

# EP16: Missing Values in Clinical Research: Multiple Imputation

## 12. Imputation with Longitudinal Data

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# The Challenge with Longitudinal Data

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- ▶ (potential) correlation between repeated measurements ignored
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**Simple summaries** to allow wide format:

- ▶ loss of information
- ▶ potential MNAR
- ▶ **bias**

# Imputation with mice

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**mice** has functions to allow imputation of longitudinal (2-level) data:

- ▶ **Level 1:**  
repeated measurements within subjects or subjects within classes
- ▶ **Level 2:**  
time-constant/baseline covariates, between subjects effects, variables on the group level

Imputation methods for **level-1** variables:

- ▶ `2l.pan`
- ▶ `2l.norm`
- ▶ `2l.lmer`
- ▶ `2l.bin`

Imputation methods for **level-2** variables:

- ▶ `2lonly.norm`
- ▶ `2lonly.pmm`
- ▶ `2lonly.mean`

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- ▶ `2lonly.norm` and `2lonly.pmm`: to impute level-2 variables
- ▶ `2lonly.mean`: imputes values with the mean of the observed values per class (only to be used in special cases!)

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The `predictorMatrix` contains extra info for multi-level imputation:

- ▶ grouping/ID variable: -2
- ▶ random effects (also included as fixed effects): 2
- ▶ fixed effects of group means: 3
- ▶ fixed effects of group means & random effects: 4

In all cases, the group identifier (“id” variable) needs to be set to -2 in the `predictorMatrix`.

# Imputation with mice

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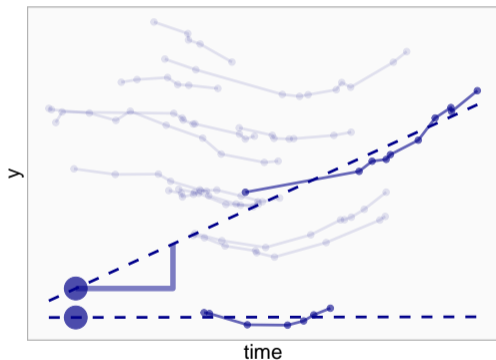
**Alternative approach:** (Erlor et al. 2016)

Get a **better summary** of the longitudinal variables!

# Imputation with mice

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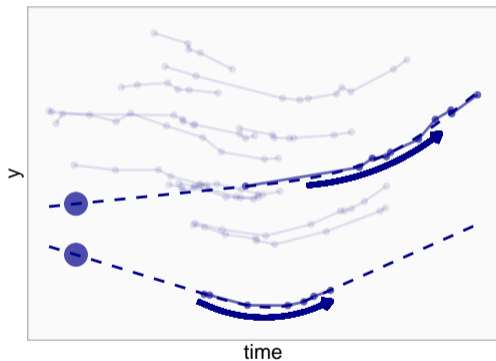
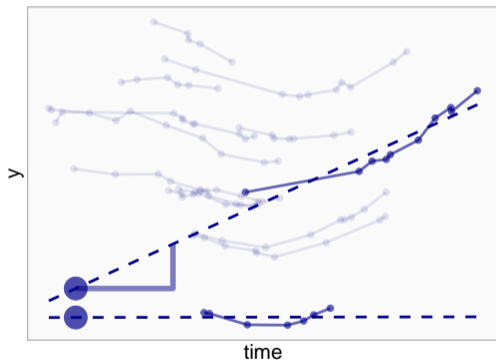
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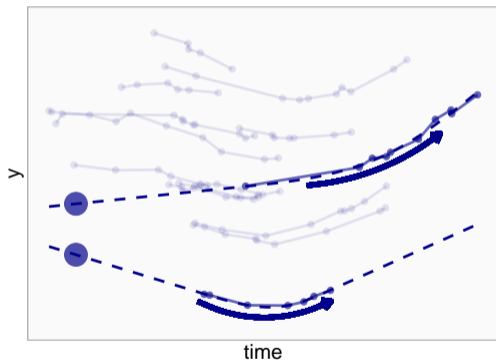
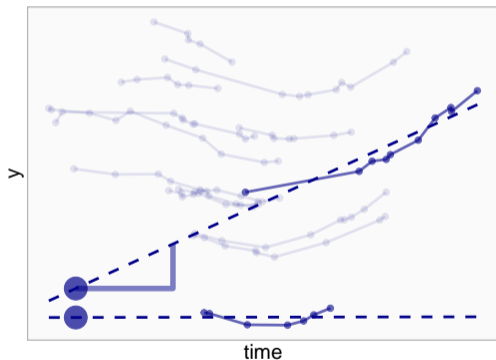
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**Approximate trajectories using random effects!**

## Imputation with mice

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**Drawback:** cannot handle **incomplete longitudinal variables**.

# Imputation with JointAI

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Example data:

- ▶  $x_1$  (complete)
- ▶  $x_2$  (binary, 30% NA)
- ▶  $x_3$  (3 categories, 30% NA)
- ▶  $x_4$  (continuous/normal, 30% NA)
- ▶  $y$  (longitudinal outcome)
- ▶  $time$  (time variable with quadratic effect)
- ▶  $id$  (id variable)

# Imputation with JointAI

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The syntax for analysing mixed models in **JointAI** is analogous the syntax used in `lme()` of the package **nlme**.

```
library("JointAI")
JointAI_long <- lme_imp(y ~ x1 + x2 + x3 + x4 + time + I(time^2),
  random = ~time|id, data = longDF2,
  n.iter = 5000)
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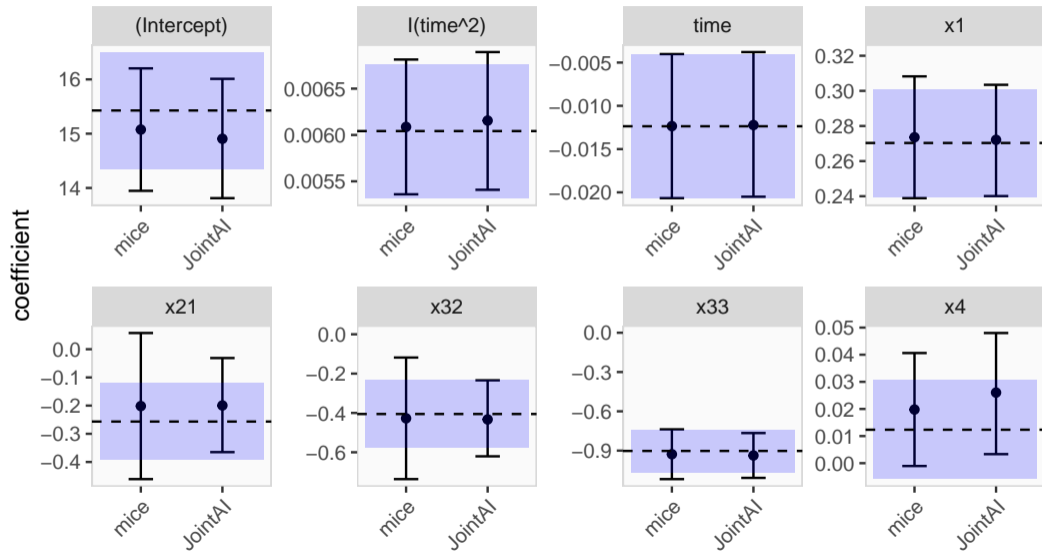
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Again, convergence of the Gibbs sampler needs to be checked before obtaining the results.

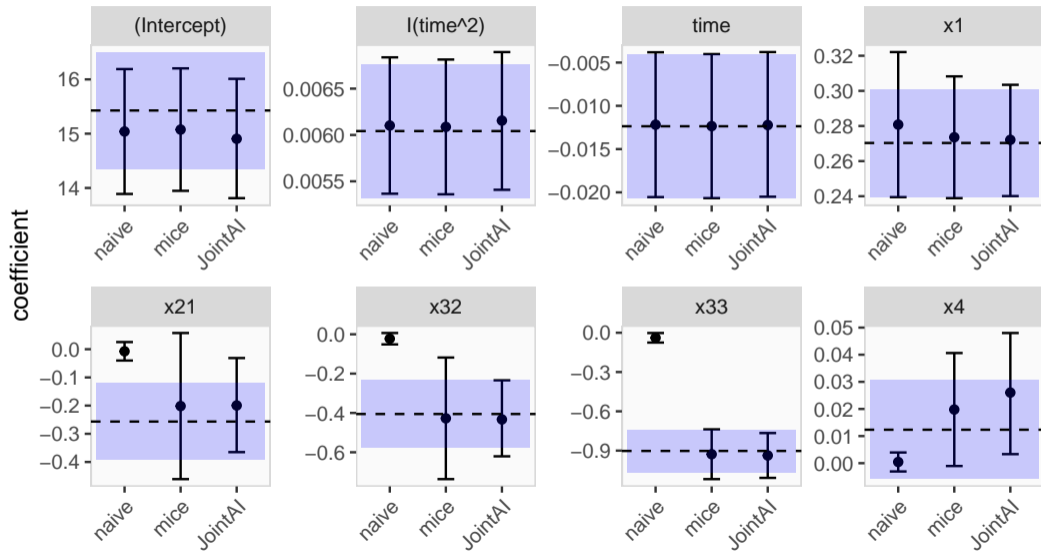
Contrary to the two-level imputation of **mice**, non-linear associations are appropriately handled.



# Comparison of Results



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## References

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Erler, Nicole S, Dimitris Rizopoulos, Joost van Rosmalen, Vincent WV Jaddoe, Oscar H Franco, and Emmanuel MEH Lesaffre. 2016. "Dealing with Missing Covariates in Epidemiologic Studies: A Comparison Between Multiple Imputation and a Full Bayesian Approach." *Statistics in Medicine* 35 (17): 2955–74. <https://doi.org/10.1002/sim.6944>.

Schafer, Joseph L, and Recai M Yucel. 2002. "Computational Strategies for Multivariate Linear Mixed-Effects Models with Missing Values." *Journal of Computational and Graphical Statistics* 11 (2): 437–57.