Imputation when data cannot be pooled

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- Context and problem
- Key idea underlying the method
- mi impute from
- Example of 100% missing confounder
- Simulation results
- Final remarks
- Collaborative efforts such as pooling or consortia projects are commonly undertaken to address complex research questions, enhance precision, and improve the generalizability of findings
- Individual data is often not pooled but harmonized and analyzed at individual sites (i.e., distributed data networks) due to regulatory constraints and the need for timely results
- Systematic (100%) missing data is likely to occur
- mi impute cannot be used without any observed data
- The variable systematically missing in one study site can be of any type (quantitative, qualitative) and any shape
- One or more study sites within the network have data to estimate an imputation model
- Files containing the estimated regression coefficients and their associated precision from the imputation model are shared across the network
- Imputations are generated by inverting the predicted conditional cumulative probabilities

It is a user-written imputation method involving two commands:

- mi impute from get receives list of files (.txt, .xlsx) containing estimated regression coefficients and returns formatted matrices. If multiple files are specified, it combines regression coefficients using an inverse-variance weighted least squares model.
- mi impute from receives the formatted regression coefficients, takes a random draw from their posterior, and generates multiple imputations

Both commands require the specification of the imputation model using the option imodel().

- qreg for modelling conditional quantiles of a quantitative variable. If p predictors, then $99(p + 1)$ regression coefficients
- mlogit for modelling conditional probabilities of a categorical variable.

If p predictors and k levels, then $k(p + 1)$ regression coefficients

• logit is used for modelling the conditional probability of a binary variable.

If p predictors, then $p + 1$ regression coefficients

How does it work conditional quantile imputation?

Consider a continuous variable z_i completely missing in Study 1.

• In another study site, saying Study 2, estimate p-quantile regression model for the continuous variable z_i conditionally on predictors w_i

$$
Q_{z_i|\mathbf{w}_i}(p) = \mathbf{w}_i \gamma(p) \quad p \in \{0.01, 0.02, \ldots, 0.99\}
$$

- \bullet Back to Study 1, draw a random value U_i from a random continuous uniform distribution $U(0, 1)$ for the *i*-th individual
- Extract the floor $f = |U_i\%|$ and modulus mod $= |U_i\%| |U\%|$
- The *m*-th imputation $z_i^{(m)}$ $i_j^{(m)}$ for the *i*-th individual is the weighted average of the f and $f + 1$ conditional predicted quantiles

$$
z_i^{(m)}=(1-\text{mod}\,)\hat{Q}_{z_i|\textbf{w}_i}(f)+\text{mod}\,\hat{Q}_{z_i|\textbf{w}_i}(f+1)
$$

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Mata function for conditional quantile imputation

```
mata:
real colvector mi_impute_cmd_from_xb_qreg(real matrix X, real rowvector b)
ſ
             p = \text{cols}(X)u = runiform(rows(X), 1, 0.01, .99) * 100pvec = J(1, 99, NULL)for (i=1; i<=99; i++) pyec[i] = \&(b[1, (i*p)-(p-1):(i*p)]vi = J(rows(X), 1, ...)for (i=1; i<=rows(X); i++) {
                 f = floor(u[i])mod = mod(u[i], 1)y1[i] = (1 - mod) * (X[i, ]* * prove[f]') + mod * (X[i, ]* * prove[f+1]')ł
return(yi)
```
Impute a Normal distribution

Impute a χ^2 distribution with 1 degree of freedom

Impute a Beta distribution

Impute a Normal distribution conditionally on a binary predictor

Impute a Beta distribution conditionally on a binary predictor

use study_1, clear

mi set wide

mi register imputed z

mi_impute_from_get , b(e_b) v(e_v) imodel(qreg) /// colnames(w _cons)

 $mat i_b = r(get_i b)$ mat $i_v = r(get_iV)$

mi impute from z, $add(1) b(i_b) v(i_v) imodel(qreg)$

Mechanisms underlying confounding effects

- Common causes of exposure and outcome
	- $C \sim \text{Bern}(0.4)$ $Z \sim \chi^2(1)$
- Exposure
	- $X \sim$ Bern(invlogit($\alpha_0 + \alpha_1 C + \alpha_2 Z$))
- Outcome
	- $Y \sim$ Bern(invlogit($\beta_0 + \beta_1 X + \beta_2 C + \beta_3 Z$))

Target of statistical inference is β_1 representing the C and Z adjusted conditional effect of the treatment X on the outcome Y .

Type of confounding

Confounders C and Z are strongly increasing the probability of being exposed $(\alpha_1 > 0, \alpha_2 > 0)$ as well as the outcome probability $(\beta_2 > 0, \beta_1 < \alpha_2 < 0)$ $\beta_3 > 0$).

The conditional effect of the exposure is a small increment in the outcome probability $\beta_1 = \ln(1.2) = 0.18$

One way to test mi impute from

- $\bullet\,$ Generate Study 1 from the confounding mechanism, estimate $\hat\beta_1$, and then set confounder Z to missing
- Generate Study 2 from the same confounding mechanism, estimate the conditional quantile imputation model
- Open Study 1, generate 10 multiple imputations using mi impute from, estimate $\bar{\beta}_1$ using mi $\,$ estimate

If mi impute from works well, we can expect that the sampling distribution of $\hat{\beta}_1$ based on fully observed data and the sampling distribution of $\bar{\beta}_1$ based on fully externally multiple imputed data should be bell-shaped and centered about the parameter β_1 set in the simulation.

Scenario 1: External imputation from identical confounding mechanism

Study 1 and Study 2 with sample size $n = 1,000$ come from the following mechanism

• Exposure

 $X \sim$ Bern(invlogit(logit(0.10) + log(3)C + log(1.3)Z))

• Outcome

 $Y \sim$ Bern(invlogit(logit(0.20) + log(1.2) $X + \log(3)C + \log(1.3)Z)$)

Comparison of simulated sampling distributions

Fully Adjusted Exposure-Outcome (log) Odds Ratio

The conditional effect of the exposure in Study 1 (under full data or 100% externally imputed) is centered about the parameter value $\beta_1 = 0.18$.

Scenario 2: External imputation from weaker confounding mechanism

Study 1 as before but Study 2 come from a weaker confounding mechanism.

• Exposure

 $X \sim$ Bern(invlogit(logit(0.10) + log(3)C + log(1.1)Z))

• Outcome

 $Y \sim$ Bern(invlogit(logit(0.20) + log(1.2) $X +$ log(3)C + log(1.1)Z))

Comparison of simulated sampling distributions

Fully Adjusted Exposure-Outcome (log) Odds Ratio

The conditional effect of the exposure in Study 1 under 100% externally imputation is, on average, twice as much the parameter value $\beta_1 = 0.18$.

Scenario 3: External imputation from stronger confounding mechanism

Study 1 as before but Study 2 come from a stronger confounding mechanism.

• Exposure

 $X \sim$ Bern(invlogit(logit(0.10) + log(3)C + log(1.6)Z))

• Outcome

Y ~ Bern(invlogit(logit(0.20) + log(1.2)X + log(3)C + log(1.6)Z))

Comparison of simulated sampling distributions

Fully Adjusted Exposure-Outcome (log) Odds Ratio

The conditional effect of the exposure in Study 1 under 100% externally imputation is, on average, at the opposite side of the parameter value $\beta_1 = 0.18$.

Scenario 4: External imputation from a heterogeneous mechanism

Study 1 as before but Study 2 come from a heterogeneous confounding mechanism.

Comparison of simulated sampling distributions

Fully Adjusted Exposure-Outcome (log) Odds Ratio

The conditional effect of the exposure in Study 1 under 100% externally imputation is, on average, centered about the parameter value $\beta_1 = 0.18$.

Back to our motivating example

```
use qreg_study_1_miss, clear
mi set wide
mi register imputed z
```

```
mi_impute_from_get , ///
b(e_b_s2 e_b_s3 e_b_s4 e_b_s5) ///
v(e_v_s2 e_v_s3 e_v_s4 e_v_s5) ///
     \text{columns}(y \ x \ c \ \_\text{cons}) \ \text{imodel}(qreg)mat ib = r(get_ib)
mat iV = r(get_iV)
```

```
mi impute from z, add(10) b(ib) v(iV) imodel(qreg) ///
rseed(240912)
```
mi estimate, post eform: logistic y x c z

 $\hat{\beta}_1 = 1.333971$ based on complete data

 $\bar{\beta}_1 = 1.304004$ based on external imputations from 4 heterogeneous studies using mi impute from

Next step in a collaborative effort would be the specification of a meta-analytical model to learn from multiple studies.

- mi impute from is based on the principle of inverting predicted conditional cumulative probabilities
- mi impute from can be used with both sporadic and systematic missing data
- mi impute from cannot be called by mi impute chained
- mi impute from using regression coefficients from imputation model estimated in data where completely different mechanisms are operating is likely to lead to the wrong inferential results
- This is an on-going joint work with Robert Thiesmeier & Matteo Bottai at Karolinska Institutet
- • Thiesmeier R, Bottai M, Orsini N. (2024). Systematically missing data in distributed data networks: multiple imputation when data cannot be pooled. Journal of Statistical Computation and Simulation. In Press.
- Thiesmeier R, Bottai M, Orsini N. (2024). Imputation when data cannot be pooled. Stata Journal. On-going.