

# Difference-in-Differences using Mixed-Integer Programming Matching Approach

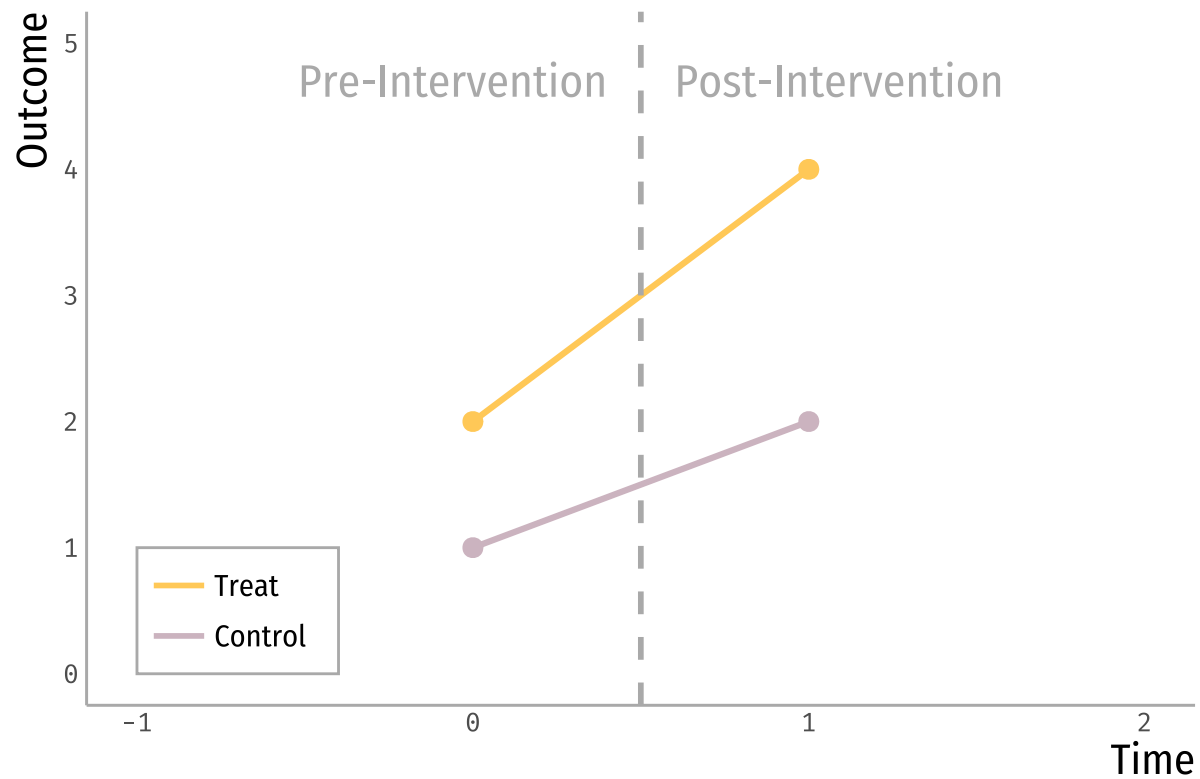
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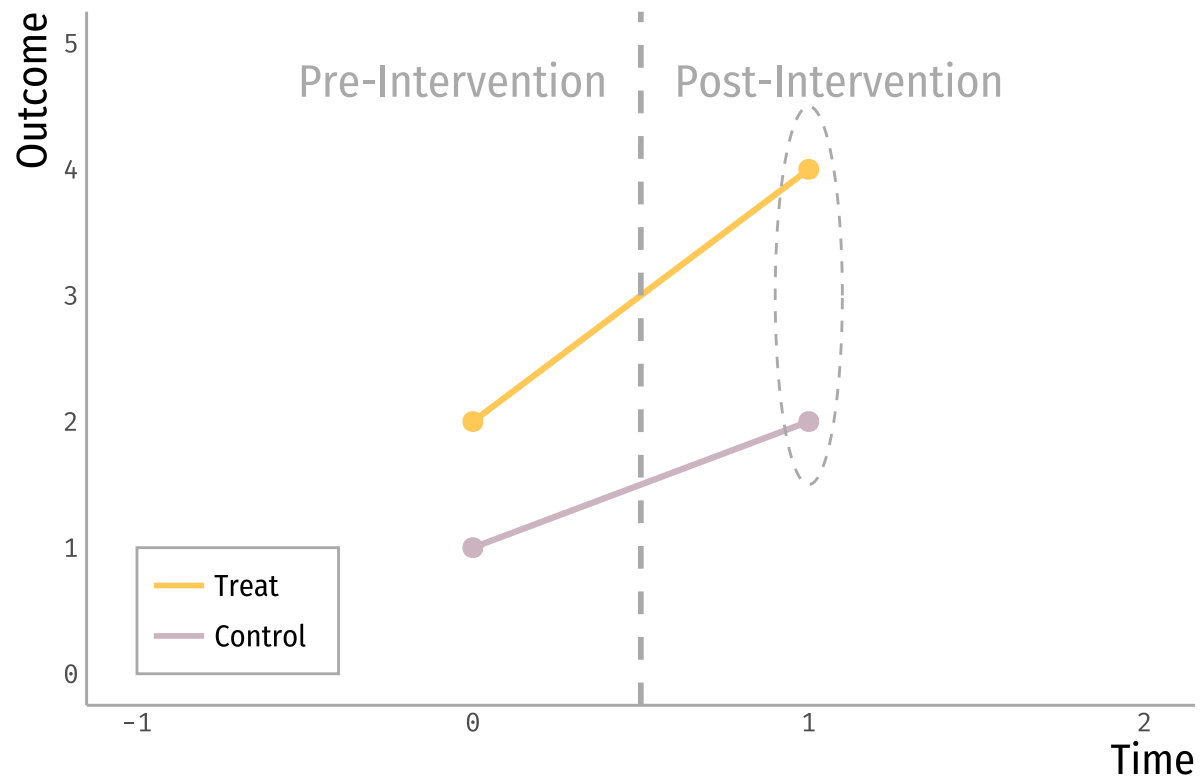
Seminario DIIS - PUC

November 20th, 2024

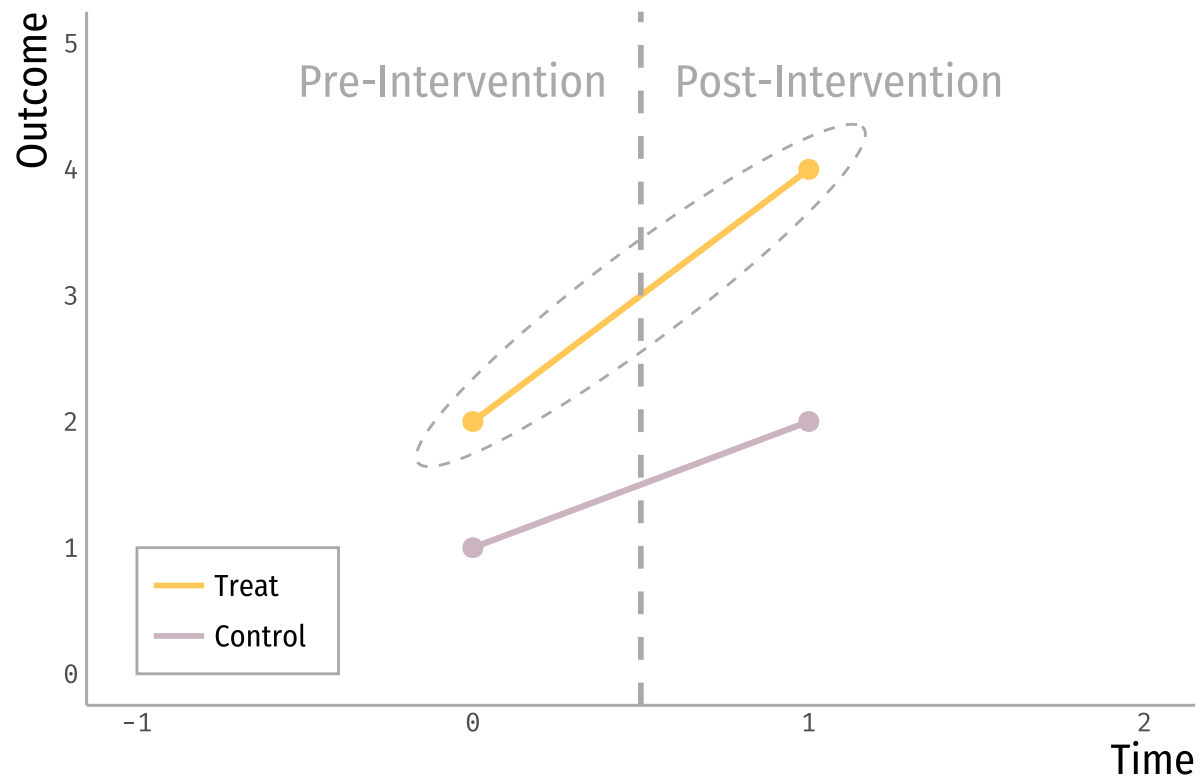
# Diff-in-Diff as an identification strategy



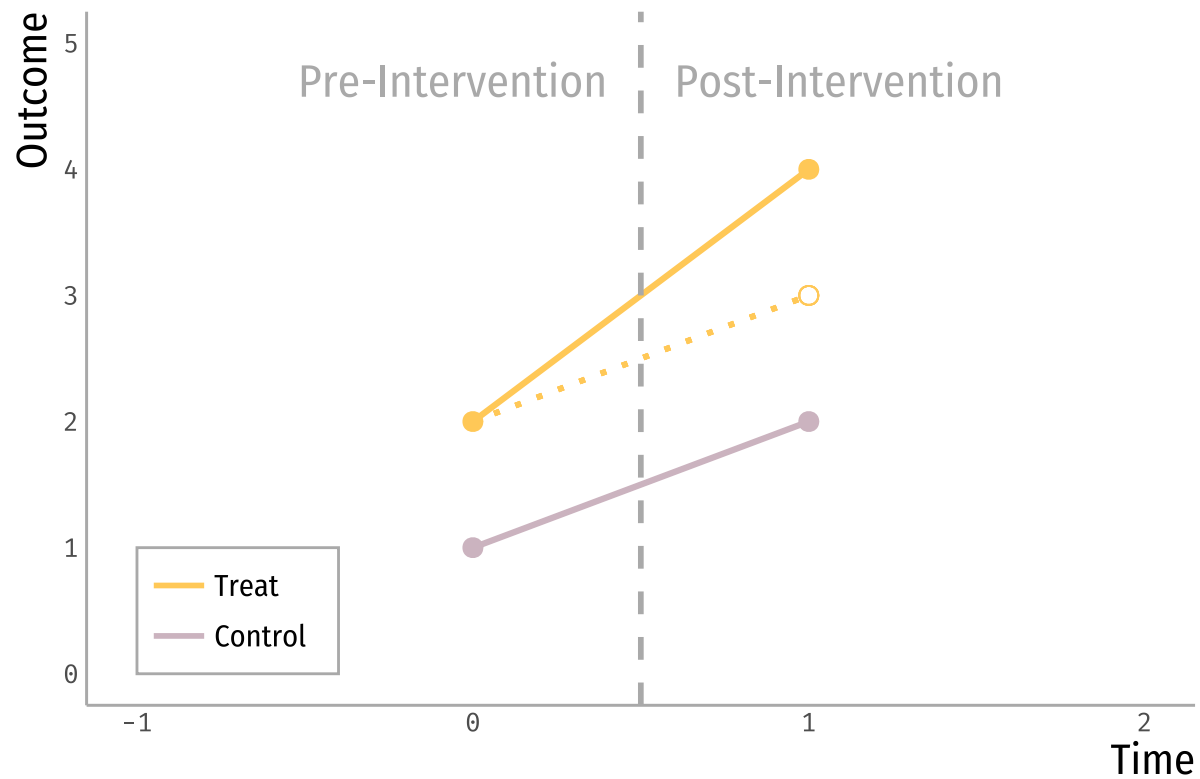
# Cannot compare treated vs control



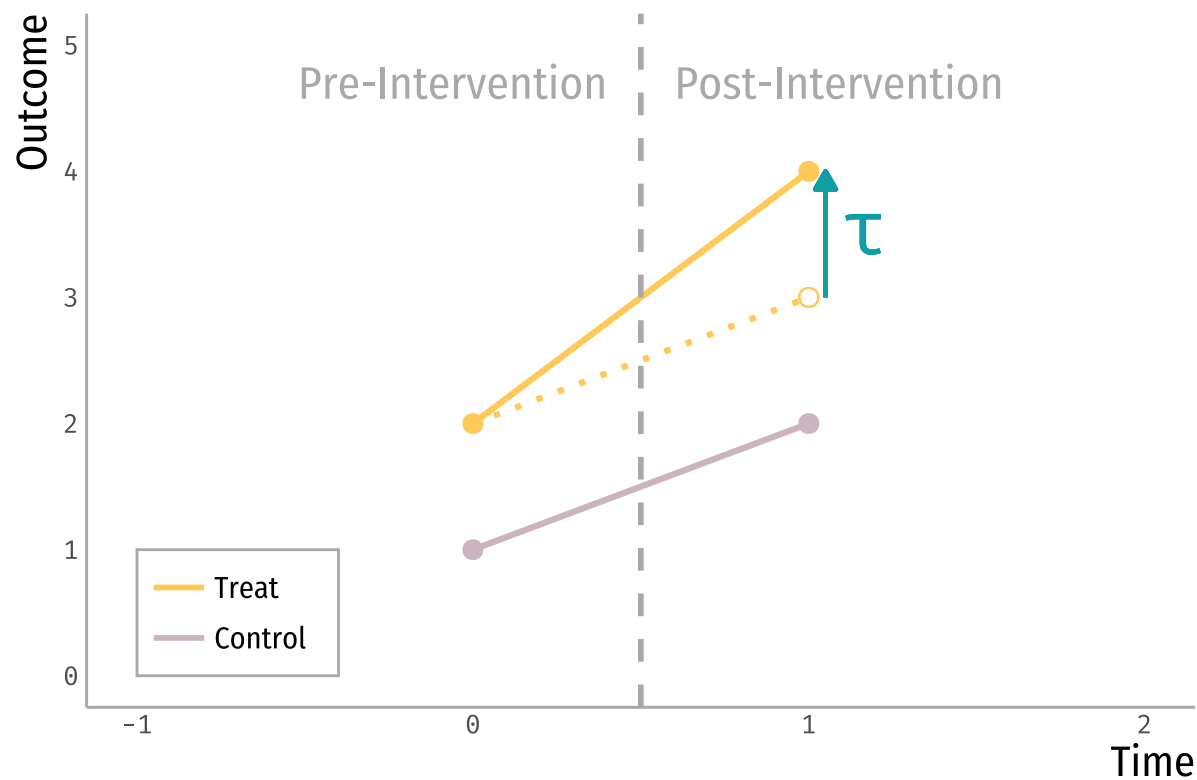
# Cannot compare before and after



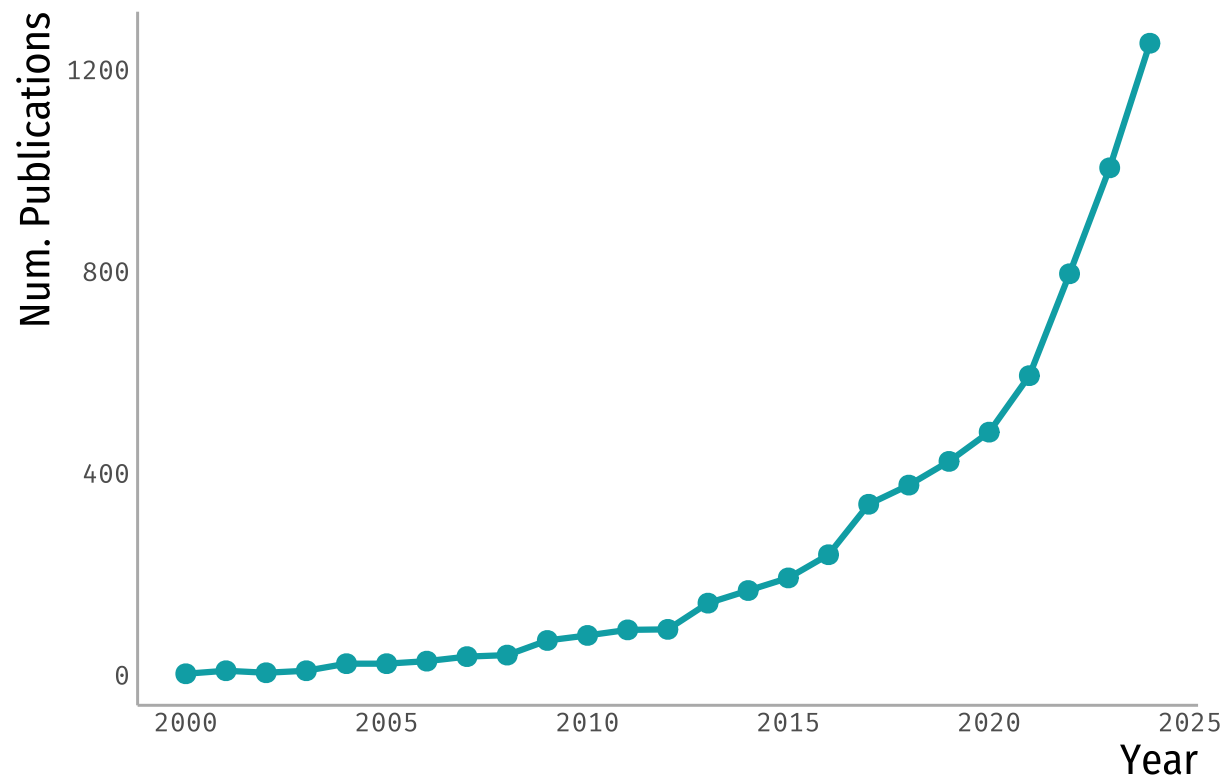
# Parallel trend assumption (PTA)



# Estimate Average Treatment Effect on the Treated (ATT)

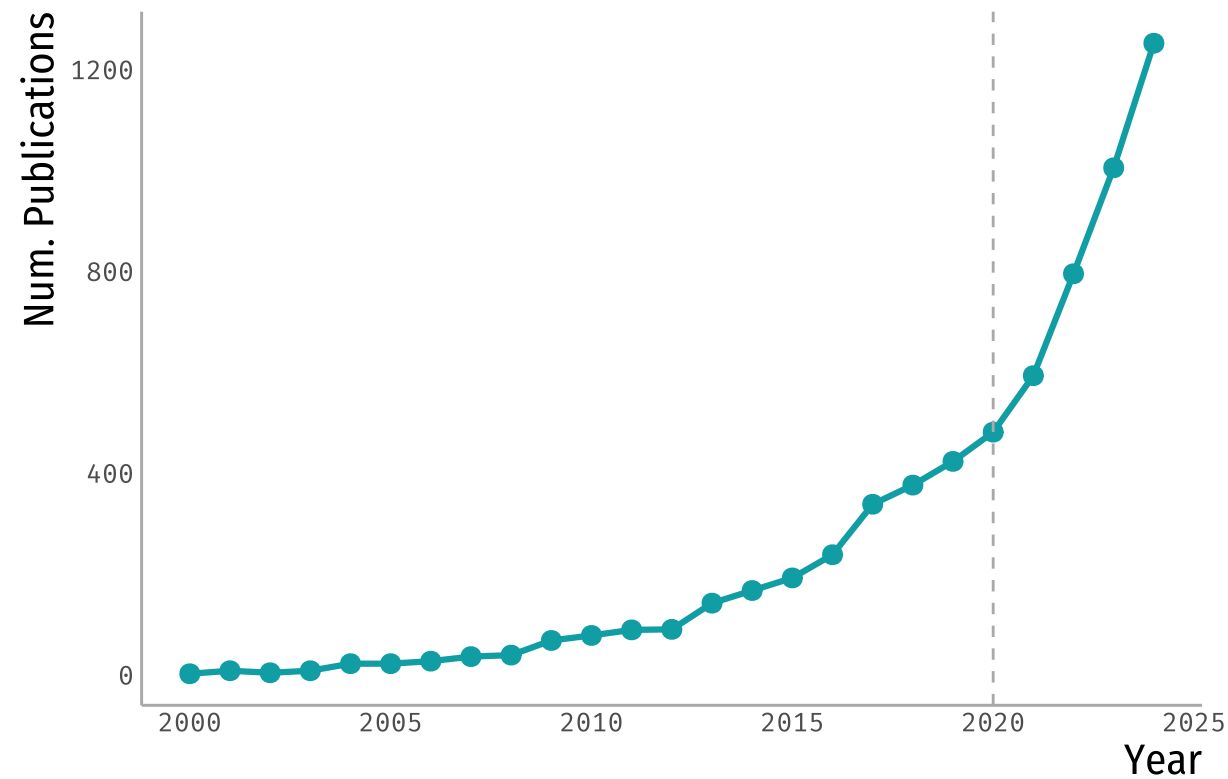


# Diff-in-Diff is very popular in Economics



Source: Web of Science (11/18/2024)

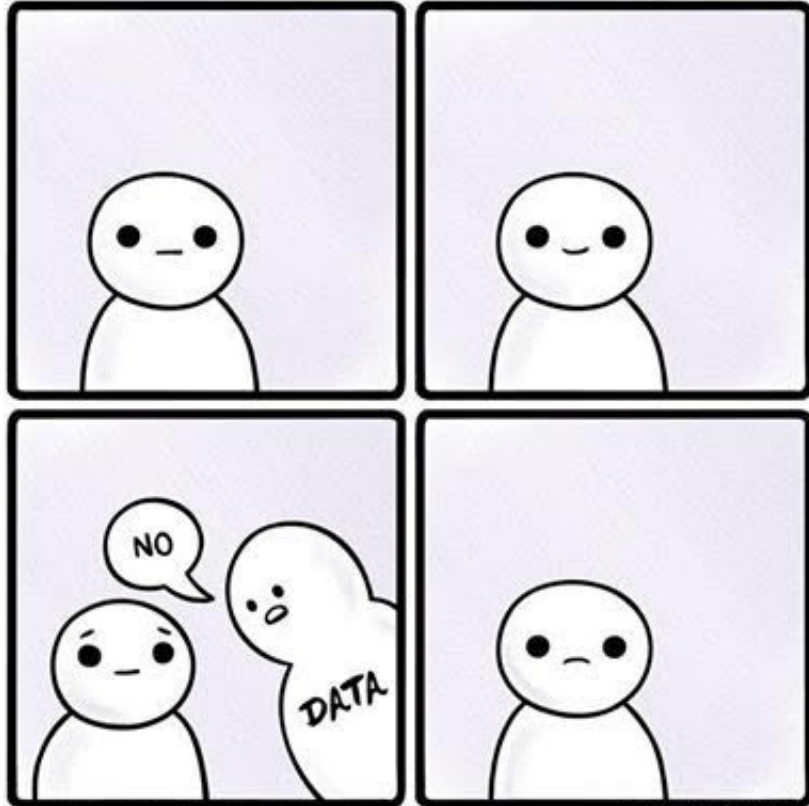
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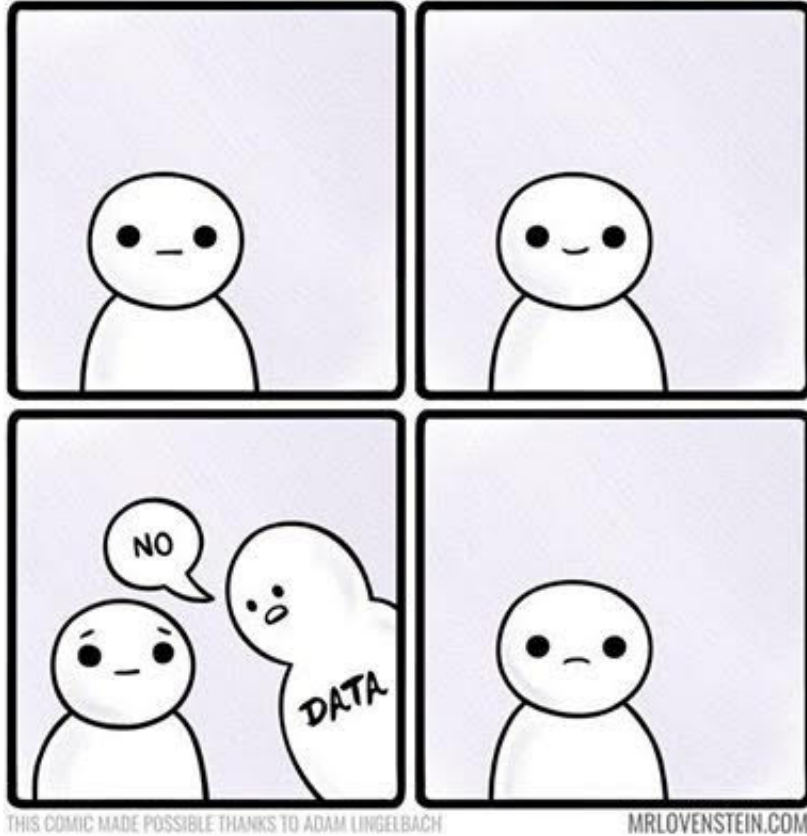
# What about parallel trends?



THIS COMIC MADE POSSIBLE THANKS TO ADAM LINGELBACH

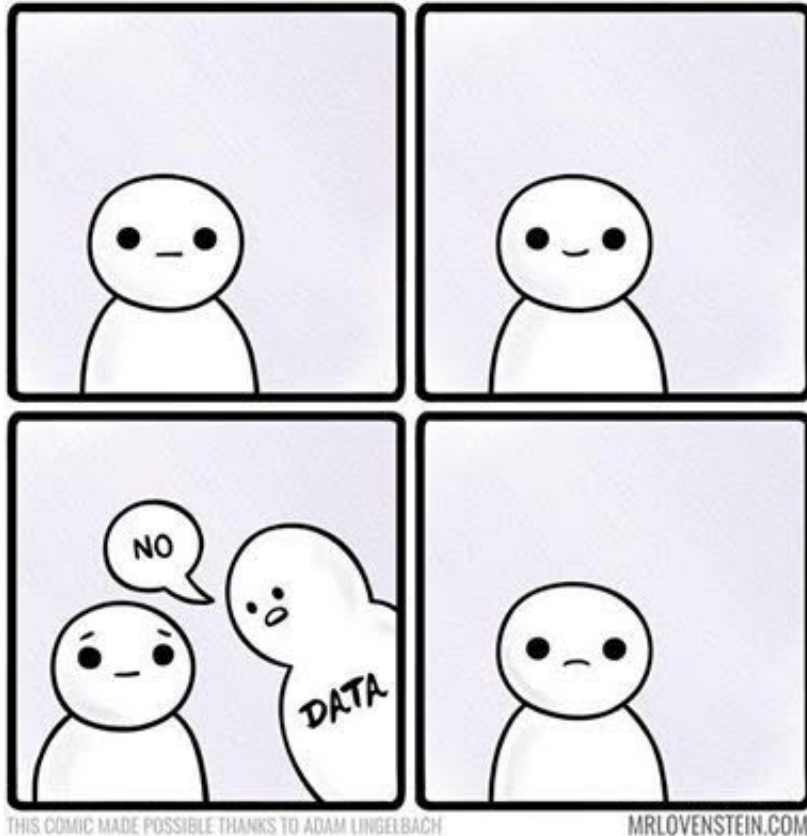
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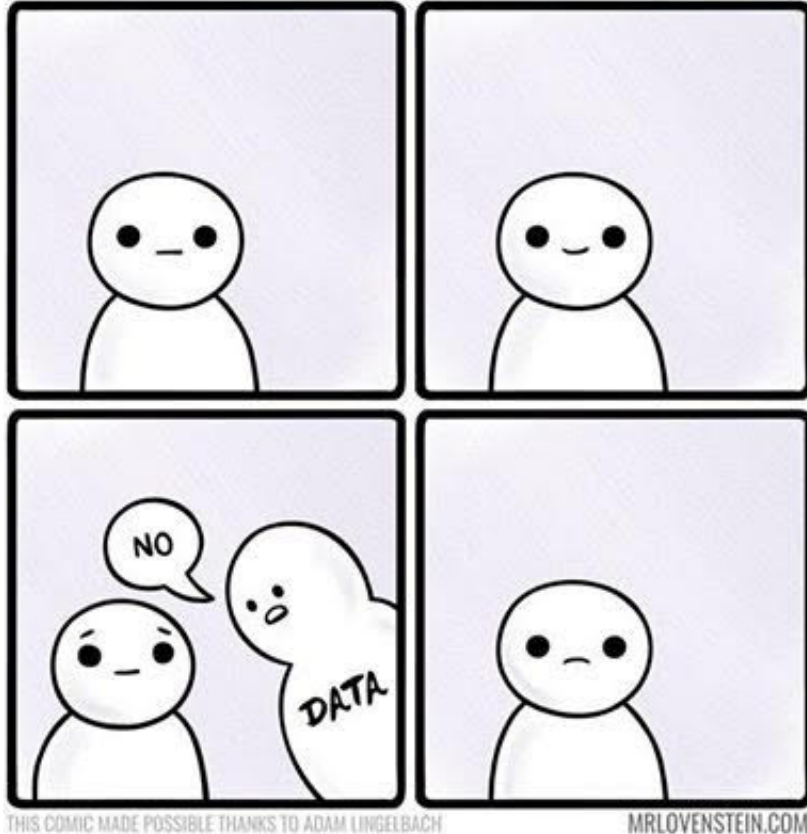
- Main identification assumption **fails**

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- Find sub-groups that potentially **follow PTA**
  - E.g. similar units in treatment and control
  - Similar to synthetic control intuition.

# What about parallel trends?



- Main identification assumption **fails**
- Find sub-groups that potentially **follow PTA**
  - E.g. similar units in treatment and control
  - Similar to synthetic control intuition.
- Can matching help?
  - It's **complicated** (Ham & Miratrix, 2022; Zeldow & Hatfield, 2021; Basu & Small, 2020; Lindner & McConnell, 2018; Daw & Hatfield, 2018 (x2); Ryan, 2018; Ryan et al., 2018)

# This paper

- Identify contexts when matching can recover causal estimates under **certain violations of the parallel trend assumption**.
  - Overall bias reduction and increase in robustness for sensitivity analysis.
- Use **mixed-integer programming matching (MIP)** to balance covariates directly.

## Simulations:

Different DGP scenarios

## Application:

School segregation & vouchers

Let's get started

# DD Setup

- Let  $Y_{it}(z)$  be the potential outcome for unit  $i$  in period  $t$  under treatment  $z$ .
- Intervention implemented in  $T_0 \rightarrow$  No units are treated in  $t \leq T_0$
- Difference-in-Differences (DD) focuses on ATT for  $t > T_0$ :

$$ATT(t) = E[Y_{it}(1) - Y_{it}(0) | Z = 1]$$

Expected difference in potential outcomes  
*if the treatment hadn't happened*  
for the treatment group

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- **Assumptions for DD:**
  - Parallel-trend assumption (PTA)
  - Common shocks

$$E[Y_{i1}(0) - Y_{i0}(0)|Z = 1] = E[Y_{i1}(0) - Y_{i0}(0)|Z = 0]$$



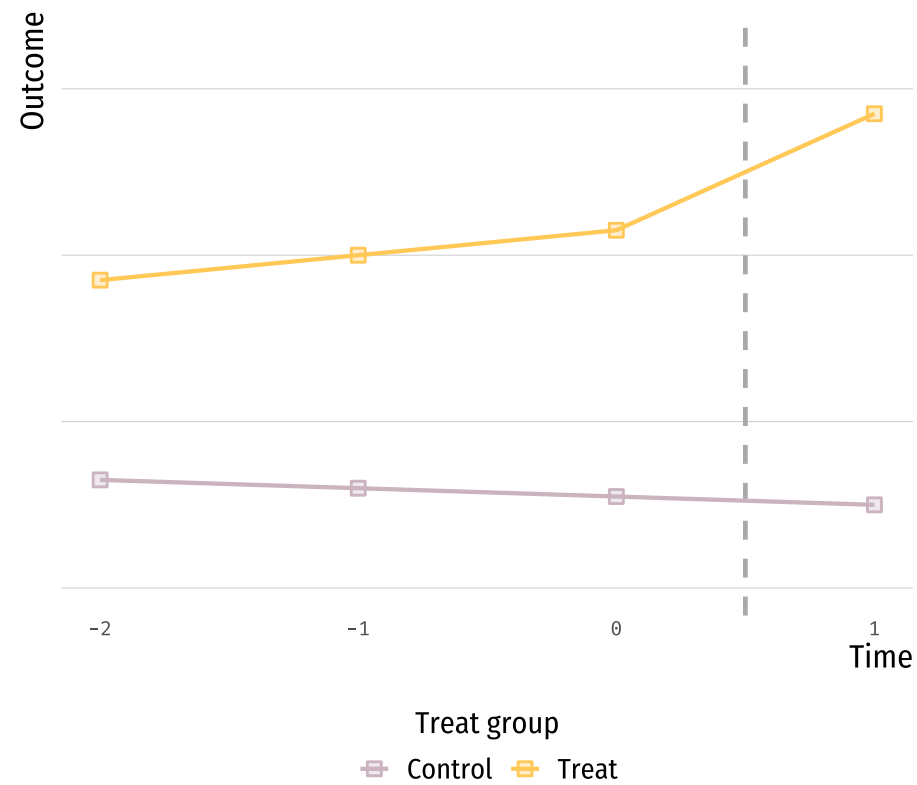
# DD Setup (cont.)

- Under these assumptions:

$$\hat{\tau}^{DD} = \overbrace{E[Y_{i1}|Z=1] - E[Y_{i1}|Z=0]}^{\Delta_{post}} - \underbrace{(E[Y_{i0}|Z=1] - E[Y_{i0}|Z=0])}_{\Delta_{pre}}$$

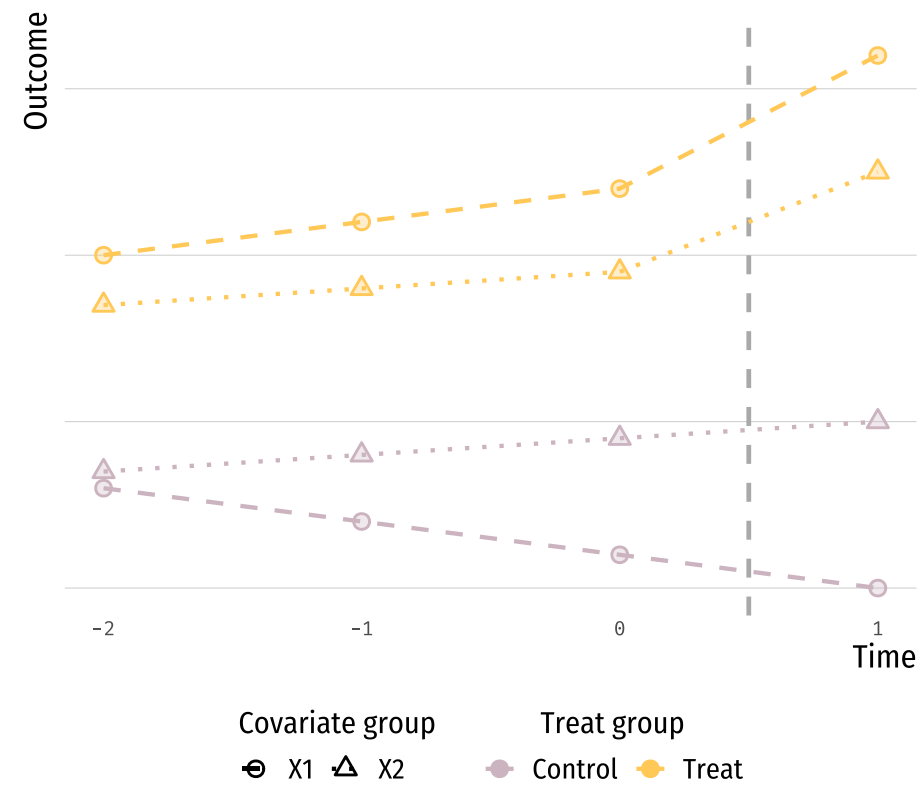
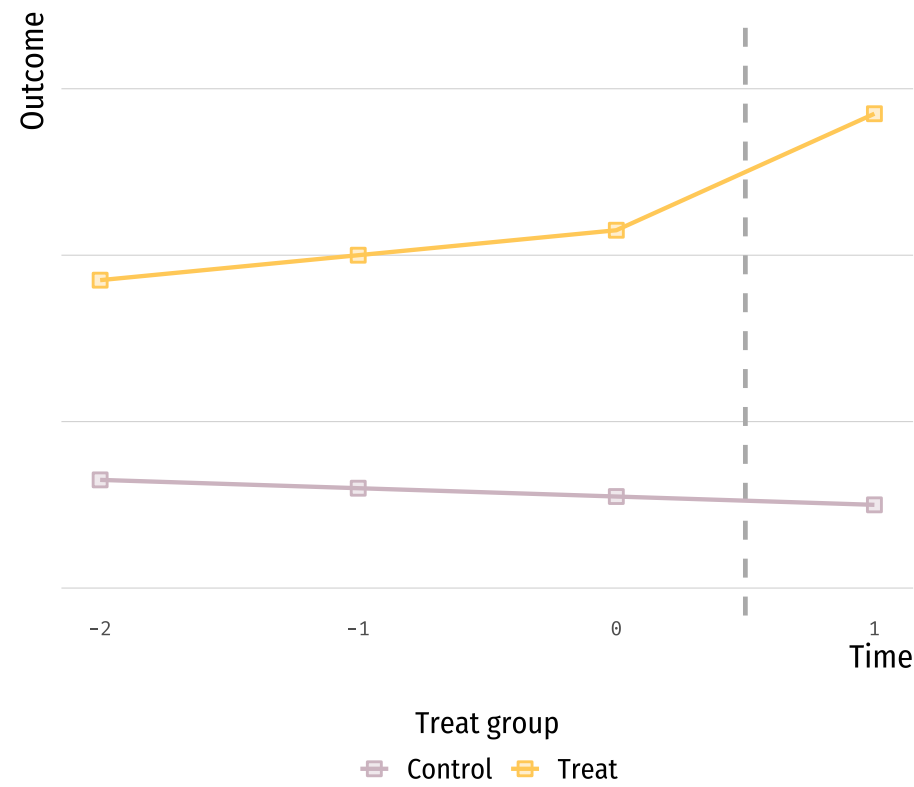
- Where  $t = 0$  and  $t = 1$  are the pre- and post-intervention periods, respectively.
- $Y_{it} = Y_{it}(1) \cdot Z_i + (1 - Z_i) \cdot Y_{it}(0)$  is the observed outcome.

# But what if the PTA doesn't hold?



# But what if the PTA doesn't hold?

We can potentially remove [part of] the bias by matching on  $X^s_{it}=X_i$



# General form of potential outcomes

We can write a general form of the potential outcomes  $Y(0)$  and  $Y(1)$  as follows:

$$Y_{it}(0) = \alpha_i + \lambda_t + \gamma_0(X_i) + \gamma_1(X_i, t) + \gamma_2(X_i, t) \cdot Z_i + u_{it}$$

$$Y_{it}(1) = Y_{it}(0) + \tau_{it} = \alpha_i + \lambda_t + \gamma_0(X_i) + \gamma_1(X_i, t) + \gamma_2(X_i, t) \cdot Z_i + \tau_{it} + u_{it}$$

Covariate distribution between groups can be **different**

$$X_i|Z \sim F_x(z)$$

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- $\gamma_1(X_i, t)$  is a time-dependent function.
- $\gamma_2(X_i, t) \cdot Z_i$  is a *differential* time-dependent function **only** for the treatment group.



# If the PTA holds...

Then, for a 2x2 DD, where  $t < T_0$  (pre) and  $t' > T_0$  (post):

$$\mathbb{E}[\gamma_1(X_i, t') + \gamma_2(X_i, t') - \gamma_1(X_i, t) - \gamma_2(X_i, t) | Z = 1] = \mathbb{E}[\gamma_1(X_i, t') - \gamma_1(X_i, t) | Z = 0]$$

One of the two conditions need to hold:

- 1) No effect or constant effect of  $X$  on  $Y$  over time:  $\mathbb{E}[\gamma_1(X, t)] = \mathbb{E}[\gamma_1(X)]$
- 2) Equal distribution of observed covariates between groups:  $X_i | Z = 1 \stackrel{d}{=} X_i | Z = 0$

in addition to:

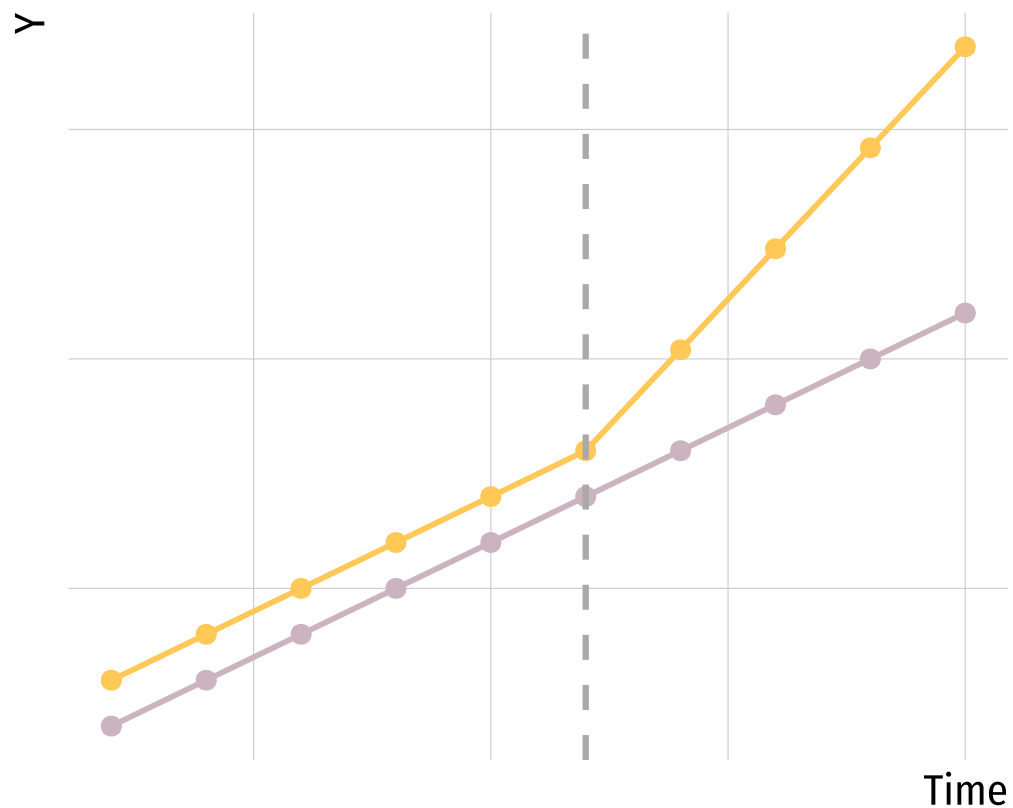
- 3) No differential time effect of  $X$  on  $Y$  by treatment group:  $\mathbb{E}[\gamma_2(X, t)] = 0$

Cond. 2 can hold through matching

Cond. 3 can be tested with sensitivity analysis

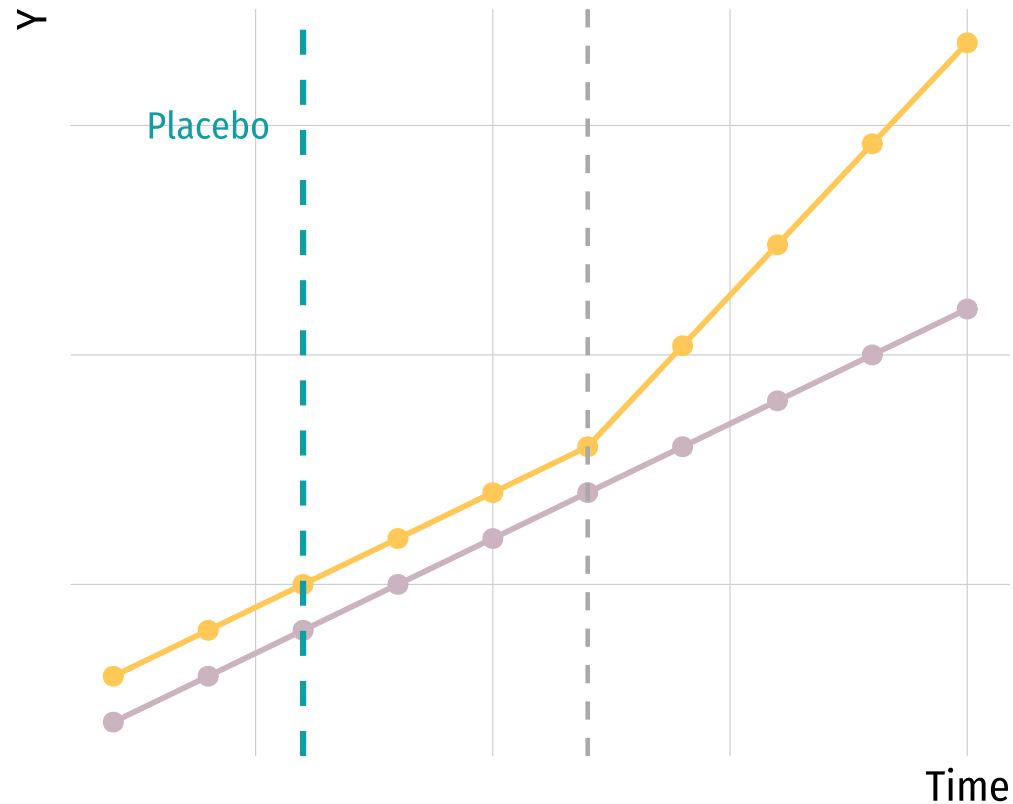
# Sensitivity analysis for Diff-in-Diff

- Use of **pre-trends** to test plausibility of the PTA:



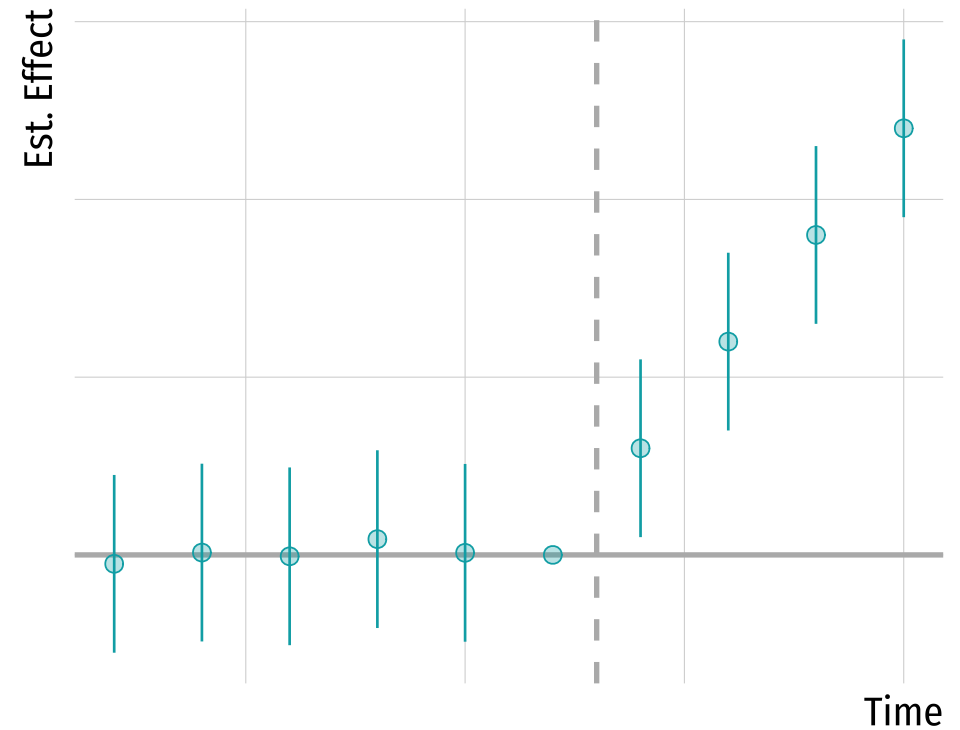
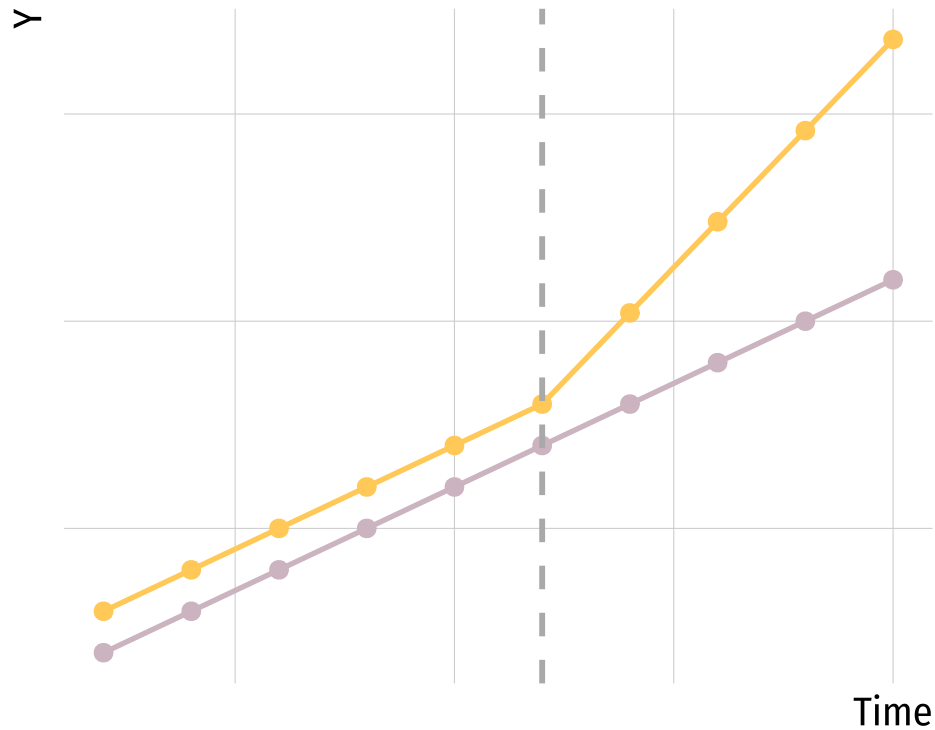
# Sensitivity analysis for Diff-in-Diff

- Using a diff-in-diff strategy, we shouldn't find an effect



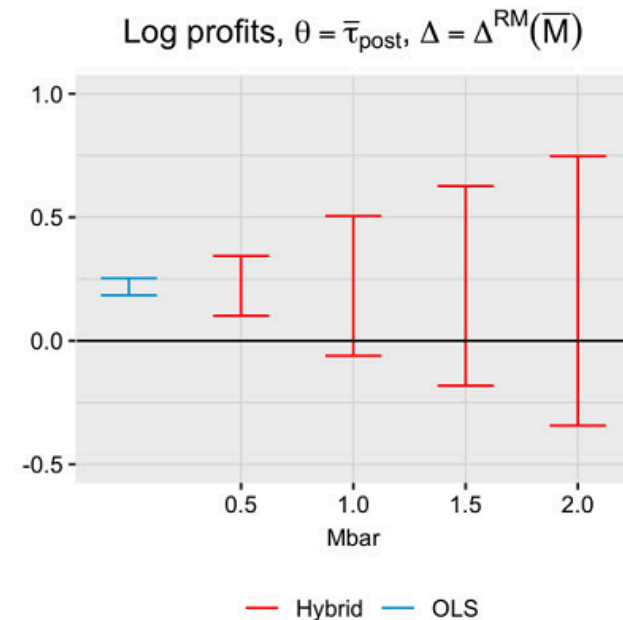
# Sensitivity analysis for Diff-in-Diff

- In an event study → null effects prior to the intervention:



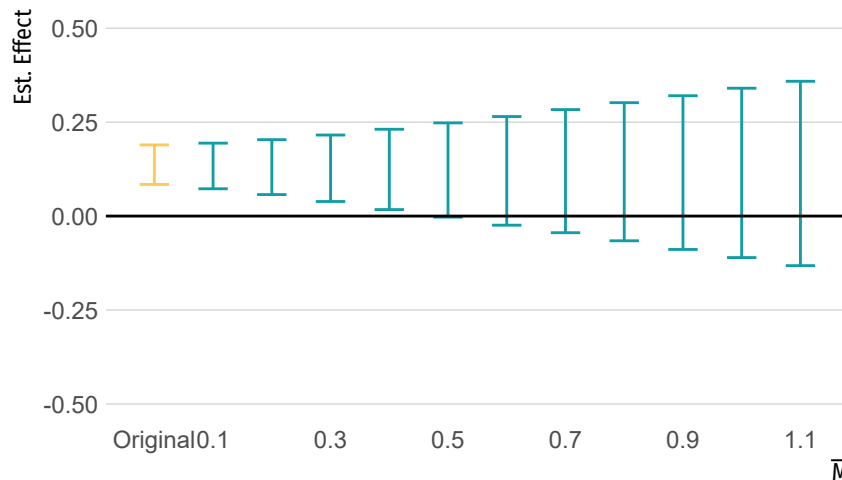
# Honest approach to test pretrends

- One main issue with the previous test → **Underpowered**
- Rambachan & Roth (2023) propose **sensitivity bounds** to allow pre-trends violations:
  - E.g. Violations in the post-intervention period can be *at most*  $M$  times the max violation in the pre-intervention period.

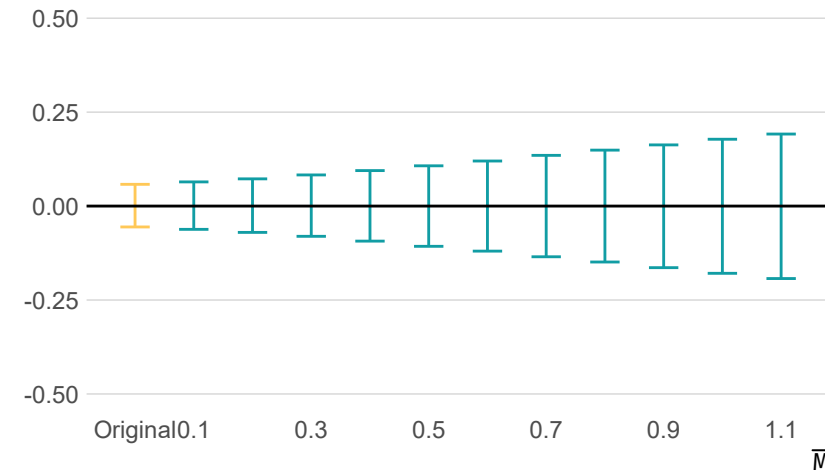


# Honest approach to test pretrends

- One drawback of the previous method is that it can **overstate** (or understate) the robustness of findings if the point estimate is biased.
  - Honest CIs depend on the **magnitude of the point estimate** as well as the **pre-trend violations**.
- Matching can **reduce the overall bias** of the point estimate



(a) Biased estimate



(b) Unbiased estimate

# How do we match?

- Match on covariates or outcomes? Levels or trends?
- Propensity score matching? Optimal matching? etc.

This paper:

- **Match on time-invariant covariates** that could make groups behave differently.
  - Use distribution of covariates to match on a template.
- Use of **Mixed-Integer Programming (MIP) Matching** (Zubizarreta, 2015; Bennett, Zubizarreta, & Vielma, 2020):
  - Balance covariates directly
  - Yield largest matched sample under balancing constraints (cardinality matching)
  - Works fast with large samples

Simulations



# Different scenarios

For linear and quadratic functions:

S1: No interaction between  $X$  and  $t$

S2: Equal interaction between  $X$  and  $t$

S3: Differential interaction between  $X$  and  $t$

Additional tests:

S1b-S3b: Including time-varying covariates

- For all scenarios, differential distribution of covariates  $X$  between groups

# Data Generating Processes

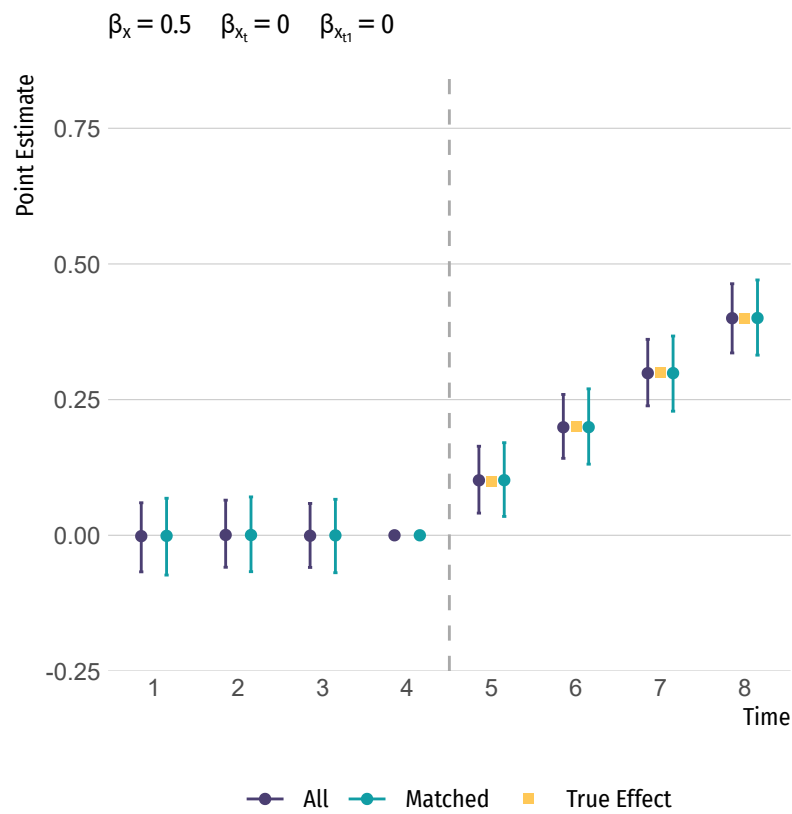
Scenarios	Functions		
<i>Linear</i>			
(1) No interaction between $X$ and $t$	$\gamma_0(X) = \beta_x \cdot X$	$\gamma_1 = \gamma_2 = 0$	
(2) Equal interaction between $X$ and $t$ by treatment	$\gamma_0(X) = \beta_x \cdot X$	$\gamma_1(X, t) = \beta_{x_t} \cdot X \cdot \frac{t}{2}$	$\gamma_2(X, t) = 0$
(3) Different interaction between $X$ and $t$ by treatment	$\gamma_0(X) = \beta_x \cdot X$	$\gamma_1(X, t) = \beta_{x_t} \cdot X \cdot \frac{t}{2}$	$\gamma_2(X, t) = \beta_{x_{t1}} \cdot X \cdot \frac{t}{5} \cdot Z$
<i>Quadratic</i>			
(1) No interaction between $X$ and $t$	$\gamma_0(X) = \beta_x \cdot X + \beta_x \cdot \frac{X^2}{10}$	$\gamma_1 = \gamma_2 = 0$	
(2) Equal interaction between $X$ and $t$ by treatment	$\gamma_0(X) = \beta_x \cdot X + \beta_x \cdot \frac{X^2}{10}$	$\gamma_1(X, t) = \beta_{x_t} \cdot X \cdot \frac{t^2}{10}$	$\gamma_2(X, t) = 0$
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# Parameters:

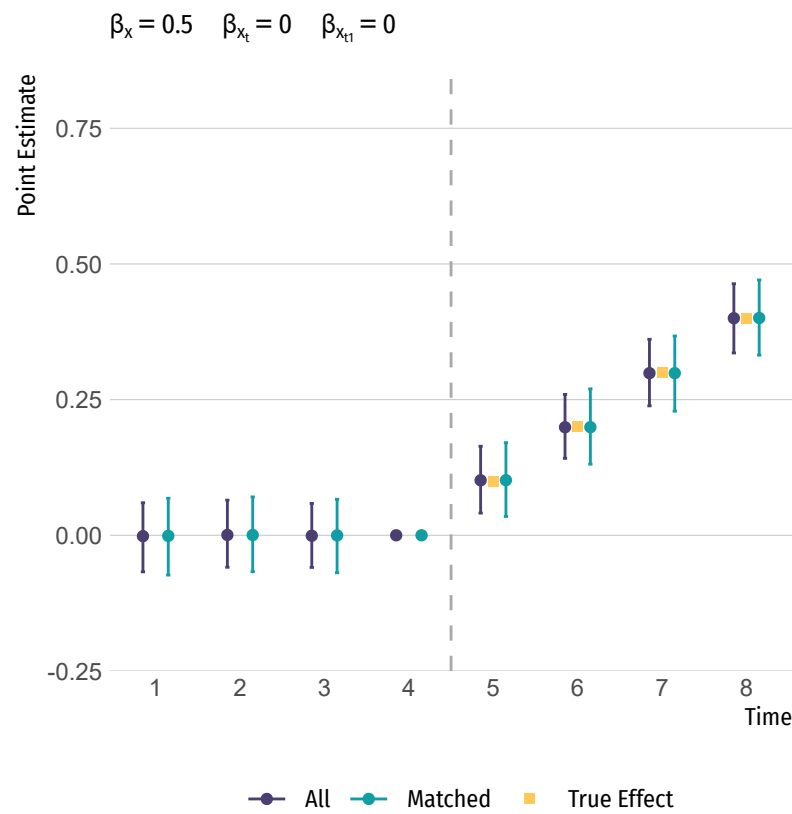
Parameter	Value
Number of obs (N)	1,000
$\Pr(Z=1)$	0.5
Time periods (T)	8
Last pre-intervention period (T_0)	4
Matching PS	Nearest neighbor (using calipers)
MIP Matching tolerance	.01 SD
Number of simulations	1,000

- Estimate compared to sample ATT (*can be different for matching*)

# S1 - No interaction between X and t

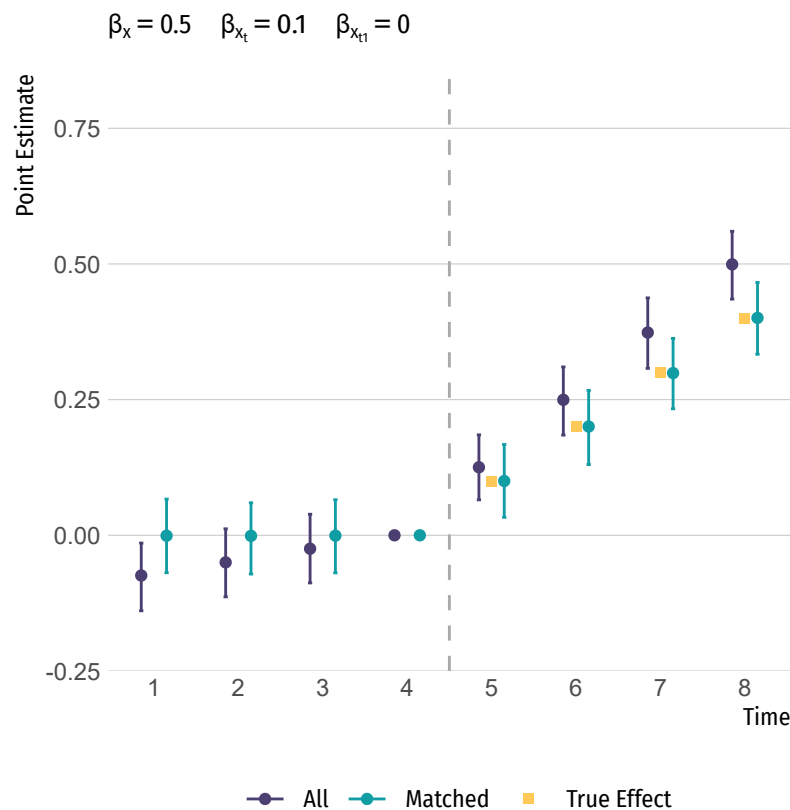


(a) Linear

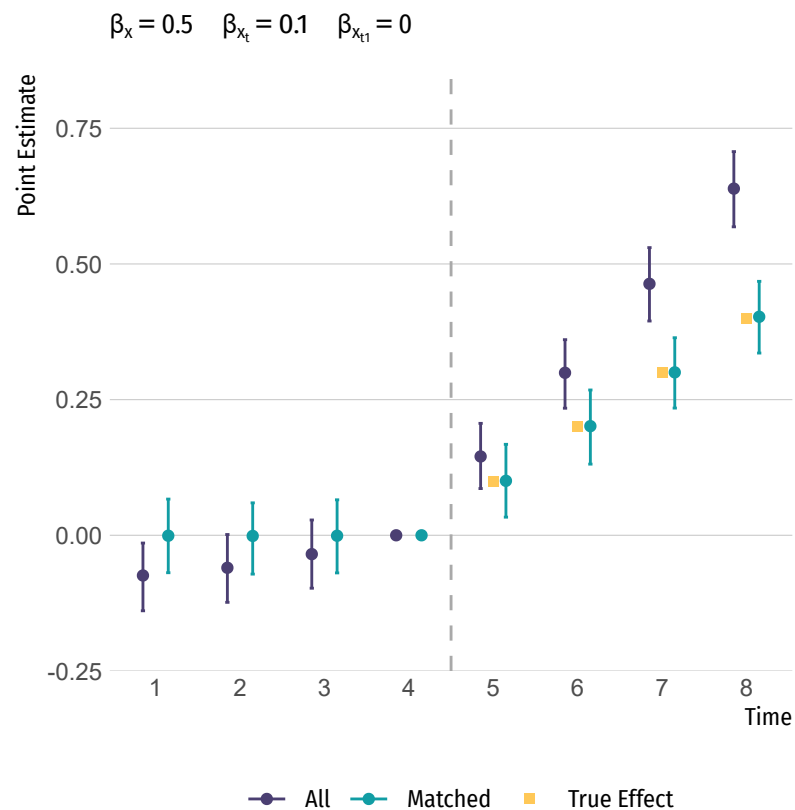


(b) Quadratic

# S2 - Equal interaction between X and t by treatment

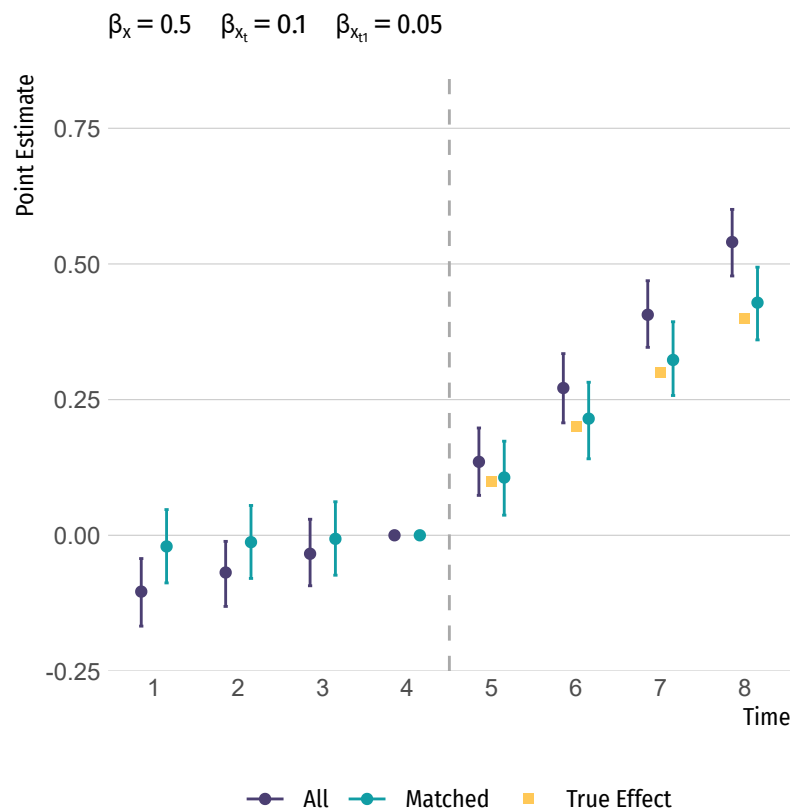


(a) Linear

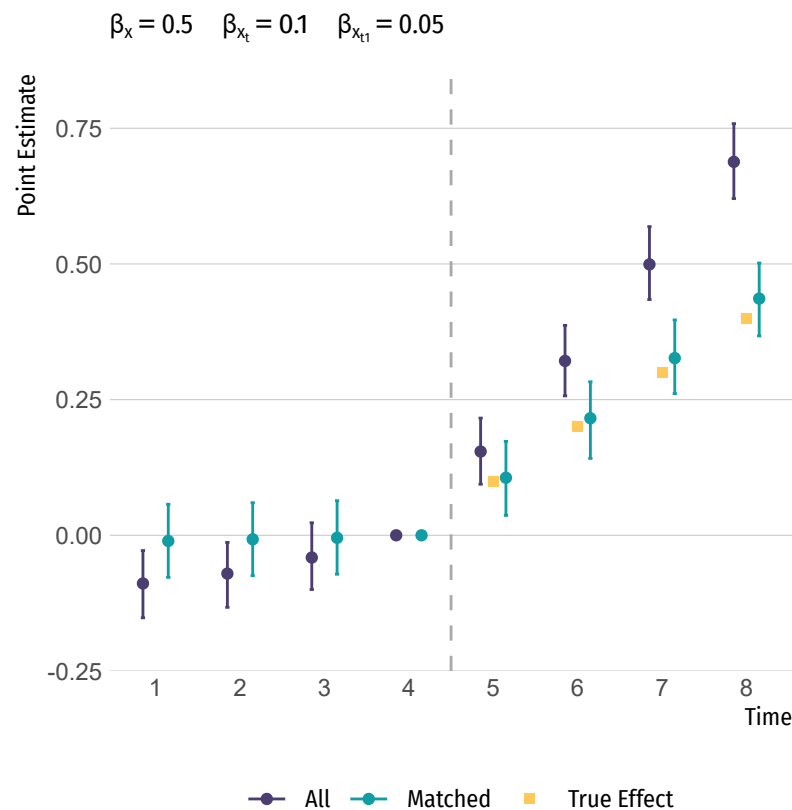


(b) Quadratic

# S3 - Differential interaction between X and t by treatment

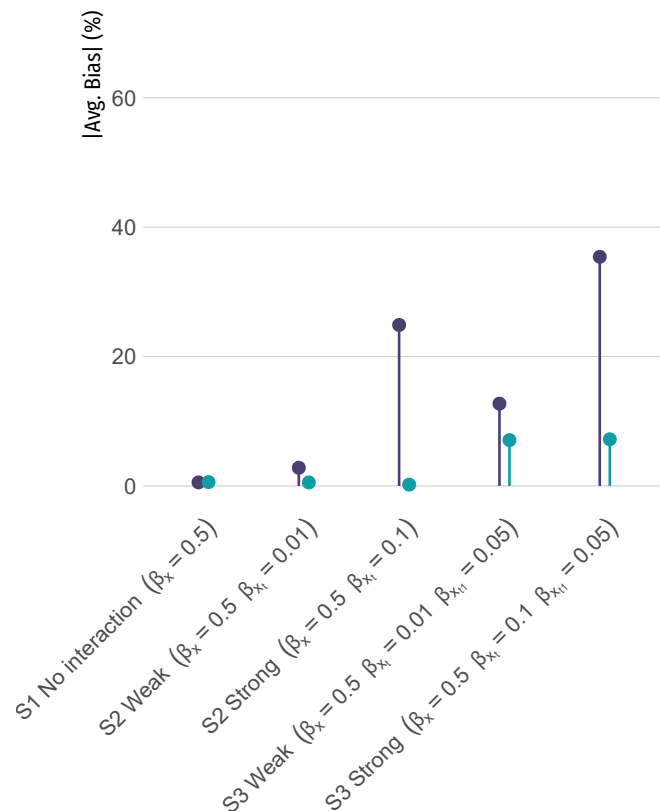


(a) Linear



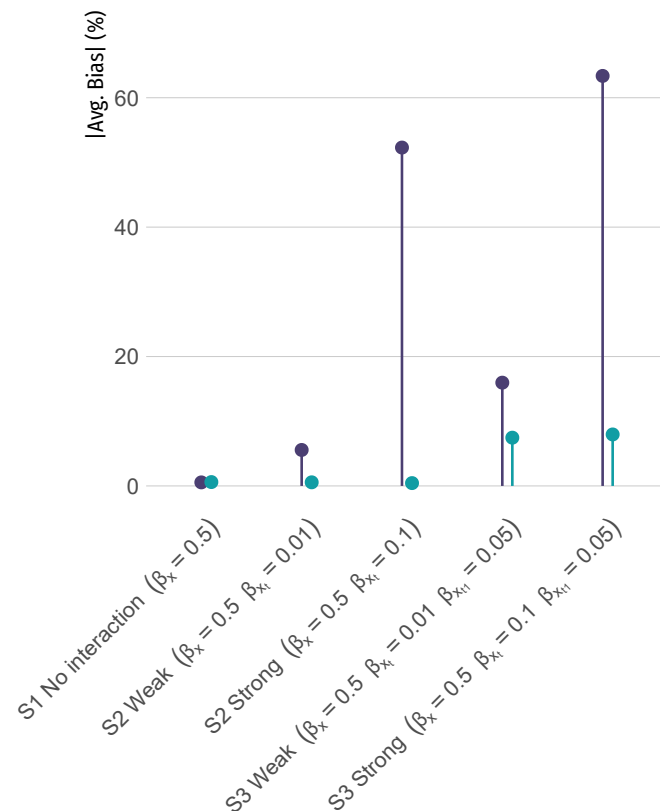
(b) Quadratic

# Matching as an adjustment method for reducing/eliminating bias



—●— All —●— Matched

(a) Linear

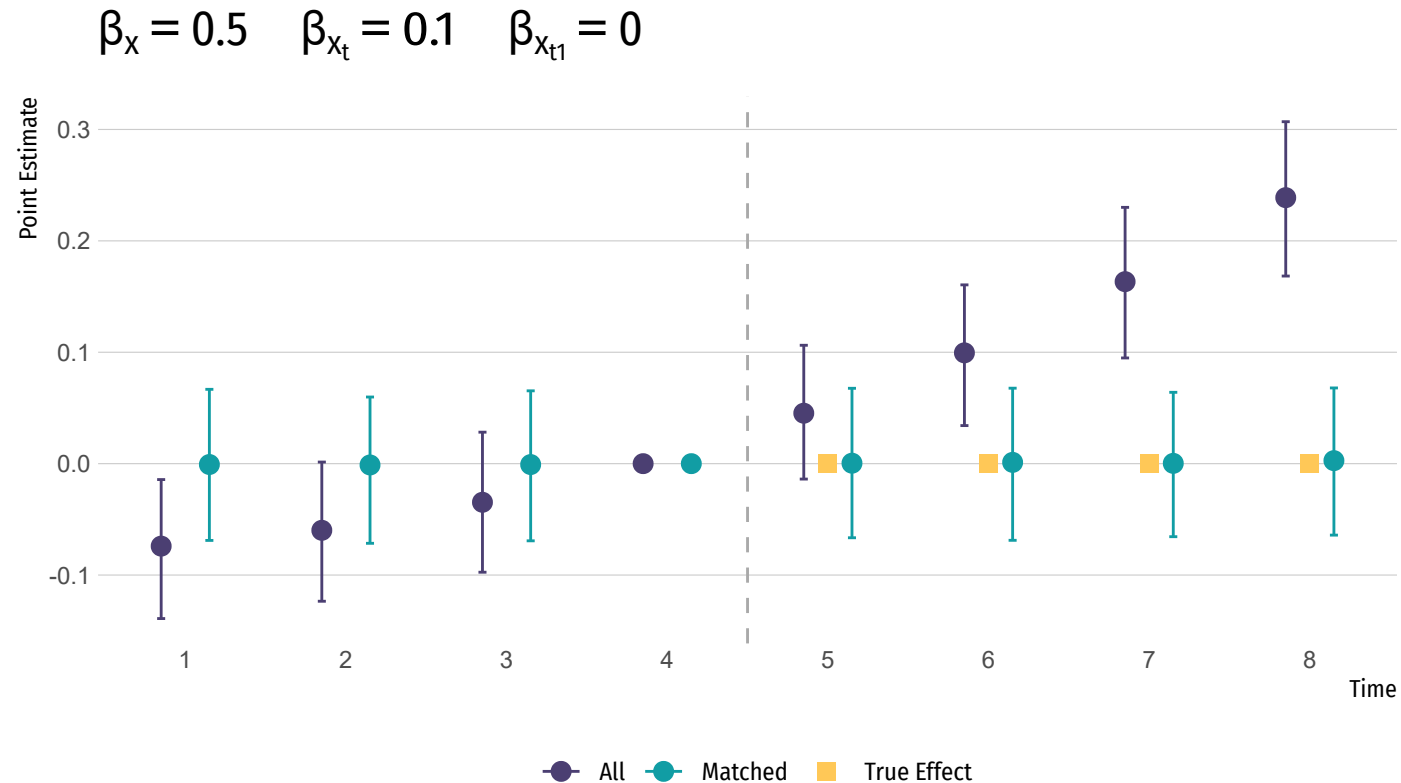


—●— All —●— Matched

(b) Quadratic

# Why is this bias reduction important?

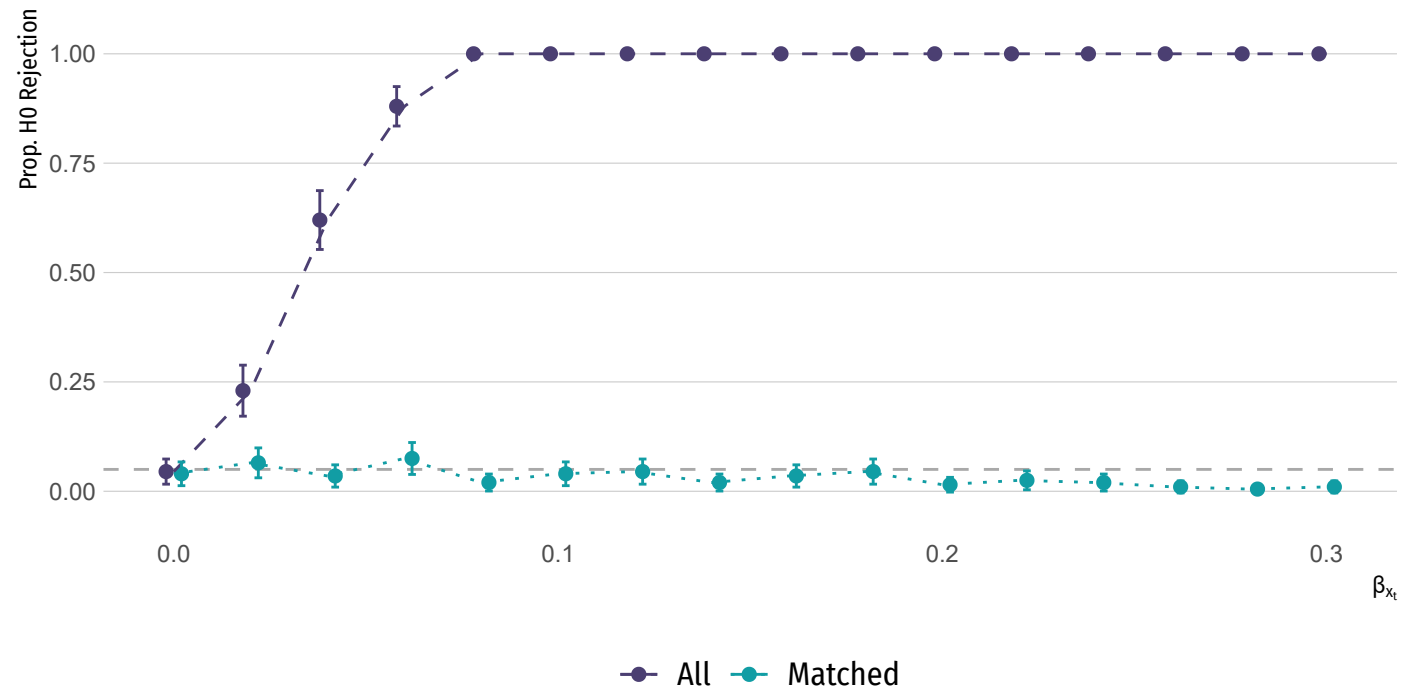
- Example of S2 (Quadratic) with no true effect:





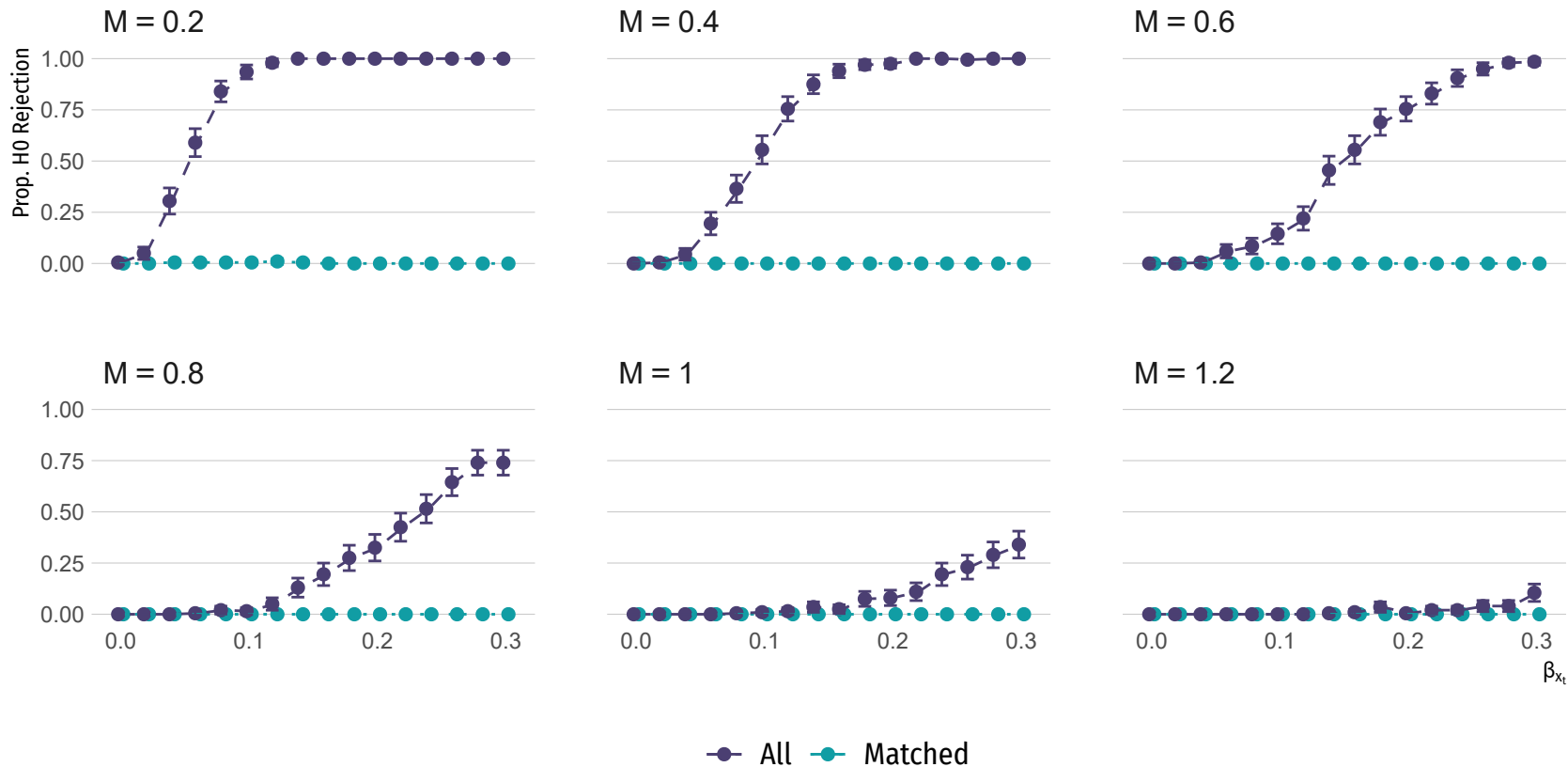
# Why is this bias reduction important?

- Even under modest bias, we would incorrectly reject the null 20% of the time.



# Why is this bias reduction important?

- Sensitivity analysis results are skewed by the magnitude of the bias.



Application

# Preferential Voucher Scheme in Chile

- Universal **flat voucher** scheme <sup>2008</sup> → Universal + **preferential voucher** scheme
- Preferential voucher scheme:
  - Targeted to bottom 40% of vulnerable students
  - Additional 50% of voucher per student
  - Additional money for concentration of SEP students.

## Students:

- Verify SEP status
- Attend a SEP school

## Schools:

- Opt-into the policy
- No selection, no fees
- Resources ~ performance

# Impact of the SEP policy

- **Mixed evidence of impact on test scores** for lower-income students (Aguirre, 2022; Feigenberg et al., 2019; Neilson, 2016; Mizala & Torche, 2013)

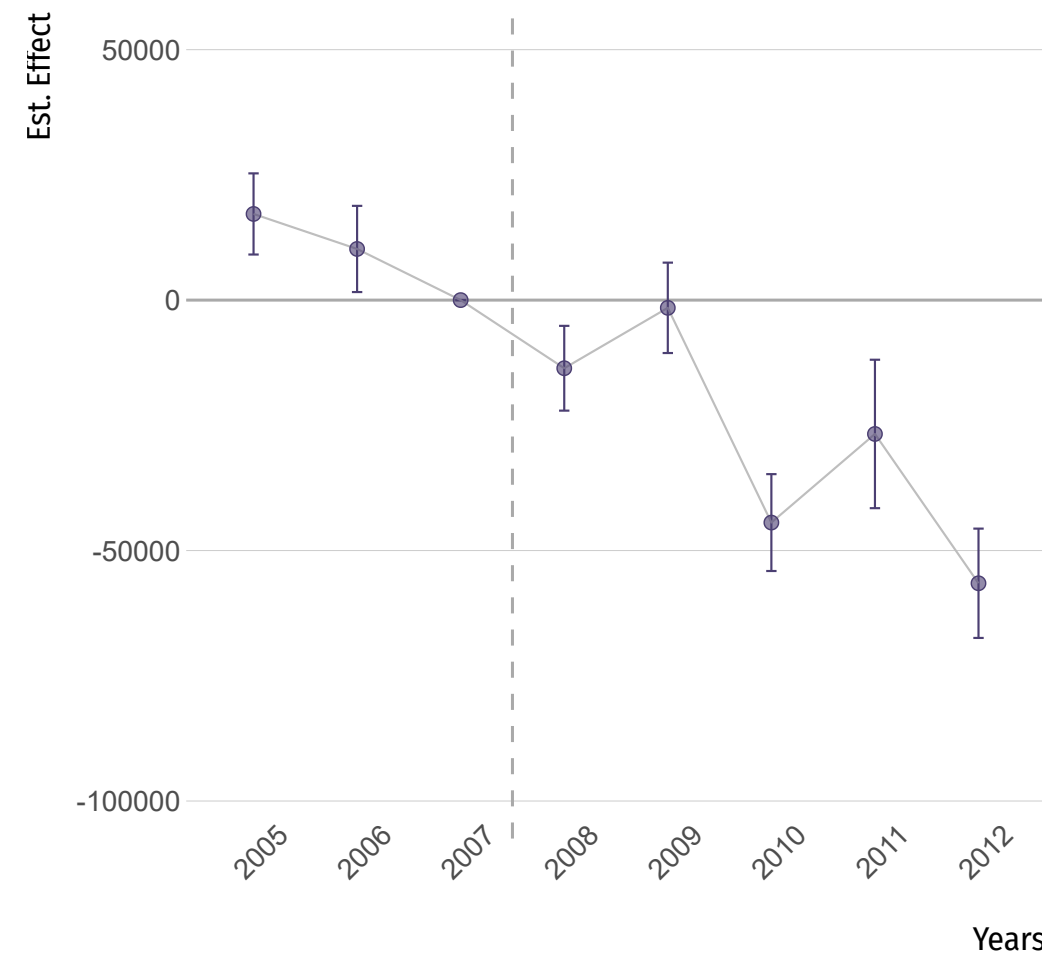
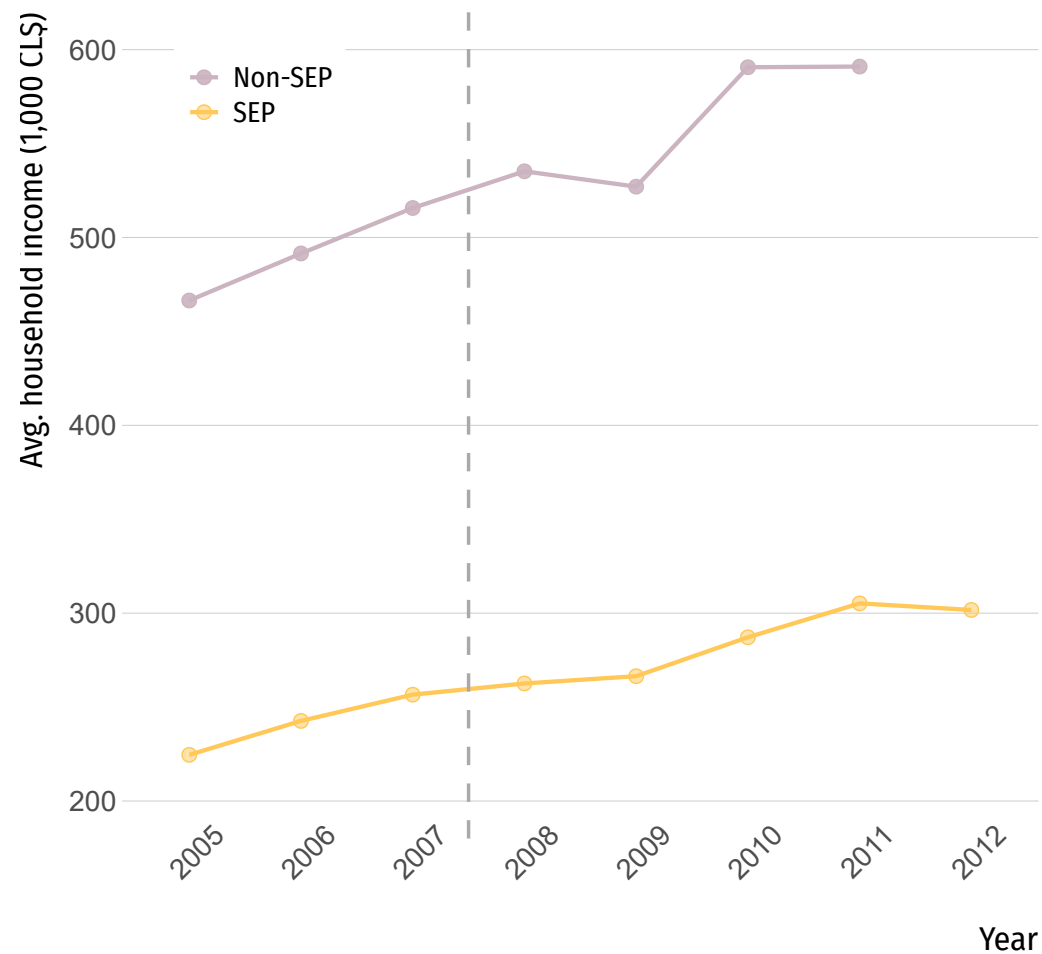
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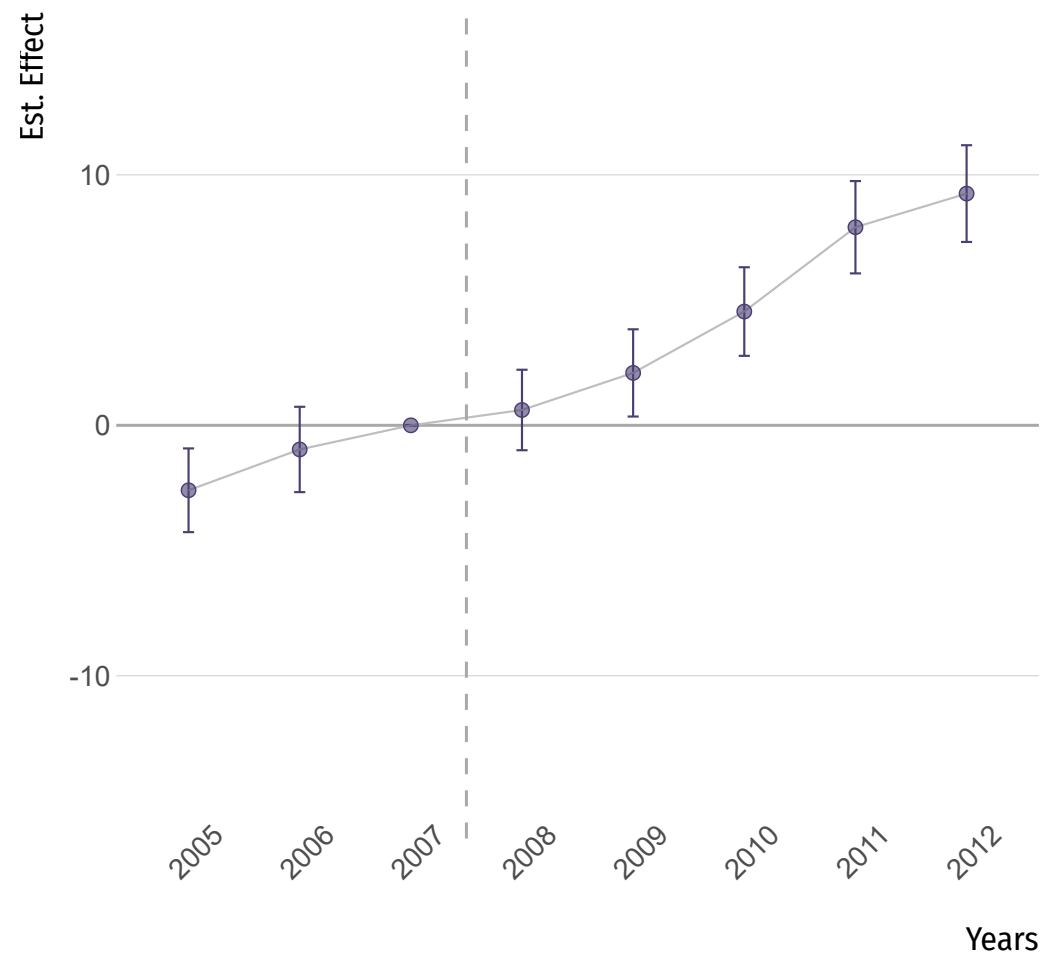
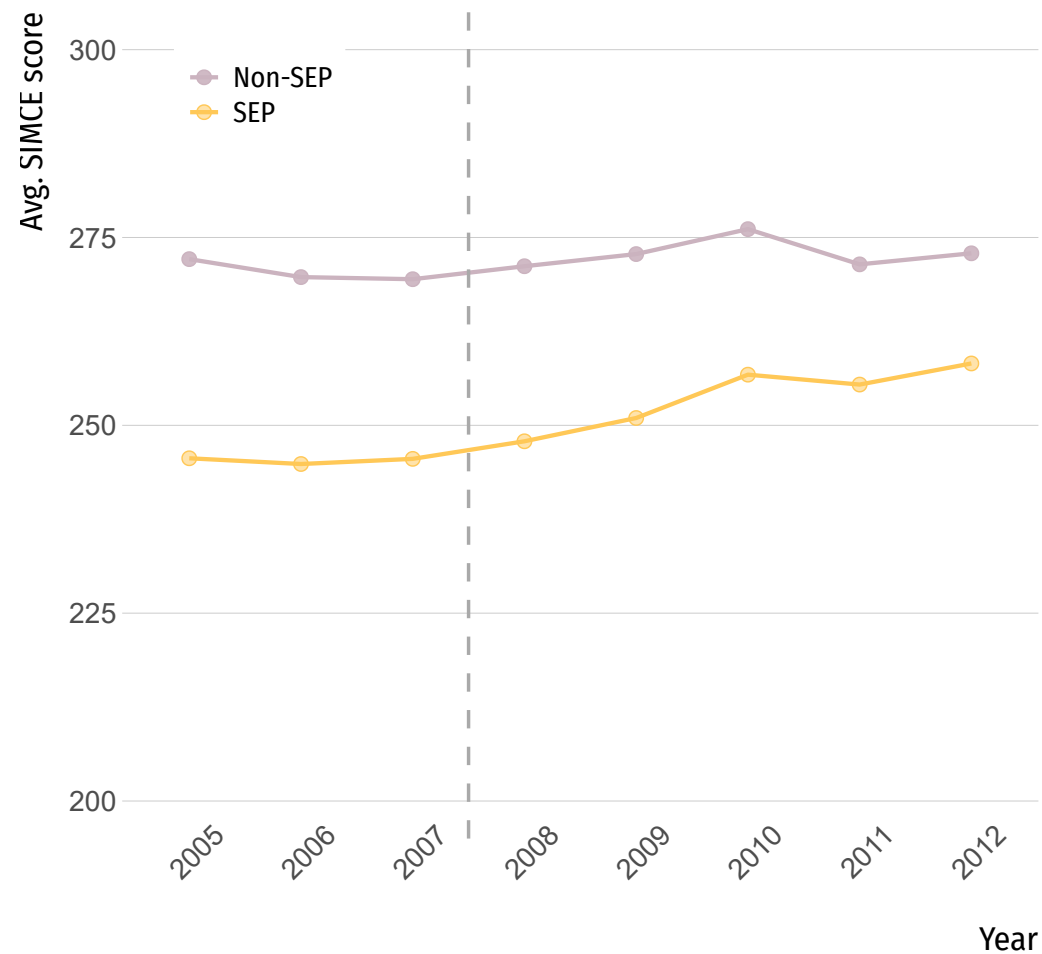
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- Design could have **increased** socioeconomic segregation (E.g. Incentives for concentration of SEP students)
- Key decision variables for schools: Performance, current SEP students, competition, add-on fees.
- **Diff-in-diff (w.r.t. 2007) for SEP and non-SEP schools:**
  - Only for **private-subsidized schools**
  - Matching using 2007 variables (similar results when using 2005-2007).
  - Outcome: Average students' household income and SIMCE score

# Before matching: Household income





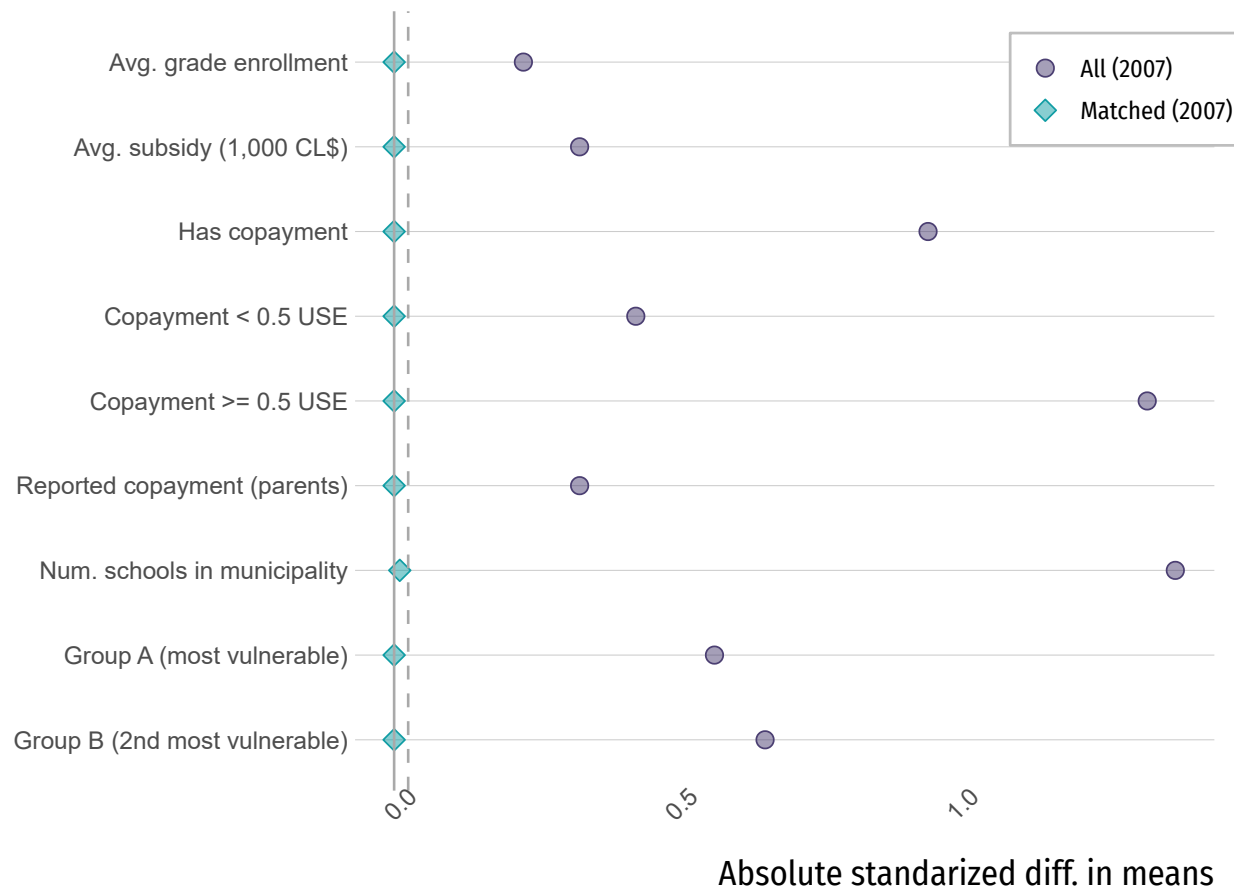
# Before matching: Average SIMCE



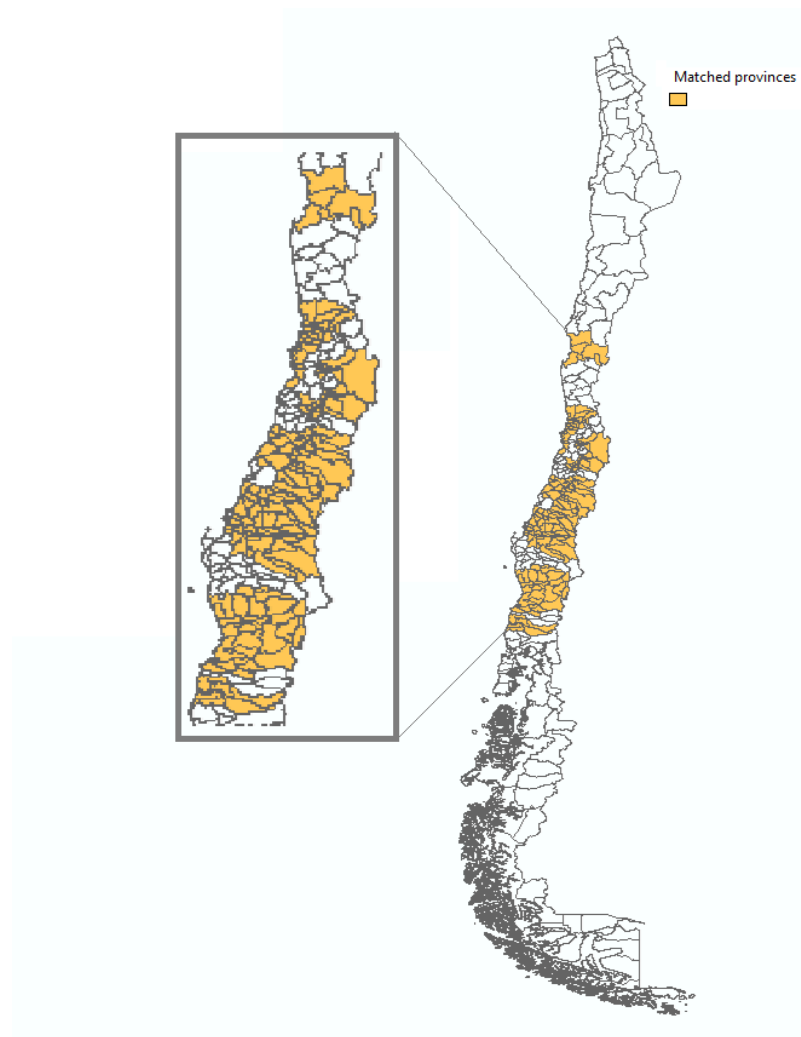
# Matching + DD

- **Prior to matching:** No parallel pre-trend
- **Different types of schools:**
  - Schools that charge high co-payment fees.
  - Schools with low number of SEP student enrolled.
- **MIP Matching** using constant or "sticky" covariates:
  - Mean balance (0.025 SD): Enrollment, average yearly subsidy, number of voucher schools in county, charges add-on fees
  - Exact balance: Geographic province

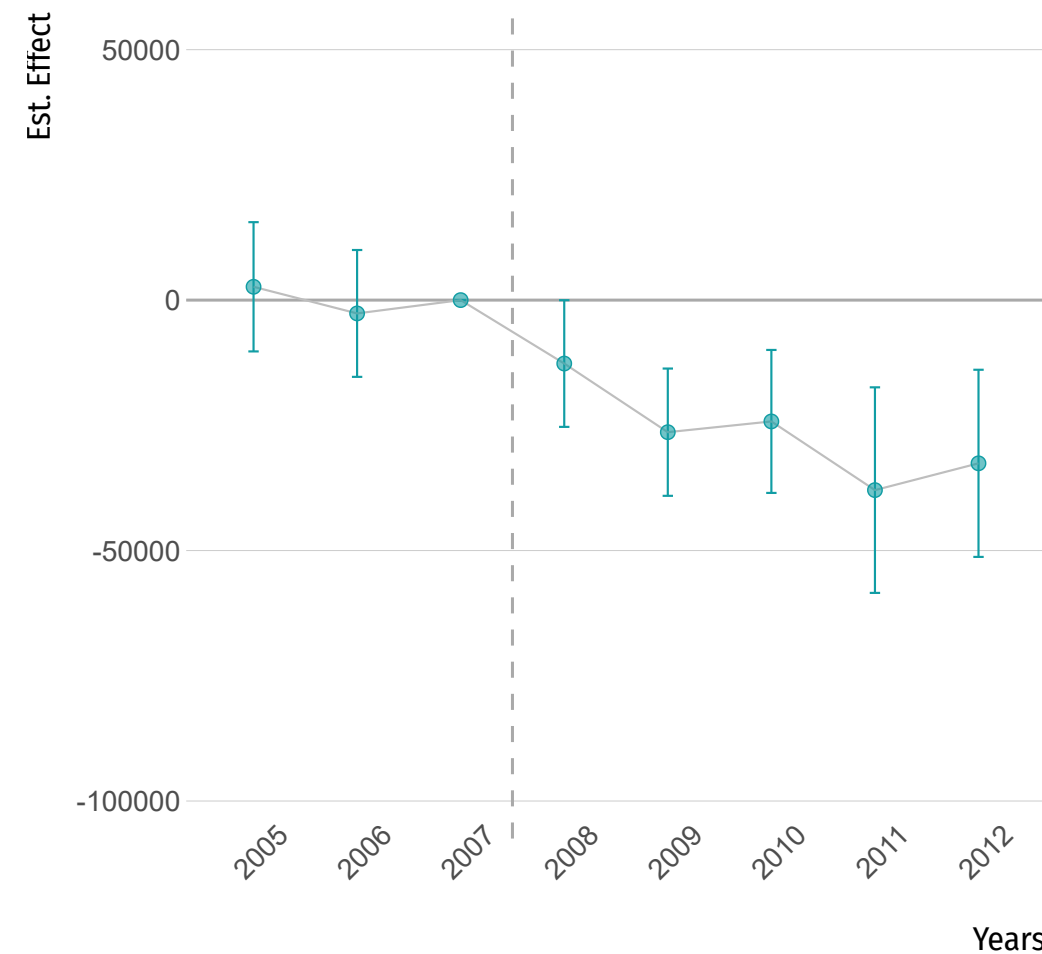
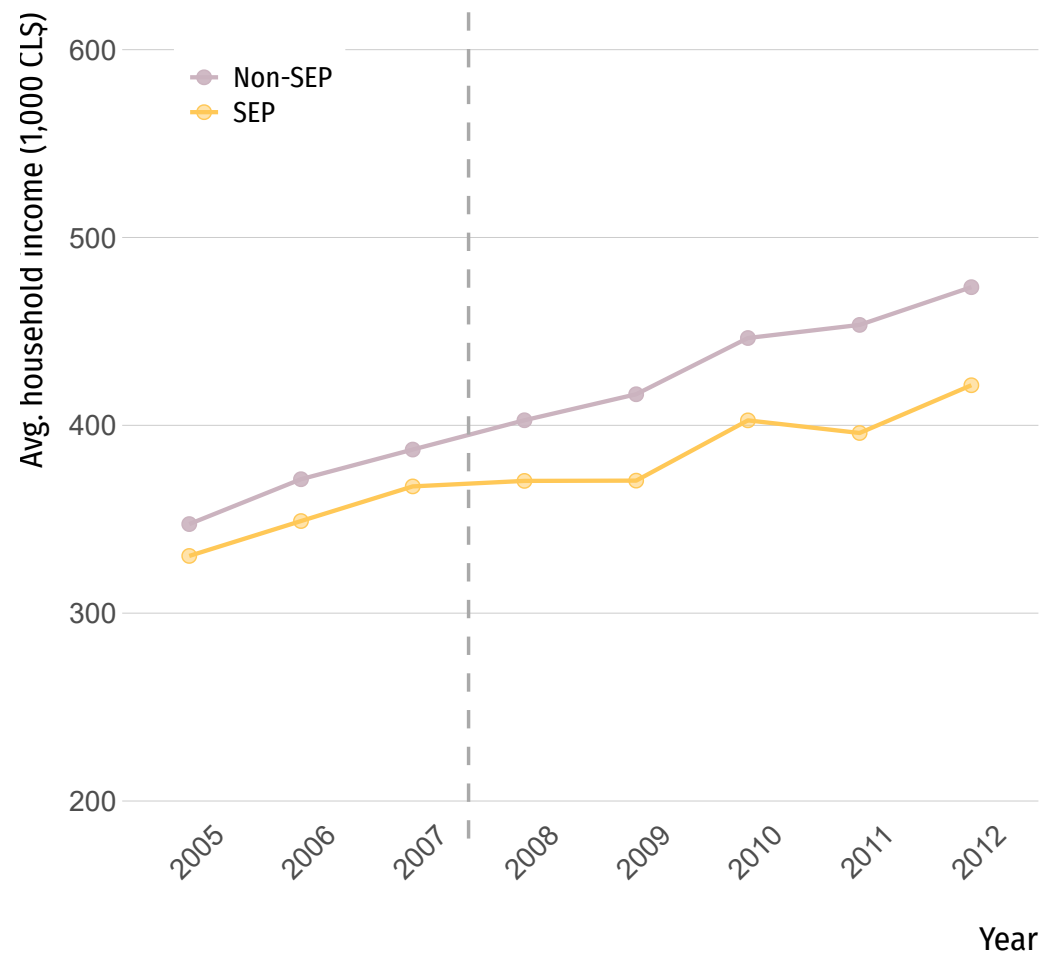
# Groups are balanced in specific characteristics



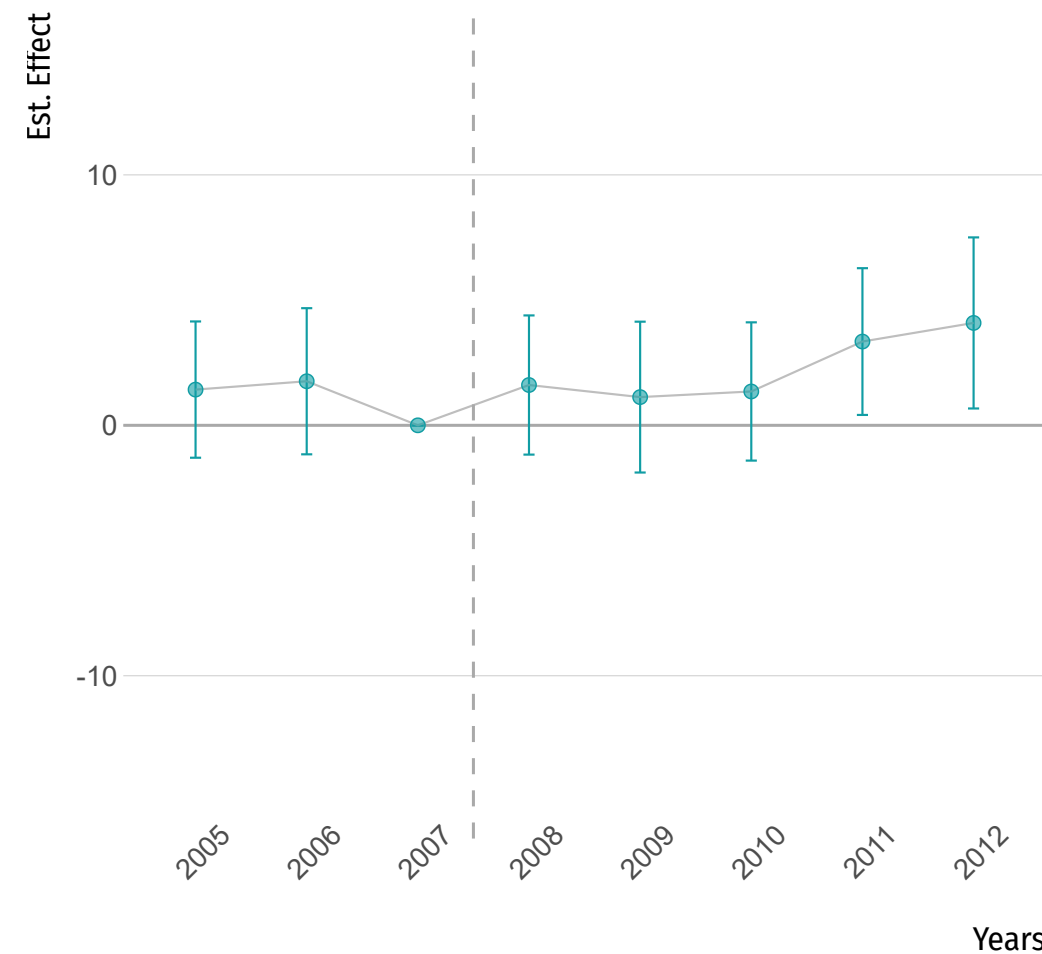
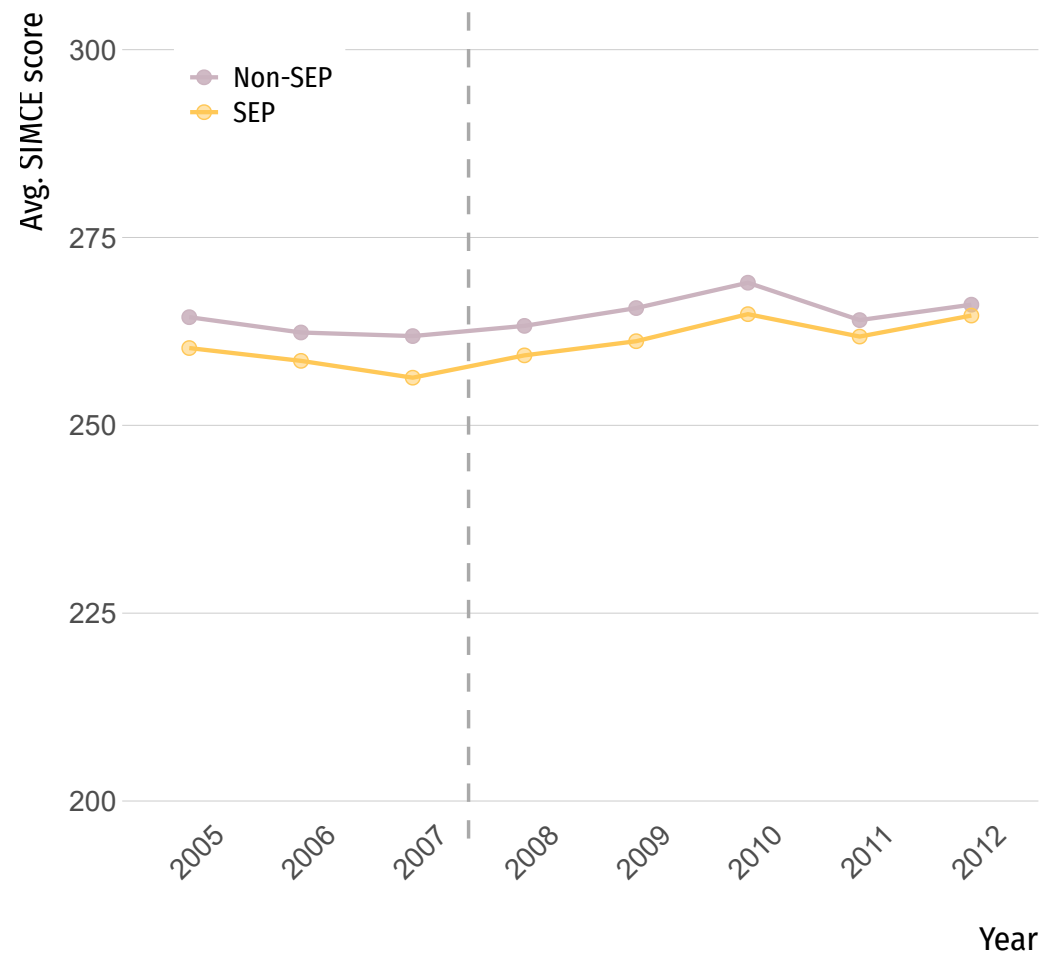
# Matching in 16 out of 53 provinces



# After matching: Household income



# After matching: Average SIMCE

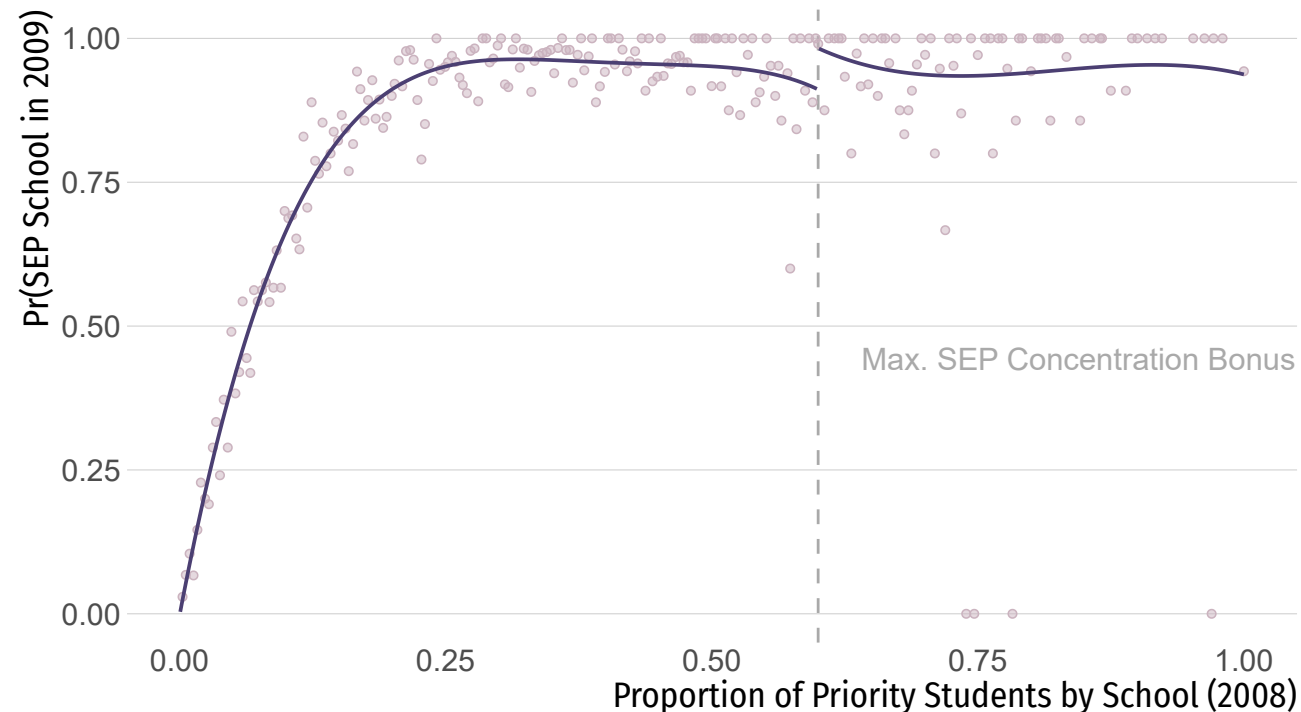


# Results

- **Matched schools:**
  - More vulnerable and lower test scores than the population mean.
- **9pp increase in the income gap** between SEP and non-SEP schools in matched DD:
  - SEP schools attracted even more vulnerable students.
  - Non-SEP schools increased their average family income.
- **No evidence of increase in SIMCE score:**
  - Could be a longer-term outcome.
- Findings in segregation are **moderately robust to hidden bias** (Keele et al., 2019):
  - $\Gamma_c = 1.76 \rightarrow$  Unobserved confounder would have to change the probability of assignment from 50% vs 50% to 32.7% vs 67.3%.
  - Allows up to 70% of the maximum deviation in the pre-intervention period ( $M = 0.7$ ) vs 50% without matching (Rambachan & Roth, 2023)

# Potential reasons?

- Increase in probability of becoming SEP in 2009 **jumps discontinuously at 60%** of SEP student concentration in 2008 (4.7 pp; SE = 0.024)





Let's wrap it up

# Conclusions and Next Steps

- Matching can be an important tool to address **violations in PTA**.
- **Bias reduction** is very important for sensitivity analysis.
- **Serial correlation** also plays an important role: Don't match on random noise.
- Next steps: Partial identification using time-varying covariates



# Difference-in-Differences using Mixed-Integer Programming Matching Approach

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# SEP adoption over time

