

# STA 235H - Introduction to Observational Studies I

Fall 2021

McCombs School of Business, UT Austin

# Announcements

## Homework 3 will be posted on Thursday

- Homework 2 answer key has been added to the course website (**check out the rubric!**)
- **No office hours today** → Moved to tomorrow (check out Calendly)
- I'll send out a poll for a **review session** for the midterm
  - If you are interested, please respond.

# Last week

- **Randomized controlled trials**
  - Why is it considered the gold standard?
  - How to analyze an RCT in practice?
  - Assumptions and limitations.



# Last week

- Randomized controlled trials
  - Assumptions for RCTs?



# Last week

- Randomized controlled trials
  - Limitations?



# Today, we're moving forward...



- **Introduction to Observational Studies:**
  - Can we identify causal effects without RCTs?
  - Assumptions
  - Matching vs OLS

No more chance[s]

# Introduction to observational studies

- Most times, we will not be able to randomize, and we need to work with **existing data**

## Observational data

- Data for which we can't manipulate the treatment assignment, e.g. data in its "natural state".

**Can we reasonably assume that the ignorability assumption holds?**



# Introduction to observational studies (cont.)



- Moving away from the core assumption of RCTs: that **"the probability of treatment assignment is a known function"** (Imbens & Rubin, 2015).

# Introduction to observational studies (cont.)



- Moving away from the core assumption of RCTs: that **"the probability of treatment assignment is a known function"** (Imbens & Rubin, 2015).
- We will maintain the assumption of **unconfoundedness** (to a certain extent).

What is that?

# Calling in the CIA

- **Unconfoundedness** means that the treatment assignment is independent from the potential outcomes.
- If you recall, the ignorability assumption assumes that:

$$Y(0), Y(1) \perp\!\!\!\perp Z$$

- What if you could assume that this holds **conditional on some covariates**?

## Conditional Independence Assumption (CIA)

$$Y(0), Y(1) \perp\!\!\!\perp Z|X$$

# An example about the CIA

- Let's think about the **fake CV example** and a real life application.
- **Causal question**: How does getting an internship affect your probability of being in the film industry 5 years later?
- A firm needs to hire interns ASAP, no time for interviews. What would this firm look at in a CV?
  - e.g. level of education, experience, name?
- *Could we assume that **conditional on education, experience, name characteristics, etc.** receiving an internship is independent from your potential outcomes?*

# The assignment mechanism

- **Key component** in causal analysis:
  - In RCTs, **assignment mechanism** is *known*.
  - But in **observational studies**?



# Selection on observables

- Units select into treatment based on characteristics **I can observe**.
- What this means in practice is that **all confounders are observable and I can adjust for them**.
  - **Overt bias**: Bias caused by observed confounders. I can remove it by adjusting by these variables.
  - **Hidden bias**: Bias caused by unobserved confounders. I can't directly remove it (I need to rely on other assumptions).

# How do we adjust for observables?

- One way we have seen so far is **regression adjustment**

$$Y_i = \beta_0 + \beta_1 Z_i + \beta_2 X_i + \varepsilon_i$$

**Under unconfoundedness, how would we interpret  $\beta_1$ ?**

# How do we adjust for observables?

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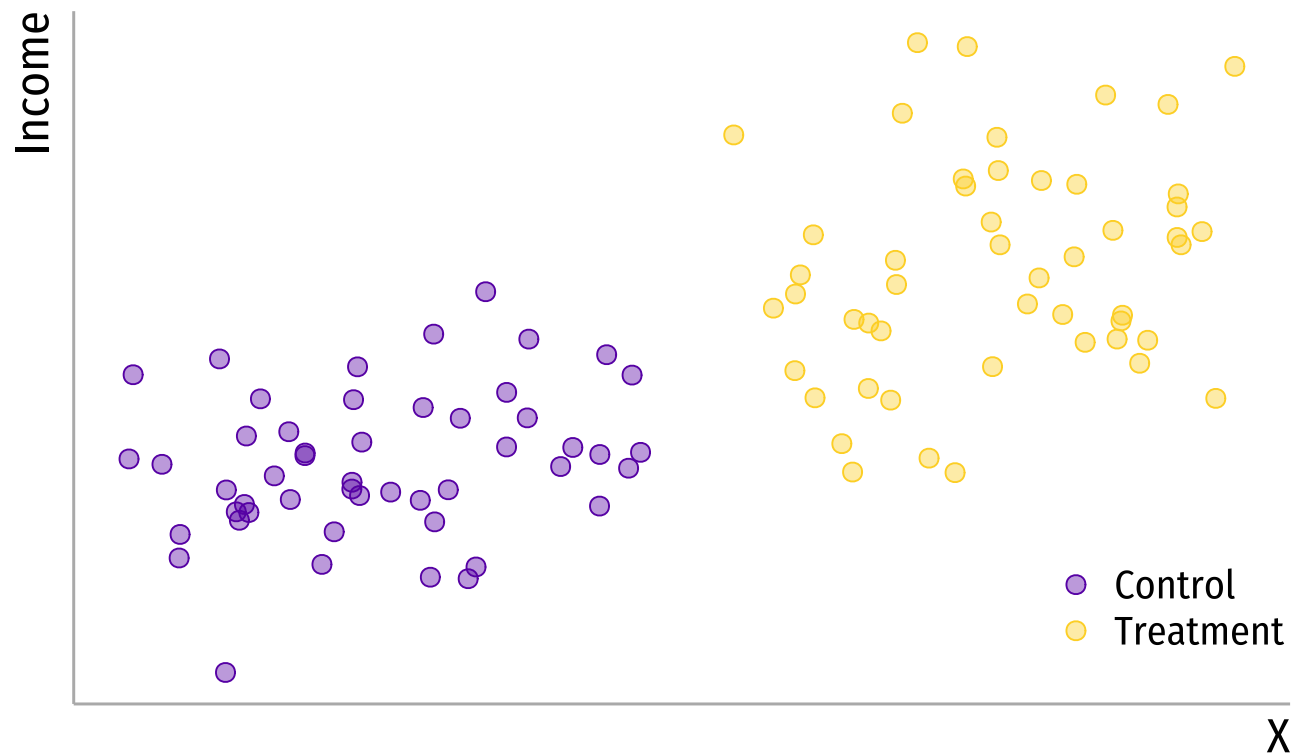
$$Y_i = \beta_0 + \beta_1 Z_i + \beta_2 X_i + \varepsilon_i$$

**$\beta_1$  is the estimated effect of Z on Y, holding X constant**



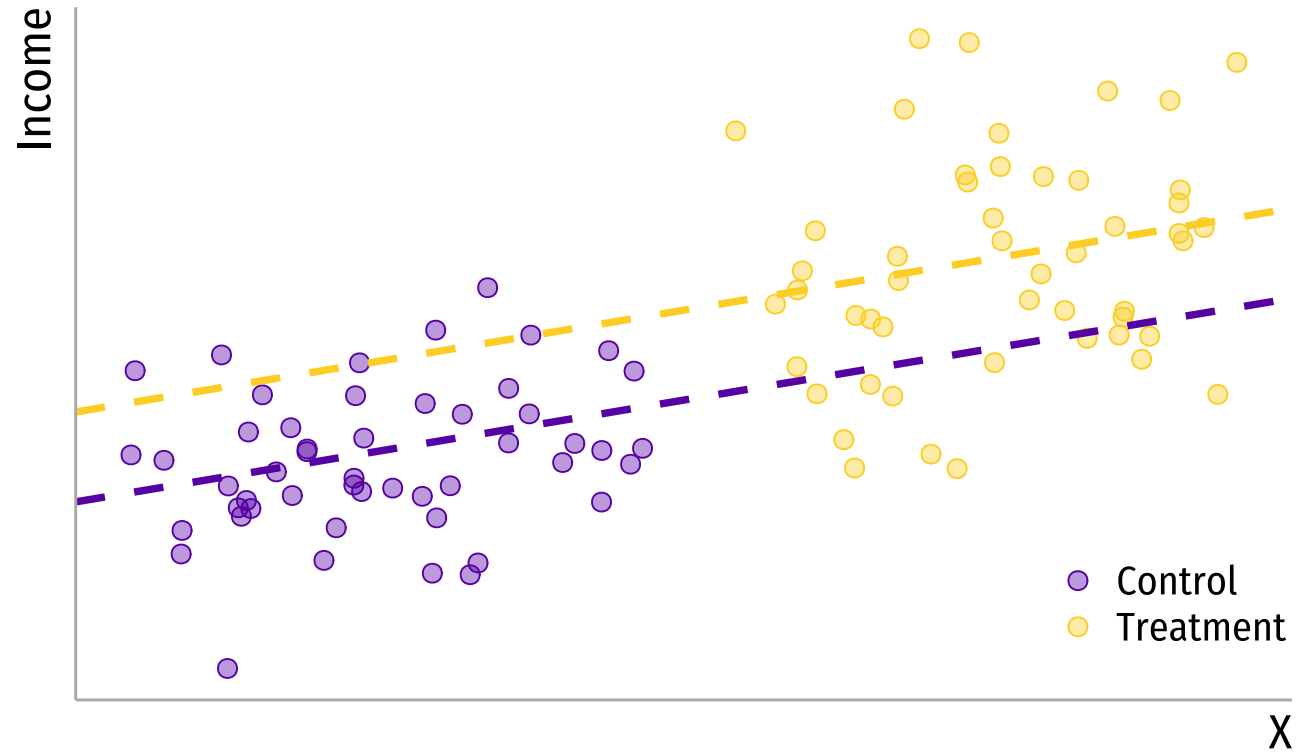
# How do we adjust for observables?

- But what if our data looks like this? Do you see a problem?



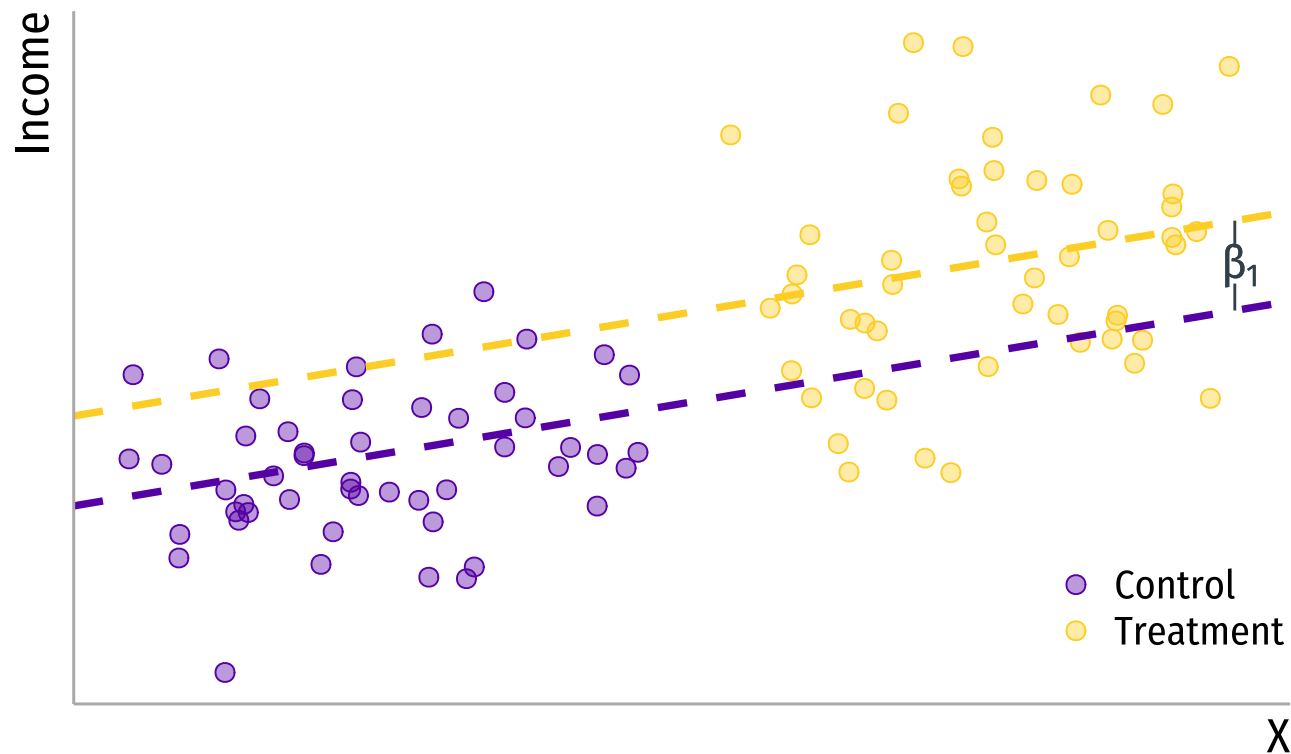
# How do we adjust for observables?

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Finding your perfect match...

# Two peas in a pod

- One other route we could take is to **find similar units** in our sample and **group them together**.
- There are different ways to do it:
  - E.g. subclassification, matching.



# Two peas in a pod

- One other route we could take is to **find similar units** in our sample and **group them together**.
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What do we gain?



# Advantages of matching methods

**Reduce model dependence**

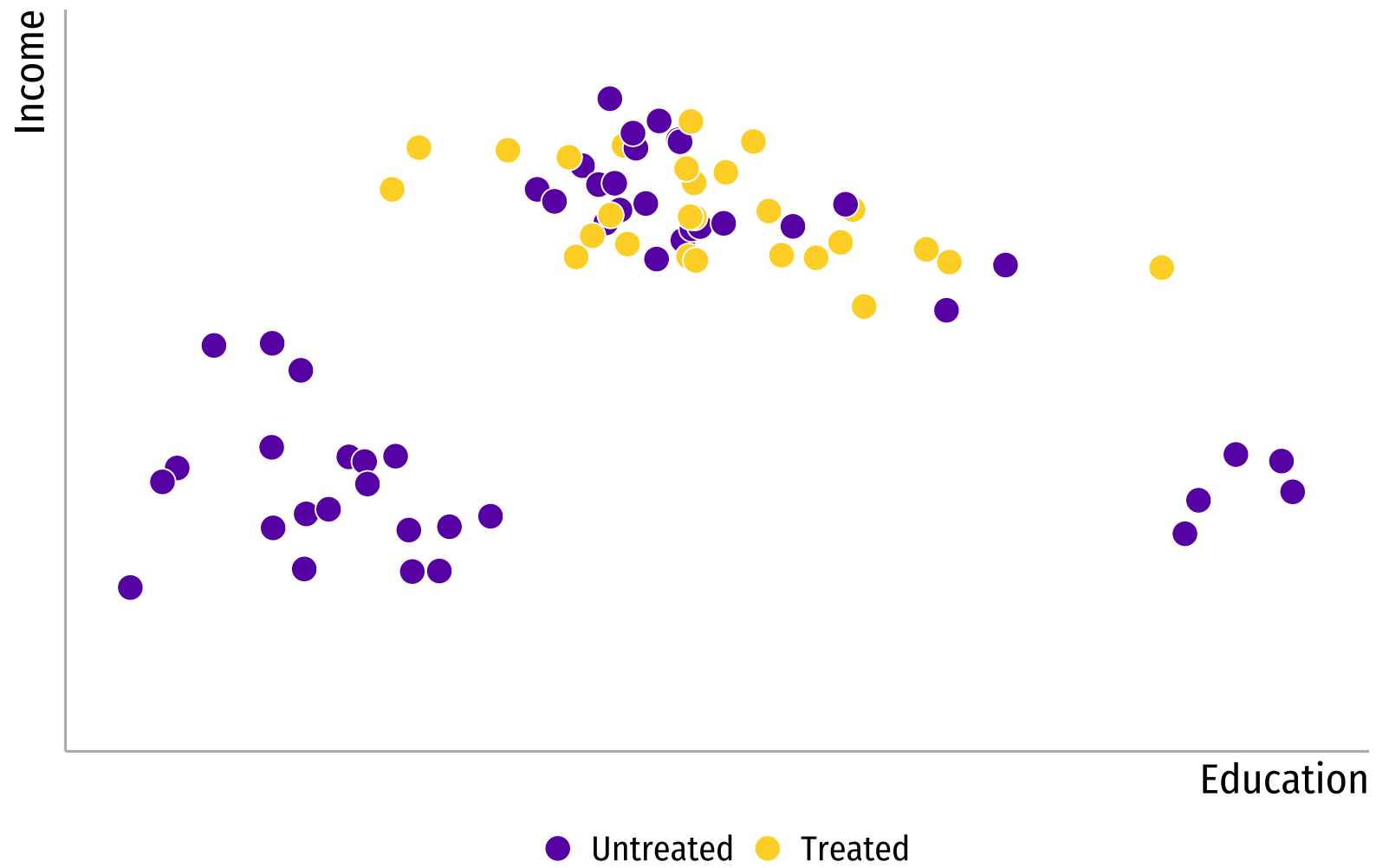
Imbalance → model dependence → researcher discretion → bias

**Compare like to like**

**No extrapolation!**

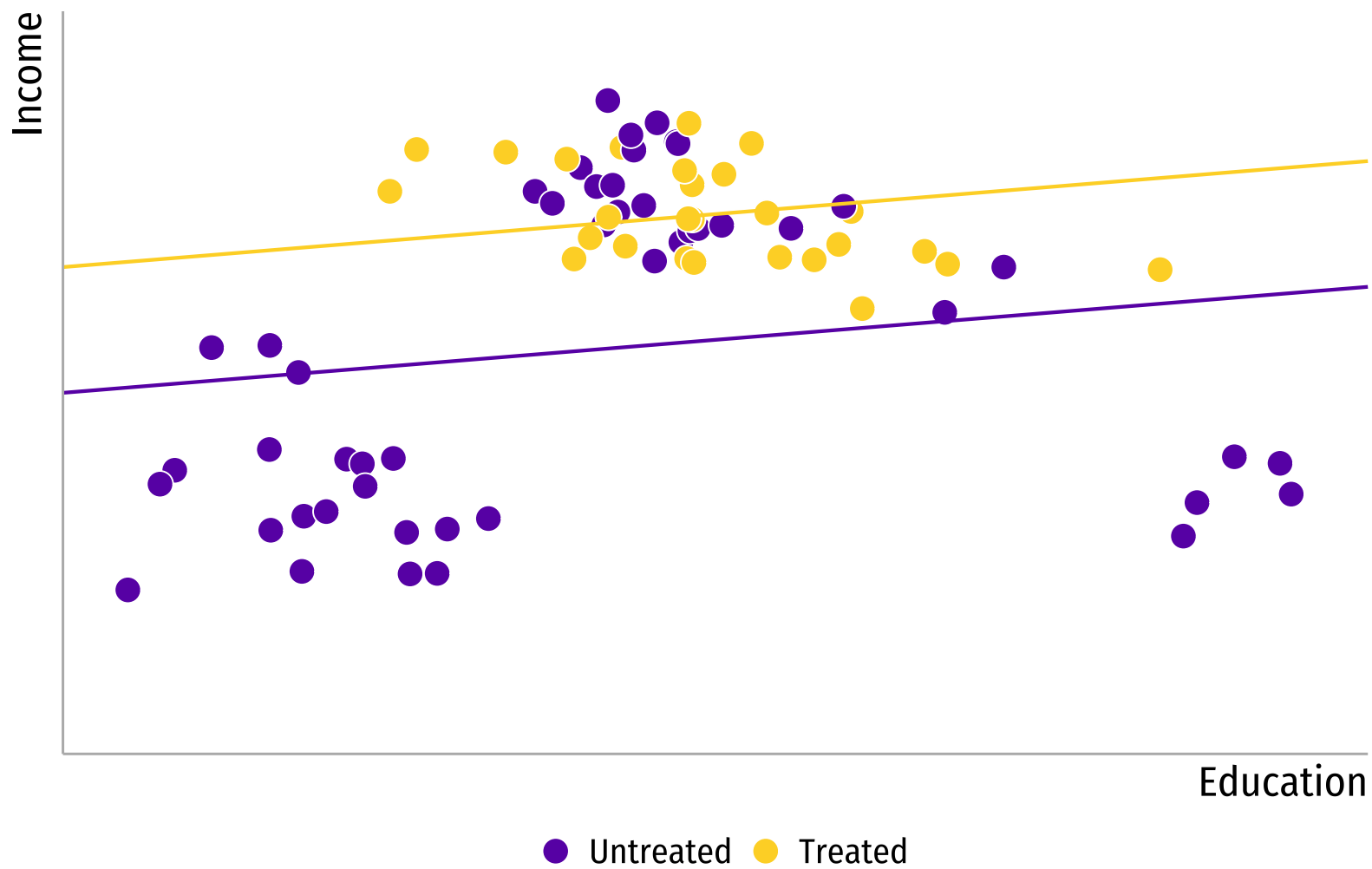
**Can adjust closely by covariates**

Exact matching, coarsened exact matching, fine balance..

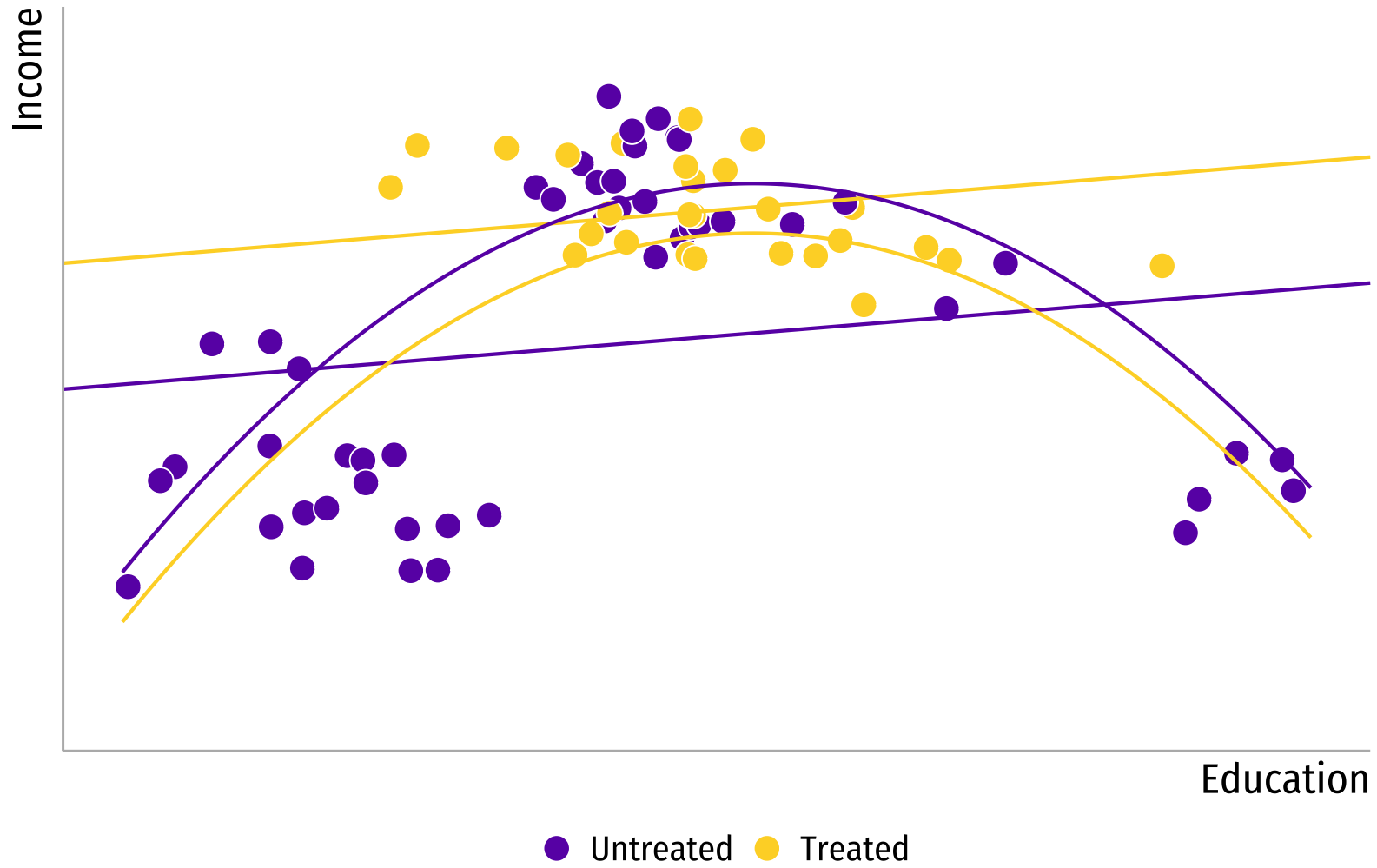


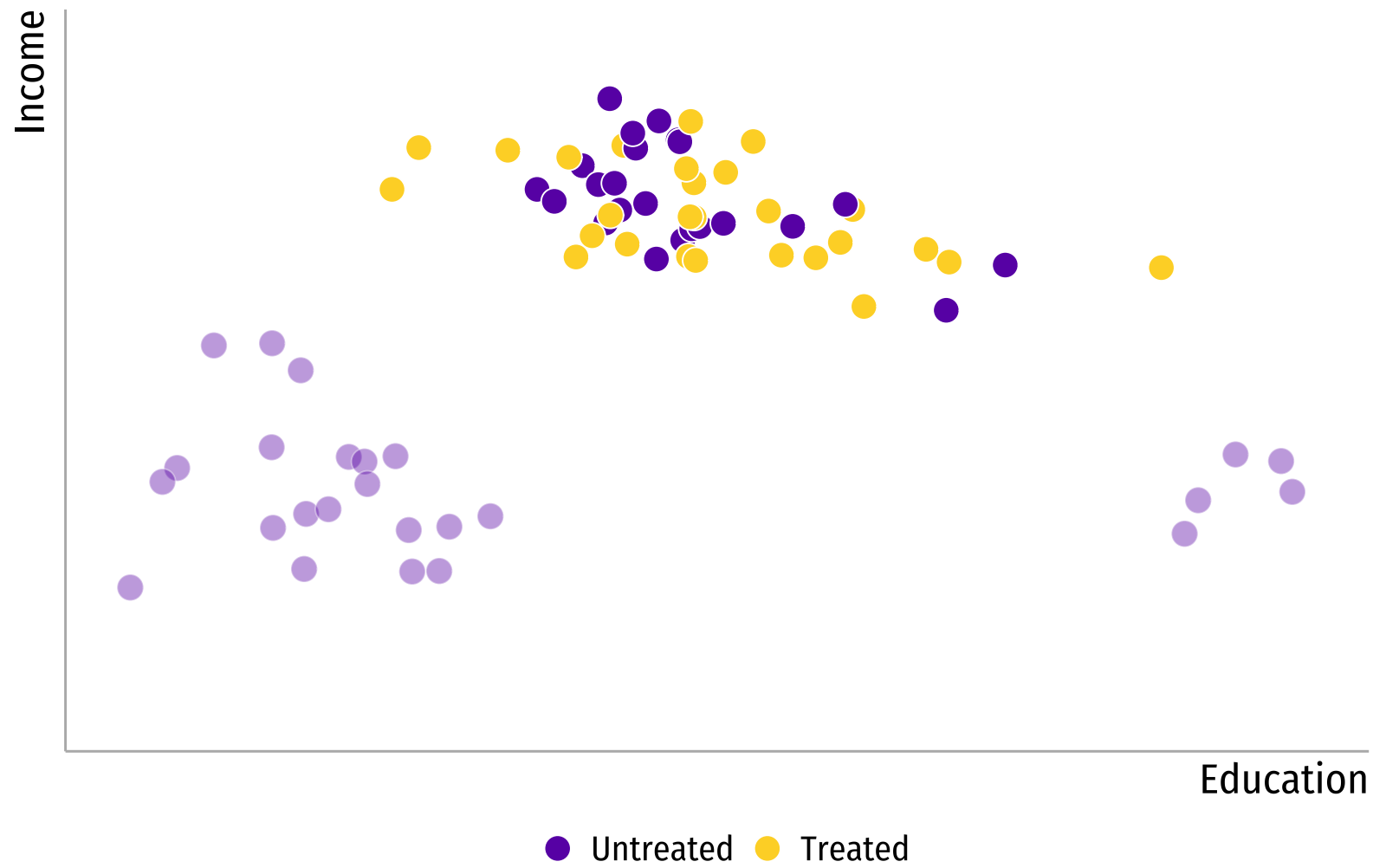


$$\text{Income} = \beta_0 + \beta_1 \text{Education} + \beta_2 \text{Treatment}$$

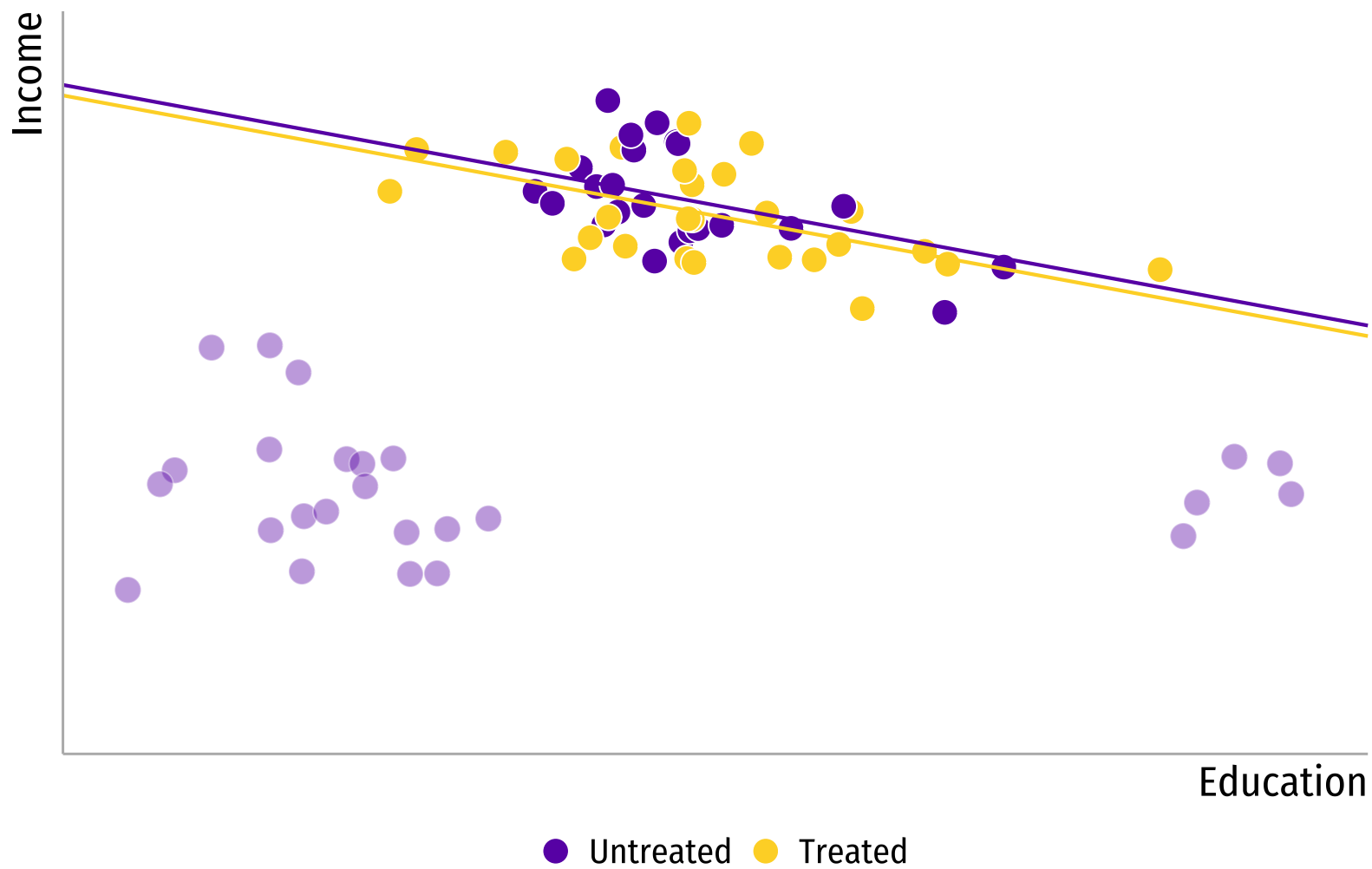


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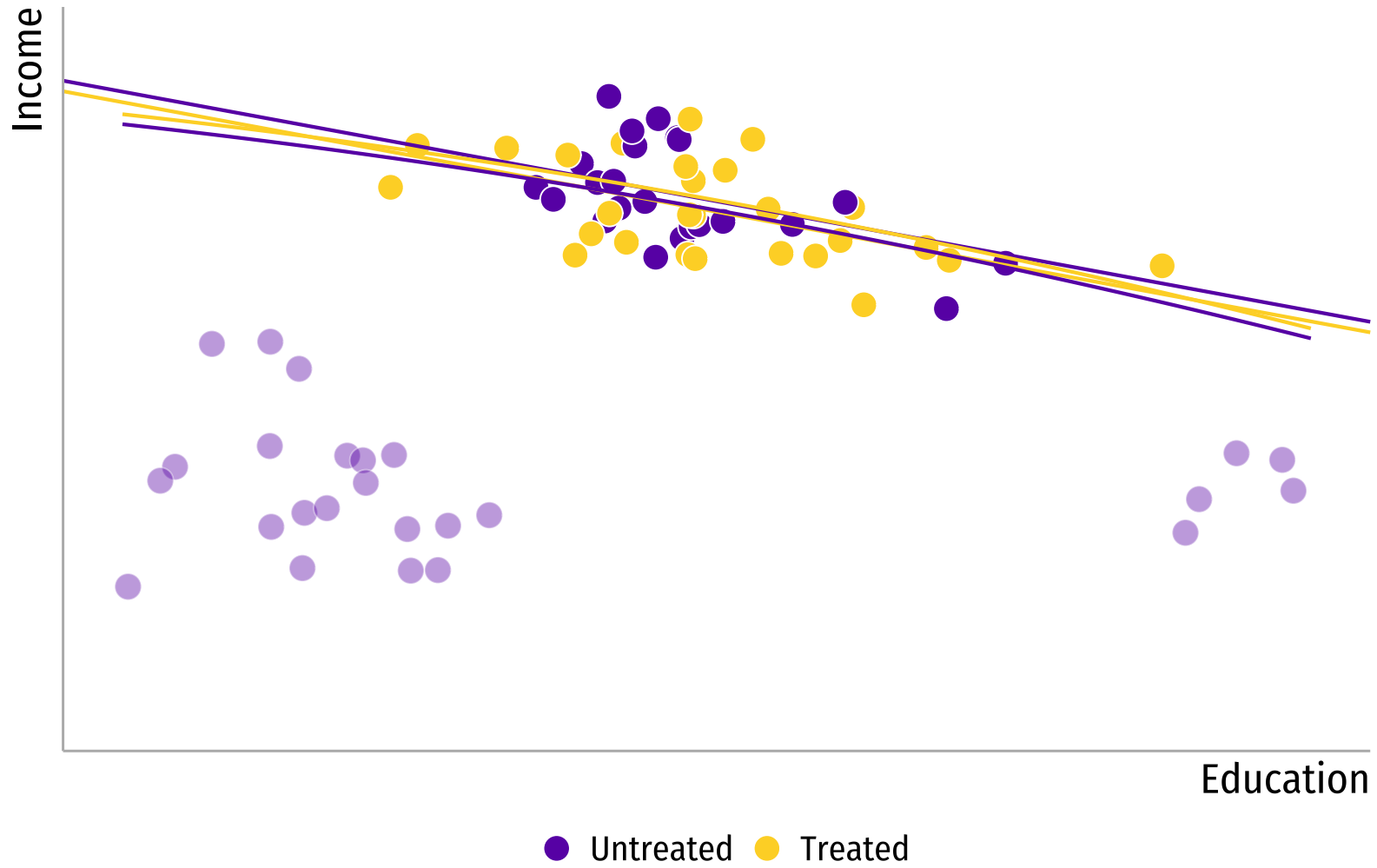




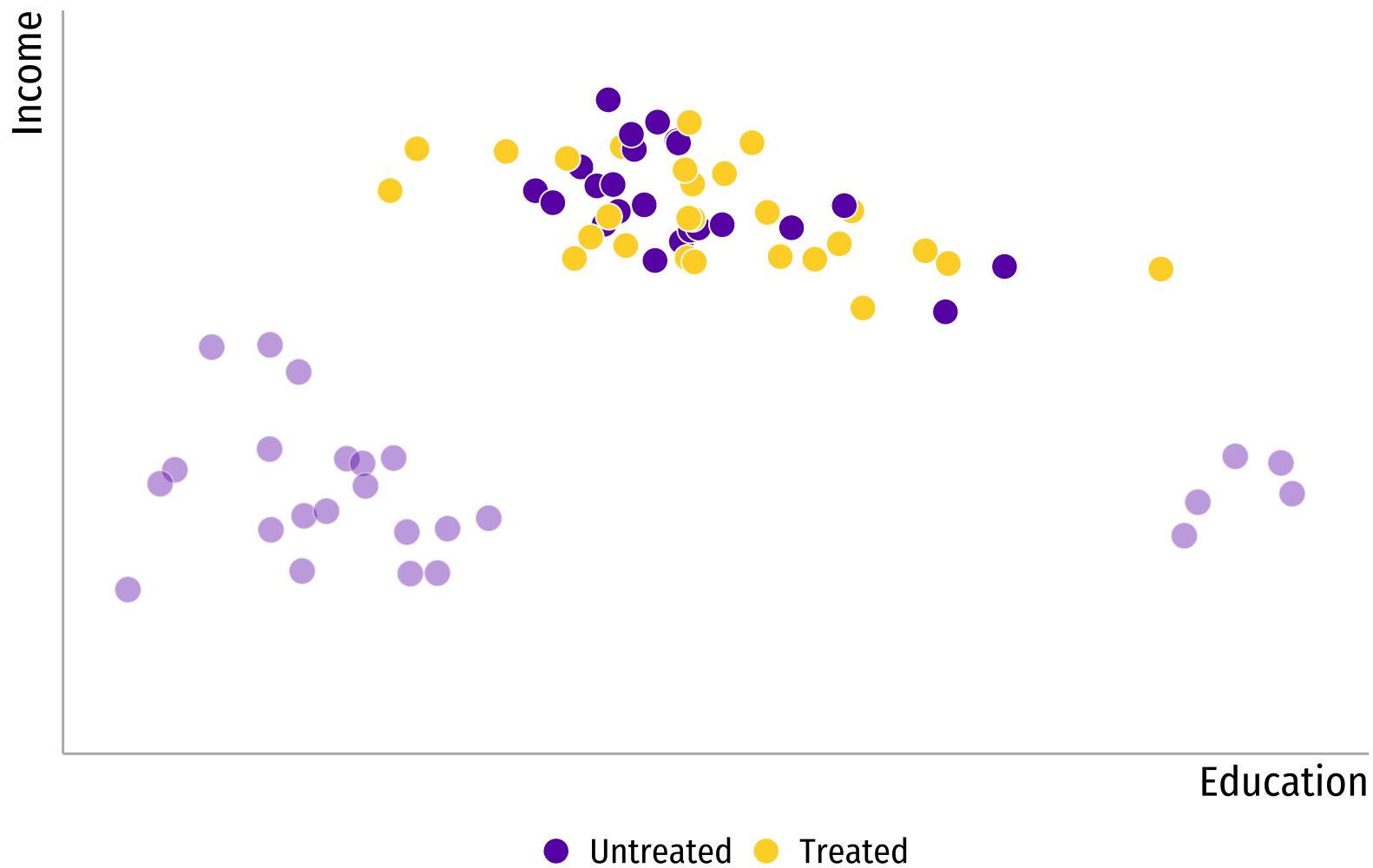
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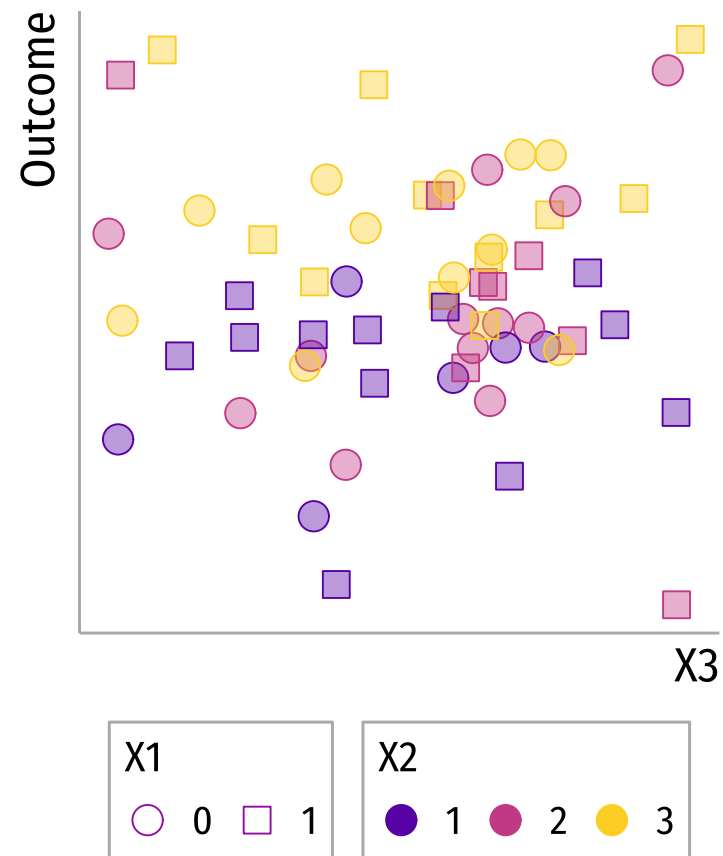


## How do we know we can remove those observations?



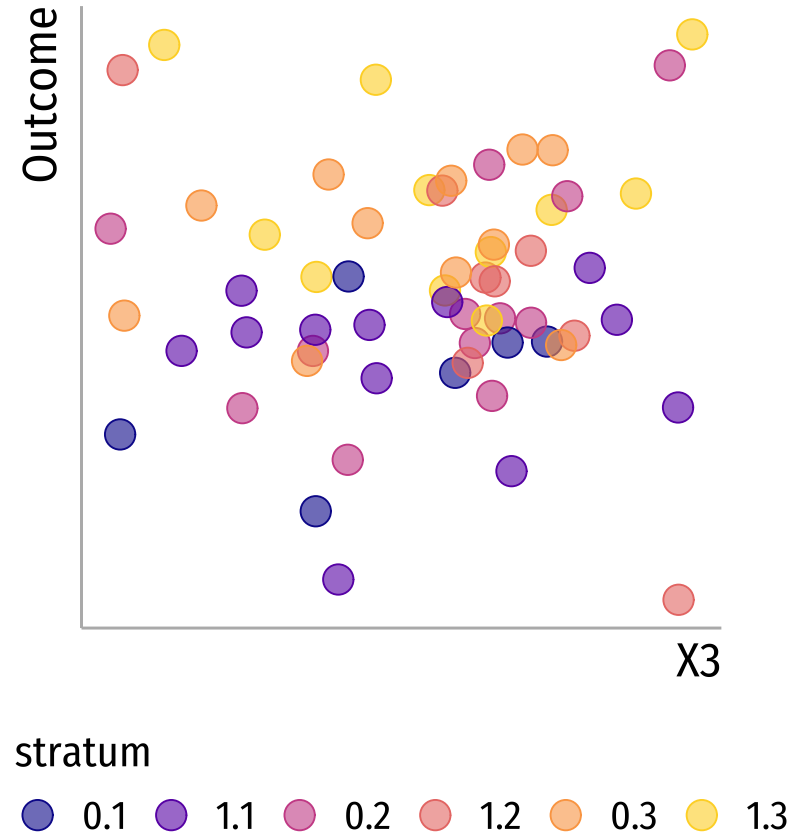
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- Very similar to **stratifying**.



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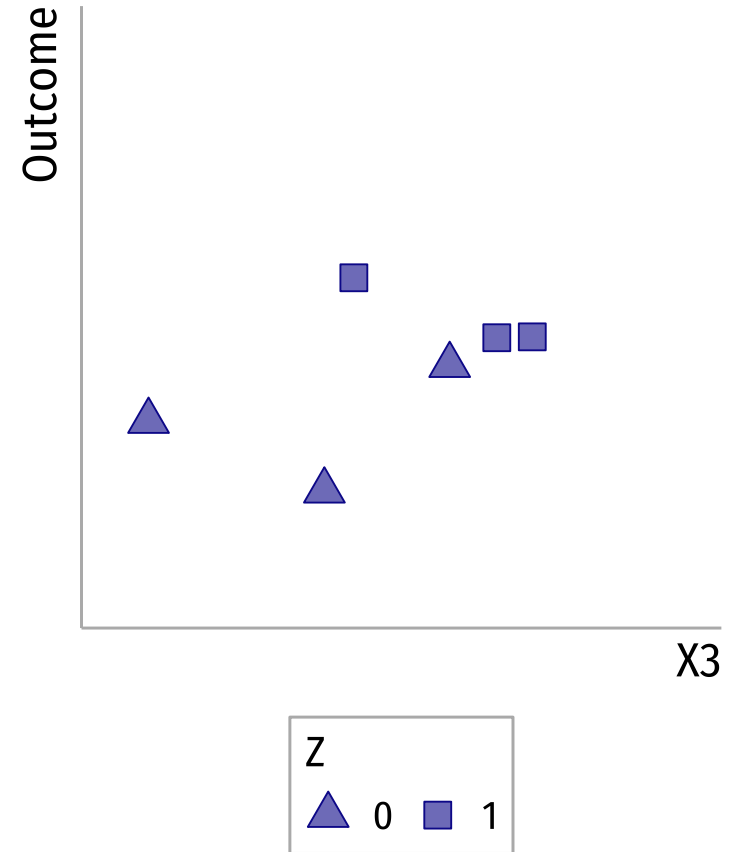
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- Build **combination** of X1 and X2 (strata).





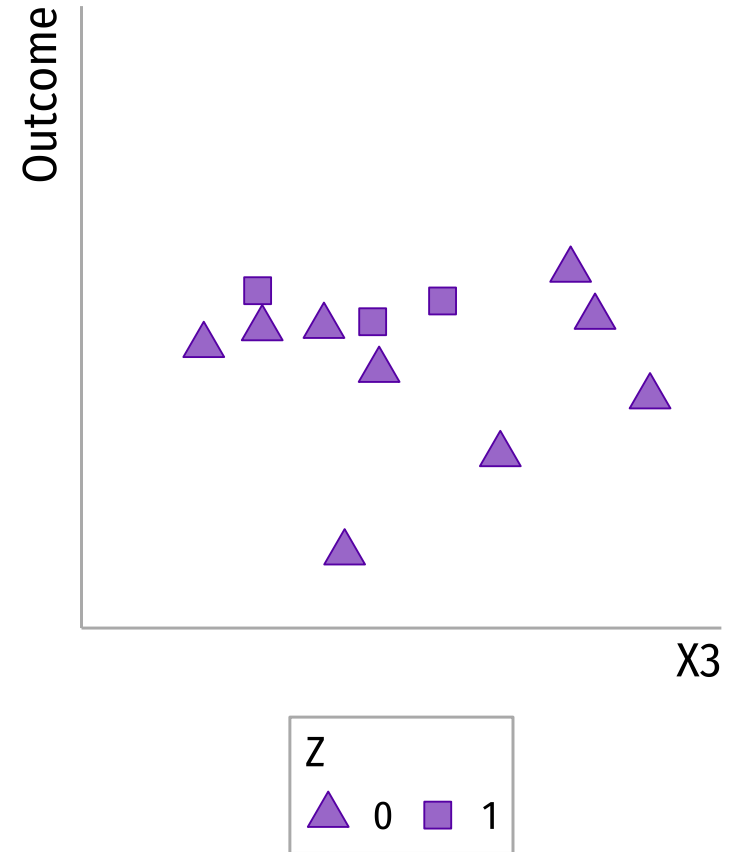
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- Compare **within stratum**.



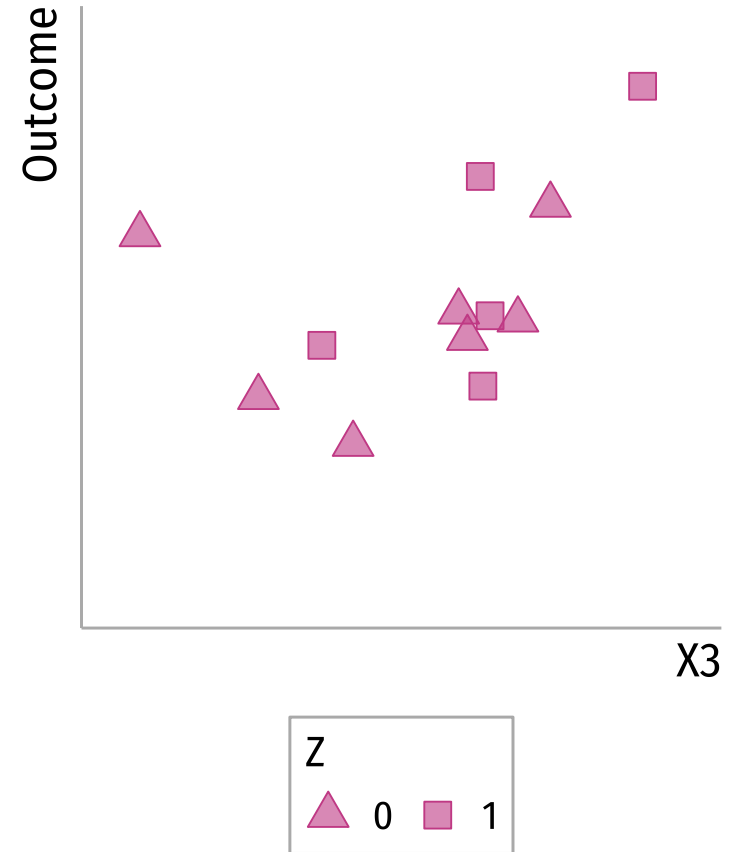
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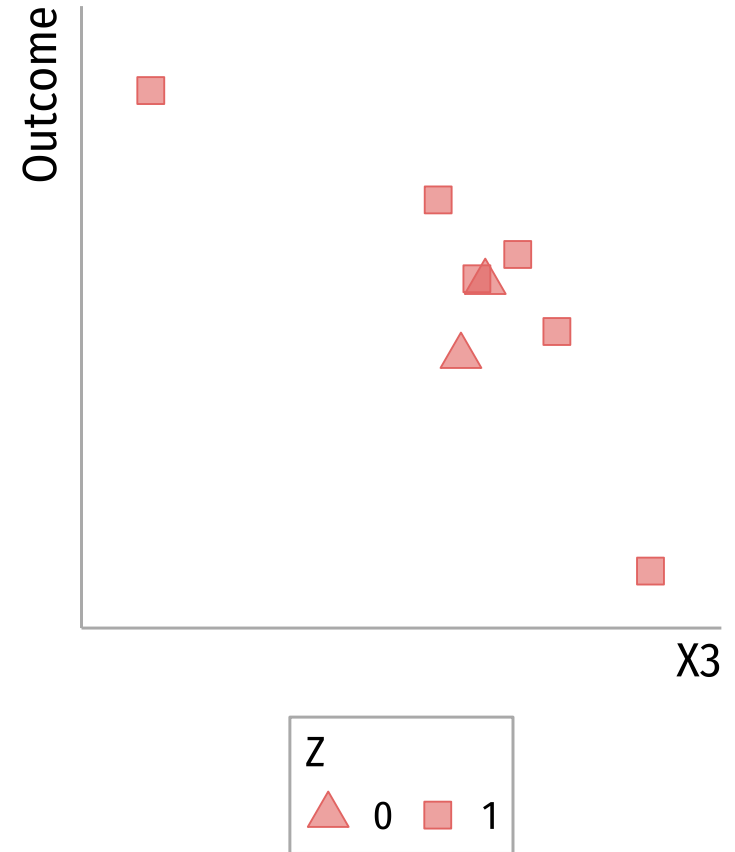
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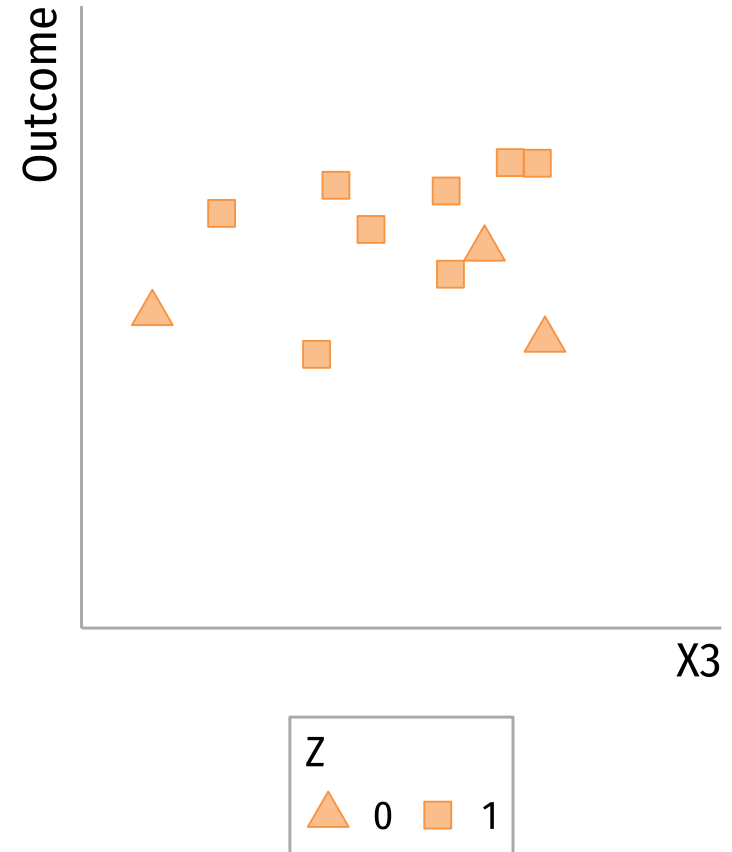
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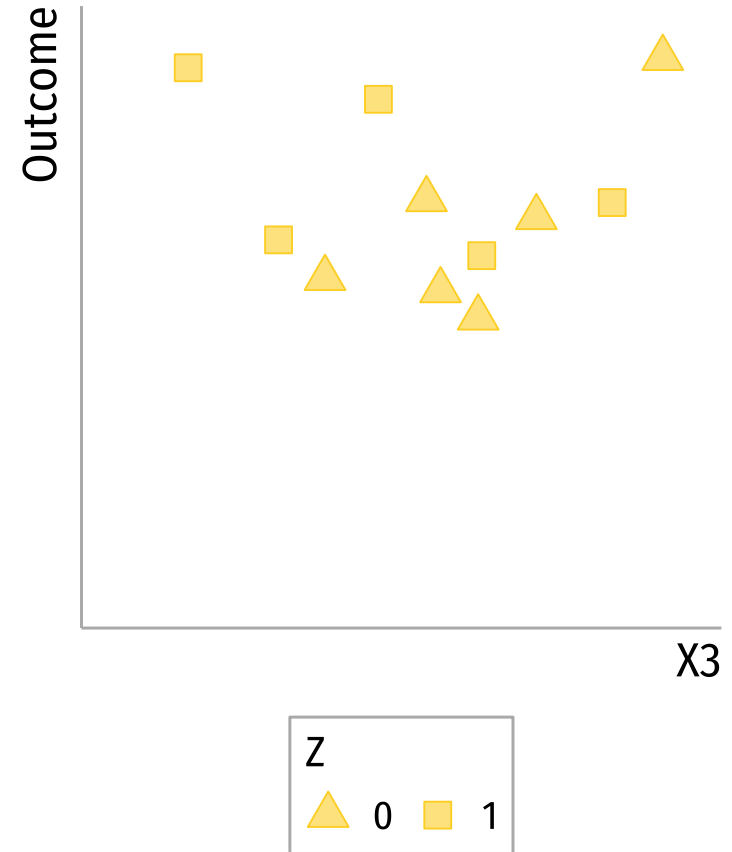
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# Subclassification

- To estimate the Average Treatment Effect, we take a **weighted average**:

$$\hat{ATE} = \sum_{s=1}^S \frac{N_s}{N} (\bar{Y}_{1s} - \bar{Y}_{0s})$$

**What happens when we have too many variables to build strata?**

# The curse of dimensionality

- When we have too many covariates, the number of strata or groups grow **exponentially**!
  - E.g. with 4 covariates, each with 5 categories, we have **625 combinations**!
- Very possible that a stratum only has treatment or control units.





# The curse of dimensionality

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What to do?



# Breaking the curse: Balancing scores

- Want to **reduce the dimensionality** of our covariates
- A balancing score  $b(x)$  is a function of the covariates such that:

$$Z_i \perp\!\!\!\perp X_i | b(X_i)$$

- This means that conditioning on the balancing score is **enough to remove bias** associated to the covariates.
- Under unconfoundedness:

$$Y_i(0), Y_i(1) \perp\!\!\!\perp Z_i | b(X_i)$$

- There are different balancing scores:
  - E.g. propensity scores, mahalanobis distance.

# Estimating balancing scores

## Propensity score

$$\log\left(\frac{p}{1-p}\right) = \beta_0 + \beta_1 X_1 + \beta_2 X_2 + \dots + \beta_p X_p + \varepsilon$$

where  $p = Pr(Z = 1)$

```
e <- predict(glm(z ~ x1 + x2 + x3, data = d, family = binomial(link="logit")),  
             type="response")
```

# Estimating balancing scores

## Propensity score

- Importance of overlap region

# Making groups comparable

- Using the previous balancing scores (or covariates directly!) we can **match observations between the treatment and control group**

## Step 1: Preprocessing

Try to model the treatment assignment

## Step 2: Estimation

Use the new trimmed/preprocessed data to build a model, calculate difference in means, etc.

# How matchy-matchy

- There are different matching methods (and different ways to use them!)

**Nearest neighbor (NN)**

Use balancing scores; Greedy algorithm

**Optimal matching**

Solves an optimization problem; slow on large samples

**Mixed Integer Programming (MIP) matching**

Balances covariates directly; can generate smaller samples

Let's go to R

# The shortcomings of matching

- Many researchers misuse matching and **confuse it with an identification strategy**
- In terms of identification, **matching still relies on selection on observables**

**You need other source of exogeneous variation!**

- Claiming that you can identify a **causal effect** just by using matching is almost the same as claiming this using a regression approach.

**Usually not a good idea...**



# Don't get it twisted

- Matching works great as an **adjustment method**.
- Combined with **other identification strategies**, it can improve results!



# Main takeaways



- Matching methods can be great tools for your analysis.
  - Create more similar groups of comparisons.
  - Reduce model dependence
  - Even help with external validity (under assumptions)

# Next week

- We will look at some **identification strategies** for observational studies:
  - Natural experiments and differences-in-differences.
- What **assumptions** need to hold?
- How do we **identify a natural experiment**?
- What does **DD** buy us?

# References

- Angrist, J. and S. Pischke. (2015). "Mastering Metrics". *Chapter 2*.
- Heiss, A. (2020). "Program Evaluation for Public Policy". *Class 7: Randomization and Matching, Course at BYU*
- Imbens, G. and D. Rubin. (2015). "Causal Inference for Statistics, Social, and Biomedical Sciences: An Introduction". *Chapter 3*
- Cunningham, S. (2021). "Causal Inference: The Mixtape". *Chapter 5*