

## EXAM OF STATISTICS (DESCRIPTIVE STATISTICS AND REGRESSION)

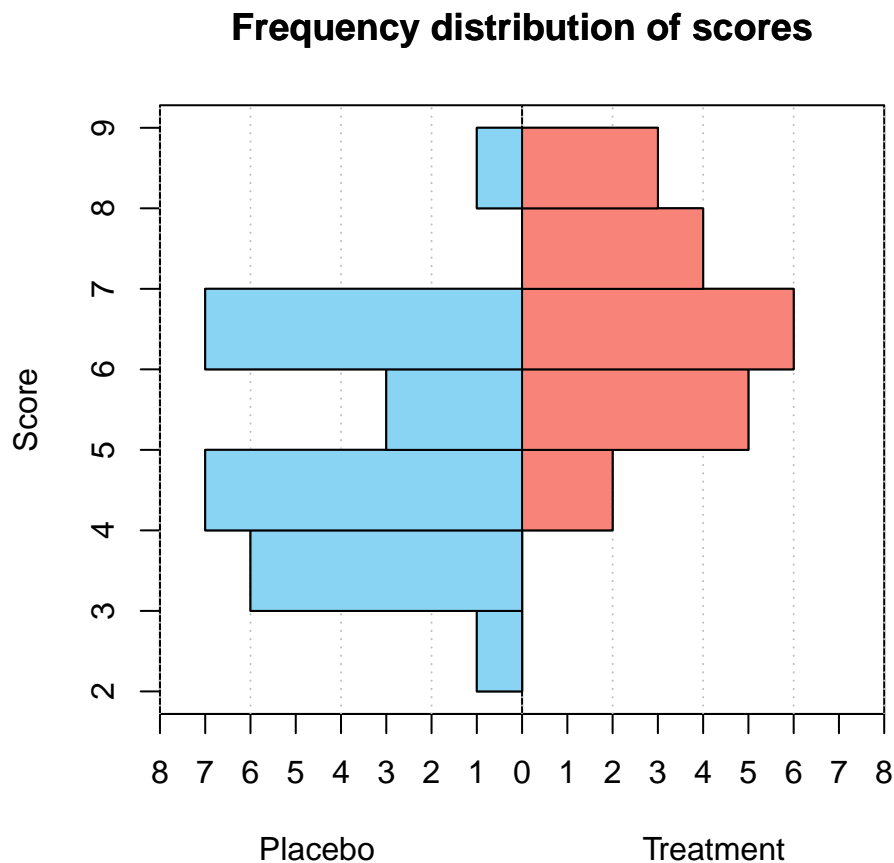
2nd Physiotherapy

Version A

June, 07 2021

**Duration:** 1 hour.

- (5 pts.) 1. To study the effectiveness of a new treatment for the polymyalgia rheumatica a sample of patients with polymyalgia was drawn and they were divided into two groups. The first group received the new treatment while the second one received a placebo. After a year following the treatment they filled out a survey. The chart below shows the distribution of the survey score of the two groups of patients (the greater the score the better the treatment).



- Construct the frequency table of the scores for the placebo group and plot the ogive.
- Compute the interquartile range of the scores for the placebo group.
- Are there outliers in the placebo group?
- In which group the score mean represents better?
- Which distribution is more normal regarding the kurtosis?
- Which score is relatively better, a score of 5 in the placebo group or a score of 6 in the treatment group?

Use the following sums for the computations:

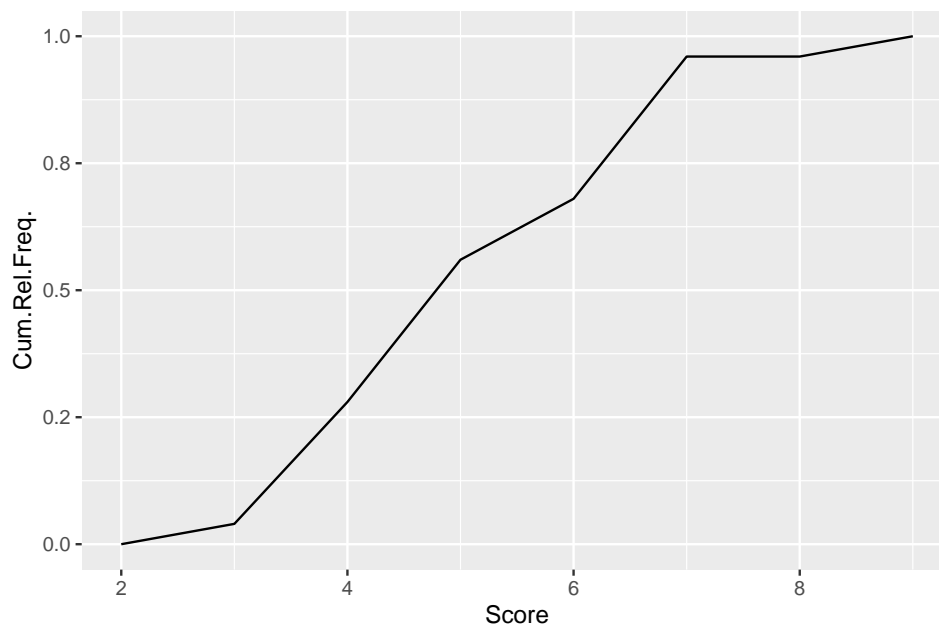
Placebo:  $\sum x_i = 125.5$ ,  $\sum x_i^2 = 680.25$ ,  $\sum (x_i - \bar{x})^3 = 27.11$  and  $\sum (x_i - \bar{x})^4 = 253.27$ .

Treatment:  $\sum x_i = 131$ ,  $\sum x_i^2 = 887$ ,  $\sum (x_i - \bar{x})^3 = 2.66$  and  $\sum (x_i - \bar{x})^4 = 88.03$ .

### Solution

(a)

Score	$n_i$	$f_i$	$N_i$	$F_i$
[2, 3]	1	0.04	1	0.0
(3, 4]	6	0.24	7	0.3
(4, 5]	7	0.28	14	0.6
(5, 6]	3	0.12	17	0.7
(6, 7]	7	0.28	24	1.0
(7, 8]	0	0.00	24	1.0
(8, 9]	1	0.04	25	1.0



(b)  $Q_1 = 3.875$ ,  $Q_3 = 6.25$  and  $IQR = 2.375$ .

(c)  $f_1 = 0.3125$  and  $f_2 = 9.8125$ . Thus, there are no outliers in the placebo sample because all the values fall between the fences.

(d) Placebo:  $\bar{x} = 5.02$ ,  $s^2 = 2.0096$ ,  $s = 1.4176$  and  $cv = 0.2824$ .

Treatment:  $\bar{x} = 6.55$ ,  $s^2 = 1.4475$ ,  $s = 1.2031$  and  $cv = 0.1837$ .

(e) Placebo:  $g_2 = -0.4914$ . Treatment:  $g_2 = -0.8992$ . Thus, the distribution of the placebo group is more normal as the coef. of kurtosis is closer to 0.

(f) Standard score for the placebo:  $z(5) = -0.0141$

Standard score for the treatment:  $z(6) = -0.4571$

As the standard score of 5 in the placebo group is greater than the standard score of 6 in the treatment group, a score of 5 in the placebo group is better.

- (5 pts.) 2. We have applied different doses of an antibiotic to a culture of bacteria. The table below shows the number of residual bacteria corresponding to the different doses.

Dose ( $\mu\text{g}$ )	0.2	0.7	1	1.5	2	2.4	2.8	3
Bacteria	40	32	28	20	18	15	12	11

- Which regression model explains better the number of residual bacteria as a function of the antibiotic dose, the linear or the exponential?
- Use the best of the two previous regression models to predict the number of residual bacteria for an antibiotic dose of  $3.5 \mu\text{g}$ . Is this prediction reliable?
- According to the linear regression model, what is the expected decrease in the number of residual bacteria per each  $\mu\text{g}$  more of antibiotic?

Use the following sums for the computations ( $X$ =Antibiotic dose and  $Y$ =Number of bacteria):

$$\begin{aligned} \sum x_i &= 13.6 \mu\text{g}, \sum \log(x_i) = 2.1362 \log(\mu\text{g}), \sum y_j = 176 \text{ bacteria}, \sum \log(y_j) = 23.9638 \log(\text{bacteria}), \\ \sum x_i^2 &= 30.38 \mu\text{g}^2, \sum \log(x_i)^2 = 6.3959 \log(\mu\text{g})^2, \sum y_j^2 = 4622 \text{ bacteria}^2, \sum \log(y_j)^2 = 73.3096 \\ &\log(\text{bacteria})^2, \\ \sum x_i y_j &= 227 \mu\text{g} \cdot \text{bacteria}, \sum x_i \log(y_j) = 37.4211 \mu\text{g} \cdot \log(\text{bacteria}), \sum \log(x_i) y_j = -17.633 \log(\mu\text{g}) \text{ bacteria}, \\ \sum \log(x_i) \log(y_j) &= 3.6086 \log(\mu\text{g}) \log(\text{bacteria}). \end{aligned}$$

---

### Solution

- $\bar{x} = 1.7 \mu\text{g}$ ,  $s_x^2 = 0.9075 \mu\text{g}^2$ .  
 $\bar{y} = 22 \text{ bacteria}$ ,  $s_y^2 = 93.75 \text{ bacteria}^2$ .  
 $s_{xy} = -9.025 \mu\text{g} \cdot \text{bacteria}$ .  
Linear coefficient of determination  $r^2 = 0.9574$ .  
 $\overline{\log(y)} = 2.9955 \log(\text{bacteria})$ ,  $s_{\log(y)}^2 = 0.1908 \log(\text{bacteria})^2$ .  
 $s_{x \log(y)} = -0.4147 \mu\text{g} \cdot \log(\text{bacteria})$ .  
Exponential coefficient of determination  $r^2 = 0.9928$ .  
 Thus, the exponential model explains better the number of residual bacteria as a function of the antibiotic dose because the exponential coef. of determination is greater.
  - Exponential regression model:  $y = e^{3.7723 + -0.4569x}$ .  
 Prediction:  $y(3.5) = 8.7845 \text{ bacteria}$ .  
 Although the coef. of determination is close to 1, the this prediction is not reliable because the sample size is very small.
  - $b_{yx} = -9.9449$ , therefore the number of bacteria decreases  $-9.9449$  per each  $\mu\text{g}$  more of antibiotic.
-