

Practical 10. Missing data

Wednesday, 22 June 2022

[Lecture 10](#)[PDF version](#)

Missing data

The file `missing_data.rds` contains a dataset with individual level data on two competing treatments. The data include $n_1 = 75$ and $n_2 = 84$ records for suitable measures of effectiveness (QALYs, e_i) and total costs (c_i) for each individual. In addition, data on the baseline utility u_i are also recorded for each individual. The data are stored in the `list` format. The file `MissingData.R` is an R script to guide you through the analysis.

- Following the script, load the data and inspect the list; for instance, you can use the command `hist` to produce a histogram of the relevant variables, e.g. `hist(data$c[[1]])` (notice that you need to subset the elements of the object `data`, because they are part of a `list`).
- Inspect the file `Normal_Normal.txt`, which codes up the bivariate Normal model. Make sure you can follow the code and match it with the slides from Lecture 10.
- Following the script, run the model assuming a marginal Normal model for the effectiveness (controlling for the baseline utility) and a conditional Normal model for the costs (given the effectiveness~variable).
- Following the script, feed the output from the Bayesian model run in `BUGS` to `BCEA`, to perform the economic analysis.
- Following the script, check the distribution of the imputed values — for example, you can consider the first individual with missing data and the variable of effectiveness as in the script, but you can explore other variables too.
- Now inspect the file `Beta_Gamma.txt`, which encodes a Beta marginal model for the QALYs and a Gamma conditional model for the costs, given the effectiveness. Make sure you can follow the code and match it with the slides in the lecture.
- Following the script, replicate the analysis using this second model and then perform the economic evaluation in `BCEA`. Check that in this case, the imputed values are within the natural range of the underlying variables (specifically, for the QALYs).

