

# Frontier Topics in Empirical Economics: Week 11

## Standard Error Issues

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# Introduction: Nonstandard Standard Error Issues

- Inference is important in practice: Data  $\Rightarrow$  Target distribution
- How accurate is our estimate? How confident are we on our results?
- In traditional inference of econometrics, we have two assumptions:
  - Uncertainty comes from random sampling, asymptotics when  $n \rightarrow \infty$
  - $y_i$  iid, sample, no correlations
- What if these two assumptions are violated?

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- In this lecture, we consider two cases
- First, when  $n$  is naturally limited (e.g. number of provinces)
- Another type of uncertainty becomes important: Design-based uncertainty
- Second, when i.i.d. fails and errors are clustered
- We have to incorporate this structure in inference
- Angrist calls them "Nonstandard Standard Error Issues"

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- In usual case, when we talk about inference, what is that?
- We have a target parameter: “*estimand*”  $\beta$  (Target)
- We want to recover it using an “*estimator*” (Method)  $\hat{\beta}$  with a sample from the population, which gives you a result called “*estimate*”  $\hat{\beta} = 0.5$  (Result)
- This process is called *estimation*, or statistical inference (Process)



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- When sample changes, your estimation result changes
- Uncertainty comes from sampling process
- Thus, you have a standard error for your estimation
- But is this the only uncertainty in empirical research?
- Today, we are going to introduce the second source of uncertainty

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- Design-based uncertainty, introduced by Abadie et al. (2020)
- It is the uncertainty coming from the treatment assignment
- Treatment  $X_i$  is no longer considered fixed
- In some cases, person 1 is treated; in other cases, person 1 is not treated
- The potential outcome you observed is different when treatment is randomly changed
- We show that this helps you to understand uncertainty of estimation when you have non-negligible sample size

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- Let's take a look at two tables from Abadie et al. (2020)
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## ■ Sampling-based uncertainty

TABLE I  
SAMPLING-BASED UNCERTAINTY (✓ IS OBSERVED, ? IS MISSING)

Unit	Actual Sample			Alternative Sample I			Alternative Sample II			...
	$Y_i$	$Z_i$	$R_i$	$Y_i$	$Z_i$	$R_i$	$Y_i$	$Z_i$	$R_i$	...
1	✓	✓	1	?	?	0	?	?	0	...
2	?	?	0	?	?	0	?	?	0	...
3	?	?	0	✓	✓	1	✓	✓	1	...
4	?	?	0	✓	✓	1	?	?	0	...
⋮	⋮	⋮	⋮	⋮	⋮	⋮	⋮	⋮	⋮	...
$n$	✓	✓	1	?	?	0	?	?	0	...



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1	✓	✓	1	?	?	0	?	?	0	...
2	?	?	0	?	?	0	?	?	0	...
3	?	?	0	✓	✓	1	✓	✓	1	...
4	?	?	0	✓	✓	1	?	?	0	...
⋮	⋮	⋮	⋮	⋮	⋮	⋮	⋮	⋮	⋮	...
$n$	✓	✓	1	?	?	0	?	?	0	...

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TABLE II  
DESIGN-BASED UNCERTAINTY (✓ IS OBSERVED, ? IS MISSING)

Unit	Actual Sample			Alternative Sample I			Alternative Sample II			...
	$Y_i^*(1)$	$Y_i^*(0)$	$X_i$	$Y_i^*(1)$	$Y_i^*(0)$	$X_i$	$Y_i^*(1)$	$Y_i^*(0)$	$X_i$	...
1	✓	?	1	✓	?	1	?	✓	0	...
2	?	✓	0	?	✓	0	?	✓	0	...
3	?	✓	0	✓	?	1	✓	?	1	...
4	?	✓	0	?	✓	0	✓	?	1	...
⋮	⋮	⋮	⋮	⋮	⋮	⋮	⋮	⋮	⋮	...
$n$	✓	?	1	?	✓	0	?	✓	0	...

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2	?	✓	0	?	✓	0	?	✓	0	...
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- Treatment is fixed, sampling observation is random
- For non-sampled individuals, we cannot observe anything
- Source of uncertainty: in each sample, we have different observations

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- Treatment is random, sampling observation is fixed (e.g., all provinces in China)
- For each individual, we only observe potential outcome in the realized status (but not counterfactual status)
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- Next, the authors construct a simple model and make the following four points:
  - 1. Show how design-based uncertainty affects the variance of the regression estimator
  - 2. Show White estimator remains conservative when we consider design-based uncertainty
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  - 4. Discuss two sources of uncertainty and external/internal validity

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- Assume that we have a **finite** population of size  $n$
- We randomly sample  $N$  from  $n$
- $R_i \in \{0, 1\}$  as an indicator of whether  $i$  is sampled or not
- There is a random binary treatment regressor  $X_i$
- $n_1, N_1$  are treated,  $n_0, N_0$  are not treated
- We have observed and potential outcome as:

$$Y_i = Y_i^*(X_i) = \begin{cases} Y_i^*(1) & \text{if } X_i = 1, \\ Y_i^*(0) & \text{if } X_i = 0 \end{cases}$$

- Potential outcomes are assumed to be non-stochastic

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- Potential outcomes are assumed to be non-stochastic



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# Design-based Uncertainty

- We use bold letters to represent vector of the whole sample  $(\mathbf{Y}, \mathbf{Y}_i^*(1), \mathbf{Y}_i^*(0), \mathbf{R})$
- We define three estimands as our proposed targets
  - Descriptive estimand: free of  $\mathbf{R}$  and potential outcomes (population mean difference)
$$\theta^{descriptive} = \frac{1}{N} \sum_{i=1}^N X_i Y_i - \frac{1}{N} \sum_{i=1}^N (1 - X_i) Y_i$$
  - Causal estimand: parameter depending on potential outcomes:  $Y_i^*(1), Y_i^*(0)$ 
$$\theta^{causal, population} = \frac{1}{N} \sum_{i=1}^N X_i (Y_i^*(1) - Y_i^*(0))$$
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- To estimate these estimands, we use a simple OLS regression of  $Y_i$  on  $X_i$  to have:

$$\hat{\theta} = \frac{1}{N_1} \sum_{i=1}^n R_i X_i Y_i - \frac{1}{N_0} \sum_{i=1}^n R_i (1 - X_i) Y_i$$

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- It is shown that:

$$E[\hat{\theta}|\mathbf{X}, N_1, N_0] = \theta^{descr}$$

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# Design-based Uncertainty

- We define the population variances as follows:

$$S_x^2 = \frac{1}{n-1} \sum_{i=1}^n \left( Y_i^*(x) - \frac{1}{n} \sum_{j=1}^n Y_j^*(x) \right)^2, \text{ for } x = 0, 1$$

$$S_\theta^2 = \frac{1}{n-1} \sum_{i=1}^n \left( Y_i^*(1) - Y_i^*(0) - \frac{1}{n} \sum_{j=1}^n (Y_j^*(1) - Y_j^*(0)) \right)^2$$

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# Design-based Uncertainty

- Based on the defined population variance, we can derive three variances

$$V^{total}(N_1, N_0, n_1, n_0) = \text{var}(\hat{\theta} | N_1, N_0) = \frac{S_1^2}{N_1} + \frac{S_0^2}{N_0} - \frac{S_{\theta}^2}{n_0 + n_1}$$

$$V^{sampling}(N_1, N_0, n_1, n_0) = E[\text{var}(\hat{\theta} | \mathbf{X}, N_1, N_0) | N_1, N_0] = \frac{S_1^2}{N_1} \left(1 - \frac{N_1}{n_1}\right) + \frac{S_0^2}{N_0} \left(1 - \frac{N_0}{n_0}\right)$$

$$V^{design}(N_1, N_0, n_1, n_0) = E[\text{var}(\hat{\theta} | \mathbf{R}, N_1, N_0) | N_1, N_0] = \frac{S_1^2}{N_1} + \frac{S_0^2}{N_0} - \frac{S_{\theta}^2}{N_0 + N_1}$$

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- $V^{total}$  is the total variance, considering both sampling-based and design-based uncertainty:  $var(\hat{\theta}|N_1, N_0)$
- It is the variance we want to capture in inference for causal estimator
- $V^{sampling}$  is the variance from only sampling-based uncertainty, by conditioning on treatment assignment:  $E[var(\hat{\theta}|\mathbf{X}, N_1, N_0)|N_1, N_0]$
- It is the variance in inference for descriptive estimator
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- In practice, we usually use White estimator of the variance matrix
- It is **calculated without considering design-based uncertainty**<sup>1</sup>

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- We can see that if we ignore design-based uncertainty in inference
- It is fine if we have a small sample compared with a massive population
- Like you have a CFPS dataset to represent all families in China
- But the positive bias will become large if we have a large sample size compared with a limited population
- Like you have a province-level regression
- In this case, traditional variance estimation can be too large and too conservative
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- What is the impact of class size on students' achievement?
- Hard to identify using observational data (selection problem)
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- Kids are randomly assigned to two kinds of classes
  - (1) Small class with 13-17 children; (2) Regular class with 22-25 children
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- One assumption we always make is i.i.d.
- However, students in the same class are of course not independently sampled
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$$y_{ig} = \beta_0 + \beta_1 x_g + e_{ig}$$

- $y_{ig}$  test score;  $x_g$  class size (randomly assigned);  $e_{ig}$  error term
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- $\rho_e$  is the error intraclass correlation,  $\sigma_e^2$  is the error variance
- Assume that we can decompose error into

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- We assume that  $\nu_g$  captures all within class correlations ( $\eta_{ig} \perp \eta_{jg}$ )
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- Let  $V_c(\hat{\beta}_1)$  be the conventional OLS variance,  $V(\hat{\beta}_1)$  be the correct variance
- Assume we have classes with equal size  $n$ , then

$$\frac{V(\hat{\beta}_1)}{V_c(\hat{\beta}_1)} = 1 + (n-1)\rho_e$$

- We call this Moulton factor
- $n, \rho_e \uparrow \Rightarrow$  Bias of conventional variance  $\uparrow$
- Larger  $n$  means fewer groups  $\Rightarrow$  less information
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  - (4) Error intraclass correlation  $\rho_e$
- The implication of (3)
  - Bias can be very large in the fixed group treatment  $x_{it}$  case
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# Clustered Standard Errors: Fix the Bias

- Now we know that std error estimation can be biased when we have correlation within classes
- What we should do? Several methods are available
  - (1) Use Moulton factor equation (2) to correct for that poor error structure assumption (homoskedasticity)
  - (2) Hsiao method (Hsiao and Fargher (1988) clustering with error correction)
    - Generally consistent as number of groups  $n \rightarrow \infty$  (in finite, correction biased)
  - (3) Fixing group-level regressions  $y_{it} = \beta_0 + \beta_1 x_{it} + \epsilon_{it}$  using WLS (group size as weights)
    - Better finite-sample properties, but  $n$  has to be group-fixed
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  - (3) Running group-level regressions  $\bar{y}_g = \beta_0 + \beta_1 x_g + \bar{e}_g$  using WLS (group size as weights)  
Better finite-sample properties, but  $x_g$  has to be group-fixed
  - Other methods: Block bootstrap, MLE...

# Clustered Standard Errors: Choosing Cluster Levels

- How to choose the level of clustering?
- In STAR experiment, why not boy/girl, black/white/asian...?
- Clustering in more dimensions/higher level gives you larger std errs
- Is that OK to always cluster in more and more dimensions (be conservative)? NO.  
You can be too conservative  $\Rightarrow$  Overestimate std err
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- This is because when you cluster in more and more dimensions
- Or at higher and higher level
- Your effective sample size compared with effective population becomes larger and larger
- As Abadie et al. (2020) has shown, it leads to overestimation of the std err
- For example, you have data of 10,000 firms in 20 provinces
- 10,000 can be a very small proportion of all firms in mainland China
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- Thus, two issues remains
  - *How to choose cluster level reasonably?*
  - *How to incorporate design-based uncertainty?*
- Abadie et al. (2023) considers clustering as a sampling/design problem
- Cluster level depends on how you get your samples/assign your treatment
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- There are three misconceptions they want to clarify
  - 1. The need for clustering hinges on the presence of a correlation between residuals
    - No. The issue is the clustering of sampling or treatment assignment.
    - Even if students' scores are correlated within classroom, there is no need to cluster when sampling and treatment are totally random.
  - 2. No harm in using clustered std err when they are not required
    - Confidence intervals will be unnecessarily conservative.
  - 3. Researchers either fully adjust for clustering by using Liang and Zeger (1986) or not do that at all
    - Not really. They propose a new estimator  $CCV/TSCE$  to correct for large effects when sample size is clustering.

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Here are some empirical suggestions from Abadie et al. (2023)

- 1. If sampling and treatment are both random
  - Do not cluster
  - In this case, if sample represents a large fraction of the population, even white-noise estimator is too conservative (Abadie et al., 2023)
- 2. If random sampling but clustered treatment assignment
  - Cluster at the treatment level
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- 3. If clustered sampling, random treatment assignment
  - a. Cluster at the sampling level, if you have small fraction of sampled clusters or small fraction of sampled units within each cluster
    - i. This is especially important in panel data analysis
    - ii. Do not cluster in other cases
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- Case 1: (*Sampling cluster*) Some household/firm survey will
  - (1) Randomly select 50/250 prefectures in China
  - (2) Randomly select 100 households/firms in each sampled prefecture
- It gives you a natural stratified data set
- Just cluster at city level (in general, first sampling stage level)
- Case 2: (*Treatment cluster*) STAR assigns treatment at class level
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# Clustered Standard Errors: DID and Serial Correlation

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- When using panel data, we usually employ time variation for identification
- You draw people, but not people in a specific year  $\Rightarrow$  serial correlation
- You are drawing samples/assign treatment clustered at individual level
- Thus, DID gives a natural clustering structure of error
- One-level-up principle:  
Cluster at individual/province/city level, but NEVER  
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- One special case we must underscore is panel data analysis
- When using panel data, we usually employ time variation for identification
- You draw people, but not people in a specific year  $\Rightarrow$  serial correlation
- You are drawing samples/assign treatment clustered at individual level
- Thus, DID gives a natural clustering structure of error
- One-level-up principle:  
Cluster at individual/province/city level, but NEVER  
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- Today we discuss two nonstandard standard error issues
  - When sample is large compared with population
  - When errors are not i.i.d. but clustered
- In the first issue, we claim that we need to consider both sampling-based and design-based uncertainty
- Using traditional inference will have too large and conservative std err

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- In the second case, we find that not adjusting for cluster will generate a too small std err
- We can use LZ estimator to fix it (consistent as  $\#groups \rightarrow \infty$ )
- Clustering at higher level is not always good
- Clustering comes from either clustered sampling or clustered treatment
- Cluster at the first sampling stage, or treatment assignment level
- Do NOT cluster if you have a totally random sample and random treatment
- In DID, cluster one level up to take care of the serial correlation

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