Robust mean and covariance matrix estimation under heterogeneous mixed-effects model with missing values

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Abstract

In this paper, robust mean and covariance matrix estimation are considered in the context of mixed-effects models. Such models are widely used to analyze repeated measures data which arise in several signal processing applications that need to incorporate possible individual variations within a common behavior of individuals. In this context, most algorithms are based on the assumption that the observations follow a Gaussian distribution. Nevertheless, in certain situations in which the data set contains outliers, such assumption is not valid and leads to a dramatic performance loss. To overcome this drawback, we design an expectation-conditional maximization either algorithm in which the heterogeneous component is considered as a part of the complete data. Then, the proposed algorithm is cast into a parallel scheme w.r.t. the individuals in order to mitigate the computational cost and a possible central processor overload. Finally, the proposed algorithm is extended to deal with missing data which refers to the situation where part of the individual responses are unobserved. Numerical simulations are conducted to assess the performance of the proposed algorithm regarding robust regression estimators, probabilistic principal component analysis and its recent robust version.

Keywords: Maximum likelihood, expectation maximization, robust mean estimation.

1. Introduction

The use of mixed-effects models has become popular in a wide range of signal processing applications [1]. Specifically, it is mainly used in applications which require to

model a common behavior of individuals with possible individual variations. This reads,

$$\mathbf{y}_{ij} = \mathbf{A}_i\mathbf{m} + \mathbf{Z}_i\mathbf{t}_{ij} + \mathbf{n}_{ij}, \quad \forall j = 1, \ldots, q \text{ and } i = 1, \ldots, s, \tag{1}$$

where $\mathbf{y}_{ij} \in \mathbb{C}^{n_i}$ denotes the complex vector of the j-th observation at the i-th individual, $\mathbf{t}_{ij} \sim \mathcal{CN}(\mathbf{0}, \mathbf{I})$ and $\mathbf{n}_{ij} \sim \mathcal{CN}(\mathbf{0}, \sigma^2\mathbf{I})$ characterize the individual errors. Finally, $\mathbf{A}_i \in \mathbb{C}^{n_i \times p}$, $\mathbf{Z}_i \in \mathbb{C}^{n_i \times q_i}$ and $\mathbf{m} \in \mathbb{C}^p$ denote the known design matrix related to the fixed-effect $\mathbf{m}$, the unknown design matrix related to the random-effect $\mathbf{t}_{ij}$ and the common unknown vector mean, respectively.

As an example in signal processing, we can cite the imaging context of the next generation of radio-interferometers. Such instruments are composed of several spaced stations. Each station, which is a collection of low band and/or high band antennas, represents one individual. In this case, the design matrix $\mathbf{A}_i$ is a known linear operator – related to the antenna geometry – that maps the image from the space domain to the visibility (observation) domain [2]. The mean $\mathbf{m}$ denotes the intensity vector, i.e. the unknown common image observed by each station. The heterogeneous mixed-effects come from that each station can be locally affected by different man-made radio frequency interferences (RFI). An additional application of the model (1) in the case of only one individual ($s = 1$) can be found in the so-called probabilistic principal component analysis literature [3].

In the aforementioned references, the authors considered normally distributed observations. Nevertheless, such assumption is not realistic in a plethora of signal processing applications as those related to high resolution sensing systems, non-homogeneous environments or in the possible presence of outliers [4, 5].

In this paper, we consider a parametric model that takes into account the possible presence of outliers by modeling the within-individual error as a mixture of a Gaussian process and a non-Gaussian distributed noise lying in a low-rank covariance matrix. The mixed-effects model gives the flexibility to assign outliers to some of the individuals only, depending on their environment. Consequently, the proposed model becomes

$$\mathbf{y}_{ij} = \mathbf{A}_i\mathbf{m} + \sqrt{\tau_{ij}}\mathbf{W}_i\mathbf{t}_{ij} + \mathbf{n}_{ij} \tag{2}$$

in which $\mathbf{t}_{ij} \sim \mathcal{CN}(0, \boldsymbol{I})$ represents the low-rank heterogeneous random effect error related to a positive texture parameter τ_{ij} and a low-rank loading matrix $\mathbf{W}_i \in \mathbb{C}^{n_i \times r_i}$ of

rank $r_i \leq n_i$. The heterogeneous component, $\sqrt{\tau_{ij}}\mathbf{W}_i\mathbf{t}_{ij}$, leads to a compound Gaussian random effect error in the case of a random texture parameter, or its counterpart referred to as heteroscedastic Gaussian random effect error if the texture parameter is assumed unknown and deterministic [6]. Such process formulation became popular since it owns a great flexibility allowing to gather several elliptically symmetric distributions, e.g., Gaussian, K- and t-distributions, Cauchy distributions, etc. Again, considering the examples given above, the non-Gaussian random effect error can model the clutter (the external noise), $\sqrt{\tau_{ij}}\mathbf{W}_i\mathbf{t}_{ij}$, plus thermal noise (the internal noise) $\mathbf{n}_{ij}$ which exists in several array/radar processing applications, or the unknown background of power-fluctuating sources present in the imaging process of the recent radio-interferometers [7, 8, 9].

To accurately tackle the estimation procedure related to model (2), we use the Expectation-Conditional Maximisation Either (ECME) algorithm. We select in a proper way the complete data, which is composed of the observed variables $\mathbf{y}_{ij}$ and the latent variables $\mathbf{t}_{ij}$ (and the missing observations in the case of missing data) which leads to closed-form expressions in the E-step and M-step. Then, the proposed estimator is cast into a parallel scheme to lower the computational cost and avoid a possible central processor overwhelm.

2. ECME-based estimation under heterogeneous mixed-effects model

Considering model (2), the unknown vector parameter reads, with some abuse of notation, $\boldsymbol{\theta} = \left[\mathbf{m}^T, \left\{\boldsymbol{\zeta}_i^T\right\}_i, \left\{\tau_{ij}\right\}_{ij}, \sigma^2\right]^T$, in which $\boldsymbol{\zeta}_i$ is the concatenation of the non-redundant elements in $\mathbf{W}_i$ (the estimation of $\mathbf{W}_i$ is known to display a rotational ambiguity which is discussed later in Section 3) and where the texture parameters $\left\{\tau_{ij}\right\}_{ij}$ are considered deterministic and unknown. The latter assumption ensures more tractability as the texture distribution is not specified, which avoids any possible model misspecification. We consider the maximum likelihood principal, that is

$$\boldsymbol{\theta}_{\mathrm{ML}} = \arg\max_{\boldsymbol{\theta}} \mathcal{L}(\{\mathbf{y}_{ij}\}_{ij}\,|\boldsymbol{\theta}) \tag{3}$$

where $\mathcal{L}(\{\mathbf{y}_{ij}\}_{ij}\,|\boldsymbol{\theta}) = -\sum_i \sum_j \log|\mathbf{C}_{ij}| - (\mathbf{y}_{ij} - \mathbf{A}_i\mathbf{m})^H \mathbf{C}_{ij}^{-1} (\mathbf{y}_{ij} - \mathbf{A}_i\mathbf{m})$ with $\mathbf{C}_{ij} = \tau_{ij}\mathbf{W}_i\mathbf{W}_i^H + \sigma^2\mathbf{I}$ in which both independence between individuals and between observations are assumed. It is clear that solving (3) is challenging due to the non-convexity of

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the objective function. Consequently, we propose hereafter the use of the ECME algorithm, which is known to be an efficient extension of the EM scheme with faster monotone convergence [10]. The ECME algorithm is an iterative algorithm whose estimates $\boldsymbol{\theta}^{(m)}$ converge, under certain mild conditions, to the maximum likelihood estimate $\boldsymbol{\theta}_{\text{ML}}$ (m denoting the iteration number). The ECME algorithm is decomposed in two steps: the E-step and the M-step. In the E-step, we derive the surrogate function $Q(.|.)$, which is the expectation of the log-likelihood of the complete data $\mathcal{L}_{\text{C}}$ conditioned on the observed data and the previously computed $\boldsymbol{\theta}^{(m-1)}$. The complete data is a combination of the observed data $\mathbf{y}_{ij}$ and the missing/latent data $\mathbf{t}_{ij}$. While the classical M-step requires to maximize Q, we also consider maximizing the loglikelihood $\mathcal{L}$ of the incomplete data [10] depending on the ease of derivation w.r.t a block of the unknown vector parameter $\boldsymbol{\theta}$. This procedure is then repeated until convergence. The E- and the M-step are now described.

E-step: First, the complete data must be specified in order to simplify the M-step while maintaining the derivation of the expectation of $\mathcal{L}_{\text{C}}$ feasible. Based on (2), it seems natural to choose the complete data as $\mathbf{x}_{ij} = [\mathbf{y}_{ij}^T, \mathbf{t}_{ij}^T]^T$. Consequently,

$$
\begin{aligned}
\mathcal{L}_{\text{C}}(\{\mathbf{x}_{ij}\}_{ij}|\boldsymbol{\theta}) &= \sum_i \sum_j \log p(\mathbf{y}_{ij}|\mathbf{t}_{ij}, \boldsymbol{\theta}) + \log p(\mathbf{t}_{ij}|\boldsymbol{\theta}) \\
&\propto -\sum_i \sum_j n_i \log \sigma^2 - ||\mathbf{r}_{ij} - \sqrt{\tau_{ij}}\mathbf{W}_i \mathbf{t}_{ij}||^2_{\sigma^2} - ||\mathbf{t}_{ij}||^2_2
\end{aligned}
\tag{4}
$$

in which $\mathbf{r}_{ij} = \mathbf{y}_{ij} - \mathbf{A}_i \mathbf{m}$ and the weighted norm reads $||\mathbf{b}||^2_{\mathbf{B}} = \mathbf{b}\mathbf{B}^{-1}\mathbf{b}^H$. Thus, the so-called Q function reads

$$
\begin{aligned}
Q(\boldsymbol{\theta}|\boldsymbol{\theta}^{(m)}) &= E_{\{\mathbf{x}_{ij}\}_{ij}|\{\mathbf{y}_{ij}\}_{ij},\boldsymbol{\theta}^{(m)}}\left\{\mathcal{L}_{\text{C}}(\{\mathbf{x}_{ij}\}_{ij}|\boldsymbol{\theta})\right\} \\
&= E_{\{\mathbf{t}_{ij}\}_{ij}|\{\mathbf{y}_{ij}\}_{ij},\boldsymbol{\theta}^{(m)}}\left\{\mathcal{L}_{\text{C}}(\{\mathbf{x}_{ij}\}_{ij}|\boldsymbol{\theta})\right\} = \sum_i \sum_j Q_{ij}(\boldsymbol{\theta}|\boldsymbol{\theta}^{(m)})
\end{aligned}
$$

where $Q_{ij}(\boldsymbol{\theta}|\boldsymbol{\theta}^{(m)}) = E_{\mathbf{t}_{ij}|\mathbf{y}_{ij},\boldsymbol{\theta}^{(m)}}\{\log p(\mathbf{x}_{ij}|\boldsymbol{\theta})\}$. First, let us derive the pdf of the latent variable $\mathbf{t}_{ij}$ conditioned on the observation and $\boldsymbol{\theta}^{(m)}$. Namely, since $p(\mathbf{t}_{ij}|\mathbf{y}_{ij}, \boldsymbol{\theta}^{(m)}) \propto p(\mathbf{y}_{ij}|\mathbf{t}_{ij}, \boldsymbol{\theta}^{(m)})p(\mathbf{t}_{ij}|\boldsymbol{\theta}^{(m)})$, after some calculus and considering the adequate normalization constant, we obtain

$$
p(\mathbf{t}_{ij}|\mathbf{y}_{ij}, \boldsymbol{\theta}^{(m)}) \sim \mathcal{CN}\left(\mathbf{V}_{ij}^{-1^{(m)}}\mathbf{u}_{ij}^{(m)}, \mathbf{V}_{ij}^{-1^{(m)}}\right)
\tag{5}
$$

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where

$$\mathbf{u}_{ij}^{(m)} = \frac{\sqrt{\tau_{ij}^{(m)}}}{(\sigma^{(m)})^2} \mathbf{W}_i^{H^{(m)}} \mathbf{r}_{ij}^{(m)}$$

$$\mathbf{V}_{ij}^{(m)} = \mathbf{I} + \frac{\tau_{ij}^{(m)}}{(\sigma^{(m)})^2} \mathbf{W}_i^{H^{(m)}} \mathbf{W}_i^{(m)}$$

Consequently, the surrogate function reads

$$Q_{ij}(\boldsymbol{\theta}|\boldsymbol{\theta}^{(m)}) = E_{\mathbf{t}_{ij}|\mathbf{y}_{ij},\boldsymbol{\theta}^{(m)}} \left\{ -n_i \log \sigma^2 - \mathrm{Tr}\left(\mathbf{t}_{ij}\mathbf{t}_{ij}^H \mathbf{V}_{ij}\right) - ||\mathbf{r}_{ij}||_{\sigma^2}^2 + 2\frac{\sqrt{\tau_{ij}}}{\sigma^2}\Re(\mathbf{r}_{ij}^H \mathbf{W}_i \mathbf{t}_{ij}) \right\}$$

$$= -n_i \log \sigma^2 - \mathrm{Tr}\left(\widehat{\mathbf{T}}_{ij}^{(m)} \mathbf{V}_{ij}\right) - ||\mathbf{r}_{ij}||_{\sigma^2}^2 + \frac{2\sqrt{\tau_{ij}}}{\sigma^2}\Re(\mathbf{r}_{ij}^H \mathbf{W}_i \widehat{\mathbf{t}}_{ij}^{(m)}) \qquad (6)$$

in which

$$\widehat{\mathbf{t}}_{ij}^{(m)} = E_{\mathbf{t}_{ij}|\mathbf{y}_{ij},\boldsymbol{\theta}^{(m)}} \left\{ \mathbf{t}_{ij} \right\} = \mathbf{V}_{ij}^{-1^{(m)}} \mathbf{u}_{ij}^{(m)} \qquad (7)$$

and

$$\widehat{\mathbf{T}}_{ij}^{(m)} = E_{\mathbf{t}_{ij}|\mathbf{y}_{ij},\boldsymbol{\theta}^{(m)}} \left\{ \mathbf{t}_{ij}\mathbf{t}_{ij}^H \right\} = \mathbf{V}_{ij}^{-1^{(m)}} + \widehat{\mathbf{t}}_{ij}^{(m)}\widehat{\mathbf{t}}_{ij}^{H^{(m)}} \qquad (8)$$

M-step: This step is carried by block coordinate descent, which has the advantage to lead to closed-form expressions of the unknown parameter when maximizing $Q(\boldsymbol{\theta}|\boldsymbol{\theta}^{(m)})$ w.r.t. $\boldsymbol{\theta}$. We recall that the EM extension used here is the ECME algorithm in which the estimates of $\left\{ \boldsymbol{\zeta}_i^T \right\}_i, \left\{ \tau_{ij} \right\}_{ij}, \sigma^2$ are obtained by maximizing $Q(\boldsymbol{\theta}|\boldsymbol{\theta}^{(m)})$ based on the complete log-likelihood, whereas the $\mathbf{m}$ estimate is obtained by maximizing the log-likelihood of the incomplete data (i.e. observations only). After some calculus, this leads to the following updates (for the sake of clarity, the iteration index m is omitted):

$$\sqrt{\tau_{ij}} = \frac{\Re(\mathbf{r}_{ij}^H \mathbf{W}_i \widehat{\mathbf{t}}_{ij})}{\mathrm{Tr}(\widehat{\mathbf{T}}_{ij}\mathbf{W}_i^H \mathbf{W}_i)} \qquad (9)$$

$$\sigma^2 = \frac{1}{\sum_i n_i} \sum_i \sum_j \mathrm{Tr}\left\{ \tau_{ij}\widehat{\mathbf{T}}_{ij}\mathbf{W}_i^H \mathbf{W}_i + \mathbf{r}_{ij}^H \mathbf{r}_{ij} - 2\sqrt{\tau_{ij}}\Re(\mathbf{r}_{ij}^H \mathbf{W}_i \widehat{\mathbf{t}}_{ij}) \right\} \qquad (10)$$

$$\mathbf{W}_i = \sum_j \sqrt{\tau_{ij}} \mathbf{r}_{ij}\widehat{\mathbf{t}}_{ij}^H \left(\sum_j \widehat{\mathbf{T}}_{ij} \right)^{-H} \qquad (11)$$

$$\mathbf{m} = \left(\sum_i \sum_j \mathbf{A}_i^H \boldsymbol{\Gamma}_{ij} \mathbf{A}_i \right)^{-1} \sum_i \sum_j \mathbf{A}_i^H \boldsymbol{\Gamma}_{ij} \mathbf{y}_{ij} \qquad (12)$$

in which $\boldsymbol{\Gamma}_{ij} = \frac{1}{\sigma^2}\left(\mathbf{I} - \mathbf{W}_i \left(\frac{\tau_{ij}}{\sigma^2}\mathbf{I} + \mathbf{W}_i^H \mathbf{W}_i \right)^{-1}\mathbf{W}_i^H \right)$. As a summary, the ECME algorithm operates as described in pseudo-code 1.

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Algorithm 1 ECME algorithm for parameter estimation under mixed-effects model
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1: Initialize mean $\mathbf{m}^{(m=0)}$ with robust multivariate mean estimate (e.g. [11, 12, 13])
   and loading matrix $\mathbf{W}_i^{(m=0)}$ with PPCA algorithm separately for each individual [3].

2: **repeat**

3:     Given current $\boldsymbol{\theta}^{(m)}$, compute $\widehat{\mathbf{t}}_{ij}^{(m)}$ and $\widehat{\mathbf{T}}_{ij}^{(m)}$ using (7) and (8)

4:     Update $\boldsymbol{\theta}^{(m+1)}$ as in (9), (10), (11) followed by (12)

5:     $m \leftarrow m + 1$

6: **until** convergence of $||\boldsymbol{\theta}^{(m+1)} - \boldsymbol{\theta}^{(m)}||_F^2$ is met.

7: **return** $\widehat{\boldsymbol{\theta}}$
---

3. Discussions and extension

3.1. Parallelization

A parallel scheme naturally appears from (9)–(12). Specifically, one can perform parallel computations on individuals indexed by i. This can be used to enhance the computational cost and avoid a central processor overload. Fig. 1 represents the operation flow and signaling exchange between a local individual's processor and the fusion center. The consensus step is enforced for deriving the common mean and homogeneous noise power. Note that the update of $\mathbf{m}$ involves the inversion of a $n_i \times n_i$ matrix in classical robust multivariate covariance and mean estimators [6]. Conversely, the ECME algorithm updates $\mathbf{m}$ locally by the inversion of a $r_i \times r_i$ matrix followed, at the fusion center, by an inversion of a $p \times p$ matrix. In array processing we commonly have $p \ll n_i$ (e.g., less sources than sensors or in radio-interferometry imaging), which makes the proposed method computationally efficient.

3.2. Rotational ambiguity of the loading matrix

It has been noted in [3] that the estimates of the loading matrix are generally not orthogonal. It means that if $\mathbf{O}_i$ denotes an orthogonal rotational matrix of an adequate size, then $\widehat{\mathbf{W}}_{i_{\mathrm{ML}}} \mathbf{O}_i$ remains the maximum likelihood estimate. Yet, it is common to be interested in the spanned subspace by the columns of $\mathbf{W}_i$ rather than the loading matrix itself. On the other hand, if needed, a post-processing of $\widehat{\mathbf{W}}_i$ is proposed in [3], which consists in computing the SVD of $\widehat{\mathbf{W}}_i^H \widehat{\mathbf{W}}_i = \mathbf{O}_i^H \boldsymbol{\Lambda}_i \mathbf{O}_i$ and rotating according to $\mathbf{O}_i \widehat{\mathbf{W}}_i$.

3.3. Extension to the missing data case

The missing data case refers to the situation where part of the individual responses are unobserved, which is a very common issue when analyzing time series, such as remote sensing data [14] or biochemical data [15]. In the following, we denote the observed responses by $\mathbf{y}_{ij}^o$ and the unobserved responses by $\mathbf{y}_{ij}^u$ for the i-th individual at the j-th observation. Furthermore, the scenario is assumed to be missing at random (i.e. the missing observations do not depend on their values) [16]. In order to use of the results obtained in the previous section, we consider the complete data as $\mathbf{x} = \{\mathbf{x}_{ij}\}_{ij}$ in which $\mathbf{x}_{ij} = \left[\mathbf{y}_{ij}^{o^T}, \mathbf{y}_{ij}^{u^T}, \mathbf{t}_{ij}^T\right]^T$. Consequently,

$$Q(\boldsymbol{\theta}|\boldsymbol{\theta}^{(m)}) = E_{\{\mathbf{y}_{ij}^u\}_{ij}|\{\mathbf{y}_{ij}^o\}_{ij},\boldsymbol{\theta}^{(m)}}\left\{E_{\{\mathbf{t}_{ij}\}_{ij}|\{\mathbf{y}_{ij}^o\}_{ij},\{\mathbf{y}_{ij}^u\}_{ij},\boldsymbol{\theta}^{(m)}}\left\{\mathcal{L}_{\mathrm{C}}(\boldsymbol{\theta}|\mathbf{x})\right\}\right\} \tag{13}$$

The inner expectation in (13) is given by (6), the outer expectation can be obtained from the classical results of Anderson [17]. Namely, it is fairly easy to see that

$$\widehat{\mathbf{y}}_{ij} \triangleq E_{\mathbf{y}_{ij}^u|\mathbf{y}_{ij}^o,\boldsymbol{\theta}^{(m)}}\left\{\mathbf{y}_{ij}\right\} = \left[\mathbf{y}_{ij}^{o^T}\widehat{\mathbf{y}_{ij}^u}^T\right]^T = \begin{bmatrix} \mathbf{y}_{ij}^o \\ \mathbf{m}^u + \boldsymbol{\Sigma}_{ij}^{uo}\boldsymbol{\Sigma}_{ij}^{uo^{-1}}\left(\mathbf{y}_{ij}^o - \mathbf{m}^o\right) \end{bmatrix} \tag{14}$$

and

$$\widehat{\mathbf{Y}}_{ij} \triangleq E_{\mathbf{y}_{ij}^u|\mathbf{y}_{ij}^o,\boldsymbol{\theta}^{(m)}}\left\{\mathbf{y}_{ij}\mathbf{y}_{ij}^H\right\} = \begin{bmatrix} \mathbf{y}_{ij}^o\mathbf{y}_{ij}^{o^H} & \mathbf{y}_{ij}^o\widehat{\mathbf{y}_{ij}^u}^H \\ \widehat{\mathbf{y}_{ij}^u}\mathbf{y}_{ij}^{o^H} & \boldsymbol{\Sigma}_{ij}^{uu} - \boldsymbol{\Sigma}_{ij}^{uo}\boldsymbol{\Sigma}_{ij}^{oo^{-1}}\boldsymbol{\Sigma}_{ij}^{ou} + \widehat{\mathbf{y}_{ij}^u}\widehat{\mathbf{y}_{ij}^u}^H \end{bmatrix} \tag{15}$$

in which the mean and the covariance matrix of $\mathbf{y}_{ij}$ are decomposed as $\mathbf{m} = \left[\mathbf{m}^{o^T}\mathbf{m}^{u^T}\right]^T$ and $\boldsymbol{\Sigma}_{ij} = \begin{bmatrix} \boldsymbol{\Sigma}_{ij}^{oo} & \boldsymbol{\Sigma}_{ij}^{ou} \\ \boldsymbol{\Sigma}_{ij}^{uo} & \boldsymbol{\Sigma}_{ij}^{uu} \end{bmatrix}$. Then, we deduce the E-step which consists in updating the following expectations (with regard to $\{\mathbf{x}_{ij}\}_{ij}|\{\mathbf{y}_{ij}^o\}_{ij},\boldsymbol{\theta}^{(m)}$):

$$\widehat{\mathbf{r}}_{ij} \triangleq E\left\{\mathbf{r}_{ij}\right\} = \widehat{\mathbf{y}}_{ij} - \mathbf{m} \tag{16}$$

$$\widehat{\mathbf{R}}_{ij} \triangleq E\left\{\mathbf{r}_{ij}\mathbf{r}_{ij}^H\right\} \triangleq \widehat{\mathbf{Y}}_{ij} + \mathbf{m}\mathbf{m}^H - 2\Re\left\{\widehat{\mathbf{y}}_{ij}\mathbf{m}^H\right\} \tag{17}$$

$$\widehat{\mathbf{t}}_{ij} = E\left\{\mathbf{t}_{ij}\right\} \triangleq \frac{\sqrt{\tau_{ij}}}{\sigma^2}\mathbf{V}_{ij}^{-1}\mathbf{W}_iE\left\{\mathbf{r}_{ij}\right\} \tag{18}$$

$$\widehat{\mathbf{r}_{ij}\mathbf{t}_{ij}^H} \triangleq E\left\{\mathbf{r}_{ij}\mathbf{t}_{ij}^H\right\} = \frac{\sqrt{\tau_{ij}}}{\sigma^2}\widehat{\mathbf{R}}_{ij}\mathbf{W}_i \tag{19}$$

$$\widehat{\mathbf{T}}_{ij} \triangleq E\left\{\mathbf{t}_{ij}\mathbf{t}_{ij}^H\right\} = \mathbf{V}_{ij}^{-1} + \frac{\tau_{ij}}{\sigma^4}\mathbf{V}_{ij}^{-1}\mathbf{W}_i^H\mathbf{R}_{ij}\mathbf{W}_i\mathbf{V}_{ij}^{-1} \tag{20}$$

Finally, by plugging (16)–(20) into (13), the M-step of the ECME reads

$$\sqrt{\tau_{ij}} = \frac{\Re\left(\mathrm{Tr}\left(\widehat{\mathbf{t}_{ij}\mathbf{r}_{ij}^H}\mathbf{W}_i\right)\right)}{\mathrm{Tr}(\widehat{\mathbf{T}}_{ij}\mathbf{W}_i^H\mathbf{W}_i)}, \sigma^2 = \frac{1}{\sum_i n_i}\sum_i\sum_j \mathrm{Tr}\left\{\tau_{ij}\widehat{\mathbf{T}}_{ij}\mathbf{W}_i^H\mathbf{W}_i+\widehat{\mathbf{R}}_{ij}-2\sqrt{\tau_{ij}}\Re\left(\mathrm{Tr}\left(\widehat{\mathbf{t}_{ij}\mathbf{r}_{ij}^H}\mathbf{W}_i\right)\right)\right\},$$

$\mathbf{W}_i = \sum_j \sqrt{\tau_{ij}}\widehat{\mathbf{r}_{ij}\mathbf{t}_{ij}^H}\left(\sum_j \widehat{\mathbf{T}}_{ij}\right)^{-H}$, and $\mathbf{m} = \left(\sum_i\sum_j \mathbf{A}_i^H\mathbf{\Gamma}_{ij}\mathbf{A}_i\right)^{-1}\sum_i\sum_j \mathbf{A}_i^H\mathbf{\Gamma}_{ij}\widehat{\mathbf{y}}_{ij}$.
As in the fully observed case, the missing data case can be cast into a parallel scheme (see Fig. 1).

4. Numerical simulations

In this section, we aim at evaluating numerically the performance of the proposed algorithm. Specifically, we consider three different scenarios with rank $r_i = 3$, $\mathbf{A}_i = \mathbf{I}$ and $\mathbf{m} \in \mathbb{C}^{10\times 1}$ following a Gaussian distribution. For the sake of simplicity, $[\mathbf{W}_i]_{1:r_i;1:r_i} = \mathbf{I} + \mathbf{K}$ with $[\mathbf{K}]_{h,q} \sim \mathcal{CN}(0,1)$ and $[\mathbf{W}]_{r_i+1:n;1:r_i} = \mathbf{0}$ such that we still have $\mathrm{rank}(\mathbf{W}_i) = r$. In the first scenario, we focus on the mean estimation, $\mathbf{m}$, by comparing the ECME proposed algorithm with the minimum covariance determinant [12], the Marona's orthogonalized Gnanadesikan-Kettenring (OGK) [13], the Olive Hawkins estimators [11], the classical probabilistic PCA [3] and the robust student-based probabilistic PCA [18]. Fig. 2 and Fig. 4 show the mean square error (MSE) of $\mathbf{m}$ estimates versus the number of observations in which we consider a 0 dB signal-to-noise ratio. Only one individual with $m \geq 2n$ is considered in order to respect restrictions given by the aforementioned algorithms [3, 11, 12, 13, 18]. In Fig. 2, the texture parameter is a realization of an inverse-Gamma distribution (leading to a t-distributed random effect component). Three strong outliers with a power 10^3 times higher than the mean amplitude are added, which represents 3% to 15% of the data. We note that the proposed ECME and the robust student-based PPCA exhibit comparable performances. Hence, the former algorithm fits well with the latter which assumes a perfect knowledge of the texture distribution. In addition, we notice from Fig. 3 that the estimation accuracy of the proposed estimator does not strongly depend on the parameter of the Student's t-distribution. It is worth mentioning that other location and scatter estimators as OGK, Olive Hawkins and the FMCD are also known to be robust to the variation of the Student's parameters. In Fig. 4, in which the texture parameter is taken as a realization of a uniform distribution over $]0, 500]$, the proposed ECME outperforms the compared algorithms since it does not

assume any distribution of texture while taking into account the low-rank structure of the within-individual error.

In the second scenario, we focus on the normalized MSE of $\mathbf{W}^H\mathbf{W}$ versus the number of observations. Again, we notice from Fig. 5 that the proposed algorithm surpasses PPCA and the robust Student's-based PPCA regarding subspace inference.

Finally, in the third scenario, we consider a multi-individual case and we show the benefit of considering the extension of the proposed algorithm to the scenario of the presence of missing data. Specifically, we plot in Fig. 6 the MSE of the mean estimate in the case of three individuals versus the percentage of missing data (i.e. missing values at individuals i and observations j). We notice that the proposed ECME extension shows a good performance since it remains close to the "clairvoyant" ECME which uses the full original data (i.e. observed and unobserved), whereas the ECME using only the observed data in all individuals exhibits a notable performance loss when the missing data ratio exceeds 10% of the total data.

5. Conclusion

In this paper we design an ECME-based algorithm for mean and covariance estimation under heterogeneous mixed-effects model. The random effects were modeled as a heteroscedastic Gaussian process allowing flexibility of the within-individual error and robustness against any possible compound Gaussian distribution. In addition, the low-rank structure of the loading matrix of each individual was taken into account, leading to a natural parallel scheme enhancing computational cost. Finally, an extension to the missing data scenario was given and numerical simulations assessing the usefulness of the proposed scheme have been presented, showing a performance gain with existing state-of-the-art estimators.

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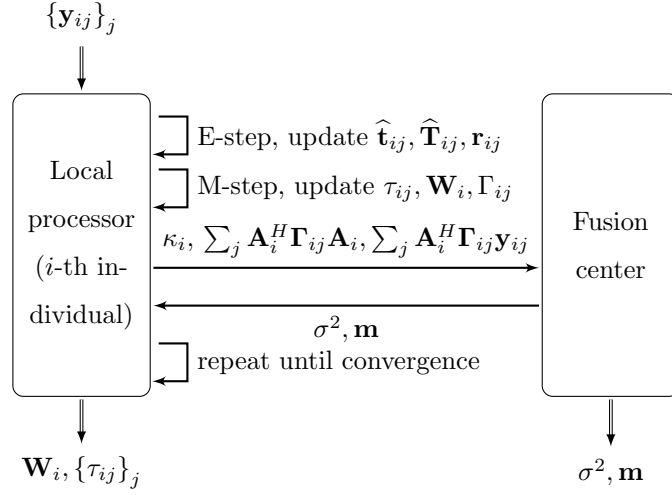
Figure 1: Operation flow and signaling exchange between a local individual's processor and the central processor, in which $\kappa_i = \sum_j \operatorname{Tr}\{\tau_{ij}\widehat{\mathbf{T}}_{ij}\mathbf{W}_i^H\mathbf{W}_i + \mathbf{r}_{ij}^H\mathbf{r}_{ij} - 2\sqrt{\tau_{ij}}\Re(\mathbf{r}_{ij}^H\mathbf{W}_i\widehat{\mathbf{t}}_{ij})$. The consensus step is enforced for deriving the common mean and homogeneous noise power.
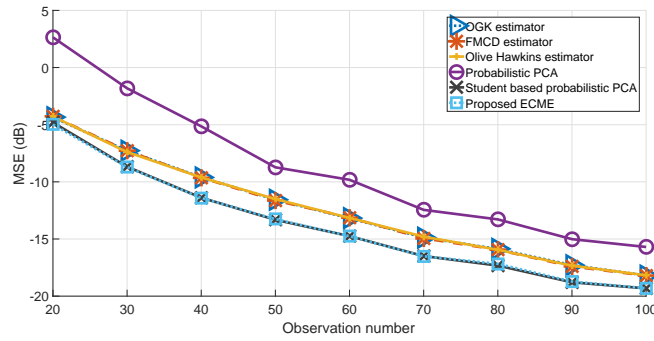

Figure 2: Mean square error vs. number of observations under t-distributed data (1000 runs mean).

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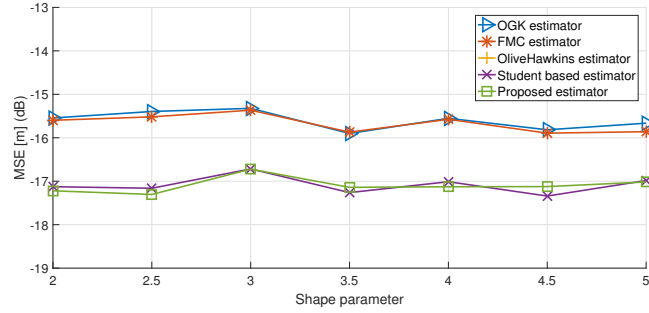
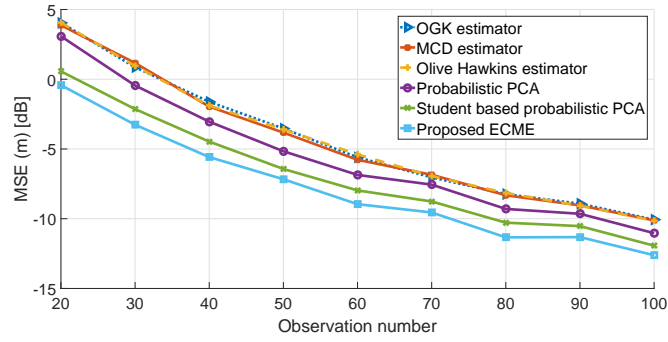Figure 3: Mean square error of the mean vs. shape parameter.

Figure 4: Mean square error vs. number of observations under a uniform distributed within-individual error (1000 runs mean).
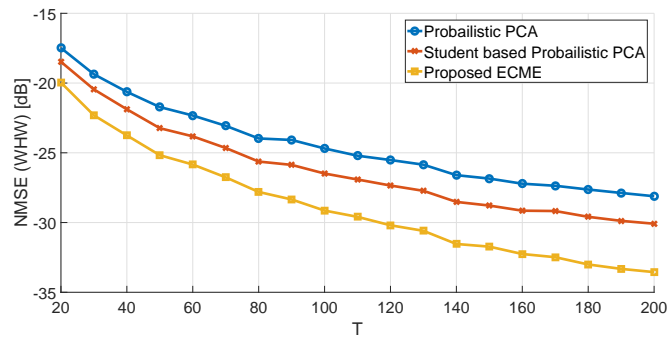

Figure 5: Normalized mean square error vs. number of observations under a uniform distributed within-individual error (1000 runs mean).
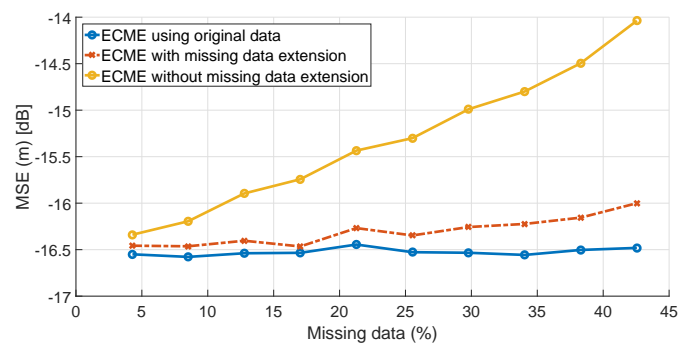
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Figure 6: Mean square error vs. percentage of unobserved data (1000 runs mean).