

# Parametric g-Formula

Cornell STSCI / INFO / ILRST 3900

Fall 2023

[causal3900.github.io](https://causal3900.github.io)

28 Sep 2023

# Learning goals for today

At the end of class, you will be able to

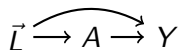
- ▶ estimate average causal effects with a parametric model
  - ▶ for the outcome  $E(Y | A, \vec{L})$
  - ▶ for the treatment  $P(A | \vec{L})$

After class:

- ▶ Hernán and Robins 2020 Chapter 12.1–12.5, 13, 15.1

# Nonparametric estimation

Causal assumptions



Nonparametric estimator

$$\hat{E}(Y^a) = \frac{1}{n} \sum_{i=1}^n \hat{E}(Y \mid \vec{L} = \vec{\ell}_i, A = a)$$

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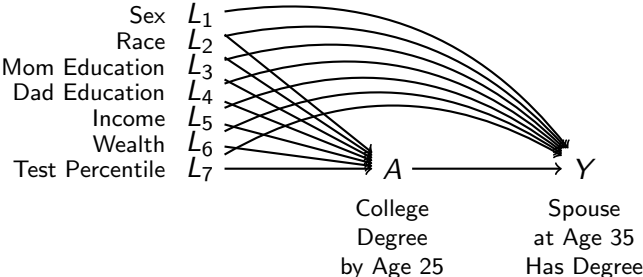
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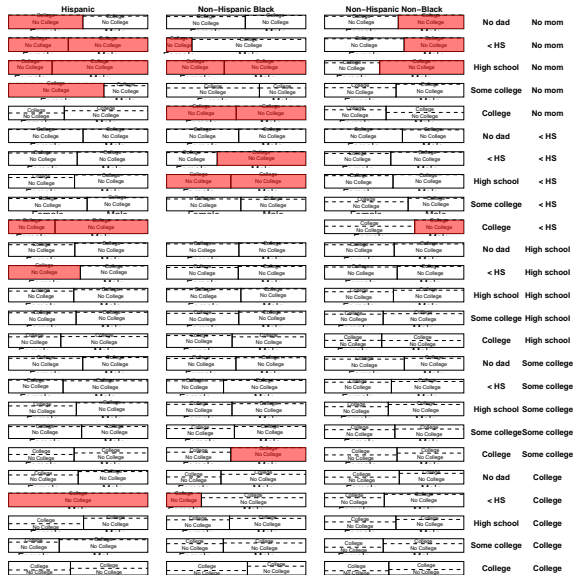
- ▶ find units who look like them on confounders  $\vec{L}$
- ▶ who actually got treatment  $A = a$
- ▶ take the average among those units

Then average over all units

# Nonparametric estimation breaks down

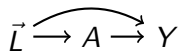


# Nonparametric estimation breaks down



# Parametric estimation: Outcome model

Causal assumptions

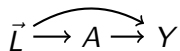


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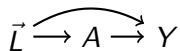
First, learn a model to predict  $Y$  given  $\vec{L}$  and  $A$

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For every unit  $i$ ,

- ▶ change the treatment value to  $a$
- ▶ predict the outcome

Then average over all units

The parametric g-formula: Connection to  $\hat{\beta}$

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Estimator for the effect  $E(Y^1) - E(Y^0)$ :

## The parametric g-formula: Connection to $\hat{\beta}$

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With OLS, the parametric g-formula collapses on the coefficient.

The parametric g-formula is more general

Suppose the effect of  $A$  depends on  $L$

$$E(Y | A, L) = \alpha + \gamma L + \beta A + \eta AL$$



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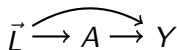
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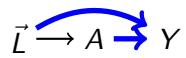
The g-formula no longer collapses to a coefficient!

## Parametric g-formula: Outcome model recap



1. Model the outcome mean  $E(Y | A, \vec{L})$
2. Change everyone's treatment to the value of interest
3. Predict for everyone
4. Take the average

$$\hat{E}(Y^a) = \frac{1}{n} \sum_{i=1}^n \hat{E}(Y | \vec{L} = \vec{\ell}_i, A = a)$$

$$\vec{L} \rightarrow A \rightarrow Y$$


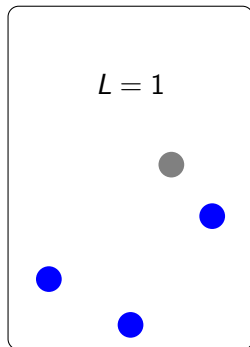
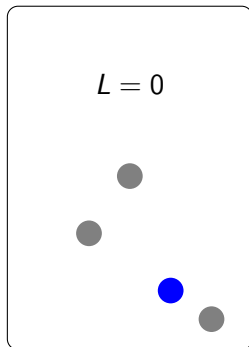
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# Inverse probability of treatment weighting

● Untreated

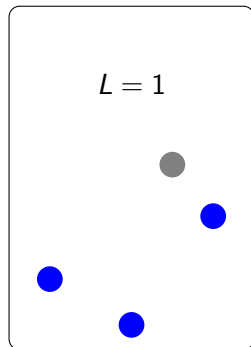
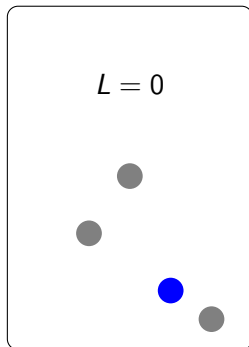
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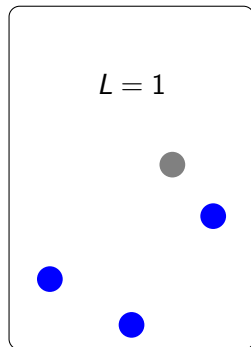
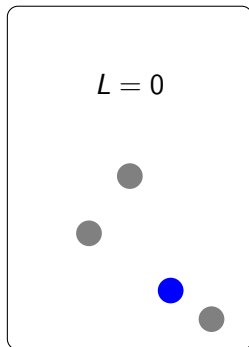
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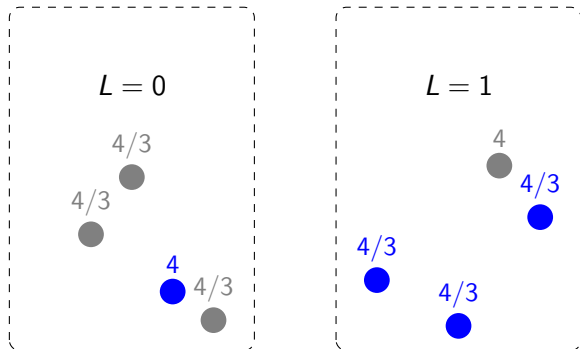
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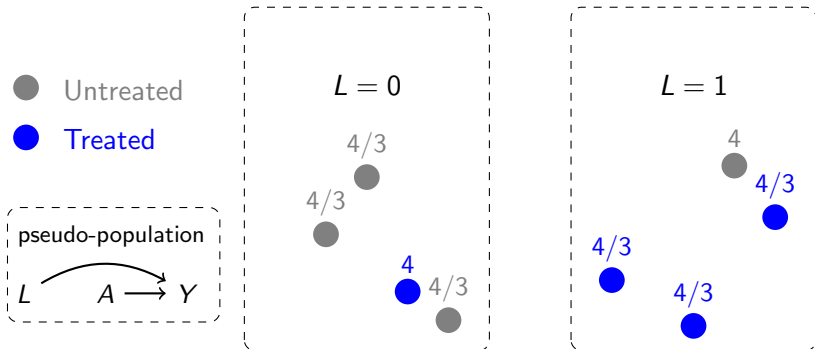
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Propensity score:  $\pi_i = P(A = A_i \mid L = L_i)$

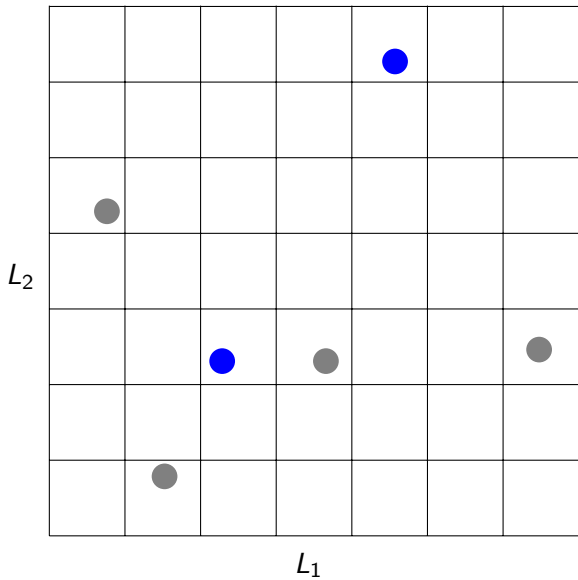
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Propensity score:  $\pi_i = P(A = A_i | L = L_i)$

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**Model** the treatment assignment

$$\hat{P}(A = 1 \mid \vec{L}) = \text{logit}^{-1} \left( \hat{\alpha} + \hat{\gamma} \vec{L} \right)$$

**Predict** the propensity score for each unit

$$\hat{\pi}_i = \begin{cases} \text{logit}^{-1} \left( \hat{\alpha} + \hat{\gamma} \vec{L} \right) & \text{if } A_i = 1 \\ 1 - \text{logit}^{-1} \left( \hat{\alpha} + \hat{\gamma} \vec{L} \right) & \text{if } A_i = 0 \end{cases}$$

**Estimate** by inverse probability weighting

$$\hat{E}(Y^a) = \frac{1}{N} \sum_{i:A_i=a} \frac{Y_i}{\hat{\pi}_i}$$

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Two solutions

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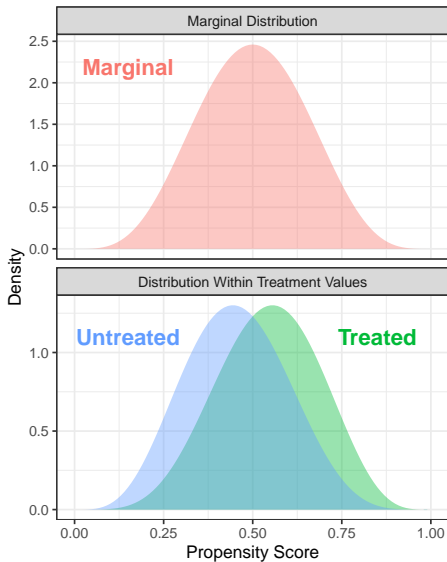
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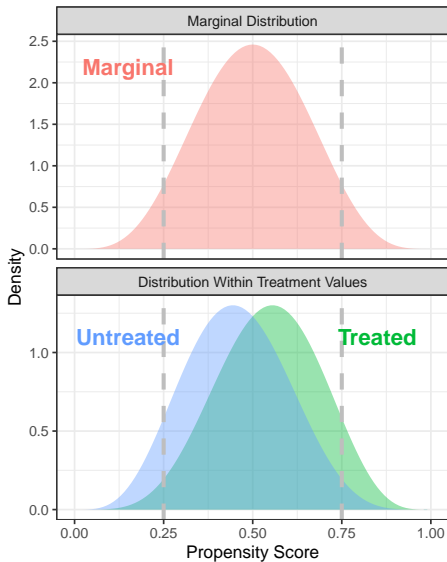
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Both solutions accept bias in order to reduce variance

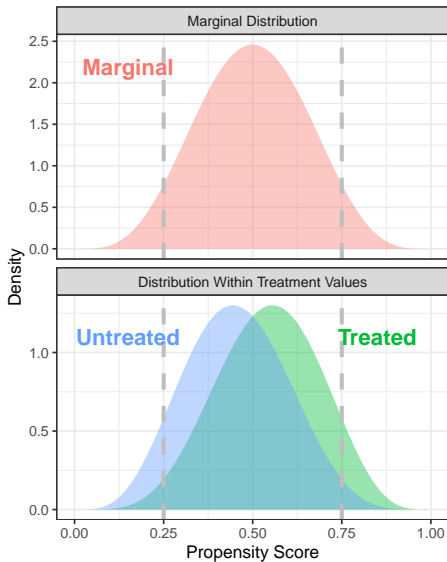
# Accepting bias to reduce variance: Trimming



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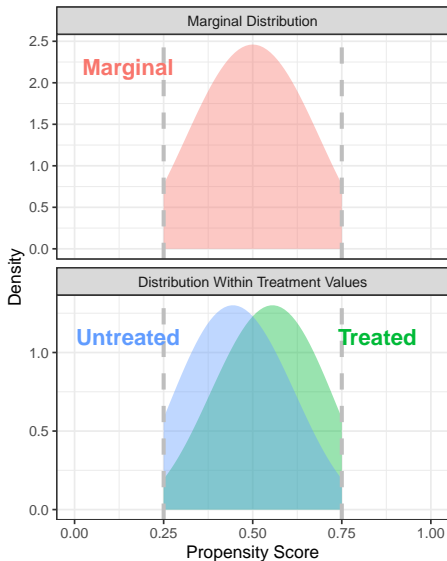


# Accepting bias to reduce variance: Trimming



Drop units with extreme weights

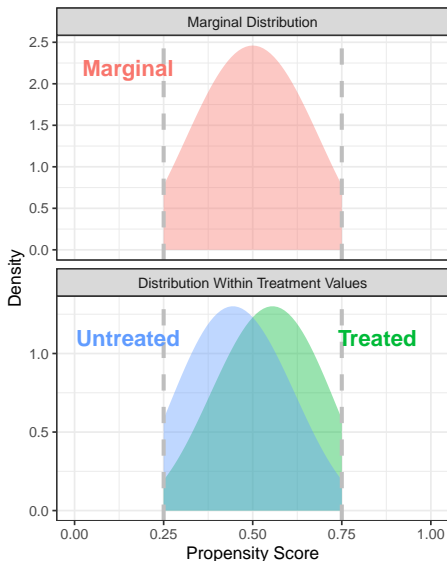
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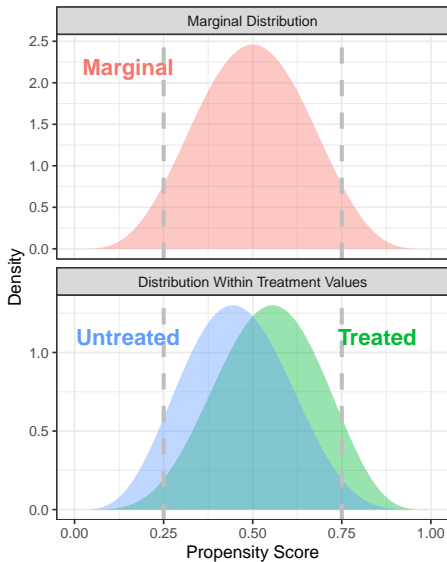
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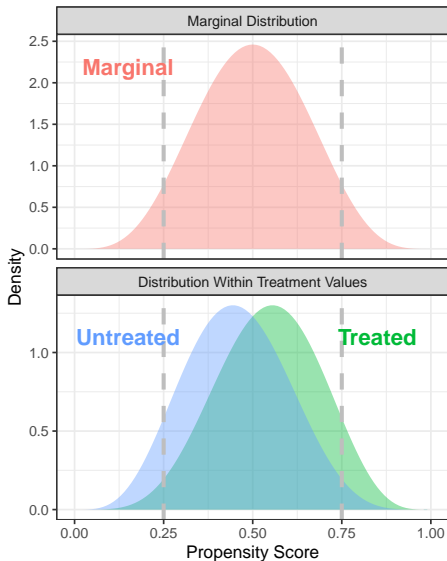
Drop units with  
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Changes target population  
— Biased for full population

# Accepting bias to reduce variance: Weight truncation

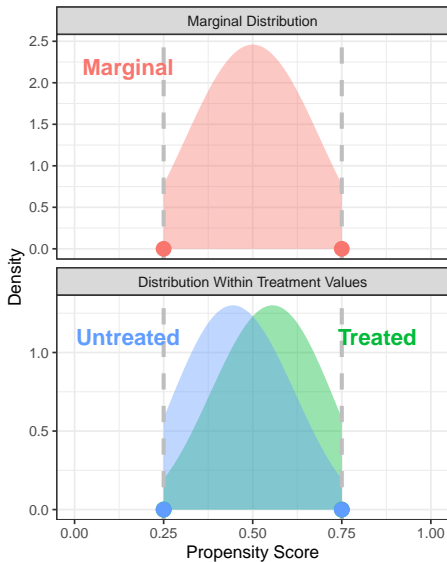


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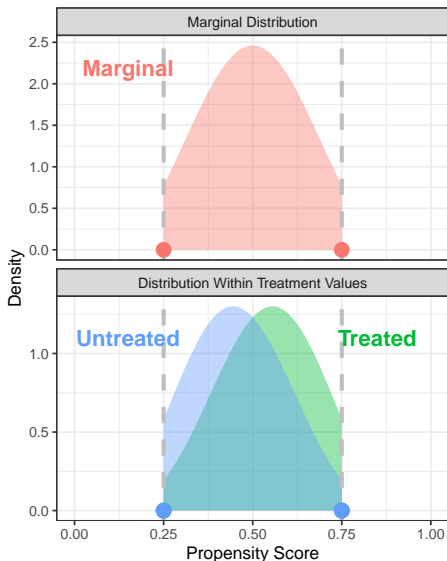
Truncate values of extreme weights

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Biased: Ignores some confounding