

How black-box use of imputation can cause bias

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Handling Missing Values is Easy!

Functions automatically exclude missing values:

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## [...]  
## Residual standard error: 2.305 on 69 degrees of freedom  
## (25 observations deleted due to missingness)  
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Imputation is super easy:

```
library("mice")  
imp <- mice(mydata)
```

However ...

Handling Missing Values Correctly is Not So Easy!

(Imputation) methods makes certain **assumptions**, e.g.:

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 - no interactions
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violation → bias

Several authors have

- investigated robustness to mis-specification (of distribution)
 - in MI using FCS / MICE
 - in joint model MI
- and/or proposed to use
 - Tukey's gh distribution
 - Fleishman polynomials
 - GAMs (in FCS)
 - Doubly-robust weighted estimating equations (instead of MI)

Joint distribution

$$\underbrace{p(y | X, b, \theta)}_{\substack{\text{analysis} \\ \text{model}}} \quad \underbrace{p(X | \theta)}_{\substack{\text{imputation} \\ \text{part}}} \quad \underbrace{p(b | \theta)}_{\substack{\text{random} \\ \text{effects}}} \quad \underbrace{p(\theta)}_{\text{priors}}$$

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Imputation part

$$p(\overbrace{x_1, \dots, x_p}^X, X_{\text{compl.}} | \theta) = p(x_1 | X_{\text{compl.}}, \theta) \\ p(x_2 | X_{\text{compl.}}, x_1, \theta) \\ p(x_3 | X_{\text{compl.}}, x_1, x_2, \theta) \\ \dots$$

Joint distribution

$$\underbrace{p(y | X, b, \theta)}_{\text{analysis model}} \quad \underbrace{p(X | \theta)}_{\text{imputation part}} \quad \underbrace{p(b | \theta)}_{\text{random effects}} \quad \underbrace{p(\theta)}_{\text{priors}}$$

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Software

Implemented in the **R** package **JointAI**

Imputation in MICE

$$p(x_1 \mid y, X_{\text{compl.}}, x_2, x_3, x_4, \dots, \theta)$$

$$p(x_2 \mid y, X_{\text{compl.}}, x_1, x_3, x_4, \dots, \theta)$$

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No issues with

- complex outcomes, e.g.:
 - multi-level
 - survival
- congeniality
- compatibility

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Potential mis-specification of

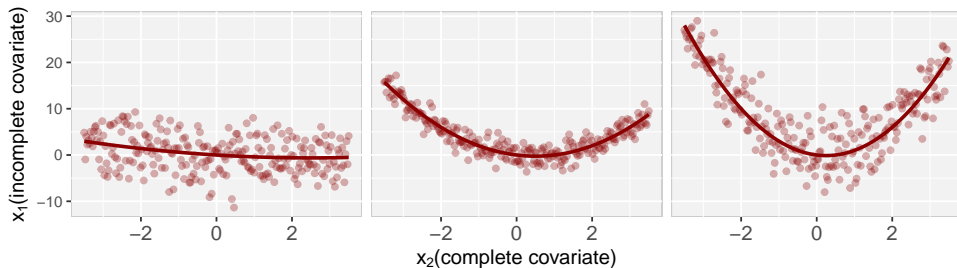
- association structure
- conditional distribution
- M(C)AR



Simulation Study: Quadratic Effect

Analysis model: $y \sim \beta_0 + \beta_1 x_1 + \beta_2 x_2 + \dots$

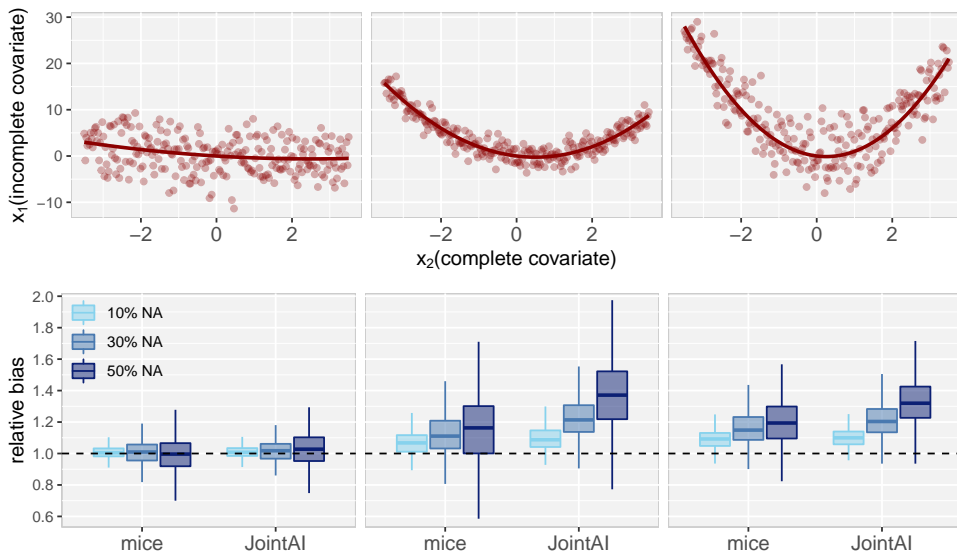
Quadratic association between covariates: $x_1 \sim \alpha_0 + \alpha_1 x_2 + \alpha_2 x_2^2 + \dots$



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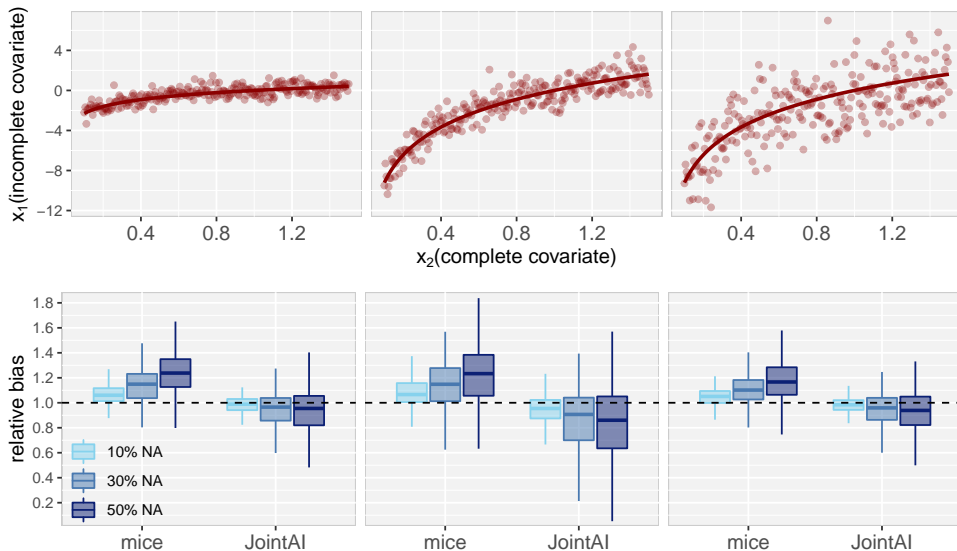
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Simulation Study: Logarithmic Effect

Analysis model: $y \sim \beta_0 + \beta_1 x_1 + \beta_2 x_2 + \dots$

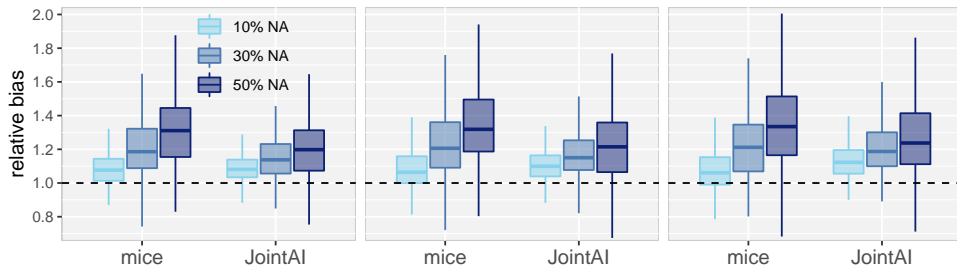
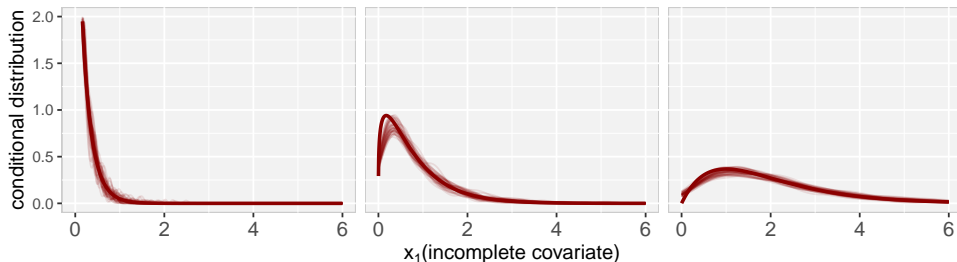
Log-association between covariates: $x_1 \sim \alpha_0 + \alpha_1 \log(x_2) + \dots$



Simulation Study: Gamma distribution

Analysis model: $y \sim \beta_0 + \beta_1 x_1 + \beta_2 x_2 + \dots$

Gamma-distributed covariate: $x_1 \mid x_2, x_3, \dots \sim Ga()$



We need more flexible imputation models!

Ideally: models that fit (almost) any distribution / association structure.

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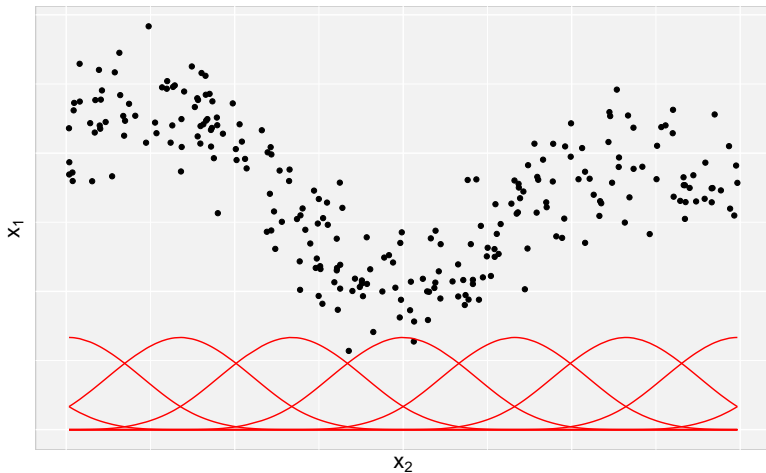
Ideally: models that fit (almost) any distribution / association structure.

Ideas:

- flexible **association** structure: **penalized splines**
- flexible residual **distribution**: **mixture of Polya-Trees**

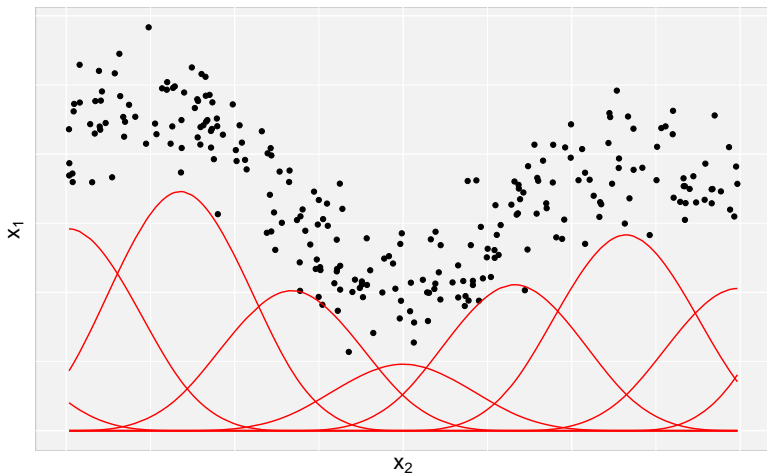
Bayesian P-Splines

Instead of $\beta_1 x_2$ we use $\sum_{\ell=1}^d \beta_{\ell} B_{\ell}(x_2)$:



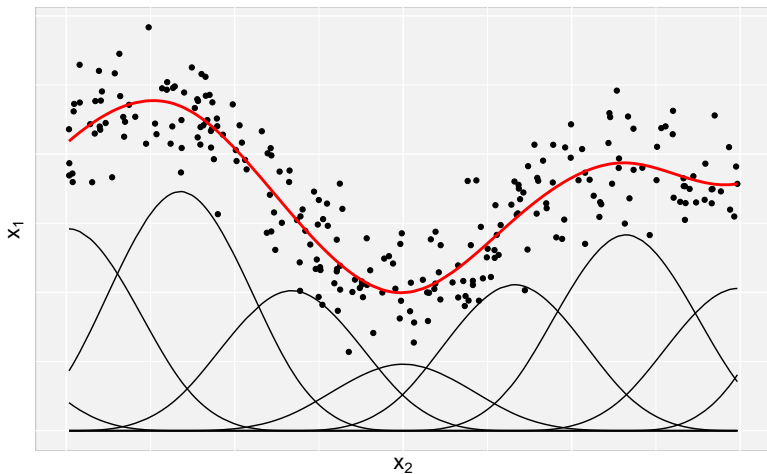
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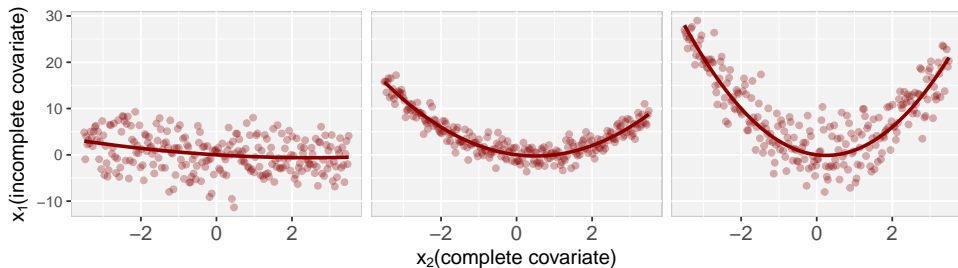
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Simulation: Bayesian P-Splines

Analysis model: $y \sim \beta_0 + \beta_1 x_1 + \beta_2 x_2 + \dots$

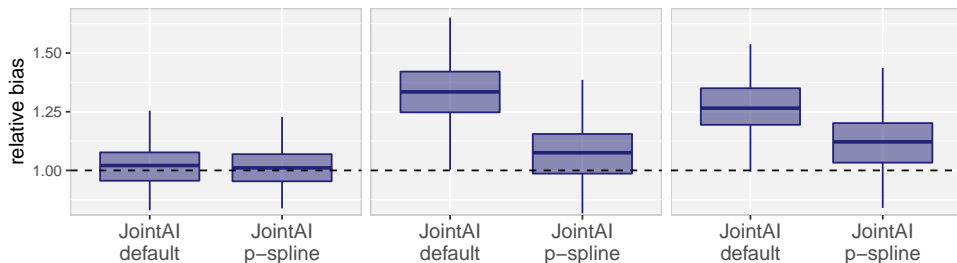
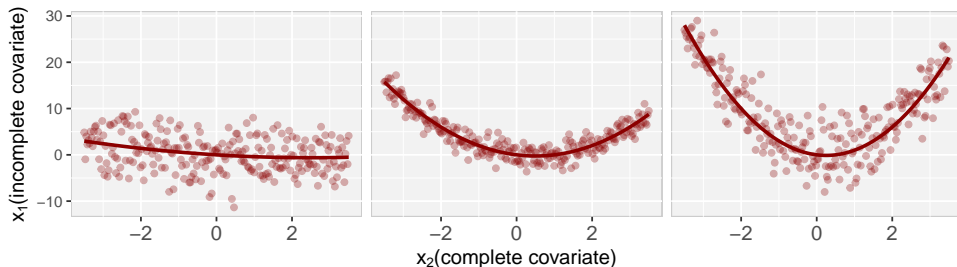
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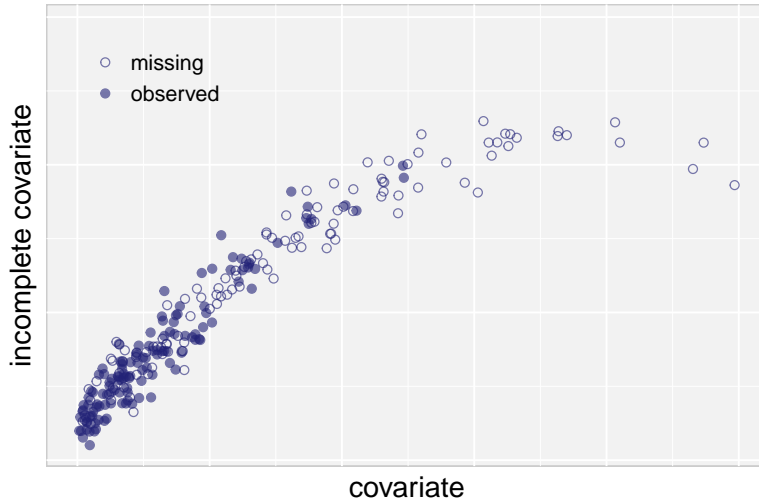
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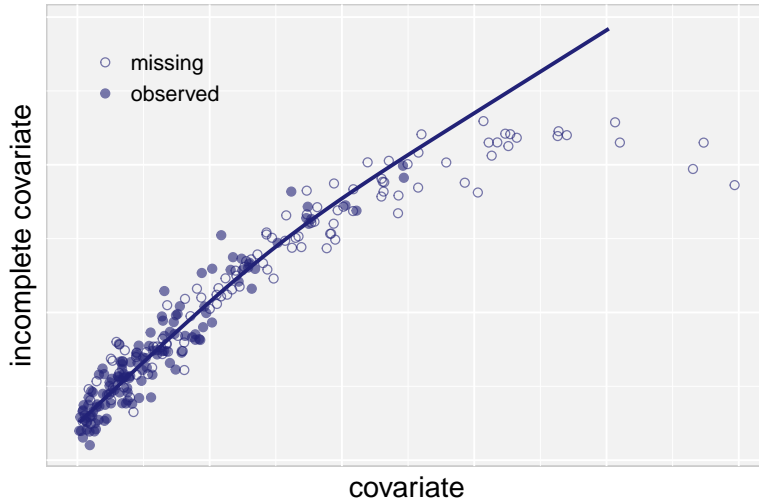
Simulation: Bayesian P-Splines

Potential Issue:

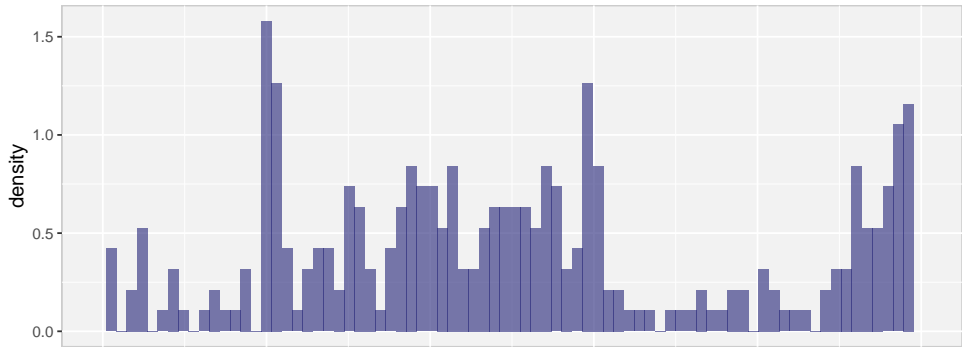


Simulation: Bayesian P-Splines

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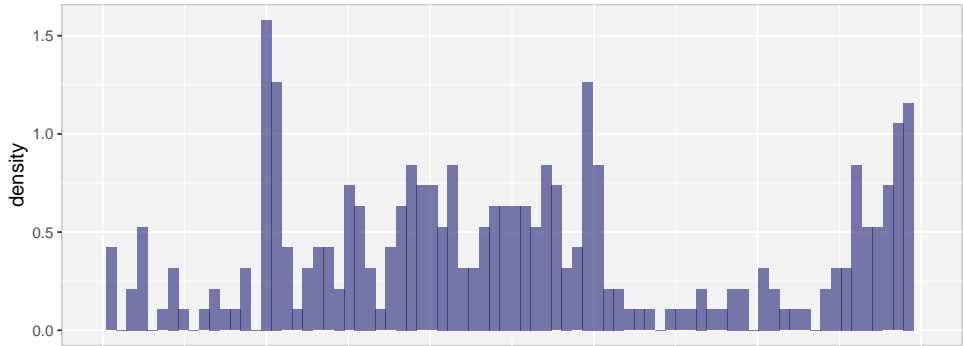


Mixture of Polya Trees



$\text{Beta}(a_0, a_0)$

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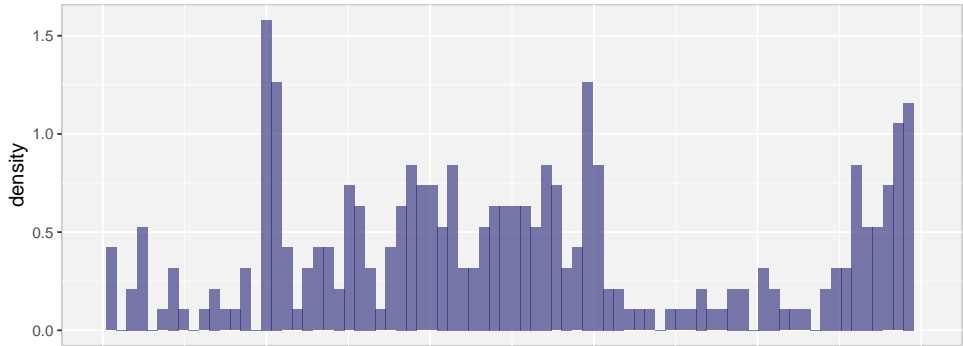


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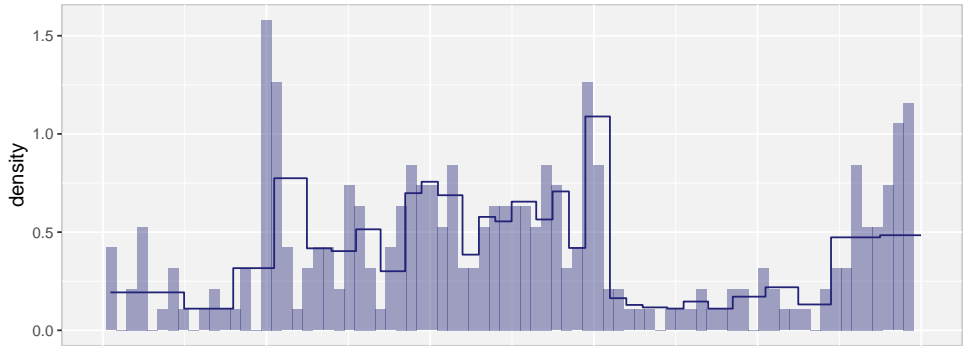
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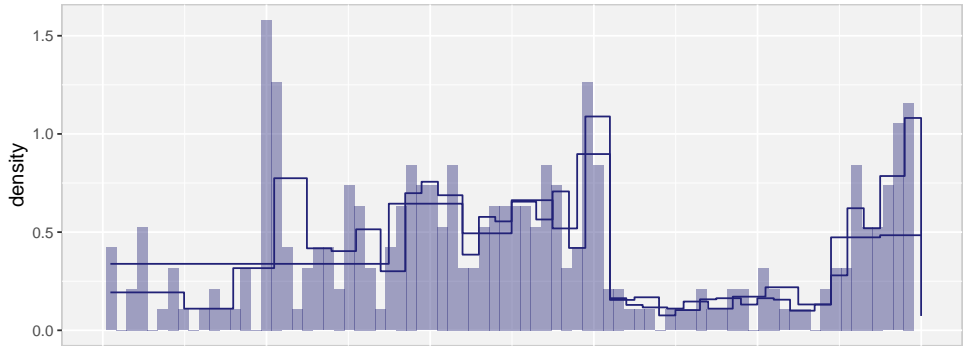
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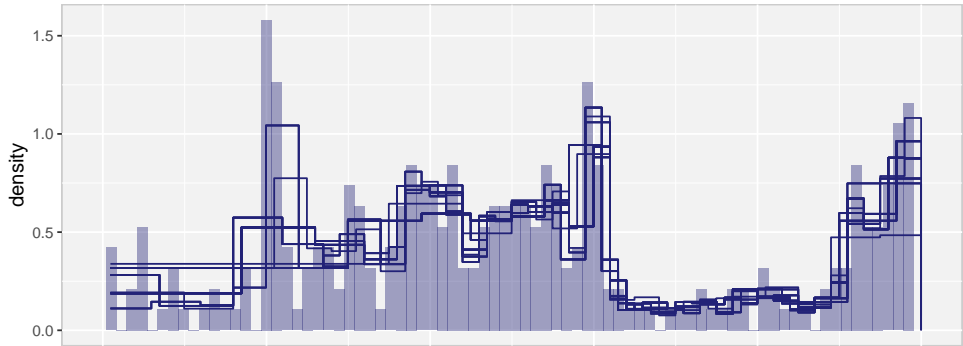
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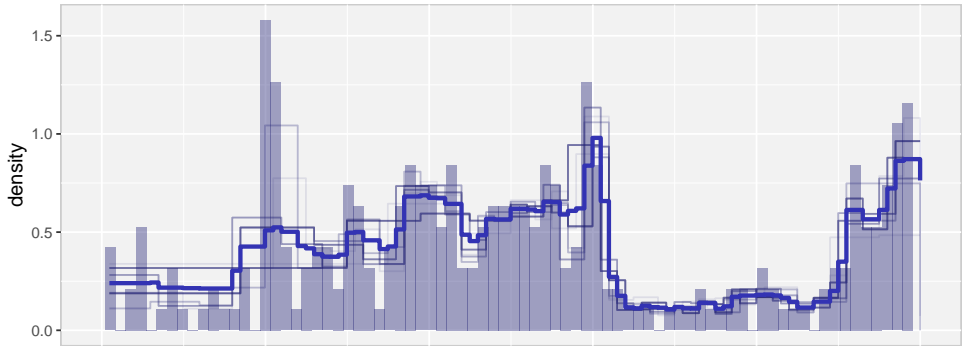
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 - χ^2 type of tests
 - Kolmogorov-Smirnoff test?
 - discordance tests?

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 - χ^2 type of tests
 - Kolmogorov-Smirnoff test?
 - discordance tests?
- feasibility checks before running the complex model

Take home message

- **assumptions** of imputation models can easily be **violated** ➔ **bias**
- **more flexible** imputation models are **needed**
- semi- / **non-parametric (Bayesian) methods** can offer a **solution**
- more **flexibility** ➔ more **complexity** ➔ need for **guidance** tools

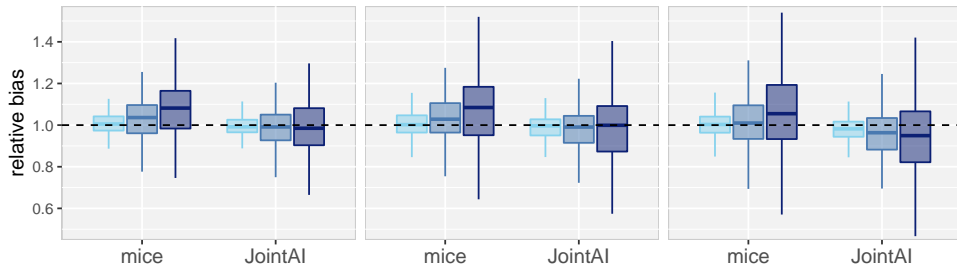
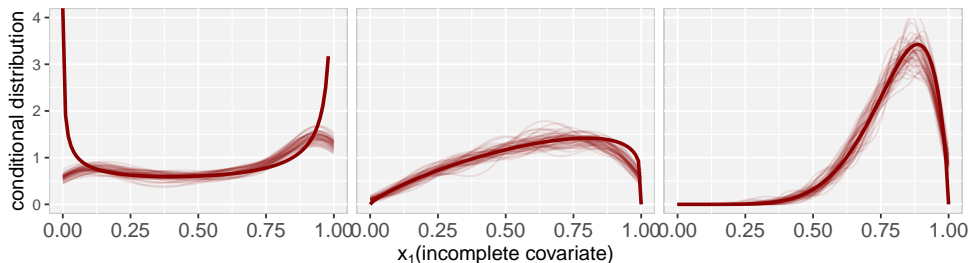
Thank you for your attention.

 n.erler@erasmusmc.nl
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 www.nerler.com

Simulation Study: Beta Distribution

Analysis model: $y \sim \beta_0 + \beta_1 x_1 + \beta_2 + \dots$

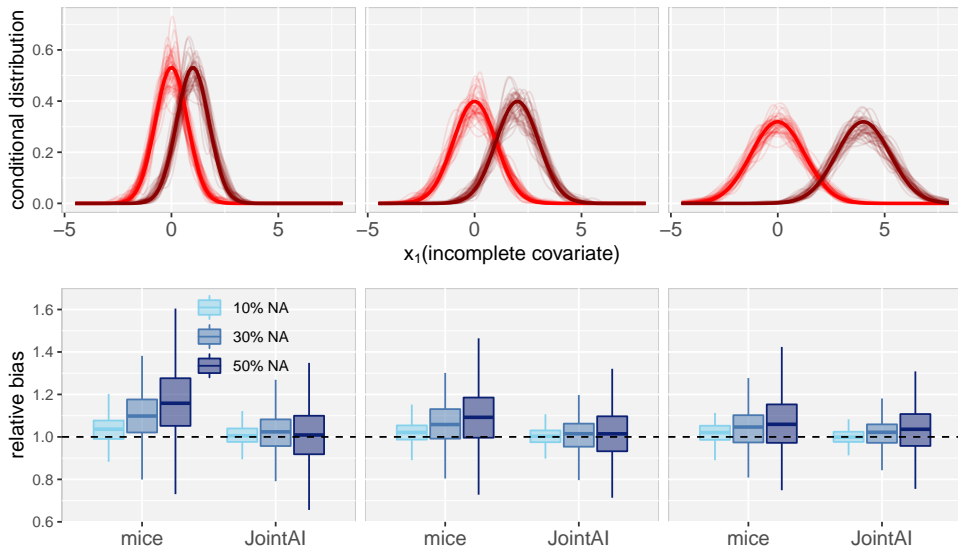
Beta-distributed covariate: $x_1 \mid x_2, x_3, \dots \sim Be()$



Simulation Study: Reverting the sequence

Analysis model: $y \sim \beta_0 + \beta_1 x_1 + \beta_2 x_2 + \dots$

Exclusion of important predictor: $x_1 \sim \alpha_0 + \alpha_1 x_2 + \alpha_2 x_3 + \dots$



Simulation Study: Ignoring an Interaction

Analysis model: $y \sim \beta_0 + \beta_1 x_1 + \beta_2 x_2 + \dots$

Interaction between covariates: $x_1 \sim \alpha_0 + \alpha_1 x_2 + \alpha_2 x_3 + \alpha_3 x_2 x_3 + \dots$

