Understanding P-values

Fatima Batool

April 16, 2024

Test of Significance Procedure

- Define the null and the alternative Hypothesis
- Compute the test statistics
- 3 Compute the p-value
- Decide whether reject or fail to reject a null hypothesis

State Hypothesis

- Testing a single value of a parameter
- Is a coin fair?
- Is a true coefficient β of linear regression equals zero?
- Does the population have mean (μ) equal to some value (μ_0) ?
- Is there a difference in corneal thickness between the Glaucoma and Normal eyes?
- Does the BMI for the covid-19 patients equal to the expected normal BMI for a given age.

$$\begin{split} H_0:\theta &= \theta_0, & H_0:\theta \leqslant \theta_0, & H_0:\theta \geqslant \theta_0, \\ H_A:\theta &\neq \theta_0. & H_A:\theta > \theta_0. & H_A:\theta < \theta_0. \end{split}$$

Test Statistics

If the parameter is population mean, and population variance σ^2 is known, $H_0: \mu = \mu_0$, and $H_A: \mu \neq \mu_0$:

$$z=\frac{\sqrt{n}(\bar{X}-\mu)}{\sigma}.$$

If σ^2 is not known we use the *t*-statistics:

$$t=\frac{\sqrt{n}(\bar{X}-\mu)}{S}.$$

If the parameter is population proportion P:

$$z = \frac{(\hat{P} - P)}{\sqrt{\frac{PQ}{n}}}.$$

The *p*-value

- The P-value is the probability value of observing a test statistics as or more extreme than the one calculated from the sample under assumption that null is true.
- A small p-value is an evidence against null, because it is a probability of seeing such an extreme (big or small) value of test statistics, if the null actually holds.

The *p*-value

- The P-value is the probability value of observing a test statistics as or more extreme than the one calculated from the sample under assumption that null is true.
- A small p-value is an evidence against null, because it is a probability of seeing such an extreme (big or small) value of test statistics, if the null actually holds.

For a Right-tailed rejection region:

$$p$$
-value = $P(t_{df} \ge t)$.

For a Left-tailed rejection region:

$$p$$
-value = $P(t_{df} \leqslant -t)$.

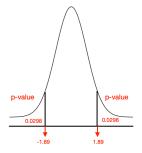
For a Two-tailed rejection region:

$$p$$
-value = $P(|t_{df}| \ge t)$.

p-value examples

■ *t*-statistics = -1.89, p-value = $P(|t_{310}| \ge 1.89)$ = 0.0597, two-tailed *t*-test,

R command: 2*pt(-1.89, df=310)

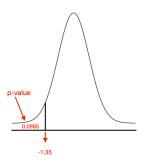


■ The p-value is 0.06, meaning if H_0 is true we will only see |t| this large 6% of the time, fails to reject null hypothesis, one possibility null hypothesis is actually true, other is we did not get evidence from this sample against null.

p-value examples contd..

■ *t*-statistics = -1.35, *p*-value = $P(t_{15} \le -1.35) = 0.0985$, left-tailed *t*-test,

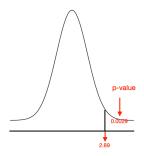
R command: pt(-1.35, df=15)



p-value examples contd..

■ *t*-statistics = 2.89, *p*-value = $P(t_{46} \ge 2.89) = 0.0029$, right-tailed *t*-test,

R command: pt(2.89, df=46, lower.tail = FALSE)



Conclusion

A small *p*-value provides an evidence against null.

- If p-value less than 0.01, strong evidence against null hypothesis.
- If p-value is between 0.01 and 0.05, some evidence against null hypothesis.
- If p-value greater than 0.05, insufficient evidence against null hypothesis.

p-value miss-interpretations

- Small *p*-value does not equate to practical or scientific significance
- p-value is not probability of null hypothesis being true
- Significant p-value does not implies cause and effect