Incomplete Data Analysis

V. Inácio de Carvalho & M. de Carvalho

University of Edinburgh





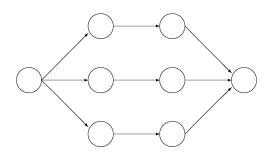












Incomplete data Imputed data Analysis results Pooled results

- → In summary:
 - **Imputation**: impute multiple times.
 - Analysis: analyse each of the datasets.
 - Pooling: combine results, taking into account additional uncertainty.

- \hookrightarrow Create a number (M > 1) of copies of the incomplete dataset, and use an appropriate procedure to impute the missing values in each of these copies.
- The imputed datasets are composed of a fixed portion the observed data– and a varying portion the imputed values. Since we do not know the true values that are missing it seems reasonable that the imputed values used in each copy should in general differ from each other.
- \hookrightarrow The choice of *M* is discussed later in the next set of slides.

- → We have created *M* imputed datasets that are now complete. How do we analyse them?
- \hookrightarrow For now, we will assume that our focus is on estimating a single (univariate) parameter, which we denote by θ .
- \hookrightarrow For instance, θ can be the mean or median of a variable, the proportion of individuals in a particular categorical (level) of a factor variable, a coefficient in a regression model, etc.
- → For each imputed dataset, perform the analysis of interest (e.g., estimating the mean or fitting the regression model) that would have been performed in the absence of missing values. In the MI literature, the model of interest is sometimes referred to as the substantive model.
- \hookrightarrow Store the parameter estimate and its variance (the squared standard error). The estimate of θ obtained from the mth $(m=1,\ldots,M)$ complete dataset is denoted by $\widehat{\theta}^{(m)}$ and its (estimated) variance (squared standard error) by $\widehat{U}^{(m)}$.

- \hookrightarrow After step 2, we have the results from M analyses, that is, we have $\widehat{\theta}^{(1)},\ldots,\widehat{\theta}^{(M)}$, and $\widehat{U}^{(1)},\ldots,\widehat{U}^{(M)}$.
- → How do we now combine them to come up with a final estimate and how to measure the
 uncertainty about such estimate?
- \hookrightarrow According to the so-called **Rubin's rules**, the multiple imputation estimate of θ , $\widehat{\theta}^{\text{MI}}$, is the average of the M individual estimates, that is,

$$\widehat{\theta}^{\mathsf{MI}} = \frac{1}{M} \sum_{m=1}^{M} \widehat{\theta}^{(m)}.$$

- \hookrightarrow To estimate the variance of $\widehat{\theta}^{\text{MI}}$, we **do not** simply average the variances from each dataset. It is slightly more complicated, but not that much complicated!
- → First, we calculate the between-imputation variance

$$B = \frac{1}{M-1} \sum_{m=1}^{M} \left(\widehat{\theta}^{(m)} - \widehat{\theta}^{MI} \right)^{2},$$

- \hookrightarrow This is simply the usual unbiased sample variance formula applied to $\widehat{\theta}^{(1)}, \dots, \widehat{\theta}^{(M)}$.
- \hookrightarrow B measures how much the estimates of θ vary across the imputed datasets.
- If there is very little missing data, the estimates from the different imputed datasets will be very similar and the between imputation variance will be small.
- → The larger the amount of missing data, the larger the variability in the estimates between the imputed datasets, and the larger the between imputation variance will be.
- \hookrightarrow B thus captures uncertainty in $\widehat{\theta}^{\mathrm{MI}}$ due to missing data.



Step 3

 \hookrightarrow Second, we calculate the **within-imputation** variance

$$\bar{U} = \frac{1}{M} \sum_{m=1}^{M} \widehat{U}^{(m)},$$

where $\widehat{U}^{(m)}$ is the estimated variance of $\widehat{\theta}^{(m)}$. This is simply the average of the individual variance estimates.

 \hookrightarrow The within imputation variance \bar{U} is measuring the uncertainty due to the fact that the sample is of finite size (i.e., we are not using the entire population). This is the usual source of uncertainty in parameter estimates.

Step 3

- \hookrightarrow It is tempting to conclude that the **total variance** V^{MI} is equal to the sum of \bar{U} and B, but that would be incorrect.
- \hookrightarrow We need to incorporate the fact that $\widehat{\theta}^{\text{MI}}$ itself is estimated using finite M, and thus only approximates $\widehat{\theta}^{\text{MI}}_{\infty}$, the estimator that would have been obtained for an infinitely large number of imputations $M=\infty$.
- \hookrightarrow Rubin (1987, eq. 3.3.5) shows that the contribution to the variance of this factor is systematic and equal to B_{∞}/M . Since B approximates B_{∞} (estimated between imputation variance for infinitely many imputations), we may write:

$$\begin{split} V^{\mathsf{MI}} &= \bar{U} + B + \frac{B}{M} \\ &= \bar{U} + \left(1 + \frac{1}{M}\right) B, \end{split}$$

for the total variance of $\widehat{\theta}^{MI}$.



- \hookrightarrow The inclusion of the term B/M is critical to make multiple imputation work at low values of M.
- → Not including it would result in p-values that are too low or confidence intervals that are too short.

- \hookrightarrow In summary, the total variance V^{MI} stems from three sources:
 - \bar{U} , the variance caused by the fact that we are taking a sample rather than observing the entire population. This is the conventional measure of variability.
 - B, the extra variance caused by the fact that there are missing values in the sample.
 - **3** B/M, the extra simulation variance caused by the fact that $\widehat{\theta}^{MI}$ itself is estimated for finite M.
- \hookrightarrow Note that if there were no missing values then B would be equal to zero and the estimated total variance V^{MI} would be \bar{U} .

Rubin's rules-toy example

- \hookrightarrow A confidence interval for θ can be constructed based on V^{MI} and $\widehat{\theta}^{\text{MI}}$.
- \hookrightarrow Specifically, the $(1 \alpha)100\%$ confidence interval is then

$$\widehat{\theta}^{\mathsf{MI}} \pm \mathit{t}_{\nu} \left(\frac{\alpha}{2}\right) \sqrt{\mathit{V}^{\mathsf{MI}}},$$

with $t_{\nu}\left(\frac{\alpha}{2}\right)$ is the $\alpha/2$ quantile of the t distribution with $\nu=(M-1)(1+1/r_{M})^{2}$, where $r_{M}=(1+1/M)B/\bar{U}$ is the relative increase in variance due to missing values.

- → Notice that r_M does not depend on the sample size of the observed data. This can lead to situations where the degrees of freedom are larger than those for the complete case analysis, which is inappropriate.
- → To avoid this problem, Barnard and Rubin (1999) proposed an improvement to calculate the degrees of freedom. This improved version is implemented in the mice package.

Rubin's rules-toy example

→ Suppose we take a survey of five people, measuring their height and weight. Only three of them disclosure their weight; the other two don't give it just because of random chance. The data are:

Height (inches)	Weight (pounds)		
65	130		
68	140		
70	150		
72	NA		
75	NA		

→ The aim of the analysis (step 2) is to regress the weight on the height, that is, our statistical model of interest is

weight =
$$\beta_0 + \beta_1$$
 height + ε , $\varepsilon \sim N(0, \sigma^2)$.



Rubin's rules-toy example

→ Suppose that five plausible values for each missing weight have been generated (represented below in blue) to create five complete datasets.

Height	Weight-1	Weight-2	Weight-3	Weight-4	Weight-5
65	130	130	130	130	130
68	140	140	140	140	140
70	150	150	150	150	150
72	157	166	155	157	156
75	171	169	167	171	168
Estimated slope $(\widehat{\beta}_1)$	4.12	4.26	3.71	4.12	3.83
$\widehat{U} = \widehat{\operatorname{var}}(\widehat{\beta}_1)$	(0.025)	(0.346)	(0.024)	(0.025)	(0.018)

Rubin's rules-toy example

 \hookrightarrow The final estimate for the slope is

$$\widehat{\beta}_1^{\text{MI}} = \frac{1}{5}(4.12 + 4.26 + 3.71 + 4.12 + 3.83) = 4.008$$

→ The within imputation variance is

$$\bar{U} = \frac{1}{5}(0.025 + 0.346 + 0.024 + 0.025 + 0.018) = 0.0876$$

$$B = \frac{1}{4} \{ (4.12 - 4.008)^2 + (4.26 - 4.008)^2 + (3.71 - 4.008)^2 + (4.12 - 4.008)^2 + (3.83 - 4.008)^2 \}$$

= 0.05227

→ Thus, the final estimate of the variance is

$$V^{\text{MI}} = 0.0876 + \left(1 + \frac{1}{5}\right) \times 0.05227 = 0.150324$$



Rubin's rules - multivariate case

- \hookrightarrow Extensions to the case where the parameter of interest is a *p*-component vector, say $\theta = (\theta_1, \dots, \theta_p)'$, are straightforward.
- → For the estimate of the parameter vector we have

$$\widehat{\theta}^{\text{MI}} = \frac{1}{M} \sum_{m=1}^{M} \widehat{\theta}^{(m)}.$$

→ In the multivariate context, the within-imputation covariance matrix is the average of the M covariance matrices, namely

$$\bar{\mathbf{U}} = \frac{1}{M} \sum_{m=1}^{M} \widehat{\mathbf{U}}^{(m)},$$

where $\widehat{\mathbf{U}}^{(m)}$ is the covariance matrix from the completed dataset m.

Rubin's rules - multivariate case

$$\mathbf{B} = \frac{1}{M-1} \sum_{m=1}^{M} \left(\widehat{\boldsymbol{\theta}}^{(m)} - \widehat{\boldsymbol{\theta}}^{\mathsf{MI}} \right) \left(\widehat{\boldsymbol{\theta}}^{(m)} - \widehat{\boldsymbol{\theta}}^{\mathsf{MI}} \right)^{\mathsf{T}},$$

where $\widehat{\theta}^{(m)}$ contains the parameter estimates from the *m*th imputed dataset, and $\widehat{\theta}^{\text{MI}}$ is the vector of pooled point estimates (i.e., the arithmetic average of the $\widehat{\theta}^{(m)}$ vectors).

- The diagonal elements of B contain the between imputation variance estimate for individual parameters, and the off-diagonal elements quantify the extent to which the between imputation fluctuation in one parameter is related to the between imputation fluctuation in another parameter.
- Considered as a whole, the between imputation covariance matrix represents the additional sampling fluctuation that results from the missing data.
- Finally, the total covariance matrix combined the within and between imputation covariance matrices as follows

$$\mathbf{V}^{\mathsf{MI}} = \mathbf{\bar{U}} + \mathbf{B} + \frac{1}{M}\mathbf{B},$$
$$= \mathbf{\bar{U}} + \left(I + \frac{1}{M}\right)\mathbf{B},$$

where I is the identity matrix.



Some remarks on the 3 steps

- → As we shall see, the only complex part of multiple imputation is step one: formulate a good imputation model.
- → The specification of an appropriate imputation model is the key issue, since if this is misspecified, there is the potential for bias.
- → The second step, producing the final estimate, is straightforward as it treats each imputed dataset as if it were a real dataset, we just have to do it M times.
- → As Schafer (1997) says, multiple imputation works by "solving an incomplete-data problem by repeatedly solving the complete-data version".
- → The third step involves simple arithmetic and typically we do not need to implement Rubin's rules manually as they are coded into most multiple imputation packages.

How many imputations?

- \hookrightarrow A natural question arising in the context of multiple imputation is how many copies of the dataset, which we have denoted by M, we should use.
- \hookrightarrow The choice of *M* does not affect the validity of our estimates and inferences.
- $\,\hookrightarrow\,$ However, it does affect their statistical efficiency and reproducibility.

How many imputations?

- \hookrightarrow Rubin originally suggested that unless the fraction of missing data was large, M=3 or M=5 would typically suffice. This advice was based on the statistical efficiency of the multiple imputation point estimate.
- \hookrightarrow Recall that the total variance estimate, based on Rubin's rules, for the multiple imputation point estimate $\widehat{\theta}^{\mathrm{MI}}$, is given by

$$V^{MI} = \bar{U} + \left(1 + \frac{1}{M}\right)B.$$

 \hookrightarrow The most efficient estimator is obtained with $M=\infty$, for which

$$V^{MI} = \bar{U} + B.$$

How many imputations?

 \hookrightarrow The ratio of the variance of the finite M estimate to the $M=\infty$ estimate is

$$\frac{\overline{U}+(1+\frac{1}{M})B}{\overline{U}+B}=1+\frac{1}{M}\frac{B}{\overline{U}+B}.$$

- \hookrightarrow When the amount of missing data is small, the term $\frac{B}{U+B}$ is close to zero, and even a small value of M means that the variance is not much increased compared to using $M=\infty$.
- This is the rationale behind the advice that usually a small M is okay. We need to take this advice with some grains of salt, as computing power in the 70s or 80s was somewhat limited.

How many imputations?

- → Multiple imputation involves generating random numbers.
- → As a result, the point estimates, standard errors, confidence intervals, p-values, etc, all have some inherent Monte-Carlo noise.
- → If someone was to re-run our code (with a different seed!), they would get slightly different results.
- → We may want to pick up a M large enough so that our results are (almost!) reproducible, in the sense that if someone re-run our code, they would get results sufficiently close enough to ours.

How many imputations?

- For instance, we can run our whole multiple imputation procedure, say 3 times, and compare results.
- \hookrightarrow If the results differ by more than what we are comfortable with, we should increase M and try again until results are close enough.
- → Further, imputing a dataset in practice often involves trial and error to adapt and refine the imputation model. Such initial explorations do not require large M.
- \hookrightarrow It is convenient to set, e.g. M=5, during model building, and increase M only after being satisfied with the model for the 'final' round of imputation.

- Remember that we have learned that stochastic regression imputation was a promising approach.
- → So, in the MI context (and for simplicity let us think about a univariate pattern of missingness), if we run stochastic regression imputation M times (i.e., for each missing value we use M draws instead of one) in step 1, is this all we have to do? Well, not exactly...But why?
- Such approach would imply that the regression coefficients and the variance of the error term are known with certainty. Such approach is termed in the literature as **improper multiple imputation**.

- → In practice, the regression coefficients and the variance of the error term are seldom known and must be estimated.
- If we had drawn a different sample from the same population, then our estimates for the regression coefficients and for the variance of the error term would be different, perhaps slightly.
- → The amount of extra variability is strongly related to the sample size, with smaller samples yielding more variable estimates.

- → The parameter uncertainty also needs to be included in the imputations.
- Therefore, to perform proper multiple imputation, we need to reflect the parameters' variability/uncertainty from one imputation to the next.
- → As an aside, the variability of the imputed values in stochastic regression imputation is composed of variability of estimation plus noise.
- → There are two main methods for taking into account the parameter uncertainty:
 - \hookrightarrow **Bayesian methods** draw the parameters directly from their posterior distributions. That is, for each copy m of the dataset, $m = 1, \ldots, M$, we would draw the parameters from the posterior distribution.
 - Bootstrap methods, in turn, resample the complete cases and re-estimate the parameters from the resampled data.



- \hookrightarrow It is useful to consider the consequences of improper multiple imputation.
- \hookrightarrow As a result, the confidence intervals based on V^{MI} would be too narrow.

Choosing the imputation model

- To provide valid estimates and inferences, MI requires data to be MAR and imputation models need to be correctly specified.
- → Of course, as George Box famously said: "All models are wrong, but some are useful". We should nevertheless ensure that our models are a good approximation to the reality.
- → If, for instance, the substantive model of interest includes interactions (between variables), then these should be preserved in the imputation model.
- → Meng (1994) introduced the concept of congeniality to refer to the relation between the imputation model and the analysis model. The imputation model should be 'congenial' with the substantive model.