

Propensity scores

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- propensity score = prob. of receiving treatment, rather than control, given covariates X
- $A=1$... treated & $A=0$... control
- denote propensity score of subject i as π_i :

$$\pi_i = P(A=1 | X_i)$$

- π_i is a function of X_i (i.e. prop. score = $f(X)$)

Example:

- if older people more likely receive treatment, then
 - o $P(A=1 | \text{age}=60) > P(A=1 | \text{age}=30)$
 - o $\pi_i > \pi_j$ if $\text{age}_i > \text{age}_j$

\Rightarrow prob. of receiving treatment is 30% if subject i has prop. score of 0.3.

Balancing scores:

- occurs when 2 subjects have the same propensity score, but they possibly have different covariate values X .
- \rightarrow both subjects are as likely to be treated
- if we restrict our data to a subpopulation of subjects w/ same value of prop. score, then there should be balance in the 2 treatment groups

\Rightarrow the propensity score is thus a balancing score

\hookrightarrow if we were to condition on the propensity score, then we will have balance btr. both groups

Example:

- (1) Consider only subjects with the same prop. score value (e.g. 0.45), then:
- (2) if we stratify that subpopulation of subjects by the factual / actual treatment received, then:
- (3) we should see the same dist. of covariates X for the treatment groups.

Formally Balancing Scores would mean:

\hookrightarrow dist. of the covariates X

$$P(X=x | \pi(X)=p, A=1) = P(X=x | \pi(X)=p, A=0)$$

Implication:

When matching on the propensity score, we should achieve balance btr. the groups $A=0$ & $A=1$.

Estimated propensity score:

- in a RT, the prop. score is known (e.g. $P(A=1)=P(A=0)=0.5$)
- \hookrightarrow in an observational study, prop. score will be unknown
- however, prop. score just involves observed data: A and X , thus we can estimate it!

\rightarrow we need to estimate $P(A=1 | X)$

\hookrightarrow the outcome here is A , a binary variable.

\rightarrow use e.g. logistic regression (LR) to estimate prop. scores

1. Fit LR model with outcome A , covariates X
2. Use fitted model to get the predicted probability (fitted values) for each subject

\Rightarrow these values are the "estimated" propensity scores.