Show me your proof: Confidence intervals and p-values

Steve Simon, P.Mean Consulting

Facebook: www.facebook.com/pmean LinkedIn: www.linkedin.com/in/pmean Main website: www.pmean.com Newsletter: www.pmean.com/news and www.facebook.com/group.php?gid=302778306676 Twitter: http://twitter.com/profmean Webinars: www.pmean.com/webinars

Why do I offer this webinar for free? I offer free statistics webinars partly for fun and partly to build up goodwill for my consulting business. For more information, go to <u>www.pmean.com/consult.html</u>.



P-values are the fundamental tools used in most inferential data analyses. They are possibly the most commonly reported statistics in the medical literature. Unfortunately, p-values are subject to frequent misinterpretations. In this presentation, you will learn the proper interpretation of p-values, and the common abuses and misconceptions about these statistics.



The pop quiz will not be graded. I do need to lay out some definitions. You may already know these terms, but it is still helpful to review them to assure that we are communicating using a common set of assumptions. I'll guide you through the first practice exercise, and then ask you to respond to the next two in the chat window. Then I'll repeat the pop quiz at the end. Please feel free to ask questions at anytime. I plan to finish no later than an hour after I have started, but will stick around if there are any questions.

<section-header><section-header><list-item><list-item><list-item><list-item><list-item><list-item><list-item><list-item><list-item><list-item>

Don't worry if you don't know the answer to this right away. I'm just trying to get a sense of how much people know in advance of the seminar. I'll repeat this pop quiz at the end as an informal evaluation of whether I have been effective at communicating my message to you.



A population is a collection of items of interest in research. The population represents a group that you wish to generalize your research to. Populations are often **defined in terms of demography, geography, occupation, time, care requirements, diagnosis, or some combination of the above**.

An example of a population would be all infants born in the state of Missouri during the 1995 calendar year who have one or more visits to the Emergency room during their first year of life.



A sample is a subset of a population. A random sample is a subset where every item in the population has the same probability of being in the sample. Usually, the size of the sample is much less than the size of the population. The primary goal of much research is to use information collected from a sample to try to characterize a certain population.



In your research, you specify a null hypothesis (typically labeled H0) and an alternative hypothesis (typically labeled Ha, or sometimes H1). By tradition, the null hypothesis corresponds to no change. A Type I error is **rejecting the null hypothesis when the null hypothesis is true**.

Example: Consider a new drug that we will put on the market if we can show that it is better than a placebo. In this context, H0 would represent the hypothesis that the average improvement (or perhaps the probability of improvement) among all patients taking the new drug is equal to the average improvement (probability of improvement) among all patients taking the placebo. A Type I error would be **allowing an ineffective drug onto the market**.



A Type II error is accepting the null hypothesis when the null hypothesis is false. Many studies have small sample sizes that make it difficult to reject the null hypothesis, even when there is a big change in the data. In these situations, a Type II error might be a possible explanation for the negative study results.

Example: Consider a new drug that we will put on the market if we can show that it is better than a placebo. In this context, H0 would represent the hypothesis that the average improvement (or perhaps the probability of improvement) among all patients taking the new drug is equal to the average improvement (probability of improvement) among all patients taking the placebo. A Type II error would be **keeping an effective drug off the market**.



The null hypothesis, traditionally represented by the symbol H0, represents the hypothesis of no change or no effect. The smaller the p-value, the more evidence we have against H0.

The p-value is also a measure of how likely we are to get a certain sample result or a result "more extreme," assuming H0 is true. The type of hypothesis (right tailed, left tailed or two tailed) will determine what "more extreme" means.

10. What is a p-value?

- Suppose that a drug company alleges that only 50% of all patients who take a certain drug will have an adverse event of some kind. You believe that the adverse event rate is much higher. In a sample of 12 patients, all twelve have an adverse event.
- P-value = 0.000244.

It is easiest to understand the p-value in a data set that is already at an extreme.

The data supports your belief because it is inconsistent with the assumption of a 50% adverse event rate. It would be like flipping a coin 12 times and getting heads each time.

The p-value, the probability of getting a sample result of **12 adverse events in 12 patients** assuming that the adverse event rate is 50%, is a measure of this inconsistency. The p-value, 0.000244, is small enough that we would reject the hypothesis that the adverse event rate was only 50%.

11. What is a p-value?
A small p-value means lots of evidence against the null hypothesis.
A large p-value means little or no evidence against the null hypothesis.
A p-value is NOT the probability that the null hypothesis is true.

The p-value is frequently misinterpreted as the probability that the null hypothesis is false. The framework in which p-values are derived assume a fixed (non-random) pair of hypotheses. Randomness appears in the use of a random sample to characterize information about a fixed population. This approach to hypothesis testing considers any statement of the probability of the null hypothesis to be absurd. An alternative approach, Bayesian statistics, places prior probabilities on hypotheses and parameters.



Perhaps the failure to reject the null hypothesis was caused by an inadequate sample size. When you see a large p-value in a research study, you should also look for one of two things:

a **power calculation** that confirms that the sample size in that study was adequate for detecting a clinically relevant difference; and/or

a confidence interval that lies entirely within the range of clinical indifference.

You should also be cautious about a small p-value, but for different reasons. In some situations, the sample size is so large that even differences that are trivial from a medical perspective can still achieve statistical significance.

13. Practice exercise: interpret the p-values shown below.

1. The Outcome of Extubation Failure in a Community Hospital Intensive Care Unit: A Cohort Study. Seymour CW, Martinez A, Christie JD, Fuchs BD. Critical Care 2004, 8:R322-R327 (20 July 2004) Introduction: Extubation failure has been associated with poor intensive care unit (ICU) and hospital outcomes in tertiary care medical centers. Given the large proportion of critical care delivered in the community setting, our purpose was to determine the impact of extubation failure on patient outcomes in a community hospital ICU. Methods: A retrospective cohort study was performed using data gathered in a 16-bed medical/surgical ICU in a community hospital. During 30 months, all patients with acute respiratory failure admitted to the ICU were included in the source population if they were mechanically ventilated by endotracheal tube for more than 12 hours. Extubation failure was defined as reinstitution of mechanical ventilation within 72 hours (n = 60), and the control cohort included patients who were successfully extubated at 72 hours (n = 93). Results: The primary outcome was total ICU length of stay after the initial extubation. Secondary outcomes were total hospital cost. Patient groups were similar in terms of age, sex, and severity of illness, as assessed using admission Acute Physiology and Chronic Health Evaluation II score (P > 0.05). Both ICU (1.0 versus 10 days; P < 0.01) and hospital length of stay (6.0 versus 17 days; P < 0.01) after initial extubation were significantly longer in reintubated patients. ICU mortality was significantly higher in patients who failed extubation (odds ratio = 12.2, 95% confidence interval [CI] = 1.5–101; P < 0.05), but there was no significant difference in hospital mortality (odds ratio = 2.1, 95% CI = 0.8–5.4; P < 0.15). Total hospital costs (estimated from direct charges) were significantly increased by a mean of US\$33, 926 (95% CI = US\$22,573–45,280; P < 0.01). Conclusion: Extubation failure in a community hospital is univariately associated with prolonged inpatient ca

14. Practice exercise: interpret the p-values shown below.

2. Elevated White Cell Count in Acute Coronary Syndromes: Relationship to Variants in Inflammatory and Thrombotic Genes. Byrne CE, Fitzgerald A, Cannon CP, Fitzgerald DJ, Shields DC. *BMC Medical Genetics 2004*, 5:13 (1 June 2004) **Background:** Elevated white blood cell counts (WBC) in acute coronary syndromes (ACS) increase the risk of recurrent events, but it is not known if this is exacerbated by pro-inflammatory factors. We sought to identify whether pro-inflammatory genetic variants contributed to alterations in WBC and C-reactive protein (CRP) in an ACS population. **Methods:** WBC and genotype of interleukin 6 (IL-6 G-174C) and of interleukin-1 receptor antagonist (IL1RN intronic repeat polymorphism) were investigated in 732 Caucasian patients with ACS in the OPUS-TIMI-16 trial. Samples for measurement of WBC and inflammatory factors were taken at baseline, i.e. Within 72 hours of an acute myocardial infarction or an unstable angina event. **Results:** An increased white blood cell count (WBC) was associated with an increased C-reactive protein (r = 0.23, p < 0.001) and there was also a positive correlation between levels of β-fibrinogen and C-reactive protein (r = 0.42, p < 0.0001). IL1RN and IL6 genotypes had no significant impact upon WBC. The difference in median WBC between the two homozygote IL6 genotypes was 0.21/mm3 (95% CI = -0.41, 0.77), and -0.03/mm3 (95% CI = -0.55, 0.86) for IL1RN. Moreover, the composite endpoint was not significantly affected by an interaction between WBC and the IL1 (p = 0.61) or IL6 (p = 0.48) genotype. **Conclusions:** Cytokine pro-inflammatory genetic variants do not influence the increased inflammatory profile of ACS patients.

15. Practice exercise: interpret the p-values shown below.

3. Is There a Clinically Significant Gender Bias in Post-Myocardial Infarction Pharmacological Management in the Older (>60) Population of a Primary Care Practice? Di Cecco R, Patel U, Upshur REG. *BMC Family Practice 2002*, 3:8 (3 May 2002) **Background**: Differences in the management of coronary artery disease between men and women have been reported in the literature. There are few studies of potential inequalities of treatment that arise from a primary care context. This study investigated the existence of such inequalities in the medical management of post myocardial infarction in older patients. **Methods**: A comprehensive chart audit was conducted of 142 men and 81 women in an academic primary care practice. Variables were extracted on demographic variables, cardiovascular risk factors, medical and non-medical management of myocardial infarction. **Results**: Women were older than men. The groups were comparable in terms of cardiac risk factors. A statistically significant difference (14.6%: 95% Cl 0.048–28.7 p = 0.047) was found between men and women for the prescription of lipid lowering medications. 25.3% (p = 0.0005, Cl 11.45, 39.65) more men than women had undergone angiography, and 14.4 % (p = 0.029, Cl 2.2, 26.6) more men than women are less likely than men to receive lipid-lowering medication which may indicate less aggressive secondary prevention in the primary care setting.

16. Repeat of pop quiz

A research paper computes a p-value of 0.45. How would you interpret this p-value?

- 1. Strong evidence for the null hypothesis
- 2. Strong evidence for the alternative hypothesis
- 3. Little or no evidence for the null hypothesis
- 4. Little or no evidence for the alternative hypothesis
- 5. None of the answers above are correct.
- 6. I do not know the answer.