

Day 2: On Weights and Clusters

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Design-Based Regression Inference
Fall 2024

Outline

1. Heterogeneous Treatment Effects
2. Clustered Standard Errors

Whose Treatment Effect is it Anyway?

- 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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- Let $x_i \in \{0,1\}$; general causal model: $y_i = \underbrace{(y_i(1) - y_i(0))}_{\beta_i} x_i + \underbrace{y_i(0)}_{\varepsilon_i}$
 - Design: $x_i \mid w, y(0), y(1) \sim F_x(w_i)$ with linear $E[x_i \mid w_i]$

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- Hence the regression a proper (convex) weighted avg. of the β_i :

$$\beta = \frac{E[\text{Var}(x_i \mid w) \beta_i]}{E[\text{Var}(x_i \mid w)]}$$

More weight put on observations with more treatment variability

Primer 2: TWFE with Staggered Adoption

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 - Units (non-randomly) adopt treatment over time: $x_{it} = \mathbf{1}[t \geq g_i]$ where $g_i \in \{1, \dots, T\} \cup \infty$ gives adoption time ($g_i = \infty$ for never treated)

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- But notice something a bit weird here: we can run this regression even if there are no never-treated units ...
 - How, then, is the regression using parallel trends in $y_{it}(0)$?

Simple Staggered Adoption

- Consider $T = 2$ and two groups: *always-treated* units (with $g_i = 1$; $x_{i1} = x_{i2} = 1$) and *switchers* (with $g_i = 2$; $x_{i1} = 0$, $x_{i2} = 1$)

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- Under PT, $E[y_{i2}(0) - y_{i1}(0) | g_i = 1] = E[y_{i2}(0) - y_{i1}(0) | g_i = 2]$ so:

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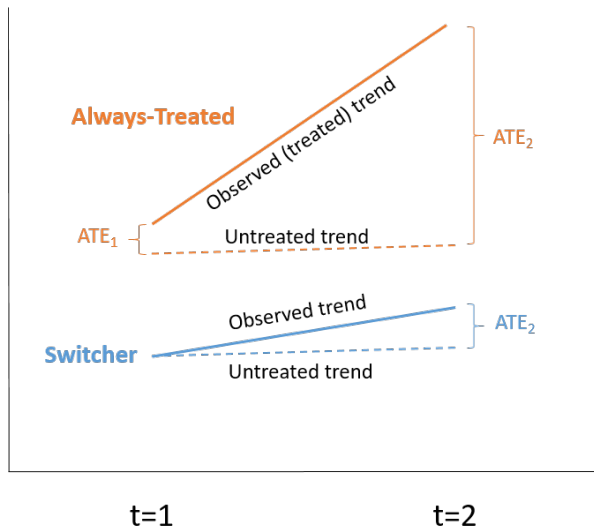
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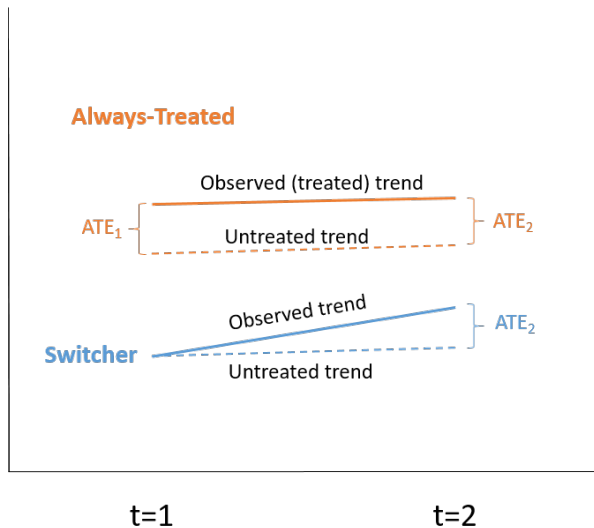
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“Forbidden Comparisons,” Illustrated



No Problem Under Constant Effects



Why So Negative?

- We can write the previous expression as a *non-convex* weighted average of $\beta_{it} = y_{it}(1) - y_{it}(0)$:

$$\begin{aligned}\beta &= E[\beta_{i2} \mid g_i = 2] + E[\beta_{i2} \mid g_i = 1] - E[\beta_{i1} \mid g_i = 1] \\ &= \frac{E[\psi_{it}\beta_{it}]}{E[\psi_{it}]} \quad \text{for } \psi_{it} = \begin{cases} +, & \text{if } t=2 \\ 0, & \text{if } t=1, g=2 \\ -, & \text{if } t=1, g=1 \end{cases}\end{aligned}$$

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 - The recent TWFE literature points this issue out in many settings and proposes alternative specifications / procedures to address it
- 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 can also write $\beta = E[\phi_i\beta_i]/E[\phi_i]$ for “ex-ante” ϕ_i which are non-negative

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- In large enough samples, OLS consistently estimates:

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

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

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- The second assumption yields a design-based OLS specification
 - Stronger (sufficient) condition: $x_i \mid (\varepsilon_i, \beta_i, w_i) \stackrel{iid}{\sim} F_x(w_i)$

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 - This can lead to sign reversals: e.g. $\beta < 0$, despite $\beta_i > 0$

Ex-Post Weights

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

$$\beta = \frac{E[\psi_i \beta_i]}{E[\psi_i]}, \quad \psi_i = \tilde{x}_i x_i$$

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 - This can lead to sign reversals: e.g. $\beta < 0$, despite $\beta_i > 0$
- The ex-post weights are the end of the story for β 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!

Ex-Ante Weights

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

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

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

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 - Could inverse-weight by $\widehat{\text{Var}}(x_i \mid w_i)$ to estimate unweighted $E[\beta_i]$
- Of course, the ϕ_i -weighted estimand may not be most of interest!
 - If $\text{Cov}(\phi_i, \beta_i) \approx 0$, 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

General Setting

- Borusyak and Hull (2024) extend ex ante / ex post weights to:
 - ① A more general causal model: potential outcomes $y_i(x)$ and $y_i = y_i(x_i)$
 - ② IV: design-based assumption is then $E[z_i | y_i(\cdot), w_i] = w_i' \lambda$

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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) where the first stage relationship need not be causal
 - We'll see more about this in tomorrow's class

Special Case: IV with Linear Heterogeneity

- Suppose $y_i = \beta_i x_i + \varepsilon_i$ (without loss of generality for binary x_i). Then:

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- Reduces to Imbens-Angrist LATE result if the first stage is causal

Multiple Treatments: Contamination Bias

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- We derive alternative estimators which avoid contamination bias while maintaining some nice properties of OLS weighting
 - Ultimately, becomes an empirical question of how important bias is

General Problem

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$$y_i = \sum_k x_{ik} \beta_k + g(w_i) + u_i$$

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- They show each regression coefficient β_k can be decomposed:

$$\beta_k = E[\lambda_{kk}(w_i) \tau_k(w_i)] + \sum_{\ell \neq k} E[\lambda_{k\ell}(w_i) \tau_\ell(w_i)]$$

for $\tau_k(w_i) = E[y_i(k) - y_i(0) | w_i]$, $\lambda_{kk} = \frac{E[\tilde{x}_{ik} x_{ik} | w_i]}{E[\tilde{x}_{ik}^2]}$, $\lambda_{k\ell} = \frac{E[\tilde{x}_{ik} x_{i\ell} | w_i]}{E[\tilde{x}_{ik}^2]}$;

\tilde{x}_{ik} is the residual from regressing x_{ik} on $g(w_i)$ and all other $x_{i,-k}$

- $E[\lambda_{kk}(w_i)] = 1$, $E[\lambda_{k\ell}(w_i)] = 0$. Further $\lambda_{kk}(w_i) \geq 0$ if $g(\cdot)$ spans p_k

Unpacking The Result

$$\beta_k = \underbrace{E[\lambda_{kk}(w_i)\tau_k(w_i)]}_{\text{Own treatment effect}} + \sum_{\ell \neq k} \underbrace{E[\lambda_{k\ell}(w_i)\tau_\ell(w_i)]}_{\text{Contamination bias}}$$

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 - FWL residual \tilde{x}_{ik} is thus not mean-zero given $(w_i, x_{i,-k})$, so it “picks up” effects of other treatments x_{ik} given w_i

Is This a Problem?

- In principle, contamination bias applies to a large number of settings:
 - ① RCTs with multiple treatments and randomization strata
 - ② Selection-on-obs with multiple treatments (e.g. “value-added” models)
 - ③ TWFE with multiple treatments (e.g. “mover” regressions)
 - ④ IV with multiple instruments (e.g. “examiner/judge” IVs)
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 - ⑤ Descriptive regressions on multiple variables (e.g. disparity analyses)
- But again, the severity of the issue is an empirical matter
 - Since the CB weights average to zero, if they're uncorrelated with effect heterogeneity there's no issue
 - The weights are identified; we can estimate them to diagnose bias

Solutions

- Contamination bias comes from the FWL auxiliary regression not controlling “flexibly enough” for $(w_i, x_{i,-k})$... but we can fix that:

$$y_i = \sum_k x_{ik} \beta_k + g(w_i) + \sum_k x_{ik} (q_k(w_i) - E[q_k(w_i)]) + u_i$$

The **blue term** captures non-linearities in (w_i, x_i)

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The **blue term** captures non-linearities in (w_i, x_i)

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- This works in principle, but in practice can fail / be really noisy
 - Key challenge: limited overlap ($p_k(w_i)$ may be close to zero or one)
 - If CB is limited, an uninteracted regression is likely more efficient...

Solutions (Cont.)

- We could of course instead just focus on one treatment at a time:

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just using observations where either $x_{ik} = 1$ or $x_{i0} = 1$

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- As before, whether any of these alternatives give a different answer than OLS is an empirical matter...

Example: Project STAR

- Krueger (1999) studies the STAR RCT, which randomized students in public elementary to one of three classroom types:
 - 1 Regular-sized (20-25 students) – Control
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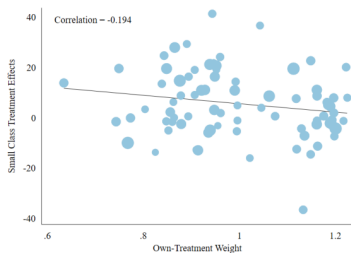
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- We find significant *potential* for contamination bias: lots of treatment effect heterogeneity and variation in contamination weights
 - But actual contamination bias is minimal: $Corr(effects, weights) \approx 0$

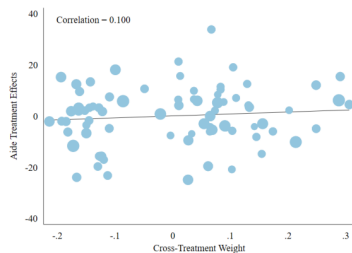
Project STAR, Revisited

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

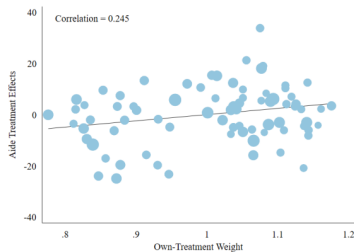
STAR Regression Weights vs. Treatment Effects



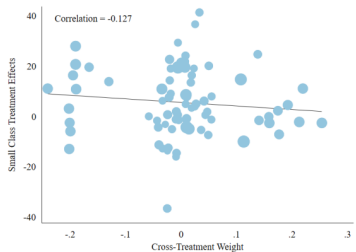
Panel A: Small Class
Own-Treatment Weight



Panel B: Aide
Cross-Treatment Weight



Panel C: Aide
Own-Treatment Weight



Panel D: Small Class
Cross-Treatment Weight

Does Contamination Bias Ever Matter?

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 - Intuitively, experimental strata are unlikely to strongly predict TE heterogeneity (variation driven by experimenter constraints, etc.)

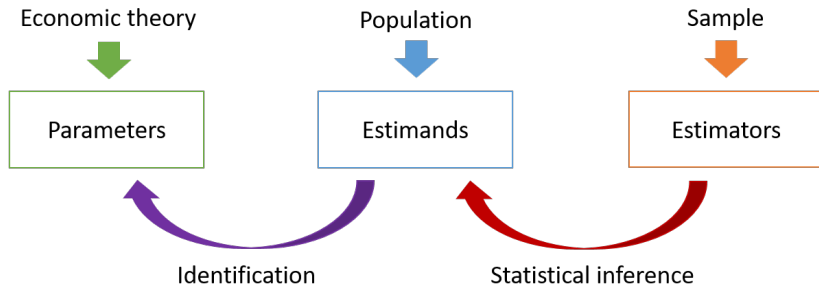
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 - Intuitively, experimental strata are unlikely to strongly predict TE heterogeneity (variation driven by experimenter constraints, etc.)
- Practical takeaway: bias diagnostics can be useful, especially in observational analyses (use our *multe* package!)

Outline

1. Heterogeneous Treatment Effects✓
2. Clustered Standard Errors

Journey to the Red Arrow...



OLS Asymptotics: Review

- 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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 - We need to zero out some covariances to make progress

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- This is what's going on under the hood when you “, *cluster(c)*”!

Easy, Right?

Types of Headaches

Migraine



Hypertension



Stress



Choosing how to
cluster your standard error



imgflip.com

Source: Khoa Vu

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- At an (unhelpfully) high level, the previous results tell us when to cluster i and j together: when we think $\text{Cov}(x_i \varepsilon_i, x_j \varepsilon_j) \neq 0$
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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

Where Intuition Can Fall Short: Paired Randomization

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- Paired randomization makes x_i and x_j *negatively* correlated in pairs
 - Clustering by pair solves this; treatment assignment is *iid* across pairs