# **Statistics**

# Lecture 9

Hypothesis testing: Parametric tests



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- z-tests for the means
  - type I and type II error
  - significance level  $\alpha$
  - power of the test  $1 \beta$
- *t*-tests for the means
- $\chi^2$ -test for the variance
- *z*-test for the population proportion
- *p*-value of the test



We presented the theorems last time:

- If  $X_1, X_2, ..., X_n \sim \mathcal{N}(\mu, \sigma^2)$  are independent, then  $\frac{\bar{X} \mu}{\sigma/\sqrt{n}} \sim \mathcal{N}(0, 1)$
- If  $X_1, X_2, ..., X_n \sim \mathcal{N}(\mu, \sigma^2)$  are independent, then  $\frac{\bar{X} \mu}{s/\sqrt{n}} \sim t_{n-1}$
- If  $X_1, X_2, ..., X_n \sim \mathcal{N}(\mu, \sigma^2)$  are independent, then  $\frac{(n-1)s^2}{\sigma^2} \sim \chi_{n-1}^2$

We used the theorems to establish the confidence intervals.



## Theorem:

• If  $X_1, X_2, ..., X_n \sim \mathcal{N}(\mu, \sigma^2)$  are independent, then  $\frac{\bar{X} - \mu}{\sigma/\sqrt{n}} \sim \mathcal{N}(0, 1)$ 

It also holds by the Lindenberg-Lévy Central Limit Theorem:

• If  $X_1, X_2, ..., X_n$  are independent and identically distributed and n is large, then  $\frac{\bar{X}-\mu}{\sigma/\sqrt{n}} \sim \mathcal{N}(0, 1)$ 

Confidence interval:

$$\mu \in \left[ \bar{x} - \frac{z(\alpha/2)\sigma}{\sqrt{n}}, \, \bar{x} + \frac{z(\alpha/2)\sigma}{\sqrt{n}} \right]$$





#### Theorem:

If 
$$X_1, X_2, ..., X_n \sim \mathcal{N}(\mu, \sigma^2)$$
 are independent, then  $\frac{\bar{X} - \mu}{s/\sqrt{n}} \sim t_{n-1}$ 

## **Confidence interval:**

$$\mu \in \left[\bar{x} - \frac{t_{n-1}(\alpha/2)s}{\sqrt{n}}, \, \bar{x} + \frac{t_{n-1}(\alpha/2)s}{\sqrt{n}}\right]$$

where  $t_{n-1}(\alpha/2) = F^{-1}\left(1 - \frac{\alpha}{2}\right)$ 

where F is the cumulative distribution function of



#### Theorem:

If 
$$X_1, X_2, ..., X_n \sim \mathcal{N}(\mu, \sigma^2)$$
 are independent, then  $\frac{(n-1)s^2}{\sigma^2} \sim \chi^2_{n-1}$ 

## **Confidence interval:**

$$\sigma^2 \in \left[\frac{(n-1)s^2}{\chi^2_{n-1}(\alpha/2)}, \frac{(n-1)s^2}{\chi^2_{n-1}(1-\alpha/2)}\right]$$

where 
$$\chi^2_{n-1}(\alpha/2) = F^{-1}\left(1 - \frac{\alpha}{2}\right)$$
 and  $\chi^2_{n-1}(1 - \alpha/2) = F^{-1}\left(\frac{\alpha}{2}\right)$ 

where F is the cumulative distribution function of

Now, we use the same theorems

- If  $X_1, X_2, ..., X_n \sim \mathcal{N}(\mu, \sigma^2)$  are independent, then  $\frac{\bar{X} \mu}{\sigma/\sqrt{n}} \sim \mathcal{N}(0, 1)$
- If  $X_1, X_2, ..., X_n \sim \mathcal{N}(\mu, \sigma^2)$  are independent, then  $\frac{\bar{X} \mu}{s/\sqrt{n}} \sim t_{n-1}$
- If  $X_1, X_2, \dots, X_n \sim \mathcal{N}(\mu, \sigma^2)$  are independent, then  $\frac{(n-1)s^2}{\sigma^2} \sim \chi_{n-1}^2$

to establish some <u>parametric</u> statistical <u>tests about the parameters</u> ( $\mu$ ,  $\sigma^2$ , ...) of the respective probability distribution.



# *z*-tests for the means



• One-sample *z*-test for

the population mean

• Paired-sample *z*-test for

the difference of the population means

• Two-sample *z*-test for

the difference of the population means

## Recall the theorem:

If 
$$X_1, X_2, ..., X_n \sim \mathcal{N}(\mu, \sigma^2)$$
 are independent, then  $\frac{\overline{X} - \mu}{\sigma/\sqrt{n}} \sim \mathcal{N}(0, 1)$ 

## Example or motivation:

Assume that  $X \sim \mathcal{N}(\mu, \sigma^2)$  is a random variable following the normal probability distribution. Assume that we do know the variance  $\sigma^2$ , but we do not know the true value of the population mean  $\mu \in \mathbb{R}$ . We conjecture / We assume / We ... / that the population mean  $\mu = \mu_0$ , i.e. the (unknown) population mean  $\mu$  is equal to some prescribed value  $\mu_0 \in \mathbb{R}$ .





Example: An archer shoots an arrow against the plane.

The sample space  $\Omega = \mathbb{R}^2 = \{ [x, y] : x, y \in \mathbb{R} \}$  is the set of all the points of the plane. The random variable X is the x-coordinate of the hit, i.e.  $X(\omega) = X([x, y]) = x.$ 

We know the archer's variance  $\sigma^2$ , but we do not know the archer's intention,

i.e. we do not know the point which the archer intends to hit,

i.e. we do not know the archer's mean  $\mu$ .

We conjecture that the archer's intention is to hit the origin, i.e.  $\mu = 0$ .



Let  $x_1 = X(\omega_1)$ ,  $x_2 = X(\omega_2)$ , ...,  $x_n = X(\omega_n)$  be the numerical results

of *n* trials of a random experiment, where  $X \sim \mathcal{N}(\mu, \sigma^2)$ ,

such as the x-coordinates of the archer's n hits.

We know the variance  $\sigma^2$ , but we do not know the mean  $\mu$ .

We state the null hypothesis (about the mean):

 $H_0: \quad \mu = \mu_0$ 

where  $\mu_0 \in \mathbb{R}$  is some number such that we conjecture that the true mean could equal the  $\mu_0$ .



## The meaning of the null hypothesis (such as $H_0: \mu = \mu_0$ in our example) is that

the observed distinct values are caused by the randomness only

(according to the assumed distribution, such as  $X \sim \mathcal{N}(\mu, \sigma^2)$  in our example)

- there are no other factors causing the distinct values
- everything is all right, no need to reconfigure anything
- all factors under the consideration are equivalent (have the same effect)

Having stated the null hypothesis

$$H_0: \quad \mu = \mu_0$$

## we also state the alternative hypothesis (denoted by $H_1$ or $H_A$ ).

There are three options how to state the alternative hypothesis:

- two-sided:  $H_1: \mu \neq \mu_0$
- one-sided:  $H_1$ :  $\mu < \mu_0$
- one-sided:  $H_1$ :  $\mu > \mu_0$







Which alternative hypothesis ( $\mu \neq \mu_0$  or  $\mu < \mu_0$  or  $\mu > \mu_0$ ) do we choose?

 $\rightarrow$  That depends upon our knowledge of the situation.

In our example:

- If we suspect that the archer's intention is to hit a point different from the given point (such as the origin), we choose  $H_1: \mu \neq \mu_0$ .
- If we conjecture that the archer's intention is to hit a point to the left of the given point (such as the origin), we choose  $H_1$ :  $\mu < \mu_0$ .
- If we conjecture that the archer's intention is to hit a point to the right of the given point (such as the origin), we choose  $H_1: \mu > \mu_0$ .



Under our assumptions  $(x_1, ..., x_n \sim \mathcal{N}(\mu, \sigma^2)$  are independent and  $\mu = \mu_0$ ),

it follows by the Theorem that

$$\frac{\bar{x}-\mu}{\sigma/\sqrt{n}} = \frac{\bar{x}-\mu_0}{\sigma/\sqrt{n}} \sim \mathcal{N}(0,1) \qquad exactly$$

Also, if  $x_1, ..., x_n$  are independent and identically distributed and n is large and  $\mu = \mu_0$ , then, by the Lindenberg-Lévy Central Limit Theorem,

$$\frac{\bar{x} - \mu}{\sigma / \sqrt{n}} = \frac{\bar{x} - \mu_0}{\sigma / \sqrt{n}} \sim \mathcal{N}(0, 1) \qquad approximately$$



Thus, having the *n* measurements  $x_1, x_2, ..., x_n$ , we calculate the statistic

$$Z = \frac{\bar{x} - \mu_0}{\sigma / \sqrt{n}}$$

#### where

- $\bar{x} = (\sum_{i=1}^{n} x_i)/n$  is the sample mean
- n is the sample size
- $\mu_0$  is the conjectured or estimated population mean
- $\sigma = \sqrt{\sigma^2}$  is the known standard deviation

We know (or assume) that  $Z \sim \mathcal{N}(0, 1)$ .

We have 
$$Z = \frac{\bar{x} - \mu_0}{\sigma / \sqrt{n}} \sim \mathcal{N}(0, 1)$$
.  
Then, if  $-\infty \le a < b \le +\infty$ , the probability that  $a < Z < b$  is

$$P(a < Z < b) = \int_{a}^{b} \varphi(x) \, \mathrm{d}x$$

where

$$\varphi(x) = \frac{1}{\sqrt{2\pi}} e^{-\frac{x^2}{2}}$$

is the density of the normalized normal distribution.



Consider the first case  $(H_1: \mu \neq \mu_0)$  first. We have:

$$\begin{array}{ll} H_0: & \mu = \mu_0 \\ \\ H_1: & \mu \neq \mu_0 \end{array} \end{array}$$

Knowing that 
$$Z = \frac{\bar{x} - \mu_0}{\sigma / \sqrt{n}} \sim \mathcal{N}(0, 1)$$
, the probability  $P(-c < Z < +c)$ 

is quite high,

so having -c < Z < +c accords with  $H_0$ , if c > 0 is large enough.





On the other hand, if  $H_0$  is true, then it is quite improbable that  $Z \notin (-c, +c)$ . Therefore, if we observe that

$$Z \leq -c \quad \text{or} \quad +c \leq Z$$

then we may conclude that  $H_0$  is probably not true,

i.e. we reject the null hypothesis  $H_0$ .

Therefore, the statistical test proceeds as follows: (see below)



Statistical one-sample z-test with two-sided alternative hypothesis ( $\mu \neq \mu_0$ ):

• choose the level of significance, a small number  $\alpha > 0$ , a very

popular value is  $\alpha = 5$  %, other popular values are 10 % or 1 % or 0.1 % etc.

• find the critical value c > 0 so that

$$\int_{-\infty}^{-c} \varphi(x) \, \mathrm{d}x = \frac{1}{2} \alpha \qquad \text{and} \qquad \int_{+c}^{+\infty} \varphi(x) \, \mathrm{d}x = \frac{1}{2} \alpha$$

for  $\alpha = 5$  %, the (two-sided) critical value is c = 1.959963 ...

- if  $Z \in (-\infty, -c] \cup [+c, +\infty)$ , the critical region, then reject the null hypothesis
- if  $Z \in (-c, +c)$ , then <u>do not reject</u> (or <u>fail to reject</u>) the null hypothesis

There are exactly four possibilities when testing the null hypothesis  $H_0$ :

- the null hypothesis  $(H_0)$  is actually true & we do not reject it OK
- the null hypothesis  $(H_0)$  is actually true & we reject it type I error
- the null hypothesis  $(H_0)$  is actually not true & we do not reject it type II error
- the null hypothesis  $(H_0)$  is actually not true & we reject it OK

The purpose is that the probability of the type I error and that of the type II error is as small as possible.





What is the probability of type I error

(the null hypothesis  $(H_0)$  is actually true & we reject it)?

The probability is equal to the significance level  $\alpha$ , usually  $\alpha = 5$  %.

<u>Recall:</u> The null hypothesis  $H_0$  is rejected if and only if  $Z \in (-\infty, -c] \cup [+c, +\infty)$ , i.e. if and only if  $|Z| \ge c$ . The critical value c is such that - if  $H_0$  holds true - then  $P(|Z| \ge c) = \alpha$ , i.e. the probability of the type I error (rejecting  $H_0$  when it is true) is  $\alpha$ .



What is the probability of type II error

(the null hypothesis  $(H_0)$  is actually false & we fail to reject it)?

The probability of type II error is denoted by  $\beta$ .

**The power of the test** is the probability  $1 - \beta$ 

It is much more difficult to calculate the probability  $\beta$  of type II error. It must be calculated for each test separately.



To calculate the probability  $\beta$  of type II error, consider that the null hypothesis

$$H_0$$
 is not true  $(\mu \neq \mu_0)$  and we fail to reject it  $(|Z| = \left| \frac{\bar{x} - \mu_0}{\sigma / \sqrt{n}} \right| < c)$ .  
By the Theorem then, we have  $\frac{\bar{x} - \mu}{\sigma / \sqrt{n}} = \frac{x - \mu_0 + (\mu_0 - \mu)}{\sigma / \sqrt{n}} \sim \mathcal{N}(0, 1)$ .

Then the probability of the type II error ( $H_0$  not true & fail to reject it) is:

$$P\left(-c < \frac{\bar{x} - \mu_0}{\sigma/\sqrt{n}} < +c\right) = P\left(-c < \frac{\bar{x} - \mu + (\mu_0 - \mu)}{\sigma/\sqrt{n}} < +c\right) =$$
$$= P\left(\frac{\mu - \mu_0}{\sigma/\sqrt{n}} - c < \frac{\bar{x} - \mu}{\sigma/\sqrt{n}} < \frac{\mu - \mu_0}{\sigma/\sqrt{n}} + c\right) =$$
$$= \beta = \int_{(\mu - \mu_0)/(\sigma/\sqrt{n}) - c}^{(\mu - \mu_0)/(\sigma/\sqrt{n}) + c} \varphi(x) \, \mathrm{d}x$$



Notice that, if the true  $\mu$  is close to the hypothesized  $\mu_0$  ( $\mu \approx \mu_0$ ), then  $\frac{\mu - \mu_0}{\sigma / \sqrt{n}} \approx 0$ , hence

$$\beta = \int_{(\mu-\mu_0)/(\sigma/\sqrt{n})-c}^{(\mu-\mu_0)/(\sigma/\sqrt{n})+c} \varphi(x) \, \mathrm{d}x \approx \int_{-c}^{+c} \varphi(x) \, \mathrm{d}x = 1 - \alpha = 95 \,\%$$

if  $\alpha = 5$  %, say.

It is recommended that  $\beta$  should be  $\leq 20$  %.

Therefore, if we wish to have  $\beta \approx 20$  % or  $\beta \leq 20$  %,

then we must not consider the true  $\mu$  close to the hypothesized  $\mu_0$ .

Consider now the second case  $(H_1: \mu < \mu_0)$ . We have:

 $H_0: \quad \mu = \mu_0$  $H_1: \quad \mu < \mu_0$ 

Knowing that  $Z = \frac{\bar{x} - \mu_0}{\sigma/\sqrt{n}} \sim \mathcal{N}(0, 1)$  and assuming that c > 0 is large enough, what is the probability that -c < Z?

If  $H_0$  is true, then it is quite improbable that  $Z \notin (-c, +\infty)$ .

Therefore, if we observe that  $Z \leq -c$ , then we may conclude that  $H_0$  is probably not true, i.e. we reject the null hypothesis  $H_0$ .





Statistical one-sample z-test with one-sided alternative hypothesis ( $\mu < \mu_0$ ):

- choose the level of significance, a small number  $\alpha > 0$ , such as  $\alpha = 5$  %, other popular values are  $\alpha = 10$  % or  $\alpha = 1$  % or  $\alpha = 0.1$  % etc.
- find the critical value c > 0 so that

$$\int_{-\infty}^{-c} \varphi(x) \, \mathrm{d}x = \alpha$$

for  $\alpha = 5$  %, the (one-sided) critical value is -c = -1.64485 ...

- if  $Z \in (-\infty, -c]$ , the critical region, then <u>reject</u> the null hypothesis
- if  $Z \in (-c, +\infty)$ , then <u>do not reject</u> (or <u>fail to reject</u>) the null hypothesis

Consider finally the third case  $(H_1: \mu > \mu_0)$ . We have:

 $H_0: \quad \mu = \mu_0$  $H_1: \quad \mu > \mu_0$ 

Knowing that  $Z = \frac{\bar{x} - \mu_0}{\sigma/\sqrt{n}} \sim \mathcal{N}(0, 1)$  and assuming that c > 0 is large enough, what is the probability that Z < +c?

If  $H_0$  is true, then it is quite improbable that  $Z \notin (-\infty, +c)$ .

Therefore, if we observe that  $+c \leq Z$ , then we may conclude that

 $H_0$  is probably not true, i.e. we reject the null hypothesis  $H_0$ .





Statistical one-sample z-test with one-sided alternative hypothesis ( $\mu > \mu_0$ ):

- choose the level of significance, a small number  $\alpha > 0$ , such as  $\alpha = 5$  %, other popular values are  $\alpha = 10$  % or  $\alpha = 1$  % or  $\alpha = 0.1$  % etc.
- find the critical value c > 0 so that

$$\int_{+c}^{+\infty} \varphi(x) \, \mathrm{d}x = \alpha$$

for  $\alpha = 5$  %, the (one-sided) critical value is +c = +1.64485 ...

- if  $Z \in [+c, +\infty)$ , the critical region, then <u>reject</u> the null hypothesis
- if  $Z \in (-\infty, +c)$ , then <u>do not reject</u> (or <u>fail to reject</u>) the null hypothesis

## Example or motivation:

Let us have a sample of n objects, e.g. n patients.

We do two measurements with each of the objects (patients)

- before some treatment
- after the treatment

The purpose it to learn whether the treatment has any effect. (Hence the null hypothesis: "The treatment has no effect.") Let  $x_1, x_2, ..., x_n$  be the values measured before the treatment, and let  $y_1, y_2, ..., y_n$  be the values measured after the treatment.





That is, the measurement  $x_i$  and  $y_i$  is done with the *i*-th object (patient) before and after the treatment for i = 1, 2, ..., n.

We assume that  $X \sim \mathcal{N}(\mu^{\text{before}}, \sigma_X^2)$ , i.e. the random variable of the measurement before the treatment follows the normal distribution, and that  $Y \sim \mathcal{N}(\mu^{\text{after}}, \sigma_Y^2)$ , i.e. the random variable of the measurement after the treatment also follows the normal distribution. We do not know the true values of the population means  $\mu^{\text{before}}$  and  $\mu^{\text{after}}$ , but we do assume that we know the variances  $\sigma_X^2$  and  $\sigma_Y^2$ .



We formulate the **null hypothesis**:

the treatment has no effect, i.e. the population means are the same

 $H_0$ :  $\mu^{\text{before}} = \mu^{\text{after}}$ 

Recall that we do not know the true population means  $\mu^{\text{before}}$  and  $\mu^{\text{after}}$ . We only test the hypothesis by having done a sample of *n* pairs of measurements.

Formulate the alternative hypothesis:

- two-sided:  $H_1: \mu^{\text{before}} \neq \mu^{\text{afer}}$
- one-sided:  $H_1$ :  $\mu^{\text{before}} < \mu^{\text{after}}$
- one-sided:  $H_1: \mu^{\text{before}} > \mu^{\text{after}}$

(the treatment has <u>some effect</u>) (the treatment <u>increases</u> / ...

... / decreases the quantity)

## Recall the theorem:

If 
$$X_1, X_2, ..., X_n \sim \mathcal{N}(\mu, \sigma^2)$$
 are independent, then  $\frac{\overline{X} - \mu}{\sigma/\sqrt{n}} \sim \mathcal{N}(0, 1)$ 

## Notice also:

If  $X_1, X_2, ..., X_n \sim \mathcal{N}(\mu^{\text{before}}, \sigma_X^2)$  and  $Y_1, Y_2, ..., Y_n \sim \mathcal{N}(\mu^{\text{after}}, \sigma_Y^2)$  are independent, then the differences

$$X_1 - Y_1, X_2 - Y_2, ..., X_n - Y_n \sim \mathcal{N}(\mu^{\text{before}} - \mu^{\text{after}}, \sigma_X^2 + \sigma_Y^2)$$

Now, the hypothesis  $\mu^{\text{before}} = \mu^{\text{after}}$  is equivalent to that the mean of the difference X - Y is  $\mu = \mu_0 = 0$ .





We have thus

reduced

## the paired-sample z-test for the difference of the population means

to

### the one-sample z-test for the population mean,

which we already know.



 $(\mu^{\text{before}} = \mu^{\text{after}})$ 

Having the *n* pairs of the measurements  $x_1, x_2, ..., x_n$  and  $y_1, y_2, ..., y_n$ , calculate the statistic

$$Z = \frac{\bar{x} - \bar{y}}{\sqrt{\sigma_X^2 + \sigma_Y^2} / \sqrt{n}} \quad \text{or} \quad Z = \frac{\bar{x} - \bar{y} - \mu_0}{\sqrt{\sigma_X^2 + \sigma_Y^2} / \sqrt{n}}$$

#### where

- $\bar{x} = (\sum_{i=1}^{n} x_i)/n$  is the sample mean of the measurements <u>before</u> the treatment
- $\bar{y} = (\sum_{i=1}^{n} y_i)/n$  is the sample mean of the measurements <u>after</u> the treatment
- n is the number of the pairs
- $\mu_0 = 0$  for no difference of the means
- $\mu_0 = \text{const.}$  for a general difference of the means ( $\mu^{\text{before}} = \mu^{\text{after}} + \text{const.}$ )

In the first case ( $H_1$ :  $\mu^{\text{before}} \neq \mu^{\text{after}}$ ), we have:

$$H_0: \quad \mu^{\text{before}} = \mu^{\text{after}}$$
$$H_1: \quad \mu^{\text{before}} \neq \mu^{\text{after}}$$

Knowing that

$$Z = \frac{\bar{x} - \bar{y}}{\sqrt{\sigma_X^2 + \sigma_Y^2} / \sqrt{n}} \sim \mathcal{N}(0, 1)$$

we fail to reject  $H_0$  iff

-c < Z < +c

where the critical value c > 0, under the assumption that  $H_0$  is true, is such that  $P(-c < Z < +c) = 1 - \alpha$  where the probability  $\alpha$  of type I error is small.




Statistical paired-sample *z*-test for the difference of the population means with two-sided alternative hypothesis ( $\mu^{\text{before}} \neq \mu^{\text{after}}$ ):

- choose the level of significance, a small number  $\alpha > 0$ , such as  $\alpha = 5 \%$
- find the critical value c > 0 so that

$$\int_{-\infty}^{-c} \varphi(x) \, \mathrm{d}x = \frac{1}{2} \alpha \qquad \text{and} \qquad \int_{+c}^{+\infty} \varphi(x) \, \mathrm{d}x = \frac{1}{2} \alpha$$

for  $\alpha = 5$  %, the (two-sided) critical value is c = 1.959963 ...

- if  $Z \in (-\infty, -c] \cup [+c, +\infty)$ , the critical region, then <u>reject</u> the null hypothesis
- if  $Z \in (-c, +c)$ , then <u>do not reject</u> (or <u>fail to reject</u>) the null hypothesis

In the second case ( $H_1$ :  $\mu^{\text{before}} < \mu^{\text{after}}$ ), we have:

 $\begin{array}{ll} H_0: & \mu^{\text{before}} = \mu^{\text{after}} \\ H_1: & \mu^{\text{before}} < \mu^{\text{after}} \end{array} \end{array}$ 

Knowing that

$$Z = \frac{\bar{x} - \bar{y}}{\sqrt{\sigma_X^2 + \sigma_Y^2} / \sqrt{n}} \sim \mathcal{N}(0, 1)$$

we fail to reject  $H_0$  iff

-c < Z

where the critical value c > 0, under the assumption that  $H_0$  is true, is such that  $P(-c < Z) = 1 - \alpha$  where the probability  $\alpha$  of type I error is small.





Statistical paired-sample *z*-test for the difference of the population means with one-sided alternative hypothesis ( $\mu^{\text{before}} < \mu^{\text{after}}$ ):

- choose the level of significance, a small number  $\alpha > 0$ , such as  $\alpha = 5 \%$
- find the critical value c > 0 so that

$$\int_{-\infty}^{-c} \varphi(x) \, \mathrm{d}x = \alpha$$

for  $\alpha = 5$  %, the (one-sided) critical value is -c = -1.64485 ...

- if  $Z \in (-\infty, -c]$ , the critical region, then <u>reject</u> the null hypothesis
- if  $Z \in (-c, +\infty)$ , then <u>do not reject</u> (or <u>fail to reject</u>) the null hypothesis

In the third case ( $H_1$ :  $\mu^{\text{before}} > \mu^{\text{after}}$ ), we have:

 $H_0: \quad \mu^{\text{before}} = \mu^{\text{after}}$  $H_1: \quad \mu^{\text{before}} > \mu^{\text{after}}$ 

Knowing that

$$Z = \frac{\bar{x} - \bar{y}}{\sqrt{\sigma_X^2 + \sigma_Y^2} / \sqrt{n}} \sim \mathcal{N}(0, 1)$$

we fail to reject  $H_0$  iff

Z < +c

where the critical value c > 0, under the assumption that  $H_0$  is true, is such that  $P(Z < +c) = 1 - \alpha$  where the probability  $\alpha$  of type I error is small.





Statistical paired-sample *z*-test for the difference of the population means with one-sided alternative hypothesis ( $\mu^{\text{before}} > \mu^{\text{after}}$ ):

- choose the level of significance, a small number  $\alpha > 0$ , such as  $\alpha = 5 \%$
- find the critical value c > 0 so that

$$\int_{+c}^{+\infty} \varphi(x) \, \mathrm{d}x = \alpha$$

for  $\alpha = 5$  %, the (one-sided) critical value is +c = +1.64485 ...

- if  $Z \in [+c, +\infty)$ , the critical region, then <u>reject</u> the null hypothesis
- if  $Z \in (-\infty, +c)$ , then <u>do not reject</u> (or <u>fail to reject</u>) the null hypothesis

### Motivation:

We have two unknown random variables X and Y. We ask (test the hypothesis) whether the population means of both random variables are the same.

We assume that both random variables are normal, i.e.  $X \sim \mathcal{N}(\mu_X, \sigma_X^2)$  and  $Y \sim \mathcal{N}(\mu_Y, \sigma_Y^2)$ , and that we know their variances  $\sigma_X^2$  and  $\sigma_Y^2$ .

We sample the variable X m-times, so we have the sample  $x_1, x_2, ..., x_m$ . We sample the variable Y n-times, so we have the sample  $y_1, y_2, ..., y_n$ .





Having the *m* observations  $x_1, x_2, ..., x_m$  of the random variable  $X \sim \mathcal{N}(\mu_X, \sigma_X^2)$  and having the *n* observations  $y_1, y_2, ..., y_n$  of the random variable  $Y \sim \mathcal{N}(\mu_Y, \sigma_Y^2)$ , we formulate the **null hypothesis**:

both samples come from the same population: the values of the population means are the same

 $H_0: \quad \mu_X = \mu_Y$ 

Recall that we do not know the true population means  $\mu_X$  and  $\mu_Y$ . We only test the hypothesis by means of two samples of m and n measurements.



Having the *m* observations  $x_1, x_2, ..., x_m$  of the random variable  $X \sim \mathcal{N}(\mu_X, \sigma_X^2)$ ,

the *n* observations  $y_1, y_2, ..., y_n$  of the random variable  $Y \sim \mathcal{N}(\mu_Y, \sigma_Y^2)$ , and

 $H_0: \quad \mu_X = \mu_Y$ 

### formulate the alternative hypothesis:

- two-sided:  $H_1: \mu_X \neq \mu_Y$  (the means are different)
- one-sided:  $H_1$ :  $\mu_X < \mu_Y$  (the first mean < the second mean)

• one-sided:  $H_1$ :  $\mu_X > \mu_Y$  (the first mean > the second mean)

### Recall the theorem:

If  $X_1, X_2, ..., X_m \sim \mathcal{N}(\mu_X, \sigma_X^2)$  and  $Y_1, Y_2, ..., Y_n \sim \mathcal{N}(\mu_Y, \sigma_Y^2)$  are independent, then

$$\frac{\overline{X} - \mu_X}{\sigma_X / \sqrt{m}} \sim \mathcal{N}(0, 1)$$
 and  $\frac{\overline{Y} - \mu_Y}{\sigma_Y / \sqrt{n}} \sim \mathcal{N}(0, 1)$ 

equivalently

$$\bar{X} \sim \mathcal{N}\left(\mu_X, \frac{\sigma_X^2}{m}\right)$$
 and  $\bar{Y} \sim \mathcal{N}\left(\mu_Y, \frac{\sigma_Y^2}{n}\right)$ 

Therefore:

$$\bar{X} - \bar{Y} \sim \mathcal{N}\left(\mu_X - \mu_Y, \frac{\sigma_X^2}{m} + \frac{\sigma_Y^2}{n}\right)$$



#### We have shown:

If  $X_1, X_2, ..., X_m \sim \mathcal{N}(\mu_X, \sigma_X^2)$  and  $Y_1, Y_2, ..., Y_n \sim \mathcal{N}(\mu_Y, \sigma_Y^2)$  are independent, then

$$\bar{X} - \bar{Y} \sim \mathcal{N}\left(\mu_X - \mu_Y, \ \frac{\sigma_X^2}{m} + \frac{\sigma_Y^2}{n}\right)$$

equivalently

$$\frac{(\bar{X}-\bar{Y})-(\mu_X-\mu_Y)}{\sqrt{\frac{\sigma_X^2}{m}+\frac{\sigma_Y^2}{n}}}\sim \mathcal{N}(0,1)$$





Having the *m* measurements  $x_1, x_2, ..., x_m$  and *n* measurements  $y_1, y_2, ..., y_n$ , calculate the statistic

$$Z = \frac{\bar{x} - \bar{y}}{\sqrt{\sigma_X^2/m + \sigma_Y^2/n}} \quad \text{or} \quad Z = \frac{\bar{x} - \bar{y} - \mu_0}{\sqrt{\sigma_X^2/m + \sigma_Y^2/n}}$$

where

- $\bar{x} = \sum_{i=1}^{m} x_i/m$  is the sample mean of the <u>first</u> sample
- $\bar{y} = \sum_{j=1}^{n} y_j / n$  is the sample mean of the <u>second</u> sample
- *m* and *n* is the size of the first and second, respectively, sample
- $\mu_0 = 0$  for no difference of the means  $(\mu_X = \mu_Y)$
- $\mu_0 = \text{const.}$  for a general difference of the means ( $\mu_X = \mu_Y + \text{const.}$ )

In the first case  $(H_1: \mu_X \neq \mu_Y)$ , we have:

$$H_0: \quad \mu_X = \mu_Y$$
$$H_1: \quad \mu_X \neq \mu_Y$$

Knowing that

$$Z = \frac{\bar{x} - \bar{y}}{\sqrt{\sigma_X^2/m + \sigma_Y^2/n}} \sim \mathcal{N}(0, 1)$$

we <u>fail to reject</u>  $H_0$  iff

-c < Z < +c

where the critical value c > 0, under the assumption that  $H_0$  is true, is such that  $P(-c < Z < +c) = 1 - \alpha$  where the probability  $\alpha$  of type I error is small.





Statistical two-sample *z*-test for the difference of the population means with two-sided alternative hypothesis ( $\mu_X \neq \mu_Y$ ):

- choose the level of significance, a small number  $\alpha > 0$ , such as  $\alpha = 5 \%$
- find the critical value c > 0 so that

$$\int_{-\infty}^{-c} \varphi(x) \, \mathrm{d}x = \frac{1}{2} \alpha \qquad \text{and} \qquad \int_{+c}^{+\infty} \varphi(x) \, \mathrm{d}x = \frac{1}{2} \alpha$$

for  $\alpha = 5$  %, the (two-sided) critical value is c = 1.959963 ...

- if  $Z \in (-\infty, -c] \cup [+c, +\infty)$ , the critical region, then reject the null hypothesis
- if  $Z \in (-c, +c)$ , then <u>do not reject</u> (or <u>fail to reject</u>) the null hypothesis

In the second case ( $H_1$ :  $\mu_X < \mu_Y$ ), we have:

$$H_0: \quad \mu_X = \mu_Y$$
$$H_1: \quad \mu_X < \mu_Y$$

Knowing that

$$Z = \frac{\bar{x} - \bar{y}}{\sqrt{\sigma_X^2/m + \sigma_Y^2/n}} \sim \mathcal{N}(0, 1)$$

we <u>fail to reject</u>  $H_0$  iff

-c < Z

where the critical value c > 0, under the assumption that  $H_0$  is true, is such that  $P(-c < Z) = 1 - \alpha$  where the probability  $\alpha$  of type I error is small.





Statistical two-sample *z*-test for the difference of the population means with one-sided alternative hypothesis ( $\mu_X < \mu_Y$ ):

- choose the level of significance, a small number  $\alpha > 0$ , such as  $\alpha = 5 \%$
- find the critical value c > 0 so that

$$\int_{-\infty}^{-c} \varphi(x) \, \mathrm{d}x = \alpha$$

for  $\alpha = 5$  %, the (one-sided) critical value is -c = -1.64485 ...

- if  $Z \in (-\infty, -c]$ , the critical region, then <u>reject</u> the null hypothesis
- if  $Z \in (-c, +\infty)$ , then <u>do not reject</u> (or <u>fail to reject</u>) the null hypothesis

In the third case  $(H_1: \mu_X > \mu_Y)$ , we have:

$$H_0: \quad \mu_X = \mu_Y$$
$$H_1: \quad \mu_X > \mu_Y$$

Knowing that

$$Z = \frac{\bar{x} - \bar{y}}{\sqrt{\sigma_X^2/m + \sigma_Y^2/n}} \sim \mathcal{N}(0, 1)$$

we <u>fail to reject</u>  $H_0$  iff

Z < +c

where the critical value c > 0, under the assumption that  $H_0$  is true, is such that  $P(Z < +c) = 1 - \alpha$  where the probability  $\alpha$  of type I error is small.





Statistical two-sample *z*-test for the difference of the population means with one-sided alternative hypothesis ( $\mu_X > \mu_Y$ ):

- choose the level of significance, a small number  $\alpha > 0$ , such as  $\alpha = 5 \%$
- find the critical value c > 0 so that

$$\int_{+c}^{+\infty} \varphi(x) \, \mathrm{d}x = \alpha$$

for  $\alpha = 5$  %, the (one-sided) critical value is +c = +1.64485 ...

- if  $Z \in [+c, +\infty)$ , the critical region, then <u>reject</u> the null hypothesis
- if  $Z \in (-\infty, +c)$ , then <u>do not reject</u> (or <u>fail to reject</u>) the null hypothesis

# *t*-tests for the means



One-sample *t*-test for

the population mean

• Paired-sample *t*-test for

the difference of the population means

• Two-sample *t*-test for

the difference of the population means

### Recall the theorem:

If 
$$X_1, X_2, ..., X_n \sim \mathcal{N}(\mu, \sigma^2)$$
 are independent, then  $\frac{\overline{X} - \mu}{\sigma/\sqrt{n}} \sim \mathcal{N}(0, 1)$ 

# Recall that, above, we have been using this theorem for the *z*-tests about the mean.

Notice, however, that we hardly ever know the variance  $\sigma^2$  of the random variable *X* in practice.

That is why we recall another theorem now.



One-sample t-test for the population mean

### Recall another theorem:

If 
$$X_1, X_2, ..., X_n \sim \mathcal{N}(\mu, \sigma^2)$$
 are independent, then  $\frac{(n-1)s^2}{\sigma^2} \sim \chi_{n-1}^2$ 

### where

•  $s^2 = \sum_{i=1}^n (X_i - \overline{X})^2 / (n-1)$  is the sample variance of the random variables •  $\chi^2_{n-1}$  is Pearson's  $\chi^2$ -distribution with n-1 degrees of freedom



Recall the corollary of the two theorems:

One-sample *t*-test for the population mean

If  $X_1, X_2, ..., X_n \sim \mathcal{N}(\mu, \sigma^2)$  are independent, then  $\frac{\bar{x} - \mu}{s/\sqrt{n}} \sim t_{n-1}$ 

where

- $\overline{X} = \sum_{i=1}^{n} X_i / n$  is the sample mean of the random variables
- $s = \sqrt{\sum_{i=1}^{n} (X_i \bar{X})^2 / (n 1)}$  is the sample standard deviation of the random variables

is Student's *t*-distribution

with n-1 degrees of freedom





Recall the corollary:

If 
$$X_1, X_2, ..., X_n \sim \mathcal{N}(\mu, \sigma^2)$$
 are independent, then  $\frac{\overline{X} - \mu}{s/\sqrt{n}} \sim t_{n-1}$ 

### Example or motivation:

Assume that  $X \sim \mathcal{N}(\mu, \sigma^2)$  is a random variable following the normal probability distribution with some mean  $\mu \in \mathbb{R}$  and with some variance  $\sigma^2 \in \mathbb{R}_0^+$ . Knowing <u>neither the variance</u>  $\sigma^2$  <u>nor the true value of the population mean</u>  $\mu \in \mathbb{R}$ , we conjecture / we assume / we ... / that the population mean  $\mu = \mu_0$ , i.e. the (unknown) population mean  $\mu$  is equal to some prescribed value  $\mu_0 \in \mathbb{R}$ .





Example: An archer shoots an arrow against the plane.

The sample space  $\Omega = \mathbb{R}^2 = \{ [x, y] : x, y \in \mathbb{R} \}$  is the set of all the points of the plane. The random variable X is the x-coordinate of the hit, i.e.  $X(\omega) = X([x, y]) = x.$ 

We do not know the archer's variance  $\sigma^2$  and we do not know the archer's intention, i.e. we do not know the point which the archer intends to hit, i.e. we do not know the archer's mean  $\mu$ .

We conjecture that the archer's intention is to hit the origin, i.e.  $\mu = 0$ .



Let  $x_1 = X(\omega_1)$ ,  $x_2 = X(\omega_2)$ , ...,  $x_n = X(\omega_n)$  be the numerical results

of *n* trials of a random experiment, where  $X \sim \mathcal{N}(\mu, \sigma^2)$ ,

such as the x-coordinates of the archer's n hits.

We do not know the variance  $\sigma^2$  and we do not know the mean  $\mu$ .

We state the null hypothesis (about the mean):

 $H_0: \quad \mu = \mu_0$ 

where  $\mu_0 \in \mathbb{R}$  is some number such that we conjecture that the true mean could equal the  $\mu_0$ .

Having stated the null hypothesis

 $H_0: \quad \mu = \mu_0$ 

### we also state the alternative hypothesis:

- two-sided:  $H_1: \mu \neq \mu_0$
- one-sided:  $H_1: \mu < \mu_0$
- one-sided:  $H_1$ :  $\mu > \mu_0$







Which alternative hypothesis ( $\mu \neq \mu_0$  or  $\mu < \mu_0$  or  $\mu > \mu_0$ ) do we choose?

 $\rightarrow$  That depends upon our knowledge of the situation.

In our example:

- If we suspect that the archer's intention is to hit a point different from the given point (such as the origin), we choose  $H_1: \mu \neq \mu_0$ .
- If we conjecture that the archer's intention is to hit a point to the left of the given point (such as the origin), we choose  $H_1$ :  $\mu < \mu_0$ .
- If we conjecture that the archer's intention is to hit a point to the right of the given point (such as the origin), we choose  $H_1: \mu > \mu_0$ .



Under our assumptions  $(x_1, ..., x_n \sim \mathcal{N}(\mu, \sigma^2)$  are independent and  $\mu = \mu_0$ ), it follows by the Theorem that

$$\frac{\bar{x}-\mu}{s/\sqrt{n}} = \frac{\bar{x}-\mu_0}{s/\sqrt{n}} \sim t_{n-1}$$

Thus, having the *n* measurements  $x_1, x_2, ..., x_n$ , we calculate the statistic

$$T = \frac{\bar{x} - \mu_0}{s / \sqrt{n}}$$

We know (or assume) that  $T \sim t_{n-1}$ .

We have 
$$T = \frac{\bar{x} - \mu_0}{s/\sqrt{n}} \sim t_{n-1}$$
.  
Then, if  $-\infty \le a < b \le +\infty$ , the probability that  $a < T < b$  is

$$P(a < T < b) = \int_{a}^{b} f(x) \, \mathrm{d}x$$

where

$$f(x) = \frac{\Gamma\left(\frac{n}{2}\right)}{\Gamma\left(\frac{n-1}{2}\right)\sqrt{\pi(n-1)}} \left(1 + \frac{x^2}{n-1}\right)^{-\frac{k}{2}}$$

is the density of Student's *t*-distribution with n-1 degrees of freedom.





$$\Gamma(z) = \int_0^{+\infty} x^z e^{-x} dx \quad \text{for } z \in \mathbb{C} \text{ such that } \operatorname{Re}(z) > 0$$

It is easy to calculate:

 $\Gamma(1) = 1$  $\Gamma(z+1) = z\Gamma(z)$ 

Therefore:

 $\Gamma(n+1) = n!$  for n = 0, 1, 2, 3, ...

## The gamma function – another definition (due to Euler)



$$\Gamma(z) = \frac{1}{z} \prod_{n=1}^{\infty} \frac{\left(1 + \frac{1}{n}\right)^z}{1 + \frac{z}{n}}$$

for 
$$z \in \mathbb{C} \setminus \{0, -1, -2, -3, ...\}$$

Consider the first case  $(H_1: \mu \neq \mu_0)$  first. We have:

$$H_0: \quad \mu = \mu_0$$
$$H_1: \quad \mu \neq \mu_0$$

Knowing that 
$$T = \frac{\bar{x} - \mu_0}{s/\sqrt{n}} \sim t_{n-1}$$
, the probability  
 $P(-c < T < +c)$ 

is quite high,

so having -c < T < +c accords with  $H_0$ , if c > 0 is large enough.





On the other hand, if  $H_0$  is true, then it is quite improbable that  $T \notin (-c, +c)$ . Therefore, if we observe that

$$T \leq -c$$
 or  $+c \leq T$ 

then we may conclude that  $H_0$  is probably not true,

i.e. we reject the null hypothesis  $H_0$ .

Therefore, the statistical test proceeds as follows: (see below)



Statistical one-sample *t*-test with two-sided alternative hypothesis ( $\mu \neq \mu_0$ ):

- choose the level of significance, a small number  $\alpha > 0$ , a very popular value is  $\alpha = 5$  %, other popular values are 10 % or 1 % or 0.1 % etc.
- find the critical value c > 0 so that

$$\int_{-\infty}^{-c} f(x) \, \mathrm{d}x = \frac{1}{2} \alpha \qquad \text{and} \qquad \int_{+c}^{+\infty} f(x) \, \mathrm{d}x = \frac{1}{2} \alpha$$

where f is the density of the t-distribution with n-1 degrees of freedom

- if  $T \in (-\infty, -c] \cup [+c, +\infty)$ , the critical region, then <u>reject</u> the null hypothesis
- if  $T \in (-c, +c)$ , then <u>do not reject</u> (or <u>fail to reject</u>) the null hypothesis



There are exactly four possibilities when testing the null hypothesis  $H_0$ :

- the null hypothesis  $(H_0)$  is actually true **&** we do not reject it **—** OK
- the null hypothesis (H<sub>0</sub>) is actually true
  & we reject it
   type I error
- the null hypothesis (H<sub>0</sub>) is actually not true & we do not reject it type II error
- the null hypothesis  $(H_0)$  is actually not true & we reject it OK

The probability of the type 1 error is the significance level a

The probability of the type II error is  $\beta$ 

The **power of the test** is the probability  $1 - \beta$ 

Consider now the second case  $(H_1: \mu < \mu_0)$ . We have:

 $H_0: \quad \mu = \mu_0$  $H_1: \quad \mu < \mu_0$ 

Knowing that  $T = \frac{\bar{x} - \mu_0}{s/\sqrt{n}} \sim t_{n-1}$  and assuming that c > 0 is large enough, what is the probability that -c < T?

If  $H_0$  is true, then it is quite improbable that  $T \notin (-c, +\infty)$ .

Therefore, if we observe that  $T \leq -c$ , then we may conclude that  $H_0$  is probably not true, i.e. we reject the null hypothesis  $H_0$ .





Statistical one-sample *t*-test with one-sided alternative hypothesis ( $\mu < \mu_0$ ):

- choose the level of significance, a small number  $\alpha > 0$ , such as  $\alpha = 5$  %, other popular values are  $\alpha = 10$  % or  $\alpha = 1$  % or  $\alpha = 0.1$  % etc.
- find the critical value c > 0 so that

$$\int_{-\infty}^{-c} f(x) \, \mathrm{d}x = \alpha$$

where f is the density of the t-distribution with n-1 degrees of freedom

- if  $T \in (-\infty, -c]$ , the critical region, then <u>reject</u> the null hypothesis
- if  $T \in (-c, +\infty)$ , then <u>do not reject</u> (or <u>fail to reject</u>) the null hypothesis
Consider finally the third case  $(H_1: \mu > \mu_0)$ . We have:

 $H_0: \quad \mu = \mu_0$  $H_1: \quad \mu > \mu_0$ 

Knowing that  $T = \frac{\bar{x} - \mu_0}{s/\sqrt{n}} \sim t_{n-1}$  and assuming that c > 0 is large enough, what is the probability that T < +c?

If  $H_0$  is true, then it is quite improbable that  $T \notin (-\infty, +c)$ .

Therefore, if we observe that  $+c \le T$ , then we may conclude that  $H_0$  is probably not true, i.e. we reject the null hypothesis  $H_0$ .





Statistical one-sample *t*-test with one-sided alternative hypothesis ( $\mu > \mu_0$ ):

- choose the level of significance, a small number  $\alpha > 0$ , such as  $\alpha = 5$  %, other popular values are  $\alpha = 10$  % or  $\alpha = 1$  % or  $\alpha = 0.1$  % etc.
- find the critical value c > 0 so that

$$\int_{+c}^{+\infty} f(x) \, \mathrm{d}x = \alpha$$

where f is the density of the t-distribution with n-1 degrees of freedom

- if  $T \in [+c, +\infty)$ , the critical region, then <u>reject</u> the null hypothesis
- if  $T \in (-\infty, +c)$ , then <u>do not reject</u> (or <u>fail to reject</u>) the null hypothesis

### Example or motivation:

Let us have a sample of n objects, e.g. n patients.

We do two measurements with each of the objects (patients)

- before some treatment
- after the treatment

The purpose it to learn whether the treatment has any effect. (Hence the null hypothesis: "The treatment has no effect.") Let  $x_1, x_2, ..., x_n$  be the values measured before the treatment, and let  $y_1, y_2, ..., y_n$  be the values measured after the treatment.





That is, the measurement  $x_i$  and  $y_i$  is done with the *i*-th object (patient) before and after the treatment for i = 1, 2, ..., n.

We assume that  $X \sim \mathcal{N}(\mu^{\text{before}}, \sigma_X^2)$ , i.e. the random variable of the measurement before the treatment follows the normal distribution, and that  $Y \sim \mathcal{N}(\mu^{\text{after}}, \sigma_Y^2)$ , i.e. the random variable of the measurement after the treatment also follows the normal distribution, for some  $\mu^{\text{before}}, \mu^{\text{after}} \in \mathbb{R}$  and for some  $\sigma_X^2, \sigma_Y^2 \in \mathbb{R}_0^+$ . We do not know the true values of the population means  $\mu^{\text{before}}$  and  $\mu^{\text{after}}$ , and we do not know the true values of the variances  $\sigma_X^2$  and  $\sigma_Y^2$ .



### We formulate the **null hypothesis**:

the treatment has no effect, i.e. the population means are the same

$$H_0: \quad \mu^{\text{before}} = \mu^{\text{after}}$$

Recall that we do not know the true population means  $\mu^{\text{before}}$  and  $\mu^{\text{after}}$ . We

only test the hypothesis by having done a sample of n pairs of measurements.

Formulate the alternative hypothesis:

- two-sided:  $H_1: \mu^{\text{before}} \neq \mu^{\text{afer}}$
- one-sided:  $H_1$ :  $\mu^{\text{before}} < \mu^{\text{after}}$
- one-sided:  $H_1$ :  $\mu^{\text{before}} > \mu^{\text{after}}$

(the treatment has <u>some effect</u>) (the treatment <u>increases</u> / ...

... / decreases the quantity)

### Recall the theorem:

If 
$$X_1, X_2, ..., X_n \sim \mathcal{N}(\mu, \sigma^2)$$
 are independent, then  $\frac{\overline{X} - \mu}{s/\sqrt{n}} \sim t_{n-1}$ 

### Notice also:

If  $X_1, X_2, ..., X_n \sim \mathcal{N}(\mu^{\text{before}}, \sigma_X^2)$  and  $Y_1, Y_2, ..., Y_n \sim \mathcal{N}(\mu^{\text{after}}, \sigma_Y^2)$  are independent, then the differences

$$X_1 - Y_1, X_2 - Y_2, ..., X_n - Y_n \sim \mathcal{N}(\mu^{\text{before}} - \mu^{\text{after}}, \sigma_X^2 + \sigma_Y^2)$$

Now, the hypothesis  $\mu^{\text{before}} = \mu^{\text{after}}$  is equivalent to that the mean of the difference X - Y is  $\mu = \mu_0 = 0$ .





We have thus

reduced

### the paired-sample t-test for the difference of the population means

to

### the one-sample t-test for the population mean,

which we already know.



Having the *n* pairs of the measurements  $x_1, x_2, ..., x_n$  and  $y_1, y_2, ..., y_n$ , calculate the statistic

$$T = \frac{\bar{x} - \bar{y}}{\sqrt{s^2} / \sqrt{n}} \quad \text{or} \quad T = \frac{\bar{x} - \bar{y} - \mu_0}{\sqrt{s^2} / \sqrt{n}}$$

where

- $\bar{x} = (\sum_{i=1}^{n} x_i)/n$  is the sample mean of the measurements <u>before</u> the treatment •  $\bar{y} = (\sum_{i=1}^{n} y_i)/n$  is the sample mean of the measurements <u>after</u> the treatment
- $\mu_0 = 0$  for no difference of the means  $(\mu^{\text{before}} = \mu^{\text{after}})$
- $\mu_0 = \text{const.}$  for a general difference of the means ( $\mu^{\text{before}} = \mu^{\text{after}} + \text{const.}$ )
- $s^2 = (\sum_{i=1}^n (x_i y_i \bar{x} + \bar{y})^2)/(n-1)$  is the sample variance of the differences

In the first case ( $H_1$ :  $\mu^{\text{before}} \neq \mu^{\text{after}}$ ), we have:

 $H_0: \quad \mu^{\text{before}} = \mu^{\text{after}}$  $H_1: \quad \mu^{\text{before}} \neq \mu^{\text{after}}$ 

Knowing that

$$T = \frac{\bar{x} - \bar{y}}{\sqrt{s^2} / \sqrt{n}} \sim t_{n-1}$$

we fail to reject  $H_0$  iff

-c < T < +c

where the critical value c > 0, under the assumption that  $H_0$  is true, is such that  $P(-c < T < +c) = 1 - \alpha$  where the probability  $\alpha$  of type I error is small.





Statistical paired-sample *t*-test for the difference of the population means with two-sided alternative hypothesis ( $\mu^{\text{before}} \neq \mu^{\text{after}}$ ):

- choose the level of significance, a small number  $\alpha > 0$ , such as  $\alpha = 5$  %
- find the critical value c > 0 so that

$$\int_{-\infty}^{-c} f(x) \, \mathrm{d}x = \frac{1}{2} \alpha \qquad \text{and} \qquad \int_{+c}^{+\infty} f(x) \, \mathrm{d}x = \frac{1}{2} \alpha$$

where f is the density of the t-distribution with n-1 degrees of freedom

- if  $T \in (-\infty, -c] \cup [+c, +\infty)$ , the critical region, then <u>reject</u> the null hypothesis
- if  $T \in (-c, +c)$ , then <u>do not reject</u> (or <u>fail to reject</u>) the null hypothesis

In the second case ( $H_1$ :  $\mu^{\text{before}} < \mu^{\text{after}}$ ), we have:

$$\begin{array}{ll} H_0: & \mu^{\text{before}} = \mu^{\text{after}} \\ H_1: & \mu^{\text{before}} < \mu^{\text{after}} \end{array} \end{array}$$

Knowing that

$$T = \frac{\bar{x} - \bar{y}}{\sqrt{s^2} / \sqrt{n}} \sim t_{n-1}$$

we fail to reject  $H_0$  iff

-c < T

where the critical value c > 0, under the assumption that  $H_0$  is true, is such that  $P(-c < T) = 1 - \alpha$  where the probability  $\alpha$  of type I error is small.





Statistical paired-sample *t*-test for the difference of the population means with one-sided alternative hypothesis ( $\mu^{\text{before}} < \mu^{\text{after}}$ ):

- choose the level of significance, a small number  $\alpha > 0$ , such as  $\alpha = 5 \%$
- find the critical value c > 0 so that

$$\int_{-\infty}^{-c} f(x) \, \mathrm{d}x = \alpha$$

where f is the density of the t-distribution with n-1 degrees of freedom

- if  $T \in (-\infty, -c]$ , the critical region, then <u>reject</u> the null hypothesis
- if  $T \in (-c, +\infty)$ , then <u>do not reject</u> (or <u>fail to reject</u>) the null hypothesis

In the third case  $(H_1: \mu^{\text{before}} > \mu^{\text{after}})$ , we have:

Knowing that

$$T = \frac{\bar{x} - \bar{y}}{\sqrt{s^2} / \sqrt{n}} \sim t_{n-1}$$

 $H_0$ :  $\mu^{\text{before}} = \mu^{\text{after}}$ 

 $H_1$ :  $\mu^{\text{before}} > \mu^{\text{after}}$ 

we fail to reject  $H_0$  iff

T < +c

where the critical value c > 0, under the assumption that  $H_0$  is true, is such that  $P(T < +c) = 1 - \alpha$  where the probability  $\alpha$  of type I error is small.





Statistical paired-sample *t*-test for the difference of the population means with one-sided alternative hypothesis ( $\mu^{\text{before}} > \mu^{\text{after}}$ ):

- choose the level of significance, a small number  $\alpha > 0$ , such as  $\alpha = 5 \%$
- find the critical value c > 0 so that

$$\int_{+c}^{+\infty} f(x) \, \mathrm{d}x = \alpha$$

where f is the density of the t-distribution with n-1 degrees of freedom

- if  $T \in [+c, +\infty)$ , the critical region, then <u>reject</u> the null hypothesis
- if  $T \in (-\infty, +c)$ , then <u>do not reject</u> (or <u>fail to reject</u>) the null hypothesis

### Motivation:

We have two unknown random variables X and Y. We ask (test the hypothesis) whether the population means of both random variables are the same.

We assume that both random variables are normal, i.e.  $X \sim \mathcal{N}(\mu_X, \sigma_X^2)$  and  $Y \sim \mathcal{N}(\mu_Y, \sigma_Y^2)$ , for some  $\mu_X, \mu_Y \in \mathbb{R}$  and for some  $\sigma_X^2, \sigma_Y^2 \in \mathbb{R}_0^+$ .

Although we do not know the means  $\mu_X$ ,  $\mu_Y$  nor the variances  $\sigma_X^2$ ,  $\sigma_Y^2$ , we assume that

$$||| ||| ||| ||| \sigma_X^2 = \sigma_Y^2 \quad ||| ||| |||$$





Having the *m* observations  $x_1, x_2, ..., x_m$  of the random variable  $X \sim \mathcal{N}(\mu_X, \sigma^2)$  and having the *n* observations  $y_1, y_2, ..., y_n$  of the random variable  $Y \sim \mathcal{N}(\mu_Y, \sigma^2)$ , we formulate the **null hypothesis**:

both samples come from the same population: the values of the population means are the same

 $H_0: \quad \mu_X = \mu_Y$ 

Recall that we do not know the true population means  $\mu_X$  and  $\mu_Y$ . We only **test the hypothesis by means of** two samples of m and n measurements with the same variance.



- Having the *m* observations  $x_1, x_2, ..., x_m$  of the random variable  $X \sim \mathcal{N}(\mu_X, \sigma^2)$ ,
  - the *n* observations  $y_1, y_2, ..., y_n$  of the random variable  $Y \sim \mathcal{N}(\mu_Y, \sigma^2)$ , and

$$H_0: \quad \mu_X = \mu_Y$$

### formulate the alternative hypothesis:

- two-sided:  $H_1: \mu_X \neq \mu_Y$  (the means are different)
- one-sided:  $H_1$ :  $\mu_X < \mu_Y$  (the first mean < the second mean)

• one-sided:  $H_1$ :  $\mu_X > \mu_Y$  (the first mean > the second mean)



### Recall the theorem:

If  $X_1, X_2, ..., X_m \sim \mathcal{N}(\mu_X, \sigma^2)$  and  $Y_1, Y_2, ..., Y_n \sim \mathcal{N}(\mu_Y, \sigma^2)$  are independent, then

$$\frac{\overline{X} - \mu_X}{\sigma/\sqrt{m}} \sim \mathcal{N}(0, 1) \quad \text{and} \quad \frac{\overline{Y} - \mu_Y}{\sigma/\sqrt{n}} \sim \mathcal{N}(0, 1)$$

equivalently

$$\bar{X} \sim \mathcal{N}\left(\mu_X, \frac{\sigma^2}{m}\right)$$
 and  $\bar{Y} \sim \mathcal{N}\left(\mu_Y, \frac{\sigma^2}{n}\right)$ 

Therefore:

$$\overline{X} - \overline{Y} \sim \mathcal{N}\left(\mu_X - \mu_Y, \frac{\sigma^2}{m} + \frac{\sigma^2}{n}\right)$$

## No.

### We have shown:

If  $X_1, X_2, ..., X_m \sim \mathcal{N}(\mu_X, \sigma^2)$  and  $Y_1, Y_2, ..., Y_n \sim \mathcal{N}(\mu_Y, \sigma^2)$  are independent, then

$$\bar{X} - \bar{Y} \sim \mathcal{N}\left(\mu_X - \mu_Y, \frac{\sigma^2}{m} + \frac{\sigma^2}{n}\right)$$

equivalently

$$\frac{(\bar{X}-\bar{Y})-(\mu_X-\mu_Y)}{\sigma\sqrt{\frac{1}{m}+\frac{1}{n}}}\sim\mathcal{N}(0,1)$$



### Recall the other theorem:

If  $X_1, X_2, ..., X_m \sim \mathcal{N}(\mu_X, \sigma^2)$  and  $Y_1, Y_2, ..., Y_n \sim \mathcal{N}(\mu_Y, \sigma^2)$  are independent, then

$$\frac{(m-1)s_X^2}{\sigma^2} \sim \chi_{m-1}^2 \quad \text{and} \quad \frac{(n-1)s_Y^2}{\sigma^2} \sim \chi_{n-1}^2$$

Therefore:

$$\frac{(m-1)s_X^2 + (n-1)s_Y^2}{\sigma^2} \sim \chi_{m+n-2}^2$$



Two-sample *t*-test for the diff. of the pop. means //  $\sigma_x = \sigma_y$ 

Recall also that, if

$$Z = \frac{(\bar{X} - \bar{Y}) - (\mu_X - \mu_Y)}{\sigma \sqrt{\frac{1}{m} + \frac{1}{n}}} \sim \mathcal{N}(0, 1)$$

and

$$Y = \frac{(m-1)s_X^2 + (n-1)s_Y^2}{\sigma^2} \sim \chi_{m+n-2}^2$$

then

$$T = \frac{Z}{\sqrt{\frac{Y}{m+n-2}}} \sim t_{m+n-2}$$

by the definition of Student's t-distribution.

### Two-sample *t*-test for the diff. of the pop. means // $\sigma_x = \sigma_y$



Therefore:

$$\frac{(\bar{X} - \bar{Y}) - (\mu_X - \mu_Y)}{\sqrt{\frac{(m-1)s_X^2 + (n-1)s_Y^2}{m+n-2}}} = \frac{\frac{(\bar{X} - \bar{Y}) - (\mu_X - \mu_Y)}{\sigma\sqrt{\frac{1}{m} + \frac{1}{n}}}}{\sqrt{\frac{(m-1)s_X^2 + (n-1)s_Y^2}{m+n-2}}} \sim t_{m+n-2}$$

where

$$s_{\bar{X}}^2 = \frac{\sum_{i=1}^m (X_i - \bar{X})^2}{m-1}$$
 and  $s_{\bar{Y}}^2 = \frac{\sum_{j=1}^n (Y_i - \bar{Y})^2}{n-1}$ 

are the sample variances.



Having the *m* measurements  $x_1, x_2, ..., x_m$  and *n* measurements  $y_1, y_2, ..., y_n$ , recall that

- $\bar{x} = \sum_{i=1}^m x_i/m$
- $\bar{y} = \sum_{j=1}^{n} y_j/n$
- $s_x^2 = \sum_{i=1}^m (x_i \bar{x})^2 / (m-1)$
- $s_y^2 = \sum_{j=1}^n (y_j \bar{y})^2 / (n-1)$

• m

• n

is the sample mean of the <u>first</u> sample is the sample mean of the <u>second</u> sample is the sample variance of the <u>first</u> sample is the sample variance of the <u>second</u> sample is the size of the <u>first</u> sample is the size of the <u>second</u> sample



Having the *m* measurements  $x_1, x_2, ..., x_m$  and *n* measurements  $y_1, y_2, ..., y_n$ , calculate the statistic

$$T = \frac{\bar{x} - \bar{y}}{\sqrt{\frac{(m-1)s_x^2 + (n-1)s_y^2}{m+n-2}}\sqrt{\frac{1}{m} + \frac{1}{n}}}$$

for no difference of the means  $(\mu_X = \mu_Y)$ 

We know (or assume) that  $T \sim t_{m+n-2}$ 



Or, having the *m* measurements  $x_1, x_2, ..., x_m$  and *n* measurements  $y_1, y_2, ..., y_n$ , calculate the statistic

$$T = \frac{\bar{x} - \bar{y} - \mu_0}{\sqrt{\frac{(m-1)s_x^2 + (n-1)s_y^2}{m+n-2}}} \sqrt{\frac{1}{m} + \frac{1}{n}}$$

for a general difference of the means  $(\mu_X = \mu_Y + \mu_0)$ 

We know (or assume) that  $T \sim t_{m+n-2}$ 

In the first case  $(H_1: \mu_X \neq \mu_Y)$ , we have:

$$H_0: \quad \mu_X = \mu_Y$$
$$H_1: \quad \mu_X \neq \mu_Y$$

Knowing that

$$T \sim t_{m+n-2}$$

we fail to reject  $H_0$  iff

-c < T < +c

where the critical value c > 0, under the assumption that  $H_0$  is true, is such that  $P(-c < T < +c) = 1 - \alpha$  where the probability  $\alpha$  of type I error is small.





Statistical two-sample *t*-test for the difference of the population means with two-sided alternative hypothesis ( $\mu_X \neq \mu_Y$ ) and with the same variances:

- choose the level of significance, a small number  $\alpha > 0$ , such as  $\alpha = 5 \%$
- find the critical value c > 0 so that

$$\int_{-\infty}^{-c} f(x) \, \mathrm{d}x = \frac{1}{2} \alpha \qquad \text{and} \qquad \int_{+c}^{+\infty} f(x) \, \mathrm{d}x = \frac{1}{2} \alpha$$

where f is the density of the t-distribution with m + n - 2 degrees of freedom

- if  $T \in (-\infty, -c] \cup [+c, +\infty)$ , the critical region, then <u>reject</u> the null hypothesis
- if  $T \in (-c, +c)$ , then <u>do not reject</u> (or <u>fail to reject</u>) the null hypothesis

In the second case ( $H_1$ :  $\mu_X < \mu_Y$ ), we have:

$$H_0: \quad \mu_X = \mu_Y$$
$$H_1: \quad \mu_X < \mu_Y$$

Knowing that

$$T \sim t_{m+n-2}$$

we fail to reject  $H_0$  iff

-c < T

where the critical value c > 0, under the assumption that  $H_0$  is true, is such that  $P(-c < T) = 1 - \alpha$  where the probability  $\alpha$  of type I error is small.





Statistical two-sample *t*-test for the difference of the population means with one-sided alternative hypothesis ( $\mu_X < \mu_Y$ ) and with the same variances :

- choose the level of significance, a small number  $\alpha > 0$ , such as  $\alpha = 5$  %
- find the critical value c > 0 so that

$$\int_{-\infty}^{-c} f(x) \, \mathrm{d}x = \alpha$$

where f is the density of the t-distribution with m + n - 2 degrees of freedom

- if  $T \in (-\infty, -c]$ , the critical region, then <u>reject</u> the null hypothesis
- if  $T \in (-c, +\infty)$ , then <u>do not reject</u> (or <u>fail to reject</u>) the null hypothesis

In the third case  $(H_1: \mu_X > \mu_Y)$ , we have:

$$H_0: \quad \mu_X = \mu_Y$$
$$H_1: \quad \mu_X > \mu_Y$$

Knowing that

$$T \sim t_{m+n-2}$$

we fail to reject  $H_0$  iff

T < +c

where the critical value c > 0, under the assumption that  $H_0$  is true, is such that  $P(T < +c) = 1 - \alpha$  where the probability  $\alpha$  of type I error is small.





Statistical two-sample *t*-test for the difference of the population means with one-sided alternative hypothesis ( $\mu_X > \mu_Y$ ):

- choose the level of significance, a small number  $\alpha > 0$ , such as  $\alpha = 5 \%$
- find the critical value c > 0 so that

$$\int_{+c}^{+\infty} f(x) \, \mathrm{d}x = \alpha$$

where f is the density of the t-distribution with m + n - 2 degrees of freedom

- if  $T \in [+c, +\infty)$ , the critical region, then <u>reject</u> the null hypothesis
- if  $T \in (-\infty, +c)$ , then <u>do not reject</u> (or <u>fail to reject</u>) the null hypothesis



Consider two normal random variables  $X \sim \mathcal{N}(\mu_X, \sigma_X^2)$  and  $Y \sim \mathcal{N}(\mu_Y, \sigma_Y^2)$ , for some  $\mu_X, \mu_Y \in \mathbb{R}$  and for some  $\sigma_X^2, \sigma_Y^2 \in \mathbb{R}_0^+$ .

We ask (test the hypothesis) whether the population means of both random variables are the same.

Once  $\sigma_X^2 = \sigma_Y^2$  is not assumed, the things get complicated. We have an approximate result only.



If the random variables  $X_1, X_2, ..., X_m \sim \mathcal{N}(\mu_X, \sigma_X^2)$  and  $Y_1, Y_2, ..., Y_n \sim \mathcal{N}(\mu_Y, \sigma_Y^2)$ are independent, then the statistic

$$T = \frac{(\bar{X} - \bar{Y}) - (\mu_X - \mu_Y)}{\sqrt{\frac{s_X^2}{m} + \frac{s_Y^2}{n}}} \sim t_v \qquad approximately$$

where

$$v = \frac{\left(\frac{s_X^2}{m} + \frac{s_Y^2}{n}\right)^2}{\frac{s_X^4}{m^2(m-1)} + \frac{s_Y^4}{n^2(n-1)}}$$



### Exercise:

Use the last Theorem (Satterthwaite's approximation) to formulate a statistical two-sample *t*-test for the difference of the population means with two-sided / one-sided alternative hypothesis (not assuming the same variance).

# $\chi^2$ -test for the variance



•  $\chi^2$ -test for the variance

### Recall the theorem:

If 
$$X_1, X_2, ..., X_n \sim \mathcal{N}(\mu, \sigma^2)$$
 are independent, then  $\frac{(n-1)s^2}{\sigma^2} \sim \chi_{n-1}^2$ 

### Example or motivation:

Assume that  $X \sim \mathcal{N}(\mu, \sigma^2)$  is a random variable following the normal probability distribution. We do not know the population mean  $\mu \in \mathbb{R}$  and we do not know the true value of the population variance  $\sigma^2 \in \mathbb{R}^+$  either. We conjecture / We assume / We ... / that the population variance  $\sigma^2 = \sigma_0^2$ , i.e. the (unknown) population variance  $\sigma^2$  is equal to some prescribed value  $\sigma_0^2 \in \mathbb{R}^+$ .


N.

Example: An archer shoots an arrow against the plane.

The sample space  $\Omega = \mathbb{R}^2 = \{ [x, y] : x, y \in \mathbb{R} \}$  is the set of all the points of the plane. The random variable X is the x-coordinate of the hit, i.e.  $X(\omega) = X([x, y]) = x.$ 

We do not know the archer's mean  $\mu$  and

we do not know the archer's variance  $\sigma^2$ .

We conjecture that the archer's variance could be  $\sigma^2 = \sigma_0^2$ 

where  $\sigma_0^2 \in \mathbb{R}^+$  is some prescribed value.



Let  $x_1 = X(\omega_1)$ ,  $x_2 = X(\omega_2)$ , ...,  $x_n = X(\omega_n)$  be the numerical results

of *n* trials of a random experiment, where  $X \sim \mathcal{N}(\mu, \sigma^2)$ ,

such as the x-coordinates of the archer's n hits.

We do not know the mean  $\mu$  and we do not know the variance  $\sigma^2$ .

We state the null hypothesis (about the variance):

$$H_0: \quad \sigma^2 = \sigma_0^2$$

where  $\sigma_0^2 \in \mathbb{R}^+$  is some number such that we conjecture that the true variance could equal the  $\sigma_0^2$ .

### $\chi^2$ -test for the population variance

#### Having stated the null hypothesis

$$H_0: \quad \sigma^2 = \sigma_0^2$$

#### we also state the alternative hypothesis:

- two-sided:  $H_1: \sigma^2 \neq \sigma_0^2$
- one-sided:  $H_1$ :  $\sigma^2 < \sigma_0^2$
- one-sided:  $H_1$ :  $\sigma^2 > \sigma_0^2$





Under our assumptions  $(x_1, ..., x_n \sim \mathcal{N}(\mu, \sigma^2)$  are independent and  $\sigma^2 = \sigma_0^2$ ), it follows by the Theorem that

$$\frac{(n-1)s^2}{\sigma^2} = \frac{(n-1)s^2}{\sigma_0^2} \sim \chi_{n-1}^2$$

Thus, having the *n* measurements  $x_1, x_2, ..., x_n$ , we calculate the statistic

$$X^2 = \frac{(n-1)s^2}{\sigma_0^2}$$

We know (or assume) that  $X^2 \sim \chi^2_{n-1}$ .

Consider the first case  $(H_1: \sigma^2 \neq \sigma_0^2)$  first. We have:

$$H_0: \quad \sigma^2 = \sigma_0^2$$
$$H_1: \quad \sigma^2 \neq \sigma_0^2$$

Knowing that  $X^2 = \frac{(n-1)s^2}{\sigma_0^2} \sim \chi_{n-1}^2$  and assuming that 0 < c < d is small enough and large enough, respectively, what is the probability that  $c < X^2 < d$ ?

If  $H_0$  is true, then it is quite improbable that  $X^2 \notin (c, d)$ .

Therefore, if we observe that  $X^2 \le c$  or  $d \le X^2$ , then we may conclude that  $H_0$  is probably not true, i.e. we reject the null hypothesis  $H_0$ .





Statistical  $\chi^2$ -test with two-sided alternative hypothesis ( $\sigma^2 \neq \sigma_0^2$ ):

- choose the level of significance, a small number  $\alpha > 0$ , such as  $\alpha = 5$  %, other popular values are  $\alpha = 10$  % or  $\alpha = 1$  % or  $\alpha = 0.1$  % etc.
- find the critical values 0 < c < d so that

$$\int_0^c f(x) \, \mathrm{d}x = \frac{\alpha}{2} \qquad \text{and} \qquad \int_d^{+\infty} f(x) \, \mathrm{d}x = \frac{\alpha}{2}$$

where f is the density of the  $\chi^2$ -distribution with n-1 degrees of freedom

- if  $X^2 \in [0, c] \cup [d, +\infty)$ , the critical region, then <u>reject</u> the null hypothesis
- if  $X^2 \in (c,d)$ , then <u>do not reject</u> (or <u>fail to reject</u>) the null hypothesis

Consider now the second case  $(H_1: \sigma^2 < \sigma_0^2)$ . We have:

$$H_0: \quad \sigma^2 = \sigma_0^2$$
$$H_1: \quad \sigma^2 < \sigma_0^2$$

Knowing that  $X^2 = \frac{(n-1)s^2}{\sigma_0^2} \sim \chi_{n-1}^2$  and assuming that 0 < c is small enough, what is the probability that  $c < X^2$  ?

If  $H_0$  is true, then it is quite improbable that  $X^2 \notin (c, +\infty)$ .

Therefore, if we observe that  $X^2 \le c$ , then we may conclude that  $H_0$  is probably not true, i.e. we reject the null hypothesis  $H_0$ .





Statistical  $\chi^2$ -test with one-sided alternative hypothesis ( $\sigma^2 < \sigma_0^2$ ):

- choose the level of significance, a small number  $\alpha > 0$ , such as  $\alpha = 5$  %, other popular values are  $\alpha = 10$  % or  $\alpha = 1$  % or  $\alpha = 0.1$  % etc.
- find the critical value c > 0 so that

$$\int_0^c f(x)\,\mathrm{d}x = \alpha$$

where f is the density of the  $\chi^2$ -distribution with n-1 degrees of freedom

- if  $X^2 \in [0, c]$ , the critical region, then <u>reject</u> the null hypothesis
- if  $X^2 \in (c, +\infty)$ , then <u>do not reject</u> (or <u>fail to reject</u>) the null hypothesis

Consider finally the third case  $(H_1: \sigma^2 > \sigma_0^2)$ . We have:

$$H_0: \quad \sigma^2 = \sigma_0^2$$
$$H_1: \quad \sigma^2 > \sigma_0^2$$

Knowing that  $X^2 = \frac{(n-1)s^2}{\sigma_0^2} \sim \chi_{n-1}^2$  and assuming that d > 0 is large enough, what is the probability that  $X^2 < d$ ?

If  $H_0$  is true, then it is quite improbable that  $X^2 \notin [0, d)$ .

Therefore, if we observe that  $d \le X^2$ , then we may conclude that  $H_0$  is probably not true, i.e. we reject the null hypothesis  $H_0$ .





Statistical  $\chi^2$ -test with one-sided alternative hypothesis ( $\sigma^2 > \sigma_0^2$ ):

- choose the level of significance, a small number  $\alpha > 0$ , such as  $\alpha = 5$  %, other popular values are  $\alpha = 10$  % or  $\alpha = 1$  % or  $\alpha = 0.1$  % etc.
- find the critical value d > 0 so that

$$\int_{a}^{+\infty} f(x) \, \mathrm{d}x = \alpha$$

where f is the density of the  $\chi^2$ -distribution with n-1 degrees of freedom

- if  $X^2 \in [d, +\infty)$ , the critical region, then <u>reject</u> the null hypothesis
- if  $X^2 \in [0, d)$ , then <u>do not reject</u> (or <u>fail to reject</u>) the null hypothesis

# *z*-test for the population proportion



- One-sample binomial test for the population proportion
- One-sample *z*-test for the population proportion

Motivation: We are tossing a coin repeatedly, and we ask:

ی ls the coin fair?

More generally:

Consider a Bernoulli trial, with the probability of the success being  $p \in (0, 1)$ ,

and with the probability of the <u>failure</u> being q = 1 - p.

#### We do not know the true probability p.

We conjecture / We assume / We ... / that the probability  $p = p_0$ , i.e. the (unknown) probability p is equal to some prescribed value  $p_0 \in (0, 1)$ , e.g., in the case of the coin, conjecture that  $p_0 = 50$  % (meaning the coin is fair).





Let  $X_1, X_2, X_3, ...$  be a sequence of random variables, each of which represents the result of a Bernoulli Trial (independent of the other trials), attaining the value  $X_i = 1$  (success) with a fixed probability  $p \in (0, 1)$  and attaining the value  $X_i = 0$  (failure) with the fixed probability q = 1 - p. (So that p + q = 1and p, q > 0.) We do not know the probability p.

We conjecture / We assume / We ... / that the probability  $p = p_0$ , i.e. the (unknown) probability p is equal to some prescribed value  $p_0 \in (0, 1)$ . Having stated the null hypothesis

 $H_0: \quad p = p_0$ 

#### we also state the alternative hypothesis:

- two-sided:  $H_1: p \neq p_0$
- one-sided:  $H_1: p < p_0$
- one-sided:  $H_1$ :  $p > p_0$







Let  $x_1 = X(\omega_1)$ ,  $x_2 = X(\omega_2)$ , ...,  $x_n = X(\omega_n)$  be the outcomes

of the *n* Bernoulli trials where the probability of success ("1") is *p* and the probability of failure ("0") is q = 1 - p.

For every  $k \in \{0, 1, ..., n\}$ , the probability that  $\sum_{i=1}^{n} x_i = k$  is  $\binom{n}{k} p^k q^{n-k}$ .

Let  $K, L \in \{0, 1, ..., n\}$  be such that K < L.

The probability that  $\sum_{i=1}^{n} x_i \in \{0, 1, \dots, K\}$  is  $\sum_{k=0}^{K} {n \choose k} p^k q^{n-k}$ .

The probability that  $\sum_{i=1}^{n} x_i \in \{L, L+1, ..., n\}$  is  $\sum_{k=L}^{n} {n \choose k} p^k q^{n-k}$ .



Consider the first case  $(H_1: p \neq p_0)$  first. We have:  $H_0: p = p_0$  $H_1: p \neq p_0$ 

- choose the level of significance, a small number  $\alpha > 0$ , such as  $\alpha = 5 \%$
- find the critical values  $K, L \in \{0, 1, ..., n\}$  so that

K is the largest number and L is the least number such that

$$\sum_{k=0}^{K} \binom{n}{k} p_0^k q_0^{n-k} \le \frac{\alpha}{2} \quad \text{and} \quad \sum_{k=L}^{n} \binom{n}{k} p_0^k q_0^{n-k} \le \frac{\alpha}{2}$$

- if  $\sum_{i=1}^{n} x_i \in \{0, \dots, K\} \cup \{L, \dots, n\}$ , the critical region, then <u>reject</u> the null hyp.
- if  $\sum_{i=1}^{n} x_i \in \{K + 1, ..., L 1\}$ , then <u>do not reject</u> (or <u>fail to reject</u>) the null hyp.



Consider now the second case  $(H_1: p < p_0)$ . We have:  $H_0: p = p_0$  $H_1: p < p_0$ 

- choose the level of significance, a small number  $\alpha > 0$ , such as  $\alpha = 5 \%$
- find the critical value  $K \in \{0, 1, ..., n\}$  so that K is the largest number such that

$$\sum_{k=0}^{K} \binom{n}{k} p_0^k q_0^{n-k} \leq \alpha$$

- if  $\sum_{i=1}^{n} x_i \in \{0, ..., K\}$ , the critical region, then <u>reject</u> the null hypothesis
- if  $\sum_{i=1}^{n} x_i \in \{K + 1, ..., n\}$ , then <u>do not reject</u> (or <u>fail to reject</u>) the null hypothesis



Consider finally the third case  $(H_1: p > p_0)$ . We have:  $H_0: p = p_0$  $H_1: p > p_0$ 

- choose the level of significance, a small number  $\alpha > 0$ , such as  $\alpha = 5 \%$
- find the critical value  $L \in \{0, 1, ..., n\}$  so that L is the least number such that

$$\sum_{k=L}^n \binom{n}{k} p_0^k q_0^{n-k} \le \alpha$$

- if  $\sum_{i=1}^{n} x_i \in \{L, ..., n\}$ , the critical region, then <u>reject</u> the null hypothesis
- if  $\sum_{i=1}^{n} x_i \in \{0, ..., L-1\}$ , then <u>do not reject</u> (or <u>fail to reject</u>) the null hypothesis

where A,



It is inconvenient to calculate the sums  $\sum_{k=0}^{K} {n \choose k} p_0^k q_0^{n-k}$  and  $\sum_{k=L}^{n} {n \choose k} p_0^k q_0^{n-k}$  if *n* is large. It is more convenient then to approximate the sums by using the de Moivre-Laplace Central Limit Theorem:

Let  $p \in (0,1)$  be the probability of success in a Bernoulli Trial, and let q = 1 - pbe the probability of failure in the trial. Whenever  $-\infty \le a < b \le +\infty$ , it then holds

$$\frac{\sum_{k=A_n}^{B_n} {n \choose k} p^k q^{n-k} - np}{\sqrt{npq}} \to \underbrace{\int_a^b \frac{1}{\sqrt{2\pi}} e^{-\frac{t^2}{2}} dt}_{\Phi(b) - \Phi(a)} \quad \text{as} \quad n \to \infty$$
$$= \left[ np + a\sqrt{npq} \right] \ge 0 \quad \text{and} \quad B_n = \left[ np + b\sqrt{npq} \right] \le n \quad \text{if} \quad n \ge \max\left(\frac{q}{n}a^2, \frac{p}{n}b^2\right)$$

<u>De Moivre-Laplace Central Limit Theorem (reformulated):</u> Let  $X \sim Bi(n,p)$  with  $p \in (0,1)$ , and put q = 1 - p for short. Whenever  $-\infty \le a < b \le +\infty$ , it then holds

$$P\left(a < \frac{X - np}{\sqrt{npq}} < b\right) \to \underbrace{\int_{a}^{b} \frac{1}{\sqrt{2\pi}} e^{-\frac{t^{2}}{2}} dt}_{\Phi(b) - \Phi(a)} \quad \text{as} \quad n \to \infty$$

and the convergence is uniform with respect to a and b.





Consider the first case  $(H_1: p = p_0)$  first. We have:  $H_0: p = p_0$  $H_1: p \neq p_0$ 

- choose the level of significance, a small number  $\alpha > 0$ , such as  $\alpha = 5 \%$
- find c > 0 so that

$$\int_{-\infty}^{-c} \frac{1}{\sqrt{2\pi}} e^{-\frac{t^2}{2}} dt = \frac{\alpha}{2} \quad \text{and} \quad \int_{+c}^{+\infty} \frac{1}{\sqrt{2\pi}} e^{-\frac{t^2}{2}} dt = \frac{\alpha}{2}$$

- if  $\sum_{i=1}^{n} x_i \le np_0 c\sqrt{np_0q_0}$  or  $np_0 + c\sqrt{np_0q_0} \le \sum_{i=1}^{n} x_i$ , the critical region, then <u>reject</u> the null hypothesis
- if  $np_0 c\sqrt{np_0q_0} < \sum_{i=1}^n x_i < np_0 + c\sqrt{np_0q_0}$ ,

then do not reject (or fail to reject) the null hypothesis



Consider now the second case  $(H_1: p < p_0)$ . We have:  $H_0: p = p_0$  $H_1: p < p_0$ 

- choose the level of significance, a small number  $\alpha > 0$ , such as  $\alpha = 5 \%$
- find c > 0 so that

$$\int_{-\infty}^{-c} \frac{1}{\sqrt{2\pi}} e^{-\frac{t^2}{2}} dt = \alpha$$

- if  $\sum_{i=1}^{n} x_i \le np_0 c\sqrt{np_0q_0}$ , the critical region, then <u>reject</u> the null hypothesis
- if  $np_0 c\sqrt{np_0q_0} < \sum_{i=1}^n x_i$ , then <u>do not reject</u> (or <u>fail to reject</u>) the null hyp.



Consider finally the third case  $(H_1: p > p_0)$ . We have:  $H_0: p = p_0$  $H_1: p > p_0$ 

- choose the level of significance, a small number  $\alpha > 0$ , such as  $\alpha = 5 \%$
- find c > 0 so that

$$\int_{+c}^{+\infty} \frac{1}{\sqrt{2\pi}} e^{-\frac{t^2}{2}} dt = \alpha$$

- if  $np_0 + c\sqrt{np_0q_0} \le \sum_{i=1}^n x_i$ , the critical region, then <u>reject</u> the null hypothesis
- if  $\sum_{i=1}^{n} x_i < np_0 + c\sqrt{np_0q_0}$ , then <u>do not reject</u> (or <u>fail to reject</u>) the null hyp.

## *p*-value of the test



The general outline

of a statistical hypothesis test

• The *p*-value of a test



- A statistical test consists in the study of the outcomes of a random experiment.
- We put down a hypothesis about the probability distribution of the outcomes of the random experiment.
- We also make up a statistic S a formula, i.e. a mathematical expression and we prove (!) as a mathematical Theorem that, under our hypotheses, the statistic S follows a certain probability distribution.
- We carry out the random experiment several (or many) times.
- We put down the results of the experiment, i.e., count positive results, count the negative results, and so on.

- We substitute the results (the counts, and so on) into the mathematical expression-statistic S, which is a random variable thus (its value depends on the results of the random experiment).
- We then choose the significance level  $\alpha$  a small probability such as  $\alpha = 5 \% = 0.05$  (i.e. "one error per twenty trials").

(Other popular choices include  $\alpha = 10 \% = 0.1$  or  $\alpha = 1 \% = 0.01$ .)

By using the mathematical Theorem, which we proved (see above),
 we find the critical region C ⊆ ℝ so that – if our hypotheses are true –
 then the probability of the event that S ∈ C is ≤ α.



- The critical region *C* is usually a closed interval or the union of two closed intervals.
- Finally, make a statistical conclusion:
- If  $S \in C$ , then **reject** the hypothesis.
- If the hypotheses are true, then it is quite improbable that *S* ∈ *C*; the probability is ≤ α. So we are making a mistake type I error,
  i.e. rejecting a hypothesis which is true about once per twenty trials, if α = 5 %.



- If  $S \notin C$ , then do not reject (or fail to reject) the hypothesis.
- The fact that we fail to reject the hypothesis is <u>not</u> a confirmation that the hypothesis is true!
- Since the statistic S is a random variable, it may happen by chance that  $S \notin C$  even if the hypothesis is false.
- This situation failing to reject a false hypothesis is a type II error.
   The probability of type II error is β, and this probability is difficult to calculate...
   If α = 5 %, then the probability β should be ≤ 20 %. (Is it ≤ 20 %?)
   The probability 1 β is the **power of the test**.

The above outline of the test is as follows:

- Choose the significance level  $\alpha$  (such as  $\alpha = 5$  %).
- Depending upon the  $\alpha$ , find the critical region  $C_{\alpha} \subseteq \mathbb{R}$  so that if the hypothesis is true – then the probability that  $(S \in C_{\alpha})$  is  $\leq \alpha$ .
- Carry out the experiment, enumerate the expression S, and see if  $S \in C_{\alpha}$ .

#### Another procedure:

- Carry out the experiment and enumerate the expression S.
- Find the least number  $p \in (0, 1)$  such that  $S \in C_p$ .
- This value p is the *p***-value of the test**.



