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Nested multiple imputation in large-scale assessments

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Abstract

Background: In order to measure the proficiency of person populations in various domains, large-scale assessments often use marginal maximum likelihood IRT models where person proficiency is modelled as a random variable. Thus, the model does not provide proficiency estimates for any single person. A popular approach to derive these proficiency estimates is the multiple imputation of plausible values (PV) to enable subsequent analyses on complete data sets. The main drawback is that all variables that are to be analyzed later have to be included in the imputation model to allow the distribution of plausible values to be conditional on these variables. These background variables (e.g., sex, age) have to be fully observed which is highly unlikely in practice. In several current large-scale assessment programs missing observations on background variables are dummy coded, and subsequently, dummy codes are used additionally in the PV imputation model. However, this approach is only appropriate for small proportions of missing data. Otherwise the resulting population scores may be biased.

Methods: Alternatively, single imputation or multiple imputation methods can be used to account for missing values on background variables. With both imputation methods, the result is a two-step procedure in which the PV imputation is nested within the background variable imputation. In the *single+multiple-imputation* (SMI), each missing value on background variables is replaced by one value. In the *multiple+multiple-imputation* (MMI), each missing value is replaced by a set of imputed values. MMI is expected to outperform SMI as SMI ignores the uncertainty due to missing values in the background data.

Results: In a simulation study, both methods yielded unbiased population estimates under most conditions. Still, the recovery proportion was slightly higher for the MMI method.

Conclusions: The advantages of the MMI method are apparent for fairly high proportions of missing values in combination with fairly high dependency between the latent trait and the probability of missing data on background variables.

Keywords: Large-scale assessment; Missing data; Imputation; Simulation; Item response theory

Background

Several large-scale assessment programs such as the National Assessment of Educational Progress (NAEP), the Trends in International Mathematics and Science Study (TIMSS), and the Programme for International Student Assessment (PISA) employ item response theory (IRT) methods to measure the achievement of populations of examinees in various domains, for example, reading and science, for the purpose of

system monitoring. The central interest is not to obtain individual proficiency scores but to obtain population and subpopulation estimates of person proficiency, denoted by θ . In addition to the item responses, the estimation process includes background information for each person (often collected via questionnaires) comprising a multitude of variables, such as demographic, social, or motivational variables. Because these variables may be associated with θ , they are of particular importance in the estimation of subpopulation differences. Large-scale assessment programs are often confronted with the problem of missing values in item responses or in background data. Planned missing data in the item responses are a direct consequence of multiple matrix sampling designs (Frey et al. 2009; Gonzalez & Rutkowski 2010). Methods of item and person parameter estimation which incorporate missing data in the item responses have long been developed and have been proved to be suitable (Lord 1974, 1983; Mislevy et al. 1992). By contrast, a considerable amount of missing data in background variables may lead to biased estimates of population parameters (Rutkowski 2011). Therefore, the present article examines the problem of missing data in background information when population estimates are of interest. We compared two methods that are based on imputation methods with regard to the bias and root mean square error (RMSE) of population parameters. The results support the implementation of multiple imputation methods in the treatment of missing data in background information.

Item and person parameter estimation

In large-scale assessments, the estimation of item and person parameters commonly uses the marginal maximum likelihood (MML) method (Embretson & Reise 2000; Tuerlinckx et al. 2004; Wilson & De Boeck 2004). Conceptually, the estimation process is composed of two parts: the *item response model*, which results from an analysis of the response patterns, and the *population model*, which results from incorporating background information (Adams & Wu 2007). von Davier et al. (2009) refer to these two parts as the likelihood function and the prior distribution function. A model that comprises both parts is called a *conditioning model* or combined model (Adams & Wu 2007).

In the item response model, the categorical response patterns to a set of test items are modeled as the dependent variable in a logistic regression with item difficulties and person proficiencies as independent variables. Adams and Wu (2007) and Adams et al. (1997) described the model in its generalized form. For the purpose of illustration, we will use the special case of a unidimensional model with dichotomous items (Rasch model):

$$\text{logit}(P(X_{ni} = 1)) = \theta_n - b_i, \quad (1)$$

where X_{ni} represents the observed response of a person n for an item i , θ_n represents the proficiency score of person n , and b_i represents the difficulty of item i . The definition of the model in the MML formulation needs to specify a density for the latent variable θ (Adams & Wu 2007). The default assumption is that persons are sampled from a normal population with mean μ and variance σ^2 . Therefore, θ is a random effect, and the population model is: $\theta_n = \mu + E_n$, with $E_n \sim N(0, \sigma^2)$. If we assume that θ depends

on background variables, the population model may be augmented to a regression model with several predictors (Adams & Wu 2007):

$$\theta_n = \mathbf{Y}_n \boldsymbol{\beta} + E_n \quad (2)$$

where \mathbf{Y}_n is a vector of u fixed and known values for a set of u variables related to each person, and $\boldsymbol{\beta}$ is the corresponding vector of regression coefficients to be estimated. E_n is assumed to be independently and identically normally distributed with mean zero and variance σ^2 ; that is, $E_n \sim N(0, \sigma^2)$. The population model now incorporates fixed and random effects and provides *a priori* information about the expected person distribution given all available background information (von Davier et al. 2009).

The conditioning model is the item response model multiplied by the population model (Adams & Wu 2007; Embretson & Reise 2000). Hence, the conditioning model captures information from the test items (expressed in the item response model) and background information (expressed in the population model). Hence, it offers the opportunity to specify a population distribution of persons consisting of several sub-populations. This is especially important if the population distribution is conditional on several demographic variables rather than being a simple (unconditional) normal distribution.

Types of missing data and their effects on the estimation process

In large-scale assessments, three types of missing data can be distinguished. The first one is rather fundamental: IRT models are usually intended to measure proficiencies of persons. As each person's individual proficiency θ_n is considered to be a latent construct, θ is an inherently unobserved (i.e., missing) variable. Missing data in the individual response pattern (the 0/1 answers of persons to test items) constitute the second type, and missing data in background information constitute the third type of missing data.

First type of missing data: inherently missing θ

The inherently missing person parameter θ_n is not a problem in MML estimation. The MML formulation of the Rasch model may be described as a random person–fixed item model (De Boeck 2008) and does not parametrize individual θ_n parameters. However, a common practice in most large-scale assessment programs is to provide individual θ_n values for subsequent analyses (e.g., ordinary least squares regression, contingency table analyses). θ_n estimates therefore have to be generated in an additional step, using the parameters of the fitted model. The most suitable method to generate θ_n estimates is plausible value imputation (Mislevy et al. 1992; von Davier et al. 2009; Wu 2005). This method is based on multiple imputation (Rubin 1987): The conditioning model defines how the observed item responses are related to θ , and the plausible-values approach provides a model-based imputation of missing θ_n values. Mislevy et al. (1992) pointed out that plausible values provide synthetic data produced by the model. Analyzing plausible values with standard techniques yields the same results as if we actually observed true person proficiencies (Rubin 1987). This does not necessarily hold for other estimates of individual θ_n values, for example Weighted Maximum Likelihood Estimates (WLEs; Warm 1989) or the expected a posteriori distribution (EAP; Bock & Aitkin 1981). Moreover, the *multiple* imputation of plausible

values allows investigators to adequately quantify the uncertainty in the parameter estimation of subsequent analyses.

The imputation of plausible values can be conditioned on background information by including some demographic variables when specifying the population model. Similar to the recommendations made for multiple imputation (Rubin 1987), it is crucial to include all variables that may be used in subsequent analyses in order to ensure unbiased estimation of their effects on achievement (Mislevy et al. 1992; von Davier et al. 2009).

Second type of missing data: missing data in the item responses

The second (and also the third) type of missing data concerns missing values in data that are expected to be observed. However, the second type of missing data likewise is not a problem for parameter estimation, if some conditions are met. Missing data in the item responses are a direct consequence of multiple matrix sampling designs (Frey et al., 2009; Gonzalez & Rutkowski 2010). As large samples of items are used to comprehensively cover the test constructs, only a subset of items is presented to each person to keep the individual workload within acceptable boundaries. Such missing responses which result from giving samples of items to samples of persons are often called *missing by design* (Enders 2010). If the test booklets are randomly distributed to the persons, these data are assumed to be missing completely at random (MCAR). MCAR means that missing values on a variable neither depend on values of other variables in the model nor on values of the variable itself (Graham 2009; Rubin 1987). By contrast, responses that a person has chosen to omit are often treated as *incorrect* responses. See Lord (1974, 1983) and Ludlow and O'leary (1999) for a discussion of this procedure.

Third type of missing data: missing data in background information

The third type of missing data—which is the focus of the present article—concerns missing values on background variables which are inherently due neither to the latent variable modeling of θ nor to the test design. In the population model (Equation 2), it is assumed that background variables are measured without error because \mathbf{Y}_n is a vector of fixed and known values (Adams & Wu 2007). In principle, the same problem applies when missing values occur in ordinary least squares regression models (e.g., Little 1992): The corresponding procedures require that all variables be fully observed.

What might be the consequences of missing values in background data? In theory, two possible effects would be conceivable: First, the parameter estimates in Equation 2 might be biased which may, in turn, result in biased estimates of subpopulation differences. Thus, the effect of missing data on the population model is comparable to the effect of missing data on ordinary least-squares regression models. Second, the uncertainty of the corresponding mean estimates might not be adequately represented which may result in biased standard errors of the subpopulation mean estimates. The extent to which these biases occur is presumably influenced by the overall proportion of missing values and by whether the probability of missing values depends on further variables, even on θ itself. The model used to estimate θ would have to take these potential dependencies into account.

Approaches to dealing with missing data in background information

Dummy coding

In several current large-scale assessment programs (Allen et al. 2001; Foy et al. 2008; OECD 2009) missing observations were dummy coded and the dummy variables were subsequently used in the population model. However, Schafer and Graham (2002) pointed out that this method merely redefines the parameters as the population model now contains two effects for a variable X in the population model: one effect for the respondents and another effect for the nonrespondents. Moreover, this approach does not take a possible missing at random (MAR) mechanism into account. MAR means that missing values on a certain variable may depend on values of other variables in the model but not on values of the variable itself. Allison (2002) demonstrated that even in the case of MCAR, dummy-variable adjustment may lead to biased estimates in ordinary least squares regression when the proportion of missing data is substantial (i.e., 50%).

Rutkowski (2011) carried out a simulation study to explore the impact of missing values on background variables on proficiency estimates when missing values on background variables were dummy coded, focusing on the estimation of differences in group means. Under an MAR condition with a maximum of 20% of missing values, the *differences* in group means were estimated without bias. However, both group means were biased in the same direction. Rutkowski termed this a *tandem shift*. Although this tandem shift preserved subgroup differences, it may lead to erroneous conclusions if the results are to be expressed on a predefined scale, for example, to estimate trend effects between different cohorts (Mazzeo & Von Davier 2008). Rutkowski (2011) pointed out that the estimation of subgroup differences can be problematic if the proportions of missing values on the background variables differ between groups. In summary, dummy coding may be appropriate if it is combined with assessment strategies that minimize the proportion of missing values in background information. If this is not the case, defining dummy codes for missing observations may be questionable from a theoretical point of view (Schafer & Graham 2002) and may also lead to biased mean estimates (Rutkowski, 2011). Hence, alternative procedures have to be applied.

Single + Multiple Imputation (SMI)

Instead of using dummy codes, missing values on background data may be imputed prior to estimating the parameters of the conditioning model (OECD 2006; Weirich et al. 2012). In order to impute the missing data, we have to construct an imputation model which includes all variables that may be related to the occurrence of missing data under an MAR assumption. Thus, the population model used to generate the plausible values includes imputed data on the background variables.

The SMI procedure contains a sequence of four steps. The first step is the specification of the item response model (often referred to as calibration). As the aim is to estimate the item parameters only, a simple MML population model without background information can be chosen as $\theta_n = \mu + E_n$, with $E_n \sim N(0, \sigma^2)$. The mean of the population distribution is often defined to be zero, i.e., $\mu = 0$. Hence, to calibrate items, missing values in background variables can be ignored.

The second step is the single imputation of missing values in background variables which requires an imputation model. If we assume that missing values might depend

on θ , we should include θ into the imputation model. However, individual θ_n values are estimated in a later step, therefore we have to use a proxy for θ , for example, the percentage of correct responses or point estimates (e.g., MLEs or WLEs) for each person from the item response model. In this study, we will use the latter one.

The third step is the estimation of the conditioning model's parameters. We jointly use the item responses and the (imputed) background information for estimation. In contrast to the item response model, $\theta_n = \mathbf{Y}_n \boldsymbol{\beta} + E_n$. \mathbf{Y}_n does not contain any missing values because all missing values were imputed in the third step. All background variables are treated as if they were fully observed. A common practice is to consider the item parameters obtained in the first step as the true item parameters (von Davier et al. 2007).

The fourth and last step is the imputation of plausible values. The imputation model for this step is the conditioning model. Hence, the process of handling different sources of missing data comprises *two* imputation procedures: a single imputation of background variables and a multiple imputation of latent person estimates (θ_n). To avoid confusion between both, we henceforth refer to the first one (step 2) as the imputation model and to the second one (step 4) as the conditioning model. Both imputation steps are not independent because the imputed data from step 2 are used in step 4.

The SMI procedure should help to minimize bias in the population model when missing values on background variables depend on θ . However, there is an uncertainty in the prediction of θ in the population model due to missing values on background information. The SMI procedure may not adequately represent this uncertainty because the population model treats observed and imputed values in the same way. For example, consider two groups with different proportions of missing values on background variables: The uncertainty in the group mean estimates should differ between the groups because we have more observed data in one group than in the other. However, this is not represented in the model if only a single imputation method is used for both groups. Although a bias in estimated group mean differences is unlikely, if the imputation model adequately captures the mechanism behind the missing data, the corresponding standard errors of the mean estimates may be underestimated (White et al. 2010), in particular if the rates of missing data in background variables are high.

Multiple + Multiple Imputation (MMI)

Nested imputation (Rubin 2003) or two-stage multiple imputation (Harel 2007; Harel & Schafer 2003; Reiter & Drechsler 2007; Reiter & Raghunathan 2007) explicitly manages two multiple imputation procedures in a dependent structure. The basic principle is described by Rubin (2003):

A few imputations of the first part are created (say M), and then for each of these, several imputations of the second part are created (say N). The standard combining rules for multiple imputation (the repeated imputation rules) have to be modified because the imputations within a nest are correlated, since they share a common set of imputed values for the first set of missing values. (p. 6)

The nested imputation also allows for different missing value mechanisms, for example, MCAR and MAR (Harel & Schafer 2003). In the context of large-scale

assessments, this may be of particular interest as we may assume different mechanisms in order to account for missing values on θ and missing values on background variables. Furthermore, MMI is preferable to SMI because MMI captures both sources of uncertainty: the uncertainty due to missing values on the background variables and the uncertainty due to the inherently missing θ values.

To implement this method in a large-scale assessment context, we would have to extend the SMI procedure described in the previous section. In the second step, we now create more than one imputation of the background data (which would result in, say, $M = 5$ data sets). Step 3 therefore has to be repeated M times, once for each of the M imputed data sets. In step 4, we would draw N plausible values from each of the M fitted conditioning models, which will result in $M \times N$ plausible values overall. We would then apply the modified combining rules by Rubin (2003) to pool statistics based on these plausible values. A more detailed illustration of how the following combining rules are applied to the sets of plausible values is provided in Additional file 1.

Let Q be the quantity of interest (i.e., our set of plausible values). $\hat{Q}^{(m,n)}$ is the mean estimate of the n th plausible value in the m th nest. The overall average \bar{Q} , then, is:

$$\bar{Q} = \frac{1}{MN} \sum_{m=1}^M \sum_{n=1}^N \hat{Q}^{(m,n)}. \quad (3)$$

\bar{Q}_m is a vector of M mean estimates across the N plausible values in each nest

$$\bar{Q}_m = \frac{1}{N} \sum_{n=1}^N \hat{Q}^{(m,n)}, \quad (4)$$

and \bar{U} is the overall average of the associated variance estimates,

$$\bar{U} = \frac{1}{MN} \sum_{m=1}^M \sum_{n=1}^N U^{(m,n)}. \quad (5)$$

Let $MS^{(b)}$ be the between-nest mean square,

$$MS^{(b)} = \frac{N}{M-1} \sum_{m=1}^M (\bar{Q}_m - \bar{Q})^2, \quad (6)$$

and let $MS^{(\omega)}$ be the within-nest mean square,

$$MS^{(\omega)} = \frac{1}{M(N-1)} \sum_{m=1}^M \sum_{n=1}^N (\bar{Q}^{(m,n)} - \bar{Q}_m)^2. \quad (7)$$

The quantity

$$T = \bar{U} + \frac{1}{N} \left(1 + \frac{1}{M} \right) MS^{(b)} + \left(1 - \frac{1}{N} \right) MS^{(\omega)} \quad (8)$$

estimates the total variance of $(Q - \bar{Q})$. Estimation of the significance levels of Q are based on a Student- t reference distribution, $T^{-0.5}(Q - \bar{Q}) \sim t_\nu$. The degrees of freedom, ν , follow from

$$v^{-1} = \left(\frac{N^{-1}(1 + M^{-1})MS^{(b)}}{T} \right)^2 \frac{1}{M-1} + \left(\frac{(1-N^{-1})MS^{(\omega)}}{T} \right)^2 \frac{1}{M(N-1)}. \quad (9)$$

This procedure is computationally more extensive than the commonly applied dummy coding procedure and, to our knowledge, has not been used in any large-scale assessment program yet. However, from a theoretical point of view, MMI strategies seem especially appropriate as they explicitly account for several types of missing data in a dependent structure, which is common in large-scale assessments. The present article therefore assesses whether using MMI leads to more precise estimates of the uncertainty of person parameters than SMI when the proportion of missing values on background variables is substantial. This question is of particular interest if it seems plausible that missing background data are MAR instead of MCAR. The study only investigates the third type of missing data and assumes that missing values in the item responses are MCAR (as a consequence of multiple matrix sampling) and missing values on background variables are MAR. Specifically, our study addressed the following research questions:

1. Does either SMI or MMI of missing values on background variables lead to unbiased estimates of subpopulation proficiency means, the mean subpopulation proficiency difference, and regression coefficients of proficiency on background variables?
2. Does either SMI or MMI of missing values on background variables lead to unbiased estimates of standard errors of subpopulation proficiency means, the mean subpopulation proficiency difference, and regression coefficients?

We expected that both methods would lead to unbiased mean estimates and regression coefficients. Moreover, the MMI method was expected to capture the uncertainty more adequately and was expected to result in slightly higher standard errors and slightly larger confidence intervals for parameters.

Method

We conducted a simulation study to compare the SMI and the MMI methods when missing proportions on background variables were substantial. Missing values on background variables were assumed to be dependent on θ (i.e., an MAR condition was assumed for the simulation).

Simulation design

In order to establish realistic conditions for the simulation study, we used item parameters from reading comprehension tasks from the German National Assessment Study 2011 in elementary schools (LV 2011) (Stanat et al. 2012), a nationally representative German large-scale assessment study. As is common in large-scale studies, the LV 2011 used a balanced multiple matrix sampling design to compile the test booklets. For the present study, 80 dichotomous reading items were grouped into nine disjoint blocks. Each block contained 6 – 16 items (9 items on average). 20 booklets with one or two blocks in each booklet were used in a rotated booklet design to mimic a balanced

incomplete block design (see e.g., Frey et al. 2009; Gonzalez & Rutkowski 2010). The booklet design is displayed in Table 1. The original item difficulties from the LV 2011 were used to simulate the item responses. The item difficulties ranged from -2.995 to 2.689, which corresponds to a probability for a correct response ranging from 92.9 percent to 9.1 percent. Item responses of $N = 2,000$ persons were simulated resulting in 400 valid responses per item.

Two uncorrelated background variables were used in the simulation. The first one, X_1 , was uniformly distributed with two levels 0 and 1 (i.e., a grouping variable splitting the population into two subpopulations). The second one, X_2 , was normally distributed. The true person proficiency distribution was simulated to depend on both background variables. The population model (see Equation 2) was

$$\theta_n = \beta_1^* X_1 + \beta_2^* X_2 + E_n, \tag{10}$$

with $E_n \sim N(0, \sigma_\theta^2)$ and $\sigma_\theta^2 = 1.44$. Equation 10 is used to both generate and analyze the data. In order to differentiate between the parameters of these models, we use asterisks for the parameters β_1^* and β_2^* of the simulation model to indicate that these are true parameters. In contrast, the parameters of the population model used to analyze the data are denoted β_1 and β_2 . For β_1^* and β_2^* , a combination of values of 0.1 and 0.4 were used, which resulted in $2 \times 2 = 4$ conditions for the population model. The coefficients were chosen to mimic conditions which can be expected in real large-scale assessments. In

Table 1 Rotated block design to form test booklets

| Booklet | Block position 1 | Block position 2 |
|---------|------------------|------------------|
| 1 | | TL 02 |
| 2 | TL 03 | TL 04 |
| 3 | TL 01 | TL 07 |
| 4 | TL 05 | |
| 5 | TL 04 | TL 01 |
| 6 | TL 06 | TL 05 |
| 7 | TL 09 | TL 01 |
| 8 | TL 07 | TL 06 |
| 9 | TL 07 | TL 03 |
| 10 | TL 09 | TL 06 |
| 11 | TL 05 | TL 09 |
| 12 | TL 08 | TL 07 |
| 13 | TL 04 | TL 02 |
| 14 | TL 08 | |
| 15 | TL 01 | TL 03 |
| 16 | | TL 02 |
| 17 | TL 02 | TL 05 |
| 18 | TL 06 | TL 08 |
| 19 | TL 08 | TL 04 |
| 20 | TL 03 | TL 09 |

Note. TL = testlet. Empty cells mean that the corresponding position in the booklet is non-occupied. For example, booklet 1 only contains testlet 2.

The booklets were used to link the items in the testlets.

the German National Assessment Study in secondary schools 2012, for example, the largest effect of socioeconomic status (SES) on proficiency is 49 points on the PISA scale with $M = 500$ and $SD = 100$ (Kuhl et al. 2013). This corresponds to a regression coefficient of about .45. For each background variable, 25% and 40% of the values were deleted, mimicking empirical findings on proportions of missing data for sensitive variables (i.e. SES) when the completion of questionnaires is voluntary (Stanat et al. 2012). The patterns of missing data of the background variables were in turn created to be dependent on true person proficiencies. For each background variable, the polyserial correlation of the missing pattern with the true person proficiencies was $-.10$ or $-.40$. To keep the number of simulations manageable, we refrained from testing against a baseline condition (i.e., 0% missing values or a zero correlation) as we did not expect to find any bias under these conditions.

The simulation was set up using a full-factorial experimental design so that all possible combinations would be represented. We varied the following factors and factor levels:

- Regression coefficients in the population model: $\beta_1^* = \{.1, .4\}$ and $\beta_2^* = \{.1, .4\}$
- Proportions of missing data for both background variables X_1 and X_2 : $m(X_1) = \{.25, .4\}$ and $m(X_2) = \{.25, .4\}$
- Correlations between the patterns of missing data in both background variables and θ^* : $d(X_1, \theta^*) = \{-.1, -.4\}$ and $d(X_2, \theta^*) = \{-.1, -.4\}$

In the next step, the 20 booklets were randomly distributed to virtual persons. Using the true person proficiency scores from Equation 10 and the known item parameters, the response probability for person n and item i was defined according to the item response model (see Equation 1):

$$P(X_{ni} = 1) = \frac{e^{\theta_n - b_i}}{1 + e^{\theta_n - b_i}}. \quad (11)$$

Each response X_{ni} was generated by sampling a value from a uniform distribution across the interval $[0, 1]$. If the sampled value was between 0 and $P(X_{ni} = 1)$, X_{ni} was set to 1. Otherwise, X_{ni} was set to 0.

We used the package TAM (Kiefer et al. 2013) in R (R Core Team 2014) to specify a marginal IRT model without conditioning on background variables, where the item parameters were fixed at their known values. This analysis was done for the sole purpose of obtaining a WLE for each person. The WLE served as a rough point estimate of the unknown true proficiency score.

Next, the missing values of both background variables were imputed using the R package MICE (van Buuren & Groothuis-Oudshoorn 2011). The imputation model consisted of three variables— X_1 , X_2 , and the WLE—in a fully conditional specification (see van Buuren 2007) to account for the process that created the missing data. Two different imputation methods were applied. For X_1 , we used a logistic regression imputation according to its binomial distribution. For X_2 , we used predictive mean matching. Overall, $M = 5$ imputed data sets were generated. From this point on, the procedure varied depending on whether missing values were imputed under a single or nested imputation method.

SMI method

The first imputed data set was used to specify the population model in order to estimate the posterior distribution of the person parameters with item parameters fixed at their known values. Using TAM, twenty plausible values were drawn from the posterior distribution. The mean proficiency estimates were pooled across these 20 imputed values according to Rubin (1987). The regression estimates were taken from the latent regression part of the population model. Therefore, pooling was not necessary.

MMI method

The method described above was repeated $M = 5$ times, which resulted in $5 \times 20 = 100$ plausible values. The mean proficiency estimates were pooled over 100 imputed values according to Rubin (2003), whereas the regression estimates were pooled over $M = 5$ imputed values according to Rubin (1987).

Measures

The estimates of interest were

- the mean proficiency estimates in both subpopulations (i.e., $\bar{\theta}_{X_1=0}$ and $\bar{\theta}_{X_1=1}$)
- the mean proficiency difference (i.e., $\bar{\theta}_{X_1=1} - \bar{\theta}_{X_1=0}$)
- the estimated coefficient β_2 in the latent regression model

The analyses were repeated 1,000 times for each of the 64 conditions. For all estimates mentioned above, three measures were of particular interest in each condition. To examine whether a bias in the corresponding estimates occurred, we computed

$$Bias(x) = N^{-1} \sum_{i=1}^N (x_i - x), \quad (12)$$

where N is the number of replications, x_i is the estimate in the i th replication, and x is the true parameter which was used for data generation. For example, if we consider the bias in the regression coefficient β_2 , x_i is the estimate of β_2 in the i th replication, and $x = \beta_2^*$. If we consider the bias in the subpopulation differences, x_i is the estimate of $\bar{\theta}_{X_1=1} - \bar{\theta}_{X_1=0}$ in the i th replication, and x is the true subpopulation difference, i.e. $x = \beta_1^*$.

The root mean square error (RMSE) between the true x and the estimated x_i parameters were computed by using

$$RMSE(x) = \sqrt{N^{-1} \sum_{i=1}^N (x_i - x)^2}. \quad (13)$$

Finally, we were interested in the recovery proportion; that is: How often does the 95% confidence interval include the true value? If the standard error of the estimate had no bias (and the estimate itself had no bias), we would expect that the 95%

confidence interval would include the true value in 95% of all cases. The 95% confidence interval for x_i was computed as

$$CI_{95}(x_i) = x_i \pm 1.96 \cdot SE(x_i). \quad (14)$$

To identify whether the bias or the RMSE depended on the simulated conditions, we calculated ANOVA effect sizes across all replications of the simulation. All ANOVA analyses contained the following seven factors, each with two levels: β_1^* , β_2^* , missing proportion on X_1 ($m(X_1)$), missing proportion on X_2 ($m(X_2)$), dependency between θ^* and the missing values on X_1 ($d(X_1, \theta^*)$), dependency between θ^* and the missing values on X_2 ($d(X_2, \theta^*)$), and the imputation method (SMI vs. MMI). For the sake of conciseness and clarity of results, we only computed main effects, two-way interactions, and three-way interactions in a common model. Hence, $7 + \binom{7}{2} + \binom{7}{3} = 63$ effects were estimated. For the same reason, only effects for which the effect size η^2 exceeded 0.005 are displayed. Moreover, we computed ANOVAs without interaction effects in order to describe whether the addition of interaction effects leads to an increase in explained variance.

To identify whether the recovery proportion depended on the simulated conditions, we conducted a logistic regression analysis. For each of the 1,000 replications of the 64 simulated conditions, we created a dummy code that equaled 1 if the confidence interval of the corresponding estimate incorporated the true value and 0 otherwise. This dummy code was used as the dependent variable in a logistic regression analysis. All regression analyses contain the seven factors mentioned above, each with two levels.

Results

The following section summarizes the results of the simulation, focusing on differences between the SMI and the MMI methods. First, we present the bias for the three estimates' mean proficiency difference, β_1 and β_2 . We examine whether the bias was influenced by conditions of the simulation using descriptive analyses and ANOVA effect size estimation. The same procedure was carried out for the RMSE and the recovery proportion.

The table for the means of all measures in all of the 64 conditions for both single and nested imputation of background variables is provided in Additional file 1 (Table B1).

Bias

Tables 2 and 3 display ANOVA effect size estimates of bias in the subpopulation differences (Table 2) and of bias in the regression coefficient β_2 (Table 3). Across all simulated conditions, the bias for all estimates was negligible (mean bias in the subpopulation differences = -0.016; mean bias for regression coefficient $\beta_2 = 0.017$). Moreover, the differences between SMI and MMI were negligible (the main effect of imputation method failed to account for a substantial amount of variance in the bias of the subpopulation differences and the regression coefficient β_2). Whereas the overall bias was small, some conditions were associated with higher bias: For example, higher correlations between the patterns of missing data in X_1 and θ^* caused a higher bias in the estimation of subpopulation differences. Higher correlations between the patterns of missing data in X_2 and θ^* caused a higher bias in the estimation of subpopulation

Table 2 ANOVA effect size table for bias in subpopulation

| Differences factor | SS | η^2 |
|---|--------|----------|
| $d(X_1, \theta^*) \times d(X_2, \theta^*)$ | 52.14 | 0.062 |
| $d(X_1, \theta^*)$ | 45.18 | 0.054 |
| $\beta_1^* \times m(X_1)$ | 21.48 | 0.025 |
| $\beta_2^* \times d(X_1, \theta^*)$ | 19.98 | 0.024 |
| $\beta_1^* \times m(X_1) \times d(X_2, \theta^*)$ | 17.65 | 0.021 |
| $m(X_1) \times m(X_2)$ | 13.54 | 0.016 |
| $\beta_2^* \times d(X_1, \theta^*) \times d(X_2, \theta^*)$ | 10.87 | 0.013 |
| $\beta_2^* \times m(X_1) \times d(X_1, \theta^*)$ | 7.24 | 0.009 |
| $\beta_2^* \times m(X_2)$ | 7.13 | 0.008 |
| $\beta_2^* \times m(X_2) \times d(X_2, \theta^*)$ | 6.96 | 0.008 |
| $\beta_1^* \times \beta_2^* \times d(X_1, \theta^*)$ | 6.08 | 0.007 |
| $\beta_1^* \times \beta_2^*$ | 5.62 | 0.007 |
| $m(X_1) \times m(X_2) \times d(X_2, \theta^*)$ | 4.81 | 0.006 |
| $m(X_1) \times d(X_2, \theta^*)$ | 4.66 | 0.006 |
| $d(X_2, \theta^*)$ | 4.56 | 0.005 |
| $\beta_2^* \times m(X_1) \times d(X_2, \theta^*)$ | 4.47 | 0.005 |
| $m(X_1) \times d(X_1, \theta^*) \times m(X_2)$ | 4.40 | 0.005 |
| Residuals | 563.55 | |
| Total | 844.02 | |

differences and in the estimation of β_2 . Higher proportions of missing data for background variable X_1 caused a bias in the estimation of β_2 . However, most ANOVA effect sizes were small. For example, the seven main effects of the ANOVA analysis displayed in Table 2 together only account for 7.3% of the explained variance, whereas all 64 effects together account for 33.2% of the explained variance. Considering Table 3, the seven main effects only account for 5.7% of the explained variance, whereas all 64 effects account for 20.4% of the explained variance. The bias was therefore rarely associated with a single factor (e.g., a large proportion of missing values alone) but rather with combinations of (adverse) conditions (e.g., a large proportion of missing values *and* a high level of dependence of the pattern of missing data on true person

Table 3 ANOVA effect size table for bias in regression coefficient β_2

| Factor | SS | η^2 |
|--|--------|----------|
| $m(X_1)$ | 12.90 | 0.087 |
| $\beta_1^* \times \beta_2^*$ | 4.13 | 0.028 |
| $d(X_2, \theta^*)$ | 3.17 | 0.022 |
| $\beta_1^* \times m(X_1)$ | 2.42 | 0.016 |
| β_1^* | 2.40 | 0.016 |
| $\beta_1^* \times \beta_2^* \times d(X_1, \theta^*)$ | 1.36 | 0.009 |
| $\beta_1^* \times d(X_2, \theta^*)$ | 1.19 | 0.008 |
| $\beta_1^* \times \beta_2^* \times m(X_2)$ | 1.04 | 0.007 |
| $\beta_2^* \times m(X_2) \times d(X_2, \theta^*)$ | 0.82 | 0.006 |
| Residuals | 110.01 | |
| Total | 147.56 | |

proficiencies). Both SMI and MMI are therefore able to compensate for a single adverse condition. As expected, the imputation method (SMI vs. MMI) did not affect the bias—all ANOVA analyses revealed only negligible main effects of the simulation methods or interaction effects of the simulation method and the other factors.

RMSE

Tables 4 and 5 display ANOVA effect size estimates for the RMSE in the subpopulation differences (Table 4) and in the regression coefficient β_2 (Table 5). Across all simulated conditions, the RMSE for the subpopulation differences was .076. The mean RMSE for the regression coefficient β_2 was .035. The RMSE differed between the two imputation methods, but only slightly and only for the regression coefficient β_2 : The RMSE was .037 for the single imputation method and .032 for the nested imputation method. This main effect was significant ($\eta^2 = 0.011$, see Table 5). Considering Table 4, the seven main effects of the ANOVA analysis only account for 4.4% of the explained variance, whereas all 64 effects account for 22% of the explained variance. Considering Table 5, the main effects account for 10% of the explained variance, whereas all 64 effects account for 25.3% of the explained variance.

Although we found a significant main effect of imputation method on the RMSE for the regression coefficient β_2 , its effect size was small and the reduction of .037–.032 = .005 in the RMSE is negligible.

Recovery proportion

We conducted logistic regression analyses to predict the probability that the confidence interval of an estimate (e.g., β_2) would include the true value. Tables B2 and B3 are provided in Additional file 1 and display the results of the logistic regression analyses for the recovery proportion of the subpopulation differences (Table B2) and regression coefficient β_2 (Table B3). Comparing the SMI and MMI methods, we found mean recovery proportions of .83 versus .95 for the subpopulation differences, and .85 versus .96 for the regression coefficient β_2 . The main effect for the imputation method was significant both for subpopulation differences and for the regression coefficient β_2 . MMI

Table 4 ANOVA effect size table for the RMSE of subpopulation differences

| Factor | SS | η^2 |
|--|--------|----------|
| $\beta_2^* \times m(X_2) \times d(X_2, \theta^*)$ | 9.37 | 0.031 |
| $\beta_1^* \times \beta_2^* \times d(X_1, \theta^*)$ | 9.31 | 0.030 |
| $\beta_1^* \times d(X_2, \theta^*)$ | 6.92 | 0.023 |
| $m(X_1)$ | 5.38 | 0.018 |
| $\beta_1^* \times \beta_2^*$ | 5.16 | 0.017 |
| $d(X_1, \theta^*)$ | 4.58 | 0.015 |
| $\beta_2^* \times d(X_1, \theta^*) \times m(X_2)$ | 2.01 | 0.007 |
| $\beta_1^* \times m(X_1) \times d(X_1, \theta^*)$ | 1.88 | 0.006 |
| $\beta_1^* \times d(X_1, \theta^*) \times m(X_2)$ | 1.85 | 0.006 |
| $m(X_1) \times m(X_2) \times d(X_2, \theta^*)$ | 1.62 | 0.005 |
| Residuals | 238.17 | |
| Total | 305.29 | |

Table 5 ANOVA effect size table for the RMSE of regression coefficient β_2

| Factor | SS | η^2 |
|---|-------|----------|
| $m(X_1)$ | 4.34 | 0.064 |
| β_2^* | 1.08 | 0.016 |
| $\beta_1^* \times \beta_2^*$ | 0.92 | 0.013 |
| $d(X_1, \theta^*) \times m(X_2) \times d(X_2, \theta^*)$ | 0.90 | 0.013 |
| <i>imp. Method</i> | 0.74 | 0.011 |
| $\beta_1^* \times d(X_1, \theta^*) \times d(X_2, \theta^*)$ | 0.73 | 0.011 |
| $m(X_1) \times d(X_1, \theta^*) \times d(X_2, \theta^*)$ | 0.67 | 0.010 |
| $\beta_2^* \times d(X_1, \theta^*) \times m(X_2)$ | 0.65 | 0.010 |
| $m(X_1) \times m(X_2)$ | 0.60 | 0.009 |
| $\beta_1^* \times m(X_1)$ | 0.56 | 0.008 |
| $\beta_1^* \times m(X_2)$ | 0.52 | 0.008 |
| $\beta_2^* \times m(X_1) \times d(X_1, \theta^*)$ | 0.52 | 0.008 |
| $\beta_2^* \times m(X_2)$ | 0.43 | 0.006 |
| $\beta_1^* \times \beta_2^* \times d(X_1, \theta^*)$ | 0.36 | 0.005 |
| Residuals | 50.82 | |
| Total | 68.01 | |

overall leads to a higher recovery proportion than SMI. Moreover, the advantage of MMI over SMI is particularly apparent in some specific conditions, for example, if the missing proportion on X_1 is 40% instead of 10% (line 23 in Table B2). This indicates that the MMI method yields a higher recovery proportion if the proportion of missing values is large, or if the correlation between the patterns of missing data in X_2 and θ^* is -0.4 instead of -0.1 . To summarize, MMI seems appropriate for (a combination of) some adverse conditions. This finding reflects the fact that the uncertainty in the imputed values, which results from some adverse conditions, is only incorporated in the MMI method.

Discussion

The goal of the present study was to compare two imputation methods—SMI and MMI—for handling missing data in background variables in large-scale assessment programs. The first key question of the present paper was whether these imputation methods would allow unbiased estimation of subpopulation differences and coefficients in the latent regression model. The second question was related to the standard errors of the estimates: How often does the confidence interval include the true value?

Concerning the first research question of unbiased population estimates, none of the simulated conditions showed differences between SMI and MMI. The mean bias was negligible for both imputation methods. Some adverse factors (and especially combinations of adverse factors, e.g., a large proportion of missing values and a high level of dependence of the missing data pattern on true person proficiency) led to biases in the range of -0.145 to 0.092 for the subpopulation differences and -0.041 to 0.061 for the regression coefficient β_2 .

Concerning the second research question, when considering the RMSE and especially the recovery proportion, the MMI method outperformed the SMI method. As the MMI method takes the between-imputation variance in the imputed background data

into account, the standard errors of the regression coefficient β_2 and of the subpopulation difference accurately reflected the uncertainty due to the amount of missing data. The estimates of the regression coefficient β_2 varied less between the replications and resulted in lower RMSEs. The differences between the two imputation methods, however, were apparent only for substantial proportions of missing data of about 40%, which may occur empirically for some sensitive variables such as SES (Stanat et al. 2012). Hence, we recommend using MMI instead of SMI if the missing data proportion is substantial *and* if researchers are interested not only in unbiased estimates (e.g., for purely descriptive purposes) but also in unbiased standard errors of these estimates (e.g., for significance testing).

In general, we conclude that imputation methods can provide a suitable alternative to dummy coding, which is flawed by some conceptual problems and may cause biased mean proficiency estimates if the proportion of missing values on the background variables is substantial (Rutkowski 2011). To date, multiple imputation methods per se are widely established in large-scale assessments to estimate θ_n values, which are considered to be inherently missing in the IRT framework, whereas the problem of missing data in background variables has rarely been tackled by imputation methods. Our analyses show that MMI yields only small practical advantages compared to SMI at the cost of higher computational effort. Therefore, SMI should be a sufficient method in most applications.

Typically, large-scale assessments comprise a multitude of background variables, which are reduced to several uncorrelated principal components prior to the estimation of the population model. Compared to this procedure, the present simulation is highly simplified. In empirical applications where background variables typically are correlated, it is plausible to assume that the more background variables are included, the less problematic is a high proportion of missing values on a single variable as other variables might compensate the loss of information. SMI and MMI methods may easily be combined with principal component analyses, which can be applied after the imputation of missing values in background variables. With MMI, this would result in several data sets, where the number of principal components may vary between the data sets.

A limitation of both SMI and MMI methods is the use of the WLE as a rough proficiency estimate for the unobserved person parameter θ_n in the imputation model. Strictly speaking, the WLE is only an unbiased estimate of the population mean when the same test items are administered to all persons (Wu 2005). In large-scale assessments, however, this requirement is not fulfilled. Moreover, WLEs suffer from scale unreliability and contain error variance (Mislevy et al. 1992). Using the WLEs in the imputation model may therefore underestimate the relation between θ and the background variables. This underestimation is expected to be more severe if only few items from the item pool are presented to each person. This would decrease the WLE reliability, which, in turn, leads to a poorer representation of the relation between θ and the background variables in the imputation model. In our simulation, the number of items and persons is constant over all conditions. Consequently, the WLE reliability remained stable at about .70. Future research should investigate the influence of scale reliability when individual θ_n estimates (e.g., WLEs or EAPs) are used in the imputation model to gain information about the level of precision (e.g., in the estimation of subpopulation differences) that could be expected given certain characteristics of the

experimental design (e.g., sample size, number of items, number of background variables, or proportion of missing values on background variables). A further improvement of the imputation approach might be to include the unreliability of the WLE as a possible source of uncertainty into the imputation model. However, as a three-level nested imputation this would result in much more computational effort which may not be desirable in empirical applications.

Conclusion

The present study showed that nested imputation methods are suitable approaches to deal with missing values in both continuous and categorical background variables. Both SMI and MMI yield virtually unbiased estimates of subpopulation differences and regression coefficients for background variables with missing data. Concerning the estimation of standard errors, MMI more accurately reflect the uncertainty due to missing data in the background variables than SMI, resulting in slightly larger standard errors. However, the differences between imputation methods are small in non-extreme conditions of missing data. Thus, both SMI and MMI can be used to impute missing values on background variables in large-scale assessments to avoid the conceptual flaws and the possible biases associated with the common approach of dummy coding missing observations.

Additional file

Additional file 1: Appendix to the paper: Nested Multiple Imputation in Large-Scale Assessments.

Competing interests

The authors declare that they have no competing interests.

Authors' contributions

SW, NH and MH reviewed the literature. SW, NH, OL and TS designed the analysis. SW conducted the analysis and evaluated the results. SW wrote the manuscript. KB and OL read and revised the manuscript. TS and MH verified the simulation program code. All authors read and approved the final manuscript.

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