

## Section 4

### Linear Regression and Randomized Experiments

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# Overview

- Logistics:
  - **Pset 4 released!** Due at 11:59 pm (ET) on Oct 6
  - Research project memo: Due at 11:59 pm (ET) on Oct 1
  - OH: Mondays **3-5pm**
- Today's topics:
  1. Linear regression and robust variance estimator
  2. Linear regression with covariates
  3. Block randomized trials
  4. Cluster randomized trials

# Recap: Linear Regression

- Using OLS to estimate ATEs
  - $\hat{\tau}_{\text{ols}} = \arg \min_{\tau} \sum_{i=1}^n (Y_i - \alpha - \tau D_i)^2 = \hat{\tau}_{\text{diff}} \leadsto$  unbiased
  - Linearity?  $\leadsto$  justified by consistency assumption

$$\begin{aligned} Y_i &= D_i Y_i(1) + (1 - D_i) Y_i(0) \\ &= \mathbb{E}[Y_i(0)] + D_i \tau + \{Y_i(0) - \mathbb{E}[Y_i(0)]\} + D_i (\tau_i - \tau) \\ &= \alpha + D_i \tau + \epsilon_i \end{aligned}$$

- Mean independent errors:  $\mathbb{E}[\epsilon_i \mid D_i] = 0?$   $\leadsto$  under randomization

# Linear regression and robust variance estimator

- Can we use "standard" variance estimator:  $\mathbb{V}[\varepsilon_i | \mathbf{D}] = \sigma^2, \forall i$ ?

- Inconsistent:  $\widehat{\mathbb{V}}_{const} - \mathbb{V}[\widehat{\tau}] \xrightarrow{P} \odot \neq 0$  unless ...

- Bias:

$$\mathbb{E}(\widehat{\mathbb{V}}_{const}) - \mathbb{V}[\widehat{\tau}]$$

$$= \mathbb{E}\left(\frac{\frac{1}{n-2} \sum_{i=1}^n \widehat{\varepsilon}_i^2}{\sum_{i=1}^n (D_i - \bar{D})^2}\right) - \left(\frac{\sigma_1^2}{n_1} + \frac{\sigma_0^2}{n_0}\right)$$

$$= \frac{(n_1 - n_0)(n - 1)}{n_1 n_0 (n - 2)} (\sigma_1^2 - \sigma_0^2)$$

- Unless

- Homoskedasticity holds:  $\sigma_1^2 = \sigma_0^2$

- Design is balanced:  $n_1 = n_0$

Constant effect

$$Y_i(1) - Y_i(0) = \text{const.}$$

$$V(Y_i(1)) = V(Y_i(0) + \text{const.})$$

$$= V[Y_i(0)]$$

$$1. \widehat{\tau}_{ols} = \arg \min_{\tau} \sum_{\text{model}} (\widehat{\varepsilon}_i - \tau)^2$$

$$2. V(\widehat{\tau}_{ols})$$

$$3. \widehat{\mathbb{V}}(\widehat{\tau}_{ols})$$

e.g. 'standard' est.  
 ELW  
 HC2 ...

# Linear regression and robust variance estimator

- Use robust variance estimator! [Pset4 Q1 (b)]
  - Eicker-Huber-White (EHW) estimator: consistent for  $\mathbb{V}(\hat{\tau}_{\text{diff}})$

$$\hat{\mathbb{V}}_{\text{EHW}} = \frac{\tilde{\sigma}_1^2}{n_1} + \frac{\tilde{\sigma}_0^2}{n_0}, \quad \text{where} \quad \tilde{\sigma}_d^2 = \frac{1}{n_d} \sum_{i:D_i=d} (Y_i - \bar{Y}_d)^2$$

- HC2 estimator: exactly the Neyman variance estimator  $\leadsto$  unbiased

$$\hat{\mathbb{V}}_{\text{HC2}} = \frac{\hat{\sigma}_0^2}{n_0} + \frac{\hat{\sigma}_1^2}{n_1}$$

# In R:

```
your_fitted_model <- lm(your_formula, data = your_data)
```

```
sandwich::vcovHC(your_fitted_model, type = 'HC2')
```

# Or

```
estimatr::lm_robust(your_formula, your_data, se_type = 'HC2')
```

# Linear regression with covariates

- What if we add covariates to increase **precision** of our estimates?
  - Intuition: less residual variation in  $Y_i$  after accounting for  $\mathbf{X}_i$
  - Use **centered** covariates:  $\tilde{\mathbf{X}}_i = \mathbf{X}_i - \bar{\mathbf{X}}$

$$(\hat{\tau}_{\text{adj}}, \hat{\alpha}_{\text{adj}}, \hat{\beta}_{\text{adj}}) = \arg \min_{\tau, \alpha, \beta} \sum_{i=1}^n (Y_i - \alpha - \tau D_i - \tilde{\mathbf{X}}_i' \beta)^2$$

- $\hat{\tau}_{\text{adj}}$  now **biased** but **consistent** for  $\tau$ .

# Linear regression with covariates

- Variance of adjustment estimator  $\rightarrow$  if  $p = \frac{1}{2} \rightarrow \frac{\sigma_{0x}^2}{np(1-p)} > 0 \rightarrow$  always help
- Usually will help precision, but can hurt (Freedman 2008):

$$\mathbb{V}[\hat{\tau}_{\text{diff}}] - \mathbb{V}[\hat{\tau}_{\text{adj}}] = \frac{\sigma_{0x} \{ \sigma_{0x} + 2(1-2p)\sigma_{1x} \}}{np(1-p)}$$

- If fully interacted, will never hurt precision (Lin 2013) [Pset4 Q1 (c)]

$$Y_i = \alpha + \tau D_i + \tilde{\mathbf{X}}_i' \beta + D_i \tilde{\mathbf{X}}_i' \gamma + \varepsilon_i$$

- Estimation: EHW robust variance estimators are consistent or asymptotically conservative for  $\mathbb{V}[\hat{\tau}_{\text{adj}}]$

# Linear regression with covariates

*# Step 1: Compute centered covariates*

```
your_data$Xtilde <- NULL
```

*# Step 2: Write down your formula*

```
your_formula <- NULL
```

*# Step 3: Fit the model using `lm()` or `estimatr::lm_robust()`*

```
your_fitted_model <- lm(your_formula, data = your_data)
```

*# Step 4: Compute robust standard errors (skip if you used `lm_robust`)*

```
your_vcov <- sandwich::vcovHC(your_fitted_model, type = 'HC2')
```

*# Step 5: Check the point and se estimate of your coefficients*

*# (look for tau hat!)*

```
est <- cbind("coef" = your_fitted_model$coef,  
            "se" = sqrt(diag(your_vcov)))
```



## Block randomized trials

- Setup: block randomized experiment with block indicators  $W_{ij}$ .

- Block “fixed effects”  $W_{ij} = 1$  if  $i$  is in block  $j$ , 0 otherwise.
- Blocks  $j \in \{1, \dots, J\}$  with sizes  $w_j = n_j/n$  and propensity scores  $p_j = n_{1,j}/n_j$

- Recall STAR project: within each school (block), classes were randomized.

$$p_j = \frac{n_{1,j}}{n_j} = \frac{\text{\# treated classes in school } j}{\text{\# total classes in school } j}$$

- Naive approach: just include the block FEs in OLS [Pset4 Q2 (a)]

$$(\hat{\tau}_{b,fe}, \hat{\alpha}_1, \dots, \hat{\alpha}_J) = \arg \min_{(\tau, \alpha_1, \dots, \alpha_J)} \sum_{i=1}^n \left( Y_i - \tau D_i - \sum_{j=1}^J \alpha_j W_{ij} \right)^2$$

- $\hat{\tau}_{b,fe}$  **not consistent** for the PATE unless ...

$$\hat{\tau}_{b,fe} \xrightarrow{p} \frac{\sum_{j=1}^J \omega_j \tau_j}{\sum_{j=1}^J \omega_j} \quad \text{where} \quad \omega_j = w_j p_j (1 - p_j) \quad p_1 = p_2 = \frac{1}{2}$$

- Propensity scores are equal across blocks:  $p_j = p$  for all  $j$ .
- ATEs are equal across strata  $\tau_j = \tau$  for all  $j$ .

## Block randomized trials: Correct analysis

1. Just use original Neyman analysis aggregating within-strata analyses. [Pset3 Q5]
2. Weight OLS by inverse of the propensity score. [Pset4 Q2 (b)]
3. Fully interact block FEs with treatment. [Pset4 Q2 (c)]
  - Check Imbens and Rubin (2015) Ch.9.6.1, second model
  - See this simulation study using DeclareDesign:  
<https://declaredesign.org/blog/biased-fixed-effects.html>

# Block randomized trials: Correct analysis

2. Weight OLS by inverse of the propensity score.

$$(\hat{\tau}_{b,w}, \hat{\alpha}_1, \dots, \hat{\alpha}_J) = \arg \min_{(\tau, \alpha_1, \dots, \alpha_J)} \sum_{i=1}^n s_i \left( Y_i - \tau D_i - \sum_{j=1}^J \alpha_j W_{ij} \right)^2$$

where  $s_i = \sum_{j=1}^J \left\{ \left( \frac{1}{p_j} \right) D_i + \left( \frac{1}{1-p_j} \right) (1 - D_i) \right\} W_{ij}$  and  $p_j = n_{1,j}/n_j$ .

# In R

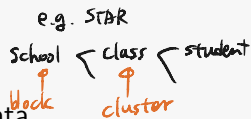
```
your_formula <- as.formula("outcome ~ treat + x_tilde1 + x_tilde2")
your_data <- data.frame(outcome, treat,
                        x_tilde1, x_tilde2,
                        weights, block)

your_fitted_model <- estimatr::lm_robust(your_formula, data = your_data,
                                         weights = weights, # s
                                         se_type = "HC2",
                                         fixed_effects = block)
```

ID of i	ID of j (school)
1	1
2	1
3	1
4	2
5	2
6	3
⋮	⋮

Not  $W_{ij}$ !

# Cluster randomized trials



- Treatment allocated at a higher level than the data.
  - Suppose schools are randomized and all the classes in same school receives same treatment
  - Now school is not a block, but cluster!
- Setup:
  - Clusters:  $k \in \{1, \dots, K\}$
  - Randomly choose  $K_1$  treatment clusters,  $K_0$  control.
  - Each cluster has units  $i \in \{1, \dots, m_k\}$  with  $\sum_{k=1}^K m_k = n$
  - Treatment assignment at cluster level:  $D_{ik} = D_k$
  - Potential outcomes  $Y_{ik}(d)$
- Cost of clustering
  - More similarity  $\leadsto$  each unit provides redundant information  $\leadsto$  less efficiency under clustering



# Cluster randomized trials

- Use **cluster-robust variance estimator**

*# In R*

```
your_formula <- as.formula("outcome ~ treat + x_tilde1 + x_tilde2")
```

```
your_data <- data.frame(outcome, treat,  
                        x_tilde1, x_tilde2,  
                        cluster)
```

```
your_fitted_model <- estimatr::lm_robust(your_formula, data = your_data,  
                                         clusters = cluster,  
                                         se_type = "CR2")
```

```
??estimatr::lm_robust # Check more options for se_type
```

*# Or*

```
your_model <- lm(your_formula, data = your_data)
```

```
your_vcov <- clubSandwich::vcovCR(your_model, cluster = your_data$cluster,  
                                  type = "CR2")
```

- You may have block and cluster design at the same time! [Pset4 Q3]