Statistical inference and randomization

Prottoy A. Akbar

Principles of Empirical Analysis (ECON-A3000) Lecture 5

Recap of last class

- Causality: how one thing affects another thing
 - requires comparing counterfactual states of the world to each other ("how would Y change if we changed X?")
 - at most, one of them is observed
- Control group in an experimental research design
 - the outcomes of the control group are used to infer what would have happened to the treatment group in the absence of the treatment
- **Selection bias** occurs when the control group is not comparable to the treatment group, i.e. $\mathbb{E}[y_{0i}|D=0] \neq \mathbb{E}[y_{0i}|D=1]$
 - = potential outcomes differ between the treatment and control groups
- Randomization eliminates selection bias
 - on expectation, the only difference between the groups is that the treatment group gets the treatment and the control group does not
 - $\,\rightarrow\,$ differences in average outcomes must be due to the treatment

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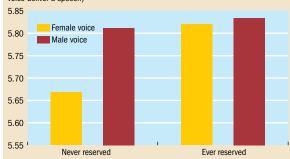
In-class discussion: Quotas in Indian local politics

- Let's start with a closer look at one of the summary figures in the summary article
 Women in Charge
 - what do we learn from this figure?

Changing minds

Indian voters perceive women leaders as less effective, but this bias diminishes with exposure to female leaders.

(rating of a *pradhan* on a scale of 1 to 10; after randomly hearing a female or male voice deliver a speech)



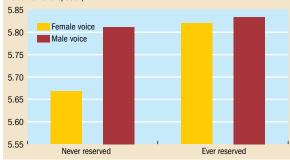
In-class discussion: Quotas in Indian local politics

- Let's start with a closer look at one of the summary figures in the summary article
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 - what do we learn from this figure?
 - would you like to have any further information before making up your mind about whether women leader truly reduce bias?

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Hypothesis testing and statistical significance

- Today's question: How likely it is that the difference between treatment and control groups could be due to chance?
 - i.e. test the null hypothesis that the treatment had no effect

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Hypothesis testing and statistical significance

- Today's question: How likely it is that the difference between treatment and control groups could be due to chance?
 - i.e. test the null hypothesis that the treatment had no effect
- Learning objectives. You understand the following concepts:
 - point estimates
 - 2 standard errors
 - g p-values
 - 4 statistical significance
 - 5 t-statistics
 - 6 critical values
 - 7 confidence intervals
 - 8 false positives and negatives (a.k.a. type I and II errors)
 - 9 statistical power (if time permits)

and how to use them to interpret basic empirical results.

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	West Bengal		
	Mean, Reserved GP	Mean, Unreserved GP	Difference
Dependent Variables	(1)	(2)	(3)
A. Village Level			<u> </u>
Number of Drinking Water Facilities	23.83	14.74	9.09
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 - ▶ first row of column (3) reports difference in averages
 - second row reports standard errors (SE)

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- This lecture: How to correctly interpret point estimates and SEs

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Point estimate and statistical significance

In the example above, we had the following sample averages

$$\bar{y}^1 = Avg[y|D=1] = 23.8$$

 $\bar{y}^0 = Avg[y|D=0] = 14.7$

where D = 1 denotes the GP being reserved for female leader

- $\bar{y}^1 \bar{y}^0 = P$ is the **point estimate**
 - the most likely impact is that, on average, P more drinking facilities are built per village when a GP is led by a woman
 - research design / identification: GPs were randomly assigned into treatment and control groups and thus selection bias is unlikely

Point estimate and statistical significance

- However, the point estimate may differ from zero because:
 - 1 female leaders are more likely to invest in drinking water
 - 2 the 54 treatment GPs just happen to invest more in drinking water (for reasons that have nothing to do with the gender of their leader)

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- Question: How likely are we to get a point estimate of at least 9.1 just due to random variation across GPs?
 - the convention is to call an estimate "statistically significant" if the likelihood of a chance finding is below 5%

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- An intuitive way to think about randomly occurring differences between groups is to create a distribution of "placebo" treatments
- Split the GPs randomly into "treatment" and "control" groups and calculate their averages
 - you can get the data here
 - ... and my simulation code on MyCourses/More Material/.

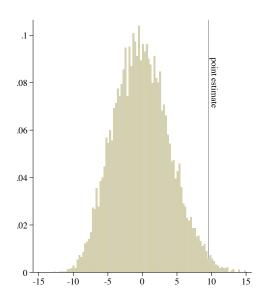
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 - ... and my simulation code on MyCourses/More Material/.
- Note that $\mathbb{E}[y|D_{pl}=1]=\mathbb{E}[y|D_{pl}=0]$
 - the "placebo" assignments D_{pl} are made-up and thus have no impact
 - but: as the table shows, with just 54 GPs in the "treatment" group, the differences can sometimes be large

$\hbox{``Treatment''}$	"Control"	Diff
15.80	19.66	-3.86
14.63	20.22	-5.59
17.10	19.03	-1.92
17.85	18.67	-0.81
13.22	20.90	-7.68
15.23	19.93	-4.70
16.91	19.12	-2.21
16.21	19.46	-3.24
21.69	16.81	4.88
19.98	17.64	2.34

10 "placebo" simulations

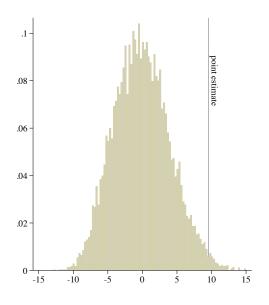
8/32

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- Simulation with 10,000 rounds
 - average: -0.099
 - standard deviation: 4.03

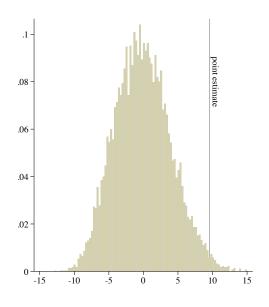
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9/32

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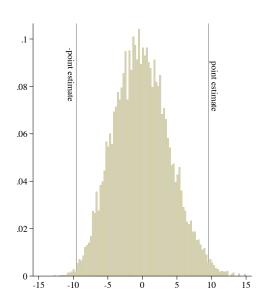


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 - the largest difference is 14.97
- However, this is quite rare:
 - difference > point estimate in 1.1% of the simulation rounds

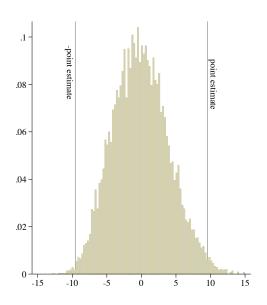
9/32

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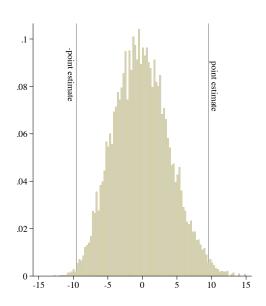
P-value



- p-value: the probability of obtaining a result at least as extreme as the result actually observed under the null hypothesis
 - here, the null hypothesis is zero treatment effect, i.e. $H_0: \mathbb{E}[y|D=1] = \mathbb{E}[y|D=0]$

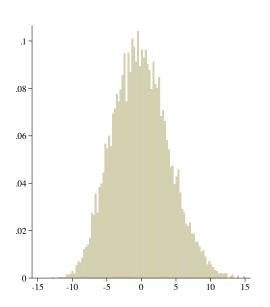


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- "2-sided" test: what is the likelihood that we'd find such a large deviation (in absolute value) from zero by chance?
 - here, the answer is 1.4%
 - by convention, estimates are called "statistically significant" (we reject the null hypothesis) if their p-value is less than 5%

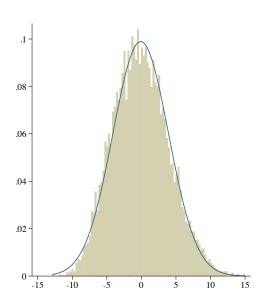


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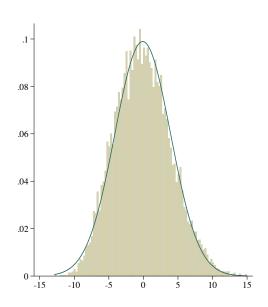
(the idea of calculating the p-value using a simulated test distribution goes back to Fisher (1935) and is now known as randomization inference)



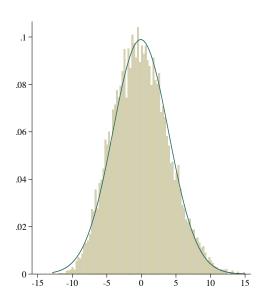
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- → We can approximate the test distribution instead of simulating it
 - saves a lot of computing time

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 - here, the statistic of interest is the treatment effect estimate (difference between treatment and control group means)

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- It summarizes the variability in the treatment effect estimate due to
 - 1 random sampling (lecture 2)
 - hence the SEs for averages in Table V
 - 2 randomness in treatment/control assignment (lecture 4)
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 - Note that even when the data includes the full population (and thus there is no random sampling), the second source of variability remains

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 We can estimate the standard error for the difference in averages between two groups with

$$\hat{SE}(\bar{y}^1 - \bar{y}^0) = S(y_i)\sqrt{\frac{1}{n_1} + \frac{1}{n_0}}$$

where $S(y_i) = \sqrt{\frac{1}{n} \sum_{i=1}^{n} (y_i - \bar{y})^2}$ is the sample standard deviation of y, and n_1 and n_0 are the number of observations in the treatment and control groups

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- many alternative estimators for SEs exists, each corresponds to different
 assumptions about the data generating process (later courses)
 (randomization inference valid for any data generating process and thus increasingly used in experimental work)
- Experiments yield more precise evidence when:
 - 1 the outcome variable has less variation [lower $S(y_i)$]
 - 2 the experiment is larger [higher n_1 and/or n_0]

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Standard errors for female leader treatment effect

• Going back to our earlier example, the corresponding numbers are

$$\hat{SE}(\bar{Y}^1 - \bar{Y}^0) = S(Y_i)\sqrt{\frac{1}{n_1} + \frac{1}{n_0}} = 18.4\sqrt{\frac{1}{54} + \frac{1}{107}} = 4.02$$

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Point estimate / standard error = the t-statistic

• Let's denote the statistic of interest with κ and its value under the null hypothesis with μ . Then the t-statistic is

$$t(\mu) = \frac{\kappa - \mu}{\mathsf{SE}(\kappa)}$$

- For treatment effects, the most common null hypothesis is H_0 : $\mu = 0$
 - under this null hypothesis, the t-value for an estimate of the average treatment effect is

$$t(0) = rac{ar{Y}^1 - ar{Y}^0}{\hat{SE}(ar{Y}^1 - ar{Y}^0)}$$

- The t-value is distributed, approximately, $t \sim \mathcal{N}(0, 1)$
 - in words: the t-value approximately follows the Normal distribution with mean zero, standard deviation one ("standard Normal distribution")

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t-statistic and significance testing

 Again, let's go back to our example and calculate the t-statistic

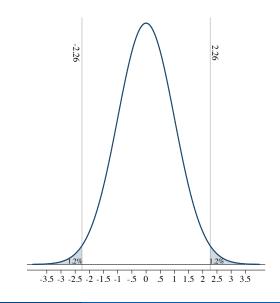
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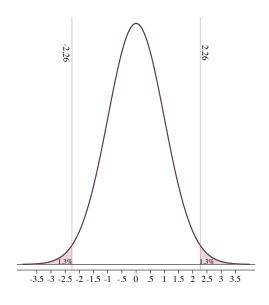
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- How exceptional would it be to draw 2.26 or more from a standard Normal distribution?
 - turns out this would happen with 1.19% probability
 - the likelihood of drawing -2.26 (or less) is also 1.19%
- \rightarrow the (two-sided) p-value is 0.0238



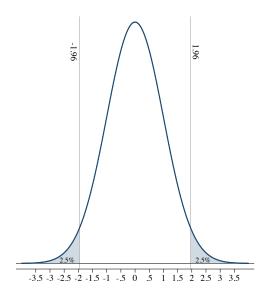
t-statistic and significance testing

- Strictly speaking, we use Student's t-distribution for calculating p-values
 - it approaches the Normal distribution when the sample size increases
- Most applications have sufficient sample size to make this distinction irrelevant
 - here, p-value increases from 0.0238 to 0.0252



Critical values and a rule-of-thumb

- Critical value is a point in the test distribution corresponding to a specific p-value
 - in large samples, a t-statistic of 1.96 corresponds to a p-value of 0.05 in a 2-sided test
- → A common rule-of-thumb is to call a result "statistically significant" if the point estimate is at least twice as large as its standard error



18 / 32

Confidence intervals

- Often the relevant question is how large/small effects we can rule out
 - instead of testing whether we can reject the null hypothesis of no effect at some confidence level (as in the previous slides)

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- We answer this using confidence intervals. For example, the 95% confidence interval is

$$[\hat{\beta} - 1.96 \times \hat{SE}, \hat{\beta} + 1.96 \times \hat{SE}]$$

where $\hat{\beta}$ is the point estimate and \hat{SE} the estimated standard error

• 1.96 corresponds to a p-value of 0.05 in a 2-sided test where the statistic (e.g. average treatment effect) is distributed $\mathcal{N}(0, 1)$ (Normal distribution with mean 0 and standard deviation 1)

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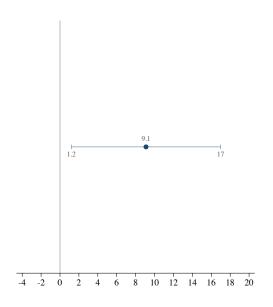
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- 1.96 corresponds to a p-value of 0.05 in a 2-sided test where the statistic (e.g. average treatment effect) is distributed $\mathcal{N}(0, 1)$ (Normal distribution with mean 0 and standard deviation 1)
- In our example, we had $\hat{\beta} = 9.1$, $\hat{SE} = 4.02 \rightarrow \text{What is the 95\% CI?}$

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19/32

Confidence intervals

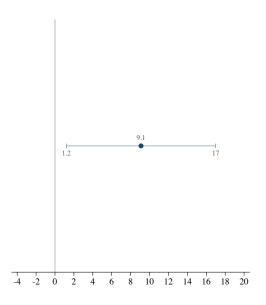


- Cls are often presented graphically
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20 / 32

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Confidence intervals



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 - e.g. the point estimate and 95% CI for our running example would look like this
- This is an informative and compact way to present results
 - but: the exact interpretation of confidence intervals is a surprisingly subtle subject
 - here, I follow Amrhein et al. (2019); most applied economists probably have this kind of an interpretation in mind

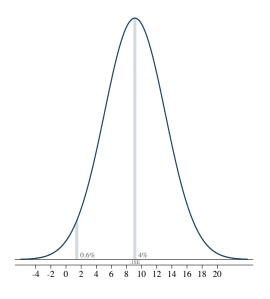
20 / 32

Interpreting confidence intervals

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- Values just outside the CI do not differ substantively from those just inside
- Not all values inside CI are equally compatible
 - point estimate is the most compatible, values near it are more compatible than those near the limits (this is the contentious part)



- The convention of dividing results to "statistically significant" and "statistically insignificant" often leads to severe misunderstandings
 - treatment is *incorrectly* thought to have been "proven to be effective" when p < .05 or "proven to have no effect" when p > .05.

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 - even if this would eventually happen, you will have to understand and interpret lots of research where statistical significance is used

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- The prevalence of such misconceptions has led to demands for abandoning the whole concept of statistical significance
 - even if this would eventually happen, you will have to understand and interpret lots of research where statistical significance is used
- No-one demands abandoning p-values and confidence intervals!
 - rather, the debate is about the misleading and unnecessary dichotomy between "significant" and "insignificant" results

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Result of an experiment

	recuity	
	Effect	No effect
Effect	True positive	False positive
No	False negative	True negative
efect		

Reality

23 / 32

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23/32

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 - also known as "type I error" or "acceptance error"
- False negative: Not finding an effect when it does exist
 - a.k.a. "type II error" or "rejection error"
- Power: the probability of finding an effect when it exists

Testing errors

Type I error (false positive) You're pregnant

Type II error (false negative) You're not pregnant

Source: Effect size FAQs

- Statistical significance testing is built to avoid false positives
 - we typically call estimates "statistically significant" if p < .05
 - i.e. if there was no effect, differences as extreme as the one we observed between treated/control would occur less than 1 out of 20 times
- Trade-off between false positives and false negatives
 - efforts to reduce one type of error increase the likelihood of other error

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 - 3 replace everyone's income in the treatment group with $y_i + \beta$, where y_i is individual i's true income and β is the simulated treatment effect

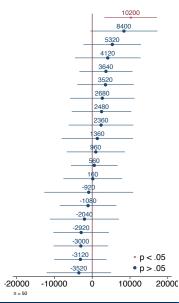
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 - 3 replace everyone's income in the treatment group with $y_i + \beta$, where y_i is individual i's true income and β is the simulated treatment effect
 - 4 calculate difference in average income between treatment and control groups and test for its statistical signficance

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 - 5 repeat many times and summarize the results

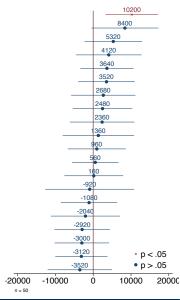
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 - 5 repeat many times and summarize the results
- ullet Let's start with the case where the treatment has no impact (eta=0)
 - question: among the false positives, how should we expect the estimated size of the effect to vary with sample size?

Prottoy A. Akbar 5: Statistical Inference Empirical Analysis 26 / 32



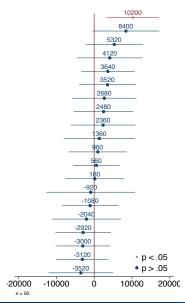
- Here are 20 simulations with n = 50
 - 25 persons in treatment, 25 in control

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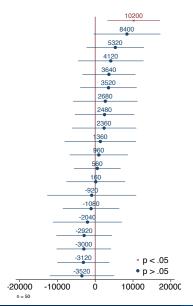


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 - 25 persons in treatment, 25 in control
- 1 out of 20 is a false positive
 - exactly what one should expect when using p < .05 as the criterion for significance

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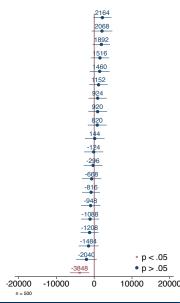
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 - the false positive result suggests that this "treatment" increased income by 10,200 euros or 0.7 standard deviations
- All confidence intervals include large effects
 - 95%Cl average width is 16,000 euros!
 - \rightarrow correct conclusion: we learn very little with n=50 (note that this is due to large variation in income; for less variable outcomes n=50 might be sufficient for meaningful analysis)

Prottoy A. Akbar 5: Statistical Inference Empirical Analysis 27 / 32

False positives with larger samples

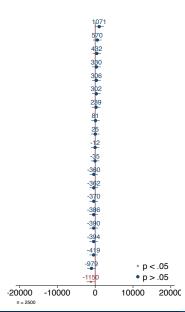


- 20 simulations with n = 500
 - again, one happens to be a false positive
- Now, the point estimate for the false positive is less spectacular
 - none of the estimates is close to 10,000

28 / 32

CI average width is 5,000 euros

False positives with larger samples

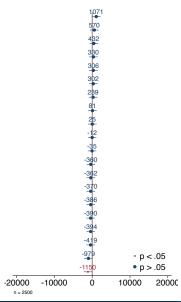


- 20 simulations with n = 2500
 - even less spectacular false positive
 - and still tighter confidence intervals (CI average width is 2,300 euros)

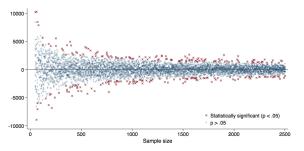
29/32

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False positives with larger samples



- 20 simulations with n = 2500
 - even less spectacular false positive
 - and still tighter confidence intervals (CI average width is 2,300 euros)
- More simulations
 - 20 rounds for 50,60,...,2500 observations
 - 0–5 false positives per round
 - overall 5.2% of simulations false positive



29 / 32

Take-aways from the first simulation

- The likelihood of a false positive does not vary with sample size
 - by definition, depends only on the p-value required for calling the esimate statistically significant (significance level)

Prottoy A. Akbar 5: Statistical Inference Empirical Analysis 30 / 32

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- The likelihood of a false positive does not vary with sample size
 - by definition, depends only on the p-value required for calling the esimate statistically significant (significance level)
- Small samples lead to large point estimates for false positives
 - small sample \rightarrow wide CI \rightarrow only large estimates significant
 - thus false positives from small samples may cause more damage
 - policy mistakes more likely if the effects are believed to be large
 - sadly, few people understand the dangers of underpowered studies

Prottoy A. Akbar 5: Statistical Inference **Empirical Analysis**

30 / 32

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 - policy mistakes more likely if the effects are believed to be large
 - sadly, few people understand the dangers of underpowered studies
 - results from small samples sometimes get huge media attention
 - unfortunately, editors and referees of scientific journals may also like spectacular and statistically signficant results

Summary

- Standard error is the standard deviation of a statistic
 - tells how *precise* our point estimate is
 - estimates become more precise (smaller SE) as the sample size increases or variation in the outcome variable decreases
- **p-value** is the probability of obtaining a result at least as extreme as the result actually observed if the null hypothesis is true
 - convention to call results "statistically significant" if p < .05
 - corresponds to |point estimate| $\geq 2 \times \text{standard error}$
- Confidence interval includes values most compatible with the data
 - the point estimate is the most compatible value
- False positives

Prottoy A. Akbar 5: Statistical Inference Empirical Analysis 31/32

Upcoming

- Pre-class assignment 5
 - Moving to Opportunity experiment!
 - Read and summarize an article

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- Deadline: Jan 24 at 13:00
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- Leave yourself time to deal with unexpected technical issues.

Prottoy A. Akbar 5: Statistical Inference Empirical Analysis 32 / 32

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- Use the course Slack channel to seek help and help others in the class
 - Quicker than waiting for private responses from the TA or me
 - Recall extra incentive: bonus points for active participation

Prottoy A. Akbar 5: Statistical Inference Empirical Analysis 32 / 32