Incomplete Data Analysis

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- → In practice, missing data may occur in several variables and the goal of this week's lecture
 is to learn about how to deal with this case in a multiple imputation context.
- → As we have seen back in week 4, when we superficially covered multivariate missingness
 in a single imputation context, there are two main routes:
 - \hookrightarrow Joint model imputation.
 - → Multiple imputation by chained equations/fully conditional specification.

Joint model approach

- → The joint model approach is the original approach for multiple imputation of multivariate missingness and it rests on specifying a multivariate (regression) model for all the variables that contain missing values.
- The main problem with this approach is that it is not always easy to set up a reasonable multivariate regression model, and this may be particularly complicated if the variables containing missing values are of mixed type (e.g., continuous and binary/categorical variables).
- → However, software exists to fit such models (e.g., norm, cat, mix, and jomo) and there are still new techniques within this line of multiple imputation being developed.

- → As we can guess by the name, this is the approach implemented by the R package mice when several variables have missing values.
- → The method requires the specification of a regression imputation model for each incomplete variable and creates imputations per variable in an iterative fashion.

Multiple imputation by chained equations

→ The following scheme illustrates the main idea behind the MICE approach.

$$\hookrightarrow Y_1 \sim Y_2 + Y_3 + Y_4 + \dots$$

$$\hookrightarrow \ Y_2 \sim Y_1 + Y_3 + Y_4 + \ldots$$

$$\hookrightarrow \ Y_3 \sim Y_1 + Y_2 + Y_4 + \ldots$$

$$\hookrightarrow Y_4 \sim Y_1 + Y_2 + Y_3 + \dots$$

- Under this imputation scheme, we can easily deal with variables of mixed types and we reduce the problem of multivariate imputation to several univariate imputation problems.
- The options that we have available in mice for each univariate regression model depends
 on the nature of the variable being imputed and we have already covered them last week.

- → Let us use the toy example from the previous slide to describe how MICE works. The
 iterative procedure is as follows:
 - → Start with initial guesses for the missing values (e.g., random draws from the observed data of the respective variable).
 - Update/impute Y₁ based on Y₂, Y₃, Y₄, and the other fully observed variables (in case they exist). Note that at this stage Y₂, Y₃, and Y₄ still have the missing values set to the initial guesses.
 - \hookrightarrow Update/impute Y_2 based on Y_1 , Y_3 , Y_4 , and the other fully observed variables. Note that now for the missing values on Y_1 we are using the updated values from the previous step, while for Y_3 and Y_4 the missing values are still set to the initial guesses.
 - $\hookrightarrow \dots$
 - Update/impute Y₁ again based on updated Y₂, Y₃, and Y₄ and on the values of the fully observed variables.
 - \hookrightarrow Repeat until convergence.



- The imputed values for Y₁, Y₂, Y₃, Y₄ from the last iteration of the algorithm are then used to replace the missing data in the original dataset.
- → Note that one run through the algorithm until convergence leads to one imputed dataset.
- \hookrightarrow We need to repeat the procedure M times to impute values in all M copies of the dataset that we have created.

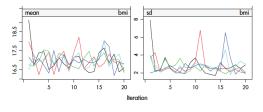
- → MICE is a Markov chain Monte Carlo (MCMC) method. More specifically, it is based on the idea of Gibbs sampling.
- → MCMC? Gibbs sampling?
- → MCMC is a technique that allows drawing samples from a complex distribution.
- → MCMC works by creating a chain of random variables (a Markov chain)! The distribution that each element in the chain is sampled from depends on the values of the previous element.
- Under certain conditions, the chain eventually stabilizes (i.e., converges to the stationary distribution).
- It can be proved that samples from the chain are then samples from our complex distribution of interest.

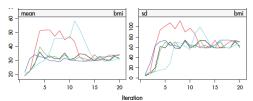
- → The sequence of imputed values for each missing value is called a (Markov!) chain.
- → The reason for why we need to iterate is because the imputed values in one variable
 depend on the imputed values of the other variables and to start the procedure we use
 random draws from the observed data for each particular variable.
- It is not unreasonable to think that these draws are far from the actual distribution and so the first draws may not be draws from the distribution of interest.

- → Note that we have several variables with missing values and each of these may contain several missing values, and therefore at the end of the iterative procedure we have a potential large number of chains of imputed values.
- → Then, because we need to repeat the procedure for all M copies of the dataset, we have a
 potential very large number of chains of imputed values multiplied by M.
- → To monitor convergence is thus infeasible to look at all produced chains.
- → Alternatively, as a way to reduce the number of chains we have to look at, and as implemented in mice, we could look at the chains of the mean and/or standard deviation (or other summary statistic) of the imputed values per variable.
- → Fortunately, as pointed by van Buuren, the number of iterations to reach convergence is typically much lower than that required in typical modern MCMC applications. The default in mice is 5, but it is quite difficult to judge convergence based on a so low number of iterations and, in practice, 20 or 30 iterations are often recommend.

Multiple imputation by chained equations

→ Figures from van Buuren (2018, p. 188 and 189). The top row shows the mice output for a
healthy convergence case, while the second row shows a traceplot that indicates non
convergence.





- → It is interesting to note that Gibbs sampling exploits the fact that a multivariate joint distribution is fully determined by its full univariate conditional distributions.
- → But MICE imputation models work the other way around. The univariate imputation models (= univariate full conditional distributions) are specified directly and there is no guarantee that a corresponding joint distribution exists.
- → This leads us to the concept of **compatibility**.
- \hookrightarrow To conditional densities $f(Y_1 \mid Y_2)$ and $f(Y_2 \mid Y_1)$ are said to be compatible if a joint distribution $f(Y_1, Y_2)$ exists that has $f(Y_1 \mid Y_2)$ and $f(Y_2 \mid Y_1)$ as its conditional densities.
- → So, the main problem of MICE is that there is no guarantee that the distributions implied by
 the different univariate imputation regression models are compatible.

- → As stated by van Buuren (2018, p. 123): "though MICE is only guaranteed to work if the conditional are compatible, these simulations suggest the results may be robust against violations of compatibility."
- van Buuren also concluded the discussion on this topic with the following: "Apart from potential feedback problem, it appears that incompatibility seems like a relatively minor problem in practice, especially if the missing data rate is modest and the imputation models fit the data well."

- → Remember that compatibility refers to the property that the conditionally specified models together specify some joint distribution from which imputations are to be drawn. It is a theoretical requirement of the Gibbs sampler.
- → Congeniality refers to the relation between the substantive model (step 2) and the imputation model.
- \hookrightarrow It is widely accepted that the imputation model should be more general than the substantive model.
- Further, interactions, quadratic terms, etc, appearing in the substantive model should be preserved in the imputation model.

- \hookrightarrow To finish we should keep in mind the assumptions made by MICE:
 - → The missing mechanism is MAR (or MCAR).
 - \hookrightarrow All associations between variables are linear.
 - \hookrightarrow Compatibility and congeniality.