### **Section 5**

#### **Observational Studies 1**

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**GOV 2003** 

Oct 7, 2021

### **Overview**

- Logistics:
  - No pset this week!
- Today's topics:
  - 1. Review session
  - 2. No unmeasured confounding + regression

#### What we learned so far

- Fisher's approach to inference: randomization inference
- Neyman's approach to inference for the ATE: diff-in-means estimator
- Analyzing experiments with regression
  - Simple OLS estimator + robust variance estimator
  - + Covariates
  - + Block design
  - + Cluster design
- This week: observational studies
  - Before we move on, let's quickly review experimental designs!

## **Experimental design**

- Types of experiments by their assignment mechanism
  - Bernoulli randomization: Each unit is assigned D<sub>i</sub> = 1 with prob.
     p independently (coin flips)
  - Completely randomized experiment: Randomly sample  $n_1$  units from the population to be treated
  - Block/stratified randomized experiment: Completely randomized experiment in each block → always efficient for PATE
  - Cluster randomized experiment: Treatment assignment at a higher level → allows for interference within clusters
- Exercise: comparing experimental designs through simulation
  - 1. Assume true potential outcomes
  - 2. Select one assignment mechanism
  - 3. Randomly generate treatment assignment
  - 4. Estimate SATE (using diff-in-means estimator)
  - 5. Repeat 3-4 multiple times
  - 6. Draw a distribution of estimates

# **Experimental design**

Setup:

• SATE = 
$$\frac{1}{16} \sum_{i=1}^{16} \tau_i = 8.5$$

Design is balanced (except for Bernoulli)

/ - \			
$Y_i(0)$	$Y_i(1)$	$ au_i$	Block/Cluster
0	1	1	А
0	2	2	Α
0	3	3	Α
0	4	4	Α
0	5	5	В
÷	÷	÷	:
0	16	16	D
	Y <sub>i</sub> (0) 0 0 0 0 0 0	0 1 0 2 0 3 0 4 0 5 : :	0 1 1 1 0 2 2 0 3 3 0 4 4 0 5 5 5 : : : :

- Q: Which design would have the largest (smallest) variance?
- Check the results here

#### Observational studies

- Problem:
  - Non-randomized treatment
  - $\rightarrow$  { $Y_i(1), Y_i(0)$ }  $\not\perp D_i$
  - → selection bias = unidentified ATT

$$\underbrace{\mathbb{E}\big[Y_i|D_i=1\big] - \mathbb{E}\big[Y_i|D_i=0\big]}_{\text{diff-in-means}} = \underbrace{\tau_t}_{\text{ATT}} + \underbrace{\mathbb{E}\big[Y_i(0)|D_i=1\big] - \mathbb{E}\big[Y_i(0)|D_i=0\big]}_{\text{selection bias}}$$

- What can we do for the identification?
  - Assume no unmeasured confounding with positivity
  - Partial identification: analysis of bounds for the ATE
  - Sensitivity analysis . . .

# Identification: No unmeasured confounding

- Identification
  - Let's begin with most common set of assumptions:
    - 1. **Overlap**/Positivity:  $0 < Pr[D_i = 1 | \mathbf{X}_i] < 1$
    - 2. No unmeasured confounding:  $\{Y_i(1), Y_i(0)\} \perp D_i \mid X_i$
  - This will identify the PATE:

$$\tau = \mathbb{E}[Y_{i}(1) - Y_{i}(0)]$$

$$= \mathbb{E}_{\mathbf{X}} \{ E[Y_{i}(1) - Y_{i}(0) \mid X_{i}] \}$$

$$= \mathbb{E}_{\mathbf{X}} \{ E[Y_{i}(1) \mid X_{i}] - \mathbb{E}[Y_{i}(0) \mid X_{i}] \}$$

$$= \mathbb{E}_{\mathbf{X}} \{ E[Y_{i}(1) \mid D_{i} = 1, X_{i}] - \mathbb{E}[Y_{i}(0) \mid D_{i} = 0, X_{i}] \}$$

$$= \mathbb{E}_{\mathbf{X}} \{ E[Y_{i} \mid D_{i} = 1, X_{i}] - \mathbb{E}[Y_{i} \mid D_{i} = 0, X_{i}] \}$$

- Estimation
  - Regression
  - Matching/Weighting (Module 7)

# **Estimation: Regression-based estimators**

Treated and control conditional expectation functions (CEFs):

$$\mu_1(\mathbf{x}) = \mathbb{E}[Y_i(1) \mid \mathbf{X}_i = \mathbf{x}], \qquad \mu_0(\mathbf{x}) = \mathbb{E}[Y_i(0) \mid \mathbf{X}_i = \mathbf{x}]$$

By consistency and no unmeasured confounding:

$$\underbrace{\mu_1(\mathbf{x})}_{\text{counterfactual}} = \underbrace{\mathbb{E}[Y_i \mid D_i = 1, \mathbf{X}_i = \mathbf{x}]}_{\text{observational}}, \qquad \mu_0(\mathbf{x}) = \mathbb{E}[Y_i \mid D_i = 0, \mathbf{X}_i = \mathbf{x}]$$

- Estimate CEFs using regression estimators  $\widehat{\mu}_1(\mathbf{x})$  and  $\widehat{\mu}_0(\mathbf{x})$ .
  - Might be linear or nonlinear models (e.g., GAMs)
  - → Regression estimator of the ATE:

$$\widehat{\tau}_{\text{reg}} = \frac{1}{n} \sum_{i=1}^{n} \widehat{\mu}_{1}(\mathbf{X}_{i}) - \widehat{\mu}_{0}(\mathbf{X}_{i})$$

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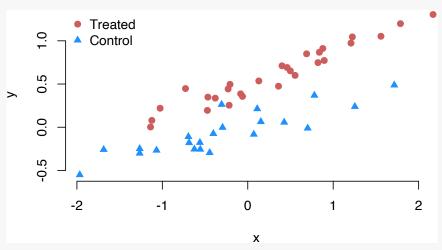
# **Estimation: Regression-based estimators**

$$\widehat{\tau}_{\text{reg}} = \frac{1}{n} \sum_{i=1}^{n} \widehat{\mu}_{1}(\mathbf{X}_{i}) - \widehat{\mu}_{0}(\mathbf{X}_{i})$$

- General procedure:
  - Obtain predicted values for all units when  $D_i = 1$ .
  - Obtain predicted values for all units when  $D_i = 0$ .
  - Take the average difference between these predicted values.
- Safest practice:
  - Estimate separate regression in each treatment group.
  - Sometimes called an imputation estimator.
  - Procedure:
    - Regress Y<sub>i</sub> on X<sub>i</sub> in the treatment group and get predicted values for all units (treated or control).
    - Regress Y<sub>i</sub> on X<sub>i</sub> in the control group and get predicted values for all units (treated or control).
    - Take the average difference between these predicted values.

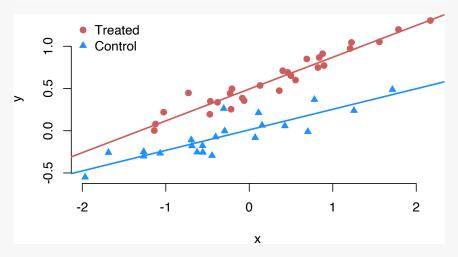
# Toy example

• Data is as follows and we will use linear regression to estimate CEFs



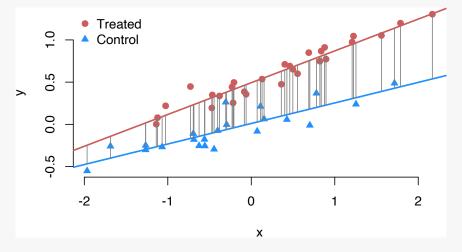
## Imputation estimator visualization

```
mod0 <- lm(y\sim x, data = toy_data, subset = d==0)
mod1 <- lm(y\sim x, data = toy_data, subset = d==1)
```



mu0.imps = predict(mod0, toy\_data); mu1.imps = predict(mod1, toy\_data)
cat("Estimate of ATE:", mean(mu1.imps - mu0.imps))

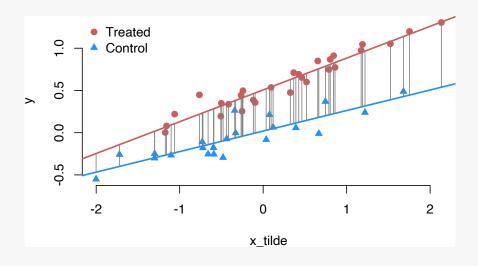
## Estimate of ATE: 0.4873975



### Fully interacted OLS visualization

- What if  $\widehat{\mu}_1(\mathbf{x})$  and  $\widehat{\mu}_0(\mathbf{x})$  are from fully interacted OLS with centered covariates?
  - Equivalent to running separate models for  $\widehat{\mu}_1(\mathbf{x})$  and  $\widehat{\mu}_0(\mathbf{x})$
  - $\widehat{\tau}_{reg} \equiv$  estimated coefficient on  $D_i$ 
    - Recall: Under linear models, \(\hat{\tau}\_{reg}\) is sometimes equivalent to a coefficient.

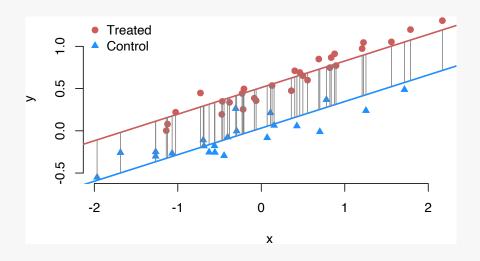
```
toy_data$x_tilde <- toy_data$x - mean(toy_data$x)</pre>
mod_full <- lm(y\sim d+x\_tilde+d*x\_tilde, data = toy\_data)
dat0 <- toy_data %>% mutate(d = 0); dat1 <- toy_data %>% mutate(d = 1)
mu0.full = predict(mod_full, dat0); mu1.full = predict(mod_full, dat1)
cat("Estimate of ATE (Fully interacted):", mean(mu1.full - mu0.full),
    "\nEstimate of ATE (Imputation):", mean(mul.imps - mu0.imps),
    "\nEstimated coefficient on Di", mod_full$coefficients["d"])
## Estimate of ATE (Fully interacted): 0.4873975
## Estimate of ATE (Imputation): 0.4873975
## Estimated coefficient on Di 0.4873975
```



#### Uninteracted OLS visualization

What if \$\hat{\mu}\_1(\mathbf{x})\$ and \$\hat{\mu}\_0(\mathbf{x})\$ are from the same OLS model Y ~ D + X?
 \$\hat{\tau}\_{reg}\$ ≡ estimated coefficient on \$D\_i\$

```
mod <- lm(y\sim d+x, data = toy_data)
mu0 = predict(mod, dat0); mu1 = predict(mod, dat1)
cat("Estimate of ATE (Uninteracted):", mean(mu1 - mu0),
    "\nEstimated coefficient on Di", mod$coefficients["d"],
    "\nEstimate of ATE (Fully interacted):", mean(mu1.full - mu0.full),
    "\nEstimate of ATE (Imputation):", mean(mul.imps - mu0.imps))
## Estimate of ATE (Uninteracted): 0.479676
## Estimated coefficient on Di 0.479676
## Estimate of ATE (Fully interacted): 0.4873975
## Estimate of ATE (Imputation): 0.4873975
```



### Variance estimation

- How do we get estimates of the variance of  $\widehat{\tau}_{reg}$ ?
- Nonparametric bootstrap
  - Recall: Source of variance is due to sampling
  - Idea: View sample (data) as "population" → in-sample "sampling"
- Procedure:
  - Randomly resample n rows of the data with replacement
  - Refit the regressions on the bootstrapped data.
  - Calculate  $\hat{\tau}_{reg}$  in each bootstrap
  - Repeat several times and use empirical variance of the bootstraps

### **Bootstrap sample codes**

```
set.seed(02138); sims<-500; tau_hat_draws<-rep(NA, sims)</pre>
for (i in 1:sims) { # Repeat the following several times
  # 1. Randomly resample n rows of the data with replacement
  sample_boot <- dplyr::slice_sample(toy_data, n = nrow(toy_data),</pre>
                                    replace = TRUE
  # 2. Refit the regressions on the bootstrapped data
  model \leftarrow lm(y \sim d + x_tilde + d*x_tilde, data = toy_data)
  dat1 <- sample_boot; dat1$d <- 1</pre>
  dat0 <- sample_boot; dat0$d <- 0</pre>
  mul_hat <- predict(model, newdata = dat1)</pre>
  mu0_hat <- predict(model, newdata = dat0)</pre>
  # 3. Calculate tau_hat in each bootstrap
  tau_hat_draws[i] <- mean(mul_hat - mu0_hat)</pre>
}
# 4. Use empirical variance of the bootstraps
var(tau_hat_draws)
   [1] 0.0003247686
```

#### DAG

- How do we know if no unmeasured confounders holds?
  - One way: use DAGs and look at back-door paths.

#### D-separation

- Can we determine conditional independence from our causal DAG?
- Yes! To verify that  $A \perp \!\!\! \perp B \mid C$  where each is a set of nodes:
  - 1. Find all paths between A and B.
  - 2. Check if each path is blocked.
  - 3. If all paths are blocked, then A is **d-separated** from B by C
- Ways to block A → B (each is a node):
  - 1.  $A \rightarrow C \rightarrow B$ , C is observed (conditioned)
  - 2.  $A \leftarrow C \leftarrow B$ , C is observed
  - 3.  $A \rightarrow C \rightarrow B$ , C is observed
  - 4.  $A \rightarrow C \leftarrow B$ , C is **not** observed
  - If C observed → collider bias
  - e.g., A=bicycle accident,B=stomachache, C=hospitalization; Sackett (1979)