A Review of Methods for Propensity Score Matched Subgroup Analysis and their Application in Peer Reviewed Research Studies



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BACKGROUND

- The United States Food and Drug Administration's (FDA) Sentinel Program is a national system for medical product monitoring involving a national distributed network of healthcare databases and a suite of routine querying tools
- □ The FDA is often interested in examining treatment effect in pre-specified subgroups.
- Trade offs of different methods using propensity score (PS) matching for subgroup analyses is not well understood

OBJECTIVE

To summarize methods papers comparing the performance of different methods to conduct PS matched subgroup analyses.

□ To describe how often different methods for PS matching in subgroup analyses are implemented in practice.

METHODS

- U We conducted separate searches for our systematic literature review of methodology and applied research papers
- □ For methodology papers, we searched PubMed for:
 - "subgroup", "effect modification", "moderation analysis", or "treatment heterogeneity" co-occurred with "propensity score" in titles or abstracts without restricting publication date.
- □ For applied research papers, we searched PubMed for:
 - "propensity score match*" and "subgroup analysis*" in any field without restriction by publication date; or published in Pharmacoepidemiology and Drug Safety (PDS) between July 1, 2015 and June 30, 2016 with "propensity score match*" in any field.

RESULTS

Methodology papers

- □ We identified 5 relevant methods papers after reviewing abstracts of 606 papers from the PubMed search (Table 1)
- □ These reported small improvements in covariate balance and bias with use of a subgroup-specific PS instead of an overall PS when the true PS

Applied papers

- 83 of 129 papers from the PubMed search met inclusion criteria after abstract review (cohort study, PS matching for main and subgroup analysis, English PDF available)
- Applied papers frequently used PS for subgroup analysis in ways not
- varied by subgroup.
- Methods papers only compared strategies that involved re-matching on PS within subgroups.
- Table 1. Methodology papers comparing performance of different ways to use propensity score matching for subgroup analyses

Paper	Data	# Subgroups	Performance	PS methods used in subgroups
Green KM, et al. 2014	Empirical	1	Balance (Abs. Std. Diff)	
Girman CJ, et al. 2014	Empirical	1	Balance (Mean Abs. Std. Diff)	1:1 greedy matching Decile adjustment
Rassen JA, et al. 2012 ¹	Empirical, Simulation	3, 1	Balance (Mahalanobis distance) Bias (difference in estimates)	1:1 nearest neighbormatchingDecile adjustmentDirect adjustment for PS
Kreif N, et al. 2012 ²	Empirical, Simulation	1, 1	Balance (weighted Std. Diff) Bias (RMSE)	1:1 Nearest neighbormatchingGenetic matchingIPTW
Radice R, et	Empirical,	1, 1	Balance (weighted	1:1 Nearest neighbor

evaluated in methods papers (Table 2)

N = number of applied research papers by strategy type

- 33% used PS to match in the overall cohort, then split the 1:1 matched cohort into subgroups without further adjustment
- 25% provided insufficient detail to clearly determine how PS matching was implemented for subgroup analysis

Table 2. Applied research papers using propensity score matching methods for subgroup analysis

Strategy	Ν	Proportion
Use overall PS to match for main analysis and do matched	14	0.17
analysis in subgroups		
Use subgroup specific PS for subgroup analysis	6	0.07
Use overall PS to match within subgroups, aggregate for main	1	0.01
analysis		
Use overall PS to match for main analysis, within matched	27	0.33
cohort, split into subgroups and do analysis without further		
matching		
Use overall PS to match for main analysis, within matched	14	0.17
cohort, split into subgroups and use multivariable or other		
adjustment in subgroups		
Unclear (lack sufficient details in publication)	21	0.25
Total	83	1.00

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al. 2012	Simulation		Std. Diff)		matching
			Bias (RMSE)		Genetic matching, IPTW

¹ Results from simulations directly adjusting for PS, no simulation results available for matching on PS
² Results from studies of cost-effectiveness, continuous outcome - cost and quality adjusted life years (QALY)
Std. Diff = standardized difference; Abs = absolute; IPTW = inverse probability of treatment weighting; PS = propensity score; RMSE = root mean squared error

CONCLUSIONS

The performance of several alternative methods for using a PS to match for subgroup analyses have been evaluated in the methods literature, however the evaluated methods do not include evaluation of the methods for PS matched subgroup analyses most commonly used in applied studies.

Further evaluation is needed to understand the relative performance of strategies for PS matching in subgroup analyses, particularly within settings with low exposure, infrequent outcomes and multiple subgroups of interest.

CONFLICTS OF INTERST

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