

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:

0. specification of the imputation model
1. **estimation** / sampling **of the parameters**
2. **drawing imputed values** from the predictive distribution

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Notation:

y: variable to be imputed

$$\mathbf{y} = \begin{matrix} \mathbf{y}_{obs} \\ \mathbf{y}_{mis} \end{matrix} \left\{ \begin{matrix} \begin{bmatrix} y_1 \\ \vdots \\ y_q \\ NA \end{bmatrix} \\ \begin{bmatrix} \vdots \\ NA \end{bmatrix} \end{matrix} \right.$$

X: design matrix of other variables

$$\mathbf{X} = \begin{matrix} \mathbf{X}_{obs} \\ \mathbf{X}_{mis} \end{matrix} \left\{ \begin{matrix} \begin{bmatrix} x_{11} & \dots & x_{1p} \\ \vdots & \dots & \vdots \\ x_{q1} & \dots & x_{qp} \end{bmatrix} \\ \begin{bmatrix} x_{q+1,1} & \dots & x_{q+1,p} \\ \vdots & \dots & \vdots \\ x_{n1} & \dots & x_{np} \end{bmatrix} \end{matrix} \right.$$

Bayesian Multiple Imputation

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

For example:

- ▶ regression coefficients β ,
- ▶ residual variance σ^2 and
- ▶ 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.

Bayesian Multiple Imputation

In **Bayesian imputation**:

1. in the **observed data**:

estimate the distribution of **the parameters** describing the association between incomplete variables and the other data

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

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

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

3. **sample values** from these distributions \rightarrow **imputation**

Bayesian Multiple Imputation

Step 1:

Specify a (Bayesian) regression model

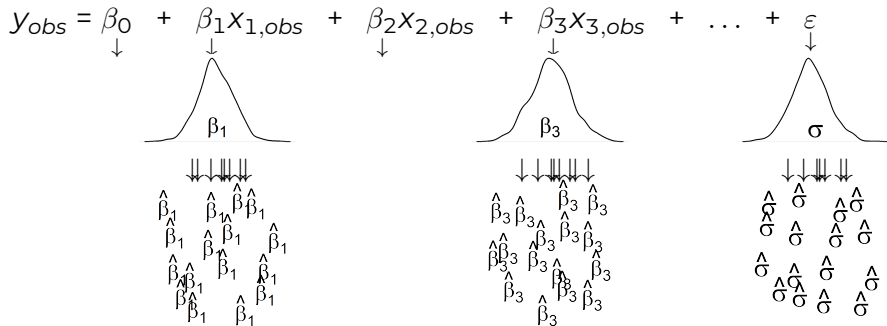
$$y_{obs} = \beta_0 + \beta_1 X_{1,obs} + \beta_2 X_{2,obs} + \beta_3 X_{3,obs} + \dots + \varepsilon$$

The diagram illustrates the Bayesian regression model equation: $y_{obs} = \beta_0 + \beta_1 X_{1,obs} + \beta_2 X_{2,obs} + \beta_3 X_{3,obs} + \dots + \varepsilon$. Below the equation, three normal distribution curves are shown, each representing a prior distribution for a parameter in the model. An arrow points from β_1 in the equation to the first curve, which is labeled β_1 . Another arrow points from β_3 in the equation to the second curve, which is labeled β_3 . A third arrow points from ε in the equation to the third curve, which is labeled σ .

Bayesian Multiple Imputation

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Bayesian Multiple Imputation

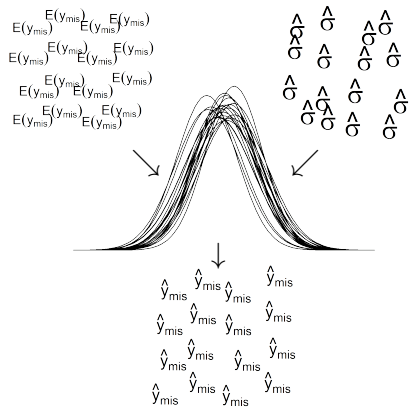
Step 2:

$$\mathbb{E}(Y_{mis}) = \hat{\beta}_0 + \hat{\beta}_1 X_{1,mis} + \hat{\beta}_2 X_{2,mis} + \hat{\beta}_3 X_{3,mis} + \dots$$

Diagram illustrating the expectation of the imputed value $\mathbb{E}(Y_{mis})$ as a linear combination of estimated coefficients and observed values. The coefficients $\hat{\beta}_0, \hat{\beta}_1, \hat{\beta}_2, \hat{\beta}_3$ are shown with multiple instances above them, indicating they are estimates. The observed values $X_{1,mis}, X_{2,mis}, X_{3,mis}$ are shown with multiple instances above them, indicating they are observed values. The expectation $\mathbb{E}(Y_{mis})$ is shown with multiple instances below it, indicating it is the expected value of the imputed value.

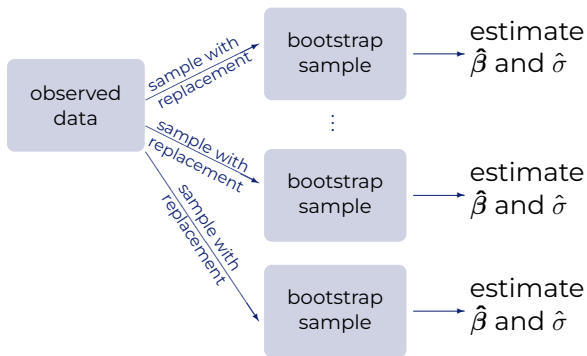
Bayesian Multiple Imputation

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

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

\downarrow \downarrow
 $\hat{\boldsymbol{\beta}}$ $\hat{\sigma}$

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

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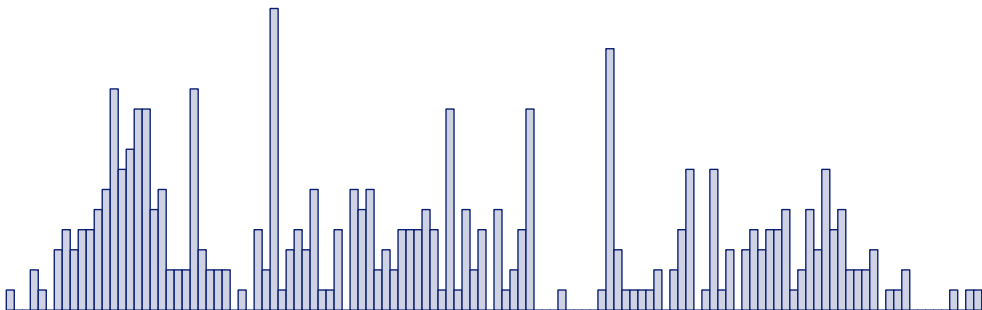
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- ▶ Imputed values are sampled from $p(\mathbf{y}_{mis} | \mathbf{X}_{mis}, \hat{\boldsymbol{\beta}}, \hat{\sigma})$ (step 2).
- ➔ Step 2 is analogous to step 3 in Bayesian multiple imputation.

Semi-parametric Imputation

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

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



Semi-parametric Imputation

Predictive Mean Matching (PMM)

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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 observed and unobserved cases are compared.

Semi-parametric Imputation

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

$$\hat{\mathbf{y}}_{obs} = \mathbf{X}_{obs}\hat{\beta}$$

3. Calculate the predicted value for the missing cases

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

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.

Semi-parametric Imputation

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

► **Case with the smallest absolute difference**

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

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

Semi-parametric Imputation

Potential issues with donor selection

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 - ▶ **predictor variables** are strongly **related to the missingness.**
- ▶ 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.

Semi-parametric Imputation

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 $\tilde{\beta}_{B/BS}$ 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.

Semi-parametric Imputation

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

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