PSAboot: An R Package for Bootstrapping Propensity Score Analysis

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Propensity score analysis [5] is an approach to estimate causal effects in observational studies where randomization is not used. However, even when balance has been achieved between treatment and control units on observed covariates, there is a risk of bias due to unobserved covariates. Rosenbaum[4] has suggested testing one hypothesis multiple times using multiple matching and/or stratification methods to avoid exaggerated bias due to method selection. The **PSAboot** extends Rosenbaum's suggestion by implementing bootstrapping [1] for PSA. Like typical bootstrapping methods, the **PSAboot** package will draw *n* random samples and provide a pooled effect size estimate and confidence intervals. However, for each bootstrap sample multiple PSA methods will be used. The package includes implementations for matching (using the **Matching**[6] and **MatchIt**[2] packages), stratification with logistic regression, and stratification using classification trees (using the **rpart**[7] and **party**[3] packages). The package emphasizes the use of visualizations for evaluating balance as well as summarizing results. Additionally, examples on extending the package for other PSA methods will be discussed.

References

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