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Dealing with missing data in model-based clustering through a MNAR model

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Take home message

- I The missing data pattern may convey some information on clustering
- **2** Embed the missingness mechanism directly within the clustering modeling step

1 Introduction

- 2 A model-based MNAR clustering approach
- **3** Inference procedures
- 4 Medical study illustration
- **5** Concluding remarks

Missing data: an inevitable event

The larger the datasets, the more missing data may appear...

Two traditional solutions (for obtaining a filled dataset)

- Discard individuals with missing data: expect to add variance into analysis
- Impute missing data: expect to add bias modeling into analysis

General guidelines

- Obtaining a non-missing dataset is not the final goal
- Missing data management should take into account the initial analysis target

Our analysis target: model-based clustering

Embed missing data management into this paradigm...

Missing data: notations

- **y** = { y_1, \ldots, y_n }: full dataset with *n* individuals
- **y**_i = $(y_i^1, \ldots, y_i^d) \in \mathbb{R}^d$: full individual $i \in \{1, \ldots, n\}$
- **c** = { c_1, \ldots, c_n }: pattern of missing data for the full dataset
- $\mathbf{c}_i = (c_i^1, \dots, c_i^d) \in \{0, 1\}^d$: pattern of missing data for individual $i \in \{1, \dots, n\}$

$$c_i^j = 1 \Leftrightarrow y_i^j$$
 is missing

- $o_i = \{j : c_i^j = 0\}$: the observed variables indexes for individual *i*
- **y**_i^o: the observed variables values for individual i
- **v** $\mathbf{y}^o = \{\mathbf{y}_1^{o_1}, \dots, \mathbf{y}_n^{o_n}\}$: the observed values in \mathbf{y}
- $m_i = \{j : c_i^j = 1\}$: the missing variables indexes for individual *i*
- **y**_{*i*}^{m_i}: the missing variables values for individual *i*
- **y**^{*m*} = { $\mathbf{y}_1^{m_1}, \ldots, \mathbf{y}_n^{m_n}$ }: the missing values in **y**

 $\mathbf{y} = {\{\mathbf{y}^o, \mathbf{y}^m\}}$ is the full dataset with its observed and missing parts

Missing data: typology of the missing mechanisms

Missing completely at random (MCAR):

$$P(\mathbf{c}|\mathbf{y}; \boldsymbol{\psi}) = P(\mathbf{c}; \boldsymbol{\psi}) \quad \forall \mathbf{y}$$

Missing at random (MAR):

$$P(\mathbf{c}|\mathbf{y}; oldsymbol{\psi}) = P(\mathbf{c}|\mathbf{y}^{o}; oldsymbol{\psi}) \quad orall \mathbf{y}^{m}$$

Missing not at random (MNAR): the mechanism is not MCAR nor MAR

Clustering: model-based approach

- Partition with K clusters: $z = (z_1, ..., z_n)$ where
 - **z**_i = $(z_i^1, \ldots, z_i^K) \in \{0, 1\}^K$ **z**_i^k = 1 if **y**_i belongs to cluster k, $z_i^k = 0$ otherwise

Gaussian mixture: $\mathbf{y}_1, \ldots, \mathbf{y}_n$ are i.i.d. from the mixture

$$f(\mathbf{y}_i; \boldsymbol{\theta}) = \sum_{k=1}^{K} \pi_k \phi_k(\mathbf{y}_i; \theta_k)$$

where:

π_k = P(z_i^k = 1)
 φ_k(.; θ_k): d-variate Gaussian pdf with mean vector and covariance matrix θ_k = (μ_k, Σ_k), d-multinomial pdf with θ_k = p_k probabilities vector or mixed pdf.
 θ = (π₁,..., π_K, θ₁,..., θ_K)

Question we address in this work

Which distribution $P(\mathbf{c}|\mathbf{y}, \mathbf{z}; \psi)$ to propose in this clustering context?

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Logistic model: a natural and flexible candidate

$$P(\mathbf{c}|\mathbf{y},\mathbf{z};\boldsymbol{\psi}) = \prod_{i=1}^{n} \prod_{j=1}^{d} P(c_{i}^{j}|\mathbf{y},\mathbf{z};\boldsymbol{\psi})$$

$$MCAR, \text{ with } \psi = \alpha_0$$

$$\logit(P(c_i^j = 1 | \mathbf{y}, \mathbf{z}; \psi)) = \alpha_0$$

$$MNARz (MNARz^j), \text{ with } \psi = (\alpha_0, \beta_1^{1...d}, \dots, \beta_K^{1...d})$$

$$\logit(P(c_i^j = 1 | \mathbf{y}, \mathbf{z}; \psi)) = \alpha_0 + \sum_{k=1}^K \beta_k^j z_i^k$$

$$MNARy, \text{ with } \psi = (\alpha_0, \alpha_1, \dots, \alpha_d)$$

$$\logit(P(c_i^j = 1 | \mathbf{y}, \mathbf{z}; \psi)) = \alpha_0 + \alpha_j y_i^j$$

$$MNARyz, \text{ with } \psi = (\alpha_0, \alpha_1, \dots, \alpha_d, \beta_1, \dots, \beta_K)$$

$$\logit(P(c_i^j = 1 | \mathbf{y}, \mathbf{z}; \psi)) = \alpha_0 + \alpha_j y_i^j + \sum_{k=1}^K \beta_k z_i^k$$

Introduction

MNARz analysis: it depends on y through z!

$$P(c_i^j = 1 | \mathbf{y}; \boldsymbol{\theta}, \boldsymbol{\psi}) = \sum_{k=1}^{K} P(c_i^j = 1 | \mathbf{y}, \mathbf{z}; \boldsymbol{\psi}) P(\mathbf{z} | \mathbf{y}; \boldsymbol{\theta})$$

Example of a univariate Gaussian model with the three components

$$0.2N(\cdot; 0, 1) + 0.3N(\cdot; 1, 2) + 0.5N(\cdot; 2, 3)$$

and with parameters of the logit expression: $\alpha_0=1, \beta_1=1, \beta_2=-1, \beta_3=1$



MNARz analysis: pattern c gives information on partition z!

Draw Bayes error of a MNARz model with two components and 20% of missing data

 $\pi_k = 0.5, \; \|oldsymbol{\mu}_2 - oldsymbol{\mu}_1\|$ varies, $\Sigma_1 = \Sigma_2 = \mathsf{I}, |eta_2 - eta_1|$ varies



Both μ_k and β_k act on the Bayes error

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Ignorable vs. non ignorable model

A missing mechanism is ignorable if likelihoods can be decomposed as

$$L(\theta, \psi; \underbrace{\mathbf{y}^{o}, \mathbf{c}}_{\text{observed data}}) = L(\psi; \mathbf{c} | \mathbf{y}^{o}) \times L(\theta; \mathbf{y}^{o})$$

Some simple algebra show that this occurs when missing mechanism is not MNAR

Inference of θ

"If the missing mechanism is ignorable then likelihood-based inferences for θ from $L(\theta; \mathbf{y}^{\circ})$ will be the same as likelihood based inference for θ from $L(\theta, \psi; \mathbf{y}^{\circ}, \mathbf{c})$." ([Little and Rubin, 2002] Section 6.2)

- MCAR is ignorable
- MNARz, MNARy and MNARyz are non ignorable

EM algorithm: looks simple

Decomposition of $Q(oldsymbol{ heta},\psi;\widehat{oldsymbol{ heta}},\widehat{\psi})$

The expected complete log-likelihood conditional related to observed data is:

$$\mathsf{E}\left[\mathsf{L}(\boldsymbol{\theta},\boldsymbol{\psi};\mathbf{c},\mathbf{y},\mathbf{z});\widehat{\boldsymbol{\theta}},\widehat{\boldsymbol{\psi}}|\mathbf{y}^{o},\mathbf{c}\right] = Q_{y}(\boldsymbol{\theta};\widehat{\boldsymbol{\theta}},\widehat{\boldsymbol{\psi}}) + Q_{c}(\boldsymbol{\psi};\widehat{\boldsymbol{\theta}},\widehat{\boldsymbol{\psi}})$$

$$\begin{aligned} Q_{\mathbf{y}}(\boldsymbol{\theta}; \widehat{\boldsymbol{\theta}}, \widehat{\boldsymbol{\psi}}) &= \sum_{i=1}^{n} \sum_{k=1}^{K} \tau_{i}^{k} E\left[\log(\pi_{k} \phi_{k}(\mathbf{y}_{i}; \theta_{k})) | \mathbf{y}_{i}^{o_{i}}, \mathbf{c}_{i}; \widehat{\boldsymbol{\theta}}, \widehat{\boldsymbol{\psi}}\right] \\ Q_{c}(\boldsymbol{\psi}; \widehat{\boldsymbol{\theta}}, \widehat{\boldsymbol{\psi}}) &= \sum_{i=1}^{n} \sum_{k=1}^{K} \tau_{i}^{k} E\left[\log(P(\mathbf{c}_{i} | z_{i}^{k} = 1, \mathbf{y}_{i}; \theta, \boldsymbol{\psi})) | \mathbf{y}_{i}^{o_{i}}, \mathbf{c}_{i}; \widehat{\boldsymbol{\theta}}, \widehat{\boldsymbol{\psi}}\right] \\ \tau_{i}^{k} &= P(z_{i}^{k} = 1 | \mathbf{c}_{i}, \mathbf{y}_{i}^{o_{i}}; \widehat{\boldsymbol{\theta}}, \widehat{\boldsymbol{\psi}}) = \frac{\widehat{\pi}_{k} \phi_{k}(\mathbf{y}_{i}^{o_{i}}; \theta_{k}^{o_{i}}) P(\mathbf{c}_{i} | z_{i}^{k} = 1, \mathbf{y}_{i}^{o_{i}}; \widehat{\boldsymbol{\psi}})}{\sum_{h=1}^{K} \widehat{\pi}_{h} \phi_{h}(\mathbf{y}_{i}^{o_{i}}) P(\mathbf{c}_{i} | z_{i}^{h} = 1, \mathbf{y}_{i}^{o_{i}}; \widehat{\boldsymbol{\psi}})} \end{aligned}$$

EM and/or SEM algorithms

$$au_i^k \propto \widehat{\pi}_k \phi_k(\mathbf{y}_i^{o_i}; heta_k^{o_j}) \prod_{j=1}^d (1 + \exp(-r_i^j \widehat{eta}_k))^{-1} ext{ where } r_i^j = \left\{egin{array}{cc} 1 & ext{if } c_i^j = 1 \ -1 & ext{otherwise} \end{array}
ight.$$

$$P(c_i^j | \mathbf{y}_i^{o_i}, z_i^k = 1; \boldsymbol{\psi}) = \begin{cases} \int_{-\infty}^{+\infty} \frac{1}{1 + \exp(-(\alpha_j y_i^j))} \phi_k(y_i^j; \theta_k^j) dy_i^j & \text{if } c_i^j = 1\\ \\ \frac{1}{1 + \exp(\alpha_j y_i^j)} & \text{otherwise} \end{cases}$$

In the Gaussian case, there is no closed form [Pirjol, 2013] (same for MNARyz)
 But SEM is still simple in that case thanks to random drawing instead of expectation

Link with some usual procedures!

Concatenation [Jones, 1996]: model equivalence $MNARz^{i}(\mathbf{y}^{o}, \mathbf{c}) \iff MCAR((\mathbf{y}^{o}|\mathbf{c}))$ $MNARz(\mathbf{y}^{o}, \mathbf{c}) \iff MCAR\left(\left(\mathbf{y}^{o}\left|\left(\sum_{j=1}^{d} \mathbf{c}_{j}^{j}\right)\right)\right)\right)$

"All Available Cases" [Little and Rubin, 2002]: estimation equivalence In case of conditional independence between variables, whatever MCAR or MNAR*:

Classical (S)EM \iff (S)EM without estimating missing y^m

... an opportunity to reduce the computing time

What about model selection?

Can select between MCAR and MNAR* with any information criterion (BIC, ICL)

Even if the missing mechanism is ignorable for MCAR...

 \ldots need to model \boldsymbol{c} to compare a MCAR and a MNAR model

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Hospital Data

- Number of patients: $n = 5 \ 146$
- Number of features: d = 7
 - Age
 - Size
 - Weight
 - Cardiac frequency
 - Hemoglobin concentration
 - Temperature
 - Minimum Diastolic and Systolic Blood Pressure
- Percentage of missing data: 6.4%

ICL comparison





- MCAR, MNARy and MNARz are equivalent until K = 3
- MNARz and MNARyz clearly indicate presence of an additional cluster (K = 4)

It seems to be an illustation of the effect of ${\bf c}$ through MNARz and MNARyz

Missing Pattern



It seems that MNARz modelling leads to a missing free cluster

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Summary

- \blacksquare Interest to put a model on ${\bf c}$
- Interest of the simple but meaningful model MNARz
- Link between our models and usual methods

Ongoing works

- Deeper analysis of the previous results with doctors...
- Implement the proposed models/algo. in the Mixmod software^a
- Use mixed data algorithms for medical study with the same MNAR* models

^ahttp://www.mixmod.org

References



Jones, M. P. (1996).

Indicator and stratification methods for missing explanatory variables in multiple linear regression.

Journal of the American statistical association, 91(433):222–230.



Little, R. J. and Rubin, D. B. (2002). Statistical Analysis with Missing Data. Wiley.



Pirjol, D. (2013).

The logistic-normal integral and its generalizations.

Journal of Computational and Applied Mathematics, 237(1):460–469.