

Robust Unmixing of Multitemporal Hyperspectral Images

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Abstract—Hyperspectral unmixing is a blind source separation problem which consists in estimating the reference spectral signatures contained in a hyperspectral image, as well as their relative contribution to each pixel according to a given mixture model. In practice, the process is further complexified by the inherent spectral variability of the observed scene and the possible presence of outliers. More specifically, multi-temporal hyperspectral images, i.e., sequences of hyperspectral images acquired over the same area at different time instants, are likely to simultaneously exhibit moderate endmember variability and abrupt spectral changes either due to outliers or to significant time intervals between consecutive acquisitions. Unless properly accounted for, these two perturbations can significantly affect the unmixing process. In this context, we propose a new unmixing model for multitemporal hyperspectral images accounting for smooth temporal variations, construed as spectral variability, and abrupt spectral changes interpreted as outliers. The proposed hierarchical Bayesian model is inferred using a Markov chain Monte-Carlo (MCMC) method allowing the posterior of interest to be sampled and Bayesian estimators to be approximated. A comparison with unmixing techniques from the literature on synthetic and real data allows the interest of the proposed approach to be appreciated.

Index Terms—Hyperspectral imagery, multitemporal images, endmember variability, Markov chain Monte-Carlo (MCMC) methods.

I. INTRODUCTION

AQUIRED in hundreds of contiguous spectral bands, hyperspectral (HS) images have received an increasing interest due to the significant spectral information they convey, which is somewhat mitigated by their lower spatial resolution in remote sensing applications. This limitation, combined with possibly complex interactions between the incident light and the observed materials, implies that the observed spectra are mixtures of several signatures corresponding to distinct materials. Spectral unmixing then consists in identifying a limited number of reference spectral signatures composing the data – referred to as *endmembers* – and their abundance fractions in each pixel according to a predefined mixture model. The choice of a specific model generally reflects the practitioners’ prior knowledge on the environmental factors

possibly affecting the acquisitions, such as declivity or multiple reflections. Traditionally, a linear mixing model (LMM) is adopted since it is appropriate to describe hyperspectral data when the declivity of the scene and microscopic interactions between the observed materials are negligible [1]. Depending on the applications, various models have also been investigated to capture higher interactions (i.e., nonlinearities) between the incident light and the observed materials (see [2], [3] for recent reviews on this topic). However, varying acquisition conditions, such as local illumination variations or the natural evolution of the scene, may significantly alter the shape and the amplitude of the acquired signatures [4], [5], thus affecting the extracted endmembers. While endmember variability has hitherto been extensively considered within a single HS image, either in a deterministic [6]–[9] or a statistical setting [10]–[12], fewer works consider temporal variability by exploiting the possibilities offered by multitemporal HS (MTHS) images.

From a hyperspectral unmixing perspective, MTHS images, i.e., sequences of HS images acquired over the same area at different time instants, can be of interest to exploit information redundancy between consecutive images (e.g., through features exhibiting moderate or smooth temporal variations as in [13], [14]) while allowing the endmember temporal evolution to be characterized. For instance, MTHS have been recently exploited to improve endmember unmixing results [15]–[17] and used in a change detection problem involving two HS images [18], [19].

Even though the approaches proposed in [15]–[17] specifically allow smooth temporal variations of some of the mixture parameters to be considered, they do not account for abrupt spectral changes either due to outliers or to possibly significant time intervals between two consecutive images. In practice, such situations can be reasonably expected, depending on the acquisition dates and possible climatic hazards, e.g., when vegetation or water is present in the observed scene. Unless specifically accounted for, this situation frequently observed in real datasets has a significant impact on the recovered endmembers, which motivates the present work. Inspired by [16], [20], [21], and based on an original interpretation of the unmixing problem under study, our contribution consists in jointly accounting for smooth endmember variations – construed as temporal endmember variability – and abrupt changes interpreted as outliers (e.g., significant variability within a single image or presence of non-linearities) using a carefully designed hierarchical Bayesian model. More precisely, we focus our analysis on scenes in which mostly the same materials are expected to be observed from an image to

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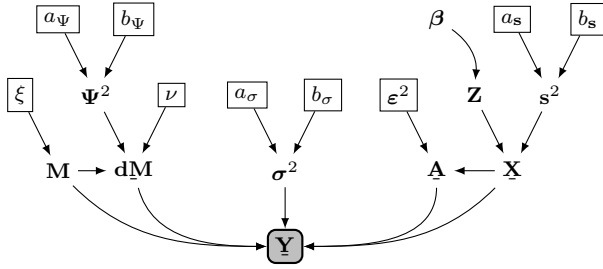


Fig. 1. Directed acyclic graph associated with the proposed Bayesian model (fixed parameters appear in boxes).

another. In this context, using the endmembers extracted from the reference scene as a starting point to unmix the whole MTHS image constitutes a reasonable attempt to generalize the analyses previously conducted for a single image. On the one hand, the endmembers identified in each single image can *in fine* be considered as time-varying instances of reference signatures shared by the different images, thus justifying the use of a modified version of the perturbed linear mixing model (PLMM) proposed in [17]. This formulation will notably allow smooth spectral variations occurring over time to be captured. On the other hand, the signatures corresponding to materials appearing in only a few images, which induce abrupt spectral changes, can be regarded as outliers with respect to the commonly shared endmembers. This paper studies a new Bayesian model allowing both spectral variability and presence of outliers to be considered in the unmixing of MTHS images. The resulting unmixing task is solved using a Markov chain Monte-Carlo (MCMC) allowing the posterior of interest to be sampled and Bayesian estimators to be approximated.

The paper is organized as follows. The mixing model considered in this paper is introduced in Section II, and the associated hierarchical Bayesian model is developed in Section III. Section IV investigates a Gibbs sampler to solve the resulting mixed integer non-linear problem. The performance of the proposed approach on synthetic and real data is studied in Sections V and VI. In particular, the results obtained with the proposed method are compared to those of the VCA/FCLS algorithm [22], [23], the SISAL/FCLS algorithm [24], the algorithm associated with the robust LMM (RLMM) proposed in [25] and the MTHS optimization method [17]. Finally, Section VII concludes this work and outlines further research perspectives.

II. PROBLEM STATEMENT

We consider a sequence of HS images acquired at T different time instants over the same area, where mostly the same materials are expected to be observed over time. In the following, at most R endmembers are assumed to be shared between the T images composing the sequence, where R is *a priori* known. Since the observed instances of a given endmember can be reasonably expected to vary from an image to another, we propose to account for smooth endmember spectral variations via a modified version of the perturbed linear mixing model (PLMM) proposed in [9], [17]. Inspired by the total least squares problem [26], the PLMM consists in representing each pixel y_{nt} by a linear combination of the R

endmembers – denoted by \mathbf{m}_r – affected by an additive error term $\mathbf{d}\mathbf{m}_{r,t}$ accounting for temporal endmember variability. However, this model shows notable limitations when the vector $\mathbf{y}_{n,t}$ is affected by abrupt changes. Consequently, this paper investigates a new unmixing model jointly accounting for endmember variability and abrupt changes possibly affecting MTHS images. To this end, the proposed model is a generalized PLMM, which includes an additional term $\mathbf{x}_{n,t}$ to capture significant deviations from the LMM, i.e., significant spatial variability or non-linearities within each image [20], [25]. The resulting observation model can thus be written

$$\mathbf{y}_{n,t} = \sum_{r=1}^R a_{r,n,t} (\mathbf{m}_r + \mathbf{d}\mathbf{m}_{r,t}) + \mathbf{x}_{n,t} + \mathbf{b}_{n,t} \quad (1)$$

for $n = 1, \dots, N$ and $t = 1, \dots, T$, where $\mathbf{y}_{n,t}$ denotes the n th image pixel at time t , \mathbf{m}_r is the r th endmember, $a_{r,n,t}$ is the proportion of the r th endmember in the n th pixel at time t , $\mathbf{d}\mathbf{m}_{r,t}$ denotes the perturbation of the r th endmember at time t , and $\mathbf{x}_{n,t}$ denotes the contribution of outliers in the n th pixel at time t . Finally, $\mathbf{b}_{n,t}$ represents an additive noise resulting from the data acquisition and the modeling errors. The so-called robust PLMM can be written

$$\mathbf{Y}_t = (\mathbf{M} + \mathbf{d}\mathbf{M}_t) \mathbf{A}_t + \mathbf{X}_t + \mathbf{B}_t \quad (2)$$

where $\mathbf{Y}_t = [\mathbf{y}_{1,t}, \dots, \mathbf{y}_{N,t}]$ is an $L \times N$ matrix containing the pixels of the t th image, \mathbf{M} denotes an $L \times R$ matrix containing the endmembers that are common to all the images of the sequence, \mathbf{A}_t is an $R \times N$ matrix composed of the abundance vectors $\mathbf{a}_{n,t}$, $\mathbf{d}\mathbf{M}_t$ is an $L \times R$ matrix whose columns contain the variability inherent to the t th image, \mathbf{X}_t is an $L \times N$ matrix whose columns are the outliers present in the image t , and \mathbf{B}_t is an $L \times N$ matrix accounting for the noise at time t . The constraints considered to reflect physical considerations are

$$\begin{aligned} \mathbf{A}_t &\succeq \mathbf{0}_{R,N}, \quad \mathbf{A}_t^T \mathbf{1}_R = \mathbf{1}_N, \quad \forall t \in \{1, \dots, T\} \\ \mathbf{M} &\succeq \mathbf{0}_{L,R}, \quad \mathbf{M} + \mathbf{d}\mathbf{M}_t \succeq \mathbf{0}_{L,R}, \quad \forall t \in \{1, \dots, T\} \\ \mathbf{X}_t &\succeq \mathbf{0}_{L,N}, \quad \forall t \in \{1, \dots, T\} \end{aligned} \quad (3)$$

where \succeq denotes a term-wise inequality. Note that the outlier term \mathbf{X}_t is intended to describe abrupt changes due to the apparition of a new endmember that was not present in the reference image. This justifies the corresponding non-negativity constraint, similar to the one imposed on the other endmembers. Given the mixture model (2), the unmixing problem considered in this work consists in inferring the abundances \mathbf{A}_t , the endmembers \mathbf{M} , the variability $\mathbf{d}\mathbf{M}_t$ and the outliers \mathbf{X}_t from the observations \mathbf{Y}_t , $t = 1, \dots, T$. In the next section, this problem is tackled in a Bayesian framework to easily incorporate all the prior knowledge available on the mixture parameters.

III. BAYESIAN MODEL

This section details the specific structure imposed on the parameters to be inferred *via* appropriate prior distributions. Note that dependencies with respect to constant parameters are omitted in the following paragraphs to simplify the notations.

A. Likelihood

Assuming the additive noise $\mathbf{b}_{n,t}$ is distributed according to a Gaussian distribution $\mathbf{b}_{n,t} \sim \mathcal{N}(\mathbf{0}_L, \sigma_t^2 \mathbf{I}_L)$, the observation model (2) leads to

$$\mathbf{y}_{n,t} | \mathbf{M}, \mathbf{dM}_t, \mathbf{A}_t, \mathbf{X}_t, \sigma_t^2 \sim \mathcal{N}\left((\mathbf{M} + \mathbf{dM}_t)\mathbf{a}_{n,t} + \mathbf{x}_{n,t}, \sigma_t^2 \mathbf{I}_L\right). \quad (4)$$

In addition, assuming prior independence between the pixels within each image and between the images \mathbf{Y}_t themselves, the likelihood function of all images $\mathbf{Y} = [\mathbf{Y}_1, \dots, \mathbf{Y}_T]$ is

$$p(\mathbf{Y} | \Theta) \propto \prod_{t=1}^T (\sigma_t^2)^{-NL/2} \times \exp\left(-\frac{1}{2\sigma_t^2} \|\mathbf{Y}_t - (\mathbf{M} + \mathbf{dM}_t)\mathbf{A}_t - \mathbf{X}_t\|_F^2\right) \quad (5)$$

where $\|\cdot\|_F$ is the Frobenius norm, $\Theta = \{\Theta_p, \Theta_h\}$ and

$$\Theta_p = \{\mathbf{M}, \mathbf{dM}, \mathbf{A}, \mathbf{X}, \sigma^2, \mathbf{Z}\}, \quad \Theta_h = \{\Psi^2, \mathbf{s}^2, \beta\} \quad (6)$$

denote the parameters and hyperparameters whose priors are defined in the following paragraphs. Note that the independence assumption between the observed images conditionally on the unknown parameters is justified by the fact that the sequence of images has been acquired by possibly different sensors at different time instants.

B. Parameter priors

1) *Abundances*: We propose to promote smooth temporal variations of the abundances between successive time instants for pixels that are not classified as outliers. To this end, we introduce a new prior defined for $n = 1, \dots, N$ as

$$\mathbf{a}_{n,1} | \mathbf{x}_{n,t} = \mathbf{0}_L \sim \mathcal{U}_{\mathcal{S}_R} \quad (7)$$

$$\mathbf{a}_{n,t} | \mathbf{x}_{n,t} \neq \mathbf{0}_L \sim \mathcal{U}_{\widetilde{\mathcal{S}}_R}, \text{ for } t = 1, \dots, T \quad (8)$$

$$p\left(\mathbf{a}_{n,t} | \mathbf{x}_{n,t} = \mathbf{0}_L, \mathbf{A}_{\setminus\{\mathbf{a}_{n,t}\}}\right) \propto \exp\left\{-\frac{1}{2\varepsilon_n^2} \left(\|\mathcal{I}_{n,t}^1 \neq \emptyset\| \|\mathbf{a}_{n,t} - \mathbf{a}_{n,\tau_{n,t}^1}\|_2^2\right)\right\} \mathbf{1}_{\mathcal{S}_R}(\mathbf{a}_{n,t}), \text{ for } t \geq 2 \quad (9)$$

where $\mathcal{U}_{\mathcal{S}_R}$ denote the uniform distribution on the set \mathcal{S}_R , $\mathbf{1}_{\mathcal{S}_R}$ is the indicator function of the set \mathcal{S}_R , $[\mathcal{P}]$ denotes the Inversion bracket applied to the logical proposition \mathcal{P} , i.e.,

$$[\mathcal{P}] = \begin{cases} 1, & \text{if } \mathcal{P} \text{ is true;} \\ 0, & \text{otherwise} \end{cases}$$

and

$$\mathcal{S}_R = \{\mathbf{x} \in \mathbb{R}^R | \forall i, x_i \geq 0 \text{ and } \mathbf{x}^T \mathbf{1}_R = 1\} \quad (10)$$

$$\widetilde{\mathcal{S}}_R = \{\mathbf{x} \in \mathbb{R}^R | \forall i, x_i \geq 0 \text{ and } \mathbf{x}^T \mathbf{1}_R \leq 1\} \quad (11)$$

$$\mathcal{I}_{n,t}^1 = \{\tau < t | z_{n,\tau} = 0\}, \quad \tau_{n,t}^1 = \max_{\tau \in \mathcal{I}_{n,t}^1} \tau. \quad (12)$$

By convention, we set $\mathcal{I}_{n,t}^1 = \emptyset$ when $t = 1$. To be more explicit, consider an image at time t and a pixel n within this image which is not corrupted by outliers (i.e., $\mathbf{x}_{n,t} = \mathbf{0}_L$). For $t = 1$, a uniform distribution defined in the unit simplex is selected to reflect the absence of specific prior knowledge while accounting for the related constraints in (3). For $t > 1$, smooth variations of $\mathbf{a}_{n,t}$ are promoted via a one-dimensional Gaussian Markov field [12], [27] penalizing

the Euclidean distance between $\mathbf{a}_{n,t}$ and the abundance of the last corresponding outlier-free pixel in the preceding images of the sequence, i.e., at time instant $\tau_{n,t}^1$. On the contrary, when outliers are present in the pixel (n, t) ($\mathbf{x}_{n,t} \neq \mathbf{0}_L$), the usual abundance sum-to-one constraint is relaxed ($\mathbf{a}_{n,t}^T \mathbf{1}_R \leq 1$) so that the prior allows cases in which the linear model does not exhaustively describe the data to be addressed. In the following, the joint abundance prior is denoted by

$$p(\mathbf{A} | \mathbf{X}) = \prod_{n=1}^N \left[\prod_{\substack{j=1 \\ t_j: \mathbf{x}_{n,t_j} \neq \mathbf{0}_L}}^{J_n} p(\mathbf{a}_{n,t_j} | \mathbf{x}_{n,t_j} \neq \mathbf{0}_L) \right] \times \left[\prod_{\substack{i=1 \\ t_i: \mathbf{x}_{n,t_i} = \mathbf{0}_L}}^{I_n} p(\mathbf{a}_{n,t_i} | \mathbf{a}_{n,t_{i-1}}, \mathbf{x}_{n,t_i} = \mathbf{0}_L) \right] \quad (13)$$

with $I_n = \sharp\{t : \mathbf{x}_{n,t} = \mathbf{0}_L\}$, $J_n = T - I_n$ and \sharp denotes the cardinal operator.

2) *Outliers \mathbf{X} and label maps \mathbf{Z}* : Similarly to [20], outliers are a priori assumed to be spatially sparse. Different approaches have been proposed in the literature to include this prior knowledge, either relying on the ℓ_1 penalty (such as the LASSO [28]) or on mixtures of probability distributions involving a Dirac mass at zero and a continuous probability distribution [29] (such as the Bernoulli-Laplace [30] or Bernoulli-Gaussian distributions [31], [32], extensively used in the literature [33]–[35]). In this work, we propose to assign the following prior to the outliers $\mathbf{x}_{n,t}$ to promote spatial sparsity

$$p(\mathbf{x}_{n,t} | z_{n,t}, s_t^2) = (1 - z_{n,t})\delta(\mathbf{x}_{n,t}) + z_{n,t}\mathcal{N}_{\mathbb{R}_+^L}(\mathbf{0}_L, s_t^2) \quad (14)$$

where $\mathcal{N}_{\mathbb{R}_+^L}$ denotes a Gaussian distribution truncated to the set \mathbb{R}_+^L . Note that $z_{n,t} = 1$ if an outlier is present in the corresponding pixel, and 0 otherwise. The proposed prior notably allows outliers to be *a priori* described by a truncated Gaussian distribution when $z_{n,t} = 1$, since the outliers $\mathbf{x}_{n,t}$ are mainly due to the apparition of new endmembers (i.e., that were not present in the reference image). With this context in mind, we further propose to promote spatial correlations between the outliers' support, since new materials are likely to appear in multiple contiguous pixels. The binary label maps $\mathbf{z}_t \in \mathbb{R}^N$ ($t = 1, \dots, T$) are consequently modeled as Ising-Markov random fields [12], [36], [37], for which the Hammersley-Clifford theorem yields

$$p(\mathbf{z}_t | \beta_t) = \frac{1}{C(\beta_t)} \exp\left(\beta_t \sum_{n=1}^N \sum_{k \in \mathcal{V}(n)} \delta(z_{n,t} - z_{k,t})\right) \quad (15)$$

where $\mathcal{V}(n)$ denotes the 4-neighbourhood of the pixel n , and $C(\beta_t)$ is the partition function [38]. With appropriate independence assumptions for the label maps \mathbf{z}_t and the outliers $\mathbf{x}_{n,t}$, we obtain

$$p(\mathbf{X} | \mathbf{Z}, \mathbf{s}^2) = \prod_{n,t} p(\mathbf{x}_{n,t} | z_{n,t}, s_t^2) \quad (16)$$

$$p(\mathbf{Z} | \beta) = \prod_{t=1}^T p(\mathbf{z}_t | \beta_t) \quad (17)$$

with $\mathbf{Z} \in \mathbb{R}^{N \times T}$ and $\beta \in \mathbb{R}^T$. Note that the prior (14) leads to the following result, which will be useful to sample the label

maps in Section IV-E

$$p(\mathbf{x}_{n,t} | \mathbf{z}_{\setminus n,t}, s_t^2, \beta_t) = (1 - \omega_{n,t})\delta(\mathbf{x}_{n,t}) + \omega_{n,t} \mathcal{N}_{\mathbb{R}_+^L}(\mathbf{0}_L, s_t^2 \mathbf{I}_L)$$

where $\mathbf{z}_{\setminus n,t}$ denotes the label map \mathbf{z}_t whose n th entry has been removed, and

$$\omega_{n,t} = \frac{1}{\#\mathcal{V}(n)} \exp\left(\beta_t \sum_{n=1}^N \sum_{k \in \mathcal{V}(n)} \delta(1 - z_{k,t})\right). \quad (18)$$

3) *Endmembers*: A non-informative prior is adopted for the endmember matrix \mathbf{M} to reflect the absence of specific prior knowledge about the spectral signatures contained in the image. More precisely, as in previous studies related to hyperspectral unmixing [12], [20], we consider the following truncated multivariate Gaussian distribution

$$\mathbf{m}_r \sim \mathcal{N}_{\mathbb{R}_+^L}(\mathbf{0}_L, \xi \mathbf{I}_L), \text{ for } r = 1, \dots, R \quad (19)$$

where ξ is set to a sufficiently large value to ensure an uninformative prior (e.g., $\xi = 1$). Assuming the endmembers \mathbf{m}_r are independent, the joint prior for the endmembers can be written as

$$p(\mathbf{M}) = \prod_{r=1}^R p(\mathbf{m}_r). \quad (20)$$

In addition, the endmembers can be *a priori* assumed to live in a subspace of dimension $K = R - 1$ [39] whose practical determination can be performed by a principal component analysis (PCA) or a robust PCA (rPCA) [40]. The PCA applied to the original data \mathbf{Y} leads to a decomposition which can be expressed as [39]

$$\mathbf{m}_r = \mathbf{U} \mathbf{e}_r + \check{\mathbf{y}}, \quad \check{\mathbf{y}} = (\mathbf{I}_L - \mathbf{U} \mathbf{U}^T) \bar{\mathbf{y}}, \quad \mathbf{U}^T \mathbf{U} = \mathbf{I}_K \quad (21)$$

where \mathbf{U} denotes a basis of the subspace of dimension K and $\bar{\mathbf{y}}$ denotes the average spectral signature obtained from \mathbf{Y} . Note that using an rPCA would result in similar expressions (modulo a simple change of notations). The projected endmembers projected \mathbf{e}_r are then assigned the following truncated Gaussian prior, which ensures the non-negativity of the endmembers

$$\mathbf{e}_r \sim \mathcal{N}_{\mathcal{E}_r}(\mathbf{0}_K, \xi \mathbf{I}_K), \text{ for } r = 1, \dots, R \quad (22)$$

with

$$\mathcal{E}_r = [e_{1,r}^-, e_{1,r}^+] \times \dots \times [e_{K,r}^-, e_{K,r}^+] \quad (23)$$

$$e_{k,r}^- = \max_{\ell \in \mathcal{U}_k^+} \left(-\frac{\check{y}_\ell + \sum_{j \neq k} u_{\ell,j} e_{j,r}^+}{u_{\ell,k}} \right) \quad (24)$$

$$e_{k,r}^+ = \min_{\ell \in \mathcal{U}_k^-} \left(-\frac{\check{y}_\ell + \sum_{j \neq k} u_{\ell,j} e_{j,r}^-}{u_{\ell,k}} \right) \quad (25)$$

$$\mathcal{U}_k^- = \{r : u_{k,r} < 0\}, \quad \mathcal{U}_k^+ = \{r : u_{k,r} > 0\}. \quad (26)$$

4) *Endmember variability*: We consider a prior for the vectors $\mathbf{d}\mathbf{m}_{r,t}$ (associated with the endmember variability) promoting smooth temporal variations while accounting for the term-wise non-negativity of the observed endmembers (i.e., $\mathbf{m}_r + \mathbf{d}\mathbf{m}_{r,t} \succeq \mathbf{0}_{L,R}$), expressed as

$$\mathbf{d}\mathbf{m}_{r,1} | \mathbf{m}_r \sim \mathcal{N}_{\mathcal{I}_r}(\mathbf{0}_L, \nu \mathbf{I}_L)$$

$$d\mathbf{m}_{\ell,r,t} | m_{\ell,r}, d\mathbf{m}_{\ell,r,(t-1)}, \psi_{\ell,r}^2 \sim \mathcal{N}_{\mathcal{I}_{\ell,r}}(d\mathbf{m}_{\ell,r,(t-1)}, \psi_{\ell,r}^2)$$

for $\ell = 1, \dots, L$, $r = 1, \dots, R$, $t = 1, \dots, T$, where $\mathcal{I}_r = \mathcal{I}_{1,r} \times \dots \times \mathcal{I}_{L,r}$ and $\mathcal{I}_{\ell,r} = [-m_{\ell,r}, +\infty)$. Assuming independence between the different endmember variabilities, the joint variability prior can finally be expressed as

$$p(\mathbf{d}\mathbf{M} | \mathbf{M}, \Psi^2) = \prod_{r=1}^R \left[p(\mathbf{d}\mathbf{m}_{r,1} | \mathbf{m}_r) \times \prod_{t=2}^T p(\mathbf{d}\mathbf{m}_{r,t} | \mathbf{m}_r, \mathbf{d}\mathbf{m}_{r,(t-1)}, \psi_r^2) \right]. \quad (27)$$

5) *Noise variance*: A non-informative inverse-gamma conjugate prior is selected for the noise variance

$$\sigma_t^2 \sim \mathcal{IG}(a_\sigma, b_\sigma) \quad (28)$$

for $t = 1, \dots, T$, with $a_\sigma = b_\sigma = 10^{-3}$ in order to ensure a weakly informative prior. Assuming the noise variances are *a priori* independent finally leads to

$$p(\sigma^2) = \prod_t p(\sigma_t^2). \quad (29)$$

C. Hyperparameters

In order to complete the description of the proposed hierarchical Bayesian model, we consider the following generic priors for the different hyperparameters.

- (i) Non-informative conjugate inverse-gamma priors for the variability variances Ψ^2 and the outlier variances \mathbf{s}^2 , i.e., for $\ell = 1, \dots, L$, $r = 1, \dots, R$ and $t = 1, \dots, T$

$$\psi_{\ell,r}^2 \sim \mathcal{IG}(a_\Psi, b_\Psi), \quad s_t^2 \sim \mathcal{IG}(a_s, b_s) \quad (30)$$

where $\mathcal{IG}(a_\Psi, b_\Psi)$ denotes the inverse gamma distribution and $a_\Psi = b_\Psi = a_s = b_s = 10^{-3}$. Classical independence assumptions for the different hyperparameters lead to

$$p(\Psi^2) = \prod_{\ell,r} p(\psi_{\ell,r}^2), \quad p(\mathbf{s}^2) = \prod_t p(s_t^2). \quad (31)$$

- (ii) A uniform prior for the granularity parameter of a Potts-Markov random field (*a fortiori* of an Ising-Markov random field). Previous studies have shown that it is reasonable to constrain the granularity parameter to belong to the interval $[0, 2]$ [41], leading to

$$\beta_t \sim \mathcal{U}_{[0,2]}, \text{ for } t = 1, \dots, T. \quad (32)$$

Assuming the granularity parameters are *a priori* independent for different time instants finally yields

$$p(\beta) = \prod_t p(\beta_t). \quad (33)$$

D. Joint posterior distribution

Applying Bayes' theorem, the joint posterior distribution of the parameters of interest is given by

$$p(\Theta | \mathbf{Y}) \propto p(\mathbf{Y} | \Theta) p(\mathbf{A} | \mathbf{X}) p(\mathbf{X} | \mathbf{Z}, \mathbf{s}^2) p(\mathbf{s}^2) \times p(\mathbf{Z} | \beta) p(\beta) p(\mathbf{d}\mathbf{M} | \mathbf{M}, \Psi^2) p(\mathbf{M}) p(\Psi^2) p(\sigma^2). \quad (34)$$

The complexity of the proposed Bayesian model summarized in the directed acyclic graph of Fig. 1 and its resulting posterior (34) prevent a simple computation of the maximum *a*

posteriori (MAP) or minimum mean square (MSE) estimators. For instance, the optimization problem associated with the determination of the MAP estimator of Θ is clearly complex, since the negative log-posterior is non-convex and parameterized by mixed continuous and discrete variables. In this context, classical matrix factorization techniques such as [25] cannot be used efficiently. An MCMC method is consequently adopted to sample the posterior (34) and to build estimators of the parameters involved in the proposed Bayesian model using the generated samples.

IV. GIBBS SAMPLER

This section studies a Gibbs sampler, which is guaranteed to produce samples asymptotically distributed according to the target distribution (34). This sampler described in Algo. 1 consists in generating samples distributed according to the conditional distribution of each parameter of interest. Section IV-A introduces the proposed sampling method, and the conditional distributions of all the parameters of interest (see Fig. 1) are detailed in the following paragraphs.

A. Bayesian inference and parameter estimation

The main steps of the proposed Gibbs sampler are summarized in Algo. 1. Similarly to [20], the sequence $\{\Theta^{(q)}\}_{q=N_{\text{bi}}+1}^{N_{\text{MC}}}$ generated by the proposed sampler (i.e., after N_{bi} burn-in iterations) is used to approximate the MMSE estimators of the different unknown parameters \mathbf{M} , \mathbf{A}_t , \mathbf{dM}_t and \mathbf{X}_t by replacing the expectations by empirical averages.

$$\widehat{\mathbf{M}}^{\text{MMSE}} \simeq \frac{1}{N_{\text{MC}} - N_{\text{bi}}} \sum_{q=N_{\text{bi}}+1}^{N_{\text{MC}}} \mathbf{M}^{(q)} \quad (35)$$

$$\widehat{\mathbf{A}}_t^{\text{MMSE}} \simeq \frac{1}{N_{\text{MC}} - N_{\text{bi}}} \sum_{q=N_{\text{bi}}+1}^{N_{\text{MC}}} \mathbf{A}_t^{(q)} \quad (36)$$

$$\widehat{\mathbf{dM}}_t^{\text{MMSE}} \simeq \frac{1}{N_{\text{MC}} - N_{\text{bi}}} \sum_{q=N_{\text{bi}}+1}^{N_{\text{MC}}} \mathbf{dM}_t^{(q)} \quad (37)$$

$$\widehat{\mathbf{X}}_t^{\text{MMSE}} \simeq \frac{1}{N_{\text{MC}} - N_{\text{bi}}} \sum_{q=N_{\text{bi}}+1}^{N_{\text{MC}}} \mathbf{X}_t^{(q)}. \quad (38)$$

Finally, the following *marginal maximum a posteriori* (mMAP) estimator is considered for the label maps

$$\widehat{z}_{n,t}^{\text{mMAP}} = \arg \max_{z_{n,t} \in \{0,1\}} p(z_{n,t} | \mathbf{y}_{n,t}, \Theta_{\setminus \{z_{n,t}\}}) \quad (39)$$

and is approximated by

$$\widehat{z}_{n,t}^{\text{mMAP}} \simeq \begin{cases} 0, & \text{if } \#\{q > N_{\text{bi}} : z_{n,t}^{(q)} = 0\} \leq \frac{N_{\text{MC}} - N_{\text{bi}}}{2} \\ 1, & \text{otherwise.} \end{cases} \quad (40)$$

B. Sampling the abundances \mathbf{A}

The likelihood function (4) combined with the prior given in Section III-B1 leads to the following conditional distribution for the abundances

$$\mathbf{a}_{n,t} | \mathbf{y}_{n,t}, \Theta_{\setminus \{\mathbf{a}_{n,t}\}} \sim \mathcal{N}_{\mathcal{S}_R}(\boldsymbol{\mu}_{n,t}^{(\mathbf{A})}, \boldsymbol{\Lambda}_{n,t}) \quad (41)$$

Algorithm 1: Proposed hybrid Gibbs sampler.

Data: $N_{\text{bi}}, N_{\text{MC}}, \mathbf{M}^{(0)}, \mathbf{A}^{(0)}, \mathbf{dM}^{(0)}, \mathbf{X}^{(0)}, \boldsymbol{\sigma}^{2(0)}, \mathbf{Z}^{(0)}, \boldsymbol{\beta}^{(0)}, \mathbf{s}^{2(0)}, \boldsymbol{\Psi}^{2(0)}.$

begin

for $q = 1$ **to** N_{MC} **do**

Sample the endmembers $\mathbf{M}^{(q)}$, cf. §IV-C ;

Sample the variability terms $\mathbf{dM}^{(q)}$, cf. §IV-D;

Sample the abundances $\mathbf{A}^{(q)}$, cf. §IV-B ;

Sample the labels and the outliers $(\mathbf{Z}^{(q)}, \mathbf{X}^{(q)})$, cf. §IV-E;

Sample the outlier variances $\mathbf{s}^{2(q)}$, cf. §IV-F;

Sample the noise variances $\boldsymbol{\sigma}^{2(q)}$, cf. §IV-G;

Sample the variability variances $\boldsymbol{\Psi}^{2(q)}$, cf. §IV-H;

Sample the granularity parameters $\boldsymbol{\beta}^{(q)}$, cf. §IV-I;

Result: $\{\mathbf{M}^{(q)}, \mathbf{dM}^{(q)}, \mathbf{A}^{(q)}, \mathbf{Z}^{(q)}, \mathbf{X}^{(q)}, \boldsymbol{\sigma}^{2(q)}, \mathbf{Z}^{(q)}, \boldsymbol{\beta}^{(q)}, \mathbf{s}^{2(q)}, \boldsymbol{\Psi}^{2(q)}\}_{q=1}^{N_{\text{MC}}}.$

$$\boldsymbol{\Lambda}_{n,t}^{-1} = \frac{1}{\sigma_t^2} \mathbf{M}_t^T \mathbf{M}_t + \frac{1}{\varepsilon_n^2} ([\mathcal{T}_{n,t}^1 \neq \emptyset] + [\mathcal{T}_{n,t}^2 \neq \emptyset]) \mathbf{I}_R \quad (42)$$

$$\mathbf{M}_t \triangleq \mathbf{M} + \mathbf{dM}_t \quad (43)$$

$$\begin{aligned} \boldsymbol{\mu}_{n,t}^{(\mathbf{A})} &= \boldsymbol{\Lambda}_{n,t} \left[\frac{1}{\sigma_t^2} \mathbf{M}_t^T (\mathbf{y}_{n,t} - \mathbf{x}_{n,t}) \right. \\ &\quad \left. + \frac{1}{\varepsilon_n^2} ([\mathcal{T}_{n,t}^1 \neq \emptyset] \mathbf{a}_{n,\tau_{n,t}^1} + [\mathcal{T}_{n,t}^2 \neq \emptyset] \mathbf{a}_{n,\tau_{n,t}^2}) \right] \end{aligned} \quad (44)$$

where $\mathcal{N}_{\mathcal{S}_R}(\boldsymbol{\mu}, \boldsymbol{\Lambda})$ denotes a Gaussian distribution truncated to the set \mathcal{S}_R and

$$\mathcal{T}_{n,t}^2 = \{\tau > t \mid z_{n,\tau} = 0\}, \quad \tau_{n,t}^2 = \min_{\tau \in \mathcal{T}_{n,t}^2} \tau \quad (45)$$

with the convention $\mathcal{T}_{n,t}^2 = \emptyset$ if $t = T$.

Samples distributed according to the above truncated multivariate Gaussian distributions can be generated by a Gibbs sampler described in [42, Section IV.B.] [43], by an Hamiltonian Monte-Carlo procedure [44], [45] or by the general method recently proposed in [46]. In this work, the Gibbs sampler [42, Section IV.B.] has been adopted to sample the parameters of interest. Note that the abundance vectors $\mathbf{a}_{n,t}$ can be sampled in parallel to accelerate the algorithm.

C. Sampling the endmembers \mathbf{M}

Combining (4) and the endmember prior given in Section III-B3 leads to

$$m_{\ell,r} | \mathbf{Y}, \Theta_{\setminus \{m_{\ell,r}\}} \sim \mathcal{N}_{[b_{\ell,r}, +\infty)}(\mu_{\ell,r}^{(\mathbf{M})}, \kappa_{\ell,r}^2) \quad (46)$$

$$b_{\ell,r} = \max\left\{0, \max_t(-dm_{\ell,r,t})\right\} \quad (47)$$

$$\begin{aligned} \mu_{\ell,r}^{(\mathbf{M})} &= \kappa_{\ell,r}^2 \sum_t \frac{1}{\sigma_t^2} [\tilde{\mathbf{y}}_{\ell,t} - \tilde{\mathbf{x}}_{\ell,t} - \tilde{\mathbf{m}}_{\ell,\setminus r} \mathbf{A}_{\setminus r,t} \\ &\quad - \widetilde{\mathbf{dM}}_{\ell,t} \mathbf{A}_t] \tilde{\mathbf{a}}_{r,t}^T \end{aligned} \quad (48)$$

$$\kappa_{\ell,r}^2 = \left[\sum_{n,t} \frac{a_{r,n,t}^2}{\sigma_t^2} + \frac{1}{\bar{\varepsilon}} \right]^{-1} \quad (49)$$

where $\mathbf{dM}_{\ell,t}$ is the ℓ th row of \mathbf{dM}_t , $\tilde{\mathbf{m}}_{\ell,\setminus r}$ is the ℓ th row of \mathbf{M} whose r th entry has been removed, $\mathbf{A}_{\setminus r,t}$ denotes the

matrix \mathbf{A}_t without its r th row and $\tilde{\mathbf{a}}_{r,t}$ is the r th row of \mathbf{A}_t . Samples distributed according to the above truncated Gaussian distributions can be efficiently generated using the algorithm described in [47]. When using a PCA as a preprocessing step (21), the projected endmembers \mathbf{e}_r , for $r = 1, \dots, R$ have a truncated multivariate Gaussian distribution [39]

$$\mathbf{e}_r \mid \mathbf{Y}, \Theta_{\setminus\{\mathbf{e}_r\}} \sim \mathcal{N}_{\mathcal{E}_r}(\boldsymbol{\mu}_r^{(\mathbf{E})}, \boldsymbol{\Lambda}_r) \quad (50)$$

where \mathcal{E}_r , $\boldsymbol{\mu}_r^{(\mathbf{E})}$ and $\boldsymbol{\Lambda}_r$ have been reported in Appendix A. Note that the rows of \mathbf{M} (resp. of the projected endmember matrix \mathbf{E}) can be sampled in parallel to decrease the computational time required by the algorithm.

D. Sampling the variability terms \mathbf{dM}

Similarly, the likelihood function (4) and the prior given in Section III-B4 lead to

$$dm_{\ell,r,t} \sim \mathcal{N}_{[-m_{\ell,r}, +\infty)}(\mu_{\ell,r,t}^{(\mathbf{dM})}, \eta_{\ell,r,t}^2) \quad (51)$$

with

$$\begin{aligned} \frac{1}{\eta_{\ell,r,t}^2} &= \frac{1}{\sigma_t^2} \sum_n a_{r,n,t}^2 + \frac{1}{\nu} [t = 1] \\ &\quad + \frac{1}{\psi_{\ell,r}^2} (1 + [1 < t < T]) \end{aligned} \quad (52)$$

$$\begin{aligned} \mu_{\ell,r,t}^{(\mathbf{dM})} &= \left[\frac{1}{\sigma_t^2} (\tilde{\mathbf{y}}_{\ell,t} - \widetilde{\mathbf{dM}}_{\ell,\setminus r,t} \mathbf{A}_{\setminus r,t} - \tilde{\mathbf{m}}_{\ell} \mathbf{a}_{n,t} - \mathbf{x}_{\ell,n,t}) \tilde{\mathbf{a}}_{r,t}^T \right. \\ &\quad \left. + \frac{1}{\psi_{\ell,r}^2} ([t < T] dm_{\ell,r,(t+1)} + [t > 1] dm_{\ell,r,(t-1)}) \right] \eta_{\ell,r,t}^2 \end{aligned} \quad (53)$$

where $\widetilde{\mathbf{dM}}_{\ell,\setminus r,t}$ denotes the ℓ th row of \mathbf{dM}_t whose r th element has been removed, $\tilde{\mathbf{m}}_{\ell}$ is the ℓ th row of \mathbf{M} and $\mathbf{A}_{\setminus r,t}$ is the matrix \mathbf{A}_t without its r th row. The rows of each variability matrix \mathbf{dM}_t can be sampled in parallel to reduce the computational time of the sampler.

E. Sampling the label maps \mathbf{Z} and the outliers \mathbf{X}

According to (4) and Section III-B2, the outliers admit the following group-sparsity promoting conditional distributions

$$\begin{aligned} p(\mathbf{x}_{n,t} \mid \mathbf{y}_{n,t}, \Theta_{\setminus\{z_{n,t}, \mathbf{x}_{n,t}\}}) &= \\ (1 - w_{n,t}) \delta(\mathbf{x}_{n,t}) + w_{n,t} \mathcal{N}_{\mathbb{R}^L}(\boldsymbol{\mu}_{n,t}, \vartheta_{n,t}^2 \mathbf{I}_L) \end{aligned} \quad (54)$$

which are mixtures of a Dirac mass at $\mathbf{0}$ and of truncated multivariate Gaussian distributions, where

$$w_{n,t} = \frac{\tilde{w}_{n,t}}{\tilde{w}_{n,t} + (1 - w_{n,t})}, \quad \vartheta_{n,t}^2 = \frac{\sigma_t^2 s_t^2}{\sigma_t^2 + s_t^2} \quad (55)$$

$$\tilde{w}_{n,t} = \frac{\omega_{n,t}}{(s_t^2)^{L/2}} (\vartheta_{n,t}^2)^{L/2} \exp\left(-\frac{1}{2\vartheta_{n,t}^2} \|\boldsymbol{\mu}_{n,t}\|^2\right) \quad (56)$$

$$\boldsymbol{\mu}_{n,t}^{(\mathbf{X})} = \frac{s_t^2}{\sigma_t^2 + s_t^2} [\mathbf{y}_{n,t} - (\mathbf{M} + \mathbf{dM}_t) \mathbf{a}_{n,t}]. \quad (57)$$

In practice, the labels $z_{n,t}$ are first sampled according to a Bernoulli distribution to select one of the two models for $\mathbf{x}_{n,t}$, with probability $\mathbb{P}[z_{n,t} = 1 \mid \mathbf{y}_{n,t}, \Theta_{\setminus\{z_{n,t}, \mathbf{x}_{n,t}\}}] = w_{n,t}$. Note that the labels $\mathbf{z}_{n,t}$ can be sampled in parallel by using a checkerboard scheme. In addition, the outliers $\mathbf{x}_{n,t}$ can be sampled in parallel to decrease the computational time.

F. Sampling the outlier variances s^2

According to Sections III-B2 and III-C, we can easily identify the conditional law of s_t^2 for $t = 1, \dots, T$ as the following inverse gamma distribution

$$\begin{aligned} s_t^2 \mid \Theta_{\setminus\{s_t^2\}} &\sim \\ \mathcal{IG}\left(a_s + \frac{\#\{n : z_{n,t} = 1\}L}{2}, b_s + \frac{1}{2} \|\mathbf{X}_t\|_F^2\right). \end{aligned} \quad (58)$$

G. Sampling the noise variances σ^2

Using Sections III-B5 and III-C, we obtain for $t = 1, \dots, T$

$$\begin{aligned} \sigma_t^2 \mid \mathbf{Y}_t, \Theta_{\setminus\{\sigma_t^2\}} &\sim \mathcal{IG}\left(a_\sigma + \frac{LN}{2}, b_\sigma + \right. \\ &\quad \left. \frac{1}{2} \|\mathbf{Y}_t - (\mathbf{M} + \mathbf{dM}_t) \mathbf{A}_t - \mathbf{X}_t\|_F^2\right). \end{aligned} \quad (59)$$

H. Sampling the variability variances Ψ^2

Similarly, Section III-B4 and III-C lead to

$$\begin{aligned} \psi_{\ell,r}^2 \mid \Theta_{\setminus\{\psi_{\ell,r}^2\}} &\sim \mathcal{IG}\left(a_\Psi + \frac{T-1}{2}, b_\Psi + \right. \\ &\quad \left. \frac{1}{2} \sum_{t=2}^T (dm_{\ell,r,t} - dm_{\ell,r,t-1})^2\right). \end{aligned} \quad (60)$$

I. Sampling the granularity parameters β_t

Provided square images are considered, the partition functions $C(\beta_t)$ have the closed-form expressions [38], [48]

$$\begin{aligned} \tilde{C}(\beta_t) &= \frac{1}{2} \log(2 \sinh \beta_t) + \frac{1}{2N} \sum_{n=1}^N \operatorname{acosh} \Delta_n(\beta_t) + \beta_t \\ \Delta_n(\beta_t) &= v(\beta_t) - C_n, \quad v(\beta_t) = \frac{\cosh^2 \beta_t}{\sinh \beta_t} \\ \tilde{C}(\beta_t) &= \frac{1}{N} \log C(\beta_t), \quad C_n = \cos\left(\frac{2n-1}{2N} \pi\right). \end{aligned}$$

The exact partition function can then be used to sample the parameters β_t using Metropolis-Hastings steps. In this work, new values of the granularity parameters have been proposed by the following Gaussian random walk

$$\beta_t^* = \beta_t^{(q)} + \varepsilon_t, \quad \varepsilon_t \sim \mathcal{N}(0, \sigma_{\beta}^2(t)) \quad (61)$$

where the parameters $\sigma_{\beta}^2(t)$ are adjusted during the burn-in iterations to yield acceptance rates in the interval $[0.4, 0.6]$. More precisely, the acceptance rates ρ_t are given by

$$\begin{aligned} \rho_t &= \max\left(1, \frac{C(\beta_t^{(q)})}{C(\beta_t^*)} \frac{\mathbb{1}_{[0,2]}(\beta_t^*)}{\mathbb{1}_{[0,2]}(\beta_t^{(q)})}\right) \\ &\quad \times \exp\left[\frac{1}{2} (\beta_t^* - \beta_t^{(q)}) \sum_{n=1}^N \sum_{k \in \mathcal{V}(n)} \delta(z_{n,t} - z_{k,t})\right]. \end{aligned} \quad (62)$$

J. Computational complexity

Assuming elementary arithmetic operations and scalar pseudo-random number generations are $O(1)$ operations, the overall computational complexity is dominated by matrix products needed to compute the parameters related to the

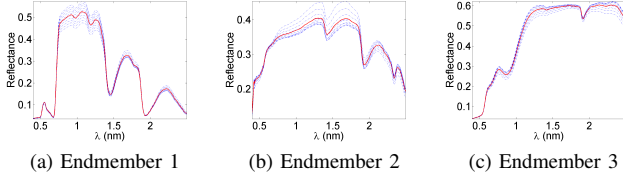


Fig. 2. Endmembers (\mathbf{m}_r , red lines) and their variants affected by variability ($\mathbf{m}_r + \mathbf{d}\mathbf{m}_{r,t}$, blue dotted lines) used to generate the synthetic mixtures with $R = 3$. Signatures corresponding to different time instants are represented in a single figure to better appreciate the variability introduced in the data.

conditional distribution of the variability vectors. Since $R \ll L \ll N$ and $T \ll L$, the per-iteration computational cost of the proposed algorithm is $O(LR^2NT)$. As detailed in the preceding paragraphs, many parameters can be sampled in parallel to reduce the computational time of the proposed algorithm.

V. EXPERIMENTS WITH SYNTHETIC DATA

The proposed method has been applied to MTHS images composed of 10 acquisitions of size 50×50 with $L = 413$ bands. The first scenario deals with the apparition of a new material in specific regions of a few images. To this end, 4 images out of the 10 have been corrupted by spatially sparse outliers, corresponding to a new endmember extracted from a spectral library. Each image of the sequence corresponds to a linear mixture of 3 endmembers affected by smooth time-varying variability, and the synthetic abundances vary smoothly from one image to another. Note that the temporal variations of the variability and the abundances have been generated from the parameters of the first image with trigonometric functions. The mixtures have finally been corrupted by an additive white Gaussian noise to ensure a resulting signal-to-noise ratio (SNR) between 25 and 30 dB. Similarly, two complementary scenarios involving 5 HS images, of size 100×100 , composed of 6 and 9 endmembers, have been considered to analyze the performance of the method in the presence of a larger number of endmembers. Note that all the images of this experiment do not satisfy the pure pixel assumption to assess the proposed method in challenging situations.

Controlled spectral variability has been introduced by using the product of reference endmembers with randomly generated piecewise-affine functions as in [9], where different affine functions have been generated for each endmember at each time instant. Typical instances of the signatures used in this experiment are depicted in Fig. 2. The robustness of the proposed method to moderate spatial variability, i.e., endmember variability occurring within single images, has also been evaluated. The corresponding results can be found in the technical report [49, Appendix B] due to space constraints.

A. Compared methods

The results of the proposed algorithm have been compared to those of several unmixing methods from the literature, some of which are specifically designed to unmix a single HS image. In the following lines, the most relevant implementation details specific to each method are briefly recalled.

TABLE I
FIXED PARAMETERS, AND INITIAL VALUES ASSOCIATED IN THE EXPERIMENTS TO PARAMETERS LATER INFERRED FROM THE MODEL.

		Synthetic data	Real data
Fixed parameters	ε_n^2	10^{-3}	10^{-2}
	ξ	1	1
	ν	10^{-3}	10^{-5}
	a_s, a_ψ, a_σ	10^{-3}	10^{-3}
	b_s, b_ψ, b_σ	10^{-3}	10^{-3}
	N_{bi}	350	450
	N_{MC}	400	500
Initial values	σ_t^2	10^{-4}	10^{-4}
	s_t^2	5×10^{-3}	5×10^{-3}
	$\psi_{\ell,r}^2$	10^{-3}	10^{-2}
	β_t	1.7	1.7

- 1) VCA/FCLS (no variability, single image): the endmembers are first extracted on each image using the vertex component analysis (VCA) [22], which requires pure pixels to be present. The abundances are then estimated for each pixel by solving a fully constrained least squares problem (FCLS) using the Alternating Direction Method of Multipliers (ADMM) [23];
- 2) SISAL/FCLS (no variability, single image): the endmembers are extracted on each image by the simplex identification via split augmented Lagrangian (SISAL) [24], and the abundances are estimated for each pixel by FCLS. The tolerance for the stopping rule has been set to 10^{-3} ;
- 3) RLMM (no variability, single image): the unmixing method associated with the robust linear mixing model (RLMM) proposed in [25] has been applied to each image of the series independently. The algorithm has been initialized with SISAL/FCLS, and the regularization parameter specific to this method is set as suggested in [25];
- 4) OU: the endmembers are estimated using the online unmixing (OU) algorithm introduced in [17] with endmembers initialized by the output of VCA applied to the first image of the sequence. The abundances are initialized by FCLS, and the variability matrices are initialized with all their entries equal to 0. The other parameters are set to the same values as those given in [17, Table I];
- 5) Proposed approach: the endmembers are initialized with VCA applied to the first image of the sequence. The abundances are initialized by FCLS, and the variability matrices and label maps are initialized with all their entries equal to 0 (i.e., the images are a priori assumed to contain no outlier). The values chosen for the other parameters are summarized in Table I. Further details related to the choice of these values can be found in Appendix B.

Performance assessment has been conducted in terms of

- (i) endmember estimation through the average spectral angle mapper (aSAM)

$$\text{aSAM}(\mathbf{M}) = \frac{1}{R} \sum_{r=1}^R \arccos \left(\frac{\mathbf{m}_r^T \hat{\mathbf{m}}_r}{\|\mathbf{m}_r\|_2 \|\hat{\mathbf{m}}_r\|_2} \right); \quad (63)$$

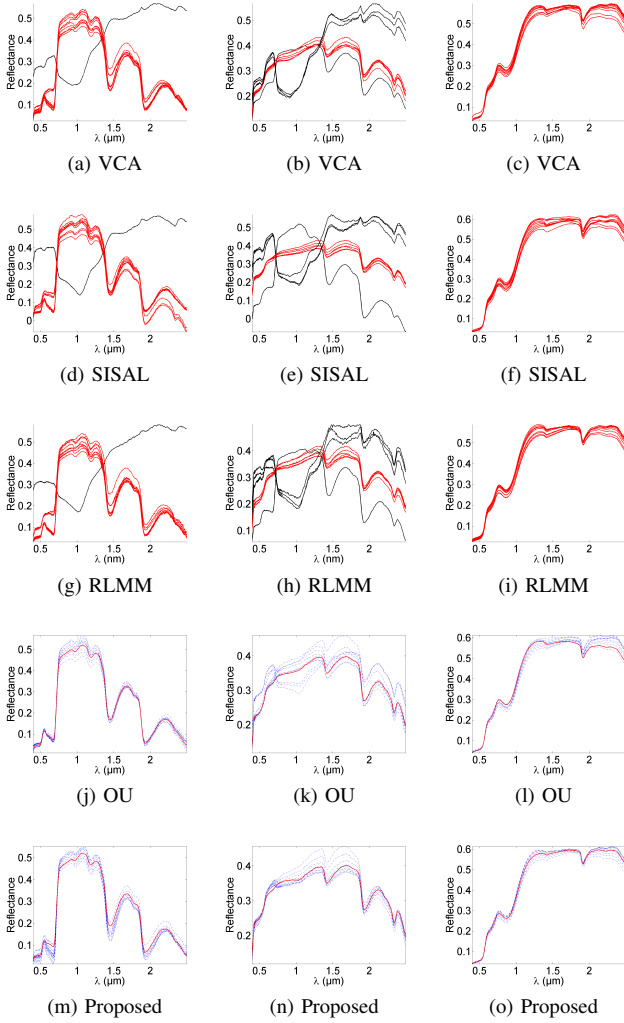


Fig. 3. Endmembers (\mathbf{m}_r , red lines) and their variants affected by variability ($\mathbf{m}_r + \mathbf{d}\mathbf{m}_{r,t}$, blue dotted lines) recovered by the different methods from the synthetic mixtures with $R = 3$. Signatures corresponding to different time instants are represented on a single figure to better appreciate the variability recovered from the data. The spectra represented in black in Figs. 3a, 3b, 3d, 3e and 3h correspond to signatures corrupted by outliers.

(ii) abundance and variability estimation through the global mean square errors (GMSEs)

$$\text{GMSE}(\mathbf{A}) = \frac{1}{TRN} \sum_{t=1}^T \|\mathbf{A}_t - \hat{\mathbf{A}}_t\|_F^2 \quad (64)$$

$$\text{GMSE}(\mathbf{dM}) = \frac{1}{TLR} \sum_{t=1}^T \|\mathbf{dM}_t - \hat{\mathbf{dM}}_t\|_F^2; \quad (65)$$

(iii) quadratic reconstruction error (RE)

$$\text{RE} = \frac{1}{TLN} \sum_{t=1}^T \|\mathbf{Y}_t - \hat{\mathbf{Y}}_t\|_F^2 \quad (66)$$

where $\hat{\mathbf{Y}}_t$ is the matrix composed of the pixels reconstructed with the estimated parameters.

B. Results

The endmembers estimated by the proposed algorithm are compared to those of VCA/FCLS, SISAL/FCLS, RLMM and OU in Fig. 3, whereas the corresponding abundance maps are

TABLE II
SIMULATION RESULTS ON SYNTHETIC DATA (ASAM(M) IN (°), GMSE(A) $\times 10^{-2}$, GMSE(dM) $\times 10^{-4}$, RE $\times 10^{-4}$, TIME IN (s)).

		aSAM(M)	GMSE(A)	GMSE(dM)	RE	time
$R = 3$	VCA/FCLS	6.07	2.32	/	3.91	1
	SISAL/FCLS	5.07	1.71	/	2.28	2
	RLMM	5.13	2.04	/	0.31	463
	OU	1.90	0.42	3.22	2.61	98
	Proposed	2.03	0.15	1.85	2.00	2530
$R = 6$	VCA/FCLS	3.81	1.57	/	3.09	2
	SISAL/FCLS	5.76	0.91	/	4.49	3
	RLMM	2.73	1.26	/	0.29	1453
	OU	2.74	0.38	3.70	1.13	420
	Proposed	1.48	0.16	2.84	0.51	8691
$R = 9$	VCA/FCLS	3.74	0.65	/	6.83	4
	SISAL/FCLS	5.91	0.36	/	5.56	5
	RLMM	2.48	0.54	/	0.31	1447
	OU	6.08	0.47	2.19	0.89	1024
	Proposed	2.23	0.15	8.38	0.82	17151

displayed in Fig. 4. Note that the abundance maps and the endmembers obtained for the mixtures of 6 and 9 endmembers are included in a separate technical report [49] due to space constraints. The unmixing performance of each method, reported in Table II, leads to the following conclusions.

- **Endmember estimation:** the proposed method shows an interesting robustness with respect to spatially sparse outliers in the sense that the estimated signatures (Figs. 3m to 3o) are very close to the corresponding ground truth (Fig. 2). In comparison, the shape of the endmembers recovered by VCA, SISAL and RLMM and the variability extracted by OU are significantly affected by outliers, as exemplified in Figs. 3b, 3e, 3h and 3k respectively. These qualitative results are confirmed by the quantitative performance measures of each method provided in Table II. Note that the endmembers recovered by the SISAL and RLMM methods are very sensitive to the VCA initialization, as illustrated by the similarity between the signatures estimated by these methods (Figs. 3a to 3i).
- **Abundance estimation:** the abundance maps estimated by FCLS, RLMM and SISAL reflect the high sensitivity of VCA (used to initialize SISAL and RLMM) to the presence of outliers (see the figures delineated in red in Fig. 4). On the contrary, the abundances recovered by OU and the proposed approach are much closer to the ground truth. These observations are confirmed by the abundance estimation performance reported in Table II.
- **Overall performance:** the performance measures reported in Table II are globally favorable to the proposed approach. It is important to mention that the price to pay with the good performance of the proposed method is its computational complexity, which is common with MCMC methods.

As a complementary output, the proposed algorithm is able to recover the location of the outliers within each image, as illustrated in Fig. 5. Up to a few false detections, the estimated labels are very close to the ground truth.

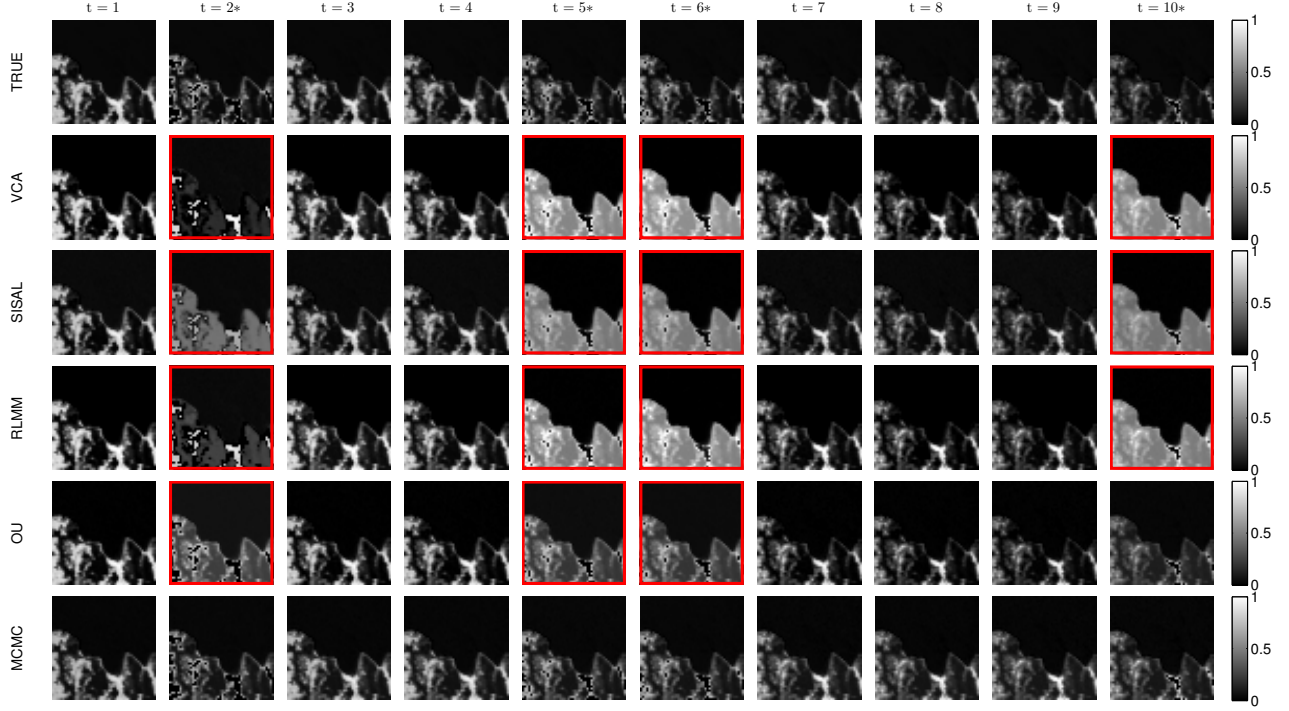


Fig. 4. Abundance map of the first endmember recovered by the different methods (in each row) at each time instant (given in column) for the experiment with $R = 3$ [the different rows correspond to the true abundances, VCA/FCLS, SISAL/FCLS, RLMM, OU and the proposed method]. The images delineated in red show that several methods are highly sensitive to the presence of outliers, and the time instants represented with * denote images containing outliers. Due to space restrictions, the abundance maps of the two other endmembers have been included in the technical report [49].

VI. EXPERIMENT WITH REAL DATA

A. Description of the dataset

We consider a real sequence of AVIRIS HS images acquired over the Lake Tahoe region (California, United States of America) between 2014 and 2015¹. The scene of interest (100×100), composed of a lake and a nearby field, has been unmixed with $R = 3$ endmembers based on the results of the noise-whitened eigengap algorithm (NWEGA) [50] applied to each image of the series (see Table III). This choice is further supported by results obtained from a previous analysis conducted on the same dataset [51, Appendix E]. For $R = 4$ and 5, the signatures of water, soil and vegetation were split into two or more components by the different algorithms, suggesting $R = 3$ is more appropriate for this study. Note that prior studies led in [17] revealed that this dataset contains outliers (area delineated in red in Fig. 6e). After removing the seemingly corrupted bands and the water absorption bands, 173 out of the 224 spectral bands were finally exploited. The initial parameters used for the proposed algorithm are given in Table I. The other methods have been run with the same parameters as in Section V.

B. Results

In the absence of any ground truth, the performance of the unmixing methods is assessed in terms of RE (Table IV) while taking into account the consistency of the estimated abundance

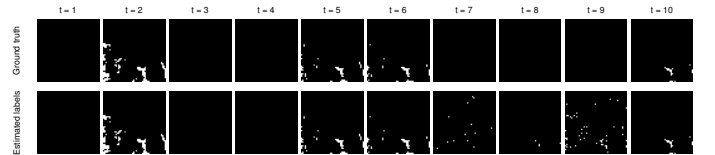


Fig. 5. Outlier labels z_t estimated for each image of the synthetic dataset with 3 endmembers (the different rows correspond to the true labels, and the estimated labels) [0 in black, 1 in white].

maps reported in Figs. 8, 9 and 10. More precisely, the abundances associated with the vegetation area are expected to be very high for $t = 1, 3, 5$ (corresponding to Figs. 6a, 6c and 6e) where the vegetation visually appears to be sufficiently irrigated (hence well represented). On the contrary, the abundance coefficients are supposed to be much lower for $t = 2, 4, 6$ (corresponding to Figs. 6b, 6d and 6f), where the vegetation is visually drier or almost absent. Concerning the presence of water in the bottom left-hand corner of the images, the latent variables introduced in Section III-B2 are expected to reflect the abrupt variations in the presence of water observed at $t = 3, 4$ and 5. These observations, combined with the extracted signatures (Fig. 7) and the estimated abundances (Figs. 8 to 10) lead to the following comments.

- **Endmember estimation:** the signature recovered for the soil by VCA, SISAL and RLMM at time $t = 5$ shows an amplitude significantly greater than one, and a shape incompatible with what can be expected based on physical considerations (see the black signatures in Figs. 7a, 7d and 7g). This is a clear indication that outliers are present in the corresponding image. A similar

¹The images used in this experiment are freely available from the online AVIRIS flight locator tool at http://aviris.jpl.nasa.gov/alt_locator/.



Fig. 6. Scenes used in the experiment, given with their respective acquisition date. The area delineated in red in Fig. 6e highlights a region known to contain outliers (this observation results from a previous analysis led on this dataset in [17]).

observation can be made for the vegetation signature obtained by VCA, SISAL and RLMM at time $t = 5$. On the contrary, the endmembers recovered by OU and the proposed approach are much more consistent from this point of view.

- **Abundance estimation:** the estimated abundances globally reflect the previous comments made on the extracted endmembers. Notably, the abundance coefficients estimated at $t = 5$ by VCA, SISAL and RLMM (delineated in red in Figs. 8 to 10) are visually inconsistent with the temporal evolution of the materials observed in the true color composition given in Fig. 6. More explicitly, the soil is not supposed to be concentrated on a few pixels as suggested by the corresponding abundance maps in Fig. 8. Similarly, the water is not supposed to be present in high proportions in all the pixels of the image as indicated in Fig. 9. These results, in contradiction with Fig. 6, suggest that outliers are present at $t = 5$. In addition, the abundance maps estimated at $t = 4$ and 6 by FCLS for the soil and the water (delineated in green in Figs. 8 and 9) suggest that the water contribution has been split into two spectra. The corresponding signatures are represented in green in Figs. 7a and 7c. On the contrary, the results reported for OU and the proposed method are consistent with the expected evolution of water and vegetation over time (abundance values close to 1 at time $t = 1, 3, 5$, lower values at time $t = 2, 4, 6$). Finally, the vegetation abundance maps estimated by the proposed method globally presents a better contrast than those obtained with OU (Fig. 10).

The previous comments, along with the lower reconstruction error reported in Table IV, suggest that the proposed approach is robust to spatially sparse outliers while allowing smooth temporal variations to be exploited. Indeed, the pixels corresponding to abrupt variations of the water signature have been properly detected. Furthermore, the outliers previously detected in this dataset [17] for $t = 5$ (highlighted in red in Fig. 6e) are well captured by the latent variables \mathbf{Z} (see Fig. 11). In addition, the spatial distribution of the estimated outlier labels (Fig. 11) is in agreement with the results of the RLMM (in terms of the spatial distribution of the outlier energy) and with the non-linearity detector [52] applied to each image of the sequence with the SISAL-estimated endmembers

TABLE III
ENDMEMBER NUMBER R ESTIMATED BY NWEGA [50] ON EACH IMAGE OF THE REAL DATASET.

	04/10/2014	06/02/2014	09/19/2014	11/17/2014	04/29/2015	10/13/2015
NWEGA	3	3	3	4	3	4

TABLE IV
SIMULATION RESULTS ON REAL DATA ($RE \times 10^{-4}$).

		RE	time (s)
$R = 3$	VCA/FCLS	45.05	1
	SISAL/FCLS	1.65	2
	rLMM	2.51	390
	OU	2.50	508
	Proposed	0.34	23608

(see Fig. 12). Concentrated on regions where non-linear effects can be reasonably expected, the active latent variables \mathbf{Z} tend to capture the spatial distribution of the non-linearities possibly occurring in the observed scene.

VII. CONCLUSION AND FUTURE WORK

This paper introduced a Bayesian model accounting for both smooth and abrupt variations possibly occurring in multitemporal hyperspectral images. The adopted model was specifically designed to handle datasets in which mostly the same materials were expected to be observed at different time instants, thus allowing information redundancy to be exploited. An MCMC algorithm was derived to solve the resulting unmixing problem in order to precisely assess the performance of the proposed approach on multitemporal HS images of moderate size (i.e., moderate spatial and temporal dimensions). This algorithm was used to sample the posterior of the proposed hierarchical Bayesian model and to use the generated samples to build estimators of the unknown model parameters. Future research perspectives include the use of relaxation methods to the Ising field to tackle similar problems with online optimization techniques, and the development of distributed unmixing procedures to efficiently unmix larger datasets. Designing unmixing methods scaling with the problem dimension while simultaneously accounting for temporal and spatial endmember variability is another interesting prospect.

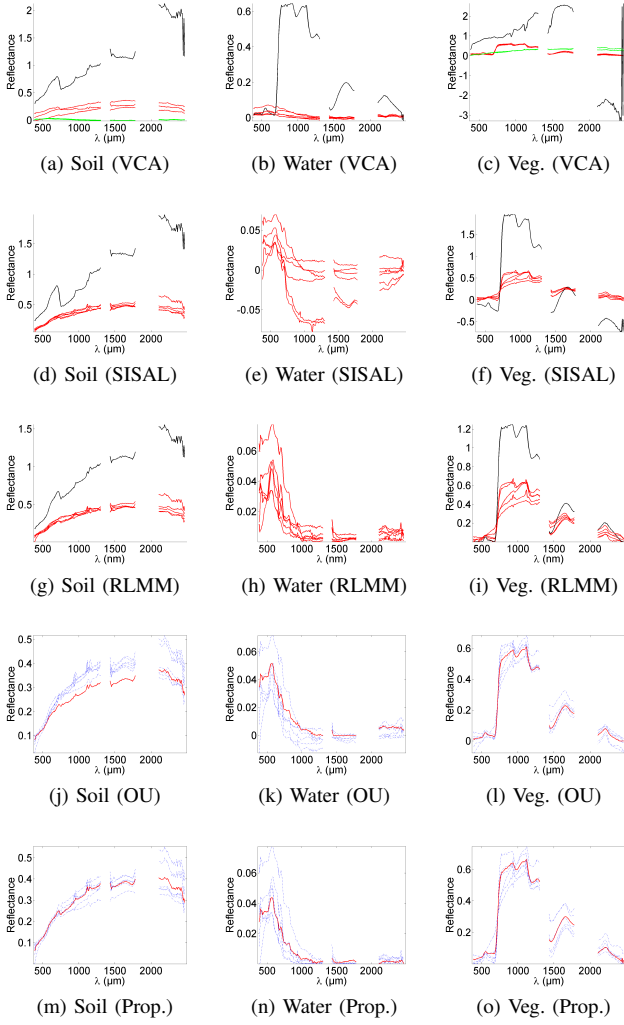


Fig. 7. Endmembers (\mathbf{m}_r , red lines) and their variants affected by variability ($\mathbf{m}_r + \mathbf{d}\mathbf{m}_{r,t}$, blue dotted lines) recovered by the different methods from the real dataset depicted in Fig. 6. Signatures corresponding to different time instants are represented in a single figure to better appreciate the variability recovered from the data. The spectra represented in black correspond to signatures corrupted by outliers, while those given in green represent endmembers which have been split into several components by the associated estimation procedure.

APPENDIX A SAMPLING THE PROJECTED ENDMEMBERS \mathbf{E}

When using a PCA as a preprocessing step, the projected endmembers \mathbf{e}_r , for $r = 1, \dots, R$, are distributed according to the following truncated Gaussian distributions

$$\mathbf{e}_r \mid \mathbf{Y}, \Theta_{\setminus \{\mathbf{e}_r\}} \sim \mathcal{N}_{\mathcal{E}_r}(\boldsymbol{\mu}_r^{(\mathbf{E})}, \boldsymbol{\Lambda}_r) \quad (67)$$

with, for $k = 1, \dots, K$

$$\mathcal{E}_r = [c_{1,r}, d_{1,r}] \times \dots \times [c_{K,r}, d_{K,r}] \quad (68)$$

$$c_{k,r} = \max_{\ell \in \mathcal{U}_k^+} \left(-\frac{\tilde{y}_\ell + \sum_{j \neq k} u_{\ell,j} e_{j,r} + b_{\ell,r}}{u_{\ell,k}} \right) \quad (69)$$

$$d_{k,r} = \min_{\ell \in \mathcal{U}_k^-} \left(-\frac{\tilde{y}_\ell + \sum_{j \neq k} u_{\ell,j} e_{j,r} + b_{\ell,r}}{u_{\ell,k}} \right) \quad (70)$$

$$b_{\ell,r} = \min \left\{ 0, \min_t (d\mathbf{m}_{\ell,r,t}) \right\} \quad (71)$$

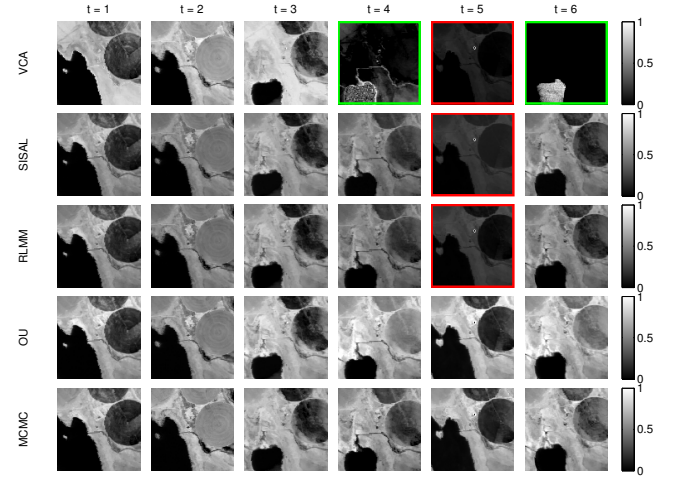


Fig. 8. Soil abundance map recovered by the different methods (in each row) at each time instant (given in column) for the experiment on the real dataset [the different rows correspond to VCA/FCLS, SISAL/FCLS, RLMM, OU, and the proposed method]. On the one hand, the images delineated in red suggest that some of the methods are particularly sensitive to the presence of outliers. On the other hand, the images delineated in green represent the abundance maps associated with signatures which have been split into two components by the corresponding unmixing procedures.

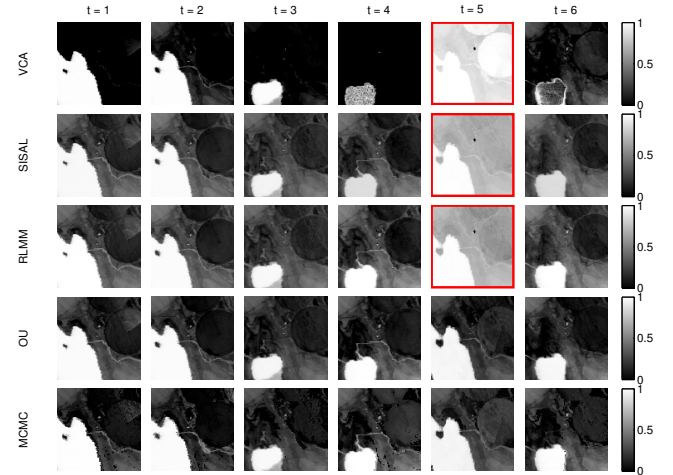


Fig. 9. Water abundance map recovered by the different methods (in each row) at each time instant (given in column) for the experiment on the real dataset [the different rows correspond to VCA/FCLS, SISAL/FCLS, RLMM, OU, and the proposed method]. The images delineated in red suggest that some of the methods are particularly sensitive to the presence of outliers.

$$\boldsymbol{\Lambda}_r^{-1} = \left[\frac{1}{\xi} + \sum_{n,t} \frac{a_{r,n,t}^2}{\sigma_t^2} \right] \mathbf{I}_{R-1} \quad (72)$$

$$\boldsymbol{\mu}_r^{(\mathbf{E})} = \boldsymbol{\Lambda}_r \mathbf{U}^T \left[\sum_{t,n} \frac{1}{\sigma_t^2} \left(\mathbf{y}_{n,t} - \mathbf{x}_{n,t} - \mathbf{d}\mathbf{M}_t \mathbf{a}_{n,t} - \tilde{\mathbf{y}} a_{r,n,t} - \sum_{j \neq r} a_{j,n,t} \mathbf{m}_j \right) a_{r,n,t} \right]. \quad (73)$$

APPENDIX B DETAILS ON THE VALUES CHOSEN IN TABLE I

The initial values chosen for the parameters in Table I are based on the following considerations.

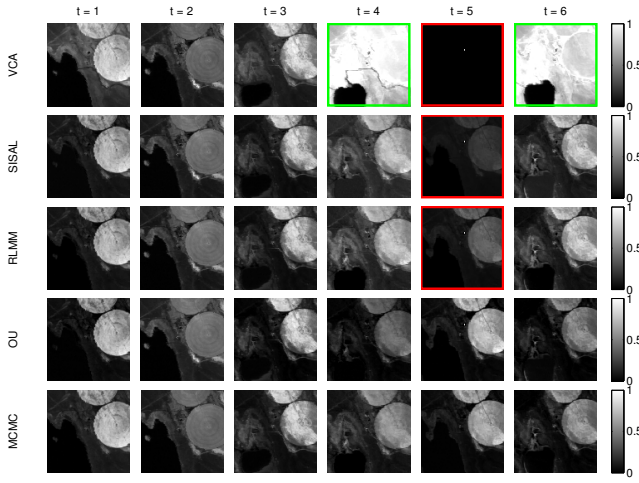


Fig. 10. Vegetation abundance map recovered by the different methods (in each row) at each time instant (given in column) for the experiment on the real dataset [the different rows correspond to VCA/FCLS, SISAL/FCLS, RLMM, OU, and the proposed method]. On the one hand, the images delineated in red suggest that some of the methods are particularly sensitive to the presence of outliers. On the other hand, the images delineated in green represent the abundance maps associated with signatures which have been split into two components by the corresponding unmixing procedures.

- The initial noise variance σ_t^2 reflects a typical order of magnitude of the SNR (HS images are typically known to have an SNR around 30 dB).
- The initial outlier variance s_t^2 has been taken an order of magnitude greater than σ_t^2 to ensure outlier contributions that can be captured by the algorithm.
- The initial variability variance $\psi_{\ell,r}^2$, which controls the temporal smoothness of the variability term, can be *a priori* chosen of the same order of magnitude as s_t^2 .
- The granularity parameters β_t were initially selected to reflect the practitioner's prior knowledge on the smoothness of the outlier spatial support. A value between 1 and 2 (i.e., above the phase-transition temperature of the Ising MRF [53]) is particularly appropriate for natural scenes, in which the observed materials exhibit a relatively smooth spatial distribution.

Similarly, the values of the fixed parameters given in Table I are selected as follows.

- ε_n^2 , which controls the confidence given to the abundance smoothness prior, has been fixed by cross-validation (i.e., based on the estimation results obtained by multiple runs for different values);
- since no specific prior knowledge is available on the endmembers, the endmember variance ξ is chosen sufficiently large (typically equal to 1) to ensure the endmember prior is weakly informative;
- ν , which controls the energy of the variability captured by the algorithm, has been set by cross validation;
- given the absence of specific prior knowledge on the outlier variances s_t^2 , the variability variances $\psi_{\ell,r}^2$ and the noise variances σ_t^2 , the hyperparameters $a_s, a_\psi, a_\sigma, b_s, b_\psi, b_\sigma$ are set to a small value (typically 10^{-3}) to ensure the chosen conjugate inverse-gamma priors are uninformative [20];
- the number of Monte-Carlo and burn-in iterations, re-

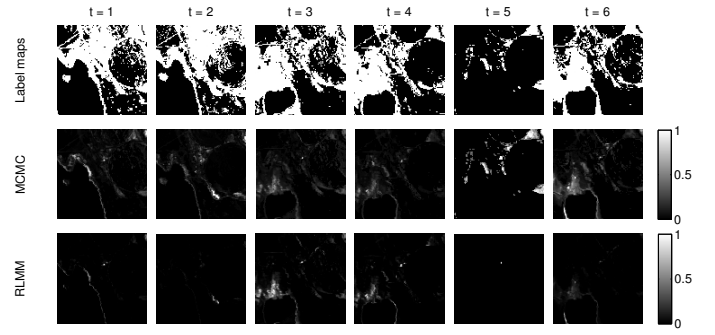


Fig. 11. mMAP estimates of the label maps recovered by the proposed approach, displayed at each time instant (the different rows correspond to: the estimated label map (pixels detected as outliers appear in white), the outlier energy map re-scaled in the interval $[0, 1]$ obtained by the proposed method, and by RLMM).

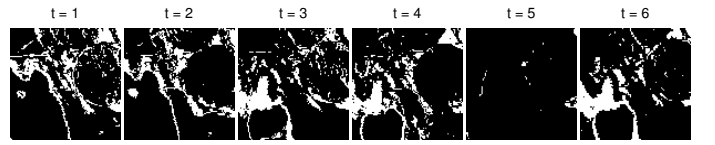


Fig. 12. Non-linearity maps estimated by the detector [52] applied to each image with the SISAL-extracted endmembers, with a probability of false alarm of 10^{-3} (pixels detected as non-linearities appear in white).

spectively denoted by N_{MC} and N_{bi} , are set according to a classical convergence diagnosis, namely the potential scale reduction factor (PSRF) [54].

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