EP16: Missing Values in Clinical Research: Multiple Imputation

4. A Closer Look at the Imputation Step

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The Imputation Step

The imputation step consists itself of two (or three) steps:

- specification of the imputation model
- 1. estimation / sampling of the parameters
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Notation:

y: variable to be imputed

$$\mathbf{y} = \begin{cases} \mathbf{y}_{obs} \\ \vdots \\ \mathbf{y}_{q} \\ NA \\ \vdots \\ NA \end{cases}$$

X: design matrix of other variables

$$\mathbf{y} = \begin{cases} \begin{cases} y_1 \\ \vdots \\ y_q \\ NA \end{cases} \\ \vdots \\ NA \end{cases}$$

$$\mathbf{X} = \begin{cases} \mathbf{x}_{11} & \dots & \mathbf{x}_{1p} \\ \vdots & \dots & \vdots \\ \mathbf{x}_{q1} & \dots & \mathbf{x}_{qp} \\ \mathbf{x}_{q+1,1} & \dots & \mathbf{x}_{q+1,p} \\ \vdots & \dots & \vdots \\ \mathbf{x}_{n1} & \dots & \mathbf{x}_{np} \end{cases}$$

In the **Bayesian framework: everything unknown** or unobserved is considered a **random variable**.

For example:

- ightharpoonup regression coefficients β ,
- ightharpoonup residual variance σ^2 and
- ightharpoonup missing values \mathbf{y}_{mis} .

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Random variables have a **probability distribution**.

- ► The **expectation** of that distribution quantifies which **values** of the random variable are **most likely**.
- ▶ The **variance** is a measure of the **uncertainty** about the values.

In Bayesian imputation:

 in the observed data: estimate the distribution of the parameters describing the association between incomplete variables and the other data

$$p(\mathbf{y}_{obs} \mid \mathbf{X}_{obs}, \beta, \sigma) \Rightarrow p(\beta \mid \mathbf{y}_{obs}, \mathbf{X}_{obs}), p(\sigma \mid \mathbf{y}_{obs}, \mathbf{X}_{obs})$$

2. use these estimates to obtain the the probability distribution of incomplete variables given the other data

$$p(\mathbf{y}_{mis} \mid \mathbf{X}_{mis}, \boldsymbol{\beta}, \sigma)$$

3. sample values from these distributions **⇒ imputation**

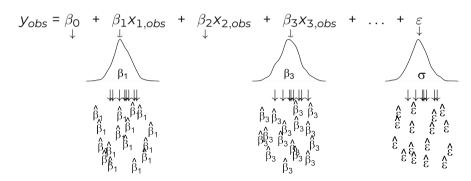
Step 1:

Specify a (Bayesian) regression model



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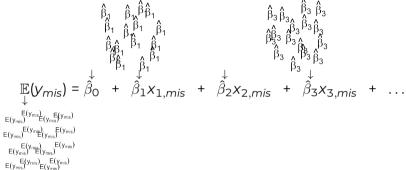
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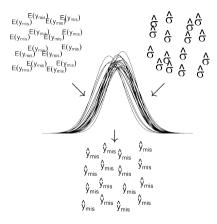
Step 2:

$$\hat{\beta}_{1}, \hat{\beta}_{1}, \hat{\beta}_{2}, \hat{\beta}_{3}, \hat{\beta$$

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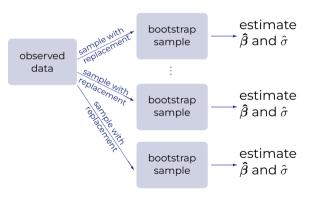


Step 3:



Bootstrap Multiple Imputation

Alternative approach to capture the uncertainty: **bootstrap**



Bootstrap samples can contain some **observations multiple times** and some **observations not at all.**

Bootstrap Multiple Imputation

In bootstrap multiple imputation,

- per imputation: one bootstrap sample of the observed data
- ► the (least squares or maximum likelihood) estimates of the parameters are calculated from

$$\mathbf{y}_{obs} = \mathbf{X}_{obs} \boldsymbol{\beta} + \varepsilon_{obs}$$
 (step 1).

▶ Imputed values are sampled from $p(\mathbf{y}_{mis} \mid \mathbf{X}_{mis}, \hat{\boldsymbol{\beta}}, \hat{\sigma})$ (step 2).

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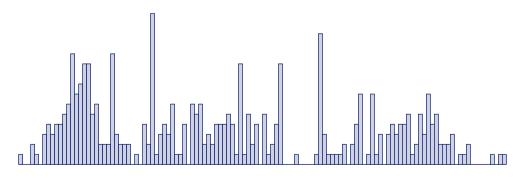
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→ Step 2 is analogous to step 3 in Bayesian multiple imputation.

Both Bayesian and bootstrap multiple imputation sample imputed values from a distribution $p(\mathbf{y}_{mis} \mid \mathbf{X}_{mis}, \hat{\boldsymbol{\beta}}, \hat{\sigma})$.

Sometimes, the empirical distribution can not be adequately approximated by a known probability distribution.



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Idea:

- find cases in the observed data that are similar to the cases with missing values
- fill in the missing value with the observed value from one of those cases

To find similar cases, the predicted values of complete and incomplete cases are compared.

The steps in PMM:

- **1.** Obtain parameter estimates for $\hat{\beta}$ and $\hat{\sigma}$ (see later)
- 2. Calculate the predicted values for the observed cases

$$\hat{m{y}}_{obs}$$
 = $m{X}_{obs}m{\hat{eta}}$

3. Calculate the predicted value for the missing cases

$$\hat{\boldsymbol{y}}_{mis}$$
 = $\mathbf{X}_{mis}\hat{\boldsymbol{\beta}}$

- **4.** For each missing value, find *d* donor candidates that fulfil a given criterion (details on the next slide).
- 5. Randomly select one of the donors.

Several **criteria to select donors** (donor candidates) have been proposed:

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$$|\hat{y}_{mis,i} - \hat{y}_{obs,j}|, j = 1,\ldots,q.$$

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- ▶ Select candidates like in 2. or 3., but select the donor from the candidates with probability that depends on $|\hat{y}_{mis,i} \hat{y}_{obs,j}|$. (Siddique & Belin, 2008)

Potential issues with donor selection

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- ▶ Therefore, using d = 1 (selection criterion 1.) is not a good idea. On the other hand, using too many candidates can lead to bad matches.
- ➤ Schenker & Taylor (1996) proposed an adaptive procedure to select d, but it is not used much in practice.

For the **sampling of the parameters** (step 1), different approaches have been introduced in the literature:

Type-0 $\hat{\beta}_{LS/ML}$ (least squares or maximum likelihood) are used in both prediction models

Type-I $\hat{\beta}_{LS/ML}$ to predict \hat{y}_{obs} ; $\tilde{\beta}_{B/BS}$ (Bayesian or bootstrapped) to predict \hat{y}_{mis} Type-II $\tilde{\beta}$ to predict \hat{y}_{obs} as well as \hat{y}_{mis}

Type-III different draws $\tilde{\beta}_{B/BS}^{(1)}$ and $\tilde{\beta}_{B/BS}^{(2)}$ to predict \hat{y}_{obs} and \hat{y}_{mis} , respectively

The use of Type-0 and Type-I matching **underestimates the uncertainty** about the regression parameters.

Another point to consider: the **choice** of the set of data used to train the prediction models

By default, the same set of data (all cases with observed y) is used to train the model and to produce predicted values of y_{obs} .

The predictive model will likely fit the observed cases better than the missing cases, and, hence, **variation will be underestimated**.

Alternatives:

- ▶ the model could be trained on the whole data (using previously imputed values)
- use a leave-one-out approach on the observed data

What is Implemented in Software?

mice (in R):

- ► PMM via mice.impute.pmm()
 - specification of number of donors d (same for all variables)
 - ▶ Type-0, Type-I, Type-II matching
- ► PMM via mice.impute.midastouch()
 - allows leave-one-out estimation of the parameters
 - distance based donor selection
 - Type-0, Type-I, Type-II matching
- ▶ bootstrap linear regression via mice.impute.norm.boot()
- bootstrap logistic regression via mice.impute.logreg.boot()
- ▶ Bayesian linear regression via mice.impute.norm()
- **.**..

References

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