# EP16: Missing Values in Clinical Research: Multiple Imputation

# 11. Imputation with Non-linear Functional Forms

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For imputation of variables that have non-linear associations

- ▶ PMM often works better than imputation with a normal model,
- ▶ the Just Another Variable approach can reduce bias in interactions,
- passive imputation
- quadratic can help to impute variables with quadratic association.

For demonstration, we use a simulated example dataset DFnonlin:

- y continuous outcome
- x continuous (normal) covariate (50% missing values MCAR)
- z binary covariate (complete)

#### We assume a

- **quadratic effect** of x on y, and
- ▶ an interaction between x and z

```
head(DF_nonlin)
```

```
## y x z
## 1 -0.4002016 -0.42298398 1
## 2 0.7883355 -1.54987816 0
## 3 0.1900922 -0.06442932 0
## 4 0.3321608 0.27088135 0
## 5 4.6146593 1.73528367 0
## 6 0.3705739 NA 0
```

```
dim(DF_nonlin)
```

## [1] 200

## Imputation with mice: JAV

#### Just Another Variable (JAV) approach:

- pre-calculate the non-linear form (or interaction term) in the incomplete data,
- > add it as a column to the dataset, and
- impute it as if it was just another variable.

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```
DF2 <- DF_nonlin  # copy of the data, only for this example
DF2$xx <- DF2$x^2  # pre-calculate the quadratic term

DF2$xz <- DF2$x * DF2$z  # pre-calculate the interaction
```

```
# JAV imputation (using pmm and full predictor matrix)
impJAV <- mice(DF2, maxit = 20, printFlag = FALSE)</pre>
```

#### Imputation with mice: JAV

To **relax the assumption** of linear associations even more, we could introduce **additional interactions** with the outcome.

In this example, we can add an interaction between z and y:

```
# JAV imputation with additional interaction
impJAV2 <- mice(DF3, maxit = 20, printFlag = FALSE)</pre>
```

## Imputation with mice: Passive Imputation

Alternative: impute all non-linear terms and interactions passively:

```
# adapt the imputation method (we re-use the vector from impJAV2 here) meth_passive <- impJAV2$method meth_passive[c("xx", "xz", "yz")] <- c("~I(x^2)", "~I(x*z)", "~I(y*z)")
```

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```
# adapt the predictor matrix (we re-use the matrix from impJAV2 here)
pred_passive <- impJAV2$predictorMatrix
pred_passive['x', 'xx'] <- 0
pred_passive[c('x', 'z'), 'xz'] <- 0
pred_passive[c('y', 'z'), 'yz'] <- 0</pre>
```

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# Imputation with mice: Polynomial Combination

The imputation method quadratic uses the "polynomial combination" method to impute covariates that have a quadratic association with the outcome (Van Buuren 2012 pp. 139–141; Vink and van Buuren 2013).

- $\rightarrow$  ensure the **imputed values** for x and  $x^2$  are **consistent**
- $\rightarrow$  reduce bias in the subsequent analysis that uses x and  $x^2$

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```
# adapt the imputation method (we re-use the vector from impJAV here)
methqdr <- impJAV$meth
methqdr[c("x", "xx", "xz")] <- c("quadratic", "~I(x^2)", "~I(x*z)")</pre>
```

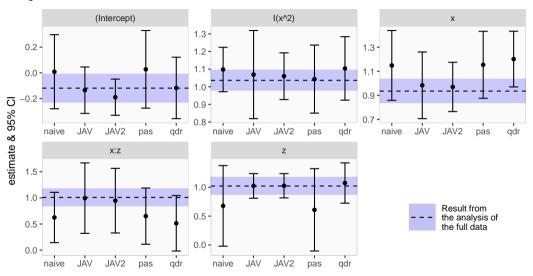
 $\rightarrow$  Here we use passive imputation for  $x^2$  and the interaction.

## Imputation with mice: polynomial combination

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For comparison, we also run a naive version (using defaults):

```
# naive imputation, using only y, x, z
impnaive <- mice(DF_nonlin, printFlag = FALSE)</pre>
```



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The syntax we use to analyse and impute the current example using **JointAI** is similar to the specification of a standard linear model using lm().

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Convergence of the Gibbs sampler can be checked using a traceplot.

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traceplot(JointAI_nonlin, ncol = 3, use_ggplot = TRUE)
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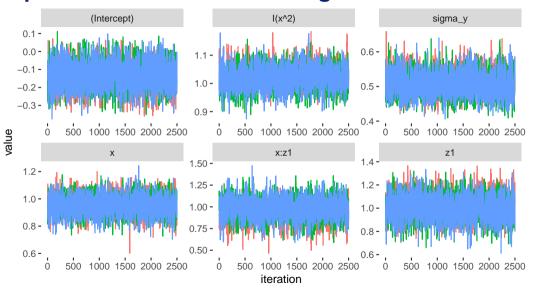
Convergence of the Gibbs sampler can be checked using a traceplot.

```
traceplot(JointAI_nonlin, ncol = 3, use_ggplot = TRUE)
```

Results (no separate analysis & pooling is necessary) can be obtained with the summary() function:

```
summary(JointAI_nonlin)
```

#### Imputation with JointAI: Convergence

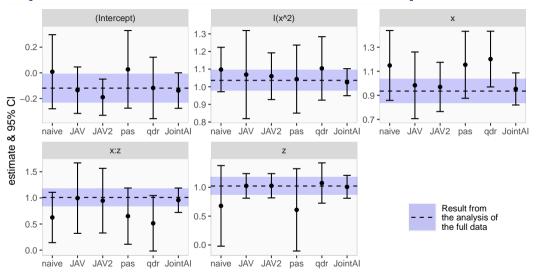


# Imputation with JointAI: Model Summary

```
##
   Linear model fitted with JointAI
##
## Call:
## lm_imp(formula = y \sim x * z + I(x^2), data = DF_nonlin. n.iter = 2500.
##
      seed = 1234)
##
## Posterior summary:
##
               Mean
                       SD 2.5% 97.5% tail-prob. GR-crit
## (Intercept) -0.138 0.0697 -0.276 0.000259
                                            0.0512
                                                     1.09
## x
          0.954 0.0683 0.820 1.086675 0.0000 1.02
## 21
      1.007 0.1005 0.810 1.207309 0.0000 1.10
## I(x^2) 1.026 0.0393 0.949 1.102465 0.0000 1.32
## x:z1 0.957 0.1189 0.722 1.188642
                                          0.0000 1.28
##
## Posterior summary of residual std. deviation:
##
          Mean
                  SD 2.5% 97.5% GR-crit
## sigma_y 0.507 0.0334 0.447 0.576 1.01
##
##
   [...]
```

## Imputation with JointAI: Model Summary

```
## [...]
##
## MCMC settings:
## Iterations = 101:2600
## Sample size per chain = 2500
## Thinning interval = 1
## Number of chains = 3
##
## Number of observations: 200
```



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#### polynomial combination

- consistent imputed values
- only available for quadratic association
- often numeric instabilities (warning messages)

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To use JointAI appropriately and to interpret the results correctly requires more knowledge about the underlying method than can be covered in this course.

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#### Note:

The example used here only serves to demonstrate the different approaches. We cannot use these results to conclude which approach works better in general.

#### References

Van Buuren, Stef. 2012. Flexible Imputation of Missing Data. Chapman & Hall/Crc Interdisciplinary Statistics. Taylor & Francis. https://stefvanbuuren.name/fimd/.

Vink, Gerko, and Stef van Buuren. 2013. "Multiple Imputation of Squared Terms." Sociological Methods & Research 42 (4): 598–607.