# Tools for Planning and Analyzing Randomized Controlled Trials and A/B Tests

Johann Gagnon-Bartsch, Adam Sales, Duy Pham, Charlotte Mann, and Jaylin Lowe

Department of Statistics, University of Michigan & Department of Mathematical Sciences, Worcester Polytechnic Institute

Educational Data Mining

July 14, 2024





Follow the steps at

https://tinyurl.com/edmrct-setup







Workshop Resources

Github site

to get everything ready to follow along in RStudio!



- 1:30-1:45 Part I: Conceptual Overview
- 1:45-3:00 Part II: Estimating Effects with RCT Data
- 3:00-3:30 Part III: Incorporating Auxiliary Data
- 3:30-3:45 Break 15 min
- 3:45-4:15 Part IV: Treatment Effect Heterogeneity
- 4:15–5:00 Part V: Planning Experiments



- Tutorial website: https://tinyurl.com/edmrct
- RStudio
- Clone repo from Github: https://github.com/manncz/edm-rct-tutorial/

We will be alternating between:

- Conceptual descriptions of the methods
- Detailed walk-throughs of the software
- Opportunities for you to run analyses yourself, with our help

#### Please feel free to ask questions at any time!

- Calling out (unmute yourself if on Zoom)
- Zoom chat
- Any other way you can think of to get our attention



# **Conceptual Overview**

Estimating Effects with RCT Data

Incorporating Auxiliary Data

Break

Treatment Effect Heterogeneity

Planning Experiments

# **Experiments in Education Research**

"Experiment" = "RCT" = "Randomized Controlled Trial"





- Randomize subjects (students? teachers? schools?) between condition
- Expose subjects to their randomized conditions
- Measure outcome(s) of interest

# **Experiments in Education Research**

"Experiment" = "RCT" = "Randomized Controlled Trial"





- Randomize subjects (students? teachers? schools?) between condition
- Expose subjects to their randomized conditions
- Measure outcome(s) of interest
- Associations between condition and outcomes are causal

# **Experiments in Education Research**

"Experiment" = "RCT" = "Randomized Controlled Trial"





- Randomize subjects (students? teachers? schools?) between condition
- Expose subjects to their randomized conditions
- Measure outcome(s) of interest
- Associations between condition and outcomes are causal
- Typical examples:
  - A/B tests in online learning
  - Field trials of (say) new curriculum vs. business as usual

### **Example 1: ASSISTments ETrials**







- Question: Text or video hints?
- Outcome: Complete skill builder?
- n = 683 middle school students



- Question: Text or video hints?
- Outcome: Complete skill builder?
- n = 683 middle school students

Results,

- Video: 205/337 (61%) completed
- Text: 193/346 (56%) completed



# **Example II: Cognitive Tutor Effectiveness Trial**

- 73 High Schools & 74 Middle Schools in 7 states
- Similar schools paired
- In each pair, one school randomized to treatment, one to control
- Algebra I students in Trt school used CTAI, Control school used business as usual
- All students took a posttest at the end of the year







#### Average Posttest

	Middle		High	
	Year 1	Year 2	Year 1	Year 2
Control	17.4	16.9	10.3	9.7
Treatment	14.3	15.2	10.1	10.6

1. What is the average effect of [intervention] on [outcome]?



2. How Does the effect vary?

- 1. What is the average effect of [intervention] on [outcome]?
  - "Intervention" AKA "Treatment" (the thing you're randomizing)
    - Contrast between 2+ conditions
    - E.g. access to ChatGPT hint vs teacher-written hint vs no hint
    - For today: focus on 2 conditions, "Treatment" vs "Control"
    - (those labels may be arbitrary)



2. How Does the effect vary?

- 1. What is the average effect of [intervention] on [outcome]?
  - "Intervention" AKA "Treatment" (the thing you're randomizing)
    - Contrast between 2+ conditions
    - E.g. access to ChatGPT hint vs teacher-written hint vs no hint
    - For today: focus on 2 conditions, "Treatment" vs "Control"
    - (those labels may be arbitrary)
  - "Outcome"
    - Scalar quantity that the intervention might affect
    - E.g. student correctness on the next problem (0 or 1)
- 2. How Does the effect vary?



- 1. What is the average effect of [intervention] on [outcome]?
  - "Intervention" AKA "Treatment" (the thing you're randomizing)
    - Contrast between 2+ conditions
    - E.g. access to ChatGPT hint vs teacher-written hint vs no hint
    - For today: focus on 2 conditions, "Treatment" vs "Control"
    - (those labels may be arbitrary)
  - "Outcome"
    - Scalar quantity that the intervention might affect
    - E.g. student correctness on the next problem (0 or 1)
  - "Average Effect" ...to be defined soon!
- 2. How Does the effect vary?



- 1. What is the average effect of [intervention] on [outcome]?
  - "Intervention" AKA "Treatment" (the thing you're randomizing)
    - Contrast between 2+ conditions
    - E.g. access to ChatGPT hint vs teacher-written hint vs no hint
    - For today: focus on 2 conditions, "Treatment" vs "Control"
    - (those labels may be arbitrary)
  - "Outcome"
    - Scalar quantity that the intervention might affect
    - E.g. student correctness on the next problem (0 or 1)
  - "Average Effect" ...to be defined soon!
- 2. How Does the effect vary?
  - From one (type of) student to the next
  - From one context to the next



1. Get the most out of your data: more data  $\rightarrow$  better estimation!!

2. ...Without making unnecessary assumptions

3. Easily

4. Design better experiments to start with



- 1. Get the most out of your data: more data  $\rightarrow$  better estimation!!
  - Baseline covariate data
  - Historical user data
- 2. ...Without making unnecessary assumptions

3. Easily

4. Design better experiments to start with



- 1. Get the most out of your data: more data  $\rightarrow$  better estimation!!
  - Baseline covariate data
  - Historical user data
- 2. ...Without making unnecessary assumptions
  - "Design-based" methods
  - NO assumptions about confounding, models, etc. etc.
- 3. Easily

4. Design better experiments to start with



- 1. Get the most out of your data: more data  $\rightarrow$  better estimation!!
  - Baseline covariate data
  - Historical user data
- 2. ...Without making unnecessary assumptions
  - "Design-based" methods
  - NO assumptions about confounding, models, etc. etc.
- 3. Easily
  - i.e. without a PhD in statistics
  - Use our software package :)
- 4. Design better experiments to start with





- Fixed at baseline
- Unaffected by treatment



- Fixed at baseline
- Unaffected by treatment

Uses:

- More precise estimates
- Explore effect variation



- Log data. For each previous skillbuilder,
  - Completed skill builder?
  - # problems attempted / completed?
  - Time to mastery
  - • •
- Demographic data



- Log data. For each previous skillbuilder,
  - Completed skill builder?
  - # problems attempted / completed?
  - Time to mastery
  - • •
- Demographic data

#### Don't use post-treatment variables!

- Covariate and outcome data from other subjects
- Often: historical data



# **Auxiliary Data**

- Covariate and outcome data from other subjects
- Often: historical data
- Requirements
  - Separate sample from RCT
  - (some of the) same covariate data as for RCT subjects
  - similar outcome data as RCT



# **Auxiliary Data**

- Covariate and outcome data from other subjects
- Often: historical data
- Requirements
  - Separate sample from RCT
  - (some of the) same covariate data as for RCT subjects
  - similar outcome data as RCT

Uses:

- More precise estimates
- Planning experiments





Estimate treatment effects



Estimate treatment effects Using all our data



Estimate treatment effects Using all our data

- Covariates (even high-dimensional)
- Auxiliary/historical data



Estimate treatment effects Using all our data

- Covariates (even high-dimensional)
- Auxiliary/historical data

Without bias or extra assumptions

**Conceptual Overview** 

# Estimating Effects with RCT Data

Incorporating Auxiliary Data

Break

**Treatment Effect Heterogeneity** 

**Planning Experiments** 

Consider a randomized experiment with:

- N participants
- One treatment group, one control group








If the coin had landed the other way, the outcome may have been different.



If the coin had landed the other way, the outcome may have been different.

• Each subject has two **potential outcomes**.



If the coin had landed the other way, the outcome may have been different.

• Each subject has two **potential outcomes**.

One for treatment, one for control.



If the coin had landed the other way, the outcome may have been different.

- Each subject has two **potential outcomes**. One for treatment, one for control.
- We only ever observe **one** potential outcome.



If the coin had landed the other way, the outcome may have been different.

- Each subject has two **potential outcomes**. One for treatment, one for control.
- We only ever observe **one** potential outcome. The other is a counterfactual.

















• For each participant i there are two potential outcomes,  $y_i^t$  and  $y_i^c$ 



- For each participant i there are two potential outcomes,  $y_i^t$  and  $y_i^c$
- Potential outcomes are **fixed** values, not random



- For each participant i there are two potential outcomes,  $y_i^t$  and  $y_i^c$
- Potential outcomes are **fixed** values, not random
- Let  $T_i$  be the treatment assignment of unit i

 $T_i = \begin{cases} 1, & \text{Unit } i \text{ is assigned to treatment} \\ 0, & \text{Unit } i \text{ is assigned to control} \end{cases}$ 



- For each participant i there are two potential outcomes,  $y_i^t$  and  $y_i^c$
- Potential outcomes are **fixed** values, not random
- Let  $T_i$  be the treatment assignment of unit i

 $T_i = \begin{cases} 1, & \text{Unit } i \text{ is assigned to treatment} \\ 0, & \text{Unit } i \text{ is assigned to control} \end{cases}$ 

 Let Y<sub>i</sub> be the observed outcome for unit i. If unit i is assigned to treatment, we observe y<sup>t</sup><sub>i</sub>; otherwise, we observe y<sup>c</sup><sub>i</sub>:

$$Y_i = \begin{cases} y_i^c & \text{ if } T_i = 0\\ y_i^t & \text{ if } T_i = 1 \end{cases}$$



• The individual treatment effect is

$$\tau_i = y_i^t - y_i^c$$



• The individual treatment effect is

$$\tau_i = y_i^t - y_i^c$$



• The individual treatment effect is never observed.

• The individual treatment effect is

$$\tau_i = y_i^t - y_i^c$$



- The individual treatment effect is never observed.
- The average treatment effect (ATE) is

$$\bar{\tau} = \frac{1}{N} \sum_{i=1}^{N} \tau_i$$

• The individual treatment effect is

$$\tau_i = y_i^t - y_i^c$$



- The individual treatment effect is **never observed**.
- The average treatment effect (ATE) is

$$\bar{\tau} = \frac{1}{N} \sum_{i=1}^{N} \tau_i$$

• The average treatment effect can be estimated.

• The individual treatment effect is

$$\tau_i = y_i^t - y_i^c$$



- The individual treatment effect is **never observed**.
- The average treatment effect (ATE) is

$$\bar{\tau} = \frac{1}{N} \sum_{i=1}^{N} \tau_i$$

- The average treatment effect can be estimated.
- Also: average effects for subgroups of subjects (more later)

- We only observe one potential outcome for each subject
  - For treatment subjects  $y^t$
  - For control,  $\boldsymbol{y}^c$



- We only observe one potential outcome for each subject
  - For treatment subjects  $y^t$
  - For control,  $y^c$
- One potential outcome is always missing
- We need to impute the missing potential outcome



- We only observe one potential outcome for each subject
  - For treatment subjects  $y^t$
  - For control,  $y^c$
- One potential outcome is always missing
- We need to impute the missing potential outcome
- Two approaches to imputation:
  - 1. Use randomization: unbiased, but imprecise



- We only observe one potential outcome for each subject
  - For treatment subjects  $y^t$
  - For control,  $y^c$
- One potential outcome is always missing
- We need to impute the missing potential outcome
- Two approaches to imputation:
  - 1. Use randomization: unbiased, but imprecise
  - 2. Use covariates & and model: biased, but precise



- We only observe one potential outcome for each subject
  - For treatment subjects  $y^t$
  - For control,  $y^c$
- One potential outcome is always missing
- We need to impute the missing potential outcome
- Two approaches to imputation:
  - 1. Use randomization: unbiased, but imprecise
  - 2. Use covariates & and model: biased, but precise
  - 3. Our approach: use both!



Train algorithms to predict  $y^c$ ,  $y^t$  as a function of covariates

 $f^c: \mathbf{X} o y^c$  (use data from ctl group)  $f^t: \mathbf{X} o y^t$  (use data from trt group)



Train algorithms to predict  $y^c$ ,  $y^t$  as a function of covariates

 $f^c: \mathbf{X} \to y^c$  (use data from ctl group)  $f^t: \mathbf{X} \to y^t$  (use data from trt group)

### Step 2:

Use algorithms to get imputations:

$$\hat{y}_i^c = f^c(X_i)$$
$$\hat{y}_i^t = f^t(X_i)$$



Train algorithms to predict  $y^c$ ,  $y^t$  as a function of covariates

 $f^c: \mathbf{X} \to y^c$  (use data from ctl group)  $f^t: \mathbf{X} \to y^t$  (use data from trt group)

#### Step 2:

Use algorithms to get imputations:

$$\hat{y}_i^c = f^c(X_i)$$
$$\hat{y}_i^t = f^t(X_i)$$

**Step 3:** Calculate  $\hat{m}_i$ : weighted average of  $\hat{y}_i^c$  and  $\hat{y}_i^t$ 



Train algorithms to predict  $y^c$ ,  $y^t$  as a function of covariates

 $f^c: \mathbf{X} \to y^c$  (use data from ctl group)  $f^t: \mathbf{X} \to y^t$  (use data from trt group)

#### Step 2:

Use algorithms to get imputations:

$$\hat{y}_i^c = f^c(X_i)$$
$$\hat{y}_i^t = f^t(X_i)$$

**Step 3:** Calculate  $\hat{m}_i$ : weighted average of  $\hat{y}_i^c$  and  $\hat{y}_i^t$  **Step 4:** 

Use randomization-based method to estimate effects on  $Y-\hat{m}$  instead of Y



 $\hat{m}_i$  independent of  $T_i$ 



 $\hat{m}_i$  independent of  $T_i$ 

Since  $Y_i$  is a function of  $T_i$ , that means we need:

 $\hat{y}^c$  and  $\hat{y}^t$  independent of  $Y_i$ 



 $\hat{m}_i$  independent of  $T_i$ 

Since  $Y_i$  is a function of  $T_i$ , that means we need:

 $\hat{y}^c$  and  $\hat{y}^t$  independent of  $Y_i$ 

We can't use i's data to train  $f^c$  and  $f^t$ !



 $\hat{m}_i$  independent of  $T_i$ 



 $\hat{y}^c$  and  $\hat{y}^t$  independent of  $Y_i$ 

# We can't use *i*'s data to train $f^c$ and $f^t$ ! Solution: re-train $f^c$ and $f^t$ for each subject *i*, leaving out *i*'s data



```
\hat{m}_i independent of T_i
```

Since  $Y_i$  is a function of  $T_i$ , that means we need:

 $\hat{y}^c$  and  $\hat{y}^t$  independent of  $Y_i$ 

# We can't use *i*'s data to train $f^c$ and $f^t$ ! Solution: re-train $f^c$ and $f^t$ for each subject *i*, leaving out *i*'s data

"Leave-One-Out Potential Outcomes" or "LOOP"



# Sticks and Stones May Break my Bones, but Bad Models Won't Hurt Me



• What if  $f^c$  and  $f^t$  are totally wrong and bad??

# Sticks and Stones May Break my Bones, but Bad Models Won't Hurt Me



- What if  $f^c$  and  $f^t$  are totally wrong and bad??
- Estimate will still be unbiased!

# Sticks and Stones May Break my Bones, but Bad Models Won't Hurt Me



- What if  $f^c$  and  $f^t$  are totally wrong and bad??
- Estimate will still be unbiased!
- Standard errors, p-values, and confidence intervals will still be valid!
## Sticks and Stones May Break my Bones, but Bad Models Won't Hurt Me



- What if  $f^c$  and  $f^t$  are totally wrong and bad??
- Estimate will still be unbiased!
- Standard errors, p-values, and confidence intervals will still be valid!
- (core of inference is based on randomization)

## Sticks and Stones May Break my Bones, but Bad Models Won't Hurt Me



- What if  $f^c$  and  $f^t$  are totally wrong and bad??
- Estimate will still be unbiased!
- Standard errors, p-values, and confidence intervals will still be valid!
- (core of inference is based on randomization)
- Covariate adjustment won't help much

## Sticks and Stones May Break my Bones, but Bad Models Won't Hurt Me



- What if  $f^c$  and  $f^t$  are totally wrong and bad??
- Estimate will still be unbiased!
- Standard errors, p-values, and confidence intervals will still be valid!
- (core of inference is based on randomization)
- Covariate adjustment won't help much
- In moderate/large samples, it won't hurt either!

Regression method: Fit model:



$$Y_i = \beta_0 + \beta_1 T_i + \beta_2 X_{1i} + \beta_3 X_{2i} + \dots$$

Estimated effect:  $\hat{\beta}_2$ 

Regression method: Fit model:

$$Y_i = \beta_0 + \beta_1 T_i + \beta_2 X_{1i} + \beta_3 X_{2i} + \dots$$

Estimated effect:  $\hat{\beta}_2$ **Problem:** What if the model is false?

- E.g. Y isn't linear in covariates
- E.g. What if there should be interactions?



Regression method: Fit model:

$$Y_i = \beta_0 + \beta_1 T_i + \beta_2 X_{1i} + \beta_3 X_{2i} + \dots$$

Estimated effect:  $\hat{\beta}_2$ **Problem:** What if the model is false?

- E.g. *Y* isn't linear in covariates
- E.g. What if there should be interactions?

**Good news**:  $\hat{\beta}$  is approximately unbiased in large samples



Why our method?

- 1. Exactly unbiased in any sample
- 2. Use any algorithm for  $f^c$ ,  $f^t$ 
  - High dimensional covariates
  - Flexible for non-linearity, interactions



Why our method?

- 1. Exactly unbiased in any sample
- 2. Use any algorithm for  $f^c$ ,  $f^t$ 
  - High dimensional covariates
  - Flexible for non-linearity, interactions
  - $\Rightarrow$  better imputations
  - $\bullet \ \Rightarrow \text{better effect estimates}$





Why our method?

- 1. Exactly unbiased in any sample
- 2. Use any algorithm for  $f^c$ ,  $f^t$ 
  - High dimensional covariates
  - Flexible for non-linearity, interactions
  - $\Rightarrow$  better imputations
  - $\bullet \ \Rightarrow \mathsf{better} \ \mathsf{effect} \ \mathsf{estimates}$
  - We recommend random forest





- 1. Randomized treatment variable
- 2. Outcome variable
- 3. Covariates
- 4. What is the experimental design?

This is not a promise.



- 1. Randomized treatment variable
- 2. Outcome variable
- 3. Covariates
- 4. What is the experimental design?

One last digression<sup>1</sup>: experimental designs

This is not a promise.



- 1. Who or What is being randomized?
- 2. How are they being randomized?



- Individual randomization
- Cluster or Group randomization



- What's the probability each unit is assigned to treatment?
- How does one unit's assignment affect other units?



- Individual randomization
  - Bernoulli
  - Paired
- Cluster randomization
  - Paired

#### Examples

- ASSISTments E-Trials A/B test
  - Students are randomized individually
  - Students are randomized independently
  - $\bullet \ \Rightarrow \mathsf{Bernoullli}$



- ASSISTments E-Trials A/B test
  - Students are randomized individually
  - Students are randomized independently
  - $\Rightarrow$  Bernoullli
- Cognitive Tutor Effectiveness Study
  - Schools are randomized
  - Randomization is within pairs
  - (if your school is randomized to treatment, its pair *must* be randomized to control)
  - $\bullet \ \Rightarrow \mathsf{paired \ cluster \ design}$



## **Other Designs**

To be implemented (hopefully) soon:

• "Completely randomized design"



- At the outset, fix # randomized to treatment, # randomized to control
- Now  $T_i$  and  $T_j$  are dependent!
- Block-randomized design
  - e.g. a separate completely randomized experiment in each classroom
  - Paired designs are a special case

## **Other Designs**

To be implemented (hopefully) soon:

• "Completely randomized design"



- At the outset, fix # randomized to treatment, # randomized to control
- Now  $T_i$  and  $T_j$  are dependent!
- Block-randomized design
  - e.g. a separate completely randomized experiment in each classroom
  - Paired designs are a special case

Probably won't get to for a while:

- Bandit designs
  - Probability *i* is assigned to treatment depends on previous subjects' outcomes



# **Estimating Effects in Practice**

# Installation:



- You will need to install the package from Github using the *devtool* package in *R*
- e.g. install\_github("manncz/dRCT")

# **Primary Functions:**

# loop(Y, Tr, Z, pred, p, ...)

- Y: outcome vector
- Tr : treatment assignment vector
- Z: matrix of covariates
- *pred* : interpolation algorithm
- *p*: probability of treatment
- ...: optional inputs for interpolation algorithm



# pred

- loop\_rf
- loop\_ols
- loop\_glm

# p\_loop(Y, Tr, Z, pred, P, n, ...)

- Y: outcome vector
- Tr : treatment assignment vector
- Z: matrix of covariates
- *pred* : interpolation algorithm
- P: vector of pair assignments
- *n*: optional vector of cluster sizes
- ...: optional inputs for interpolation algorithm



#### pred

- p\_ols\_po
- *p\_ols\_v12*
- p\_ols\_interp
- p\_rf\_po
- *p\_rf\_v12*
- p\_rf\_interp



#### Real Data Example: Texas School Data

- AEIS: School-level data from Texas Education Agency from 2003-2011
- > 3,000 schools
- TAKS (standardized test) passing rates
- Thousands of additional possible covariates







• Inspired by the Cognitive Tutor Algebra I study (Pane et al. 2014)

- Inspired by the Cognitive Tutor Algebra I study (Pane et al. 2014)
- RCT Sample: 50 Texas middle schools
- Treatment: Alternative 8th grade mathematics curriculum
- **Design:** Schools randomly assigned to implement new curriculum or continue standard in the 2007/8 school year



- Inspired by the Cognitive Tutor Algebra I study (Pane et al. 2014)
- RCT Sample: 50 Texas middle schools
- Treatment: Alternative 8th grade mathematics curriculum
- **Design:** Schools randomly assigned to implement new curriculum or continue standard in the 2007/8 school year
- Outcome: 2008 8th grade math TAKS passing rate



- Inspired by the Cognitive Tutor Algebra I study (Pane et al. 2014)
- RCT Sample: 50 Texas middle schools
- Treatment: Alternative 8th grade mathematics curriculum
- **Design:** Schools randomly assigned to implement new curriculum or continue standard in the 2007/8 school year
- Outcome: 2008 8th grade math TAKS passing rate
- Pretest: 2007 8th grade math TAKS passing rate





- 1. Follow along while we talk through 01-explore-aeis-data.Rmd
- 2. Work through 02-effect-est.Rmd
  - Effect estimate for Bernoilli randomized trial
  - Effect estimate for paired randomed trial
  - Effect esitmate for paired cluster randomed trial
- 3. Flag any of us down as you have questions!

Conceptual Overview

Estimating Effects with RCT Data

# Incorporating Auxiliary Data

Break

Treatment Effect Heterogeneity

Planning Experiments

## **Auxiliary Data**

By "Auxiliary Data" we mean a dataset that meets these criteria:

- 1. Doesn't include data from RCT participants
- 2. Includes covariate data
- 3. Includes outcome data



## **Auxiliary Data**

By "Auxiliary Data" we mean a dataset that meets these criteria:

- 1. Doesn't include data from RCT participants
- 2. Includes covariate data
- 3. Includes outcome data

# Examples:

- A/B test: historical log data from users who worked on similar modules before the experiment started
- Field trial: Administrative (e.g. SLDS) data from students in schools that were not part of the RCT



## **Auxiliary Data**

By "Auxiliary Data" we mean a dataset that meets these criteria:

- 1. Doesn't include data from RCT participants
- 2. Includes covariate data
- 3. Includes outcome data

# Examples:

- A/B test: historical log data from users who worked on similar modules before the experiment started
- Field trial: Administrative (e.g. SLDS) data from students in schools that were not part of the RCT

Note: we have sometimes called this the "remnant"



- Already imputing potential outcomes with  $f^c$  and  $f^t$  in LOOP
- $f^c$  and  $f^t$  can be flexible, high dimensional
- They are fit to representative data


- Already imputing potential outcomes with  $f^{c}$  and  $f^{t}$  in LOOP
- $f^c$  and  $f^t$  can be flexible, high dimensional
- They are fit to representative data

Limits on  $f^{c} \mbox{ and } f^{t}$ 

- RCT sample size might be too small to fit really good models
- Human-adaptive modeling: no good!



## **Example 1: ASSISTments**

Covariates:

- Log data. For each previous skillbuilder,
  - Completed skill builder?
  - # problems attempted / completed?
  - Time to mastery
- Demographic data



## **Example 1: ASSISTments**

#### Covariates:

- Log data. For each previous skillbuilder,
  - Completed skill builder?
  - # problems attempted / completed?
  - Time to mastery
- Demographic data

## Auxiliary Data:

- Observational
- Students who were not randomzied
  - Previous users
  - Current users not assigned to that skillbuilder
- Same covariates available



## RCT

Control



## Step 1: Train Model $\hat{y}(\cdot): {m x} ightarrow Y$ With auxiliary data





## Step 1: Train Model $\hat{y}(\cdot): {m x} ightarrow Y$ With auxiliary data

#### Step 2:

Extrapolate With fitted model & RCT data





## Step 1: Train Model $\hat{y}(\cdot): {m x} ightarrow Y$ With auxiliary data

#### Step 2:

Extrapolate With fitted model & RCT data

# Step 3: Use $\hat{y}(\boldsymbol{x})$ as a "super-covariate"

## RCT Control $\hat{y}(\boldsymbol{x}_i)$ Treatment $\hat{y}(\boldsymbol{x}_{i})$







- The function  $\hat{y}(\cdot)$  is fit on auxiliary data
- The covariates x are pre-treatment
- $\Rightarrow \hat{y}({m x})$  is invariant to treatment assignment



- The function  $\hat{y}(\cdot)$  is fit on auxiliary data
- The covariates x are pre-treatment
- $\Rightarrow \hat{y}({m{x}})$  is invariant to treatment assignment
- $\hat{y}(\boldsymbol{x})$  might be an amazing covariate



- The function  $\hat{y}(\cdot)$  is fit on auxiliary data
- The covariates x are pre-treatment
- $\Rightarrow \hat{y}({m{x}})$  is invariant to treatment assignment
- $\hat{y}(oldsymbol{x})$  might be an amazing covariate
- ...or it might not



- If  $\hat{y}(\boldsymbol{x})$  predicts Y really well, we would expect a linear relationship
  - $\bullet \ \Rightarrow {\rm fit} \ {\rm OLS} \ {\rm models} \ {\rm within} \ {\rm LOOP}$



- + If  $\hat{y}(\pmb{x})$  predicts Y really well, we would expect a linear relationship
  - $\bullet \ \Rightarrow {\rm fit} \ {\rm OLS} \ {\rm models} \ {\rm within} \ {\rm LOOP}$
- Maybe  $\hat{y}(x)$  isn't that much better than other covariates (or, maybe it's useless)
  - $\bullet \ \Rightarrow {\sf use \ random \ forest \ within \ LOOP}$



- + If  $\hat{y}(\pmb{x})$  predicts Y really well, we would expect a linear relationship
  - $\bullet \ \Rightarrow {\rm fit} \ {\rm OLS} \ {\rm models} \ {\rm within} \ {\rm LOOP}$
- Maybe  $\hat{y}(x)$  isn't that much better than other covariates (or, maybe it's useless)
  - $\bullet \ \Rightarrow {\sf use \ random \ forest \ within \ LOOP}$
- Let the data decide!
  - pred=reloop



## **Incorporating Auxiliary Data in Practice**



- Y: outcome vector
- Tr : treatment assignment vector
- Z: matrix of covariates
- *pred* = *reloop* : specify auxiliary data interpolation algorithm
- *p*: probability of treatment
- yhat : vector of auxiliary predictions
- ...: optional inputs for interpolation algorithm

- AEIS data includes thousands of schools not in our RCT
- A great setting for integrating auxiliary and RCT data







- 1. Work through 03-integrate-aux.Rmd
  - We fit an auxiliary model and generate predictions to input as *yhat*
- 2. Apply what you learned in 04-effect-estABtest.Rmd
- 3. Flag any of us down as you have questions!

Conceptual Overview

Estimating Effects with RCT Data

Incorporating Auxiliary Data

## Break

Treatment Effect Heterogeneity

Planning Experiments



## Take a 15 minute break!

## See you back at 3:45 pm

**Conceptual Overview** 

Estimating Effects with RCT Data

Incorporating Auxiliary Data

Break

## Treatment Effect Heterogeneity

**Planning Experiments** 



- Traditionally, most problems in causal inference focus on the ATE as the estimand of interest.
- However, it is not unreasonable that the same treatment might have different effects on different individuals.
- As an average, the ATE **cannot** account for such potential variations.

Example



Example



• The conditional average treatment effect (CATE) is

$$\tau(x) = \mathbb{E}[\tau_i | X_i = x] = \mathbb{E}[y_i^t - y_i^c | X_i = x]$$



## **Conditional Average Treatment Effect**

• The conditional average treatment effect (CATE) is

$$\tau(x) = \mathbb{E}[\tau_i | X_i = x] = \mathbb{E}[y_i^t - y_i^c | X_i = x]$$

• The expected treatment effect conditional on having a specific set of covariate values.



• The conditional average treatment effect (CATE) is

 $\tau$ 

$$\mathbf{T}(x) = \mathbb{E}[\tau_i | X_i = x] = \mathbb{E}[y_i^t - y_i^c | X_i = x]$$

- The expected treatment effect conditional on having a specific set of covariate values.
- Unlike the ATE, the CATE accounts for the different characteristics of individuals as reflected by the covariates  $(X_i = x)$ .



#### **Estimating the Conditional Average Treatment Effect**

• Consider the following decomposition:



$$\tau(x) = \mathbb{E}[\tau_i | X_i = x] = \mathbb{E}[y_i^t - y_i^c | X_i = x]$$
$$= \mathbb{E}[y_i^t | X_i = x] - \mathbb{E}[y_i^c | X_i = x]$$
$$= \mathbb{E}[Y_i | X_i = x, T_i = 1] - \mathbb{E}[Y_i | X_i = x, T_i = 0]$$

#### **Estimating the Conditional Average Treatment Effect**

• Consider the following decomposition:



$$\begin{aligned} \tau(x) &= \mathbb{E}[\tau_i | X_i = x] = \mathbb{E}[y_i^t - y_i^c | X_i = x] \\ &= \mathbb{E}[y_i^t | X_i = x] - \mathbb{E}[y_i^c | X_i = x] \\ &= \mathbb{E}[Y_i | X_i = x, T_i = 1] - \mathbb{E}[Y_i | X_i = x, T_i = 0] \end{aligned}$$

- Regress observed outcomes on the covariates and treatment assignments.
- Estimate the CATE as the difference between estimated treatment ( $T_i = 1$ ) and control ( $T_i = 0$ ) outcomes.

#### Interaction Term(s) with Treatment Assignment

• Consider the following linear regression model:



$$\mu(x,t) = \mathbb{E}[Y_i|X_i = x, T_i = t] = \beta_0 + \beta_1 x + \beta_2 t + \beta_3 (x \cdot t)$$

#### Interaction Term(s) with Treatment Assignment

• Consider the following linear regression model:

$$\mu(x,t) = \mathbb{E}[Y_i|X_i = x, T_i = t] = \beta_0 + \beta_1 x + \beta_2 t + \beta_3 (x \cdot t)$$



• Thus, we can estimate the CATE as:

$$\hat{\tau}(x) = \hat{\mu}(x,1) - \hat{\mu}(x,0) = (\hat{\beta}_0 + \hat{\beta}_1 x + \hat{\beta}_2 + \hat{\beta}_3 x) - (\hat{\beta}_0 + \hat{\beta}_1 x) = \hat{\beta}_2 + \hat{\beta}_3 x$$

#### Interaction Term(s) with Treatment Assignment

• Consider the following linear regression model:

$$\mu(x,t) = \mathbb{E}[Y_i|X_i = x, T_i = t] = \beta_0 + \beta_1 x + \beta_2 t + \beta_3 (x \cdot t)$$



• Thus, we can estimate the CATE as:

$$\hat{\tau}(x) = \hat{\mu}(x,1) - \hat{\mu}(x,0) = (\hat{\beta}_0 + \hat{\beta}_1 x + \hat{\beta}_2 + \hat{\beta}_3 x) - (\hat{\beta}_0 + \hat{\beta}_1 x) = \hat{\beta}_2 + \hat{\beta}_3 x$$

- Linear parametric model: Ease of interpretation and statistical inference.
- However, strict linearity is also restrictive and thus potentially imprecise.

## Machine Learning for Estimating Heterogeneous Treatment Effects



- Broadly speaking, there are two categories:
  - **Meta-learners:** Methods that leverage off-the-shelf machine learning algorithms to **indirectly** estimate the CATE by learning its components. *Example:* S-, T-, & X-Learner (Künzel et al. 2019).
  - **Direct Estimators:** Methods specifically designed from the ground up to **directly** estimate the CATE. *Example:* Causal Forest (Wager and Athey 2018).

## Machine Learning for Estimating Heterogeneous Treatment Effects

- Broadly speaking, there are two categories:
  - **Meta-learners:** Methods that leverage off-the-shelf machine learning algorithms to **indirectly** estimate the CATE by learning its components. *Example:* S-, T-, & X-Learner (Künzel et al. 2019).
  - **Direct Estimators:** Methods specifically designed from the ground up to **directly** estimate the CATE. *Example:* Causal Forest (Wager and Athey 2018).
- More flexible machine learning methods mean **potentially more precise** estimates but at the cost of **interpretability and inference**.



- **Accuracy** is undeniably important. Knowing how much a treatment will help or harm someone to the best of our ability is always good...
- ...but so is **Interpretability**, especially since we are conditioning on and considering specific values of covariates.
- In other words, how to have our cake and eat (a bit of) it too?

## Leveraging Estimates of Individual Treatment Effects

• Recall the definition of the CATE:



$$\tau(x) = \mathbb{E}[\tau_i | X_i = x] = \mathbb{E}[y_i^t - y_i^c | X_i = x]$$

• If we know the true individual treatment effects, we can fit a parametric model estimating  $\tau_i$  with  $X_i$  to estimate the CATE.
• Recall the definition of the CATE:



$$\tau(x) = \mathbb{E}[\tau_i | X_i = x] = \mathbb{E}[y_i^t - y_i^c | X_i = x]$$

- If we know the true individual treatment effects, we can fit a parametric model estimating  $\tau_i$  with  $X_i$  to estimate the CATE.
- Unfortunately,  $\tau_i$  is **never available** as we exclusively observe **either**  $y_i^t$  **or**  $y_i^c$ .

• Recall the definition of the CATE:



- If we know the true individual treatment effects, we can fit a parametric model estimating τ<sub>i</sub> with X<sub>i</sub> to estimate the CATE.
- Unfortunately,  $\tau_i$  is **never available** as we exclusively observe **either**  $y_i^t$  **or**  $y_i^c$ .
- **However**, we have the estimate  $\hat{\tau}_i$  from the LOOP estimator:

$$\hat{\tau}_i = \{Y_i - [(1-p)\hat{y}_i^t + p\hat{y}_i^c]\}\frac{T_i - p}{p(1-p)}$$



• Given a Bernoulli randomization,  $\hat{\tau}_i$  is an **unbiased** estimate of  $\tau_i$  ( $\mathbb{E}[\hat{\tau}_i] = \tau_i$ ) In addition, if the imputations  $y_i^t$  and  $y_i^c$  are accurate,  $\hat{\tau}_i$  will also be a **precise** estimate ( $\hat{\tau}_i \approx \tau_i$ ) (Wu and Gagnon-Bartsch 2018).

- Given a Bernoulli randomization,  $\hat{\tau}_i$  is an **unbiased** estimate of  $\tau_i$  ( $\mathbb{E}[\hat{\tau}_i] = \tau_i$ ) In addition, if the imputations  $y_i^t$  and  $y_i^c$  are accurate,  $\hat{\tau}_i$  will also be a **precise** estimate ( $\hat{\tau}_i \approx \tau_i$ ) (Wu and Gagnon-Bartsch 2018).
- We first estimate the true individual treatment effects using the LOOP estimator.

- Given a Bernoulli randomization,  $\hat{\tau}_i$  is an **unbiased** estimate of  $\tau_i$  ( $\mathbb{E}[\hat{\tau}_i] = \tau_i$ ) In addition, if the imputations  $y_i^t$  and  $y_i^c$  are accurate,  $\hat{\tau}_i$  will also be a **precise** estimate ( $\hat{\tau}_i \approx \tau_i$ ) (Wu and Gagnon-Bartsch 2018).
- We first estimate the true individual treatment effects using the LOOP estimator.
- We can regress  $\hat{\tau}_i$  on  $X_i$  and estimate the CATE as  $\tau(x) = \mathbb{E}[\hat{\tau}_i | X_i = x]$ , using the estimated effect  $\hat{\tau}_i$  as a proxy for the true effect  $\tau_i$ :

$$\hat{\tau}(x) = \hat{\alpha}_1 + \hat{\alpha}_2 x$$

- Given a Bernoulli randomization,  $\hat{\tau}_i$  is an **unbiased** estimate of  $\tau_i$  ( $\mathbb{E}[\hat{\tau}_i] = \tau_i$ ) In addition, if the imputations  $y_i^t$  and  $y_i^c$  are accurate,  $\hat{\tau}_i$  will also be a **precise** estimate ( $\hat{\tau}_i \approx \tau_i$ ) (Wu and Gagnon-Bartsch 2018).
- We first estimate the true individual treatment effects using the LOOP estimator.
- We can regress  $\hat{\tau}_i$  on  $X_i$  and estimate the CATE as  $\tau(x) = \mathbb{E}[\hat{\tau}_i | X_i = x]$ , using the estimated effect  $\hat{\tau}_i$  as a proxy for the true effect  $\tau_i$ :

$$\hat{\tau}(x) = \hat{\alpha}_1 + \hat{\alpha}_2 x$$

• Notice that this formulation is (roughly) equivalent to the interaction model:

$$\hat{\tau}(x) = \hat{\mu}(x,1) - \hat{\mu}(x,0) = (\hat{\beta}_0 + \hat{\beta}_1 x + \hat{\beta}_2 + \hat{\beta}_3 x) - (\hat{\beta}_0 + \hat{\beta}_1 x) = \hat{\beta}_2 + \hat{\beta}_3 x$$

• If the LOOP estimator's requirements are satisfied, the estimates will be unbiased – **even with poor-fitting models**.



• If the LOOP estimator's requirements are satisfied, the estimates will be unbiased – **even with poor-fitting models**.



• Thus, we do not have to worry about bias from the regression in the second stage carrying over or, worse, amplifying bias from the first.

• If the LOOP estimator's requirements are satisfied, the estimates will be unbiased – even with poor-fitting models.



- Thus, we do not have to worry about bias from the regression in the second stage carrying over or, worse, amplifying bias from the first.
- Flexible model(s) to estimate the ITE in the first: More precise than a strictly linear model with interaction(s).
- Parametric model to estimate the CATE in the second: More interpretable than a powerful but non-parametric model.





• This function will work for an estimator built with or without auxiliary data, which allows us to improve precision further.

- This function will work for an estimator built with or without auxiliary data, which allows us to improve precision further.
- However, it is currently only for Bernoulli-randomized experiments.

- This function will work for an estimator built with or without auxiliary data, which allows us to improve precision further.
- However, it is currently only for Bernoulli-randomized experiments.
- Once you have retrieve the estimates, choose your favorite model and do some regressing!



- 1. Work through 05-heterogeneousEffects.Rmd
  - We fit retrieve ITE estimates from the model in 04-effect-estABtest.Rmd.
  - We then estimate the CATE by regressing these estimates on the covariates.
- 2. Flag any of us down as you have questions!

Conceptual Overview

Estimating Effects with RCT Data

Incorporating Auxiliary Data

Break

Treatment Effect Heterogeneity

### **Planning Experiments**

- We'll be using the *dRCTpower* package to plan experiments
- Main function is *run\_app*
- You can download the package in R using the following commands:

install.packages("devtools")
devtools::install\_github("jaylinlowe/dRCTpower")

• We will be using the *aux\_dat\_small.csv* file from the Github repo





# How to choose a sample size for our experiment, particularly if auxiliary data will be incorporated?



• Incorporating auxiliary data in our analysis can improve precision, meaning we can have a smaller sample size with the same power



- Incorporating auxiliary data in our analysis can improve precision, meaning we can have a smaller sample size with the same power
- Gain in precision is determined by how predictive a model fit on the auxiliary data is for the RCT



- Incorporating auxiliary data in our analysis can improve precision, meaning we can have a smaller sample size with the same power
- Gain in precision is determined by how predictive a model fit on the auxiliary data is for the RCT
- But....we don't have the RCT data!



#### 1. Break auxiliary dataset into subgroups



- 1. Break auxiliary dataset into subgroups
- 2. For each subgroup, treat it as the RCT and the rest of the data as the auxiliary data



- 1. Break auxiliary dataset into subgroups
- 2. For each subgroup, treat it as the RCT and the rest of the data as the auxiliary data
- 3. Calculate the required sample size under this framework



Large auxiliary dataset that:

• is substantially larger than the RCT will be



Large auxiliary dataset that:

- is substantially larger than the RCT will be
- has covariates



Large auxiliary dataset that:

- is substantially larger than the RCT will be
- has covariates
- has the same outcome of interest as the RCT



• Method is only plausible if it's reasonable to assume the RCT looks like some subgroup of the auxiliary data, even if we don't know what subgroup that is



- Method is only plausible if it's reasonable to assume the RCT looks like some subgroup of the auxiliary data, even if we don't know what subgroup that is
- Dangerous to assume RCT looks like any one subgroup



- Method is only plausible if it's reasonable to assume the RCT looks like some subgroup of the auxiliary data, even if we don't know what subgroup that is
- Dangerous to assume RCT looks like any one subgroup
- Dangerous to choose most optimistic option

$$n = 2\sigma^2 \frac{(\xi_{1-\alpha/2} + \xi_{1-\beta})^2}{\Delta_A^2}$$



$$n = 2\sigma^2 \frac{(\xi_{1-\alpha/2} + \xi_{1-\beta})^2}{\Delta_A^2}$$



•  $\xi_{1-\alpha/2}$  is the critical value obtained from a normal distribution for Type I error equal to  $\alpha$ .

$$n = 2\sigma^2 \frac{(\xi_{1-\alpha/2} + \xi_{1-\beta})^2}{\Delta_A^2}$$



- $\xi_{1-\alpha/2}$  is the critical value obtained from a normal distribution for Type I error equal to  $\alpha$ .
- $\xi_{1-\beta}$  is the critical value from a normal distribution for Type II error rate  $\beta$ .

$$n = 2\sigma^2 \frac{(\xi_{1-\alpha/2} + \xi_{1-\beta})^2}{\Delta_A^2}$$



- $\xi_{1-\alpha/2}$  is the critical value obtained from a normal distribution for Type I error equal to  $\alpha$ .
- $\xi_{1-\beta}$  is the critical value from a normal distribution for Type II error rate  $\beta$ .
- $\Delta_A$  is the effect size, typically 20% of the standard deviation of the outcome in the population

$$n = 2\sigma^2 \frac{(\xi_{1-\alpha/2} + \xi_{1-\beta})^2}{\Delta_A^2}$$



- $\xi_{1-\alpha/2}$  is the critical value obtained from a normal distribution for Type I error equal to  $\alpha$ .
- $\xi_{1-\beta}$  is the critical value from a normal distribution for Type II error rate  $\beta$ .
- $\Delta_A$  is the effect size, typically 20% of the standard deviation of the outcome in the population
- $\sigma^2$  is the true variance of the outcome in the population, typically replaced with an estimate from a sample



• We replace  $\sigma^2$  with an estimate from each subgroup



- We replace  $\sigma^2$  with an estimate from each subgroup
- Shiny app gives two estimates, one if you were to use auxiliary data in analysis, and one without



- We replace  $\sigma^2$  with an estimate from each subgroup
- Shiny app gives two estimates, one if you were to use auxiliary data in analysis, and one without
- "Without auxiliary data" estimate is variance of outcome for that subgroup



- We replace  $\sigma^2$  with an estimate from each subgroup
- Shiny app gives two estimates, one if you were to use auxiliary data in analysis, and one without
- "Without auxiliary data" estimate is variance of outcome for that subgroup
- "With auxiliary data" estimate is variance of the residuals,  $(y_i \hat{y}_i)$ , where  $\hat{y}_i$  are out-of-bag predictions from model

Three options:

- 1. Categorical Variable
  - Divide based on levels of categorical variable
  - Can create your own categorical variables



Three options:

- 1. Categorical Variable
  - Divide based on levels of categorical variable
  - Can create your own categorical variables
- 2. Numerical Variable
  - Divide into 10 (adjustable) equally sized groups
  - May need to divide into fewer if there isn't enough variation



Three options:

- 1. Categorical Variable
  - Divide based on levels of categorical variable
  - Can create your own categorical variables
- 2. Numerical Variable
  - Divide into 10 (adjustable) equally sized groups
  - May need to divide into fewer if there isn't enough variation
- 3. Best-Worst Case Scenario
  - Divide based on how predictive we expect the auxiliary model to be for that group
  - Good starting point





# Shiny App Demo



## References

Gagnon-Bartsch, Johann A., Adam C. Sales, Edward Wu, Anthony F. Botelho, John A. Erickson, Luke W. Miratrix and Neil T. Heffernan. 2023. "Precise unbiased estimation in randomized experiments using auxiliary observational data." *Journal of Causal Inference* 11(1):20220011.

URL: https://www.degruyter.com/document/doi/10.1515/jci-2022-0011/html

Künzel, Sören R, Jasjeet S Sekhon, Peter J Bickel and Bin Yu. 2019. "Metalearners for estimating heterogeneous treatment effects using machine learning." *Proceedings of the National Academy of Sciences* 116(10):4156–4165.

Lowe, Jaylin, Charlotte Mann, Jiaying Wang, Adam Sales and Johann Gagnon-Bartsch. Forthcoming. "Power Calculations for Randomized Controlled Trials with Auxiliary Observational Data." *EDM 2024*.



Mann, Charlotte, Jiaying Wang, Adam Sales and Johann Gagnon-Bartsch. Forthcoming. "Using Publicly Available Auxiliary Data to Improve Precision of Treatment Effect Estimation in a Randomized Efficacy Trial." *EDM* 2024 .

Pane, John F., Beth Ann Griffin, Daniel F. McCaffrey and Rita Karam. 2014. "Effectiveness of Cognitive Tutor Algebra I at Scale." *Educational Evaluation* and Policy Analysis 36(2):127–144.

**URL:** https://doi.org/10.3102/0162373713507480

- Pham, Duy, Kirk Vanacore, Adam Sales and Johann Gagnon-Bartsch. Forthcoming. "LOOL: Towards Personalization with Flexible Robust Estimation of Heterogeneous Treatment Effects." *EDM* 2024 .
- Sales, Adam C, Ethan B Prihar, Johann A Gagnon-Bartsch and Neil T Heffernan. 2023. "Using Auxiliary Data to Boost Precision in the Analysis of A/B Tests on an Online Educational Platform: New Data and New Results." *arXiv preprint arXiv*:2306.06273.



Wager, Stefan and Susan Athey. 2018. "Estimation and Inference of Heterogeneous Treatment Effects using Random Forests." *Journal of the American Statistical Association* 113(523):1228–1242.

Wu, Edward and Johann A. Gagnon-Bartsch. 2018. "The LOOP Estimator: Adjusting for Covariates in Randomized Experiments." *Evaluation Review* 42(4):458–488. Publisher: SAGE Publications Inc.
URL: https://doi.org/10.1177/0193841X18808003

Wu, Edward and Johann A. Gagnon-Bartsch. 2021. "Design-Based Covariate Adjustments in Paired Experiments." *Journal of Educational and Behavioral Statistics* 46(1):109–132. Publisher: American Educational Research Association.

URL: https://doi.org/10.3102/1076998620941469

