# EP16: Missing Values in Clinical Research: Multiple Imputation

## 5. Know Your Data

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To demonstrate the **work flow** when performing **multiple imputation** with the **mice** package, we use data from the National Health and Nutrition Examination Survey (NHANES).

There are several packages in  $\mathbf{R}$  that provide functions to investigate **the** missing data pattern.

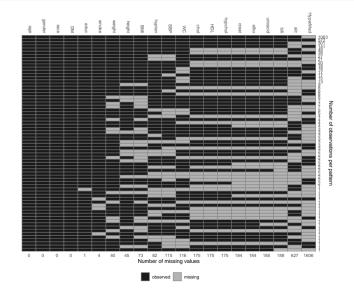
Examples are:

mice, JointAI, VIM, Amelia, visdat, naniar, ...

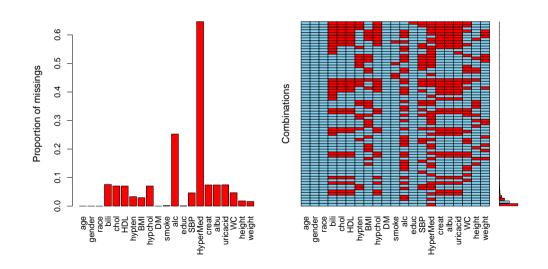
```
mdp <- mice::md.pattern(NHANES, plot = FALSE)
head(mdp[, -c(7:14)]) # omit some columns to fit it on the slide</pre>
```

```
tail(mdp[, -c(7:14)])
```

JointAI::md\_pattern(NHANES)



VIM::aggr(NHANES, prop = TRUE)



We are also interested in the number and proportion of (in)complete cases ...

... and the proportion of missing values per variable:

```
# NA % NA
                              # NA % NA
                                                           % NA
##
                     ##
                                           ##
                                                      # NA
           0.00
                               45 1.81
                                                      175 7.05
## age
                     ## height
                                           ## hypchol
## gender
           0.00
                     ## BMI
                               73 2.94
                                           ## creat
                                                      184 7.41
## race
           0.00
                               82 3.30
                                           ## albu
                                                      184 7.41
                     ## hypten
## DM
           0 0.00
                     ## SBP
                               115 4.63
                                           ## uricacid
                                                      185 7.45
## educ
           1 0.04
                     ## WC
                               116 4.67
                                           ## bili
                                                      188 7.57
## smoke 4 0.16
                     ## chol
                              175 7.05
                                           ## alc
                                                      627 25.25
## weight
         40 1.61
                     ## HDL
                               175 7.05
                                           ## HyperMed 1606 64.68
```

#### See also:

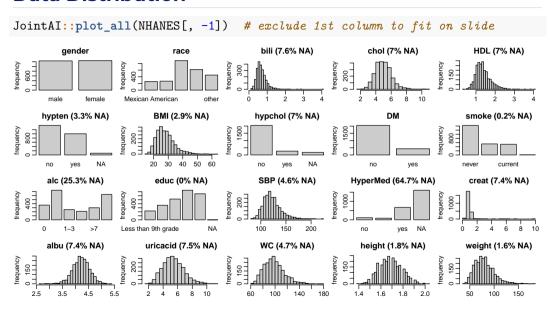
```
mice::md.pattern()
visdat::vis_miss()
visdat::vis_dat()
```

visuat..vis\_dat(

**•** ...

```
naniar::prop_miss_case(),
naniar::pct_miss_case()
naniar::prop_complete_case(),
naniar::pct_complete_case()
naniar::miss_var_summary()
mice::md.pairs()
```

#### **Data Distribution**



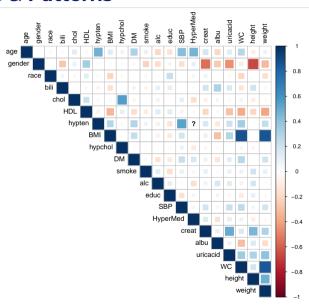
#### **Correlations & Patterns**

A quick (and dirty) way to check for strong correlations between variables is:

#### Note:

We only use the correlation coefficient for categorical variables for visualization, not as a statistical result!

## **Correlations & Patterns**



### **Correlations & Patterns**

Check out what the problem is with hypertension and HyperMed:

```
## HyperMed

## hypertension no previous yes <NA>

## no 0 0 0 1397

## yes 114 90 673 127

## <NA> 0 0 0 82
```

## Why are values missing?

Knowing your data also means being able to answer these questions:

- ▶ Do missing values in multiple variables always occur together? (e.g. blood measurements)
- ► Are there **structural missing values**? (e.g. pregnancy status in men)
- ▶ Are there **patterns** in the missing values? (e.g. only patients with hypertension have observations of HyperMed)
- ► Are values **missing by design**?
- ► Is the **assumption of ignorable missingness** (MAR or MCAR) justifiable?

## **Auxiliary Variables**

**Auxiliary variables** are variables that are not part of the analysis but **can help during imputation**.

#### Good auxiliary variables

- are related to the probability of missingness in a variable, or
- are related to the incomplete variable itself,
- do not have many missing values themselves and
- are (mostly) observed when the incomplete variable of interest is missing.

### **Your Turn!**

#### **Practical**

Getting to know your data html