

On fast supervised learning for normal mixture models with missing information

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Abstract

It is an important research issue to deal with mixture models when missing values occur in the data. In this paper, computational strategies using auxiliary indicator matrices are introduced for efficiently handling mixtures of multivariate normal distributions when the data are missing at random and have an arbitrary missing data pattern, meaning that missing data can occur anywhere. We develop a novel EM algorithm that can dramatically save computation time and be exploited in many applications, such as density estimation, supervised clustering and prediction of missing values. In the aspect of multiple imputations for missing data, we also offer a data augmentation scheme using the Gibbs sampler. Our proposed methodologies are illustrated through some real data sets with varying proportions of missing values.

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1. Introduction

Finite mixture models are known as powerful and flexible tools, which have been fully developed and applied in various theoretic and real problems as they are capable of modelling a wide range of densities, see for example Refs. [1–3]. However, missing values frequently appear in many real-world multivariate data sets that complicate data analyses and statistical inferences for practitioners. Missing data imputation techniques under the assumption of multivariate normal model have been well studied by Refs. [4,5]. Recently, learning mixture models from incomplete data becomes an important research issue in multivariate analysis. The work on the use of Gaussian component was pioneered by Ghahramani and Jordan [6], denoted by GJ hereafter. They present how to implement the expectation–maximization (EM) algorithm [7] to compute maximum likelihood (ML) estimates from

multivariate data with arbitrary pattern of missingness. They also compare the performance of EM imputation with a common mean imputation (MI) heuristic for the supervised classification of incomplete features.

Due to rapid advance of computational developments, Bayesian sampling-based approaches are usually considered as an alternative way in dealing with mixture models. There are plenty of papers in the literature to address the problem of fitting normal mixture models under Bayesian treatments. For example, Diebolt and Robert [8] employ the data augmentation (DA) technique of Tanner and Wong [9] as an approximation method for evaluating the posterior distribution and show a duality principle. Escobar and West [10] present a nonparametric Bayesian density estimation for Dirichlet process mixture models. Richardson and Green [11] and Zhang et al. [12] propose a full Bayesian inference for a normal mixture model with unknown number of components using the reversible jump MCMC algorithm proposed by Green [13]. Stephens [14] and Fruhwirth-Schnatter [15] demonstrate Bayesian strategies for the elimination of label switching problems.

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In this paper, we offer an efficient EM algorithm for the fitting of a likelihood-based normal mixture model using partially observed data. To reduce computational burden during the EM iterations, we incorporate two types of auxiliary binary indicator matrices corresponding to the observed and unobserved components of each datum. With strategies similar to EM, we also offer a DA computational technique for efficiently imputing missing values and learning parameters using the Gibbs sampler [16], which constructs a Markov chain that converges to a tractable posterior distribution. The feature of the chosen prior distributions is weakly informative to avoid mathematical and computational pitfalls of using improper priors in mixture model, see Celeux et al. [17].

The rest of the paper proceeds as follows. In the next section, we describe the model and its notations, and present some important statistical properties based on the missing information framework. In Sections 3 and 4, two efficient EM and DA algorithms are developed to cope with ML and Bayesian estimation, respectively. We also investigate two issues regarding classification and prediction of incomplete features from both ML and Bayesian perspectives. In Section 5, some real data sets are utilized to illustrate our proposed methodologies with varying proportions of artificially missing values. Also, empirical comparisons between ML and Bayesian approaches in terms of classification and prediction accuracies for incomplete features are demonstrated. Finally, some concluding remarks are given in Section 6.

2. A normal mixture model with missing information

In the normal mixture model, we assume that $Y = (Y_1, \dots, Y_n)$ form a p -dimensional random sample from a population with g subclasses $\mathcal{C}_1, \dots, \mathcal{C}_g$, and each Y_j has the density

$$f(Y_j | \Theta) = \sum_{i=1}^g w_i \phi_p(Y_j | \mu_i, \Sigma_i), \quad w_i \geq 0, \quad \sum_{i=1}^g w_i = 1, \quad (1)$$

where w_i 's are mixing probabilities, $\phi_p(\cdot | \mu, \Sigma)$ denotes a p -dimensional multivariate normal component density with mean μ and covariance matrix Σ , and $\Theta = (w_1, \dots, w_g, \mu_1, \dots, \mu_g, \Sigma_1, \dots, \Sigma_g)$ is the vector of mixture model parameters subject to $\sum_{i=1}^g w_i = 1$ and Σ_i 's are positive definite matrices. Thus, there are $g(p+1)(p+2)/2 - 1$ distinct parameters in model (1).

Typically, in the EM framework, mixture models can be characterized as having an incomplete data structure. It is convenient to formalize the missing part as a set of membership labels $Z = (Z_1, \dots, Z_n)$ with each label $Z_j = (Z_{1j}, \dots, Z_{gj})$ being a binary vector such that $Z_{ij} = 1$ if Y_j belongs to component i and $Z_{ij} = 0$ otherwise. Given the mixing probabilities ω , Z_1, \dots, Z_n independently

follow a multinomial distribution. We shall write $Z_j \sim \mathcal{M}(1; w_1, \dots, w_g)$.

For notational simplicity, let

$$\Delta_{ij} = (Y_j - \mu_i)^\top \Sigma_i^{-1} (Y_j - \mu_i), \quad (2)$$

denote the Mahalanobis distance for Y_j with respect to mean μ_i and covariance matrix Σ_i . The complete likelihood function for Θ is

$$L_c(\Theta | Y, Z) \propto \prod_{j=1}^n \prod_{i=1}^g \left(w_i |\Sigma_i|^{-1/2} \exp\left(-\frac{1}{2} \Delta_{ij}\right) \right)^{Z_{ij}}. \quad (3)$$

We consider the maximum likelihood estimation problem of model (1) when Y are not completely observed. We further assume that the patterns of missingness are arbitrary and missing at random (MAR), see Refs. [18,19] for more details. Generally speaking, MAR refers to the missingness depending only on observed values but not on missing values.

Let Y_j be partitioned into two components (Y_j^o, Y_j^m) , where Y_j^o ($p_j^o \times 1$) and Y_j^m ($(p - p_j^o) \times 1$) denote the observed and missing components of Y_j , respectively. To facilitate the EM algorithm, it is advantageous to introduce two types of binary indicator matrices, denoted by O_j and M_j hereafter, corresponding to Y_j such that $Y_j^o = O_j Y_j$ and $Y_j^m = M_j Y_j$, respectively. Notice that O_j and M_j are $p_j^o \times p$ and $(p - p_j^o) \times p$ matrices extracted from a p -dimensional identity matrix I_p corresponding to row-positions of Y_j^o and Y_j^m in Y_j , respectively. We then have the following two propositions.

Proposition 1. Suppose Y_j is partitioned into two components (Y_j^o, Y_j^m) , where $Y_j^o = O_j Y_j$ and $Y_j^m = M_j Y_j$. We thus have

$$Y_j = \begin{cases} Y_j^o, & \text{if } p_j^o = p; \\ O_j^\top Y_j^o + M_j^\top Y_j^m, & \text{if } 1 \leq p_j^o < p, \end{cases}$$

$$\text{and } O_j^\top O_j + M_j^\top M_j = I_p.$$

Proof. The proof is straightforward and hence is omitted. □

Proposition 2. Let $Y_j \sim \sum_{i=1}^g w_i \phi_p(Y_j | \mu_i, \Sigma_i)$, and let Y_j^o and Y_j^m be the observed and missing components corresponding Y_j , respectively. The marginal distribution of Y_j^o is denoted by $Y_j^o \sim \sum_{i=1}^g w_i \phi_{p_j^o}(Y_j^o | \mu_{ij}^o, \Sigma_{ij}^{oo})$, where

$$\phi_{p_j^o}(Y_j^o | \mu_{ij}^o, \Sigma_{ij}^{oo}) = (2\pi)^{-p_j^o/2} |\Sigma_{ij}^{oo}|^{-1/2} \exp\left(-\frac{1}{2} \Delta_{ij}^o\right),$$

and

$$\begin{aligned} \mu_{ij}^o &= O_j \mu_i, \quad \Sigma_{ij}^{oo} = O_j \Sigma_i O_j^\top, \\ \Delta_{ij}^o &= (Y_j^o - \mu_i)^\top \Sigma_{ij}^{oo} (Y_j^o - \mu_i), \\ \Sigma_{ij}^{oo} &= O_j^\top (O_j \Sigma_i O_j^\top)^{-1} O_j. \end{aligned} \quad (4)$$

Consequently, $\mathbf{Y}_j^m | \mathbf{Y}_j^o \sim \sum_{i=1}^g w_{ij}^* \phi_{p-p_j^o}(\mathbf{Y}_j^m | \boldsymbol{\mu}_{ij}^{m,o}, \boldsymbol{\Sigma}_{ij}^{mm,o})$, where

$$\phi_{p-p_j^o}(\mathbf{Y}_j^m | \boldsymbol{\mu}_{ij}^{m,o}, \boldsymbol{\Sigma}_{ij}^{mm,o}) = (2\pi)^{-(p-p_j^o)/2} |\boldsymbol{\Sigma}_{ij}^{mm,o}|^{-1/2} \times \exp(-\frac{1}{2} \Delta_{ij}^{m,o}),$$

and

$$w_{ij}^* = w_i \phi_{p_j^o}(\mathbf{Y}_j^o | \boldsymbol{\mu}_{ij}^o, \boldsymbol{\Sigma}_{ij}^{oo}) / \sum_{h=1}^g w_h \phi_{p_j^o}(\mathbf{Y}_j^o | \boldsymbol{\mu}_{hj}^o, \boldsymbol{\Sigma}_{hj}^{oo}),$$

$$\begin{aligned} \boldsymbol{\mu}_{ij}^{m,o} &= \mathbf{M}_j(\boldsymbol{\mu}_i + \boldsymbol{\Sigma}_i \mathbf{S}_{ij}^{oo}(\mathbf{Y}_j - \boldsymbol{\mu}_i)), \\ \boldsymbol{\Sigma}_{ij}^{mm,o} &= \mathbf{E}_{ij} \boldsymbol{\Sigma}_i \mathbf{M}_j^\top, \\ \mathbf{E}_{ij} &= \mathbf{M}_j(\mathbf{I}_p - \boldsymbol{\Sigma}_i \mathbf{S}_{ij}^{oo}), \\ \Delta_{ij}^{m,o} &= (\mathbf{Y}_j - \boldsymbol{\mu}_i)^\top \mathbf{S}_{ij}^{mm,o}(\mathbf{Y}_j - \boldsymbol{\mu}_i), \\ \mathbf{S}_{ij}^{mm,o} &= \mathbf{E}_{ij}^\top (\mathbf{E}_{ij} \boldsymbol{\Sigma}_i \mathbf{M}_j^\top)^{-1} \mathbf{E}_{ij}. \end{aligned} \quad (5)$$

Proof. The sketch of the proof is given in Appendix A. \square

To enhance the computational efficiency for estimation, we suggest to rearrange \mathbf{Y} according to unique missing patterns of the data. The procedure can be implemented as follows:

- (a) Build a binary $n \times p$ indicator matrix, $\mathbf{R} = [r_{ij}]$, with each entry $r_{ij} = 1$ if Y_{ij} is missing and $r_{ij} = 0$ otherwise.
- (b) Build a $p \times 1$ vector $\mathbf{z} = \mathbf{R}\mathbf{b}$, where $\mathbf{b} = (2^1, 2^2, \dots, 2^p)^\top$. Notice that the number of unique missing patterns is equal to the number of unique elements in \mathbf{z} .
- (c) Rank \mathbf{z} in an ascending or descending order, denoted by \mathbf{z}^* . Rearrange \mathbf{Y} according to the row positions of \mathbf{z}^* in \mathbf{z} . This will yield clustering of identical patterns of missingness in \mathbf{Y} which are adjacent to each other.

3. An efficient EM procedure for ML estimation

Let $\mathbf{Y}^o = (\mathbf{Y}_1^o, \dots, \mathbf{Y}_n^o)$ and $\mathbf{Y}^m = (\mathbf{Y}_1^m, \dots, \mathbf{Y}_n^m)$ denote the observed portion and missing portion of the data, respectively. The complete-data log-likelihood function can be reexpressed by

$$\begin{aligned} \ell_c(\boldsymbol{\Theta} | \mathbf{Y}^o, \mathbf{Y}^m, \mathbf{Z}) &= \ell_{c_1}(\mathbf{w} | \mathbf{Y}^o, \mathbf{Y}^m, \mathbf{Z}) + \ell_{c_2}(\boldsymbol{\Psi} | \mathbf{Y}^o, \mathbf{Y}^m, \mathbf{Z}) \\ &= \sum_{i=1}^g \sum_{j=1}^n Z_{ij} \log w_i \\ &\quad + \frac{1}{2} \sum_{i=1}^g \left(\log |\boldsymbol{\Sigma}_i^{-1}| \sum_{j=1}^n Z_{ij} - \sum_{j=1}^n Z_{ij} (\Delta_{ij}^o + \Delta_{ij}^{m,o}) \right), \end{aligned} \quad (6)$$

where $\boldsymbol{\omega} = (w_1, \dots, w_g)$ and $\boldsymbol{\Psi} = (\boldsymbol{\mu}_1, \dots, \boldsymbol{\mu}_g, \boldsymbol{\Sigma}_1, \dots, \boldsymbol{\Sigma}_g)$. From Eq. (5), it is easy to verify that $\boldsymbol{\Sigma}_i^{-1} = \mathbf{S}_{ij}^{oo} + \mathbf{S}_{ij}^{mm,o}$ and $\mathbf{O}_j^\top \mathbf{O}_j (\mathbf{I}_p - \boldsymbol{\Sigma}_i \mathbf{S}_{ij}^{oo}) = \mathbf{0}$. Hence, we have the following result.

Proposition 3. The conditional expectation of Eq. (6) is given by

$$\begin{aligned} Q(\boldsymbol{\Theta} | \hat{\boldsymbol{\Theta}}^{(k)}) &= E(\ell_c(\boldsymbol{\Theta} | \mathbf{Y}^o, \mathbf{Y}^m, \mathbf{Z}) | \mathbf{Y}^o, \hat{\boldsymbol{\Theta}}^{(k)}) \\ &= Q_1(\mathbf{w} | \hat{\boldsymbol{\Theta}}^{(k)}) + Q_2(\boldsymbol{\Psi} | \hat{\boldsymbol{\Theta}}^{(k)}). \end{aligned}$$

It follows that

$$Q_1(\mathbf{w} | \hat{\boldsymbol{\Theta}}^{(k)}) = \sum_{i=1}^g \sum_{j=1}^n \hat{Z}_{ij}^{(k)} \log w_i, \quad (7)$$

$$\begin{aligned} Q_2(\boldsymbol{\Psi} | \hat{\boldsymbol{\Theta}}^{(k)}) &= \frac{1}{2} \sum_{i=1}^g \left(\log |\boldsymbol{\Sigma}_i^{-1}| \sum_{j=1}^n \hat{Z}_{ij}^{(k)} \right. \\ &\quad \left. - \text{tr} \left(\boldsymbol{\Sigma}_i^{-1} \sum_{j=1}^n \boldsymbol{\Omega}_{ij}^{(k)} \right) \right), \end{aligned} \quad (8)$$

where

$$\begin{aligned} \boldsymbol{\Omega}_{ij}^{(k)} &= \hat{Z}_{ij}^{(k)} ((\hat{\mathbf{Y}}_{ij}^{(k)} - \boldsymbol{\mu}_i)(\hat{\mathbf{Y}}_{ij}^{(k)} - \boldsymbol{\mu}_i)^\top \\ &\quad + (\mathbf{I}_p - \hat{\boldsymbol{\Sigma}}_i \hat{\mathbf{S}}_{ij}^{oo(k)}) \hat{\boldsymbol{\Sigma}}_i), \end{aligned} \quad (9)$$

$$\hat{Z}_{ij}^{(k)} = \frac{\hat{w}_i^{(k)} \phi_{p_j^o}(\mathbf{Y}_j^o | \hat{\boldsymbol{\mu}}_{ij}^{o(k)}, \hat{\boldsymbol{\Sigma}}_{ij}^{oo(k)})}{\sum_{h=1}^g \hat{w}_h^{(k)} \phi_{p_j^o}(\mathbf{Y}_j^o | \hat{\boldsymbol{\mu}}_{hj}^{o(k)}, \hat{\boldsymbol{\Sigma}}_{hj}^{oo(k)})}, \quad (10)$$

$$\hat{\mathbf{Y}}_{ij}^{(k)} = \hat{\boldsymbol{\mu}}_i^{(k)} + \hat{\boldsymbol{\Sigma}}_i^{(k)} \hat{\mathbf{S}}_{ij}^{oo(k)} (\mathbf{Y}_j - \hat{\boldsymbol{\mu}}_i^{(k)}), \quad (11)$$

and $\hat{\mathbf{S}}_{ij}^{oo(k)}$ is \mathbf{S}_{ij}^{oo} given in Eq. (4) with $\boldsymbol{\Sigma}_i$ replaced by $\hat{\boldsymbol{\Sigma}}_i^{(k)}$.

Proof. The detailed proof is shown in Appendix B. \square

By these propositions, a modification of GJ's EM algorithm can be implemented as follows:

E-step: Given $\boldsymbol{\Theta} = \hat{\boldsymbol{\Theta}}^{(k)}$, impute $\hat{Z}_{ij}^{(k)}$ and $\hat{\mathbf{Y}}_{ij}^{(k)}$ for $i = 1, \dots, g$ and $j = 1, \dots, n$, using Eqs. (10) and (11).

M-Step:

1. Update $\hat{w}_i^{(k)}$ by maximizing Eq. (7) over w_i subject to their sum is unity, which gives

$$\hat{w}_i^{(k+1)} = \frac{1}{n} \sum_{j=1}^n \hat{Z}_{ij}^{(k)}.$$

2. Fix $\boldsymbol{\Sigma}_i$ at $\hat{\boldsymbol{\Sigma}}_i^{(k)}$, update $\hat{\boldsymbol{\mu}}_i^{(k)}$ by maximizing Eq. (8) over $\boldsymbol{\mu}_i$, which leads to

$$\hat{\boldsymbol{\mu}}_i^{(k+1)} = \frac{\sum_{j=1}^n \hat{Z}_{ij}^{(k)} \hat{\mathbf{Y}}_{ij}^{(k)}}{\sum_{j=1}^n \hat{Z}_{ij}^{(k)}}.$$

3. Fix μ_i at $\hat{\mu}_i^{(k+1)}$, update $\hat{\Sigma}_i^{(k)}$ by maximizing constrained Eq. (8) over Σ_i , which leads to

$$\hat{\Sigma}_i^{(k+1)} = \frac{\sum_{j=1}^n \hat{\Omega}_{ij}^{(k)}}{\sum_{j=1}^n \hat{Z}_{ij}^{(k)}},$$

where $\hat{\Omega}_{ij}^{(k)}$ is $\Omega_{ij}^{(k)}$ in Eq. (9) with μ_i replaced by $\hat{\mu}_i^{(k+1)}$. We remark two major advantages of the above EM algorithm:

- With auxiliary matrices \mathbf{O}_j 's obtained at the initiation, there is no need to take care of the associated row positions of missing values at each iteration.
- The implementation of the M-step has low computational cost as it is similar to the case of no missing values. Therefore, the modified EM algorithm is more straightforward than the version of GJ.

Applying Bayes' theorem, the posterior probability of the Y_j belonging to \mathcal{C}_i can be estimated by

$$\begin{aligned} \hat{w}_{ij}^* &= \Pr(Z_{ij} = 1 | Y^o, \hat{\Theta}) \\ &= \frac{\hat{w}_i \phi_{p_j^o}(Y_j | \hat{\mu}_{ij}^o, \hat{\Sigma}_{ij}^{oo})}{\sum_{h=1}^g \hat{w}_h \phi_{p_j^o}(Y_j | \hat{\mu}_{hj}^o, \hat{\Sigma}_{hj}^{oo})}. \end{aligned} \quad (12)$$

By the ML classification theory [20], Y_j is assigned to \mathcal{C}_s if $\hat{w}_{sj}^* > \hat{w}_{ij}^*$ ($i = 1, \dots, g$; $i \neq s$).

Consequently, an ML predictor for the missing component Y_j^m is given by

$$\begin{aligned} \hat{Y}_j^m &= E(Y_j^m | Y^o, \hat{\Theta}) \\ &= \mathbf{M}_j \sum_{i=1}^g \hat{w}_{ij}^* (\hat{\mu}_i + \hat{\Sigma}_i \hat{S}_{ij}^{oo} (Y_j - \hat{\mu}_i)). \end{aligned} \quad (13)$$

4. A data augmentation scheme for Bayesian sampling

The DA algorithm [9] is a general and effective algorithm for producing multiple imputation of missing data. The DA has been broadly applied in a variety of missing data problems, see Refs. [4,19] and references therein. In this section, we construct an efficient DA algorithm that combines the latent variables \mathbf{Z} and unobserved data Y^m for simulating the posterior density of Θ .

The DA algorithm consists of the imputation step (I-step) and the posterior step (P-step). At the k th iteration of the DA algorithm, the I-step is defined by drawing imputations of $Z_j^{(k)}$ and $Y_j^{m(k)}$ from the predictive distributions $p(Z_j | Y^o, \Theta^{(k)})$ and $p(Y_j^m | Y^o, Z_j, \Theta^{(k)})$, respectively for all j , and the P-step refers to generating $\Theta^{(k+1)}$ from $p(\Theta | Y^o, Y^{m(k+1)}, Z^{(k+1)})$. If iterations are performed by a sufficiently long burn-in period, then the simulations $Z_j^{(k)}$, $Y_j^{m(k)}$ and $\Theta^{(k)}$ are distributed according to $p(Z_j | Y^o)$, $p(Y_j^m | Y^o)$ and $p(\Theta | Y^o)$, respectively for all k . To perform the

Bayesian inference for mixture models, it is necessary to choose a proper prior distribution for each parameter to avoid yielding improper posterior distributions [17].

In various mixture contexts, a vague Dirichlet distribution, denoted by $\mathcal{D}(\delta, \dots, \delta)$, is the most natural prior for mixing probabilities \mathbf{w} . Its density is proportional to $w_1^{\delta-1} \dots w_{g-1}^{\delta-1} (1 - w_1 - \dots - w_{g-1})^{\delta-1}$. For component mean vectors μ_i , it is standard to adopt conjugate Gaussian priors. As for the inverse covariance matrix Σ_i^{-1} , the Wishart distribution is often chosen as a conjugate prior. A p -dimensional Wishart distribution with parameters v and \mathbf{A} ($p \times p$) is denoted by $\mathcal{W}_p(v, \mathbf{A})$, and for $v \geq p$ the density is

$$f(\mathbf{U} | \mathbf{A}) \propto |\mathbf{A}|^{-v/2} |\mathbf{U}|^{(v-p-1)/2} \exp\left(-\frac{1}{2} \text{tr}(\mathbf{U}\mathbf{A}^{-1})\right),$$

where \mathbf{A} is called a *hyperparameter* matrix if \mathbf{U} is considered random in Bayesian treatments.

Following the suggestion of [11,14] who base their recommendation on the use of conjugate priors, our chosen priors are

$$\begin{aligned} \mathbf{w} &\sim \mathcal{D}(\delta, \dots, \delta), \\ \mu_i &\sim \mathcal{N}_p(\xi, \kappa^{-1}) \quad (i = 1, \dots, g), \\ \Sigma_i^{-1} | \mathbf{B} &\sim \mathcal{W}_p(2\alpha, (2\mathbf{B})^{-1}) \quad (i = 1, \dots, g), \\ \mathbf{B} &\sim \mathcal{W}_p(2\gamma, (2\mathbf{H})^{-1}), \end{aligned}$$

where \mathbf{B} is a hyperparameter matrix with a conjugate Wishart distribution, and $(\kappa, \mathbf{H}, \xi, \alpha, \delta, \gamma)$ are fixed as appropriate quantities to reflect the flatness of priors. The joint prior distribution function of Θ and \mathbf{B} is

$$\begin{aligned} \pi(\Theta, \mathbf{B}) &\propto |\mathbf{B}|^{g\alpha + (2\gamma - p - 1)/2} \exp(-\text{tr}(\mathbf{H}\mathbf{B})) \prod_{i=1}^g w_i^{\delta-1} \\ &\quad \times \prod_{i=1}^g |\Sigma_i^{-1}|^{(2\alpha - p - 1)/2} \exp \\ &\quad \times \left(-\frac{1}{2} (\mu_i - \xi)^\top \kappa (\mu_i - \xi) - \text{tr}(\mathbf{B}\Sigma_i^{-1})\right). \end{aligned} \quad (14)$$

Upon multiplying Eqs. (3) and (14), we have the following joint posterior density:

$$\begin{aligned} p(\Theta, \mathbf{B}, Y^m, \mathbf{Z} | Y^o) &\propto w_1^{\delta-1} \dots w_g^{\delta-1} |\mathbf{B}|^{g\alpha + (2\gamma - p - 1)/2} \exp(-\text{tr}(\mathbf{H}\mathbf{B})) \\ &\quad \times \prod_{i=1}^g \exp\left(-\frac{1}{2} (\mu_i - \xi)^\top \kappa (\mu_i - \xi)\right) \\ &\quad \times |\Sigma_i^{-1}|^{(2\alpha - p - 1)/2} \exp(-\text{tr}(\mathbf{B}\Sigma_i^{-1})) \\ &\quad \times \prod_{j=1}^n \prod_{i=1}^g \left(w_i |\Sigma_i^{-1}|^{1/2} \exp\left(-\frac{1}{2} (\Delta_{ij}^o + \Delta_{ij}^{m:o})\right)\right)^{Z_{ij}}, \end{aligned} \quad (15)$$

where Δ_{ij}^o and $\Delta_{ij}^{m:o}$ are given in Eqs. (4) and (5), respectively.

Proposition 4. The full conditional posteriors of Θ , \mathbf{B} , \mathbf{Z} and \mathbf{Y}^m are as follows (the symbol “ $|\dots$ ” denotes conditioning on all other variables):

$$p(\mathbf{Z}_j | \mathbf{Y}^o, \Theta) \propto \prod_{i=1}^g (w_i \phi_{p_j^o}(\mathbf{Y}_j^o | \boldsymbol{\mu}_{ij}^o, \boldsymbol{\Sigma}_{ij}^{oo})^{Z_{ij}},$$

$$p(\mathbf{Y}_j^m | Z_{ij} = 1, \dots) \propto \exp\left(-\frac{1}{2}(\mathbf{Y}_j^m - \boldsymbol{\mu}_{ij}^{m-o})^\top \boldsymbol{\Sigma}_{ij}^{mm-o^{-1}} (\mathbf{Y}_j^m - \boldsymbol{\mu}_{ij}^{m-o})\right),$$

$$p(\mathbf{w} | \dots) \propto \prod_{i=1}^g w_i^{\sum_{j=1}^n Z_{ij} + \delta - 1},$$

$$p(\boldsymbol{\mu}_i | \dots) \propto \exp\left(-\frac{1}{2}(\boldsymbol{\mu}_i - \boldsymbol{\mu}_i^*)^\top \boldsymbol{\Sigma}_i^{*-1} (\boldsymbol{\mu}_i - \boldsymbol{\mu}_i^*)\right),$$

$$p(\mathbf{B} | \dots) \propto |\mathbf{B}|^{(2(g\alpha + \gamma) - p - 1)/2} \times \exp\left(-\text{tr}\left(\mathbf{B}\left(\mathbf{H} + \sum_{i=1}^g \boldsymbol{\Sigma}_i^{-1}\right)\right)\right),$$

$$p(\boldsymbol{\Sigma}_i^{-1} | \dots) \propto |\boldsymbol{\Sigma}_i^{-1}|^{(\alpha^* - p - 1)/2} \exp\left(-\frac{1}{2}\text{tr}(\boldsymbol{\Sigma}_i^{-1} \mathbf{A}_i)\right),$$

where $\boldsymbol{\mu}_{ij}^{m-o}$ and $\boldsymbol{\Sigma}_{ij}^{mm-o}$ are given by Eq. (5), and

$$\boldsymbol{\Sigma}_i^* = \left(\boldsymbol{\Sigma}_i^{-1} \sum_{j=1}^n Z_{ij} + \boldsymbol{\kappa}\right)^{-1}, \tag{16}$$

$$\boldsymbol{\mu}_i^* = \boldsymbol{\Sigma}_i^* \left(\boldsymbol{\Sigma}_i^{-1} \sum_{j=1}^n Z_{ij} \mathbf{Y}_j + \boldsymbol{\kappa} \boldsymbol{\xi}\right), \tag{17}$$

$$\alpha_i^* = \sum_{j=1}^n Z_{ij} + 2\alpha, \tag{18}$$

$$\mathbf{A}_i = 2\mathbf{B} + \sum_{j=1}^n Z_{ij} (\mathbf{Y}_j - \boldsymbol{\mu}_i) (\mathbf{Y}_j - \boldsymbol{\mu}_i)^\top, \tag{19}$$

for $i = 1, \dots, g$ and $j = 1, \dots, n$.

Proof. The proof is straightforward and hence is omitted. \square

In the simulation process, samples for \mathbf{Z} , \mathbf{Y}^m , \mathbf{B} and Θ are alternately generated, the DA algorithm using the Gibbs sampler can be implemented as follows:

I-Step:

1. Given Θ , \mathbf{Y}^m and \mathbf{Y}^o , generate \mathbf{Z}_j from $\mathcal{M}(1; r_{1j}, \dots, r_{gj})$, where

$$r_{ij} = \frac{w_i \phi_{p_j^o}(\mathbf{Y}_j^o | \boldsymbol{\mu}_{ij}^o, \boldsymbol{\Sigma}_{ij}^{oo})}{\sum_{h=1}^g w_h \phi_{p_j^o}(\mathbf{Y}_j^o | \boldsymbol{\mu}_{hj}^o, \boldsymbol{\Sigma}_{hj}^{oo})}.$$

2. Generate \mathbf{Y}_j^m given $Z_{ij} = 1$, Θ and \mathbf{Y}^o , from $N_{p-p_j^o}(\boldsymbol{\mu}_{ij}^{m-o}, \boldsymbol{\Sigma}_{ij}^{mm-o})$, where $\boldsymbol{\mu}_{ij}^{m-o}$ and $\boldsymbol{\Sigma}_{ij}^{mm-o}$ are as in Eq. (5).

P-Step:

1. Generate \mathbf{w} given \mathbf{Z} from $\mathcal{D}(n_1 + \delta, \dots, n_g + \delta)$, where $n_i = \sum_{j=1}^n Z_{ij}$.
2. Generate $\boldsymbol{\mu}_i$ given \mathbf{Z} , $\boldsymbol{\Sigma}_i$, \mathbf{Y}^o and \mathbf{Y}^m from $\mathcal{N}_p(\boldsymbol{\mu}_i^*, \boldsymbol{\Sigma}_i^*)$ with $\boldsymbol{\mu}_i^*$ and $\boldsymbol{\Sigma}_i^*$ given in Eqs. (17) and (16), respectively.
3. Generate \mathbf{B} given $\boldsymbol{\Sigma}_1, \dots, \boldsymbol{\Sigma}_g$ from $\mathcal{W}_p(2\gamma^*, (2\mathbf{H}^*)^{-1})$, where $\gamma^* = g\alpha + \gamma$ and $\mathbf{H}^* = \mathbf{H} + \sum_{i=1}^g \boldsymbol{\Sigma}_i^{-1}$.
4. Generate $\boldsymbol{\Sigma}_i^{-1}$ given \mathbf{Z} , $\boldsymbol{\mu}_i$, \mathbf{Y}^o and \mathbf{Y}^m from $\mathcal{W}_p(\alpha_i^*, \mathbf{A}_i^{-1})$, where α_i^* and \mathbf{A}_i are given in Eqs. (18) and (19), respectively.

To satisfy the “Principle of Stable Estimation” of Edwards et al. [21] in the Bayesian treatment, we need to specify $(\boldsymbol{\xi}, \boldsymbol{\kappa}, \alpha, \gamma, \mathbf{H})$ so as to be insensitive to changes of the prior. Specifically, it is often to choose $\delta = 1$. For $\boldsymbol{\xi}$ and $\boldsymbol{\kappa}$, we let $\boldsymbol{\xi}$ be the empirical mean vector and $\boldsymbol{\kappa}^{-1} = (1 - \eta)^{-1} \text{diag}\{R_1^2, \dots, R_p^2\}$, where η is the percentage of missing values of the data which is used to adjust the flatness and R_i is the range of the observed values for the i th attribute. This specification makes a weak prior information for $\boldsymbol{\mu}_i$. As a generalization of [11], we take $\alpha = p + 1$, $\gamma = (p + 1)/10$ and $\mathbf{H} = 10\boldsymbol{\kappa}$.

We are interested in the classification and prediction problems for incomplete features. Under certain conditions, quantities based on Rao-Blackwellization [22] often greatly improve the precision of Monte Carlo estimates. Given a set of converged Monte Carlo DA samples $\Theta^{(\ell)}$ ($\ell = 1, \dots, L$), a Bayesian predictor for \mathbf{Y}_j^m is given by

$$\begin{aligned} \tilde{\mathbf{Y}}_j^m &= \frac{1}{L} \sum_{\ell=1}^L E(\mathbf{Y}_j^m | \mathbf{Y}_j^o, \Theta^{(\ell)}) \\ &= \mathbf{M}_j \frac{1}{L} \sum_{\ell=1}^L \left(\sum_{i=1}^g r_{ij}^{(\ell)} (\boldsymbol{\mu}_i^{(\ell)} + \boldsymbol{\Sigma}_i^{(\ell)} \mathbf{S}_{ij}^{oo(\ell)} (\mathbf{Y}_j - \boldsymbol{\mu}_i^{(\ell)})) \right), \end{aligned} \tag{20}$$

where

$$r_{ij}^{(\ell)} = \frac{w_i^{(\ell)} \phi_{p_j^o}(\mathbf{Y}_j^o | \boldsymbol{\mu}_{ij}^{o(\ell)}, \boldsymbol{\Sigma}_{ij}^{oo(\ell)})}{\sum_{h=1}^g w_h^{(\ell)} \phi_{p_j^o}(\mathbf{Y}_j^o | \boldsymbol{\mu}_{hj}^{o(\ell)}, \boldsymbol{\Sigma}_{hj}^{oo(\ell)})}.$$

Consequently, a Bayesian classifier for \mathbf{Y}_j can be estimated by averaging over the draws of $\Theta^{(\ell)}$

$$\hat{r}_{ij}^* = \Pr(Z_{ij} = 1 | \mathbf{Y}_j^o) \approx \frac{1}{L} \sum_{\ell=1}^L r_{ij}^{(\ell)}. \tag{21}$$

By the Bayesian classification rule, \mathbf{Y}_j is assigned to \mathcal{C}_s if $\hat{r}_{sj}^* > \hat{r}_{ij}^*$ ($i = 1, \dots, g; i \neq s$).

Table 1

A comparison of CPU time (in seconds) and relative reduced time (RRT) between GJ-EM algorithm (old) and the proposed procedure (new) under various missing rates (Replications = 500)

Data	$\eta = 10\%$			$\eta = 20\%$			$\eta = 30\%$		
	Old	New	RRT (%)	Old	New	RRT (%)	Old	New	RRT (%)
<i>Iris</i>	12.47	1.22	90.2	21.51	1.61	92.5	56.21	3.61	93.6
<i>Crabs</i>	34.72	3.27	90.6	78.77	6.78	91.4	265.01	20.68	92.2

$$RRT = (\text{old} - \text{new})/\text{old} \times 100\%$$

Table 2

A comparison of prediction accuracies for MI, EM and DA imputations with the standard deviations in parentheses for the *iris* data set (Replications=500)

η (%)	MAE			MARE			RMSE		
	MI	EM	DA	MI	EM	DA	MI	EM	DA
10	0.812 (0.081)	0.213 (0.026)	0.210 (0.026)	0.697 (0.186)	0.100 (0.027)	0.099 (0.027)	1.062 (0.096)	0.285 (0.050)	0.280 (0.050)
20	0.816 (0.053)	0.237 (0.025)	0.233 (0.025)	0.675 (0.129)	0.114 (0.031)	0.113 (0.031)	1.071 (0.065)	0.331 (0.060)	0.326 (0.060)
30	0.820 (0.046)	0.268 (0.023)	0.259 (0.022)	0.684 (0.097)	0.138 (0.033)	0.132 (0.032)	1.078 (0.058)	0.395 (0.061)	0.380 (0.060)
40	0.819 (0.035)	0.301 (0.030)	0.278 (0.026)	0.683 (0.082)	0.161 (0.038)	0.154 (0.036)	1.077 (0.041)	0.448 (0.065)	0.428 (0.063)
50	0.817 (0.029)	0.346 (0.031)	0.325 (0.028)	0.675 (0.084)	0.198 (0.043)	0.188 (0.041)	1.074 (0.036)	0.522 (0.063)	0.495 (0.060)

Table 3

A comparison of prediction accuracies for MI, EM and DA imputations with the standard deviations in parentheses for the *crabs* data set (Replications=500)

η (%)	MAE			MARE			RMSE		
	MI	EM	DA	MI	EM	DA	MI	EM	DA
10	4.063 (0.337)	0.421 (0.055)	0.415 (0.050)	0.202 (0.018)	0.024 (0.003)	0.023 (0.003)	5.391 (0.427)	0.611 (0.114)	0.598 (0.105)
20	4.008 (0.227)	0.484 (0.041)	0.474 (0.037)	0.200 (0.012)	0.027 (0.002)	0.026 (0.002)	5.343 (0.305)	0.714 (0.090)	0.693 (0.083)
30	4.037 (0.169)	0.568 (0.044)	0.550 (0.041)	0.202 (0.009)	0.030 (0.002)	0.029 (0.002)	5.384 (0.225)	0.846 (0.096)	0.812 (0.091)
40	4.036 (0.138)	0.662 (0.044)	0.632 (0.042)	0.203 (0.007)	0.035 (0.002)	0.033 (0.002)	5.381 (0.188)	0.977 (0.092)	0.932 (0.094)
50	4.039 (0.108)	0.768 (0.052)	0.728 (0.050)	0.202 (0.006)	0.039 (0.002)	0.037 (0.002)	5.386 (0.142)	1.120 (0.102)	1.058 (0.100)

5. Experimental results

For illustration purposes, we start to apply results developed in Sections 2–4 to two famous multivariate data sets. One is the *iris* data taken from Anderson [23] or Fisher [24]. It consists of four-dimensional measurements in centimeters on the attributes of petal length, petal width, sepal length and sepal width for 50 flower specimens of each of three species: *setosa*, *versicolor*, and *virginica*. The other is the *crabs* data of Campbell and Mahon [25] on the *gensus Leptograpsus*. It consists of

Table 4

A comparison of average misclassification rates (%) between ML and Bayesian classifiers (replicates = 500)

η (%)	Iris		Crabs	
	ML	Bayesian	ML	Bayesian
0	3.33	3.00	7.50	7.30
10	3.85	3.75	9.75	9.50
20	5.20	5.00	13.66	13.55
30	6.90	6.10	19.22	18.80
40	10.15	9.20	26.75	25.20
50	13.42	12.30	35.21	33.00

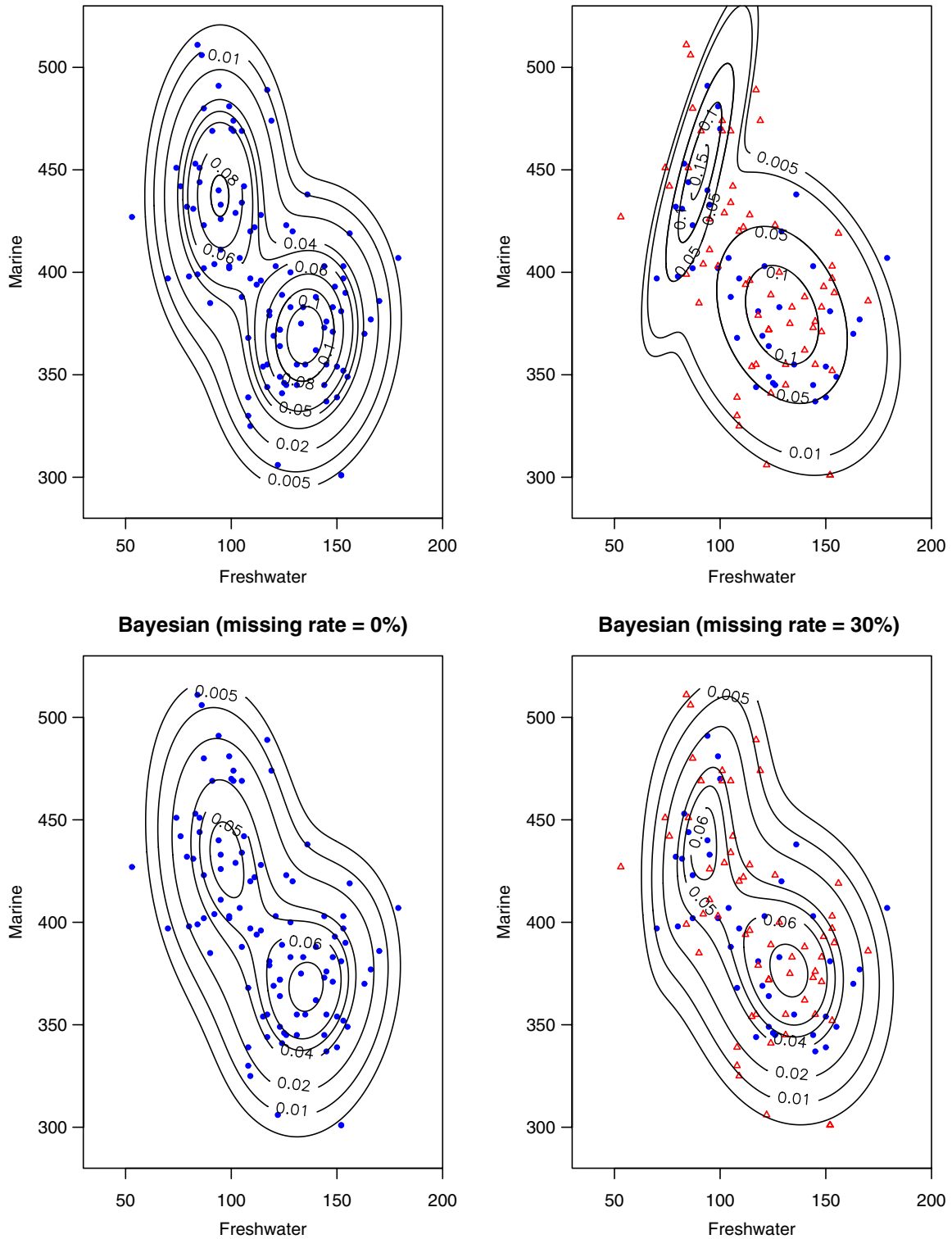


Fig. 1. ML and Bayesian density estimation for the two-component salmon data (●, both attributes are completely observed; △, one of the two attributes is missing).

five-dimensional morphological measurements on the attributes of width of frontal lip, rear width, length along the mid-line of the carapace, maximum width of the carapace and body depth for 50 crabs of each of four groups: blue

male, blue female, orange male and orange female. Both data sets are included as a part of the R package, which is freely available at the web site <http://cran.r-project.org>.

To conduct experimental studies, we first generate 500 artificially missing data sets by deleting at random from the three data sets under various specified missing rate η (proportion of missing values) while we maintain each datum to have at least one observed attribute. Table 1 presents the computation times of our developed EM algorithm and those of using GJ-EM. All computations are solely carried out by R package in the environment of a desktop PC (CPU: 3Gb-MHz/Intel Pentium 4 Processor; RAM: 1Gb/DDR-400). Since the programming implementations have many characteristics (e.g., vector or matrix subroutines instead of loops), the CPU times in Table 1 might not be directly comparable, but provide a sense of their actual performances in a practical setting. As seen in the table, all computation times are dramatically reduced over 90% by using the new EM procedure.

To exemplify the predictive performance for the EM and DA imputation methods, see Eqs. (13) and (20), together with the traditional mean imputation (MI) method, known as “filling-in” with the sample mean of the associated attribute, we utilize the pseudo-cross-validation (PSV) of Stone [26] to evaluate these three approaches. A relative tolerance of 10^{-8} for the log-likelihood function and parameter estimates are used as the convergence criterion for the EM algorithm. As for the DA algorithm, we take the ML estimates as the initialization and carry out 2000 iterations with the first 1000 iterations as burn-in and the remaining 1000 iterations as inference samples. It is noted that our chosen burn-in number is much larger than needed based on checking the *multivariate potential scale reduction factor* (MPSRF) of Brooks and Gelman [27]. As for discrepancy measures, we use the mean absolute error (MAE), the mean absolute relative error (MARE) and root mean square error (RMSE). Comparison results are listed in Tables 2 and 3. As seen in the tables, we found that both EM and DA substantially outperform MI for all cases. Furthermore, DA imputation exhibits considerable promising accuracy in the prediction of missing values when compared to the EM imputation, especially as the size of observed values becomes small (i.e., missing rate increases).

As another illustration, we attempt to explore classification accuracies between the ML classifier Eq. (12) and the Bayesian classifier Eq. (21) via PSV. Experimental results in Table 4 indicate that both classifiers are comparable at low-level missing, but Bayesian classifier yields lower misclassification rates as the missing rate increases, though improvements are not substantial.

Finally, we are interested in comparing behaviors of density estimation from both ML-fitted and Bayesian posterior predictive aspects. To illustrate this, we use the *salmon* data taken from Johnson and Wichern [28]. This data set has two attributes, the diameter of rings for the first-year freshwater growth and the diameter of rings for the first-year marine growth (both measured in hundredths of an inch), for each of 50 Alaskan-born and Canadian-born salmon fishes. The ML-fitted density estimation is obtained by

plugging the ML estimates into Eq. (1). As for Bayesian predictive density, it can be approximated by the use of Rao-Blackwellization

$$\begin{aligned} p(\mathbf{y}|\mathbf{Y}^o) &= \int p(\mathbf{y}|\mathbf{Y}^o, \Theta) p(\Theta|\mathbf{Y}^o) d\Theta \\ &\approx \frac{1}{L} \sum_{\ell=1}^L p(\mathbf{y}|\Theta^{(\ell)}) \\ &= \frac{1}{L} \sum_{\ell=1}^L \left(\sum_{i=1}^g w_i^{(\ell)} \left((2\pi)^{-p/2} |\Sigma_i^{(\ell)}|^{-1/2} \right. \right. \\ &\quad \left. \left. \times \exp\left(-\frac{1}{2} \Delta_{ij}^{(\ell)}\right) \right) \right), \end{aligned} \quad (22)$$

where $\Delta_{ij}^{(\ell)} = (\mathbf{y} - \mu_i^{(\ell)})^\top \Sigma_i^{(\ell)-1} (\mathbf{y} - \mu_i^{(\ell)})$ and $\Theta^{(\ell)}$ ($\ell = 1, \dots, L$) is a set of converged Monte Carlo samples generated from the DA algorithm.

The contour plots obtained by the ML-fitting and Bayesian predictive densities Eq. (22) for both completely observed data ($\eta=0\%$) and partially observed data ($\eta=30\%$) are depicted in Fig. 1, respectively. Both look similar when data are not missing but using Eq. (22) seems to have a relatively smoother appearance. In addition, we found that the ML-fitted contour shapes tend to be distorted at high-level missing and even for moderate-level missing ($\eta=30\%$). However, the distortion rarely happened while using Eq. (22). This indicates that Bayesian learning is more resistant to missing values.

6. Conclusions

In this paper, two novel EM and DA computational algorithms for learning normal mixture models under a missing information framework are presented. It should be emphasized that our proposed procedures offer neat ways to program with low-cost computation. Experimental results indicate that Bayesian treatment is a worthwhile tool for mixture modelling under a considerable extent of missing information.

Recently, Bayesian and non-Bayesian robust mixture model modelling using the t distribution has received notable attentions, see Refs. [29–32]. Future work will make some kind of comparisons theoretically or empirically among various competitive models.

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Appendix A. Proof of Proposition 2

Suppose $Y \sim N_p(\mu, \Sigma)$, then for any $q \times p$ matrix A with rank $q (q \leq p)$, we can obtain $AY \sim N_p(A\mu, A\Sigma A^T)$. With similar arguments, the marginal distributions of Y_j^o and Y_j^m are:

$$Y_j^o = O_j Y_j \sim \sum_{i=1}^g w_i \phi_{p_j^o}(\mu_{ij}^o, \Sigma_{ij}^{oo}), \quad \mu_{ij}^o = O_j \mu_i,$$

$$\Sigma_{ij}^{oo} = O_j \Sigma_i O_j^T,$$

$$Y_j^m = M_j Y_j \sim \sum_{i=1}^g w_i \phi_{p-p_j^o}(\mu_{ij}^m, \Sigma_{ij}^{mm}), \quad \mu_{ij}^m = M_j \mu_i,$$

$$\Sigma_{ij}^{mm} = M_j \Sigma_i M_j^T.$$

Note that the Δ_{ij} in Eq. (2) can be reexpressed as

$$\Delta_{ij} = \begin{bmatrix} Y_j^o - \mu_{ij}^o \\ Y_j^m - \mu_{ij}^m \end{bmatrix}^T \begin{bmatrix} \Sigma_{ij}^{oo} & \Sigma_{ij}^{om} \\ \Sigma_{ij}^{mo} & \Sigma_{ij}^{mm} \end{bmatrix}^{-1} \begin{bmatrix} Y_j^o - \mu_{ij}^o \\ Y_j^m - \mu_{ij}^m \end{bmatrix}, \quad (23)$$

where $\Sigma_{ij}^{om} = O_j \Sigma_i M_j^T$ and $\Sigma_{ij}^{mo} = M_j \Sigma_i O_j^T$. Also, the second and third factors on the right hand side of Eq. (23) can be represented by

$$\begin{bmatrix} \Sigma_{ij}^{oo} & \Sigma_{ij}^{om} \\ \Sigma_{ij}^{mo} & \Sigma_{ij}^{mm} \end{bmatrix}^{-1} = \begin{bmatrix} I & -\Sigma_{ij}^{oo^{-1}} \Sigma_{ij}^{om} \\ \mathbf{0} & I \end{bmatrix} \begin{bmatrix} \Sigma_{ij}^{oo^{-1}} & \mathbf{0} \\ \mathbf{0} & \Sigma_{ij}^{mm-o^{-1}} \end{bmatrix} \times \begin{bmatrix} I & \mathbf{0} \\ -\Sigma_{ij}^{mo} \Sigma_{ij}^{oo^{-1}} & I \end{bmatrix},$$

and

$$\begin{bmatrix} Y_j^o - \mu_{ij}^o \\ Y_j^m - \mu_{ij}^m \end{bmatrix} = \begin{bmatrix} O_j(Y_j - \mu_i) \\ M_j(Y_j - \mu_i) \end{bmatrix} = \begin{bmatrix} O_j \\ M_j \end{bmatrix} (Y_j - \mu_i).$$

We then have the following standard results:

$$\Sigma_{ij}^{mm-o} = \Sigma_{ij}^{mm} - \Sigma_{ij}^{mo} \Sigma_{ij}^{oo^{-1}} \Sigma_{ij}^{om}$$

$$= M_j \Sigma_i M_j^T - M_j \Sigma_i O_j^T (O_j \Sigma_i O_j^T)^{-1} O_j \Sigma_i M_j^T$$

$$= M_j (I_p - \Sigma_i S_{ij}^{oo}) \Sigma_i M_j^T = E_{ij} \Sigma_i M_j^T,$$

where $E_{ij} = M_j (I_p - \Sigma_i S_{ij}^{oo})$, $S_{ij}^{oo} = O_j^T (O_j \Sigma_i O_j^T)^{-1} O_j$.

Since

$$-\Sigma_{ij}^{mo} \Sigma_{ij}^{oo^{-1}} O_j + M_j = M_j - M_j \Sigma_i O_j^T (O_j \Sigma_i O_j^T)^{-1} O_j$$

$$= M_j (I_p - \Sigma_i S_{ij}^{oo})$$

$$= E_{ij}$$

and

$$\begin{bmatrix} I & \mathbf{0} \\ -\Sigma_{ij}^{mo} \Sigma_{ij}^{oo^{-1}} & I \end{bmatrix} \begin{bmatrix} O_j \\ M_j \end{bmatrix} = \begin{bmatrix} O_j \\ E_{ij} \end{bmatrix},$$

it suffices to show that

$$\mu_{ij}^{m-o} = \mu_{ij}^m + \Sigma_{ij}^{mo} \Sigma_{ij}^{oo^{-1}} (Y_j^o - \mu_{ij}^o)$$

$$= M_j \mu_i + M_j \Sigma_i O_j^T (O_j \Sigma_i O_j^T)^{-1} O_j (Y_j - \mu_i)$$

$$= M_j (\mu_i + \Sigma_i S_{ij}^{oo} (Y_j - \mu_i)).$$

Hence,

$$\Delta_{ij} = (Y_j^o - \mu_{ij}^o)^T \Sigma_{ij}^{oo^{-1}} (Y_j^o - \mu_{ij}^o)$$

$$+ (Y_j^m - \mu_{ij}^{m-o})^T \Sigma_{ij}^{mm-o^{-1}} (Y_j^m - \mu_{ij}^{m-o})$$

$$= (Y_j - \mu_i)^T (S_{ij}^{oo} + S_{ij}^{mm-o}) (Y_j - \mu_i)$$

$$= \Delta_{ij}^o + \Delta_{ij}^{m-o},$$

where

$$\Delta_{ij}^o = (Y_j^o - \mu_{ij}^o)^T \Sigma_{ij}^{oo^{-1}} (Y_j^o - \mu_{ij}^o)$$

$$= (Y_j - \mu_i)^T S_{ij}^{oo} (Y_j - \mu_i),$$

$$\Delta_{ij}^{m-o} = (Y_j^m - \mu_{ij}^{m-o})^T \Sigma_{ij}^{mm-o^{-1}} (Y_j^m - \mu_{ij}^{m-o})$$

$$= (Y_j - \mu_i)^T S_{ij}^{mm-o} (Y_j - \mu_i),$$

$$S_{ij}^{mm-o} = E_{ij}^T (E_{ij} \Sigma_i M_j^T)^{-1} E_{ij}.$$

Using the fact that $|\Sigma_i| = |\Sigma_{ij}^{oo}| |\Sigma_{ij}^{mm-o}|$ and above results, we have

$$f(Y_j^m | Y_j^o) = \frac{f(Y_j)}{f(Y_j^o)}$$

$$= \frac{\sum_{i=1}^g w_i \phi_{p_j^o}(Y_j^o | \mu_{ij}^o, \Sigma_{ij}^{oo}) \phi_{p-p_j^o}(Y_j^m | Y_j^o, \mu_{ij}^{m-o}, \Sigma_{ij}^{mm-o})}{\sum_{i=1}^g w_i \phi_{p_j^o}(Y_j^o | \mu_{ij}^o, \Sigma_{ij}^{oo})}$$

$$= \sum_{i=1}^g w_{ij}^* \phi_{p-p_j^o}(Y_j^m | Y_j^o, \mu_{ij}^{m-o}, \Sigma_{ij}^{mm-o}),$$

where $w_{ij}^* = w_i \phi_{p_j^o}(Y_j^o | \mu_{ij}^o, \Sigma_{ij}^{oo}) / \sum_{h=1}^g w_h \phi_{p_j^o}(Y_j^o | \mu_{hj}^o, \Sigma_{hj}^{oo})$.

Appendix B. Proof of Proposition 3

Letting $\hat{Z}_{ij}^{(k)} = E(Z_{ij} | Y^o, \hat{\Theta}^{(k)})$, $\xi_{ij}^{(k)} = E(Z_{ij} Y_j | Y^o, \hat{\Theta}^{(k)})$ and $\hat{\Theta}_{ij}^{(k)} = E(Z_{ij} Y_j Y_j^T | Y^o, \hat{\Theta}^{(k)})$, we can show that

$$\hat{Z}_{ij}^{(k)} = \Pr(Z_{ij} = 1 | Y_j^o, \hat{\Theta}^{(k)})$$

$$= \frac{\hat{w}_i^{(k)} \phi_{p_j^o}(Y_j^o | \hat{\mu}_{ij}^{o(k)}, \hat{\Sigma}_{ij}^{oo(k)})}{\sum_{h=1}^g \hat{w}_h^{(k)} \phi_{p_j^o}(Y_j^o | \hat{\mu}_{hj}^{o(k)}, \hat{\Sigma}_{hj}^{oo(k)})},$$

$$\xi_{ij}^{(k)} = \Pr(Z_{ij} = 1 | Y^o, \hat{\Theta}^{(k)}) E[Y_j | Z_{ij} = 1, Y^o, \hat{\Theta}^{(k)}]$$

$$= E(Z_{ij} | Y^o, \hat{\Theta}^{(k)}) E(Y_j | Z_{ij} = 1, Y^o, \hat{\Theta}^{(k)})$$

$$= \hat{Z}_{ij}^{(k)} \hat{Y}_{ij}^{(k)},$$

and

$$\begin{aligned} \hat{\Phi}_{ij}^{(k)} &= E(Z_{ij}Y_jY_j^\top | Y^o, \hat{\Theta}^{(k)}) \\ &= E(Z_{ij}|Y^o, \hat{\Theta}^{(k)})E(Y_jY_j^\top | Z_{ij} = 1, Y^o, \hat{\Theta}^{(k)}) \\ &= \hat{Z}_{ij}^{(k)}(\hat{Y}_{ij}^{(k)}\hat{Y}_{ij}^{(k)\top} + \text{Cov}(Y_j|Z_{ij} = 1, Y^o, \hat{\Theta}^{(k)})) \\ &= \hat{Z}_{ij}^{(k)}(\hat{Y}_{ij}^{(k)}\hat{Y}_{ij}^{(k)\top} + (I_p - \hat{\Sigma}_i^{(k)}\hat{S}_{ij}^{\text{oo}(k)})\hat{\Sigma}_i^{(k)}). \end{aligned}$$

Since $Y_j = O_j^\top Y_j^o + M_j^\top Y_j^m$ and $O_j^\top O_j(I_p - \Sigma_i S_{ij}^{\text{oo}}) = \mathbf{0}$, we have

$$\begin{aligned} \hat{Y}_{ij}^{(k)} &= E(Y_j | Z_{ij} = 1, Y^o, \hat{\Theta}^{(k)}) \\ &= E(O_j^\top Y_j^o + M_j^\top Y_j^m | Z_{ij} = 1, Y^o, \hat{\Theta}^{(k)}) \\ &= O_j^\top Y_j^o + M_j^\top E(Y_j^m | Z_{ij} = 1, Y^o, \hat{\Theta}^{(k)}) \\ &= O_j^\top Y_j^o + M_j^\top \hat{\mu}_{ij}^{\text{m-o}(k)} \\ &= O_j^\top O_j Y_j + M_j^\top M_j (\hat{\mu}_i^{(k)} + \hat{\Sigma}_i^{(k)} \hat{S}_{ij}^{\text{oo}(k)} (Y_j - \hat{\mu}_i^{(k)})) \\ &= O_j^\top O_j Y_j + (I_p - O_j^\top O_j) (\hat{\mu}_i^{(k)} \\ &\quad + \hat{\Sigma}_i^{(k)} \hat{S}_{ij}^{\text{oo}(k)} (Y_j - \hat{\mu}_i^{(k)})) \\ &= \hat{\mu}_i^{(k)} + \hat{\Sigma}_i^{(k)} \hat{S}_{ij}^{\text{oo}(k)} (Y_j - \hat{\mu}_i^{(k)}) \\ &\quad + O_j^\top O_j (I_p - \hat{\Sigma}_i^{(k)} \hat{S}_{ij}^{\text{oo}(k)}) (Y_j - \hat{\mu}_i^{(k)}) \\ &= \hat{\mu}_i^{(k)} + \hat{\Sigma}_i^{(k)} \hat{S}_{ij}^{\text{oo}(k)} (Y_j - \hat{\mu}_i^{(k)}), \end{aligned}$$

and

$$\begin{aligned} \text{Cov}(Y_j | Z_{ij} = 1, Y^o, \hat{\Theta}^{(k)}) &= \text{Cov}(O_j^\top Y_j^o + M_j^\top Y_j^m | Z_{ij} = 1, Y^o, \hat{\Theta}^{(k)}) \\ &= M_j^\top \text{Cov}(Y_j^m | Z_{ij} = 1, Y^o, \hat{\Theta}^{(k)}) M_j \\ &= M_j^\top \hat{\Sigma}_{ij}^{\text{mm-o}(k)} M_j \\ &= M_j^\top \hat{E}_{ij} \hat{\Sigma}_i \hat{M}_j \\ &= M_j^\top M_j (I_p - \hat{\Sigma}_i^{(k)} \hat{S}_{ij}^{\text{oo}(k)}) \hat{\Sigma}_i \hat{M}_j \\ &= (I_p - O_j^\top O_j) (I_p - \hat{\Sigma}_i^{(k)} \hat{S}_{ij}^{\text{oo}(k)}) \hat{\Sigma}_i (I_p - O_j^\top O_j) \\ &= (I_p - \hat{\Sigma}_i^{(k)} \hat{S}_{ij}^{\text{oo}(k)}) \hat{\Sigma}_i (I_p - O_j^\top O_j) \\ &= (I_p - \hat{\Sigma}_i^{(k)} \hat{S}_{ij}^{\text{oo}(k)}) \hat{\Sigma}_i^{(k)}. \end{aligned}$$

Therefore,

$$\begin{aligned} \Omega_{ij}^{(k)} &= \hat{\Phi}_{ij}^{(k)} - \hat{\xi}_{ij}^{(k)} \mu_i^\top - \mu_i \hat{\xi}_{ij}^{(k)\top} + \hat{Z}_{ij}^{(k)} \mu_i \mu_i^\top \\ &= \hat{Z}_{ij}^{(k)} (\hat{Y}_{ij}^{(k)} \hat{Y}_{ij}^{(k)\top} + (I_p - \hat{\Sigma}_i^{(k)} \hat{S}_{ij}^{\text{oo}(k)}) \hat{\Sigma}_i^{(k)}) \\ &\quad - 2 \hat{Z}_{ij}^{(k)} \hat{Y}_{ij}^{(k)} \mu_i^\top + \hat{Z}_{ij}^{(k)} \mu_i \mu_i^\top \\ &= \hat{Z}_{ij}^{(k)} (\hat{Y}_{ij}^{(k)} - \mu_i) (\hat{Y}_{ij}^{(k)} - \mu_i)^\top \\ &\quad + (I_p - \hat{\Sigma}_i^{(k)} \hat{S}_{ij}^{\text{oo}(k)}) \hat{\Sigma}_i^{(k)}. \end{aligned}$$

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