Day 2: On Weights and Clusters

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Design-Based Regression Inference Spring 2024

Outline

- 1. Heterogeneous Treatment Effects
- 2. Clustered Standard Errors

- On Monday we contrasted design vs. outcome-model strategies in a constant-effect world (i.e. with a causal model of $y_i = \beta x_i + \varepsilon_i$)
 - Of course the real world is messier: more realistic is $y_i = \beta_i x_i + \varepsilon_i$ (or more complicated forms of effect heterogeneity)

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 - Bottom line: design avoids recent concerns over "negative weights"...
 - ... at least as long as you don't have multiple treatments!

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- Why is this a concern? The possibility of sign reversals:
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- ullet It turns out that such ψ_i also arise in design-based specifications, and they can also be negative
 - But sign reversals are impossible in design-based specs: then we also have $\beta = E[\phi_i \beta_i]/E[\phi_i]$ for "ex-ante" ϕ_i which are always non-negative

Simple Setup

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$$y_i = \beta x_i + w_i' \gamma + e_i$$

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- Assume appropriate asymptotics for OLS to consistently estimate:

$$\beta = \frac{E[\tilde{x}_i y_i]}{E[\tilde{x}_i^2]} = \frac{E[\tilde{x}_i x_i \beta] + E[\tilde{x}_i \varepsilon_i]}{E[\tilde{x}_i^2]}$$

where \tilde{x}_i are residuals from the population regression of x_i on w_i

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ASSUMPTION 1: $E[\varepsilon_i \mid x_i, w_i] = w_i' \gamma$

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ASSUMPTION 2: $E[x_i \mid \varepsilon_i, \beta_i, w_i] = w_i' \lambda$

- Treatment is conditionally mean-independent of potential outcomes, with a linear expected treatment $E[x_i \mid w_i]$ (e.g. the propensity score)
- E.g. a stratified experiment, where x_i is randomly assigned within strata dummied out in w_i
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- The second assumption yields a design-based OLS specification
 - Stronger (sufficient) condition: $x_i \mid (\varepsilon_i, \beta_i, w_i) \stackrel{iid}{\sim} G(w_i)$

• Since $E[\tilde{x}_i \varepsilon_i] = 0$, the OLS estimand has an average-effect representation under either assumption:

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- ullet The ex-post weights are the end of the story for eta under Assumption
 - 1. But in design-based specifications we can take one more step
 - In experiments, who is in the effective control group is *random*. Before treatment is drawn, everyone expects the same weight!

• Using the law of iterated expectations, we can also write:

$$\beta = \frac{E[E[\psi_i \mid w_i, \beta_i]\beta_i]}{E[E[\psi_i \mid w_i, \beta_i]]} \equiv \frac{E[\phi_i \beta_i]}{E[\phi_i]}$$

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- Hence: sign reversals cannot occur in design-based OLS specifications

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 - Could inverse-weight by $\widehat{Var}(x_i \mid w_i)$ to estimate unweighted $E[\beta_i]$
- Of course, the ϕ_i -weighted estimand may not be most of interest!
 - If $Cov(\phi_i,\beta_i) \approx 0$, we'll still get something close to $E[\beta_i]$
 - Otherwise, ϕ_i -weighting has desirable efficiency properties (Goldsmith-Pinkham et al. 2024)
 - Large class of alternative propensity-score-based estimators for other estimands under the stronger design assumption

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 - **1** A more general causal model: potential outcomes $y_i(x)$ and $y_i = y_i(x_i)$
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- For convex ex-ante weights in IV we require first-stage monotonicity: that x_i is non-decreasing in z_i for all units regardless of $y_i(\cdot)$
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 - Ex post weights are still potentially non-convex under monotonicity
- Framework is general, allowing for "formula" IVs (e.g. shift-share)
 - We'll see more about this in Friday's class

Multiple Treatments: Contamination Bias

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 - A weighted average of treatment j's effects, with convex weights in design-based specifications √
 - A non-convex combination of effects from other treatments k ("contamination bias") X
- They derive alternative estimators which avoid contamination bias while maintaining the nice efficiency properties of OLS weighting
 - Ultimately, becomes an empirical question of how important bias is

Example: Project STAR

- Krueger (1999) studies the STAR RCT, which randomized 12k students in 80 public elementary schools in Tennessee to one of 3 classroom types:
 - Regular-sized (20-25 students) Control
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- Kids were randomized within schools, so the propensity of assignment to each treatment varied by school
 - Krueger thus estimates: $TestScore_i = \alpha_{school(i)} + \beta_1 D_{i1} + \beta_2 D_{i2} + \varepsilon_i$

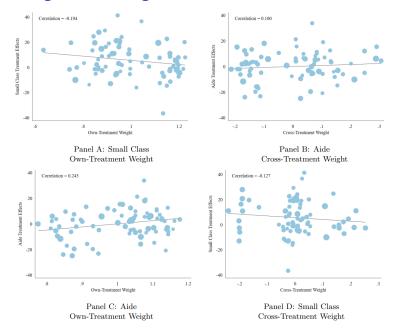
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- We find significant potential for contamination bias: lots of treatment effect heterogeneity and variation in contamination weights
 - But actual contamination bias is minimal: $Corr(effects, weights) \approx 0$

Project STAR, Revisited

	A. Contamination Bias Estimates				
	Regression	Own	Bias	Worst-Case Bias	
	Coefficient	Effect		Negative	Positive
	(1)	(2)	(3)	(4)	(5)
Small Class Size	5.357	5.202	0.155	-1.654	1.670
	(0.778)	(0.778)	(0.160)	(0.185)	(0.187)
Teaching Aide	0.177	0.360	-0.183	-1.529	1.530
	(0.720)	(0.714)	(0.149)	(0.176)	(0.177)
	B. Treatment Effect Estimates				
		Unweighted	Efficiently-Weighted		
		(ATE)	One-at-a-time	Common	
		(1)	(2)	(3)	
Small Class Size		5.561	5.295	5.563	
		(0.763)	(0.775)	(0.764)	
		[0.744]	[0.743]	[0.742]	
Teaching Aide		0.070	0.263	-0.003	
		(0.708)	(0.715)	(0.712)	
		[0.694]	[0.691]	[0.695]	

STAR Regression Weights vs. Treatment Effects



Outline

1. Heterogeneous Treatment Effects ✓

2. Clustered Standard Errors

• Where do SEs come from? OLS $\hat{\beta} = (\mathbf{X}'\mathbf{X})^{-1}\mathbf{X}'\mathbf{Y}$ can be rewritten:

$$\sqrt{N}(\hat{\beta} - \beta) = \left(\frac{\mathbf{X}'\mathbf{X}}{N}\right)^{-1} \left(\frac{\mathbf{X}'\boldsymbol{\varepsilon}}{\sqrt{N}}\right)$$

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$$\beta^* \sim N(\beta, \frac{V}{N}), \quad V = E\left[\frac{1}{N}\sum_i X_i X_i'\right]^{-1} Var\left(\frac{1}{\sqrt{N}}\sum_i X_i \varepsilon_i\right) E\left[\frac{1}{N}\sum_i X_i X_i'\right]^{-1}$$

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$$\beta^* \sim \mathrm{N}(\beta, \frac{V}{N}), \quad V = E\left[\frac{1}{N}\sum_i X_i X_i'\right]^{-1} \mathit{Var}\left(\frac{1}{\sqrt{N}}\sum_i X_i \varepsilon_i\right) E\left[\frac{1}{N}\sum_i X_i X_i'\right]^{-1}$$

• SEs come from $\hat{V} = \left(\frac{1}{N}\sum_i X_i X_i'\right)^{-1} \widehat{Var} \left(\frac{1}{\sqrt{N}}\sum_i X_i \varepsilon_i\right) \left(\frac{1}{N}\sum_i X_i X_i'\right)^{-1}$

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 - We need to zero out some covariances to make progress

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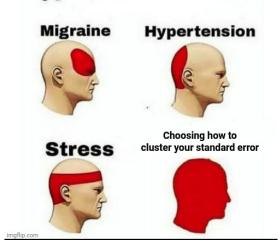
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This is what's going on under the hood when you ", cluster(c)"!

Easy, Right?

Types of Headaches



Source: Khoa Vu (of course)

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 - Suppose $(X_1, \ldots, X_N) \mid (\varepsilon_1, \ldots, \varepsilon_N)$ is mean-zero with $X_i \perp X_j$ whenever $c(i) \neq c(j)$ (e.g. village-level RCT with c(i) giving i's village)

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- So we only need to cluster by c(i): the design tells us what to do!
- This leads to the popular (and sometimes misused) heuristic: cluster at the level of treatment / identifying variation
 - See Abadie et al. (2023) for a more complete version of this argument

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