

## 7. Positivity: The problem of empty cells

Ian Lundberg

Cornell Info 6751: Causal Inference in Observational Settings  
Fall 2022

13 Sep 2022

# Learning goals for today

At the end of class, you will be able to:

1. Define positivity
2. Understand how positivity relates to the adjustment set
3. Make estimates for the feasible subpopulation
4. Begin translating ideas to actual data

# Positivity

Assumption that  $P(A = a \mid \vec{L} = \vec{\ell}) > 0$  (i.e., is positive) for all treatment values  $a$  in all population strata  $\vec{\ell}$  defined by confounders

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**Why this matters:**

# Positivity

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## Why this matters:

If there is a subgroup  $\vec{L} = \vec{\ell}$  where the treatment  $A = a$  never happens, then we could never learn from data about the outcome in that subgroup under that treatment.

# Running example for today's discussion

Government Assistance  
Protects Low-Income  
Families from Eviction

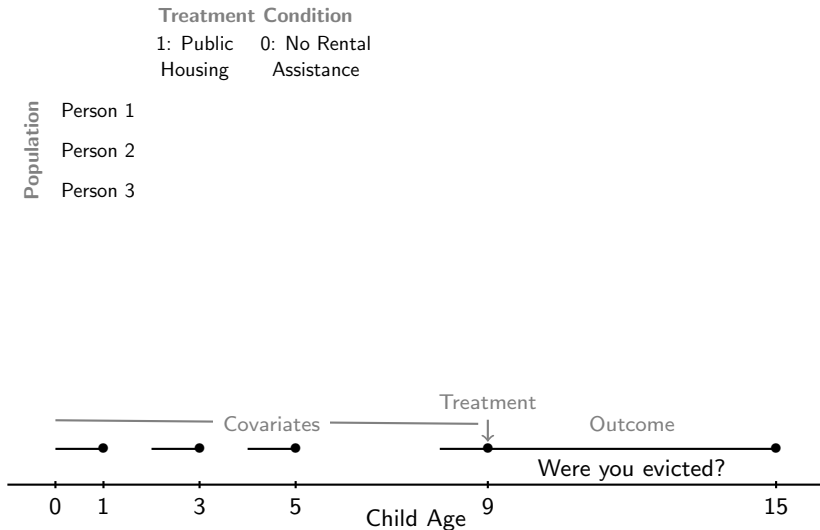
*Ian Lundberg*  
*Sarah L. Gold*  
*Louis Donnelly*  
*Jeanne Brooks-Gunn*  
*Sara S. McLanahan*

Journal of Policy Analysis and Management  
2021

## Running example for today's discussion

Does public housing protect families from eviction?

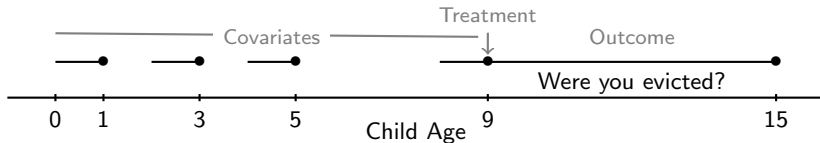
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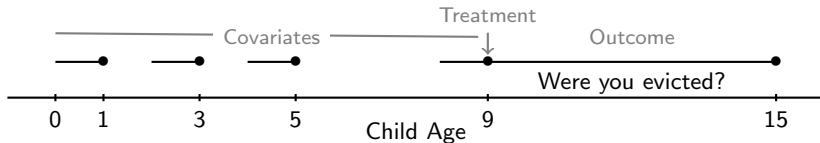
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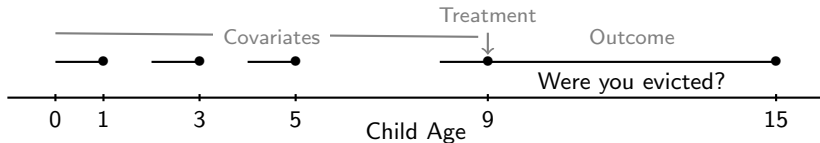
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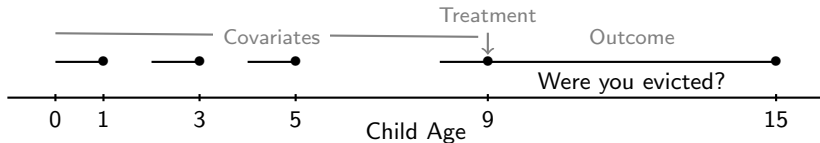
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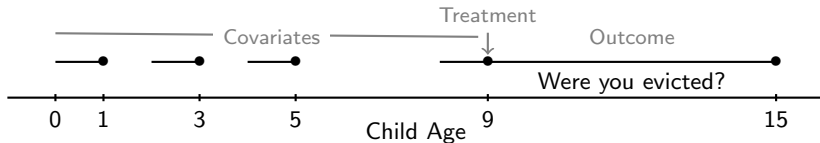
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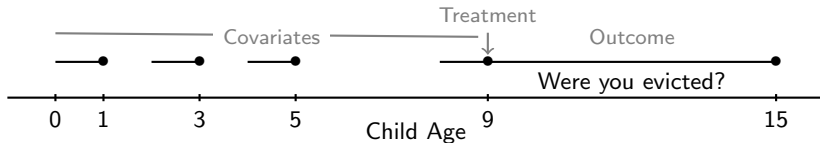
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Robins 1986  
Hahn 1998



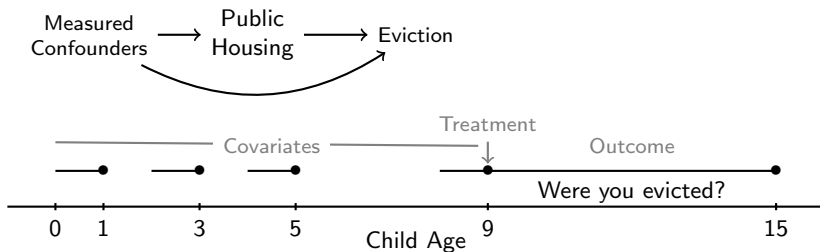
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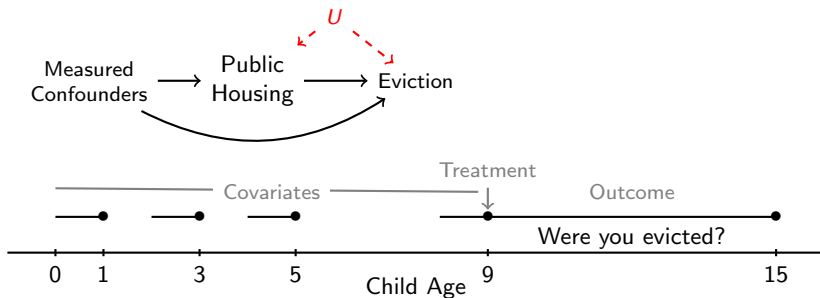
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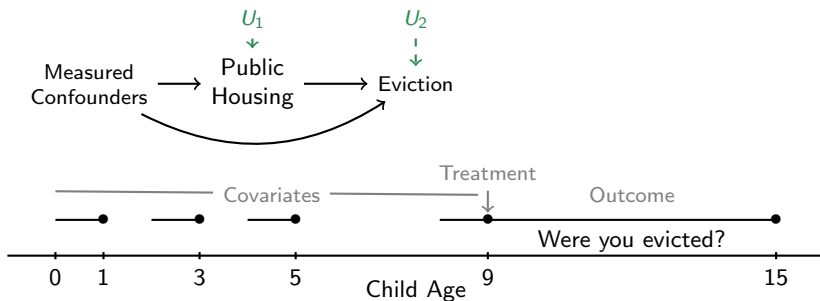
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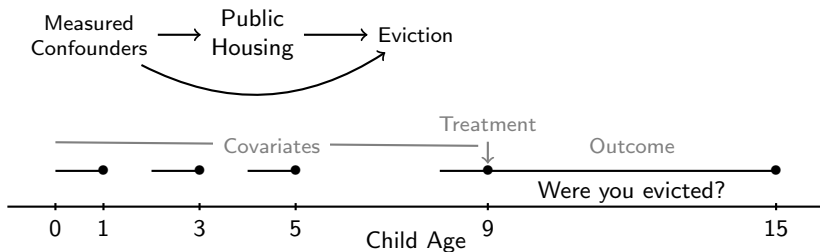
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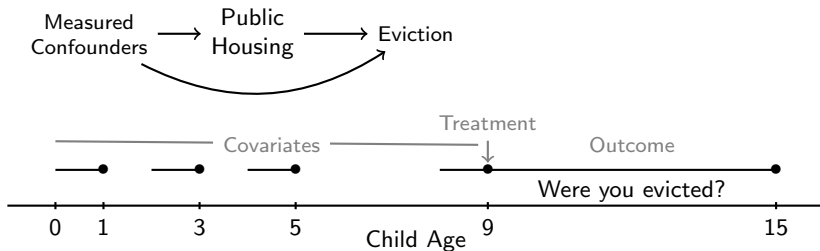
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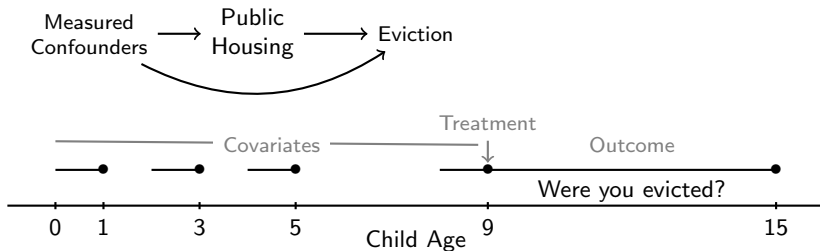
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<b>Average</b>		<b>3%</b>	<b>11%</b>



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— Caveat: Your method must be nonparametric (no regression)



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1. Among those with (lagged eviction = TRUE), estimate the effect
2. Among those with (lagged eviction = FALSE), estimate the effect
3. Take a weighted average of the two estimates, weighted by the number of cases in each group

Nonparametric estimation: In R code

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```
# Load packages
library(tidyverse)

# Simulate some data
sim_data <- data.frame(x1 = rbinom(n,1,.5),
                      x2 = rbinom(n,1,.5),
                      x3 = rbinom(n,1,.5)) %>%

# Generate the treatment
mutate(a = rbinom(n,1,plogis(x1 + x2 + x3))) %>%
# Generate the outcome
mutate(y = rnorm(n, x1 + x2 + x3 + a))
```

# Nonparametric estimation: In R code

```
# Define the confounders
confounders <- c("x1", "x2", "x3")

# Count cases in each stratum
strata_counts <- sim_data %>%
  # Group by the confounders
  group_by(across(all_of(confounders))) %>%
  # Count the number of cases
  summarize(cases = n(),
            .groups = "drop")
```

# Nonparametric estimation: In R code

```
# Estimate effect in each stratum
strata_effects <- sim_data %>%
  # Group by the confounders and treatment
  group_by(a, across(all_of(confounders))) %>%
  # Estimate the mean outcome
  summarize(ybar = mean(y),
            .groups = "drop") %>%
  # Prepare to make the data wider by re-valuing the treatment
  mutate(a = paste0("ybar_",a)) %>%
  # Make the data wide
  pivot_wider(names_from = "a", values_from = "ybar") %>%
  # Estimate the effect
  mutate(conditional_effect = ybar_1 - ybar_0)
```

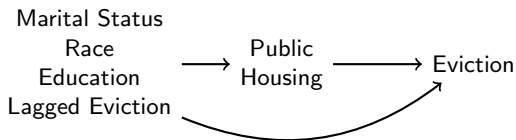
The `pivot_wider` step makes a conversion like this:

Stratum	a	ybar		Stratum	ybar_1	ybar_0
1	ybar_1	3.6	→	1	3.6	3.2
1	ybar_0	3.2		2	3	2.8
2	ybar_1	3		⋮	⋮	⋮
2	ybar_0	2.8				
⋮	⋮	⋮				

# Nonparametric estimation: In R code

```
# Aggregate over strata
strata_counts %>%
  # Merge the effects into the counts data frame
  full_join(strata_effects, by = confounders) %>%
  # Stop working within strata. Average the effect
  ungroup() %>%
  summarize(average_effect = weighted.mean(conditional_effect,
                                           w = cases))
```

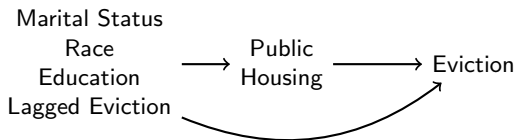
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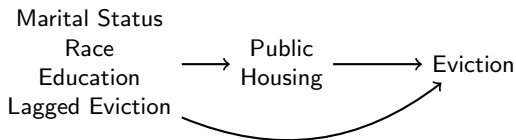
**Question:** How would you estimate in this setting?

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1. In each subgroup defined by the covariates, estimate the effect
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**PROBLEM:** In some subgroups, the treatment does not vary

# To discuss

**Situation:** In some subgroups, the treatment does not vary

**Example:** (oversimplified for concreteness)

When a mother has a college degree,  
her family is never seen in public housing

**Discuss:** Does causal inference make sense for this subgroup

1. if this happens in the sample but not the population?
2. if this happens in the sample and in the population?

If it makes sense, how might you go about it?

## Positivity assumption (in the population)

$$P(A = a \mid \vec{L} = \vec{\ell}) > 0$$

for all treatment values  $a$  in the causal estimand  
and

for every covariate stratum  $\vec{\ell}$  in the population of interest

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Guarantees that in an infinite sample you will eventually see the needed treatment values

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You can theorize about these questions.

But they will never happen—in an infinite sample, you'd never learn the answer.

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Note:

- ▶ Empirical positivity implies theoretical positivity
  - ▶ If we saw all treatments in this subgroup in our sample, they must exist in the population
- ▶ A lack of empirical positivity does not imply a lack of theoretical positivity
  - ▶ All the treatment values may exist in this subgroup, and our sample just happened to miss them

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We will go do this in R.

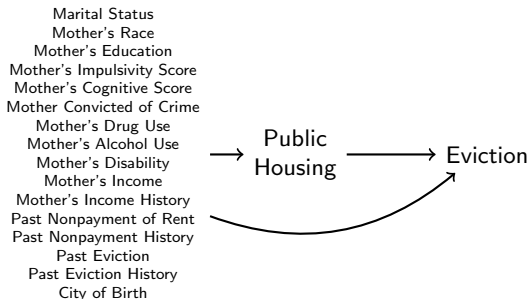
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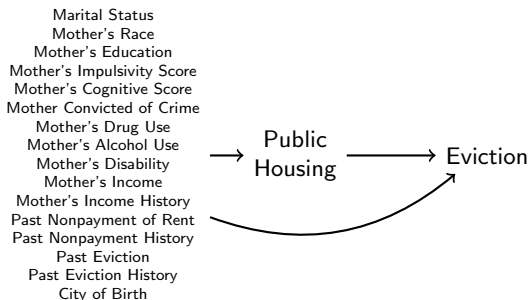
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Let's see how many strata have both

- ▶ Cases in public housing and
- ▶ Cases with no assistance



Conclusion: Positivity is deceptively hard to satisfy

---

<sup>1</sup>D'Amour, A., Ding, P., Feller, A., Lei, L., & Sekhon, J. (2021). Overlap in observational studies with high-dimensional covariates. *Journal of Econometrics*, 221(2), 644-654.

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Positivity  $P(A = a \mid \vec{L} = \vec{\ell}) > 0$  may seem straightforward.

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  - ▶ Tons of strata
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  - ▶ Tons of strata
  - ▶ Hard to populate all treatments in all of them
- ▶ You end up leaning on a model to extrapolate (next class)

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Let me know what you are thinking

[tinyurl.com/CausalQuestions](https://tinyurl.com/CausalQuestions)

Office hours TTh 11am-12pm and at  
[calendly.com/ianlundberg/office-hours](https://calendly.com/ianlundberg/office-hours)  
Come say hi!