Introduction to biostatistics

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Introduction to (bio)statistics

What statistics are (not) all about



"Data don't make any sense, we will have to resort to statistics."



Introduction to statistics

Branches of statistics

Descriptive statistics

 Describe, summarize, order or graphically represent empirical data

Exploratory data analysis

• Identify patterns or structures in the data

Inferential statistics

 Predict, estimate and generalize about populations based on data derived from samples



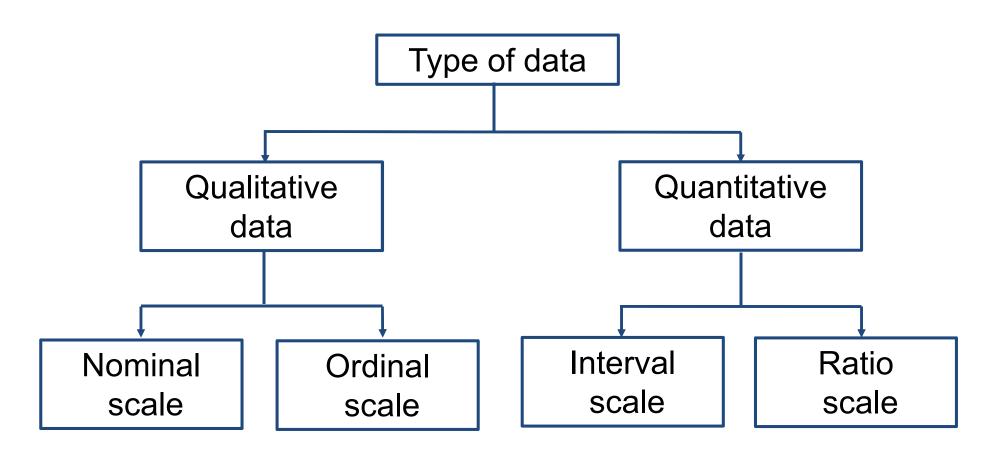




Introduction to descriptive statistics

- Descriptive statistics provide the basis of quantitative data analysis
- The focus is to describe, summarize and graphically represent the data as it is without making any assumptions or generalizations
- Descriptive statistics provide measures to:
 - Describe the central tendency of data
 - Mean, median, modus
 - Describe the dispersion of the data
 - Variance, standard deviation, range
 - Describe how different data are related to each other
 - Correlation coefficient







- Nominal scale
 - The lowest level of measurement
 - Data belong to mutually exclusive categories
 - The only possible comparison between elements is "equal to" or "not equal to"
 - Examples include:
 - Blood type
 - Gender
 - Color of your eyes
 - Nucleotides in the DNA sequence
 - University field of studies



- Ordinal scale
 - Data belong to mutually exclusive categories which can be ordered (<,> comparisons allowed)
 - Differences between categories are not allowed
 - Examples include:
 - Exam grades (A, B, C, D)
 - Questionnaire options ("strongly agree", "agree", disagree", strongly disagree")
 - Levels of happiness, satisfaction, etc.
 - Some clinical scores



- Interval scale
 - A metric scale which allows building differences between values but not ratios
 - Examples include:
 - Celsius temperature
 - IQ score
 - Time on a clock
 - No "true" zero value is defined on an interval scale
 - 0 degrees Celsius does not mean there is no temperature
 - 0:00 does not mean time does not exist



- Ratio scale
 - A metric scale which allows building differences and ratios between values
 - A value of zero indicates non-existence
 - Examples include:
 - Age
 - Height
 - Weight
 - Number of children
 - Distance
 - Blood pressure
- Note: higher levels of measurements can be reduced to lower scales but not the other way around



Measures of central tendency

- Central tendency, center or location of the data is a central or typical value
- Mode
 - The element if the data which appears most often
 - The mode can be determined for variables measured on any scale

Out of 40 participants in a statistics course, 10 are studying bioinformatics, 25 are studying biomedicine and 5 are studying biochemistry. What is the mode for the the variable field of studies of the course takers?



Measures of central tendency

- Median
 - The "middle" value in an ordered data set, half of the data are smaller and half of the data are larger than the median
 - Can be calculated for data measured at least on an ordinal scale
- Arithmetic mean
 - The arithmetic mean only makes sense for variables on a metric scale

$$\bar{x} = \frac{1}{n} \sum_{i=1}^{n} x_i$$



Measures of central tendency

- Mean or median?
- Consider a small sample containing the following values {3.5; 4; 2.5; 6; 7; 5.5}

$$\bar{x} = \frac{3.5 + 4 + 2.5 + 6 + 7 + 5.5}{6} = 4.75$$

$$Median = 4.75$$

• Consider adding the value 500 to the set {3.5; 4; 2.5; 6; 7; 5.5; 500}

$$\bar{x} = \frac{3.5 + 4 + 2.5 + 6 + 7 + 5.5 + 500}{7} = 75.5$$

$$Median = 5.5 - a robust statistic$$



Measures of dispersion

• Empirical variance

$$s^{2} = \frac{1}{n} \sum_{i=1}^{n} (x_{i} - \bar{x})^{2}$$

• Sample variance

$$s^{2} = \frac{1}{n-1} \sum_{i=1}^{n} (x_{i} - \bar{x})^{2}$$

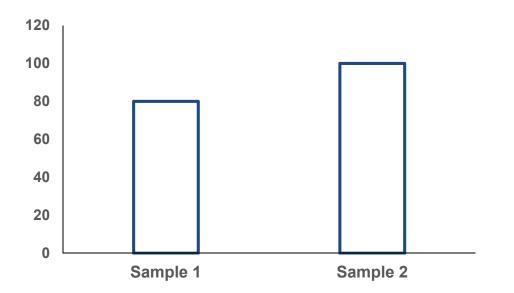
• Standard deviation

$$s = \sqrt{s^2}$$



Graphical representation of location and dispersion

 Suppose that blood pressure was measured in two samples of 30 patients each. Mean diastolic blood pressure for the first sample turned out to be 80, whereas mean pressure in the second sample was 100 mmHg.

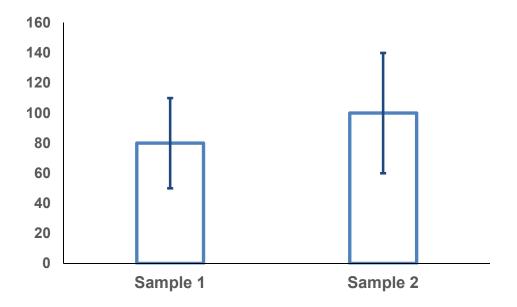


• Do peaple in the second sample have higher diastolic blood pressure?



Graphical representation of location and dispersion

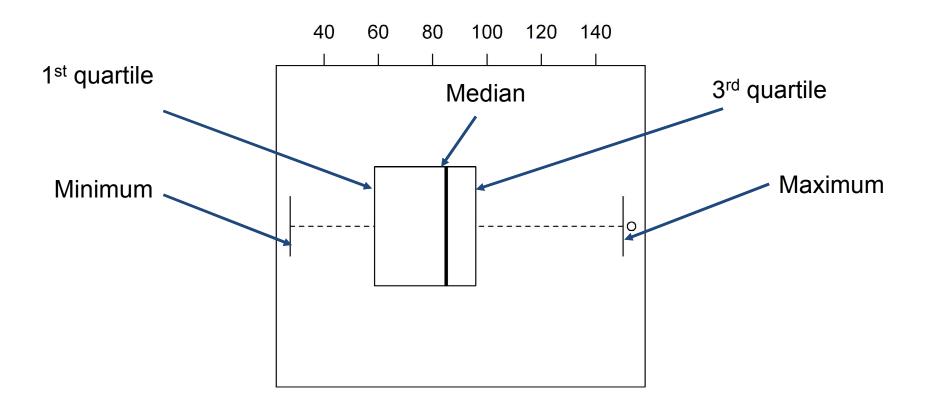
• Suppose that blood pressure was measured in two samples of 30 patients each. Mean diastolic blood pressure for the first sample turned out to be 80 ± 30 , whereas mean pressure in the second sample was 100 ± 40 mmHg.





Boxplot

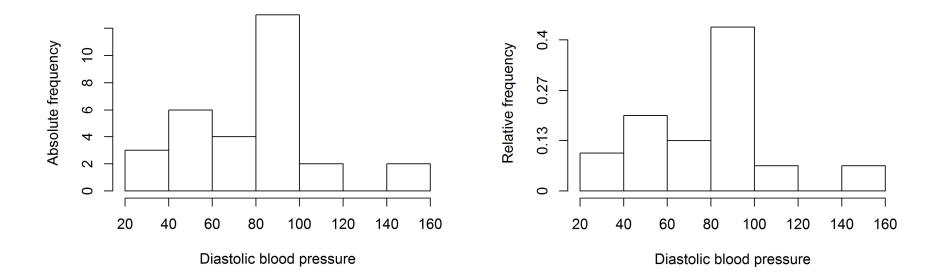
• Consider the first sample of 30 patients with mean diastolic blood pressure of 80, with a standard deviation of 30





Histograms

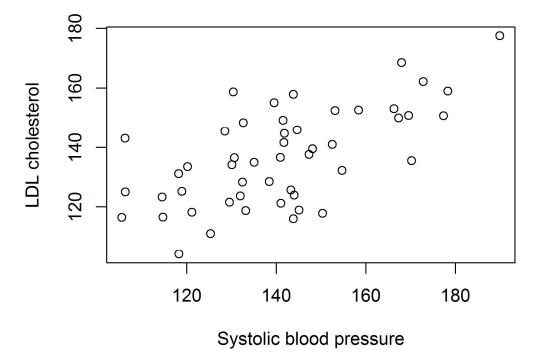
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Association between metric variables

- In a study of 50 patients, systolic blood pressure and LDL cholesterol were measured. Mean blood pressure was 137.5 mmHg with a standard deviation of 18.27. Average LDL cholesterol was 134 mg/dL with a standard deviation of 14.14.
 - Is higher blood pressure associated with higher LDL cholesterol levels?





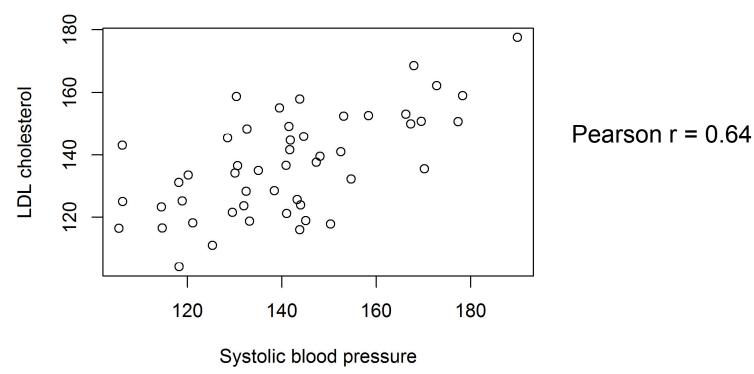
Association between metric variables

- Pearson correlation coefficient
 - Measures the linear relationship between metric variables
 - Values range from -1 to 1
 - -1 perfect negative correlation
 - 0 no correlation
 - +1 perfect positive correlation



Association between metric variables

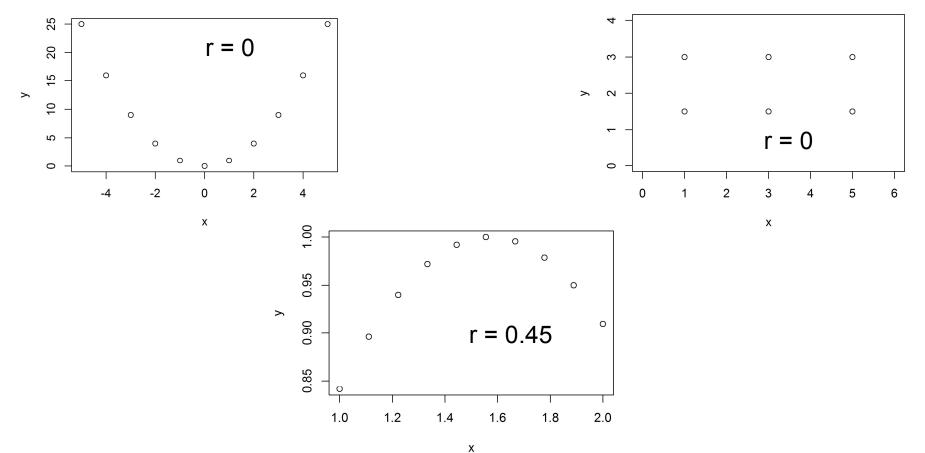
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Pearson correlation

• Pearson correlation coefficient is only appropriate if the relationship between the variables is linear



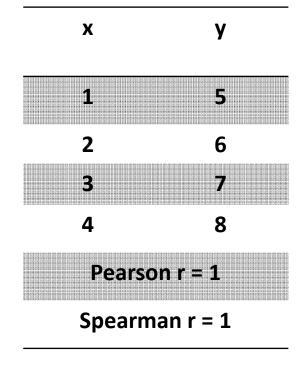
Spearman correlation

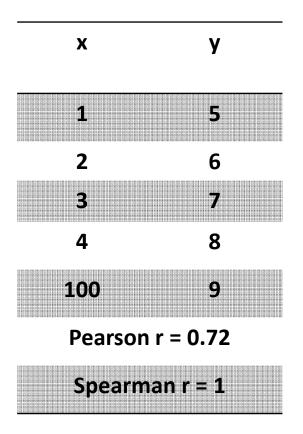
- The Spearman correlation coefficient is an alternative to the Pearson correlation coefficient appropriate when:
 - The relationship between the variables is monotonically increasing or decreasing but not linear
 - Data are measured on an ordinal scale
- Values range from -1 to 1, 0 corresponds to no correlation
- Data are arranged in an increasing order and every data point is assigned a rank
- Spearman correlation belongs to the category of robust statistics because it is robust in the presence of outliers



Pearson vs. Spearman correlation

• Influence of outliers







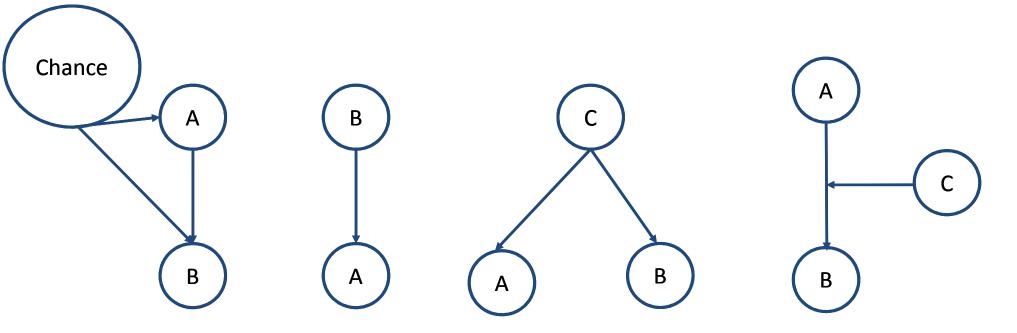
Correlation versus causality

- Causality is often the focus of research
 - We try to explain why phenomena occur and what causes them
- Caution: statistics only provides measures of association. The question of what is cause and what is consequence remains open.
- Causality, therefore, requires careful theoretical consideration and appropriate experimental designs
 - Causality can often be masked by random confounding factors
 or latent factors



Correlation versus causality

- Consider two variables A and B which are highly correlated with each other.
- Different causal relationships are possible







Random processes

- Random process a phenomenon with an uncertain outcome. Examples include:
 - A coin toss or throwing a dice
 - The gender of an unborn child
 - Result from an exam
 - A scientific experiment
- The set of all possible outcomes of a random process is called a sample space Ω.
 - E.g. The colour of a gummy bear we take out of a bag with blue, red, yellow and green gummy bears is a random experiment with the following sample space:

```
\Omega = \{, blue", , red", , yellow", , green"\}
```



Radom events

- The outcome of a **random experiment** is called a **random event**
 - **Simple event** an event which contains only a single outcome
 - E.g. A green gummy bear
 - An event is a unification of simple events

E.g. A green gummy bear or a yellow gummy bear;

Not a blue gummy bear

- Although the single outcome of a random process is uncertain, outcomes follow certain distributions if the random experiment is repeated many times
- The chance of a random outcome occurring is described by the probability



Frequency versus probability

- Consider the following random experiment:
 - A pot contains 50 white and 50 black balls. What is the probability of drawing a white ball if we draw if replacement (we put the ball back in the pot after drawing it)
 - An R simulation was performed investigating relative frequency of white balls after repeating the experiment 10 000 times by drawing 1 to 10 000 balls.



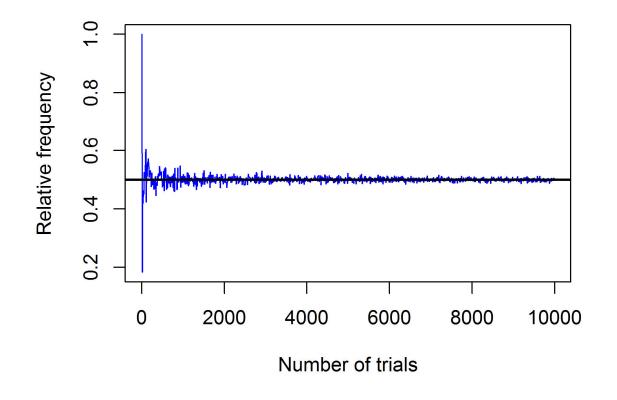
Frequency versus probability

Number of draws	Number of white balls	Relative frequency of white balls
1	1	1
11	2	0.182
101	61	0.604
501	238	0.475
1001	494	0.494
9991	4985	0.499

• The relative frequency approaches 0.5 as the number of trials increases



Frequency versus probability



• The **relative frequency** converges against a limit value as the number of trials **n** increases. The limit is called **probability**



Laplace probability

- In a Laplace experiment all simple events have the same probability
- Let P(A) describe the probability that event A occurs. The Laplace probability is then:

 $P(A) = \frac{Number of elements in A}{Number of all elements}$

E.g. 50 students are taking part in a statistics course, 15 of them are studying bioinformatics, 25 are studying biomedicine and 10 are studying biophysics. What is the probability that a randomly selected student is studying biomedicine?

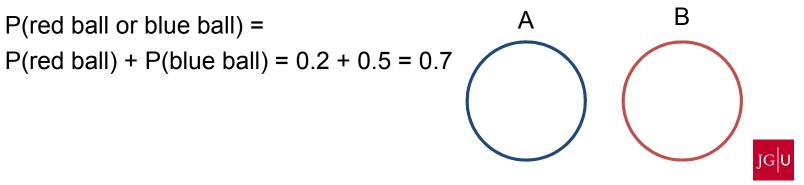
$$P = \frac{25}{50} = 0.5$$



Rules for probabilities

- The probability must satisfy the following conditions:
 - 1. $P(A) \ge 0$, the probability is always positive
 - P(Ω) = 1, every random experiment must have an outcome, therefore the sum of the probabilities of all possible outcomes is 1
 - 3. P(A or B) = P(A) + P(B) if A and B are **disjoint**. Two events are disjoint if they cannot occur together

E.g. There are red, blue and yellow balls in a pot and the probabilities of drawing a red, blue or yellow balls are 0.2, 0.5 and 0.3, respectively. What is the probability of drawing a red or a blue ball?

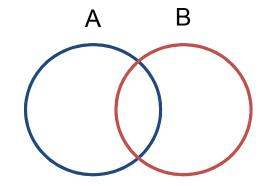


Rules for probabilities

• P(A or B) = P(A) + P(B) - P(A and B) if A and B are not disjoint

E.g. The probability of suffering from hypertension over the age of 65 is 60% and the probability of having a heart arrhythmia is 10 %. 30% of the population over 65 has both hypertension and a heart arrhythmia. What is the probability of having at least one of these conditions over the age of 65?

$$P(H \text{ or } A) = P(H) + P(A) - P(H \text{ and } A)$$
$$P(H \text{ or } A) = 0.6 + 0.1 - 0.3 = 0.4$$





Independent events

- Two events are stochastically independent if the occurrence of one of the events does not influence the probability of the other event occurring
 - Flipping a coin multiple times
 - Throwing a dice multiple times
 - Drawing balls from a pot with replacement
- The probability that two independent events A and B occur simultaneously is the product of the respective probabilities for A and B:

P(A and B) = P(A)P(B)

E.g. the probability two get first tail and then head after flipping a coin twice is:

P(first tail and then head) = P(tail)P(head) = 0.5 * 0.5 = 0.25

 What is the probability of getting head and tail after flipping a coin twice?

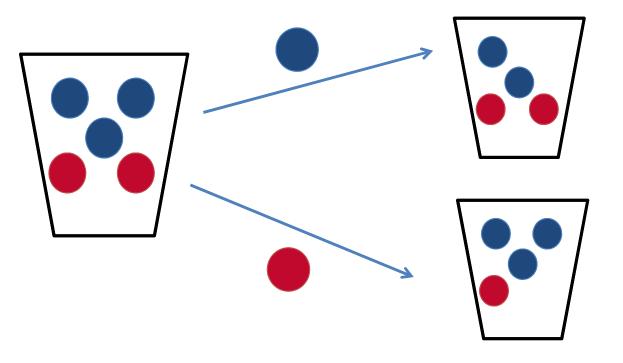


Dependent events

• Two events are **dependent** if the occurrence of one of the events influences the probability of the second event.

For example, suppose we are drawing balls from a pot without replacement. The pot contains 3 blue and 2 red balls. What is the probability that the second ball we draw is red?

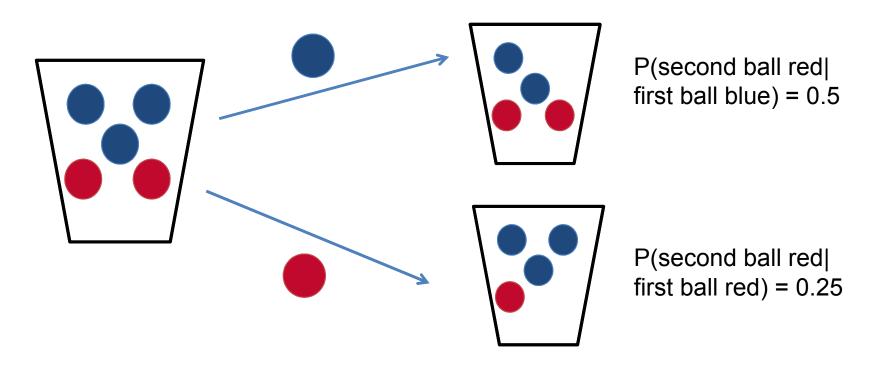
> Depends on the color of the first ball we draw.





Conditional probability

 The conditional probability describes the probability of an event **A** when we know that an event **B** has occurred *P*(*A*|*B*)





Conditional probability

• The probability that two **dependent events A** and **B** occur simultaneously:

P(A and B) = P(A|B)P(B) = P(B|A)P(A)

- The temporal sequence in which events occur is irrelevant
- The conditional probability is therefore:

$$P(A|B) = \frac{P(A \text{ and } B)}{P(B)} = \frac{P(B|A)P(A)}{P(B)}$$



Conditional probability

Example: A test T used to detect a specific disease D is positive in 99% of the cases when a patients truly suffers from the disease. Altogether, the test is positive in 3 % of tested patients and the prevalence of the disease in the investigated population is 1 %. What is the probability that a patient truly has the disease when the test result is positive?

- P(T) probability that the test is positive
- P(D) prevalence of the disease

P(T|D) – probability that the test is positive when a patient truly has the disease

P(D|T) – probability that the patient truly suffers from the disease knowing that the test result is positive



Conditional probability

Example: A test T used to detect a specific disease D is positive in 99% of the cases when a patients truly suffers from the disease. Altogether, the test is positive in 3 % of tested patients and the prevalence of the disease in the investigated population is 1 %. What is the probability that a patient truly has the disease when the test result is positive?

P(T) = 0.03P(D) = 0.01P(T|D) = 0.99P(D|T) = ?

$$P(D|T) = \frac{P(T|D)P(D)}{P(T)} = \frac{0.99 * 0.01}{0.03} = 0.33$$



Random variables and probability distributions



Random variables

Definition of a random variable

- A random variable is a numerical representation of a random phenomenon. The values of the random variable correspond to the outcomes of the random process
- The values of the random variable are called realizations
 - For example, flipping a coin is a random variable X with values
 0 (in case of heads) and 1 (in case of tails)
 - The color of a ball we draw from a pot with blue, red and green balls is a random variable Y with values 0,1 and 2, respectively
 - The outcome of throwing a dice is a random variable Z with values {1,2,3,4,5,6}
- Notation:
 - Random variables X,Y,Z
 - Realizations x_i, y_i, z_i



Random variables

Discrete random variables

- Random variables with a finite, countable set of possible values are called **discrete random variables**
 - The number of heads after flipping a coin n times
 - The number of children in a family
 - The gender of a randomly selected student from the class
 - The field of studies from a randomly selected student on the campus at a given time point



Random variables

Probability (mass) function

- Each outcome of a random experiment is associated with a specific probability, therefore each value of a random variable has a corresponding probability.
- Let X be a discrete random variable. Then the function

 $f(x_i) = P(X = x_i)$

is called the **probability (mass) function** of \boldsymbol{X}

- Following the rules for probabilities:
 - $0 \le f(x_i) \le 1$
 - $\sum_{i=1}^{n} f(x_i) = 1$



Cumulative distribution function

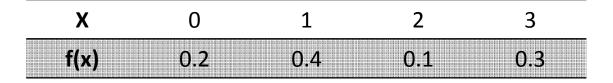
• Let X be a discrete random variable. Then the function $F(x_i) = P(X \le x_i)$

is called the **cumulative disctribution function** of X.

- Properties of F(x):
 - $0 \le F(x) \le 1$
 - A monotonically increasing function:
 - $F(x_i) < F(x_j)$ if $x_i < x_j$
 - $P(a < X \le b) = F(b) F(a)$

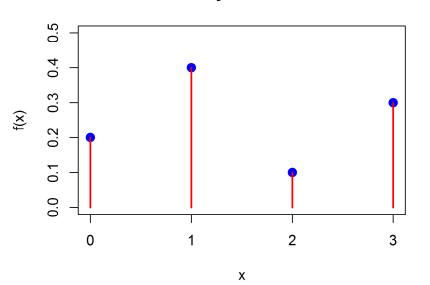


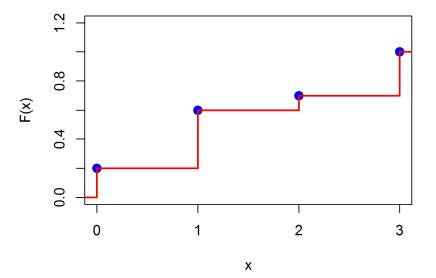
Graphical representation of f(x) and F(x)



Probability mass function

Cumulative distribution function







Expected value of a discrete random variable

 Let X be a discrete random variable with a probability function f(x). The expected value E(X) is:

$$E(X) = \mu = \sum_{i=1}^{n} x_i f(x_i)$$

- E(X) is a sum of the values of X weighted by their probability
- E(X) is the value we would expect X to take if we repeat the experiment numerous times.



Expected value of a discrete random variable



$$E(X) = \mu = \sum_{i=1}^{n} x_i f(x_i)$$

= 0 * 0.2 + 1 * 0.4 + 2 * 0.1 + 3 * 0.3
= 1.5



Continuous random variables

- A random variable which can take an infinite number of values in an interval [a; b] is called a continuous random variable
 - Height and weight of a sample of citizens
 - Blood pressure of patients in a study to test anti-hypertension therapy
 - The time needed to solve an exercise during an exam
 - The daily revenue of a supermarket



Probability density function

 Since there are infinitely many possible values for a continuous random variable, the probability that X takes exactly a specific value is 0

$$P(X=x)=0$$

- Continuous random variables are characterized by their (probability) density function f(x).
- f(x) is used to specify the probability of X falling in a particular range of values



Cumulative distribution function

• Let X be a continuous random variable. The function $F(x) = P(X \le x)$

is called **cumulative distribution function**.

• F(x) is the area under the density function curve

$$F(a) = \int_{-\infty}^{a} f(x) dx$$

 Following the rules of probability, the total area under the density curve must be 1

$$\int_{-\infty}^{\infty} f(x) dx = 1$$

- $P(X \ge a) = 1 P(X \le a) = 1 F(a)$
- $P(a \le X \le b) = F(b) F(a)$



Graphical representation of f(x) and F(x)

 Let X be a continuous random variable taking values from -1 to 0. The density function is given by:

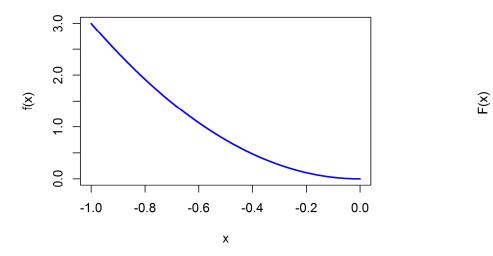
$$f(x) = \begin{cases} 3x^2, x \in [-1; 0] \\ 0, otherwise \end{cases}$$

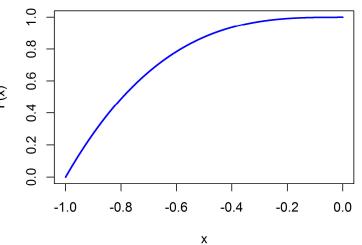
• The cumulative distribution function is therefore:

$$F(x) = \int_{-1}^{x} 3x^2 dx = x^3 + 1$$

Probability density function









Graphical representation of f(x) and F(x)

 Let X be a continuous random variable taking values from -1 to 0. The density function is given by:

$$f(x) = \begin{cases} 3x^2, x \in [-1; 0] \\ 0, otherwise \end{cases}$$

• What is the probability of X < -0.4?

$$P(X < -0.4) = \int_{-1}^{-0.4} 3x^2 dx = F(-0.4) = 0.934$$





-0.8

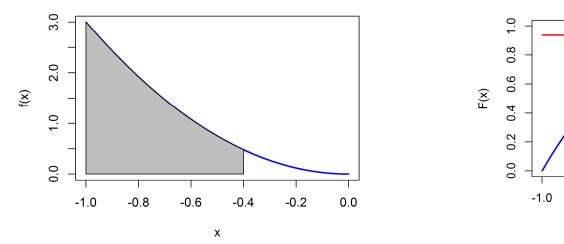
-0.6

х

-0.4

-0.2

0.0



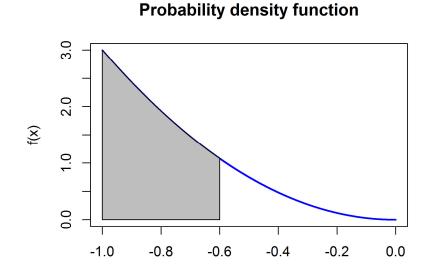


Graphical representation of f(x) and F(x)

 Let X be a continuous random variable taking values from -1 to 0. The density function is given by:

$$f(x) = \begin{cases} 3x^2, x \in [-1; 0] \\ 0, otherwise \end{cases}$$

- What is the probability that $X \ge 0.6$
- $P(X \ge -0.6) = P(X \ge -0.6) = 1 P(X \le -0.6) = 1 F(-0.6)$



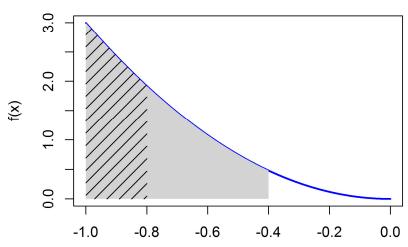


Graphical representation of f(x) and F(x)

 Let X be a continuous random variable taking values from -1 to 0. The density function is given by:

$$f(x) = \begin{cases} 3x^2, x \in [-1; 0] \\ 0, otherwise \end{cases}$$

- What is the probability that X lies between -0.8 and -0.4?
- P(-0.8 < X < -0.4) = F(-0.4) F(-0.8)





Expected value of a continuous random variable

• Remember, E(X) for a discrete random variable is

$$E(X) = \mu = \sum_{i=1}^{n} x_i f(x_i)$$

- Since a continuous random variable can take infinitely many values, we cannot build a weighted sum of xf(x) to calculate E(X)
- Instead, the expected value is the total area under the function xf(x):

$$E(X) = \mu = \int_{-\infty}^{\infty} x f(x) dx$$



Expected value of a continuous random variable

 Let X be a continuous random variable taking values from -1 to 0. The density function is given by:

$$f(x) = \begin{cases} 3x^2, x \in [-1; 0] \\ 0, otherwise \end{cases}$$

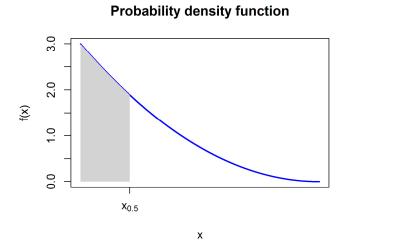
• The expected value is therefore:

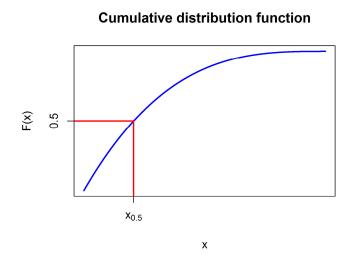
$$\int_{-1}^{0} x \, 3x^2 dx = -\frac{3}{4}$$



Quantiles

- Quantiles are values of a random variable corresponding to specific values of the cumulative probability function
 - $q_{0.1} = a \rightarrow F(a) = 0.1$
 - $q_{0.3} = b \rightarrow F(b) = 0.3$
 - $q_{0.5} = c \rightarrow F(c) = 0.5$
 - Values which are not exceeded with a specific probability or values corresponding to specific areas under the density function curve







Motivation for the normal distribution

- Many **empirical distributions** can be approximated by a normal distribution (e.g. height of adults, duration of pregnancy)
- Represents the **limit distribution** of many other probability distributions (e.g. binomial, poisson, t-distribution)
- **Sampling distributions** asymptotically approximate the normal distribution for large sample sizes
- The normal distribution provides the theoretical basis for numerous models in statistics

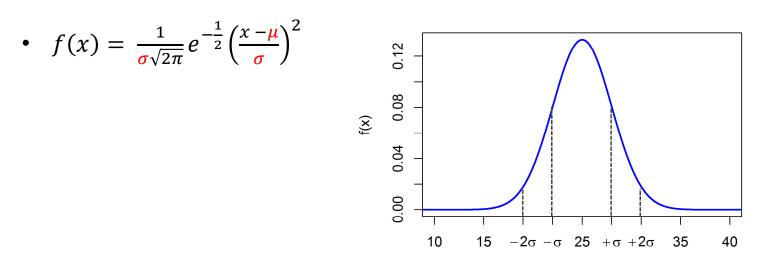


Features of the normal distribution

Completely defined by its two paramers expected value μ and variance σ²:

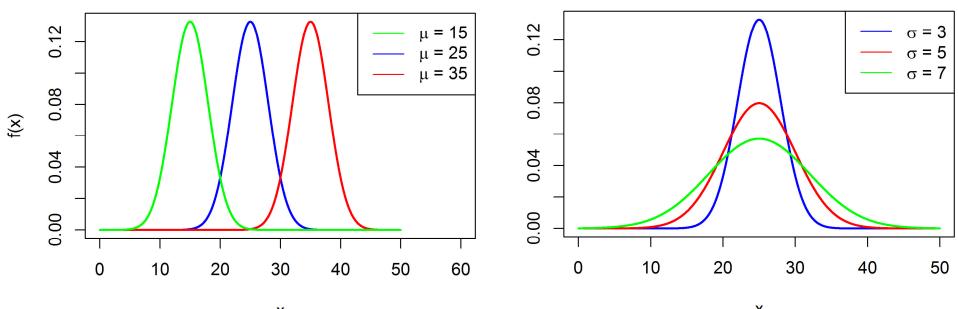
$$X \sim \mathcal{N} (\mu; \sigma^2)$$

- Density function is unimodal with characteristic "bell" shape
- Density is maximal at $X = \mu$ and symmetrical around μ with inflection points $+\sigma$ and $-\sigma$





Normal distributions for different parameters



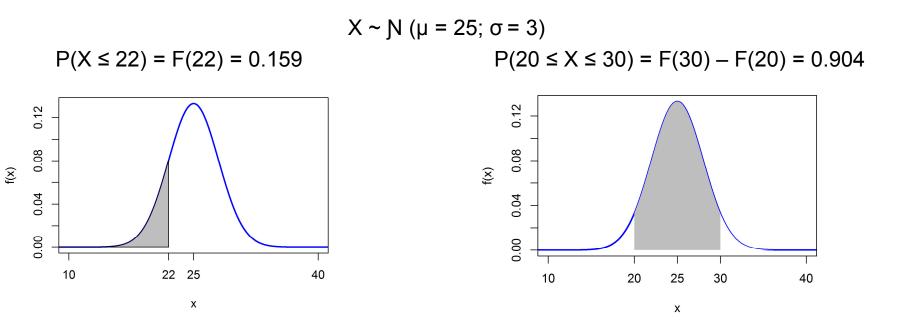
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Cumulative distribution function

- F(a) = P(X ≤ a) corresponds to the area under the density function curve
- Total area under the curve = 1, therefore $P(X \ge a) = 1 P(X \le a)$
- $P(a \le X \le b) = F(b) F(a)$

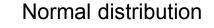




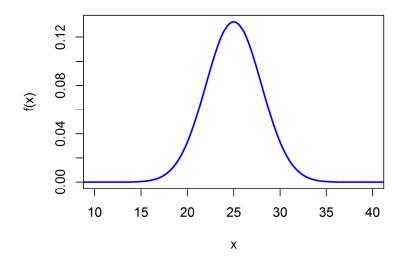
Standard normal distribution

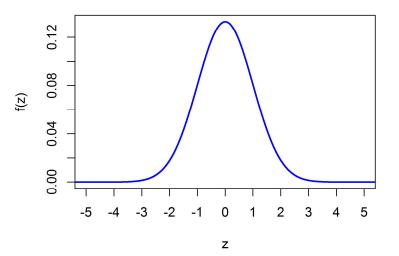
- Expected value $E(Z) = \mu = 0$
- Variance Var(X) = $\sigma^2 = 1$
- Every normal distribution can be transformed in the standard normal distribution using the following transformation

$$Z = \frac{X - \mu}{\sigma} \sim \mathcal{N}(0, 1)$$



Standard normal distribution



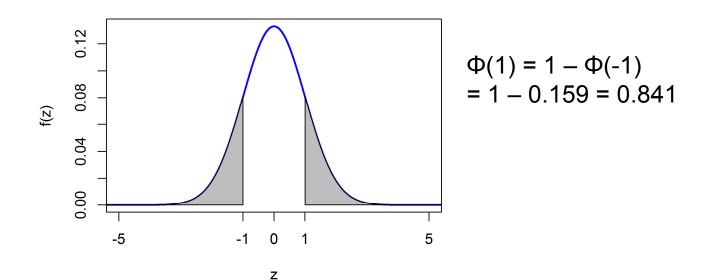




Cumulative distribution function

•
$$F(a) = P(X \le a) = P(\frac{X-\mu}{\sigma} \le \frac{a-\mu}{\sigma}) = \Phi(\frac{a-\mu}{\sigma})$$

- $\Phi(z)$ is the cumulative distribution function of the standard normal distribution.
- Values for $\Phi(z)$ are available in statistical tables
- Due to the symmetry of the distribution $\Phi(-z) = 1 \Phi(z)$





Quantiles of the standard normal distribution

- Quantiles of the standard normal distribution with the corresponding cumulative distribution function values are summarized in statistical tables
- Since there are infinitely many normal distributions, they are usually transformed into a standard normal distribution

z	0.00	0.01	0.02	0.03	0.04	0.05	0.06	0.07	0.08	0.09
0.0	.5000	.5040	.5080	.5120	.5160	.5199	.5239	.5279	.5319	.5359
0.1	.5398	.5438	.5478	.5517	.5557	.5596	.5636	.5675	.5714	.5753
0.2	.5793	.5832	.5871	.5910	.5948	.5987	.6026	.6064	.6103	.6141
0.3	.6179	.6217	.6255	.6293	.6331	.6368	.6406	.6443	.6480	.6517
0.4	.6554	.6591	.6628	.6664	.6700	.6736	.6772	.6808	.6844	.6879
0.5	.6915	.6950	.6985	.7019	.7054	.7088	.7123	.7157	.7190	.7224
0.6	.7257	.7291	.7324	.7357	.7389	.7422	.7454	.7486	.7517	.7549
0.7	.7580	.7611	.7642	.7673	.7704	.7734	.7764	.7794	.7823	.7852
0.8	.7881	.7910	.7939	.7967	.7995	.8023	.8051	.8078	.8106	.8133
0.9	.8159	.8186	.8212	.8238	.8264	.8289	.8315	.8340	.8365	.8389
1.0	.8413	.8438	.8461	.8485	.8508	.8531	.8554	.8577	.8599	.8621



Inferential statistics and hypothesis testing

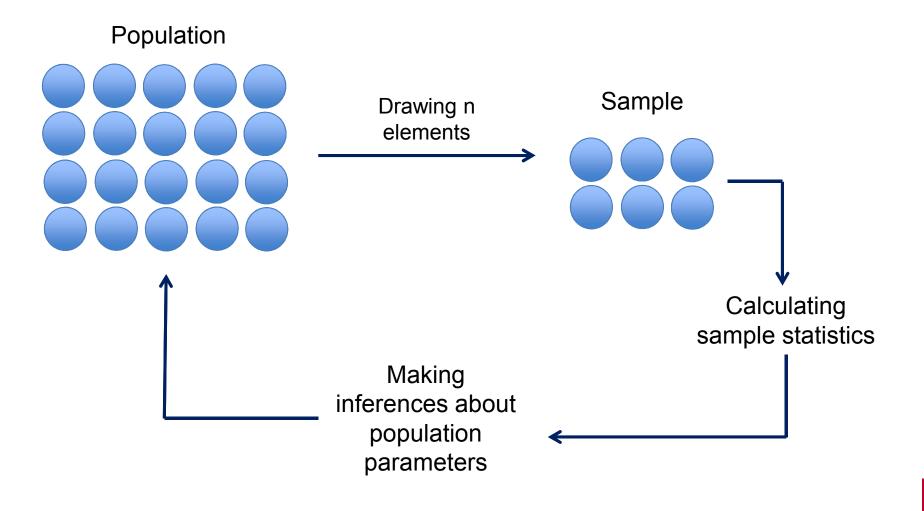


Population vs. sample

- Statistical population a set of similar elements of interest in a reasearch question or experiment
 - E.g. All patients undergoing cardiac surgery over the age of 65
 - Usually it is not possible to investigate a whole population due to time or cost reasons
- Statistical sample a subset of a statistical population chosen by following a specific stragegy or criterion
 - E.g. All patients undergoing heart valve surgery at the University Medicine Mainz in the period between January 2017 and December 2017



Population vs. sample





Population parameters and sample statistics

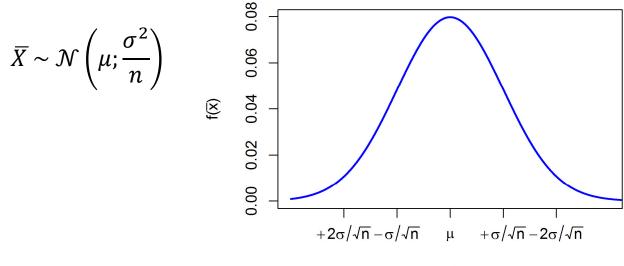
- Population parameters are unknown because the entire population is not available
- The measures which characterize samples are called statistics directly calculated from the data
- Sample statistics are used to infer population parameters

Population parameter	Sample statistic				
Expected value μ	Sample mean \overline{x}				
Variance o²	Sample variance s²				
Standard deviation σ	Sample standard deviation <i>s</i>				
Correlation <i>p</i>	Correlation coefficient <i>r</i>				



Sampling distributions

- Sample statistics are random variables because they are based on random samples
- Therefore, sample statistics have their own probability distributions called **sampling distributions**
 - E.g. Let the height of Germans in cm be a normally distributed random variable X with mean μ and variance σ². Let X be the mean height in a sample of size n.
 - \overline{X} is a random variable with the following distribution:



Confidence intervals

- The sample mean, variance, standard deviation, etc. are so called point estimates for the respective population parameters because they provide only a single value
- The probability that the point estimate is equal to the true parameter is equal to 0 for continuous random variables
- Interval estimates provide a range of values which could contain the population parameter and are therefore more informative
- Confidence intervals are calculated from the sample data at predefined confidence level 1 - α

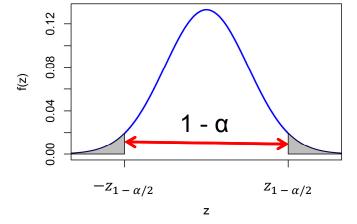


Confidence intervals

- Confidence interval for normally distributed parameters, e.g. *µ*
 - Let C_u and C_o be the lower and upper limits of a 1- α confidence interval for μ

$$\begin{aligned} 1 - \alpha &= P(C_u \leq \bar{X} \leq C_o) \\ 1 - \alpha &= P(\frac{C_u - \mu}{\sigma} \sqrt{n} \leq \frac{\bar{X} - \mu}{\sigma} \sqrt{n} \leq \frac{C_o - \mu}{\sigma} \sqrt{n}) \\ 1 - \alpha &= P(-z_{1 - \alpha/2} \leq \frac{\bar{X} - \mu}{\sigma} \sqrt{n} \leq z_{1 - \alpha/2}) \\ 1 - \alpha &= P(\bar{X} - z_{1 - \frac{\alpha}{2}} \frac{\sigma}{\sqrt{n}} \leq \mu \leq \bar{X} + z_{1 - \alpha/2} \frac{\sigma}{\sqrt{n}}) \end{aligned}$$

• Typical values for α include 0.05, 0.01, 0.001



Confidence intervals

Calculate the 95 % confidence interval for the average total bilirubin concentration of 2.7 mg/dL in a sample of 36 patients with moderate liver injury with a standard deviation of 0.4

$$1 - \alpha = 95 \% = 0.95 \rightarrow \alpha = 0.05$$

$$\pm z_{1-\alpha/2} = \pm z_{1-\frac{0.05}{2}} = \pm z_{0.975} = \pm 1.96$$

$$95\% CI = \bar{X} \pm z_{1-\alpha/2} \frac{\sigma}{\sqrt{n}}$$

$$95\% CI = 2.7 \pm 1.96 \frac{0.4}{\sqrt{36}}$$

$$95\% CI = [2.57; 2.83]$$

- Interpretation of CI
 - If confidence intervals were repeatedly calculated for many samples, then 1 - α of the intervals would countain the true population parameter



- Researchers are often interested in yes/no questions regarding population parameters
- E.g. A new medicine to prevent cold was tested against a control treatment. In a small pilot experiment, 20 subjects were randomized to the control and 20 to the new treatment. 11 subjects in the control group got a cold compared to 7 subjects in the treatment group. *Does the new treatment prevent getting a cold?*

	Control	Treatment
Got a cold	11	7
Remained healthy	9	13



Hypothesis tests

- Results based on sample statistics initially only apply to the investigated sample
- Hypothesis tests are needed to decide if the observed results are due to population differences or simply the result of chance.
- The basis of hypothesis test is establishing the **hypothesis pair**:
 - Null hypothesis: Represents the status quo, absence of a difference – usually the hypothesis we want to disprove

 H_0 : The new therapy does not reduce the risk of getting a cold.

Alternative hypothesis: Represents the alternative to H₀ – usually represents the desired outcome

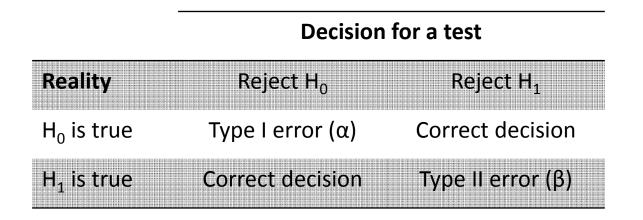
 H_1 : The new therapy prevents getting a cold.



- The null and alternative hypothesis make assumptions about an unknown population parameter Θ (e.g. population mean µ or population proportion p)
- Types of hypothesis pairs:
 - $H_0: \Theta = \Theta_0$ vs. $H_1: \Theta \neq \Theta_0$: **two-tailed** test
 - E.g. Expression levels of gene X under condition A are different compared to condition B
 - $H_0: \Theta = \Theta_0 \text{ vs. } H_1: \Theta < \Theta_0: \text{ left-tailed test}$
 - E.g. Therapy A reduces blood pressure compared to therapy B
 - $H_0: \Theta = \Theta_0 \text{ vs. } H_1: \Theta > \Theta_0: \text{ right-tailed test}$
 - E.g. Therapy A increases survival time in cancer patients compared to control.



- At the end of a specific test we always retain one of the hypotheses and reject the other. However, we never know if this decision is correct.
- What we do instead is try to reduce the probability of a false decision
- The following possibilities exist:





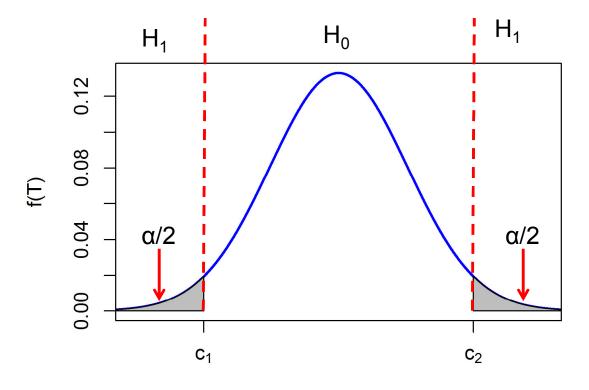
- Researchers typically control the type I error probability, each statistical test is performed at a predefined **significance level** α.
- Typical values include $\alpha = 0.05$, $\alpha = 0.01$, $\alpha = 0.001$
- In order to decide which hypothesis to retain, a test statistic T is calculated.
- **T** is a function of the sample statistic, e.g. the sample mean \bar{X}
 - Remember that hypothesis are defined for the unknown population parameters
- **T** is therefore a random variable with a probability distribution
- In order to decide which hypothesis to retain/reject, the distribution of T under the null hypothesis is evaluated.
- The null hypothesis is rejected if T exceeds critical values which depend on α.



Hypothesis tests

Distribution of the test statistic **T**, two-tailed test

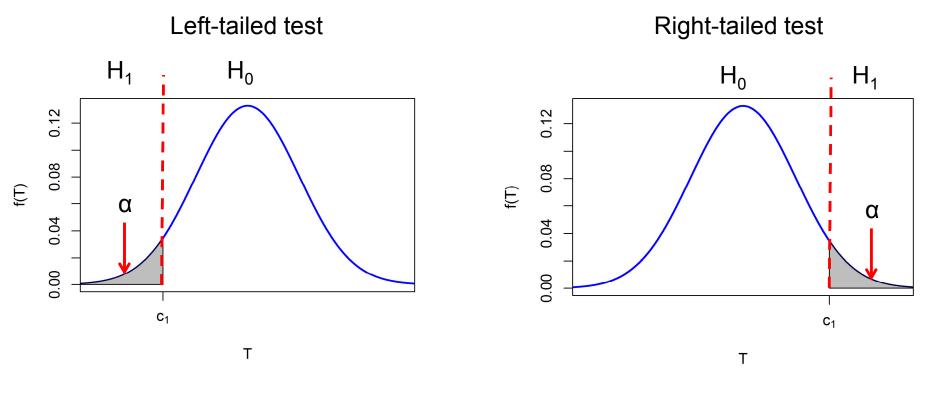
- Reject H_0 if $T < c_1$ or if $T > c_2$
- c₁ and c₂ are the so called critical values



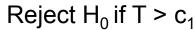


Hypothesis tests

Distribution of the test statistic **T**, left- and right- tailed tests



Reject H_0 if $T < c_1$





Steps in hypothesis testing

- 1. Evaluation of test assumptions
- 2. Definition of the null and alternative hypothesis
- 3. Definition of significance level
- 4. Calculation of the test statistic
- 5. Definition of the rejection region based on the probability distribution of the test statistic, critical values specification
- 6. Comparison of the test statistic with critical values
- 7. Decision to retain/reject null hypothesis and interpretation



One sample z test

• The one sample z-test investigates if a population mean μ significantly deviates from a given value μ_0 when the population variance is known.

E.g. The average note in the final biostatistics exam of biomedicine students at the University Mainz from the last 5 years was 2.2 with a known variance of 0.9. The average note of the 49 students who took the exam this year was 1.8. Does the result for this year's exam represent a systematic change or only random fluctuation?

Investigate hypothesis with the help of the one sample ztest using the 7-point scheme outlined on the previous slide



One sample z test example

The average note in the final biostatistics exam of biomedicine students at the University Mainz from the last 5 years was 2.2 with a known variance of 0.9. The average note of the 49 students who took the exam this year was 1.8. Does the result for this year's exam represent a systematic change or only random fluctuation?

- 1. Evaluation of test assumptions
 - Normally distributed random variable and known population variance
- 2. Definition of the null and alternative hypothesis
 - Null hypothesis: The average note of biomedicine students in statistics is 2.2

 $H_0: \mu = 2.2$

 Alternative: The average note of biomedicine students in statistics is not 2.2.

 $H_1: \mu \neq 2.2 \longrightarrow$ two-sided test



One sample z test example

The average note in the final biostatistics exam of biomedicine students at the University Mainz from the last 5 years was 2.2 with a known variance of 0.9. The average note of the 49 students who took the exam this year was 1.8. Does the result for this year's exam represent a systematic change or only random fluctuation?

- 3. Definition of significance level
 - α = 0.05 (type I error = 5 %)
- 4. Calculation of the test statistic

$$T = \frac{\overline{X} - \mu_0}{\sigma} \sqrt{n}$$

 \bar{X} = 1.8, μ_0 = 2.2, σ = $\sqrt{0.9}$, n $\,$ = 49 $\,$

$$T = \frac{1.8 - 2.2}{\sqrt{0.9}}\sqrt{49} = -2.951$$



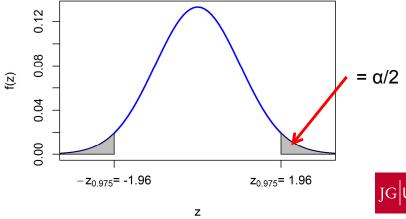
One sample z test example

The average note in the final biostatistics exam of biomedicine students at the University Mainz from the last 5 years was 2.2 with a known variance of 0.9. The average note of the 49 students who took the exam this year was 1.8. Does the result for this year's exam represent a systematic change or only random fluctuation?

5. Definition of the rejection region based on the probability distribution of the test statistic, critical values specification

- When assumptions of the test are met, the test statistic follows a standard normal distribution, T ~ N(0,1).
- Critical values for a two sided test:

$$\pm z_{1-\alpha/2} = \pm z_{1-0.025} = \pm z_{0.975} = \pm 1.96$$



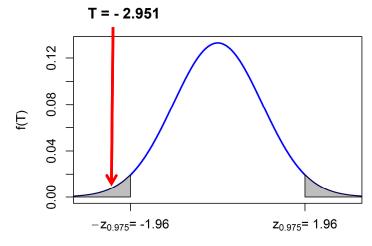
One sample z test example

The average note in the final biostatistics exam of biomedicine students at the University Mainz from the last 5 years was 2.2 with a known variance of 0.9. The average note of the 49 students who took the exam this year was 1.8. Does the result for this year's exam represent a systematic change or only random fluctuation?

6. Comparison of the test statistic with critical values

$$T = -2.951 < -z_{0.975} = -1.96$$

Test statistic exceeds the critical value





One sample z test example

The average note in the final biostatistics exam of biomedicine students at the University Mainz from the last 5 years was 2.2 with a known variance of 0.9. The average note of the 49 students who took the exam this year was 1.8. Does the result for this year's exam represent a systematic change or only random fluctuation?

- 7. Decision to retain/reject null hypothesis and interpretation
 - Null hypothesis is rejected
 - Interpretation:

The average note of biomedicine students in statistics this year is significantly different from the 5 year-average.

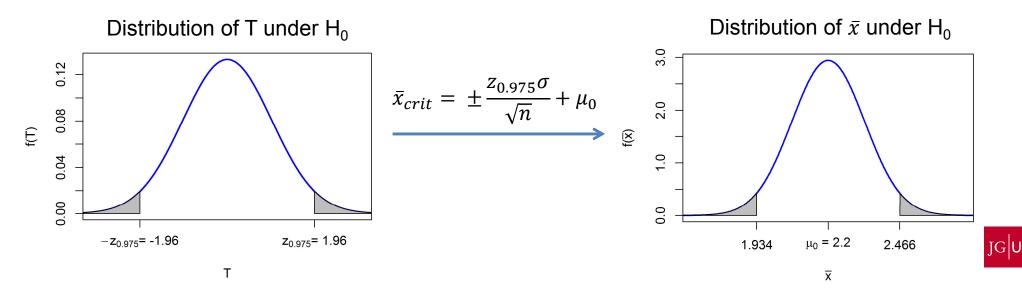


- Usually, the desired outcome of a hypothesis test is to find sufficient empirical evidence to reject the null hypothesis.
- Even if H₁ is true, we still might fail to reject H₀ (Type II error: false negative)
- Type II errors can have serious consequences especially in the field of clinical research
- Power = 1β , probability to reject a false H₀
- Ways to increase power of a test:
 - Increase the significance level
 - Increase sample size

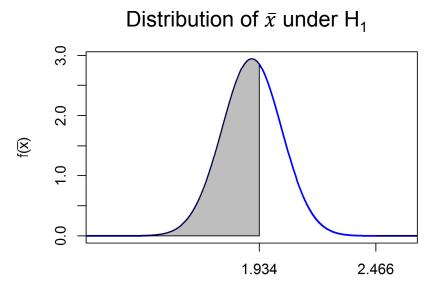


- Power can only be calculated for specific values of the parameters under the alternative hypothesis.
- The average note in the final biostatistics exam of biomedicine students at the University Mainz from the last 5 years was 2.2 with a known variance of 0.9. The average note of the 49 students who took the exam this year was 1.8. What is the power of a two-sided z-test given the alternative hypothesis that $\mu_1 = 1.9$?

Power = P(T <
$$-z_{0.975}$$
 or T > $z_{0.975}$ | μ_1 = 1.9) = ?



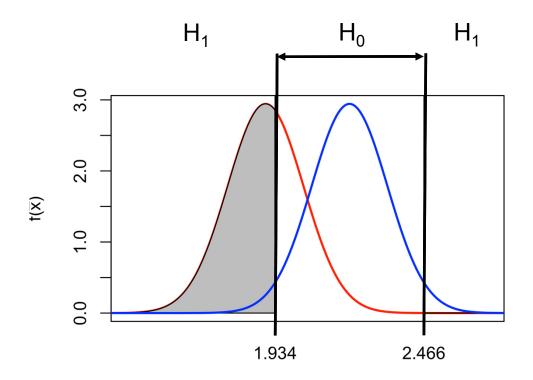
- Reject H_0 if $\bar{x} < 1.934$ or $\bar{x} > 2.466$
- Power = P($\bar{x} < 1.934$ or $\bar{x} > 2.466 \mid \mu_1 = 1.9$) = 0.599







- Reject H_0 if $\bar{x} < 1.934$ or $\bar{x} > 2.466$
- Power = P($\bar{x} < 1.934$ or $\bar{x} > 2.466 \mid \mu_1 = 1.9$) = 0.599





Effect size

- Rejecting a null hypothesis does not automatically imply that results have practical relevance
- In large samples, even trivial differences might be statistically significant
- This calls for standard measures to evaluate effect size, e.g.:
 - Standardized mean difference
 - Percentage of variance explained by the model
- Effect sizes should be reported with results from significance tests



Effect size example

• Cohen's *d* is a measure of effect size evaluating standardized mean difference

• Cohen's
$$d = \frac{|\mu_1 - \mu_2|}{\sigma}$$

- Reference values:
 - d = 0.2 small effect
 - d = 0.5 medium effect
 - d > 0.8 large effect
- E.g. The average note of biomedicine students from this year's exam $\bar{x} = 1.8$ was shown to be significantly different from the 5-year average of 2.2. What is the effect size?

$$d = \frac{|1.8 - 2.2|}{\sqrt{0.9}} = 0.422$$



Power and effect size in controlled experiments

- Controlled clinical experiments are associated with significant costs and time
- Failing to detect a true alternative hypothesis could be detrimental
- From a statistical point of view, studies are planned to maximize the chances of proving a study hypothesis which is also of practical relevance:
 - Sample size is based on the level of power desired to be achieved, typically 80%
 - Effect size is based on theoretical knowledge or pilot investigations

