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Conditional Independence Assumption

Comparison of Matching and Regression

Propensity Score Matching (PSM)

The Two PSM Lemmas Identification Estimation Strategy PSM Estimation Algorithm PSM Estimation Algorithm

Estimation of ATT by Stratification on the Score

Nearest Neighbor, Kernel and Radius

Weighting on the Score

Econometrics Matching

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Motivation

- Matching or selection on observables can help if controlled randomization impossible and natural experiments unavailable
- Underlying conditional independence assumption often criticized

But

- observations selected without reference to the outcome
- and might reduce selection bias

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Conditional Independence Assumption

Conditional Independence Assumption

$$y(1), y(0) \perp D|x$$

x := pretreatment variables

Under the CIA

$$\mathbb{E}[y_i(0)|D_i = 0, x] = \mathbb{E}[y_i(0)|D_i = 1, x] = \mathbb{E}[y_i(0)|x]$$

 $\mathbb{E}[y_i(1)|D_i = 0, x] = \mathbb{E}[y_i(1)|D_i = 1, x] = \mathbb{E}[y_i(1)|x]$

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$$\delta_{x} = ATE$$

= $\mathbb{E}[\Delta_{i}|x]$
= $\mathbb{E}[y_{i}|D_{i} = 1, x] - \mathbb{E}[y_{i}|D_{i} = 0, x]$

$$\tau = ATT$$

$$= \mathbb{E}[\Delta_i | D_i = 1]$$

$$= \mathbb{E}[\mathbb{E}[\Delta_i | D_i = 1, x] | D_i = 1]$$

$$= \mathbb{E}[\mathbb{E}[y_i | D_i = 1, x] - \mathbb{E}[y_i | D_i = 0, x] | D_i = 1]$$

$$= \mathbb{E}[\delta_x | D_i = 1] \quad \text{By LIE}$$

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Estimation Strategy under CIA

Estimation strategy for ATE and ATT under CIA:

- Stratify data by x-values.
- Compute ATE for each stratification.
- Average ATEs in population of treated units.

Essential difference between matching and regression are different weighting schemes.

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Comparison of Matching and Regression

$$\begin{aligned} ATT &= \Delta_M = \mathbb{E}[y_i(1) - y_i(0) | D_i = 1] \\ &= \delta_0 Pr(x_i = 0 | D_i = 1) + \delta_1 Pr(x_i = 1 | D_i = 1) \\ &= \delta_0 \frac{Pr(D_i = 1 | x_i = 0) Pr(x_i = 0)}{Pr(D_i = 1)} \\ &+ \delta_1 \frac{Pr(D_i = 1 | x_i = 1) Pr(x_i = 1)}{Pr(D_i = 1)} \end{aligned}$$

Where weights are proportional to probability of treatment at each covariate value.

$$\delta_0 = \mathbb{E}[y_i(1) - y_i(0)|D_i = 1, x_i = 0]$$

$$\delta_1 = \mathbb{E}[y_i(1) - y_i(0)|D_i = 1, x_i = 1]$$

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Comparison of Matching and Regression

This compares to (fully saturated) regression $y_i = \alpha + \beta x_i + \Delta r D_i + \varepsilon_i$

$$\begin{aligned} \Delta r &= \frac{\mathbb{E}[(D_i - \mathbb{E}[D_i|x_i])y_i]}{\mathbb{E}[(D_i - \mathbb{E}[D_i|x_i)D_i]} \\ &= \delta_0 \frac{Pr(D_i = 1|x_i = 0)[1 - Pr(D_i = 1|x_i = 0)]Pr(x_i = 0)}{\mathbb{E}[Pr(D_i = 1|x_i)[1 - Pr(D_i = 1|x_i)]]} \\ &+ \delta_1 \frac{Pr(D_i = 1|x_i = 1)[1 - Pr(D_i = 1|x_i = 1)]Pr(x_i = 1)}{\mathbb{E}[Pr(D_i = 1|x_i)[1 - Pr(D_i = 1|x_i)]]} \end{aligned}$$

Where weights are proportional to variance of D at each value of covariate \Rightarrow Highest variance of probability of D = 0.5

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Comparison of Matching and Regression

Feasibility of matching and regression estimators?

- ▶ Regression: Curse of dimensionality ⇒ non-saturated model not a solution
- Matching: Problem if in some stratification only treated or only controls
- \Rightarrow Propensity score mitigates dimensionality problem.

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Propensity Score Matching (PSM) The Two PSM Lemmas

$$\rho(x) = \Pr(D = 1|x) = \mathbb{E}[D|x]$$

Lemma 1:

$$D \perp x | \rho(x)$$

Proof:

$$Pr(D = 1|x, \rho(x)) = \mathbb{E}[D|x, \rho(x)]$$
$$= \mathbb{E}[D|x]$$
$$= Pr(D = 1|x)$$
$$= \rho(x)$$

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Propensity Score Matching (PSM) The Two PSM Lemmas

$$Pr(D = 1|\rho(x)) = \mathbb{E}[D|\rho(x)]$$
$$= \mathbb{E}[\mathbb{E}[D|x, \rho(x)]|\rho(x)]$$
$$= \mathbb{E}[\rho(x)|\rho(x)]$$
$$= \rho(x)$$

$$\Rightarrow \Pr(D = 1 | x, \rho(x)) = \Pr(D = 1 | \rho(x)) \blacksquare$$

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Propensity Score Matching (PSM) The Two PSM Lemmas Lemma 2:

$$y(1), y(0) \perp D | x \Rightarrow y(1), y(0) \perp D | \rho(x)$$

Proof:

$$\begin{aligned} \Pr(D = 1|y(1), y(0), \rho(x)) &= \mathbb{E}[D|y(1), y(0), \rho(x)] & \text{The Two PSM} \\ &= \mathbb{E}[\mathbb{E}[D|x, y(1), y(0)]|y(1), y(0), \rho(x)] \\ &= \mathbb{E}[\mathbb{E}[D|x]|y(1), y(0), \rho(x)] & \text{PSM Lemmas} \\ &= \mathbb{E}[\mathbb{E}[D|x]|y(1), y(0), \rho(x)] & \text{PSM Lemmas} \\ &= \mathbb{E}[\rho(x)|y(1), y(0), \rho(x)] = \rho(x) & \text{Stratification of A Stratification A Stratification of A Stratificati$$

 $Pr(D=1|\rho(x)) = \rho(x)$ From Lemma 1

$$\Rightarrow \Pr(D = 1 | y(1), y(0), \rho(x)) = \Pr(D = 1 | \rho(x)) \blacksquare$$

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Propensity Score Matching (PSM) Identification

$$\mathbb{E}[y_i(0)|D_i = 0, \rho(x_i)] = \mathbb{E}[y_i(0)|D_i = 1, \rho(x_i)] = \mathbb{E}[y_i(0)|\rho(x_i)]$$

$$\mathbb{E}[y_i(1)|D_i = 0, \rho(x_i)] = \mathbb{E}[y_i(1)|D_i = 1, \rho(x_i)] = \mathbb{E}[y_i(1)|\rho(x_i)]$$

$$\begin{split} \delta_{\rho(x)} &= \mathbb{E}[\Delta_i | \rho(x_i)] \\ &= \mathbb{E}[y_i(1) - y_i(0) | \rho(x_i)] \\ &= \mathbb{E}[y_i(1) | D_i = 1, \rho(x_i)] - \mathbb{E}[y_i(0) | D_i = 0, \rho(x_i)] \\ &= \mathbb{E}[y_i | D_i = 1, \rho(x)_i] - \mathbb{E}[y_i | D_i = 0, \rho(x_i)] \end{split}$$

$$\begin{aligned} ATT &= \tau = \mathbb{E}[\Delta_i | D_i = 1] \\ &= \mathbb{E}[\mathbb{E}[\Delta_i | D_i = 1, \rho(x_i)]] \\ &= \mathbb{E}[\mathbb{E}[y_i(1) | D_i = 1, \rho(x_i)] - \mathbb{E}[y_i(0) | D_i = 0, \rho(x_i)] | D_i = 1] \\ &= \mathbb{E}[\delta_{\rho(x)} | D_i = 1] \end{aligned}$$

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Propensity Score Matching (PSM) Estimation Strategy

- 1. Estimate $\rho(x)$
- 2. Estimate ATE given $\rho(x)$; actual matching
 - ► Stratification on p(x)
 - Nearest neighbour
 - Radius matching
 - Kernel matching
 - ▶ weighting based on p(x)

 \Rightarrow No dimensionality problem, thanks to Lemma 1.

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Propensity Score Matching (PSM)

PSM Estimation Algorithm

- 1. Start with parsimonious logit or probit.
- 2. Sort data from low to high.
- 3. Sorting all observations into blocks such that treated and controls not statistically different.
- 4. Test whether balancing property holds in all blocks for all covariates.
 - a) Test whether means of all covariates in each block different from zero for treated and controls.
 - b) If one covariate not balanced in ONE block, split the block.
 - c) If one covariate not balanced in ALL blocks, add interactions and/or higher-order terms to logit.

Tips:

- $\rho(x)$ -ranges for treated and controlled ideally the same.
- Frequency of treated and controls in all blocks the same.

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Estimation of ATT by Stratification on the Score

$$\tau_f^s = \frac{\sum_{i \in I(f)} y_i^T}{N_f^T} - \frac{\sum_{j \in I(f)} y_j^C}{N_f^C} \qquad f \text{ is index for strate}$$

$$\tau^{s} = \sum_{f=1}^{Q} \tau_{f}^{s} \frac{\sum_{i \in I(f)} D_{i}}{\sum_{\forall i} D_{i}}$$

$$Var[\tau_{f}^{s}] = \frac{Var[y_{f}^{T}]}{N_{f}^{T}} + \frac{Var[y_{f}^{C}]}{N_{f}^{C}}$$

$$\begin{aligned} & \textit{Var}[\tau^{s}] = \frac{1}{N^{T}} \left[\textit{Var}[y_{i}^{T}] + \sum_{f=1}^{Q} \frac{N_{f}^{T}}{N^{T}} \frac{N_{f}^{T}}{N_{f}^{C}} \textit{Var}[y_{j}^{C}] \right] \\ & \stackrel{\text{from}}{=} \sum_{f=1}^{Q} \left(\frac{N_{f}^{T}}{N^{T}} \right)^{2} \textit{Var}[\tau_{f}^{s}] \end{aligned}$$

• Irrelevant controls for ATT: discard controls with $\rho(x) > \max$ or $< \min$. Penalty for unequal number of treated and controls in a block by increasing variance.

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Nearest Neighbor, Kernel and Radius

Nearest Neighbor, Kernel and Radius

Nearest neighbor closest to ideal, but what if nearest neighbor already used?

- Nearest neighbor or radius matching with replacement.
- Kernel matching: every treated is matched with weighted average of all controls with weights that are inversely proportional to the distance Q between the two.

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Weighting on the Score Lemma 1

Lemma 1: (ATE)

 $y(1), y(0) \perp D|x$

$$\Rightarrow w = \mathbb{E}[y_i(1)] - \mathbb{E}[y_i(0)] \\ = \mathbb{E}[y_i D_i | \rho(x_i)] - \mathbb{E}[y_i(1 - D_i) | (1 - \rho(x_i))]$$

Proof:

$$\mathbb{E}[y_i D_i | \rho(x_i)] - \mathbb{E}[y_i (1 - D_i) | (1 - \rho(x_i))]$$

$$= \mathbb{E}\left[\mathbb{E}\left[\frac{y_i D_i}{\rho(x_i)}|x\right] - \mathbb{E}\left[\frac{y_i (1 - D_i)}{1 - \rho(x_i)}|x\right]\right]$$

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Weighting on the Score

Weighting on the Score Lemma 1

Using definition of $\rho(x)$ and fact that CIA makes conditioning on treatment irrelevant in two internal expectations, we have:

$$\mathbb{E}[\mathbb{E}[y_i(1)|x] - \mathbb{E}[y_i(0)|x]] = \mathbb{E}[y_i(1)] - \mathbb{E}[y_i(0)] \blacksquare$$

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Weighting on the Score

Weighting on the Score Lemma 2

Lemma 2: (ATT)

 $y(1), y(0) \perp D|x$

$$\Rightarrow \tau = \mathbb{E}[y_i(1)|D_i = 1] - \mathbb{E}[y_i(0)|D_i = 1]$$
$$= \mathbb{E}[y_iD_i] - \mathbb{E}\left[y_i(1-D_i)\frac{\rho(x_i)}{1-\rho(x_i)}\right]$$

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Weighting on the Score

Weighting on the Score Lemma 1 **Proof:**

$$\begin{split} \mathsf{E}[\mathsf{y}_i D_i] &- \mathbb{E}\left[y_i(1-D_i)\frac{\rho(\mathsf{x}_i)}{1-\rho(\mathsf{x}_i)}\right] \\ &= \mathsf{E}\left[\mathbb{E}[y_i D_i|\mathsf{x}] - \mathbb{E}\left[y_i(1-D_i)\frac{\rho(\mathsf{x}_i)}{1-\rho(\mathsf{x}_i)}|\mathsf{x}\right]\right] \\ &= \mathsf{E}[\mathbb{E}[y_i(1)|D_i = 1,\mathsf{x}]Pr(D_i = 1|\mathsf{x}) \\ &- \mathsf{E}\left[y_i(0)\frac{\rho(\mathsf{x}_i)}{1-\rho(\mathsf{x}_i)}|D_i = 1,\mathsf{x}\right]Pr(D_i = 0|\mathsf{x}) \end{split}$$

Using the definition of $\rho(x_i)$ and CIA makes conditioning irrelevant,

$$\mathbb{E}[\mathbb{E}[y_i(1)|D_i = 1, x] - \mathbb{E}[y_i(0)|D_i = 1, x]|D_i = 1] \\ = \mathbb{E}[y_i(1)|D_i = 1] - \mathbb{E}[y_i(0)|D_i = 0] \blacksquare$$

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Weighting on the Score

- Problem with Weighting: Sensitive to method p(x) is estimated
- Advantage of Weighting: Does not rely on stratification or matching
- Automated covariate balancing *i* genetic matching and entropy balancing.

 \Rightarrow Overall: Matching generally preferable to OLS

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