

# Econometrics II

## Lecture 5: Matching Estimators

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# Literature

- 1 **"Causal Inference for Statistics, Social and Biomedical Sciences: An Introduction"**, Imbens and Rubin  
Chapters 12.1-12.3
- 2 **"Mostly Harmless Econometrics"**, Angrist and Pischke  
Chapter 3.3.1-3.3.3

These notes draw on those books. All mistakes are mine.

# Plan for Today

- 1 Conditional Independence Assumption and Balance
- 2 Propensity Score
- 3 Achieving Balance
  - Matching
  - Propensity Score Matching
  - Reweighting
- 4 Matching and Regressions
- 5 Practical Issues

# Conditional Independence

In Lecture 1 we discussed that we require

$$\mathbb{E}[Y_i(0)|D_i = 1] - \mathbb{E}[Y_i(0)|D_i = 0] = 0$$

for causal inference about ATT. **This is fundamentally not testable.**

Maybe plausible:  $\{Y_i(0), Y_i(1)\} \perp D_i | X_i$ , and  $X_i$  are observable (CIA).<sup>1</sup>

Conditional Independence Assumption implies:

$$\begin{aligned}\mathbb{E}[Y|D = 1, X = x] - \mathbb{E}[Y|D = 0, X = x] \\ = \mathbb{E}[Y(1) - Y(0)|X = x] \equiv \tau_x\end{aligned}$$

Suggests one could condition analysis on  $X$ . But also...

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<sup>1</sup>Fundamentally not testable either! Requires substantial information.

# Why Covariate Balance?

...when does a simple difference in means comparison give  $\mathbb{E}_X[\tau_x]$ ?

$$\begin{aligned} & \mathbb{E}[Y|D=1] - \mathbb{E}[Y|D=0] \\ = & \int \mathbb{E}[Y|D=1, X=x] f_{X|D=1} dx - \int \mathbb{E}[Y|D=0, X=x] f_{X|D=0} dx && \text{[LIE]} \\ = & \int (\mathbb{E}[Y|D=1, X=x] - \mathbb{E}[Y|D=0, X=x]) f_X dx && \text{[If } f_{X|D=1} = f_{X|D=0}! \text{]} \\ = & \mathbb{E}_X[\tau_x] \end{aligned}$$

Balanced covariate distribution is important!

# Why Covariate Balance?

## Today:

Check whether (confounding) covariates  $X_i$  are balanced.

Exactly analogous to balance test of randomised experiment.

Make sure the (confounding) covariates  $X_i$  are balanced.

Exactly analogous to stratification in randomised experiment.

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# Propensity Score

Building Intuition, Step 1:

*Why might imbalances in covariates be problematic?*

Imagine for some  $x_1 \neq x_2$

$$\{Y_i(0), Y_i(1)\} | X_i = x_1 \not\sim \{Y_i(0), Y_i(1)\} | X_i = x_2,$$

and we are simply comparing outcomes of treated and control observations.

*When will this yield an unbiased estimate of  $\mathbb{E}_X[\tau_x]$ ?*

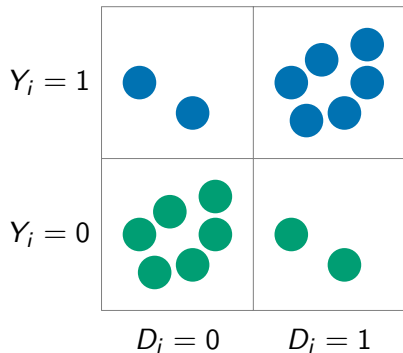


# Propensity Score

A simple example:

- $x_1 = 0, x_2 = 1$
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Suppose share treated at  $X_i = 0$  is 0.25 and at  $X_i = 1$  it is 0.75.



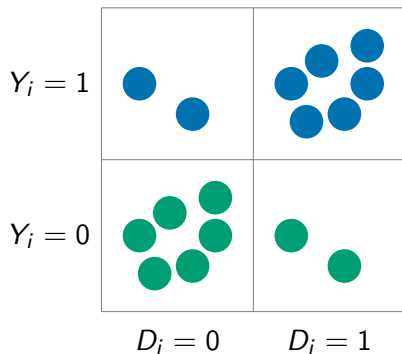
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Find positive treatment effect!



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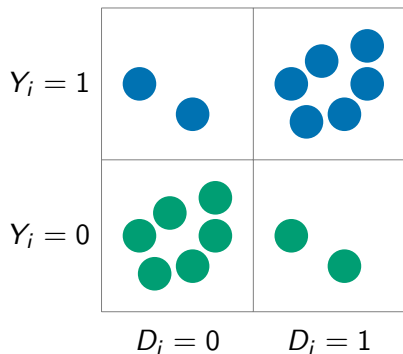
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**The problem**, somehow:

Observations with different  $(Y_i(0), Y_i(1))$  are assigned to treatment at different frequencies.



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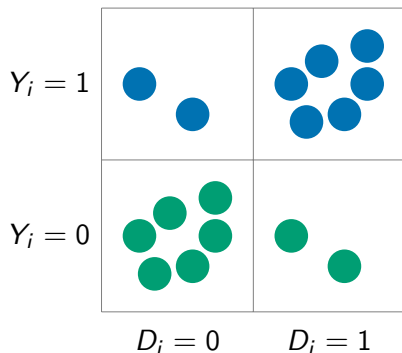
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Find positive treatment effect!

**The problem**, somehow:

Observations with different  $(Y_i(0), Y_i(1))$  are assigned to treatment at different frequencies.

Note: Find imbalanced covariate distributions.



# Propensity Score

Building Intuition, Step 2:

*How might imbalances in covariates occur?*

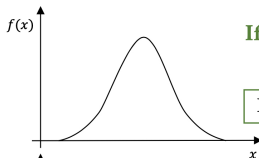
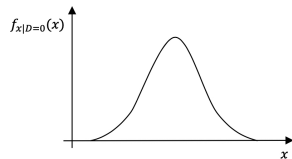
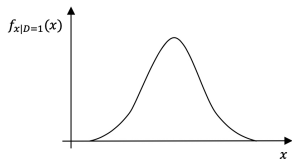
Denote with  $e(x) \equiv \Pr(D = 1|X = x)$  the probability of treatment given  $x$ .

We call this the '**propensity score**'.

*Can it be that  $e(x_1) \neq e(x_2)$  for some  $x_1$  and  $x_2$ , yet  $f_{X|D=1}(x) = f_{X|D=0}(x)$  for all  $x$ ?*

# Propensity Score

*When will the covariate distribution be balanced across t/c?*



**If and only if  $e(X)$  is a constant!**

Probably most important insight today.



**Proof?**

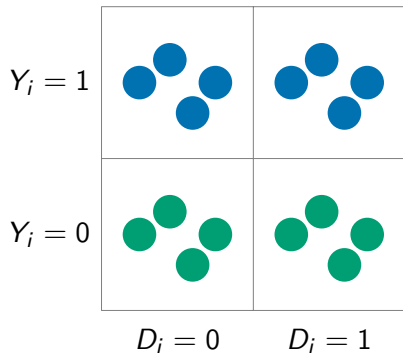
See Question 3 of Problem Set 2.

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Suppose  $e(0) = 0.5$  and  $e(1) = 0.5$ .



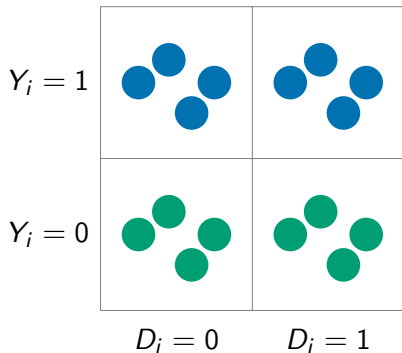
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Suppose  $e(0) = 0.5$  and  $e(1) = 0.5$ .

Find no treatment effect!





# Propensity Score

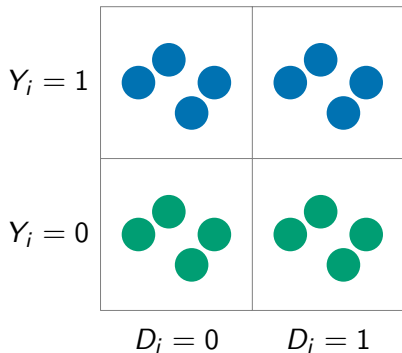
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Suppose  $e(0) = 0.5$  and  $e(1) = 0.5$ .

Find no treatment effect!

Find no imbalances in covariate distributions!



# Propensity Score

**Note:** the source of the problem is *not*...

- ... that distribution of outcomes is different at different  $X$ .
- ... that  $e(X_i) \neq 0.5$ .

**Two diagnostics:**

- $X_i = x$  is differently frequent in different treatment arms.

Motivation for standard balance checks.

- $e(X_i)$  not independent of  $x$ .

**Result:** In fact, these two diagnostics are equivalent!<sup>2</sup>

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<sup>2</sup>Probably most important insight today.

# Plan for Today

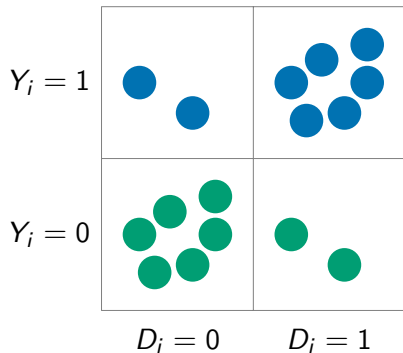
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# Achieving Balance

## Solution 1, Matching:

Control for influence of  $X$  by  
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*You see why that solves the problem?*



# Achieving Balance

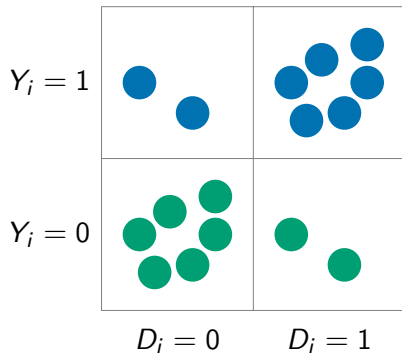
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Fundamental reason why this works:

Within each  $X$  cell,  $e(X)$  is constant.



# Achieving Balance

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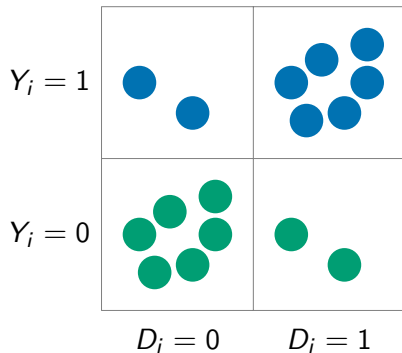
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Fundamental reason why this works:

Within each  $X$  cell,  $e(X)$  is constant.

(Typically: Average  $\tau_X$ 's by weights of interest.)



# Achieving Balance

Take a just slightly more complex example:

- $X$  has support  $\{x_1, x_2, x_3, x_4\}$ ;
- Distribution of potential outcomes is different;
- $e(x_1) = 0.5, e(x_2) = 0.25, e(x_3) = 0.5, e(x_4) = 0.5$ .

**Solution 2, Balancing Score Matching:**  $D_i \perp X_i | b(X_i)$ .

# Achieving Balance

**Solution 3, Propensity Score Matching:** Match on  $e(X)$ .<sup>3</sup>

Recall from before...

- If  $e(X)$  is constant, distribution of  $X$  in the treatment arms is the same.
- And if distribution of  $X$  is the same, simple difference in outcomes estimates the average treatment effect, by CIA!

Note:

- This is precisely why randomised trials are useful, they achieve constant  $e(X)$ .
- Stratification is about forcing realised  $\hat{e}(X)$  to be constant.  
Like with RCTs, realised  $\hat{e}(X)$  is what matters. *You see why?*

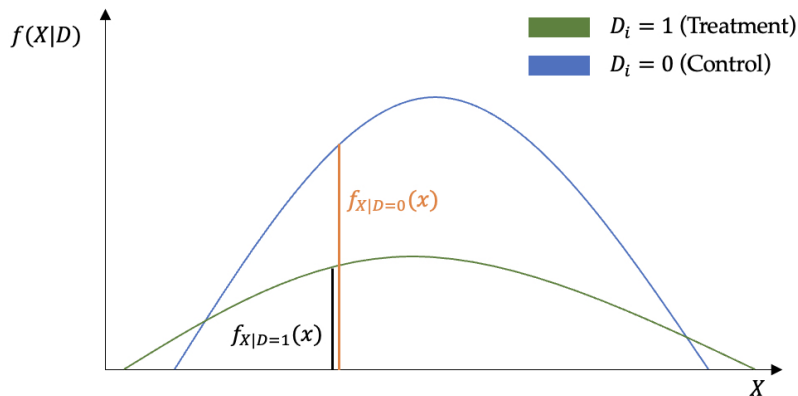
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<sup>3</sup>Propensity Score is the coarsest balancing score.



# Achieving Balance

Solution 4, **Reweighting** (Horvitz and Thomson):

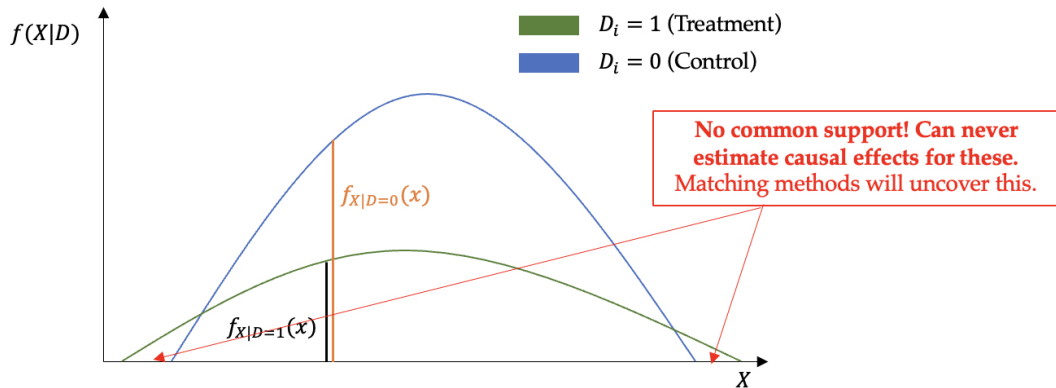


**Idea:** Find set of weights  $\omega(x)$  for all  $x$  such that

$$\omega(x) f_{X|D=0}(x) = f_{X|D=1}(x).$$

# Achieving Balance

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# Achieving Balance

By Bayes' rule (see Angrist and Pischke, ch. 3.3.1):

$$f_{X|D=1}(x) = \frac{Pr(D_i = 1|X_i = x)}{Pr(D_i = 1)} f_X(x) = \frac{e(x)}{e} f_X(x)$$
$$f_{X|D=0}(x) = \frac{Pr(D_i = 0|X_i = x)}{Pr(D_i = 0)} f_X(x) = \frac{1 - e(x)}{1 - e} f_X(x)$$

where  $e(x)$  is the propensity score and  $e = Pr(D_i = 1)$ . Then

$$\omega(x) = \frac{f_{X|D=1}(x)}{f_{X|D=0}(x)} = \frac{e(x)}{1 - e(x)} \frac{1 - e}{e} \quad (1)$$

Then the ATT can be estimated as

$$\hat{\beta}^{ATT} = \frac{1}{N_1} \sum_{i=1}^N D_i Y_i - \frac{1}{N_0} \sum_{i=1}^N (1 - D_i) Y_i \frac{\hat{e}_i(x)}{1 - \hat{e}_i(x)} \frac{1 - \hat{e}}{\hat{e}} \quad (2)$$

According to Imbens and Rubin, sensitive to the estimation of weights.

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# Matching and Regressions

## Solution 5, Regression Control:

*Upside* (see Angrist and Pischke):

Running regression on treatment, fully saturated in  $X$ , have

$$\beta_R = \frac{\mathbb{E}[\sigma_D^2(X_i)\tau_X]}{\mathbb{E}[\sigma_D^2(X_i)]}, \text{ where } \sigma_D^2(X_i) \equiv \mathbb{E}[(D_i - \mathbb{E}[D_i|X_i])^2|X_i]$$

and a standard ATT matching estimator would be

$$\beta_M = \mathbb{E}[\tau_X \Pr(X_i|D_i = 1)].$$

- 1 Matching estimator weights high  $X$ s with many treatment observations; OLS gives weight to observations with equal treatment shares (max. variance in  $D_i$ ).
- 2 If  $\tau_X$  varies little across  $X$  cells, makes little difference.
- 3 None gives any weight to  $X$  cells with no or all treatment.

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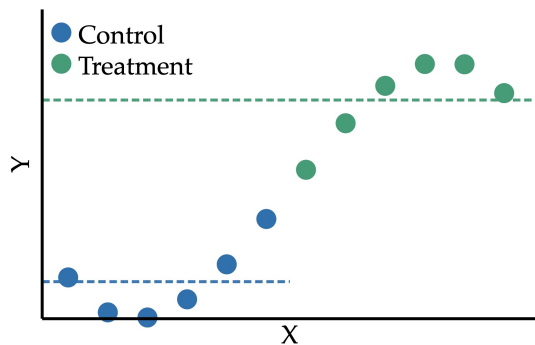
*Downside* (see Imbens and Rubin):

Their argument is that regression requires:

- 1 unconfoundedness assumption,  
and additionally:
- 2 functional form assumptions for  $X$ .

Is **strong** and **unnecessary**.

# Matching and Regressions

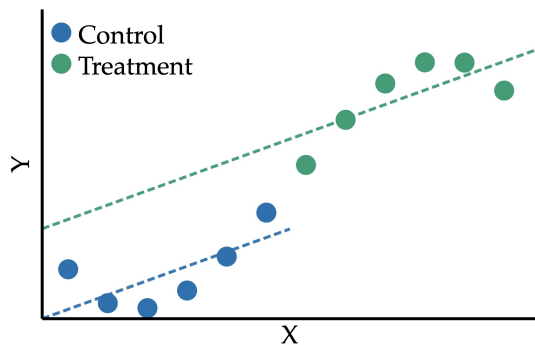


Suppose you run regression of  $Y$  on treatment status  $D$  and...

- **no control for  $X$ ,**
- linear control for  $X$ , or
- quadratic control for  $X$  interacted with  $D$ .

*What is the coefficient estimate on  $D$ ?*

# Matching and Regressions

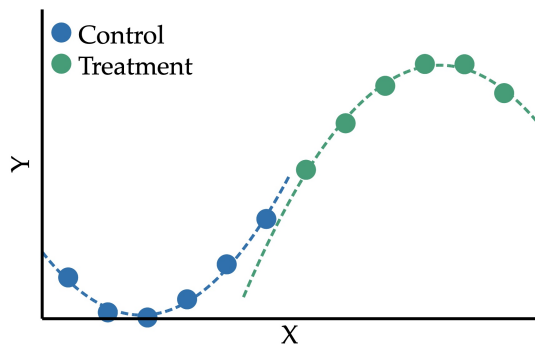


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# Practical Issues

*Distinguish:*

## 1 Design Stage

### 1 Assessing Overlap in Distributions

- How similar are distributions? [Univariate/Multivariate tests.]
- Do similar observations with opposite level of treatment exist?

### 2 Estimate Propensity Score

- Estimated propensity score matters; goal is to achieve balance in sample.
- Machine learning methods can potentially be helpful.

### 3 Create Balanced Sample: unbiased/robust inference; power.

- Match estimation sample: 1 to 1; 1 to many; on PS or X...
- Trim sample. Two competing forces for power: sample size vs. match quality.

# Practical Issues

*Distinguish:*

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- Match estimation sample: 1 to 1; 1 to many; on PS or X...
- Trim sample. Two competing forces for power: sample size vs. match quality.

No need for any  $Y$  here!

Can go back and forth, “play around”, until balanced sample is found...

... just do not condition this on  $Y$  data.

# Practical Issues

## 2 Assessment Stage

- Use pseudo outcomes to validate the approach (see example on next slide).

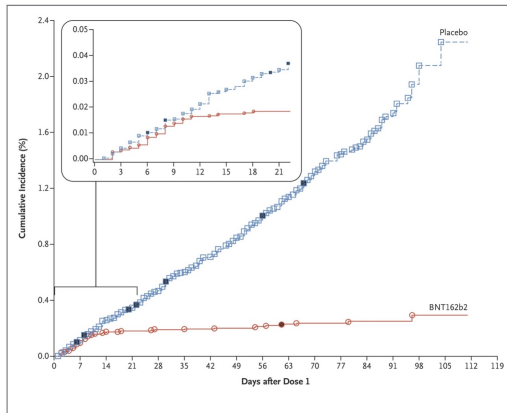
## 3 Analysis Stage

- Create PS blocks, or create exact matches, or inexact matches, and estimate mean outcome difference within those.
- Covariate adjustment might increase precision, see Imbens and Rubin, Part III.
- Inference easier with exact matches.



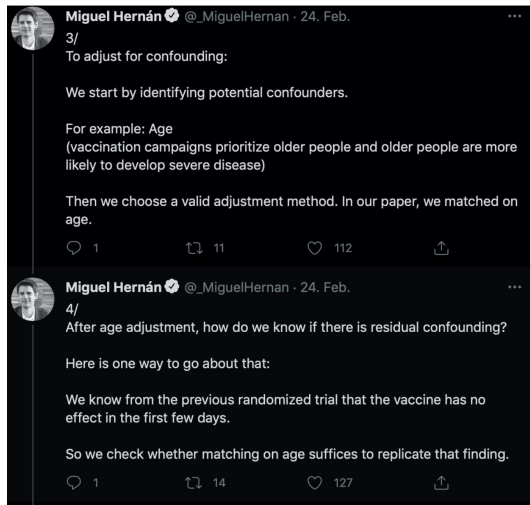
# Pseudo Outcomes: Example

Pfizer/BioNTech released results from an RCT in 12/2020 (NEJM,  $N = 43548$ )



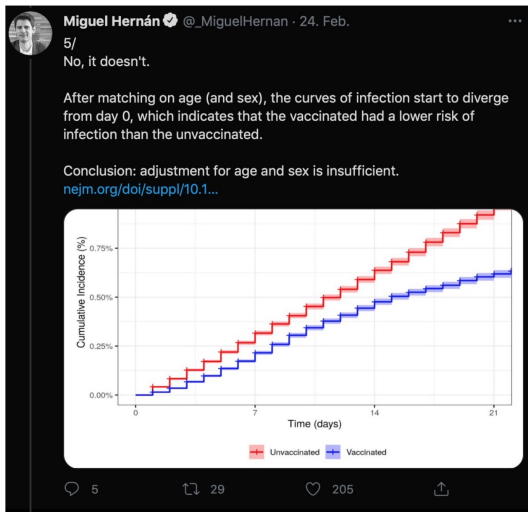
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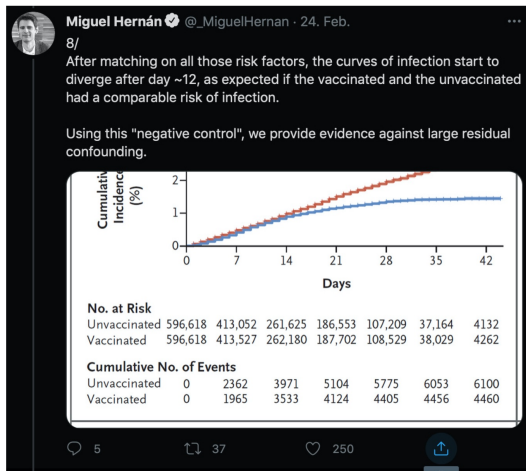
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# Making Matching Persuasive

David McKenzie writes about statistical and "rhetorical" plausibility.<sup>4</sup>

"Rhetorical" plausibility talks explicitly about why some individuals were treated and others were not.

Examples:

- Separate decision-maker with limited information decides on treatment.
- Capacity limits.
- Treatment consequence of randomization.
- Decision maker caress about different outcome than evaluator.

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<sup>4</sup><https://blogs.worldbank.org/impactevaluations/what-do-you-need-do-make-matching-estimator-convincing-rhetorical-vs-statistical>

# Summary

- Role of balanced covariates distribution.
- Covariates distributions are balanced iff  $e(X)$  is constant.
- Obtain balance forcing  $X$  to be the same (*matching*),  
or forcing  $e(X)$  to be the same (*propensity score matching*).
- Many practical choices how to implement those ideas.

Questions?