1^2 + (n_0 - 1)s_0^2}{n_1 + n_0 - 2} \right]}$$

### Example: 7.2 in Kirkwood & Sterne

We return to the data of **birth weights**. The **test statistic** is given by

$$t = \frac{3.1743 - 3.6267}{0.4121\sqrt{(1/14 + 1/15)}} = -\frac{0.4524}{0.1531} = -2.95$$

The corresponding **P-value** calculated from the  $t$  distribution with  $(14 + 15 - 2) = 27$  degrees of freedom is given as:

$$p = 0.006$$

Therefore, *the data suggest that smoking during pregnancy reduces the birthweight of the baby*

## Test statistic

- In the one sample t-test we had

$$T = \frac{\bar{X}_n - \mu_0}{s} \sqrt{n}$$

- Now  $T = \frac{\bar{X}_1 - \bar{X}_2}{s_p} \sqrt{n_p}$

$S_p$  is the pooled standard deviation  $\sqrt{\frac{(n_1-1)s_1^2 + (n_2-1)s_2^2}{n_1+n_2-2}}$  and

$$n_p = \frac{n_1 \cdot n_2}{n_1 + n_2}.$$



## Test statistic (cont.)

- $T \sim t_{n_1+n_2-2}$  under the null hypothesis  $H_0$
- Rejection and conclusion:

$H_0$	$\mu_1 = \mu_2$
Rejection, if	$ T_0 $ large
Rejection, if	$P = 2 P_{H_0}(t >  T_0 ) < \alpha$
Conclusion	$\mu_1 \neq \mu_2$

## Small samples, unequal standard deviations

- When the population standard deviations,  $\sigma_1$  and  $\sigma_0$ , of the two groups are different, and the sample size,  $n$ , is not large, the main possibilities are:
  - ▶ Use a **transformation** on the data which makes the standard deviations similar so that methods based on the  $t$  distribution can be used
  - ▶ Use **non-parametric** methods based on ranks
  - ▶ Use either the **Fisher-Behrens** or the **Welch** tests, which allow for unequal deviations
  - ▶ Estimate the difference between the means using the original measurements, but use **bootstrap** methods to derive confidence intervals

## How to check for normal distribution

- Box-plot
- Histograms
- Q-Q plot

## What if the data does not look normal?

- Try to find a meaningful transformation
- Use a test which does not assume normally distributed data

→ Lecture on transformations and non-parametric methods in day 1 of week 2