When is statistical significance not significant?

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The article provides a non-technical introduction to the *p value* statistics. Its main purpose is to help researchers make sense of the appropriate role of the *p value* statistics in empirical political science research. On methodological grounds, we use replication, simulations and observational data to show when statistical significance is not significant. We argue that: (1) scholars must always graphically analyze their data before interpreting the *p value*; (2) it is pointless to estimate the *p value* for non-random samples; (3) the *p value* is highly affected by the sample size, and (4) it is pointless to estimate the *p value* when dealing with data on population.

**Keywords:** *p value* statistics; statistical significance; significance tests.
The basic problem with the null hypothesis significance test in political science is that it often does not tell political scientists what they think it is telling them. (J. Gill)

The statistical difficulties arise more generally with findings that are suggestive but not statistically significant. (A. Gelman and D. Weakliem)

The research methodology literature in recent years has included a full frontal assault on statistical significance testing. (J. E. McLean and J. M. Ernest)

Statistical significance testing has involved more fantasy than fact. (R. Carver)

Introduction

What is the fate of a research paper that does not find statistically significant results? According to Gerber, Green and Nickerson (2001: 01), “articles that do not reject the null hypothesis tend to go unpublished” Likewise, Sigelman (1999: 201) argues that “statistically significant results are achieved more frequently in published than unpublished studies. Such publication bias is generally seen as the consequence of a widespread prejudice against non significant results”. Conversely, Henkel (1976: 07) argues that significance tests “are of little or no value in basic social science research, where basic research is identified as that which is directed toward the development and validation of theory”. Similarly, McLean and Ernest (1998: 15) point out that significance tests provide no information about the practical significance of an event, or about whether or not the result is replicable. More directly, Carver (1978; 1993) argues that all forms of significance test should be abandoned. Considering this controversy, what is the appropriate role of the p value statistic in empirical political science research? This is our research question.

This paper provides a non-technical introduction to the p value statistic. Our main purpose is to help students in making sense of the appropriate role of the p value statistic in empirical political science research. On methodological grounds, we use observational data from the Quality of Government Institute simulations and replicate results from Anscombe (1973), Cohen (1988) and Hair et al., (2006) to show what can be learned from the p value statistic. There are situations where interpretation of the p value requires caution and we suggest four warnings: (1) scholars must always graphically analyze their data.
before interpreting the p value; (2) it is pointless to estimate the p value for non-random samples; (3) the p value is highly affected by the sample size, and (4) it is pointless to estimate the p value when dealing with data from population⁵.

The remainder of the paper consists of three sections. Firstly, we outline the underlying logic of null hypothesis significance tests, and we define what p value is and how it should be properly interpreted. Next, we replicate Anscombe (1973), Cohen (1988) and Hair et al., (2006) data, using basic simulation and analyze observational data to explain our view regarding the proper role of the p value statistic. We close with a few concluding remarks on statistical inference in political science.

What the p value is, what it means and what it does not

Statistical inference is based on the idea that it is possible to generalize results from a sample to the population⁶. How can we assure that relations observed in a sample are not simply due to chance? Significance tests are designed to offer an objective measure to inform decisions about the validity of the generalization. For example, one can find a negative relationship in a sample between education and corruption, but additional information is necessary to show that the result is not simply due to chance, but that it is “statistically significant”. According to Henkel (1976), hypothesis testing is:

Employed to test some assumption (hypothesis) we have about the population against a sample from the population (…) the result of a significance test is a probability which we attach to a descriptive statistic calculated from a sample. This probability reflects how likely it is that the statistic could have come from a sample drawn from the population specified in the hypothesis (Henkel, 1976: 09)⁷.

In the standard approach to significance testing, one has a null hypothesis (H₀) and an alternative hypothesis (H₁), which describe opposite and mutually exclusive patterns regarding some phenomena⁸. Usually while the null hypothesis (H₀) denies the existence of a relationship between X and Y, the alternative hypothesis (H₁) supports that X and Y are associated. For example, in a study about the determinants of corruption, while the null hypothesis (H₀) states that there is no correlation between education and corruption, the alternative hypothesis (H₁) states that these variables are correlated, or more specifically indicates the direction of the relationship; that education and corruption are negatively associated⁹.

Usually, scholars are interested in rejecting the null hypothesis in favor of the alternative hypothesis, since the alternative hypothesis represents the corroboration of the theoretical expectations of the researcher. Also, as identified by Gerber, Green and Nickerson
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(2001), there is a publication bias that favors papers that successfully reject the null hypothesis. Therefore, scholars have both substantial and practical incentives to prefer statistically significant results.

McLean and Ernest (1998: 16) argue that “a null hypothesis (H₀) and an alternative hypothesis (H₁) are stated, and if the value of the test statistic falls in the rejection region the null hypothesis is rejected in favor of the alternative hypothesis. Otherwise the null hypothesis is retained on the basis that there is insufficient evidence to reject it”. In essence, the main purpose of hypothesis test is to help the researcher to make a decision about two competing views of the reality. According to Henkel (1976),

Significance testing is assumed to offer an advantage over subjective evaluations of closeness in contexts such as that illustrated above where there are no specific criteria for what constitutes enough agreement (between our expectations and our observations) to allow us to continue to believe our hypothesis, or constitutes great enough divergence to lead us to suspect that our hypothesis is false. In a general sense, tests of significance, as one approach to assessing our beliefs or assumptions about reality, differ from the common sense approach only in the degree to which the criterion for closeness of, or correspondence between, observed and expected results are formalized, that is, specific and standardized across tests. Significance testing allows us to evaluate differences between what we expect on the basis of our hypothesis, and what we observe, but only in terms of one criterion, the probability that these differences could have occurred by ‘chance’ (Henkel, 1976: 10).

In theory, the p value is a continuous measure of evidence, but in practice it is typically trichotomized approximately into highly significant, marginally significant, and not statistically significant at conventional levels, with cutoffs at p≤0.01, p≤0.05 and p>0.10 (Gelman, 2012: 2). According to Cramer and Howitt (2004),

The level at which the null hypothesis is rejected is usually set as 5 or fewer times out of 100. This means that such a difference or relationship is likely to occur by chance 5 or fewer times out of 100. This level is generally described as the proportion 0.05 and sometimes as the percentage 5%. The 0.05 probability level was historically an arbitrary choice but has been acceptable as a reasonable choice in most circumstances. If there is a reason to vary this level, it is acceptable to do so. So in circumstances where there might be very serious adverse consequences if the wrong decision were made about the hypothesis, then the significance level could be made more stringent at, say, 1% (Cramer and Howitt, 2004: 151).

Figure 1 illustrates the logic of null hypothesis significance testing.
Figure 1. Null Hypothesis Significance Testing illustrated

We know that the area under the curve equates to 1 and can be represented by a probability density function. As we standardize the variable to a standard normal, we have a mean of zero and the spread is described by the standard deviation. Importantly, given that this curve’s standard deviation equals 1, we know that 68.26% of all observations are between -1 and +1 standard deviation, 95.44% of all observations will fall between -2 and +2 standard deviation and 99.14% of all cases are between -3 and +3 standard deviation. The shaded area represents the probability of observing a result from a sample as extreme as we observed, assuming the null hypothesis in population is true. For example, in a regression of $Y$ on $X$ the first step is to state the competing hypotheses:

$H_0$: $b_x = 0$

$H_a$: $b_x \neq 0$

While the null hypothesis states that the effect of $X$ on $Y$ is zero ($b_x = 0$), the alternative hypothesis states that the effect is different from zero ($b_x \neq 0$). The second step is to compare our estimate with the parameters specified under the null hypothesis. The more our estimate approximates to the parameters specified by the null hypothesis, the less confidence we have in rejecting it. The more distant our estimate is from the parameters specified by the null hypothesis, the more confidence we have in rejecting $H_0$ in favor of $H_a$. The p value statistic is a conditional probability, the probability of obtaining the observed or more extreme result given that the null hypothesis is true. To estimate the p value or the probability value, we should proceed as follows; (1) write down both the null ($H_0$) and the alternative hypothesis ($H_a$); (2) calculate the difference between the expected value under the null hypothesis and the observed value based on sample data; (3) standardize the difference into Z scores, and (4) estimate the probability of the alternative hypothesis assuming that the null hypothesis is true. Algebraically,
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\[ Z = \frac{\bar{x} - \mu_0}{\frac{\sigma}{\sqrt{n}}} \]

Where \( \bar{x} \) represents the observed value, \( \mu_0 \) represents the value under the null, \( \sigma \) represents the variance of the distribution and \( n \) represents the sample size (number of observations). When the difference between the observed value and the value under the null increases, all other things constant, higher is the \( Z \). Similarly, when the sample size gets bigger, all other things constant, the variance is lower and the \( Z \) is higher. The \( Z \) score is higher, and the \( p \) value statistic is lower. Therefore, the \( p \) value depends not only upon the effect magnitude but is, by definition, determined by the sample size.

To make the estimation of the \( p \) value statistic more practical to political science scholars, we use observational data from the Quality of Government Institute\(^\text{11}\). The first step is to state both the null hypothesis and the alternative hypothesis.

\[ H_0: r \geq 0 \]
\[ H_a: r < 0 \]

Theoretically, we expect a negative correlation between schooling years and corruption \((r < 0)\). Alternatively, the \( H_0 \) (null hypothesis) states that the correlation between schooling years and corruption will be higher or equal to zero \((r \geq 0)\). The \( p \) value statistic will inform the probability that the observed value is due to chance, assuming that the null hypothesis is true. We know from basic statistics classes the rule that “when the \( p \) value is low, the null hypothesis must go”. In other words, the lower the \( p \) value, the higher our confidence in rejecting the null hypothesis in favor of the alternative hypothesis. Figure 2 summarizes our empirical results.

**Figure 2. Average schooling years and Corruption**
The results suggest a strong negative (-0.799) correlation between average schooling years and corruption. The relationship is statistically significant (p value < 0.000) with a sample of 113 country-cases. Following the 0.05 criteria, we should reject the null hypothesis of no or positive association between the variables. Then, we should conclude that there is a negative relationship between average schooling years and the level of corruption.

Most statistics handbooks present a rule of thumb of 0.1, 0.05 and 0.01 significance levels. It is important to stress that these are highly arbitrary cutoffs, and that the scholar should choose between them, preferably before analyzing the data. Fisher (1923) argues that:

> If one in twenty does not seem high enough, we may, if we prefer it, draw the line at one in fifty (2 per cent point), or one in one hundred (the 1 per cent point). Personally, the writer prefers to set a low standard of significance at the 5 per cent point, and ignore entirely all results which fail to reach this level. A scientific fact should be regarded as experimentally established only if a properly designed experiment rarely fails to give this level of significance (Fisher, 1923: 85).

Through analyzing the relationship between schooling years and corruption, we established a 0.05 cutoff. Our p value is less than 0.000. To be sure, the p value is never exactly zero, but it is usual to report only three digits, therefore we interpret it as less than 0.01. So, given the null hypothesis that the correlation between schooling years and corruption is higher or equal to zero, the p value of less than 0.000 means that the probability of finding a correlation as extreme as -0.799 is less than 0.000. Therefore we should reject the null hypothesis in favor of the alternative hypothesis.

After defining what the p value means, it is important to stress what it does not. It does not mean that the likelihood that the observed results are due to chance is less than 1%. Similarly, a p value of .01 does not mean that there is a 99% chance that your results are true, and a p value of .01 does not mean that there is a 1% chance that the null hypothesis is true. Additionally, you cannot interpret it as 99% evidence that the alternative hypothesis is true12. Let’s follow Fisher’s classic definition: the p value is the probability, under the assumption of no effect (the null hypothesis $H_0$), of obtaining a result equal to or more extreme than what was actually observed (Fisher, 1925).

The next section considers situations where the p value interpretation requires caution, and presents four warnings to properly interpret the p value statistic.
Four Warnings on Significance Testing

Anscombe (1973) demonstrated the importance of exploring data graphically before drawing inferences from it. He showed that different relationships between $X$ and $Y$ can be summarized by the same statistics (F, standard error, b, beta etc). Figure 3 replicates his data, presenting four different relationships between variables with the same significant p value.

All graphs in Figure 3 gave the same p value statistic of 0.002. However, the graphical analysis shows that the nature of the relationship between the variables is markedly different. For example, the upper-right graph shows a non-linear relationship. As long the scholar only examines the p value he would never get this information. Therefore, if you rely only on statistical significance you would reject the null hypothesis for all cases,
arguing that the relationship found using sample data could be generalized to the population. In addition, the researcher may fail to reject the null hypothesis after finding significant coefficient estimates, but this could be due to an outlier effect (for example the bottom right graph in Figure 3). Basic statistics handbooks teach us that outliers can influence the probability of making Type I and Type II errors. Thus, our first warning is that before interpreting the p value statistic, scholars must graphically analyze their data. To make our case more explicitly, we examine the correlation between the Human Development Index (HDI) and corruption.

Figure 4. Correlation between Human Development Index and Corruption

In both cases the correlation is statistically significant (p value < 0.000). However, when the researcher examines only the p value he would fail to acknowledge that the relationship between the variables is best described by a quadratic function ($r^2 = 0.673$) rather than by a linear function ($r^2 = 0.457$). The practical consequence of functional form misspecification in this case is the underestimation of the magnitude of relationship between the variables. Functional form misspecification can also lead to Type I and Type II errors.

To summarize, the careful graphical depiction of data is an essential step in empirical research. Scholars must avoid interpreting the p value statistic without graphically analyzing their data first. Breaking this important rule can lead to incorrect conclusions about political phenomena.

Our second warning is that it is pointless to estimate a p value for non-random samples. According to Moore and McCabe (2006: 250), “a simple random sample (SRS) of
size n consists of n individuals from the population chosen in such a way that every set of n individuals has an equal chance to be the sample actually selected”. There are different sampling methods, considering the complexity of the population and the representativeness of different characteristics. However, regardless of the method, if the sample is not random, the underlying assumptions of both the normal distribution and central limit theorem do not hold. Thus, sample statistics are no longer unbiased and efficient estimates of population parameters. According to Smith (1983: 394), “the arguments for randomization are twofold. The first, and most important for science, is that randomization eliminates personal choice and hence eliminates the possibility of subjective selection bias. The second is that the randomization distribution provides a basis for statistical inference”.

Henkel (1976: 23) argues that “the manner in which we select samples from our populations is critical to significance testing, since the sampling procedure determines the manner in which chance factors affect the statistic(s) we are concerned with, and consequently affect the sampling distribution of the statistic”18. If the sample is random, the data is subject to the laws of probability and the behavior of estimated statistics as described by the sampling distribution. According to Moore and McCabe (2006), when you use systematic random samples to collect your data, the values of the estimated statistic neither consistently overestimate nor consistently underestimate the value of the population parameters.

Figure 5. Bias and variability

- High bias, Low variability
- High bias, High variability
- Low bias, High variability
- Low bias, Low variability
While randomization minimizes bias, the larger sample size reduces variability; the researcher is interested in producing estimates that are both unbiased and efficient (low variability) (see bottom right example).

To make this claim more practical, we intentionally selected a biased sample and then we estimated the relationship between Human Development Index (HDI) and corruption. Figure 6 summarizes this information.

**Figure 6. Correlation between Human Development Index and Corruption**

When dealing with the biased sample (n = 14), the scholar would conclude that Human Development Index (HDI) and corruption are independent variables (r = 0.018 with a p value of 0.950). However, when we consider the full sample (N = 172), we observe a negative (-0.676) and statistically significant correlation (p value < 0.000). This is to say that if we use the biased sample to infer about the population, we would not reject the null hypothesis when we should reject it (type II error). It is natural to blame the sample size when explaining a lack of statistical significance. However, this is not always the case; as long the pattern of correlation between the variables is stable, as we use a random sample, it is more likely to detect the presence of the association. Figure 7 illustrates this argument.
When dealing with a random sample of the same size (n=14), we observe a negative (-0.760) and statistically significant (p value = 0.002) correlation between the variables. In this case, when working with the random sample the scholar would reach the same conclusion based on population data.

Another potential problem is the use of a biased sample to reject the null hypothesis when it should not be rejected (type I error). Figure 8 illustrates this problem.

**Figure 8.** Correlation between Area and Average schooling years

We analyzed the correlation between the geographic area and average schooling years. While the left graph shows the raw data, the right one displays the transformed data.
In both cases, the correlation is not statistically significant. In other words, we cannot reject the null hypothesis that the estimated coefficient is equal to zero. So, we should conclude that geographical area and education are statistically independent. The following figure replicates this correlation with an income biased sample.

**Figure 8a. Correlation between Area and Average schooling years**

The result is markedly different. When dealing with the income biased sample (n=14), we observe a positive (0.671) and statistically significant (p value = 0.009) correlation between the variables. In this case, we would wrongly reject the null hypothesis (type I error). Therefore, we would conclude that geographical area and schooling years are positively associated.

Regardless of the sampling method design, we should meet the assumption of equiprobability, i.e. that each element in the population has the same chance of being selected. Non probabilistic samples cannot be used to make reliable statistical inferences. Therefore, we argue that it is pointless to interpret the p value of non-random samples.

Our third claim is that even marginal effects tend to be statistically significant when the sample size is large enough. We use basic simulation to show that, as the size of the sample increases, the power \( (b) \) of detecting a significant relationship also enhances. Hair et al. (2006: 10) argue that “power is the probability of correctly rejecting the null hypothesis when it should be rejected. Thus, power is the probability that statistical significance will be indicated if it is present”. They suggest five rules of thumb regarding statistical power analysis:
(1) Researchers should always design the study to achieve a power level of .80 at the desired significance level;

(2) More severe significance levels require larger samples to achieve the desired power level;

(3) Conversely, power can be increased by choosing a less severe alpha. 4. Smaller effects sizes always require large sample sizes to achieve the desired power, and 5. Any increase in power is most likely achieved by increasing sample size.

To make our case, we partially replicate the experiment developed by The Institute for Digital Research and Education from the University of California\(^2\). We want to compare the means of two groups on a standardized mathematic test. Computationally, we used the Stata program function `fpower` to do the power analysis\(^2\). To do so, we had to define (1) the number of groups, (2) the effect size (delta)\(^3\) and (3) the alpha level. We compared two groups with two different effect sizes (0.35 and 1.2), and we varied the alpha level in the three traditional cutoffs (0.1; 0.05 and 0.01). Figure 9 summarizes this information.

**Figure 9.** Test power and sample size

![Figure 9](image)

Our simulation results show that the larger the sample size, the higher the probability of detecting statistical significance. Similarly, the smaller the effect’s size, the greater the sample’s size should be to achieve statistical significance. In particular, when sample size approaches 250 any difference/effect is statistically significant, regardless of the alpha level. To make our argument more robust, we generated three random variables. X has a mean of zero and standard deviation of 1. X and Z correlates at .75, and X and W correlates at .10. Figure 10 illustrates their relationship.
According to Hair et al., (2006: 11), “by increasing sample size, smaller and smaller effects will be found to be statistically significant, until at very large samples sizes almost any effect is significant”. What is the meaning of a minute difference that is statistically significant? The relationship between X and W is statistically significant (p-value<0.000) but there is no substantive meaning on it. As stated by Gill (1999: 658), “political scientists are more interested in the relative magnitude of effects (...) and making only binary decisions about the existence of an effect is not particularly important”. Similarly, Luskin, (1991: 1044) points out that “to know that a parameter is almost certainly nonzero is to know very little. We care much more about its magnitude. But on this score the numbers do not speak for themselves”. Rather than focusing only on the p value, scholars should be looking at the magnitude of their estimates, the difference between group means, etc. The main conclusion is that when the sample is large enough (n>300), even marginal effects/differences tend to be statistically significant. Scholars should be aware of this fact before claiming too much about their empirical findings.

In summary, whether we maintain the effect size constant and vary the sample size (figure 9), and whether we hold the number of cases constant and vary the magnitude of the observed relationship between the variables (figure 10), these procedures have a great effect on the size of the p value statistic. In either case, scholars should be cautious before extracting too much information from it, at the risk to reach wrong substantive conclusions.

Our fourth claim is that it is pointless to estimate the p value when you are dealing with population. Analytical data from the population has increased in political science in general. Examples of this in Brazilian political science include the analysis of elections, roll-call votes and budget allocations. According to Hair et al., (2006), “a census of the
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entire population makes statistical inference unnecessary, because any differences or relationship, however small, is true and does exist” (Hair et al. 2006: 09). Introductory statistics handbooks teach us that we should use samples because they are faster and cheaper. If they are properly collected, they are also reliable. Keeping in mind that the role of inferential statistics is to draw inferences about the population from sample data, if the population is contained in all observations, estimation is not needed. There is no reason to test the hypothesis since you already know your parameters.

**Figure 11.** Population, sample and statistical inference

In our view, if the sample size is equal to the size of the population, we cannot expect any error in estimating the true population parameter. Imagine a census of the population where men are shown to have an average income of X and women an average income of X+1. The mean difference between the groups is just one unit. As long we have information about all cases (the population), there is no room for estimating the expected value of the population parameter, since we already know it. The important issue is the magnitude of the difference, which is in this case very small. Our general position is that scholars should focus on the size of the expected effects, instead of worrying about the significance of the difference, especially in the situations we have defined here.

Usually, statistical handbooks present introductory statistics based on this sample-population inference approach. However, it is important to stress that this position is not unanimous in the literature. There are differing answers to the “what does statistical p value mean when the sample equals the population” question. There is an ongoing debate between frequentist-bayesian approaches. In general, Bayesian scholars tend to reject point estimation (p value) and prefer to use confidence intervals. Some of them have even argued that confidence intervals should be abolished altogether. For example, Gelman (2012b, n.p.) states “I’m thinking more and more that we have to get rid of statistical
significance, 95% intervals, and all the rest, and just come to a more fundamental acceptance of uncertainty” (Gelman, 2012a, n.p.). Regarding the sample-population debate, Bayesian scholars also tend to believe that we should always consider the population as a sample produced by an underlying generative process. The rationale is the following: the population you examined can be seen as a result of a more complex and dynamic process that can always generate a different population in some other moment in time. However, if you ask the same question to a frequentist researcher, he will likely answer that there is no meaning in the interpretation of the p value when the sample equals the population, since we already know the true parameter value.

As long as our main purpose is to introduce the logic of the p value statistic and is based on the frequentist position, we do not examine this debate more deeply. We encourage readers to follow the references to obtain more information about this issue. Additionally, we advise students to adopt the following guidelines: (1) define your own position on the role of the p value statistic when the sample size equals the population; (2) once you pick one, you should be consistent with it, and (3) regardless of your technical position, you should always report not only the p value but also all the estimates in order to facilitate other people’s assessment of your work. As a rule, you should always report all the information since your reader may not share the same opinions as you.

Conclusion

And if my estimated coefficient is not statistically significant? God help you! Unfortunately, many scholars still share this opinion. The evidence of this claim can be found in publication bias phenomena in various fields of knowledge. Gill (1999: 669) argues that “from the current presentation of null hypothesis significance testing in published work it is very easy to confuse statistical significance with theoretical or substantive importance”. In other words, political science students should be aware of the difference between statistical significance and practical significance. The p value cannot inform us about the magnitude of the effect of X on Y. Similarly, the p value cannot help us to choose which variable explains the most. The p value cannot be compared across samples of different sizes. The p value cannot, by itself, answer the questions scholars are interested in. As noted by Moore and McCabe (2006), critical thinking about the use of significance tests is a sign of statistical maturity, however, scholars cannot make this decision when they do not fully understand the role of significance tests. Through this essay, we hope to help political science students make sense of the appropriate role of the p value statistic in empirical research.

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**APPENDIX 1**

According to King, Keohane and Verba (1994: 27), “scholars should always record the exact methods, rules, and procedures used to gather information and draw inferences so that another researcher can do the same thing and draw the same conclusion”. The main purpose of this section is to describe how the variables were measured.
Table 01. Variable Description

<table>
<thead>
<tr>
<th>Name</th>
<th>Description</th>
<th>Source</th>
<th>QOG Code</th>
</tr>
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<tbody>
<tr>
<td>Corruption</td>
<td>CPI Score relates to perceptions of the degree of corruption as seen by business people, risk analysts and the general public and ranges between 10 (highly clean) and 0 (highly corrupt). The index was inverted so that greater values equal more corruption.</td>
<td><a href="http://www.transparency.org">www.transparency.org</a></td>
<td>ti_cpi</td>
</tr>
<tr>
<td>Average Schooling Years</td>
<td>Average schooling years in the total population aged 25 and over.</td>
<td><a href="http://www.cid.harvard.edu/ciddata.ciddata.html">www.cid.harvard.edu/ciddata.ciddata.html</a></td>
<td>bl_asyt25</td>
</tr>
<tr>
<td>HDI</td>
<td>Composite index that measures the average achievements in a country in three basic dimensions of human development: a long and healthy life, as measured by life expectancy at birth, knowledge, as measured by the adult literacy rate and the combined gross enrolment ratio for primary, secondary and tertiary schools, and a decent standard of living, as measured by GDP per capita in purchasing power parity (PPP) US dollars.</td>
<td><a href="http://www.hdr.undp.org">www.hdr.undp.org</a> (UNDP, 2004)</td>
<td>undp_hdi</td>
</tr>
<tr>
<td>Area</td>
<td>A country’s total area, excluding area under inland water bodies, national claims to continental shelf, and exclusive economic zones. In most cases the definition of inland water bodies includes major rivers and lakes.</td>
<td>Food and Agriculture Organization.</td>
<td>wdi_area</td>
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Table 02. Descriptive Statistics

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<th>min</th>
<th>max</th>
<th>mean</th>
<th>standard deviation</th>
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<td>1.20</td>
<td>9.70</td>
<td>4</td>
<td>2.07</td>
</tr>
<tr>
<td>Average schooling years</td>
<td>103</td>
<td>0.76</td>
<td>12.25</td>
<td>6.02</td>
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<td>Human Development Index</td>
<td>175</td>
<td>0.27</td>
<td>0.96</td>
<td>0.70</td>
<td>0.18</td>
</tr>
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<td>Area</td>
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<td>2</td>
<td>17.098.240</td>
<td>702.337.27</td>
<td>1,937,230.60</td>
</tr>
</tbody>
</table>

Figure 12. Variable Histogram
Figure 13. Variable Histogram (log)

Notes

1. As the paper provides a non-technical introduction to the p value statistic, we minimized mathematical application of all concepts presented. Readers interested in more sophisticated approaches should follow the references. In addition, we are mute on the frequentist-bayesian
debate, since it falls outside of the context of this paper. For an online introduction to the frequentist-bayesian debate, see Jordan (2009). Readers interested in the Bayesian statistics should check Gelman, Carlin, Stern and Rubin (2003), Gill (2007) and Gelman (2007).

Publication bias represents the trend of both referees and scholars to overestimate the importance of statistical significant findings. According to Scargle (2000: 91), “publication bias arises whenever the probability that a study is published depends on the statistical significance of its results. This bias, often called the file-drawer effect because the unpublished results are imagined to be tucked away in researchers’ file cabinets, is a potentially severe impediment to combining the statistical results of studies collected from the literature”. Similarly, Everitt and Skrondal (2010: 346) define it as “the possible bias in published accounts of, for example, clinical trials, produced by editors of journals being more likely to accept a paper if a statistically significant effect has been demonstrated”. To get more information on publication bias see Greenwald (1975), Mahoney (1977), Coursoul and Wagner (1986), Simes (1986) and Begg and Berlin (1988), De Long and Lang (1992). In political science see Sigelman (1999) and Gerber, Green and Nickerson (2001). To see a simple definition see The cocharane Collaboration (n.d).

For a rebuttal of Carver arguments see Sawilowsky (2003).

Most data in political science comes from observational research designs rather than from experimental ones. For this reason, we use observational data to show the interpretation of the p value statistic when dealing with real political science research problems. As we know, observational data suffers from all sorts of shortcomings compared to experimental data. Scholars working with observational data face more challenges in making causal claims compared to those working with experiments. In addition, it is easier for a novice to understand examples based on observational data than from basic simulation.

A more systematic way of examining the role of the p value statistic in Brazilian political science literature is to survey all empirical papers and analyze how scholars have been interpreting the statistical significance of their empirical findings. The downside of this approach is the potential personal damage of exposing eventual mistakes. To minimize conflict, we prefer to focus on a more pedagogical approach.

Everitt (2006: 306) defines population as “any finite or infinite collection of ‘units’, which are often people but may be, for example, institutions, events, etc. Similarly, he defines sample as “a selected subset of a population chosen by some process usually with the objective of investigating particular properties of the parent population”.


Van Evera (1997: 09) defines a hypothesis as “A conjectured relationship between two phenomena. Like Laws, hypothesis can be of two types: causal (I surmise that A causes B) and noncausal (I surmise that A and B are caused by C; hence A and B are correlated but neither causes the other).”

The hypothesis formulation is fundamental in any empirical research. For this reason, it should be clearly stated at the beginning of the study.
Regarding figure 1, Gill (1999) argues that the test procedure assigns one of two decisions (D₀, D₁) to all possible values in the sample space of T, which correspond to supporting either H₀ or H₁ respectively. The *p value* (associated probability) is equal to the area in the tail (or tails) of the assumed distribution under H₀ which starts at the point designated by the placement of T on the horizontal axis and continues to infinity. If a predetermined α level has been specified, then H₀ is rejected for *p values* less than α, otherwise the *p value* itself is reported. Thus decision D₁ is made if the test statistic is sufficiently atypical given the distribution under the H₀ (Gill, 1999).

The description of the variables, basic descriptive statistics and distributions are presented in the appendix.

Too see this example in a cartoon see Youtube (2010).

A Type I error is the rejection of a true null hypothesis. Simply put, it is the chance of the test showing statistical significance when it is not present (false positive). The Type II error is the probability of failing to reject the null hypothesis when you should reject it (false positive) (Hair et al., 2006: 10). A more intuitive way of thinking about type I and type II errors is the following: imagine a man in a courtroom. He is not guilty (H₀). If he is convicted, the jury mistakenly rejected a true null hypothesis (type I error). Contrary, if the man is guilty (H₁) and the jury let him free, it means that they failed to reject a false null hypothesis (type II error). More details about this example can be found at Rogers (n.d).

Everitt and Skrondal (2010: 280) define misspecification as “a term applied to describe assumed statistical models which are incorrect for one of a variety of reasons, for example using the wrong probability distribution, omitting important covariates, or using the wrong link function. Such errors can produce inconsistent or inefficient estimates of parameters”.

“a probability sample is a sample chosen by chance. We must know what samples are possible and what chance, or probability, each possible sample has (…) the use of chance to select the sample is the essential principle of statistical sampling” (Moore and McCabe, 2006: 250-251).

Draw a random sample of size n from any population with mean μ and standard deviation. When n is large, the sampling distribution of the sample mean is approximately normal. Then, all properties of normal distribution apply to drawing inferences about population using sample data. According to Moore and McCabe (2006: 398), “the central limit theorem allows us to use normal probability calculations to answer questions about sample means from many observations even when the population distribution is not normal”.

By unbiased we mean that the sampling distribution of the statistic is equal to the true value of the parameter we are interested in. By efficient we mean that the estimated statistic has the lowest variability of all unbiased estimates.

The probability distribution of a statistic calculated from a random sample of a particular size. For example, the sampling distribution of the arithmetic mean of samples of size n, taken from a normal distribution with mean μ and standard deviation σ, is a normal distribution also with mean but with standard deviation \( \frac{\sigma}{\sqrt{n}} \) (Everitt, 2006: 350).

The biased sample is an intentional selection of the 20 countries with the highest Gross Domestic Product (GDP).

There are two main types of sample: the probabilistic sample and the non-probabilistic sample. Non-probability sampling is a sampling technique where the samples are gathered in a process
that does not give all the individuals in the population an equal chance of being selected. According to Davidson (2006: 15), “forms of sampling that do not adhere to probability methods. Probability methods choose samples using random selection and every member of the population has an equal chance of selection. Some types of nonrandom sampling still aim to achieve a degree of representativeness without using random methods. Several different techniques are associated with this approach, for example accidental or convenience sampling: snowball sampling; volunteer sampling; quota sampling, and theoretical sampling. Convenience samples are also known as accidental or opportunity samples. The problem with all of these types of samples is that there is no evidence that they are representative of the populations to which the researchers wish to generalize”. Simple Random Sampling: A simple random sample (SRS) of size n is produced by a scheme which ensures that each subgroup of the population of size n has an equal probability of being chosen as the sample. Stratified Random Sampling: Divide the population into “strata”. There can be any number of these. Then choose a simple random sample from each stratum. Combine those into the overall sample; this is a stratified random sample. (Example: Church A has 600 women and 400 men as members. One way to get a stratified random sample size of 30 is to take an SRS of 18 women from the 600 women and another SRS of 12 men from the 400 men.) Multi-Stage Sampling: Sometimes the population is too large and scattered for it to be practical to make a list of the entire population from which to draw an SRS. For instance, when a polling organization samples US voters, they do not do an SRS. Since voter lists are compiled by counties, they might first do a sample of the counties and then sample within the selected counties. This illustrates two stages. In some instances, they might use even more stages. At each stage, they might do a stratified random sample on sex, race, income level, or any other useful variable on which they could get information before sampling. See: http://www.ma.utexas.edu/users/parker/sampling/srs.htm. Since sampling procedures design is not the focus of this paper, we restrict ourselves to discussing the role of the p value statistic for probabilistic samples. To get more information about sampling see http://www.sagepub.com/upm-data/40803_5.pdf

21 The full description of the original simulation is available at http://www.ats.ucla.edu/stat/stata/dae/fpower.htm

22 Power analysis is the name given to the process of determining the sample size for a research study. The technical definition of power is that it is the probability of detecting a “true” effect when it exists.

23 According to Everitt and Skrondal (2010: 148), “most commonly the difference between the control group and experimental group population means of a response variable divided by the assumed common population standard deviation. Estimated by the difference of the sample means in the two groups divided by a pooled estimate of the assumed common standard deviation”.

24 According to Everitt and Skrondal (2010: 154) “The process of providing a numerical value for a population parameter on the basis of information collected from a sample. If a single figure is calculated for the unknown parameter the process is called point estimation. If an interval is calculated which is likely to contain the parameter, then the procedure is called interval estimation”.