Statistics

Lecture 9

Hypothesis testing: Parametric tests



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Outline of the lecture



- z-tests for the means
 - type I and type II error
 - significance level α
 - power of the test 1β
- *t*-tests for the means
- χ^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 $\frac{\overline{X} - \mu}{\sigma/\sqrt{n}} \sim \mathcal{N}(0, 1)$

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

Confidence interval:

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

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

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 $\frac{\overline{X} \mu}{\sigma/\sqrt{n}} \sim \mathcal{N}(0, 1)$
- 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 $\frac{(n-1)s^2}{\sigma^2} \sim \chi_{n-1}^2$

• ...

to establish some <u>parametric</u> statistical <u>tests about the parameters</u> (μ , σ^2 , ...) of the respective probability distribution.





- 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 $\frac{\bar{X} - \mu}{\sigma/\sqrt{n}} \sim \mathcal{N}(0, 1)$

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 σ^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 μ 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.

$$X(\omega) = X([x,y]) = x.$$



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

i.e. we do not know the archer's mean μ .

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 σ^2 , but we do not know the mean μ .

We state the **null hypothesis** (<u>about the mean</u>):

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

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

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

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 $(\mu \neq \mu_0 \text{ or } \mu < \mu_0 \text{ or } \mu > \mu_0)$ do we choose?

→ 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 H₁: μ ≠ μ₀.
- 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 H₁: μ < μ₀.
- 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 H₁: μ > μ₀.



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

• μ_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:

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

$$H_1$$
: $\mu \neq \mu_0$

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

$$Z \leq -c$$
 or $+c \leq Z$

then we may conclude that H_0 is probably not true,

i.e. we <u>reject the null hypothesis</u> 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 <u>reject</u> the null hypothesis
- if $Z \in (-c, +c)$, then **do not reject** (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 α , usually $\alpha = 5$ %.

Recall: 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 α .



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 β .

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

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



To calculate the probability β 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 $\frac{\bar{x}-\mu}{\sigma/\sqrt{n}} = \frac{x-\mu_0 \div (\mu_0-\mu)}{\sigma/\sqrt{n}} \sim \mathcal{N}(0,1)$.

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

$$P\left(-c < \frac{\bar{x} - \mu_0}{\sigma/\sqrt{n}} < +c\right) = P\left(-c < \frac{\bar{x} - \mu + (\mu_0 - \mu)}{\sigma/\sqrt{n}} < +c\right) =$$

$$= P\left(\frac{\mu - \mu_0}{\sigma/\sqrt{n}} - c < \frac{\bar{x} - \mu}{\sigma/\sqrt{n}} < \frac{\mu - \mu_0}{\sigma/\sqrt{n}} + c\right) =$$

$$= \beta = \int_{(\mu - \mu_0)/(\sigma/\sqrt{n}) - c}^{(\mu - \mu_0)/(\sigma/\sqrt{n}) - c} \varphi(x) dx$$



Notice that, if the true μ is close to the hypothesized μ_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 β should be ≤ 20 %.

Therefore, if we wish to have $\beta \approx 20 \%$ or $\beta \leq 20 \%$, then we must not consider the true μ close to the hypothesized μ_0 .



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

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

$$H_1$$
: $\mu < \mu_0$

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 **do not reject** (or <u>fail to reject</u>) the null hypothesis



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

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

$$H_1$$
: $\mu > \mu_0$

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 $+c \le Z$, 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 do not reject (or fail to reject) 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 μ^{before} and μ^{after} , but we do assume that we know the variances σ_X^2 and σ_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 μ^{before} and μ^{after} . We only test the hypothesis by having done a sample of n pairs of measurements. Formulate the alternative hypothesis:

• two-sided: H_1 : $\mu^{\text{before}} \neq \mu^{\text{afer}}$ (the treatment has some effect)

• one-sided: H_1 : $\mu^{\text{before}} < \mu^{\text{after}}$ (the treatment increases / ...

• one-sided: H_1 : $\mu^{\mathrm{before}} > \mu^{\mathrm{after}}$... / <u>decreases</u> the quantity)



Recall the theorem:

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

Notice also:

If $X_1, X_2, ..., X_n \sim \mathcal{N}\left(\mu^{\text{before}}, \sigma_X^2\right)$ and $Y_1, Y_2, ..., Y_n \sim \mathcal{N}\left(\mu^{\text{after}}, \sigma_Y^2\right)$ 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 $\mu^{\rm before} = \mu^{\rm after}$ 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.



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

• $\mu_0 = \text{const.}$ for a general difference of the means $(\mu^{\text{before}} = \mu^{\text{after}} + \text{const.})$



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

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

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

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 α 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 **do not reject** (or <u>fail to reject</u>) the null hypothesis



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

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

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

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

$$-c < Z$$

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 α 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 **do not reject** (or <u>fail to reject</u>) the null hypothesis



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

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

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

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 α 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 do not reject (or fail to reject) 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 σ_X^2 and σ_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 <u>null hypothesis</u>:

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

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

Recall that we do not know the true population means μ_X and μ_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

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

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

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

equivalently

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

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 first sample

•
$$\bar{y} = \sum_{j=1}^{n} y_j/n$$
 is the sample mean of the second 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:

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

$$H_1$$
: $\mu_X \neq \mu_Y$

Knowing that

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

we 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 α 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 **do not reject** (or <u>fail to reject</u>) the null hypothesis



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

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

$$H_1$$
: $\mu_X < \mu_Y$

Knowing that

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

we 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 α 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 **do not reject** (or <u>fail to reject</u>) the null hypothesis



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

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

$$H_1$$
: $\mu_X > \mu_Y$

Knowing that

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

we 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 α 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 **do not reject** (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 $\frac{\bar{X} - \mu}{\sigma/\sqrt{n}} \sim \mathcal{N}(0, 1)$

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 σ^2 of the random variable X in practice.

That is why we recall another theorem now.



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

• $s^2 = \sum_{i=1}^n (X_i - \bar{X})^2/(n-1)$ is the sample variance of the random variables

• χ^2_{n-1} is Pearson's χ^2 -distribution with n-1 degrees of freedom



Recall the corollary of the two theorems:

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

•
$$\bar{X} = \sum_{i=1}^n X_i/n$$

• $s = \sqrt{\sum_{i=1}^{n} (X_i - \bar{X})^2 / (n-1)}$

is the sample mean of the random variables

is the sample standard deviation of the random variables

• t_{n-1}

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 $\frac{\bar{X} - \mu}{s/\sqrt{n}} \sim t_{n-1}$

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> σ^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 μ 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 σ^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 μ .

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 σ^2 and we do not know the mean μ .

We state the null hypothesis (about the mean):

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

where $\mu_0 \in \mathbb{R}$ is some number such that we conjecture that the true mean could equal the μ_0 .



Having stated the **null hypothesis**

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

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 $(\mu \neq \mu_0 \text{ or } \mu < \mu_0 \text{ or } \mu > \mu_0)$ do we choose?

→ 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 H₁: μ ≠ μ₀.
- 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 H₁: μ < μ₀.
- 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 H₁: μ > μ₀.



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.

The gamma function



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

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:

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

$$H_1$$
: $\mu \neq \mu_0$

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 <u>reject the null hypothesis</u> 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

$$\int_{-\infty}^{-c} f(x) dx = \frac{1}{2}\alpha \quad \text{and} \quad \int_{+c}^{+\infty} f(x) dx = \frac{1}{2}\alpha$$

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 **do not reject** (or <u>fail to reject</u>) the null hypothesis

Type I and Type II error



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

- the null hypothesis (H_0) is actually true \mathbb{R} we do not reject it \mathbb{R} 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 probability of the type I error is the significance level a

The probability of the type II error is β

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



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

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

$$H_1$$
: $\mu < \mu_0$

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 **do not reject** (or <u>fail to reject</u>) the null hypothesis

One-sample t-test for the population mean



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

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

$$H_1$$
: $\mu > \mu_0$

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 .

One-sample t-test for the population mean



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 **do not reject** (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 μ^{before} , $\mu^{\text{after}} \in \mathbb{R}$ and for some σ_X^2 , $\sigma_Y^2 \in \mathbb{R}_0^+$. We do not know the true values of the population means μ^{before} and μ^{after} , and we do not know the true values of the variances σ_X^2 and σ_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 μ^{before} and μ^{after} . We only test the hypothesis by having done a sample of n pairs of measurements. Formulate the alternative hypothesis:

• two-sided: H_1 : $\mu^{\text{before}} \neq \mu^{\text{afer}}$ (the treatment has some effect)

• one-sided: H_1 : $\mu^{\text{before}} < \mu^{\text{after}}$ (the treatment increases / ...

• one-sided: H_1 : $\mu^{\mathrm{before}} > \mu^{\mathrm{after}}$... / <u>decreases</u> the quantity)



Recall the theorem:

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

Notice also:

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

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

where

- $\bar{x} = (\sum_{i=1}^{n} x_i)/n$ is the sample mean of the measurements before 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:

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

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

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 α 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

$$\int_{-\infty}^{-c} f(x) dx = \frac{1}{2}\alpha \quad \text{and} \quad \int_{+c}^{+\infty} f(x) dx = \frac{1}{2}\alpha$$

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 **do not reject** (or <u>fail to reject</u>) the null hypothesis



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

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

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

Knowing that

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

we fail to reject H_0 iff

$$-c < T$$

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 α 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 **do not reject** (or <u>fail to reject</u>) the null hypothesis



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

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

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

Knowing that

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

we fail to reject H_0 iff

$$T < +c$$

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 α 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 **do not reject** (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 μ_X , μ_Y nor the variances σ_X^2 , σ_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 <u>null hypothesis</u>:

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

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

Recall that we do not know the true population means μ_X and μ_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

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

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

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

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:

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



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

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

Therefore:

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



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.



Therefore:

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

where

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

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>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:

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

$$H_1$$
: $\mu_X \neq \mu_Y$

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 α 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

$$\int_{-\infty}^{-c} f(x) dx = \frac{1}{2}\alpha \quad \text{and} \quad \int_{+c}^{+\infty} f(x) dx = \frac{1}{2}\alpha$$

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 **do not reject** (or <u>fail to reject</u>) the null hypothesis



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

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

$$H_1$$
: $\mu_X < \mu_Y$

Knowing that

we fail to reject H_0 iff

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

$$-c < T$$

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



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

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

$$H_1$$
: $\mu_X > \mu_Y$

Knowing that

we fail to reject H_0 iff

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

$$T < +c$$

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 α 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 **do not reject** (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.

Theorem (Satterthwaite's approximation):



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).

χ^2 -test for the variance



• χ^2 -test for the variance

χ^2 -test for the population 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 σ^2 is equal to some prescribed value $\sigma_0^2 \in \mathbb{R}^+$.

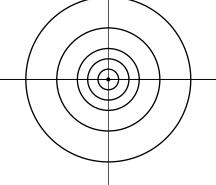


Example: An archer shoots an arrow against the plane.

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 mean μ and we do not know the archer's variance σ^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 μ and we do not know the variance σ^2 .

We state the **null hypothesis** (<u>about the variance</u>):

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

where $\sigma_0^2 \in \mathbb{R}^+$ is some number such that we conjecture that the true variance could equal the σ_0^2 .



Having stated the **null hypothesis**

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

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 $X^2 \sim \chi_{n-1}^2$.



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

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

$$H_1$$
: $\sigma^2 \neq \sigma_0^2$

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 χ^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 χ^2 -distribution with n-1 degrees of freedom

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



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

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

$$H_1$$
: $\sigma^2 < \sigma_0^2$

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 χ^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 χ^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:

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

$$H_1$$
: $\sigma^2 > \sigma_0^2$

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 $d \leq X^2$, then we may conclude that

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



Statistical χ^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

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

where f is the density of the χ^2 -distribution with n-1 degrees of freedom

- if $X^2 \in [d, +\infty)$, the critical region, then reject the null hypothesis
- if $X^2 \in [0,d)$, then **do not reject** (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 Is the coin fair?

More generally:

Consider a Bernoulli trial, with the probability of the <u>success</u> 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 = 1 and 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**

$$H_0$$
: $p=p_0$

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

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

- 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 **do not reject** (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 do not reject (or fail to reject) the null hypothesis



It is inconvenient to calculate the sums $\sum_{k=0}^{K} \binom{n}{k} p_0^k q_0^{n-k}$ and $\sum_{k=L}^{n} \binom{n}{k} p_0^k q_0^{n-k}$ 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 - p be the probability of failure in the trial. Whenever $-\infty \le a < b \le +\infty$, it then holds

$$\frac{\sum_{k=A_n}^{B_n} \binom{n}{k} p^k q^{n-k} - np}{\sqrt{npq}} \to \underbrace{\int_a^b \frac{1}{\sqrt{2\pi}} e^{-\frac{t^2}{2}} dt}_{\Phi(b)-\Phi(a)} \quad \text{as} \quad n \to \infty$$

where $A_n = \left\lceil np + a\sqrt{npq} \right\rceil \ge 0$ and $B_n = \left\lceil np + b\sqrt{npq} \right\rceil \le n$ if $n \ge \max\left(\frac{q}{p}a^2, \frac{p}{q}b^2\right)$.



De Moivre-Laplace Central Limit Theorem (reformulated):

Let $X \sim \text{Bi}(n, p)$ 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 <u>fail to reject</u>) 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 **do not reject** (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 **do not reject** (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 α a small probability such as α = 5 % = 0.05 (i.e. "one error per twenty trials").
 (Other popular choices include α = 10 % = 0.1 or α = 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, <u>make a statistical conclusion</u>:
- If S∈ 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 ∉ 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 ∉ 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 *p*-value of the test



The above outline of the test is as follows:

- Choose the significance level α (such as $\alpha = 5$ %).
- Depending upon the α, find the critical region C_α ⊆ ℝ so that –
 if the hypothesis is true then the probability that (S ∈ C_α) is ≤ α.
- 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.