Discussion of Papers on the Extensions of Propensity Score

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August 3, 2010

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Generalized Propensity Score

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The Theme and Papers of this Panel

- Theme: Extensions of the Propensity Score Method
- Wang: Extend propensity score to continuous treatment regimes
- Huang: Extend propensity score to mismeasured covariates
- Hong: Extend propensity score to causal mediation analysis
- Hansen: Provide a theoretical justification for some propensity score matching methods

The Wang Paper

- Extend propensity score to continuous treatment regimes
- Imai and van Dyk (JASA, 2004)
 - Parametric model for conditional probability of treatment: $p_{\psi}(A \mid X)$
 - Generalized propensity function: $e(\cdot \mid X) = p_{\psi}(\cdot \mid X)$
 - Suppose θ uniquely represents $e(\cdot \mid \theta_{\psi}(X))$
 - $e(\cdot \mid X)$ depends on X only through $\theta_{\psi}(X)$
- Two main theoretical results:
 - Propensity function as balancing score
 - Ignorable treatment assignment given propensity function
- Practical implications:
 - Causal effects estimation:
 - **(1)** Estimate the propensity function $p_{\psi}(A \mid X)$
 - 2 Subclassify on $\hat{\theta}(X)$
 - Estimate causal effects within each subclass and aggregate
 - Model diagnostics: check independence between A and X after conditioning on
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The Wang Paper (continued)

- Main theoretical results:
 - Depending on the outcome model, one can further reduce the dimension of the propensity function
 - If the interactions between treatment and some covariates exist in the outcome model, these covariates need to be adjusted in addition to the propensity function
- Two main advantages of the propensity score method:
 - Robustness: when the knowledge of outcome model is lacking
 - Diagnostics: when the knowledge of propensity model is lacking
- What are the practical implications of these two theoretical results in light of these advantages?
 - How do these results help analysts if they do not possess the knowledge of outcome model?
 - 2 Do these results suggest new diagnostics about propensity or outcome model specification?
 - Models with interactions: Don't we know the propensity model is incorrect if covariates aren't balanced?

The Huang Paper

- Use of propensity score with mismeasured covariates
- Some existing works on mismeasured treatments (e.g., Lewbel; Imai and Yamamoto) but little is done on mismeasured covariates
- Non-differential measurement error: conditionally independent of potential outcomes given (true) covariates
- In addition, measurement error is assumed to be independent of treatment status
- These are reasonable assumptions
- Nonparametric identification analysis in a simple situation (e.g., binary treatment, covariate, and outcome) may be illuminating

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The Huang Paper (continued)

- Theorem 2: true propensity score balances true covariates and measurement error as well as mismeasured covariates
- Identification via the restriction on propensity model
- How can we diagnose propensity model specification? Is balancing mismeasured covariates sufficient?
- Finite mixture model that combines outcome model, propensity model, measurement error model
- This is nice but how does one conduct diagnostics within this approach?
- Are there additional advantages for simultaneously modeling propensity score and exposure effect?

The Hong Paper

- Use of propensity score in causal mediation analysis
- Exploration of causal mechanisms require the estimation of natural direct and indirect effects
- Under standard designs, the mediator is not randomized
- Under sequential ignorability, natural direct/indirect effects are *nonparametrically* identified (Imai *et al.* Stat. Sci. 2010):

$$\{ Y_i(t', m), M_i(t) \} \quad \bot \quad T_i \mid X_i = x,$$

$$Y_i(t', m) \quad \bot \quad M_i(t) \mid T_i = t, X_i = x,$$

- Nonlinear structural estimation: Imai *et al.* (Psy. Meth. in-press) and Pearl (Working paper, 2010)
- Marginal structural estimation: VanderWeele (Epidemiology, 2009)

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The Hong Paper (continued)

- A new approach based on propensity score weighting that can handle post-treatment confounders
- Treatment-by-mediator interaction effects are also handled but this is not a problem for existing methods so long as post-treatment confounders do not exist
- The outcome model is nonparametric
- Robins' no-interaction effect assumption in the presence of post-treatment confounder is difficult to justify
- What are the key identifying assumptions in this paper? And how they should be interpreted by substantive researchers?

$$\begin{array}{cccc} Y_i(t,m) & \bot & M_i(t) \mid T_i = t, X_i = x, L_i(t) = I \\ Y_i(t,m) & \bot & M_i(t') \mid T_i = t, X_i = x, L_i(t) = I \\ L_i(t) & \bot & M_i(t') \mid T_i = t, X_i = x \end{array}$$

• Need for empirical and simulation examples: what happens if mediator is continuous and/or has skewed distributions?

The Hansen Paper

- Novel analytical results: even if one cannot match exactly on propensity score, in a large sample correct inference can be made so long as covariate balance is "good enough"
- Formal definitions of informally used concepts; adjustability, crude balance
- Formal results showing the conditions under which matching can be justified
- What are the implications for practice?
 - Conduct balance test and then estimate causal effects?
 - Can balance tests be used to diagnose the misspecification of propensity score model? If so, how? The possibility of multiple testing?
 - If two different matching methods give the "same" result in terms of balance tests, which one should one choose?

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The Hansen Paper (continued)

- A different perspective for the purpose of discussion (Imai et al. *JRSSA*; Ho et. al. *Pol. Anal.*)
- "balance tests" are often conducted on different matched samples to diagnose the misspecification of propensity model
- multiple testing, different sample size can be problematic
- balance should be maximized without limit for better inference
- the gold standard is the experiment with matched-pair design rather than the experiment with simple randomization
- matching with pre-determined balance
 - matching with fine balance
 - coarsened exact matching
 - maximum entropy matching
- matching as nonparametric preprocessing for making parametric inference robust

- A great set of papers extending the propensity score methods to various situations of practical importance
- Common challenges:
 - Development of diagnostics for propensity model specification
 - Connecting interesting theoretical results to practice
- Look forward to seeing future versions of the papers

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