

Matching

$$D_i = \begin{cases} 1 & \text{treated} \\ 0 & \text{untreated / control} \end{cases} \quad \text{e.g. college educ (D=1), no college educ (D=0)}$$

$y_L(1)$ potential outcome of L with treatment

$y_L(0)$ " " " without "

$$\Delta_L = y_L(1) - y_L(0) = \text{treatment effect for obs } L$$

↓
never observed

$$\Delta^{ATE} = E [y(1) - y(0)] \quad \text{average treatment effect (ATE)}$$

Observed for L y_L, D_L, x_L

$$y_L = D_L \gamma_L(1) + (1 - D_L) \gamma_L(0) = \text{observed outcome for } L$$

Assumptions to estimate $\Delta^{A:F}$

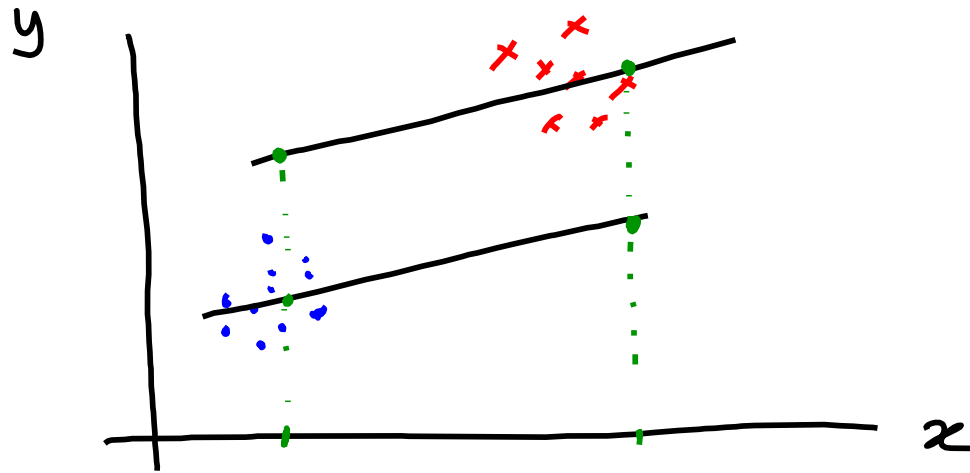
$$\gamma(0), \gamma(1) \perp D \mid x$$

$$0 < P(D=1 \mid x) < 1$$

Random sample $\{y, D, x\}$

Matching \rightarrow when comparing treated and control obs, more weight given to similar obs

No functional form ass^ums (such as linear regression)



$$\hat{\Delta}^{ATE} = \frac{1}{N} \sum [\gamma_i(1) - \gamma_i(0)]$$

↳ infeasible → missing data / counterfactual

$\gamma_i(0)$ observed if $D_i = 0$; $\gamma_i(1)$ unobserved if $D_i = 1$

Replace $\gamma_i(0)$ with a close match to $\gamma_i(1)$

but with $D = 0$ (e.g. $\gamma_d(0)$ where $D_d = 0$)

Various matching approaches to determine what weight to attach to obs from the opposite group

One approach relies on the propensity score, $p(x)$

$$p(x) = P_r(D=1 | x)$$

↳ typically estimated using logit/probit

$$Y(0), Y(1) \perp D | x \Rightarrow Y(0), Y(1) \perp D | p(x)$$

Single nearest neighbor matching $y_i(1)$

If $D_i = 1$ & $y_i(0)$ is unobserved use $y_d(0) = y_i$

$D_i = 0$ & $y_i(1)$ where $D_d = 0$ & p_d is $D_d = 1$
closest to p_i