# Handling Missing Data in Epidemiology Research Exploring Multiple Imputation as a Missing Data Strategy

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# Background



#### Introduction

Why is missing data important?

- ▶ Poor reporting, analysis, and handling of missing data [5]
- ▶ Impacts the results of a study
- Occurs everywhere

# Types of Missing Data

#### **MCAR**

Data is said to be missing completely at random if the probability of being missing is the same for all variables. [1]

#### **MAR**

Data is said to be missing at random if the probability of being missing is the same only within groups defined by the *observed* data. [1]

#### **MNAR**

Data is said to be missing not at random if the probability of being missing varies for unknown reasons and depends on unobserved values. [1]

### Data Context

- National Health and Nutrition Examination Survey (public dataset)
- Around 5,000 participants ages 0-80+ every two years
- NHANES oversamples to obtain **nationally representative** data (people aged 60 and over, African Americans, and Hispanic-identifying individuals)
- Constantly evolving to meet emerging health/nutrition needs [2]

Background

## Variables

For the purpose of this project, relevant variables were selected from NHANES 2018 data:

Variable	Description
age	Patient age in years
sex	Sex of patient $(M/F)$
race_ethnicity	Self identified patient race
combined_education	Patient education level
reference_education	Reference person education level
fipr	Federal income poverty ratio (0-5)
tap₋water	Amount of tap water drank, day 1, grams

#### Motivating Question

Can fipr help predict tap\_water consumption?

#### **Definition**

fipr: A ratio of family income to poverty guidelines, where a 1 represents a family at the federal poverty line. (Range from 0-5) [2]

#### Data

- ▶ Sample size: 8704
- ► Complete cases: 4056
- Notice by ommitting NA's, we lose 53% of our data
- Missing fipr cases: 1070
- ▶ For the scope of this project, we only had time to focus on missing data in fipr variable

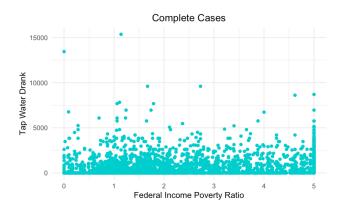
# Data Cleaning

- Created combined\_education variable
- Created new\_race\_ethnicity variable
- ▶ Recoded "don't know" or "refused" as NA values [2]

# Ad-hoc Solutions to Handling Missing Data

#### Complete Cases

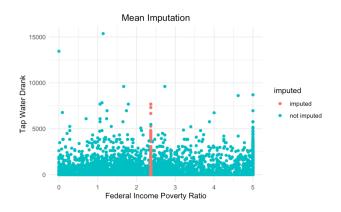
Also known as listwise deletion, the complete cases method omits all rows that contain a missing value in any category. [1]



# Ad-hoc Solutions to Handling Missing Data

## Mean Imputation

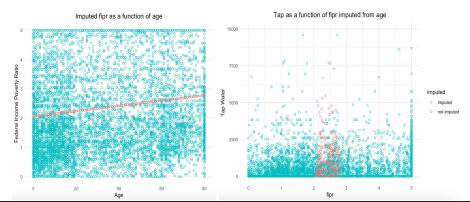
Replaces missing data values with the mean of that variable. [1]



# Ad-hoc Solutions to Handling Missing Data

### Regression Imputation

A model is built from the observed data first, then predictions for missing values are calculated under the fitted model. The missing values are assigned the "most likely" values on the regression line. [1]



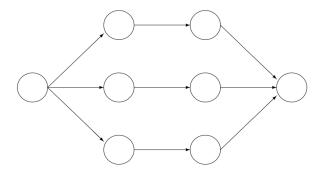
## Research Question

How can multiple imputation be used to handle missing data?

# Overview of Multiple Imputation

- ▶ MI deals with the uncertainty of the imputations themselves by imputing *several* values for each missing case (*g* between 5-10) [4]
- Imputed datasets are identical for observed data entries and differ in imputed values
- ▶ Imputed values are averaged into a pooled dataset

## Visualization



Incomplete data Imputed data Analysis results Pooled result

Flexible Imputation of Missing Data, van Buuren, 2018.

ground NHANE

# Multiple Regression

We begin with a multiple regression model:

$$Y = \beta_0 + \beta_1 x_1 + \beta_2 x_2 + \dots + \beta_k x_k + E$$

#### Where

Background

- Y is first fipr, and later tap\_water
- $\triangleright$   $\beta_0$  is the intercept, or constant
- $\triangleright$   $\beta_1, \beta_2, ..., \beta_k$  are the coefficients of our predictor variables
- $\triangleright$   $x_1, x_2, ... x_k$  are the predictor variables
- E is the random error term that represents the part of Y that cannot be explained by the predictor variables [3]

# Model Prediction using R Statistical Software

We use multiple linear regression to predict a model for fipr based on the complete cases of the following variables:

#### Random Noise

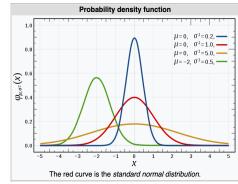
We impute our dataset g times, adding randomly generated noise E to Y based on the standard deviation of the residuals in fipr\_model that assumes a normal distribution:

$$\textit{E} \sim \textit{N}\left(0, \sigma^2\right)$$

based on n-m complete cases:

$$\sigma^2 = \frac{\sum_{i=m+1}^{n} \left(Y - \hat{Y}\right)^2}{n - m}$$

# Normal distribution



"Normal Distribution." Wikipedia.

# fipr Models

We fit 5 models to impute the missing fipr values as follows:

▶ If fipr is missing, replace NA value with

$$\beta_0 + \beta_1 x_1 + \dots + \beta_{11} x_{11} + E_i$$

Where

- $\beta_1, \beta_2, ..., \beta_{11}$  are the regression coefficients
- $x_1, x_2, ... x_{11}$  are the indicator variables
- E is the random noise
- ▶ Else, use given fipr

Note: the pmin()pmax() functions were used to adjust negative fipr values to 0, and fipr values > 5 to 5.

# Tap Models

Now we have 5 complete datasets across our education, race, age, sex, and fipr variables. We use these predictor variables to model tap water consumption.

### Coefficients and Standard Errors

#### From these models we aquire:

- $\blacktriangleright \ \beta_0^\ell, \beta_1^\ell, \beta_2^\ell, ..., \beta_k^\ell \ \text{for} \ \ell = 1...g$
- $\blacktriangleright$   $SE(\beta_0^\ell), SE(\beta_1^\ell), ..., SE(\beta_k^\ell)$  for  $\ell=1...g$

#### Where

- k represents each of our 11 predictor variables, and
- $\ell$  represents each of our 5 imputed datasets.

## Regression Coefficients

Averaging across our 5 imputed datasets produces point estimates of our regression coefficients...

$$ilde{eta}_j \equiv rac{\sum_{\ell=1}^{\mathcal{g}}eta_j^\ell}{oldsymbol{g}}$$

### Standard Errors

Standard errors of estimated coefficients are obtained by combining info about within and between imputation variation:

$$ilde{SE}\left( ilde{eta}_{j}
ight) \equiv \sqrt{V_{j}^{(W)} + rac{g+1}{g}V_{j}^{(B)}}$$

Within imputations:

$$V_{j}^{(W)} \equiv rac{\sum_{\ell=1}^{g} \mathit{SE}^{2}\left(\beta_{j}^{\ell}
ight)}{\mathit{g}}$$

Between imputations:

$$V_j^{(B)} \equiv rac{\sum_{\ell=1}^{g} \left(eta_j^{\ell} - ilde{eta}_j
ight)^2}{g-1}$$

Background

### Final Model

... And we are left with a final model:

```
t\hat{a}p = 570.1 + (-86.3)ref_some_college +
    (-62.9)ref_college_grad + (-164.9)com_some_college +
(-361.9)com_less_hs + (184.1)race_asian + (17.2)race_black +
    (286.9)race_white + (254.0)race_other + (-0.8)age +
               (-12.9)sex_female + (38.8)fipr
```

Note that there is uncertainty around each of our  $\beta$  values, for example:

- ▶ The standard error for fipr is 19.29, with a 95% confidence interval between 0.19 and 75.8.
- ▶ The standard error for sex\_female is 23.44, with a 95% confidence interval between -58.78 and 33.08.

Background

# Example with Toy Data

•	fipr ‡	race ‡	age ‡	tap_water ‡
1	0.5	white	44	400
2	4.0	black	37	200
3	2.5	white	74	150
4	3.0	white	20	600
5	NA	black	25	550
6	NA	white	61	500
7	4.7	black	40	1000

Figure: Before

# **I**mputations

imputed_fipr1 ‡	imputed_fipr2 ‡	imputed_fipr3 ‡	imputed_fipr4 ‡	imputed_fipr5 ‡
0.5	0.5	0.5	0.5	0.5
4.0	4.0	4.0	4.0	4.0
2.5	2.5	2.5	2.5	2.5
3.0	3.0	3.0	3.0	3.0
5.0	4.3	5.0	2.5	5.0
2.6	1.7	3.4	0.0	3.5
4.7	4.7	4.7	4.7	4.7

Figure: During

# Average Imputed Values

*	imputed_fipr_avg <sup>‡</sup>	race ‡	age ‡	tap_water ‡
1	0.5	white	44	400
2	4.0	black	37	200
3	2.5	white	74	150
4	3.0	white	20	600
5	4.3	black	25	550
6	2.2	white	61	500
7	4.7	black	40	1000

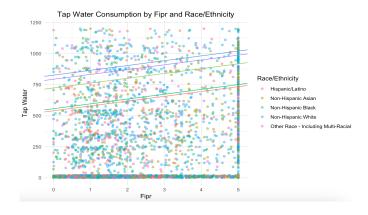
Figure: After

```
example_model <- lm(fipr ~ race + age, data = example data)
    set.seed(16)
    imputed fipr1 <- ifelse(
6
      is.na(example data$fipr).
      pmin(pmax(
8
        4.558427 + (-2.309397) * example data$race ind + (-0.005414) * example data$age +
10
          rnorm(1, mean = 0, sd = 1.36), 0), 5),
      example_data$fipr)
    imputed_fipr5 <- ifelse(</pre>
14
      is.na(example data$fipr).
      pmin(pmax(
16
        4.558427 + (-2.309397) * example data$race ind + (-0.005414) * example data$age +
          rnorm(1, mean = 0, sd = 1.36), 0), 5),
      example_data$fipr)
18
19
20
    imputed_fiprs <- cbind(imputed_fipr1, imputed_fipr2, imputed_fipr3, imputed_fipr4, imputed_fipr5)
24
    average fiprs <- rowMeans(imputed fiprs, na.rm = TRUE)
26
    example data$imputed fipr avg <- example data$fipr
28
    example_data$imputed_fipr_avg[is.na(example_data$fipr)] <- average_fiprs[is.na(example_data$fipr)]
```

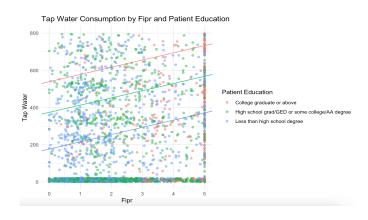
Multiple Imputation

References

## Visualizations



## Visualizations



### Limitations

- ▶ We only worked with missing fipr values, rather than missing values across our whole dataset
- new\_race\_ethnicity decision during data cleaning
- ▶ We simplified the MI process, by only adding random error to the regression outputs (we didn't also add random error to the coefficients)
- ▶ We looked at all tap\_water cases, we did not remove 0's. This could be a direction of future research...

Background

## **Takeaways**

- Multiple imputation as a missing data strategy provides a more sophisticated way of handling missing data that accounts for randomness and provides plausible imputed values.
- ▶ In response to our motivating question, we can conclude that fipr is a statistically significant predictor of tap water consumption.

## Acknowledgements

- ▶ Dr. Justice
- ▶ Dr. Simic-Muller
- ▶ Dr. Edgar

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# **Questions?**

