

Thesis Proposal

Methodology and Statistics for the Behavioural, Biomedical and Social Sciences

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2019-09-27

ShinyMICE: an Evaluation Suite for Multiple Imputation

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Word count: 775

Introduction

At some point, any (social) scientist conducting statistical analyses will run into a missing data problem (Allison 2001). Simply ignoring the missingness or using ad hoc solutions can yield wildly invalid inferences (Van Buuren 2018). A framework to circumvent the *ubiquitous* problem of missing information is multiple imputation (MI; (Rubin 1987)). This technique – that is growing in popularity (Van Buuren 2018) – entails ‘guessing’ the missing values in an incomplete data set several times. The variability between the completed data sets represents how much uncertainty in the inferences is due to missingness (Rubin 1987).

With MI, many assumptions are made about the nature of the observed and missing parts of the data, and their relation to the ‘true’ data generating model. Without proper evaluation of the imputations and the underlying assumptions, any drawn inference may erroneously be deemed valid (**source?**). Such evaluation measures are currently missing or under-developed in MI software. Therefore, I aim to answer the following question: ‘Which measures are vital for evaluating the validity of multiply imputed data?’.

The goal is to develop an MI evaluation suite for the world leading R package **MICE** (Van Buuren and Groothuis-Oudshoorn 2011). It will feature novel assessment tools (e.g., a measure to flag algorithmic non-convergence), and interactive adaptations of (partially) implemented modules (e.g., plots to compare the incomplete and completed data sets). This research project will aid applied researchers in drawing valid inference from incomplete data sets. Simultaneously, a contribution to the scientific literature is made by developing novel methodology and guidelines for evaluating MI data methods.

Literature Review

The numerous assumptions underlying MI algorithms can principally not be verified from the incomplete observed data (**source?**). Instead, imputers are designated to evaluate the (plausibility of) multiply imputed data, and the robustness of results to the assumptions. The latter can be assessed by performing sensitivity analyses, see e.g. Nguyen, Carlin, and Lee (2017) for practical guidelines. Methodology for evaluating the MI data however, is still largely missing. In the only available review of diagnostics, Abayomi, Gelman, and Levy (2008) overlook the most vital state to be evaluated: convergence of the algorithm. Without convergence, any ‘deeper’ assumption and resulting inference is invalid (**source?**).

While the convergence properties of iterative MI algorithms are still under debate (Takahashi 2017), specific procedures like *predictive mean matching* pose entirely open questions (Murray 2018). Van Buuren (2018) summarizes the issues with diagnosing convergence as follows: “No method works best in all circumstances. The consensus is to assess convergence with a combination of tools. The added value of using a combination of convergence diagnostics for missing data imputation has not yet been systematically studied” (Van Buuren (2018), §6.5). Meanwhile, applied researchers have to rely on visual inspection of the algorithm’s iterations (Van Buuren 2018). Convergence is said to be reached when parameters (e.g., means of completed variables) are stable across iterations (White, Royston, and Wood 2011). And additionally, the variation between imputation chains should be no larger than the variation within each individual chain (Van Buuren 2018). Conceptually, the visual inspection procedure is equivalent to Gelman, & Rubin’s (1992) convergence statistic \hat{R} (Li et al. 2014). In practice, however, \hat{R} cannot be applied directly to MI data, and is prone to producing false negatives (Lacerda, Ardington, and Leibbrandt 2007). The limitations in assessing convergence of MI algorithms may be overcome by combining \hat{R} with other diagnostics like auto-correlation and MC error.

Add: which other assumptions could be checked? The assumption of MAR. Ideally, we would want to study all possible combinations of variables: univariate, bivariate, etc. And to include both plots and stats.

Methods

This research project will be supervised by the **MICE** developers. I will develop novel methodology for evaluating MI data, and implement these methods in an interactive evaluation device in R Shiny (Chang et

al. 2019): ‘ShinyMICE’. This approach does not require the use of any empirical data. Therefore, the project is not subject to the school’s ethical approval process.

The research report will consist of an investigation into algorithmic convergence of MI algorithms. Ideally, this will result in a single summary indicator to flag non-convergence.

In the second stage of the research project I will focus on other evaluation measures to implement in ShinyMICE. The application will at least consist of the following: 1) one or more measures to assess algorithmic convergence; 2) data visualizations (e.g., scatter-plots, densities, and cross-tabulations of the data pre- and post-imputation); and 3) statistical evaluation of relations between variables pre- versus post-imputation (i.e., χ^2 -tests or t-tests).

A working beta version of ShinyMICE will be considered a sufficient milestone to proceed with writing a technical paper on the methodology and the software. I aim to submit this for publication in *Journal of Statistical Software*. Finally, ShinyMICE will be integrated into the existing MICE environment, and a vignette for applied researchers will be written.

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