

Learning Nonnegative Factors From Tensor Data: Probabilistic Modeling and Inference Algorithm

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Abstract—Tensor canonical polyadic decomposition (CPD) with nonnegative factor matrices, which extracts useful latent information from multidimensional data, has found wide-spread applications in various big data analytic tasks. Currently, the implementation of most existing algorithms needs the knowledge of tensor rank. However, this information is practically unknown and difficult to acquire. To address this issue, a probabilistic approach is taken in this paper. Different from previous works, this paper firstly introduces a sparsity-promoting nonnegative Gaussian-gamma prior, based on which a novel probabilistic model for the CPD problem with nonnegative and continuous factors is established. This probabilistic model further enables the derivation of an efficient inference algorithm that accurately learns the nonnegative factors from the tensor data, along with an integrated feature of automatic rank determination. Numerical results using synthetic data and real-world applications are presented to show the remarkable performance of the proposed algorithm.

Index Terms—Tensor decomposition, nonnegative factors, variational inference, automatic rank determination.

I. INTRODUCTION

MODERN society generates large amounts of data, much of which has multiple attributes, and tensors provide a natural representation for such data. Because these data often consist of latent components, tensor canonical polyadic decomposition (CPD) [1], which is defined as a linear combination of rank-1 tensors, has been very popular and successful in achieving state-of-the-art performance for various big data mining tasks

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including social group clustering [2]–[4], text mining [5], [6], drug discovery [7], environmental and biomedical analysis [8], [9], and channel estimation in the next generation communications [10], [11]. Compared to the standard matrix decomposition which provides a “flat-world view,” tensor CPD retains the information along all the data dimensions by virtue of its multi-linear structure, and is capable of producing unique and physically meaningful latent components both theoretically [1], [12] and empirically [2]–[11]. That is why tensor CPD has witnessed increasing popularity and adoption in applications mentioned above and beyond.

Despite the inherent uniqueness of the basic CPD, incorporating side information into the CPD model could further improve its identifiability and interpretability [13]. This has triggered a number of recent studies on developing constrained tensor CPD algorithms for various applications [14], [15], [24], wherein tensor CPD with nonnegative factors plays an important role [16], [17]. Due to the nonnegative constraint imposed on each element of the factor matrices, the combination of latent rank-1 components only allows addition but not subtraction. This leads to a parts-based representation of the data, in the sense that each extracted latent component is a part of the data, thus further enhancing the interpretability of various data analytic results [13]–[17].

However, imposing nonnegative constraints on the CPD model is not easy, since it further complicates the originally non-convex CPD problem [1]. To tackle this, the most popular solution is the nonnegative alternating least-squares (NALS) method [16]. The NALS algorithm iteratively optimizes one factor matrix at a time while holding other factor matrices fixed, and in each iteration, the optimization with respect to a single factor matrix is a nonnegative least-squares problem [16], for which recent parallel algorithms [18], [19] have been proposed. In addition to the NALS framework, recent derivative-based approaches [20], [21], which update all the nonnegative factor matrices in each iteration via first-order optimization methods, were also developed to achieve scalability.

While deriving tensor CPD algorithms with nonnegative factors is possible from a nonlinear programming perspective, their implementations require the knowledge of tensor rank. The tensor rank is defined as the smallest number of rank-one tensors that could be linearly combined to generate the original tensor [1]. Its physical meaning is the number of latent components inside the data. For example, in social group analysis [2], the tensor rank corresponds to the number of clusters of people, or in biomedical data analysis [9], the tensor

rank represents how many different types of chemical species are in the given sample. This knowledge might be obtained from problem-specific domain information in some cases, but most of the time it is unknown and has to be estimated.

Unfortunately, determining the tensor rank is known to be non-deterministic polynomial-time hard (NP-hard) [1]. A common approach is to run multiple algorithms assuming different tensor ranks, and choose the best one in data interpretation. Although this manual tuning procedure is widely adopted in tensor data mining research, heavy computational burden is inevitable. To learn the tensor rank automatically, recent advances in probabilistic inference [5], [6], [22]–[26], [31] integrate the tensor rank learning into its hyper-parameter inference steps, and the Bayesian theory provides a natural recipe for automatic rank determination. However, existing probabilistic CPD models [23]–[26], [31] only considered factor matrices with no constraints or with orthogonal constraints. Although recent works [5], [6], [22] have considered the tensor CPD with nonnegative constraints, their algorithms are tailored to handle count-valued data and factor matrices, which frequently occurs in text mining and link prediction. However, continuous data and factor matrices are prevalent in many applications such as biomedical data analysis [8], [9] and Gaussian noise corrupted image/video/speech processing [23], [24], [36]. Even though some datasets are originally count-valued, any pre-processing procedure such as normalization and centralization would destroy the discrete nature [2]–[4]. Therefore, it is essential to design a probabilistic tensor CPD model that is tailored to continuous data and incorporating nonnegative constraints on its continuous factor matrices.

Designing a probabilistic model is an art. It needs to trade off the expressive power of the model and the tractability of the inference algorithm. A good model should be flexible enough to incorporate information of the problem while simple enough to enable an efficient inference algorithm, and this forms the core of modern research of probabilistic inference [22]–[33], [38]–[40], [49]–[51]. In previous probabilistic CPD models [23]–[26], the Gaussian-gamma prior distribution is used as the primary building block due to its appealing sparsity-promoting and exponentially conjugacy property, which has enabled automatic relevance determination in relevance vector machine [27], [28] and low-rank matrix decomposition [29], [30]. However, it is with an unbounded support and thus cannot model non-negativeness. In this paper, we inspect its nonnegative variant, i.e., the nonnegative Gaussian-gamma prior distribution, and show that it inherits all the desired properties of the Gaussian-gamma prior distribution. Using the nonnegative Gaussian-gamma prior, a novel probabilistic CPD model with explicit nonnegative constraints on its continuous factor matrices is proposed. Then, under the framework of variational inference [32], [33], an efficient inference algorithm with no high-dimensional multiple integration is developed. The resulting algorithm, which includes the NALS algorithm as a special case, is guaranteed to converge and is very flexible in incorporating the most recent advances in optimization [18], [41] for scalability improvement.

The remainder of this paper is organized as follows. Section II presents the formulation for the CPD problem with

nonnegative factors and the challenges ahead. In Section III, a probabilistic CPD model with nonnegative factors is established, based on which an efficient inference algorithm is derived in Section IV. Numerical results with synthetic data and real-world data are reported in Section V. Finally, conclusions are drawn in Section VI.

Notation: Boldface lowercase and uppercase letters will be used for vectors and matrices, respectively. Tensors are written as calligraphic letters. $\mathbb{E}[\cdot]$ denotes the expectation of its argument. Superscript T denotes transpose, and the operator $\text{Tr}(\mathbf{A})$ denotes the trace of a matrix \mathbf{A} . $\|\cdot\|_F$ represents the Frobenius norm of the argument. $\mathcal{N}(\mathbf{x}|\mathbf{u}, \mathbf{R})$ stands for the probability density function of a Gaussian vector \mathbf{x} with mean \mathbf{u} and covariance matrix \mathbf{R} . The $N \times N$ diagonal matrix with diagonal components a_1 through a_N is represented as $\text{diag}\{a_1, a_2, \dots, a_N\}$, while \mathbf{I}_M represents the $M \times M$ identity matrix. The $(i, j)^{\text{th}}$ element, the i^{th} row, and the j^{th} column of a matrix \mathbf{A} are represented by $\mathbf{A}_{i,j}$, $\mathbf{A}_{i,:}$ and $\mathbf{A}_{:,j}$, respectively.

II. CPD WITH NONNEGATIVE FACTORS AND CHALLENGES IN RANK ESTIMATION

Tensor CPD with nonnegative factors has found applications in many different fields [13]–[17]. For illustration, fluorescence data analysis [34], [35] and e-mail data mining applications [2]–[4] are presented in Appendix A. The general problem, which decomposes a N dimensional tensor $\mathcal{Y} \in \mathbb{R}^{J_1 \times J_2 \times \dots \times J_N}$ into a set of nonnegative factor matrices $\{\Xi^{(n)}\}_{n=1}^N$, is formulated as:

$$\min_{\{\Xi^{(n)}\}_{n=1}^N} \left\| \mathcal{Y} - \underbrace{\sum_{r=1}^R \Xi_{:,r}^{(1)} \circ \Xi_{:,r}^{(2)} \circ \dots \circ \Xi_{:,r}^{(N)}}_{\triangleq [\Xi^{(1)}, \Xi^{(2)}, \dots, \Xi^{(N)}]} \right\|_F^2$$

$$\text{s.t. } \Xi^{(n)} \geq \mathbf{0}_{J_n \times R}, n = 1, 2, \dots, N, \quad (1)$$

where symbol \circ denotes vector outer product.

In problem (1), there are two significant challenges. Firstly, nonnegative factor matrices $\{\Xi^{(n)}\}_{n=1}^N$ are complicatedly coupled, resulting in a difficult non-convex optimization problem. To tackle this challenge, alternating optimization is one of the most commonly used techniques. In each iteration, after fixing all but one factor matrices, problem (1) will become a standard nonlinear least-square (LS) subproblem, for which there are various off-the-shelf algorithms for solving it [42], including interior point methods and augmented Lagrangian methods. To scale the solution of each subproblem to handle big tensor data, first-order methods, such as Nesterov-type projected gradient descent [44, page 81 and 90], has been proposed to replace the interior point methods in [18], [19].

Although pioneering works [18], [19] allow the learning of nonnegative factors from big multidimensional data, they still face the second critical challenge of problem (1): how to automatically learn the tensor rank R from the data? With the physical meaning of tensor rank being the number of components/groups inside the data (see Appendix A), this value is usually unknown in practice and its estimation has been shown to be NP-hard [1]. For tensor CPD with no constraints on at least

one factor matrix [23]–[26], this problem has been well solved via a probabilistic inference approach by the employment of the Gaussian-gamma prior distribution. In particular, without loss of generality, consider a machine learning model with parameter $\mathbf{w} \in \mathbb{R}^{M \times 1}$. The model parameter \mathbf{w} consists of S non-overlapped blocks, each of which is denoted as $\mathbf{w}_s \in \mathbb{R}^{M_s \times 1}$. The Gaussian-gamma prior can be expressed as [27]–[30]:

$$p(\mathbf{w}|\{\alpha_s\}_{s=1}^S) = \prod_{s=1}^S p(\mathbf{w}_s|\alpha_s) = \prod_{s=1}^S \mathcal{N}(\mathbf{w}_s|\mathbf{0}_{M_s \times 1}, \alpha_s^{-1} \mathbf{I}_{M_s}), \quad (2)$$

$$p(\{\alpha_s\}_{s=1}^S) = \prod_{s=1}^S \text{gamma}(\alpha_s|a_s, b_s), \quad (3)$$

where α_s is the precision parameter (i.e., the inverse of variance, also called weight decay rate) that controls the relevance of model block \mathbf{w}_s in data interpretation, and $\{a_s, b_s\}_{s=1}^S$ are pre-determined hyper-parameters. There are two important properties of Gaussian-gamma prior that leads to its success and prevalence in a variety of applications [23]–[30], [49]–[51]. Firstly, after integrating the gamma hyper-prior, the marginal distribution of model parameter $p(\mathbf{w})$ is a student's t distribution, which is strongly peaked at zero and with heavy tails, thus promoting sparsity. Secondly, the gamma hyper-prior (3) is conjugate¹ to the Gaussian prior (2). This conjugacy permits the closed-form solution of the variational inference [32], [33], which has recently come up as a major tool in inferring complicated probabilistic models with inexpensive computations.

However, the Gaussian-gamma prior in (2) and (3) cannot be used in the CPD problem with nonnegative factors, since the support of Gaussian probability density function (pdf) in (2) is not restricted to the nonnegative region. This calls for a different prior distribution modeling. An immediate idea might be to replace the Gaussian distribution in Gaussian-gamma prior by the truncated Gaussian distribution with a nonnegative support. However, a closer inspection is needed since there is no existing work discussing whether a gamma distribution is conjugate to a truncated Gaussian distribution with a nonnegative support (see Appendix B for related discussions).

III. PROBABILISTIC MODELING FOR CPD WITH NONNEGATIVE FACTORS

A. Properties of Nonnegative Gaussian-Gamma Prior

The support of the Gaussian pdf in (2) is unbounded, and thus cannot model non-negativeness. On the other hand, the truncated Gaussian prior with a nonnegative support for each model block \mathbf{w}_s can be written as:

$$p^+(\mathbf{w}|\{\alpha_s\}_{s=1}^S) = \prod_{s=1}^S p^+(\mathbf{w}_s|\alpha_s)$$

¹In Bayesian theory, a probability density function (pdf) $p(x)$ is said to be conjugate to a conditional pdf $p(y|x)$ if the resulting posterior pdf $p(x|y)$ is in the same distribution family as $p(x)$.

$$= \prod_{s=1}^S \frac{\mathcal{N}(\mathbf{w}_s|\mathbf{0}_{M_s \times 1}, \alpha_s^{-1} \mathbf{I}_{M_s})}{\int_{\mathbf{0}_{M_s \times 1}}^{\infty} \mathcal{N}(\mathbf{w}_s|\mathbf{0}_{M_s \times 1}, \alpha_s^{-1} \mathbf{I}_{M_s})} \text{U}(\mathbf{w}_s \geq \mathbf{0}_{M_s \times 1}), \quad (4)$$

where the function $\text{U}(\mathbf{w}_s \geq \mathbf{0}_{M_s \times 1})$ is a unit-step function with value one when $\mathbf{w}_s \geq \mathbf{0}_{M_s \times 1}$, and with value zero otherwise. Together with the gamma distributions (3) for modeling the precision parameters $\{\alpha_s\}_{s=1}^S$, we have the nonnegative Gaussian-gamma prior. Even though it is clear that nonnegative Gaussian-gamma prior can model the non-negativeness of model parameters due to the unit-step function $\text{U}(\mathbf{w}_s \geq \mathbf{0}_{M_s \times 1})$, whether it enjoys the advantages of the vanilla Gaussian-gamma prior needs further inspection. In the following, two properties of the nonnegative Gaussian-gamma prior are presented.

Property 1: The gamma distribution in (3) is conjugate to the nonnegative Gaussian distribution in (4). ■

Proof: See Appendix B.

Property 2: After integrating out the precision parameters $\{\alpha_s\}_{s=1}^S$, the marginal pdf of model parameter \mathbf{w} is

$$\begin{aligned} p^+(\mathbf{w}) &= \int p^+(\mathbf{w}|\{\alpha_s\}_{s=1}^S) p(\{\alpha_s\}_{s=1}^S) d\{\alpha_s\}_{s=1}^S \\ &= \prod_{s=1}^S 2^{M_s} \left(\frac{1}{\pi} \right)^{\frac{M_s}{2}} \frac{\Gamma(a_s + M_s/2)}{(2b_s)^{-a_s} \Gamma(a_s)} (2b_s + \mathbf{w}_s^T \mathbf{w}_s)^{-a_s - M_s/2} \\ &\quad \times \text{U}(\mathbf{w}_s \geq \mathbf{0}_{M_s \times 1}). \end{aligned} \quad (5)$$

It is a product of multivariate truncated student's t distributions, each of which is with a nonnegative support.

Proof: See Appendix B. ■

The shape of the marginal distribution $p^+(\mathbf{w})$ is determined by the hyper-parameters $\{a_s, b_s\}_{s=1}^S$. Usually, their values are chosen to be a very small value (e.g., $\epsilon = 10^{-6}$), since as $a_s \rightarrow 0$ and $b_s \rightarrow 0$, a Jeffrey's non-informative prior $p(\alpha_s) \propto \alpha_s^{-1}$ [37] can be obtained. After letting the hyper-parameters $\{a_s, b_s\}_{s=1}^S$ in (5) go to zero, it is easy to obtain the following property.

Property 3: If $a_s \rightarrow 0$ and $b_s \rightarrow 0$, the marginal pdf of the model parameter \mathbf{w} becomes

$$p^+(\mathbf{w}) \propto \prod_{s=1}^S 2^{M_s} \left(\frac{1}{\|\mathbf{w}_s\|_2} \right)^{M_s} \text{U}(\mathbf{w}_s \geq \mathbf{0}_{M_s \times 1}), \quad (6)$$

which is highly peaked at zeros.

As an illustration for Property 2 and 3, univariate marginal pdfs with different hyper-parameters $\{a_s, b_s\}_{s=1}^S$ are plotted in Fig. 1, from which it is clear that the nonnegative Gaussian-gamma prior is sparsity-promoting. Further with the conjugacy property revealed in Property 1, the nonnegative Gaussian-gamma prior is a good candidate for probabilistic modeling with nonnegative model parameters.

B. Probabilistic Modeling of CPD With Nonnegative Factors

In the CPD problem with nonnegative factors in (1), the l th column group $\{\Xi_{:,l}^{(n)}\}_{n=1}^N$, which consists of the l th column of all the factor matrices, can be treated as a model block, since their outer product contributes a rank-1 tensor. Therefore, with the principle of nonnegative Gaussian-gamma prior in the previous subsection, the l th column group $\{\Xi_{:,l}^{(n)}\}_{n=1}^N$ can be modeled

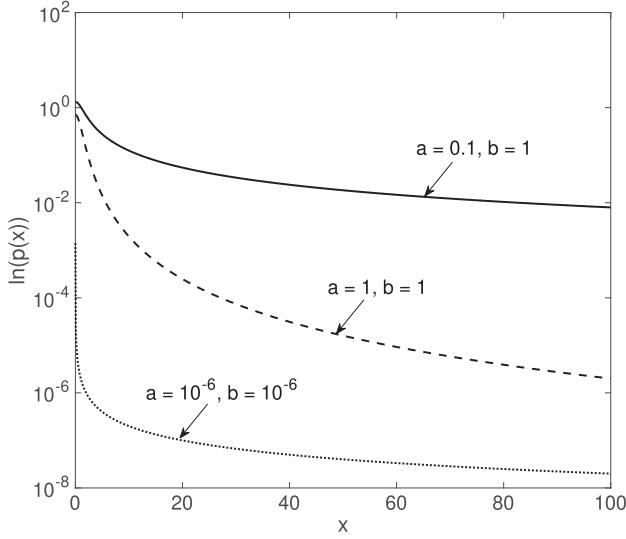


Fig. 1. Univariate marginal probability density functions in (5) with different parameters.

using (4), but with \mathbf{w}_s replaced by $\{\Xi_{:,l}^{(n)}\}_{n=1}^N$ and α_s replaced by γ_l . Considering the independence among different column groups in $\{\Xi^{(n)}\}_{n=1}^N$, we have

$$\begin{aligned} p(\{\Xi^{(n)}\}_{n=1}^N | \{\gamma_l\}_{l=1}^L) \\ = \prod_{n=1}^N \prod_{l=1}^L \frac{\mathcal{N}(\Xi_{:,l}^{(n)} | \mathbf{0}_{J_n \times 1}, \gamma_l^{-1} \mathbf{I}_{J_n})}{\int_{\mathbf{0}_{J_n \times 1}}^{\infty} \mathcal{N}(\Xi_{:,l}^{(n)} | \mathbf{0}_{J_n \times 1}, \gamma_l^{-1} \mathbf{I}_{J_n}) d\Xi_{:,l}^{(n)}} \\ \times \text{U}(\Xi_{:,l}^{(n)} \geq \mathbf{0}_{J_n \times 1}), \end{aligned} \quad (7)$$

$$p(\{\gamma_l\}_{l=1}^L | \boldsymbol{\lambda}_\gamma) = \prod_{l=1}^L \text{gamma}(\gamma_l | c_l^0, d_l^0), \quad (8)$$

where the precision γ_l is modeled as a gamma distribution. From discussions below Property 2, c_l^0 and d_l^0 can be chosen to be a very small value (e.g., $\epsilon = 10^{-6}$) to approach a non-informative prior of precision parameter γ_l .

On the other hand, the least-square objective function in the nonnegative tensor CPD problem (1) motivates the use of a Gaussian likelihood [27]–[30]:

$$\begin{aligned} p(\mathcal{Y} | \Xi^{(1)}, \Xi^{(2)}, \dots, \Xi^{(N)}, \beta) \\ \propto \exp\left(-\frac{\beta}{2} \|\mathcal{Y} - [\Xi^{(1)}, \Xi^{(2)}, \dots, \Xi^{(N)}]\|_F^2\right), \end{aligned} \quad (9)$$

in which the parameter β represents the inverse of noise power. Since there is no information about it, a gamma distribution $p(\beta | \boldsymbol{\alpha}_\beta) = \text{gamma}(\beta | \epsilon, \epsilon)$ with very small ϵ is employed, making $p(\beta | \boldsymbol{\alpha}_\beta)$ approach a Jeffreys non-informative prior.

The Gaussian likelihood function in (9) is with an unbounded support over the real space, and thus it is suitable for applications such as fluorescence data analysis [34], [35] and blind speech separation [36], in which the observed data \mathcal{Y} could be both positive and negative [34]–[36]. On the other hand, if the data \mathcal{Y}

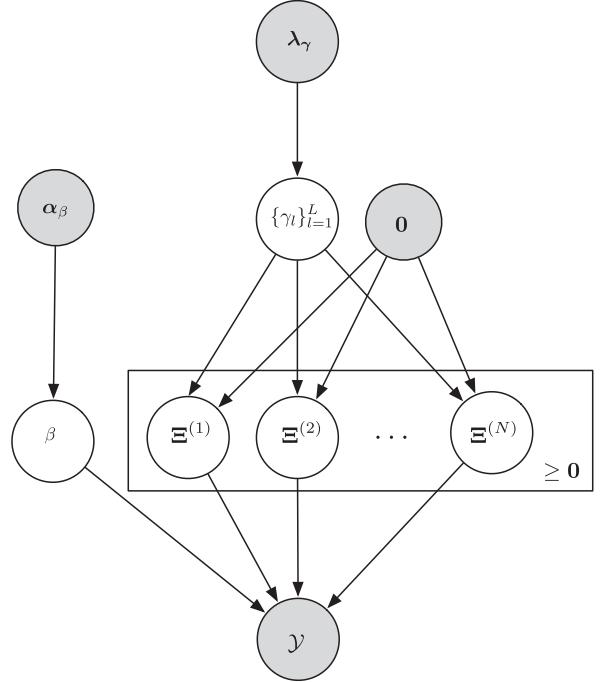


Fig. 2. Probabilistic model for tensor CPD with nonnegative factors.

are all nonnegative and continuous (e.g., the email dataset [2]–[4] after pre-processing), a truncated Gaussian likelihood can be used to model the data:

$$\begin{aligned} p(\mathcal{Y} | \Xi^{(1)}, \Xi^{(2)}, \dots, \Xi^{(N)}, \beta) \\ \propto \exp\left(-\frac{\beta}{2} \|\mathcal{Y} - [\Xi^{(1)}, \Xi^{(2)}, \dots, \Xi^{(N)}]\|_F^2\right) \text{U}(\mathcal{Y} \geq 0). \end{aligned} \quad (10)$$

Finally, the complete probabilistic model is a three-layer Bayes network and is illustrated in Fig. 2.

Remark 1: For applications in Appendix A (i.e., fluorescence spectroscopy and social group clustering), the desired factor matrices are with continuous and nonnegative elements, and thus the prior distribution should have a nonnegative support. On the other hand, in order to automatically identify the inherent component/cluster number, the prior distribution needs to enjoy the sparsity promoting property. Therefore, the choice of nonnegative Gaussian pdf together with a gamma hyper-prior suits these applications.

IV. INFERENCE ALGORITHM FOR TENSOR CPD WITH NONNEGATIVE FACTORS

The unknown parameter set Θ includes the factor matrices $\{\Xi^{(n)}\}_{n=1}^N$, the noise power β^{-1} and the precision parameter $\{\gamma_l\}_{l=1}^L$. The aim of Bayesian inference is to infer the posterior distribution $p(\Theta | \mathcal{Y}) = p(\Theta, \mathcal{Y}) / \int p(\Theta, \mathcal{Y}) d\Theta$. However, the proposed probabilistic model is too complicated to enable an analytical solution since multiple integrations are involved. To tackle this, variational inference [32], [33] has recently been

widely used for inferring parameters of a complicated probabilistic model. The key idea is to approximate the true posterior distribution by a variational pdf $Q(\Theta)$ that minimizes the Kullback-Leibler (KL) divergence $\text{KL}(Q(\Theta) \parallel p(\Theta | \mathcal{Y})) \triangleq -\mathbb{E}_{Q(\Theta)}\{\ln \frac{p(\Theta | \mathcal{Y})}{Q(\Theta)}\}$, thus recasting the probabilistic inference problem into a functional optimization problem. To facilitate the optimization, the variational pdf $Q(\Theta)$ is usually restricted to the mean-field family $Q(\Theta) = \prod_{k=1}^K Q(\Theta_k)$, where Θ is partitioned into mutually disjoint non-empty subsets Θ_k (i.e., Θ_k is a part of Θ with $\bigcup_{k=1}^K \Theta_k = \Theta$ and $\bigcap_{k=1}^K \Theta_k = \emptyset$). Under the mean-field assumption, each optimal variational pdf $Q^*(\Theta_k)$ that minimizes the KL divergence is obtained by solving the following problem with other $\{Q(\Theta_j)\}_{j \neq k}$ fixed [32]:

$$\min_{Q(\Theta_k)} \int Q(\Theta_k) (-\mathbb{E}_{\prod_{j \neq k} Q(\Theta_j)} [\ln p(\Theta, \mathcal{Y})] + \ln Q(\Theta_k)) d\Theta_k. \quad (11)$$

For this convex problem, the Karush-Kuhn-Tucker (KKT) condition gives the optimal solution as [32, page 132]:

$$Q^*(\Theta_k) = \frac{\exp\left(\mathbb{E}_{\prod_{j \neq k} Q(\Theta_j)} [\ln p(\Theta, \mathcal{Y})]\right)}{\int \exp\left(\mathbb{E}_{\prod_{j \neq k} Q(\Theta_j)} [\ln p(\Theta, \mathcal{Y})]\right) d\Theta_k}. \quad (12)$$

Nevertheless, even under the mean-field family assumption, the unknown parameter $\Xi^{(k)}$ is still difficult to be inferred since its moments cannot be easily computed. In particular, in the proposed probabilistic model, if no functional assumption is made for variational pdf $Q(\Xi^{(k)})$, after using (12), a multivariate truncated Gaussian distribution would be obtained, of which the moments are known to be very difficult to be computed due to the multiple integrations involved [46]. In this case, the variational pdf $Q(\Xi^{(k)})$ could be further restricted to be a Dirac delta function $Q(\Xi^{(k)}) = \delta(\Xi^{(k)} - \hat{\Xi}^{(k)})$, where $\hat{\Xi}^{(k)}$ is the point estimate of the parameter $\Xi^{(k)}$. After substituting this functional form into problem (11), the optimal point estimate $\hat{\Xi}^{(k)*}$ is obtained by [32, page 164]:

$$\hat{\Xi}^{(k)*} = \arg \max_{\prod_{j \neq k} Q(\Theta_j)} \mathbb{E}_{\prod_{j \neq k} Q(\Theta_j)} [\ln p(\Theta, \mathcal{Y})]. \quad (13)$$

This is indeed the framework of variational expectation maximization (EM), in which the factor matrices $\{\Xi^{(k)}\}_{k=1}^N$ are treated as global parameters and other variables are treated as latent variables.

In (12) and (13), the log of the joint pdf $\ln p(\Theta, \mathcal{Y})$ needs to be evaluated. If the Gaussian likelihood function (9) is adopted, it is expressed as

$$\begin{aligned} \ln p^\dagger(\Theta, \mathcal{Y}) &= \sum_{n=1}^N \ln \left(\text{U}(\Xi^{(n)} \geq \mathbf{0}_{J_n \times L}) \right) + \frac{\prod_{n=1}^N J_n}{2} \ln \beta \\ &\quad - \frac{\beta}{2} \|\mathcal{Y} - [\Xi^{(1)}, \Xi^{(2)}, \dots, \Xi^{(N)}]\|_F^2 + \sum_{n=1}^N \frac{J_n}{2} \sum_{l=1}^L \ln \gamma_l \\ &\quad - \sum_{n=1}^N \frac{1}{2} \text{Tr} \left(\Xi^{(n)} \Gamma \Xi^{(n)T} \right) + \sum_{l=1}^L [(10^{-6} - 1) \ln \gamma_l - 10^{-6} \gamma_l] \end{aligned}$$

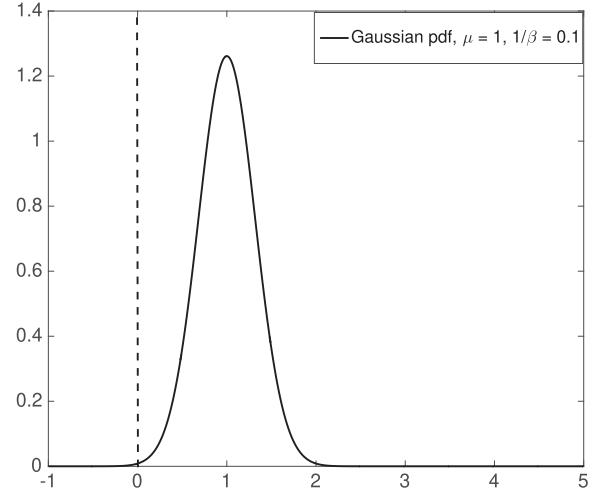


Fig. 3. Illustration of a univariate Gaussian probability density function with its mean much larger than the variance.

$$+ (10^{-6} - 1) \ln \beta - 10^{-6} \beta + \text{const}, \quad (14)$$

where the term $\Gamma = \text{diag}\{\gamma_1, \gamma_2, \dots, \gamma_L\}$. On the other hand, if the truncated Gaussian likelihood (10) is used, the log of the joint pdf $\ln p^\sharp(\Theta, \mathcal{Y})$ takes the following form

$$\ln p^\sharp(\Theta, \mathcal{Y}) = \ln p^\dagger(\Theta, \mathcal{Y}) - \ln \Phi\left(\{\Xi^{(n)}\}_{n=1}^N, \beta\right) + \text{const}. \quad (15)$$

where

$$\begin{aligned} \Phi\left(\{\Xi^{(n)}\}_{n=1}^N, \beta\right) &= \int_0^\infty \left(\frac{\beta}{2\pi} \right)^{\prod_{n=1}^N \frac{J_n}{2}} \\ &\quad \times \exp\left(-\frac{\beta}{2} \|\mathcal{Y} - [\Xi^{(1)}, \Xi^{(2)}, \dots, \Xi^{(N)}]\|_F^2\right) d\mathcal{Y}. \end{aligned} \quad (16)$$

In (15), the term $\ln \Phi(\{\Xi^{(n)}\}_{n=1}^N, \beta)$, which arises from the truncated Gaussian likelihood in (10), is very difficult to evaluate and differentiate, due to the multiple integrations involved. Fortunately, for most applications in Bayesian non-negative matrix/tensor decomposition [38]–[40], the confidence of the low-rank matrix/tensor model is relatively high, in the sense that the noise power $1/\beta$ is small compared to average element in signal tensor $[\Xi^{(1)}, \Xi^{(2)}, \dots, \Xi^{(N)}]$. Under this assumption, it is easy to see $\ln \Phi(\{\Xi^{(n)}\}_{n=1}^N, \beta) \approx \ln 1 = 0$, since Gaussian pdf $p(\mathcal{Y}) = \left(\frac{\beta}{2\pi}\right)^{\prod_{n=1}^N \frac{J_n}{2}} \exp(-\beta \|\mathcal{Y} - [\Xi^{(1)}, \Xi^{(2)}, \dots, \Xi^{(N)}]\|_F^2) d\mathcal{Y}$ decays very rapidly and thus most densities are over the region $\mathcal{Y} \geq 0$. As an illustration, a univariate Gaussian pdf with its mean much larger than the variance is plotted in Fig. 3, in which the probability density in the negative region is negligible. This suggests that the log of the joint pdf $\ln p^\sharp(\Theta, \mathcal{Y})$ in (15) can be well approximated by $\ln p^\dagger(\Theta, \mathcal{Y})$ in (14), and therefore algorithm derivations are unified for the two likelihoods. This also explains why the previous Bayesian nonnegative matrix/tensor decompositions [38]–[40] all employ the Gaussian likelihood function.

A. Derivation for Variational Pdfs

As discussed in the first paragraph of this section, the mean-field approximation is employed to enable closed-form expression for each variational pdf. For the precision parameter γ_l , by substituting (14) into (12) and only taking the terms relevant to γ_l , the variational pdf $Q(\gamma_l)$ can be found to take the same functional form as that of the gamma distribution, i.e., $Q(\gamma_l) = \text{gamma}(\gamma_l | c_l, d_l)$ with

$$c_l = \sum_{n=1}^N \frac{J_n}{2} + \epsilon, \quad (17)$$

$$d_l = \sum_{n=1}^N \frac{1}{2} \mathbb{E}_{Q(\Xi^{(n)})} \left[\Xi_{:,l}^{(n)T} \Xi_{:,l}^{(n)} \right] + \epsilon. \quad (18)$$

Since the variational pdf $Q(\gamma_l)$ is determined by parameters c_l and d_l , its update is equivalent to the update of the two parameters in (17) and (18).

Similarly, using (12) and (14), the variational pdf $Q(\beta)$ can be found to be a gamma distribution $Q(\beta) = \text{gamma}(\beta | e, f)$, where

$$e = \frac{\prod_{n=1}^N J_n}{2} + \epsilon, \quad (19)$$

$$f = \frac{1}{2} \mathbb{E}_{\prod_{\Theta_j \neq \beta} Q(\Theta_j)} \left[\|\mathcal{Y} - [\Xi^{(1)}, \Xi^{(2)}, \dots, \Xi^{(N)}]\|_F^2 \right] + \epsilon. \quad (20)$$

On the other hand, by substituting (14) into (13), the point estimate of $\hat{\Xi}^{(k)}$ can be obtained via solving the following problem:

$$\begin{aligned} \max \mathbb{E}_{\prod_{\Theta_j \neq \Xi^{(k)}} Q(\Theta_j)} & \left[\ln \left(\mathbb{U}(\Xi^{(k)} \geq \mathbf{0}_{J_k \times L}) \right) \right. \\ & \left. - \frac{\beta}{2} \|\mathcal{Y} - [\Xi^{(1)}, \Xi^{(2)}, \dots, \Xi^{(N)}]\|_F^2 - \frac{1}{2} \text{Tr} \left(\Xi^{(k)T} \Gamma \Xi^{(k)} \right) \right]. \end{aligned} \quad (21)$$

After distributing the expectations, expanding the Frobenius norm, and utilizing the fact that $\ln(0) = -\infty$, problem (21) can be equivalently shown to be:

$$\begin{aligned} \min f(\Xi^{(k)}) \\ \text{s.t. } \Xi^{(k)} \geq \mathbf{0}_{J_k \times L}, \end{aligned} \quad (22)$$

where

$$\begin{aligned} f(\Xi^{(k)}) &= \frac{1}{2} \text{Tr} \left(\Xi^{(k)T} \mathbb{E}_{\prod_{\Theta_j \neq \Xi^{(k)}} Q(\Theta_j)} \left[\beta \mathfrak{B}^{(k)T} \mathfrak{B}^{(k)} + \Gamma \right] \Xi^{(k)} \right) \\ & \quad - 2\beta \Xi^{(k)T} \mathbb{E}_{\prod_{\Theta_j \neq \Xi^{(k)}} Q(\Theta_j)} \left[\mathfrak{B}^{(k)} \right] \mathcal{Y}^{(k)T}. \end{aligned} \quad (23)$$

In (23), the term $\mathfrak{B}^{(k)} = (\diamond_{n=1, n \neq k}^N \Xi^{(n)})^T$, with the multiple Khatri-Rao products $\diamond_{n=1, n \neq k}^N \mathbf{A}^{(n)} = \mathbf{A}^{(N)} \diamond \mathbf{A}^{(N-1)} \diamond \dots \diamond \mathbf{A}^{(k+1)} \diamond \mathbf{A}^{(k-1)} \diamond \dots \diamond \mathbf{A}^{(1)}$. It is easy to see that problem (22)

is a quadratic programming (QP) problem with nonnegative constraints. Since each diagonal element γ_l in the diagonal matrix Γ is larger than zero, the Hessian matrix of the function $f(\Xi^{(k)})$, with the expression being

$$\mathbf{H}^{(k)} = \mathbb{E}_{\prod_{\Theta_j \neq \Xi^{(k)}} Q(\Theta_j)} \left[\beta \mathfrak{B}^{(k)T} \mathfrak{B}^{(k)} + \Gamma \right] \quad (24)$$

is positive definite. This implies that problem (22) is a convex problem, and its solutions have been investigated for decades [42, Chapter 2 and 4]. In particular, first-order methods have recently received much attention due to their scalability in big data applications. Within the class of first-order methods, a simple gradient projection method is introduced as follows.

In each iteration of the gradient projection method, the update equation is of the form [42, page 224]:

$$\Xi^{(k,t+1)} = \left[\Xi^{(k,t)} - \alpha_t \nabla f(\Xi^{(k,t)}) \right]_+, \quad (25)$$

where the gradient $\nabla f(\Xi^{(k,t)})$ is computed as

$$\begin{aligned} \nabla f(\Xi^{(k,t)}) &= \Xi^{(k,t)} \mathbb{E}_{\prod_{\Theta_j \neq \Xi^{(k)}} Q(\Theta_j)} \left[\beta \mathfrak{B}^{(k)T} \mathfrak{B}^{(k)} + \Gamma \right] \\ & \quad - \mathcal{Y}^{(k)} \mathbb{E}_{\prod_{\Theta_j \neq \Xi^{(k)}} Q(\Theta_j)} \left[\beta \mathfrak{B}^{(k)T} \right]. \end{aligned} \quad (26)$$

In (26), the symbol $[\cdot]_+$ denotes projecting each element of $\Xi^{(k)}$ to $[0, \infty)$ (i.e., $[x]_+ = x$ if $x \geq 0$ and $[x]_+ = 0$ otherwise) and $\alpha_t \geq 0$ is the step size. During the inference, due to the sparsity-promoting property of the nonnegative Gaussian-gamma prior, some of the precision parameters will go to very large numbers while some of them will tend to be zero. This will result in a very large condition number of the Hessian matrix $\mathbf{H}^{(k)}$. In this case, applying the diminishing rule² to the step size α_t still enjoys a good convergence performance [42, page 228] and thus is set as the default step-size rule in the proposed algorithm.

B. Summary of the Inference Algorithm

From equations (17)–(26), it can be seen that we need to compute various expectations. In particular, for expectations $\mathbb{E}_{Q(\Xi^{(n)})}[\Xi^{(n)}]$, $\mathbb{E}_{Q(\gamma_l)}[\gamma_l]$ and $\mathbb{E}_{Q(\beta)}[\beta]$, their computations are very straightforward, i.e., $\mathbb{E}_{Q(\Xi^{(n)})}[\Xi^{(n)}] = \hat{\Xi}^{(n)}$, $\mathbb{E}_{Q(\gamma_l)}[\gamma_l] = \frac{c_l}{d_l}$ and $\mathbb{E}_{Q(\beta)}[\beta] = \frac{e}{f}$. However, when updating $\hat{\Xi}^{(n)}$ using (22) and (23), there is one complicated expectation $\mathbb{E}_{\prod_{\Theta_j \neq \Xi^{(k)}} Q(\Theta_j)}[\mathfrak{B}^{(k)T} \mathfrak{B}^{(k)}]$. Fortunately, it can be shown that $\mathbb{E}_{\prod_{\Theta_j \neq \Xi^{(k)}} Q(\Theta_j)}[\mathfrak{B}^{(k)T} \mathfrak{B}^{(k)}] = \odot_{n=1, n \neq k}^N \hat{\Xi}^{(n)T} \hat{\Xi}^{(n)}$, where the multiple Hadamard products $\odot_{n=1, n \neq k}^N \mathbf{A}^{(n)} = \mathbf{A}^{(N)} \odot \mathbf{A}^{(N-1)} \odot \dots \odot \mathbf{A}^{(k+1)} \odot \mathbf{A}^{(k-1)} \odot \dots \odot \mathbf{A}^{(1)}$. Since the computation of one variational pdf needs the statistics of other variational pdfs, alternating update is needed, resulting in the iterative algorithm summarized in Algorithm 1.

²In the diminishing rule [42, page 227], the step size α_t needs to satisfy $\alpha_t \rightarrow 0$ and $\sum_{t=0}^{\infty} \alpha_t = \infty$.

Algorithm 1: Probabilistic Tensor CPD With Nonnegative Factors.

Initializations: Choose $L > R$ and initial values

$$\{\hat{\Xi}^{(n,0)}\}_{n=1}^N, \epsilon.$$

Iterations: For the s th iteration ($s \geq 0$),
Update the parameter of $Q(\Xi^{(k)})^{(s+1)}$

Set initial value $\Xi^{(k,0)} = \hat{\Xi}^{(k,0)}$.

Iterations: For the t th iteration ($t \geq 0$), compute

$$\Xi^{(k,t+1)} = \left[\Xi^{(k,t)} - \alpha_t \nabla f(\Xi^{(k,t)}) \right]_+$$

where

$$\begin{aligned} \nabla f(\Xi^{(k,t)}) &= \Xi^{(k,t)} \left[\frac{e^s}{f^s} \bigodot_{n=1, n \neq k}^N \hat{\Xi}^{(n,s)T} \hat{\Xi}^{(n,s)} \right. \\ &\quad \left. + \text{diag} \left\{ \frac{c_1^s}{d_1^s}, \dots, \frac{c_L^s}{d_L^s} \right\} \right] - \frac{e^s}{f^s} \mathcal{Y}^{(k)} \left(\bigodot_{n=1, n \neq k}^N \hat{\Xi}^{(n,s)} \right) \end{aligned}$$

and α_t is chosen by the diminishing rule [42, page 227].

Until Convergence

Set $\hat{\Xi}^{(k,s+1)} = \Xi^{(k,t+1)}$.

Update the parameter of $Q(\gamma_l)^{s+1}$

$$c_l^{s+1} = \sum_{n=1}^N \frac{J_n}{2} + \epsilon$$

$$d_l^{s+1} = \sum_{n=1}^N \frac{1}{2} \hat{\Xi}_{:,l}^{(n,s+1)T} \hat{\Xi}_{:,l}^{(n,s+1)} + \epsilon$$

Update the parameter of $Q(\beta)^{s+1}$

$$e^{s+1} = \epsilon + \frac{\prod_{n=1}^N J_n}{2}$$

$$f^{s+1} = \epsilon + \frac{1}{2} \|\mathcal{Y} - [\hat{\Xi}^{(1,s+1)}, \hat{\Xi}^{(2,s+1)}, \dots, \hat{\Xi}^{(N,s+1)}]\|_F^2$$

Until Convergence

C. Discussions and Insights

To gain further insight from the proposed inference algorithm, discussions of its convergence property, automatic rank determination, relationship to the NALS algorithm, computational complexity, and scalability improvement are presented in the following.

1) Convergence Property: The proposed algorithm is derived under the framework of mean-field variational inference, where a variational pdf $Q(\Theta) = \prod_k Q(\Theta_k)$ is sought that minimizes the KL divergence $\text{KL}(Q(\Theta)||p(\Theta|\mathcal{Y}))$. Even though this problem is known to be non-convex due to the non-convexity of the mean-field family set, it is convex with respect to a single variational pdf $Q(\Theta_k)$ after fixing other variational pdfs $\{Q(\Theta_j), j \neq k\}$ [32, page 138]. Therefore, the iterative algorithm, in which a single variational pdf is optimized in each iteration with other variational pdfs fixed, is essentially a coordinate descent algorithm in the functional space of variational pdfs. Since the subproblem in each iteration is not only convex but also has a unique solution, the limit point generated by the

coordinate descent steps over the functional space of variational pdfs is guaranteed to be at least a stationary point of the KL divergence [32, page 163].

2) Automatic Rank Determination: During the inference, the expectations of some precision parameters $\{\gamma_l\}$, i.e., $\{\frac{c_l^s}{d_l^s}\}$ will go to a very large value. It indicates that the corresponding columns in the factor matrices are close to zero vectors, thus playing no role in data interpretation. As a result, after convergence, those columns can be pruned out and the number of remaining columns in each factor matrix gives the estimate of tensor rank.

In practice, to reduce the computational complexity, the pruning would be executed during the iteration. In particular, in each iteration, after the precision estimates $\{\frac{c_l^s}{d_l^s}\}$ exceed a certain threshold (e.g., 10^6), the associated columns are safely pruned out. After every pruning, it is equivalent to starting minimization of the KL divergence of a new (but smaller) probabilistic model, with the current variational distributions acting as the initialization of the new minimization. Therefore, the pruning steps will not affect the convergence, and are widely used in recent related works [23]–[30], [49]–[51].

Usually, the hyper-parameters $\{c_l^0, d_l^0\}$ of the prior gamma distribution $\gamma_l | c_l^0, d_l^0$ are set to be a very small number $\epsilon = 10^{-6}$ to approach a non-informative prior. Otherwise, their values might affect the behavior of tensor rank estimate. For example, if we prefer a high value of the tensor rank, the initial value d_l^0 can be set to be very large while the initial value c_l^0 can be set to be small, so that the update of $\frac{c_l}{d_l}$ can be steered towards a small value in order to promote a high tensor rank. However, how to set the hyper-parameters to accurately control the degree of low-rank is challenging, and deserves future investigation.

3) Relationship to NALS Algorithm: The mean-field variational inference for tensor CPD problem could be interpreted as alternating optimizations over the Riemannian space (in which the Euclidean space is a special case). This insight has been revealed in previous works [25], [45], and can also be found in the proposed algorithm above. For example, for the precision parameters and the noise power parameter, the variational pdfs are with no constraint on the functional form, and thus the corresponding alternating optimization is over the Riemannian space due to the exponentially conjugacy property of the proposed probabilistic model [25], [45]. On the other hand, for unknown factor matrices, since the variational pdfs to be optimized are with a delta functional form, the corresponding alternating optimization is over the Euclidean space, thus is similar the conventional NALS step. However, there is a significant difference. In the proposed algorithm, there is a shrinkage term Γ appeared in the Hessian matrix in (24), and Γ will be updated together with other parameters in the algorithm. This intricate interaction is due to the employed Bayesian framework, and cannot be revealed by NALS framework. Consequently, the proposed algorithm is a generalization of the NALS algorithm, with the additional novel feature in automatic rank determination achieved via optimization in Riemannian space.

4) Computational Complexity: For each iteration, the computational complexity is dominated by computing the gradient of each factor matrix in (25), costing $O(\prod_{n=1}^N J_n L)$. From

this expression, it is clear that the computational complexity in each iteration is linear with respect to the tensor dimension product $\prod_{n=1}^N J_n$. Consequently, the complexity of the algorithm is $O(q(\prod_{n=1}^N J_n L))$ where q is the iteration number at convergence.

5) *Speeding up the Algorithm via Acceleration Schemes and Parallel Computations:* From the proposed inference algorithm, it is clear that the bottleneck of the algorithm efficiency is the update of factor matrices $\{\Xi^{(n)}\}_{n=1}^N$ via solving problem (22). Fortunately, if the problem is well conditioned, in the sense that the condition number of the Hessian matrix $\mathbf{H}^{(k)}$ in (24) is smaller than a threshold (e.g., 100), acceleration schemes, including variants of the Nesterov scheme [44, page 81 and page 90], [18], can be utilized to significantly reduce the required number of iteration for solving problem (22), thus speeding up the proposed algorithm. Besides reducing the iteration number for convergence, ideas of leveraging parallel computing architecture like message passing interface (MPI) in [18], [19] and super-computer in [43] deserve future investigation. For instance, it is easy to see that the non-negatively constrained quadratic problem (22) is separable across the rows of the factor matrix $\Xi^{(n)}$, making it possible to optimize multiple rows of the factor matrices in parallel. More sophisticated schemes on parallel computations could be found in [18], [19], [43].

V. NUMERICAL RESULTS

In this section, numerical results using synthetic data are firstly presented to assess the performance of the proposed algorithm in terms of convergence property, factor matrix recovery, tensor rank estimation and running time. Next, the proposed algorithm is utilized to analyze two real-world data sets (the amino acids fluorescence data and the ENRON email corpus), for demonstration on blind source separation and social group clustering. For all the simulated algorithms, the initial factor matrix $\hat{\Xi}^{(k,0)}$ is set as the singular value decomposition (SVD) approximation $\mathbf{U}_{:,1:L}(\mathbf{S}_{1:L,1:L})^{\frac{1}{2}}$ where $[\mathbf{U}, \mathbf{S}, \mathbf{V}] = \text{SVD}[\mathcal{Y}^{(k)}]$ and $L = \min\{J_1, J_2, \dots, J_N\}$. The parameter ϵ is set to be 10^{-6} . The algorithms are deemed to be converged when $\|\hat{\Xi}^{(1,s+1)}, \hat{\Xi}^{(2,s+1)}, \dots, \hat{\Xi}^{(N,s+1)}\| - \|\hat{\Xi}^{(1,s)}, \hat{\Xi}^{(2,s)}, \dots, \hat{\Xi}^{(N,s)}\|_F^2 < 10^{-6}$. All experiments were conducted in Matlab R2015b with an Intel Core i7 CPU at 2.2 GHz.

A. Validation on Synthetic Data

A three dimensional tensor $\mathcal{X} = [\mathbf{M}^{(1)}, \mathbf{M}^{(2)}, \mathbf{M}^{(3)}] \in \mathbb{R}^{100 \times 100 \times 100}$ with rank $R = 10$ is considered as the noise-free data tensor. Each element in factor matrix $\mathbf{M}^{(n)}$ is independently drawn from a uniform distribution over $[0,1]$ and thus is nonnegative. On the other hand, two observation data tensors are considered: 1). the data \mathcal{X} is corrupted by a noise tensor $\mathcal{W} \in \mathbb{R}^{100 \times 100 \times 100}$, i.e., $\mathcal{Y} = \mathcal{X} + \mathcal{W}$, with each element of noise tensor \mathcal{W} being independently drawn from a zero-mean Gaussian distribution with variance σ_w^2 , and this corresponds to the Gaussian likelihood model (9); 2). the data \mathcal{Y}^+ is obtained by setting the negative elements of \mathcal{Y} to

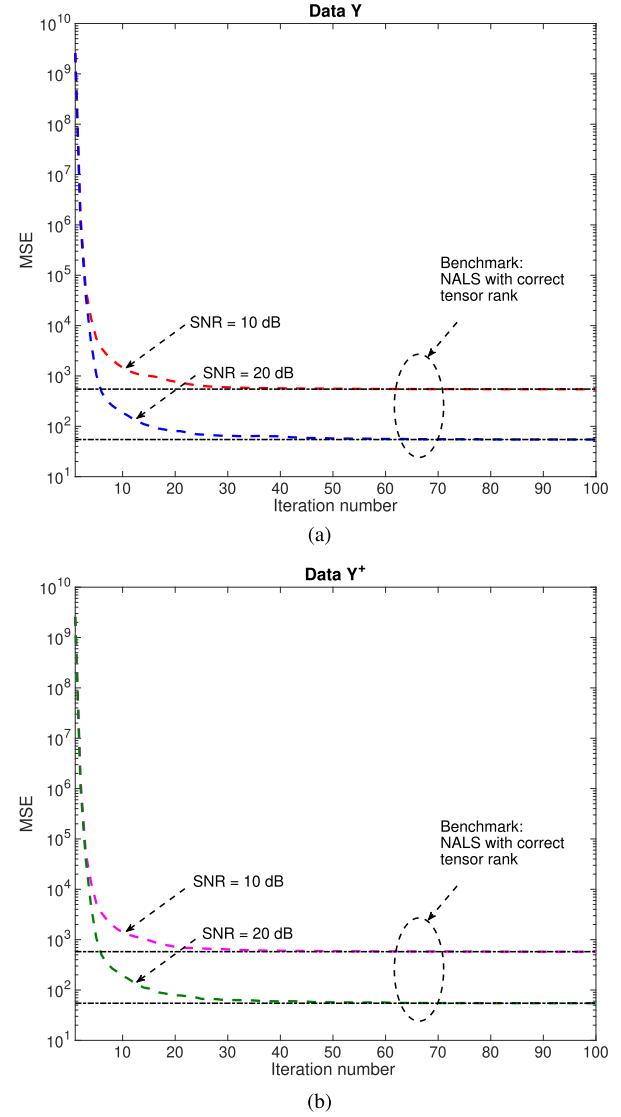


Fig. 4. Convergence of the proposed algorithm for different test data.

zero, i.e., $\mathcal{Y}_{i_1, i_2, i_3}^+ = \mathcal{Y}_{i_1, i_2, i_3} \mathbf{U}(\mathcal{Y}_{i_1, i_2, i_3} \geq 0)$, and the truncated Gaussian likelihood model (10) is employed to fit these data. The SNR is defined as $10 \log_{10}(\|\mathcal{X}\|_F^2 / \mathbb{E}_{p(\mathcal{W})}[\|\mathcal{W}\|_F^2]) = 10 \log_{10}(\|\mathcal{X}\|_F^2 / (100^3 \sigma_w^2))$. For the proposed algorithm, the step size sequence is chosen as $\alpha_t = 10^{-3}/(t+1)$ [42], and the gradient projection update is terminated when $\|f(\Xi^{(k,t)}) - f(\Xi^{(k,t-1)})\|_F \leq 10^{-3}$. Each result in this subsection is obtained by averaging 100 Monte-Carlo runs.

Fig. 4 presents the convergence performances of the proposed algorithm under different SNRs and different test data, where the mean-square-error (MSE) $\|\hat{\Xi}^{(1,s)}, \hat{\Xi}^{(2,s)}, \hat{\Xi}^{(3,s)}\| - \mathcal{X}\|_F^2$ is chosen as the assessment criterion. From Fig. 4(a), it can be seen that for test data \mathcal{Y} , the MSEs of the proposed algorithm, which assumes no knowledge of tensor rank, decrease significantly in the first few iterations and converge to the MSE of the NALS algorithm [16] (with exact tensor rank) under SNR = 10 dB and SNR = 20 dB. Similar convergence performances can be observed for the test data \mathcal{Y}^+ . This is of no surprise because

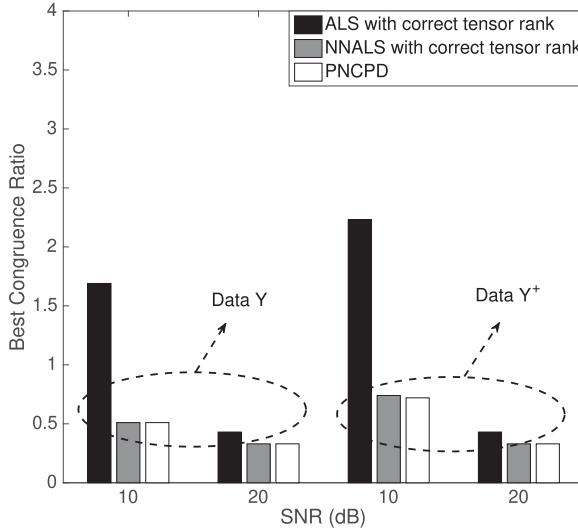


Fig. 5. Best congruence ratios of the proposed algorithm for different test data.

approximating (15) by (14) dose not make any changes on the algorithm framework of variational inference, and thus the excellent convergence performance of the proposed algorithm is expected (as discussed in Section IV-C).

The MSE measures the performance of low-rank tensor recovery. However, due to the uniqueness property of tensor CPD [1], each factor matrix can be recovered up to an unknown permutation and scaling ambiguity. To directly assess the accuracies of factor matrices recovery, the best congruence ratio (BCR), which involves computing the MSE between the true factor matrix $\mathbf{M}^{(k)}$ and the estimated factor matrix $\hat{\mathbf{\Xi}}^{(k)}$, is widely used as the assessment criterion. It is defined as

$$\sum_{k=1}^3 \min_{\Delta^{(k)}, \mathbf{P}^{(k)}} \frac{\|\mathbf{M}^{(k)} - \hat{\mathbf{\Xi}}^{(k)} \mathbf{P}^{(k)} \Delta^{(k)}\|_F}{\|\mathbf{M}^{(k)}\|_F},$$

where the diagonal matrix $\Delta^{(k)}$ and the permutation matrix $\mathbf{P}^{(k)}$ are found via the greedy least-squares column matching algorithm [48]. From Fig. 5, it is seen that both the proposed algorithm (labeled as PNCPD) and the NALS algorithm (with exact tensor rank) achieve much better factor matrix recovery than the ALS algorithm (with exact tensor rank) [1]. This shows the importance of incorporating the nonnegative constraint into the algorithm design. Furthermore, the factor matrix recovery performances of the proposed algorithm under test data \mathcal{Y} and \mathcal{Y}^+ are indistinguishable under $\text{SNR} = 20$ dB. This shows that when SNR is high, equation (14) gives a quite good approximation to equation (15), thus leading to remarkably accurate factor matrices recovery. Although the BCR of the proposed algorithm is higher for the data \mathcal{Y}^+ than that for the data \mathcal{Y} , it is with nearly the same performance as that of the NALS algorithm (with exact tensor rank).

On the other hand, the tensor rank estimates of the proposed algorithm under different SNRs are presented in Fig. 6, with each vertical bar showing the mean and the error bars showing the standard derivation of tensor rank estimates. The blue horizontal

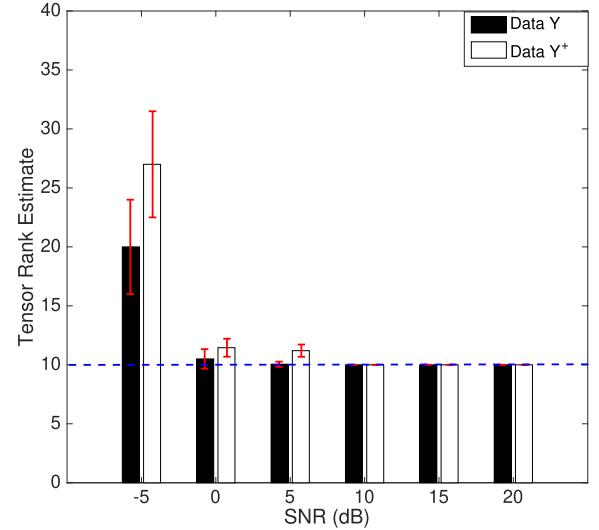


Fig. 6. Tensor rank estimates of the proposed algorithm for different test data.

TABLE I
PERFORMANCE OF TENSOR RANK ESTIMATION VERSUS DIFFERENT TRUE TENSOR RANKS FOR TENSOR DATA \mathcal{Y} WHEN $\text{SNR} = 20$ dB

True tensor rank	Mean of tensor rank estimates	Standard derivation of tensor rank estimates	Percentage of correct tensor rank estimates
10	10	0	100%
30	29.6	1.27	90%
50	28.15	18.29	25%

dashed line indicates the true tensor rank. From Fig. 6, it is seen that the proposed algorithm recovers the true tensor rank with 100% accuracy for a wide range of SNRs, in particular when SNR is larger than 10 dB. Even though the performance is not 100% accurate when SNR is 0 dB and 5 dB, the estimated tensor rank is still close to the true tensor rank with a high probability for test data \mathcal{Y} . However, under these two low SNRs, the rank estimation performances of the proposed algorithm for the data \mathcal{Y}^+ degrade significantly. This is because equation (15) cannot be well approximated by equation (14) under very low SNRs. Furthermore, the proposed algorithm fails to give correct rank estimates when SNR is lower than -5 dB for both two test data sets, since the noise with very large power masks the low-rank structure of the data.

To assess the tensor rank estimation accuracy when the tensor is with a larger true rank, we apply the the proposed algorithm to the tensor data \mathcal{Y} with the true rank $R = \{10, 30, 50\}$ and $\text{SNR} = 20$ dB. The rank estimation performance is presented in Table I. From Table I, it can be seen that the proposed algorithm recovers the rank accurately when the true rank is 10 and 30. However, when $R = 50$, the proposed algorithm fails to accurately recover the tensor rank. This could be explained by the fact that the Gaussian-gamma prior would lead to the sparsest estimation result [52], and thus fail to work well when the true rank is high. This has also been observed in a recent matrix decomposition work [53], in which a Gaussian-Wishart prior has been employed to tackle the high-rank estimation challenge. However, employing the Gaussian-Wishart prior for tensor decompositions is challenging as Wishart distribution is

TABLE II
PERFORMANCE OF TENSOR RANK ESTIMATION WHEN THE COLUMNS IN ONE FACTOR MATRIX ARE CORRELATED AND SNR = 20 dB

s	0	1	3	5	100
Percentage of correct tensor rank estimates	100%	100%	100%	25%	5%

TABLE III
PERFORMANCE OF TENSOR RANK ESTIMATION WHEN THE COLUMNS IN ALL FACTOR MATRICES ARE CORRELATED AND SNR = 20 dB

s	0	1	3	5	100
Percentage of correct tensor rank estimates	100%	40%	0%	0%	0%

inherently defined for a matrix [53]. Thus, high rank tensor decomposition, which is both important and interesting, would be left as future work.

The simulation results presented so far are for well-conditioned tensors, i.e., the columns in each of the factor matrices are independently generated. In order to fully assess the rank learning ability of the proposed algorithm, we consider another noise-free three dimensional tensor $\bar{\mathcal{X}} = [\bar{\mathbf{M}}^{(1)}, \bar{\mathbf{M}}^{(2)}, \bar{\mathbf{M}}^{(3)}] \in \mathbb{R}^{100 \times 100 \times 100}$ with rank $R = 10$. The factor matrix is set as $\bar{\mathbf{M}}^{(1)} = 0.11_{100 \times 10} + 2^{-s} \mathbf{M}^{(1)}$, and each element in factor matrices $\{\bar{\mathbf{M}}^{(n)}\}_{n=1}^3$ is independently drawn from a uniform distribution over $[0,1]$. According to the definition of the tensor condition number [54], [55], when s increases, the correlation among the columns in the factor matrix $\bar{\mathbf{M}}^{(1)}$ increases, and the tensor condition number becomes larger. In particular, when s goes to infinity, the condition number of the tensor goes to infinity too. We apply the proposed algorithm to $\bar{\mathcal{X}}$ corrupted with noise: $\bar{\mathcal{Y}} = \bar{\mathcal{X}} + \bar{\mathcal{W}}$, where each element of noise tensor $\bar{\mathcal{W}}$ is independently drawn from a zero-mean Gaussian distribution with variance $\bar{\sigma}_w^2$. Table II shows the rank estimation accuracy of the proposed algorithm when SNR = 20 dB. It can be seen that the proposed algorithm can correctly estimate the tensor rank when $s < 5$. But as the tensor conditional number increases (i.e., the columns are more correlated in the factor matrix $\bar{\mathbf{M}}^{(1)}$), the tensor rank estimation performance decreases significantly.

Next, we consider an extreme case in which the columns in all factor matrices are highly correlated: $\tilde{\mathcal{X}} = [\bar{\mathbf{M}}^{(1)}, \bar{\mathbf{M}}^{(2)}, \bar{\mathbf{M}}^{(3)}] \in \mathbb{R}^{100 \times 100 \times 100}$ with rank $R = 10$, where each factor matrix $\bar{\mathbf{M}}^{(n)} = 0.11_{100 \times 10} + 2^{-s} \mathbf{M}^{(n)}$, and each element in factor matrices $\{\bar{\mathbf{M}}^{(n)}\}_{n=1}^3$ is independently drawn from a uniform distribution over $[0,1]$. With the same observation data model as $\bar{\mathcal{Y}}$ as before and when SNR = 20 dB, the percentages of correct tensor rank estimate are shown in Table III. It can be seen that it is difficult for the proposed algorithm to accurately estimate the tensor rank even when $s = 1$.

Finally, as discussed in the Section IV-C, acceleration schemes could be incorporated to speed up the proposed algorithm. As discussed in Section IV-A, in the first few iterations, since some precision parameters are learnt to be very large

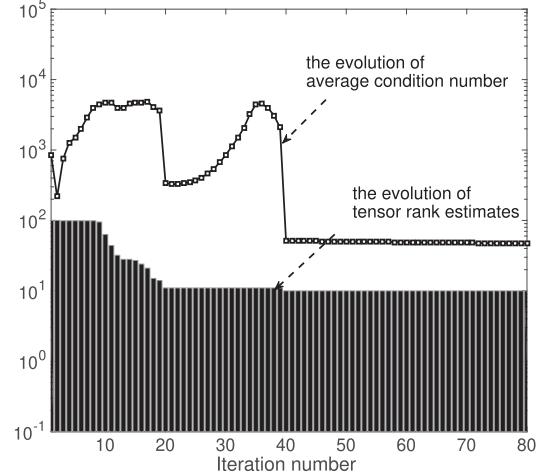


Fig. 7. The average condition number of the Hessian matrix (24) and the tensor rank estimate versus number of iterations for test data \mathcal{Y} when SNR = 20 dB.

TABLE IV
AVERAGE RUNNING TIME IN SECONDS OF THE PROPOSED ALGORITHM FOR DIFFERENT TEST DATA. THE ACCELERATED ALGORITHM IS LABELED AS PNCPD-A

Data	SNR = 10 dB		SNR = 20 dB	
	PNCPD	PNCPD-A	PNCPD	PNCPD-A
\mathcal{Y}	113.61	63.66	62.52	52.08
\mathcal{Y}^+	99.48	58.02	59.61	49.86

while some of them are with very small values, the average condition number of the Hessian matrix of problem (22), i.e., $\frac{1}{3} \sum_{k=1}^3 \text{condition_number}(\mathbf{H}^{(k)})$, is very large. After several iterations, the proposed algorithm has gradually recovered the tensor rank, and then the remaining precision parameters are with comparable values, leading to a well-conditioned Hessian matrix $\mathbf{H}^{(k)}$ of problem (22). These results can be observed in Fig. 7. Inspired by the pioneering work [18], the Nesterov scheme [44, page 90] is utilized for the problem (22) when the condition number of the Hessian matrix is smaller than 100. Consequently, even with the same MSE and BCR performances, the accelerated algorithm is much faster than the default version of the proposed algorithm³ as presented in Table IV. Besides the Nesterov scheme, other advances in first-order optimizations could be incorporated to further improve the scalability of the algorithm. However, this is not the main focus of this paper, and thus in the following, we only examine the performances of the default version of the proposed algorithm for real-world applications.

B. Fluorescence Data Analysis

In this subsection, the proposed algorithm is utilized to analyze the amino acids fluorescence data⁴ [44]. This data set consists of five laboratory-made samples. Each sample contains different amounts of tyrosine, tryptophan and phenylalanine

³The presented time for the accelerated scheme includes the time for computing the condition numbers.

⁴<http://www.models.life.ku.dk>

TABLE V
SOCIAL GROUPS WITH PEOPLE IN TOP 10 SCORES IN EACH GROUP FOR THE ENRON E-MAIL DATA USING THE PROPOSED ALGORITHMS

Legal	Government Affairs Executive
'Tana Jones (tana.jones) Employee Financial Trading Group ENA Legal', 'Sara Shackleton (sara.shackleton) Employee ENA Legal', 'Mark Taylor (mark.taylor) Manager Financial Trading Group ENA Legal', 'Stephanie Panus (stephanie.panus) Senior Legal Specialist ENA Legal', 'Marie Heard (marie.heard) Senior Legal Specialist ENA Legal', 'Mark Haedicke (mark.haedicke) Managing Director ENA Legal', 'Susan Bailey (susan.bailey) Legal Assistant ENA Legal', 'Louise Kitchen (louise.kitchen) President Enron Online', 'Kay Mann (kay.mann) Lawyer', 'Debra Perltinger (debra.perltinger) Legal Specialist ENA Legal'	'Jeff Dasovich (jeff.dasovich) Employee Government Relationship Executive', 'James Steffes (james.steffes) VP Government Affairs', 'Steven Kean (steven.kean) VP Chief of Staff', 'Richard Shapiro (richard.shapiro) VP Regulatory Affairs', 'David Delainey (david.delainey) CEO ENA and Enron Energy Services', 'Richard Sanders (richard.sanders) VP Enron Wholesale Services', 'Shelley Corman (shelley.corman) VP Regulatory Affairs', 'Margaret Carson (margaret.carson) Employee Corporate and Environmental Policy', 'Mark Haedicke (mark.haedicke) Managing Director ENA Legal', 'Vince Kaminski (vince.kaminski) Manager Risk Management Head'
Trading / Top Executive	Pipeline
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dissolved in phosphate buffered water. The samples were measured by fluorescence and were corrupted by Gaussian noise with power 0.1, resulting in $\text{SNR} = 0.16$ dB. The fluorescence excitation-emission measured (EEM) data collected is with size $5 \times 201 \times 61$, and should be representable with a CPD model with rank 3, since there are three different types of amino acid and each individual amino acid gives a rank-one CPD component.

The proposed PNCPD algorithm was run to decompose the EEM tensor data with initial rank $L = 5$. For the proposed algorithm, the step size sequence is chosen as $\alpha_t = 10^{-2}/(t + 1)$ [42], and the gradient projection update is terminated when the gradient norm is smaller than 10^{-3} . At convergence, the proposed algorithm identified the correct tensor rank⁵ $R = 3$. Furthermore, the emission spectra and the excited spectra of three amino acids, which are obtained from the decomposed factor matrices [44], are shown in Fig. 8, with the clean data spectra⁶ serving as the benchmark. From Fig. 8, it can be seen that the recovered spectra from the proposed algorithm are very close to the clean data spectra, with the MSE of the emission spectra estimation equals 1.51×10^{-4} per wavelength and the MSE of the excitation spectra estimation equals 1.08×10^{-4} per wavelength.

C. ENRON E-mail Data Mining

In this subsection, the ENRON Email corpus,⁷ [3] was analyzed. This data set is with size $184 \times 184 \times 44$, and contains e-mail communication records between 184 people within 44 months, in which each entry denotes the number of e-mail exchanged between two particular people within a particular month. Before fitting the data to the proposed algorithms, the same pre-processing as in [2], [3] is applied to the non-zero data to compress the dynamic range. Then, the proposed algorithm is utilized to fit the data into the proposed nonnegative CPD

⁵Notice that even if the initial tensor rank L is set as 20, which is much larger than the true tensor rank 3, the proposed algorithm can still recover the true tensor rank. This shows that the proposed method is not sensitive to the initial tensor rank value.

⁶The clean data spectra is obtained by decomposing the clean data [44] using the NALS algorithm with correct tensor rank $R = 3$.

⁷The original source of the data is from [3] and we greatly appreciate Prof. Vagelis Papalexakis for sharing the data with us.

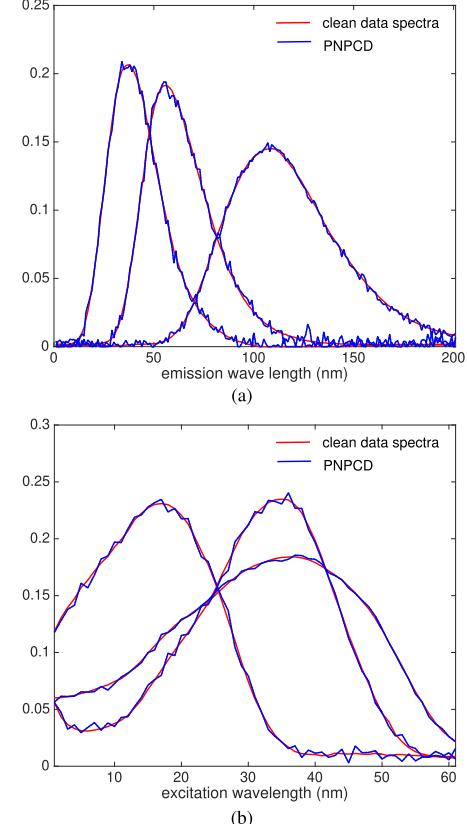


Fig. 8. The estimates of (a) emission spectra and (b) excitation spectra using the proposed algorithms, with the clean data spectra serving as the benchmark.

model, with the initial rank set as $L = 44$, the step size sequence being $\alpha_t = 1/(t + 1)$ [42], and the gradient projection update terminated when the gradient norm is smaller than 10^{-3} . As introduced in the Appendix A, the estimated tensor rank has the physical meaning of the number of underlying social groups, and each element in the first factor matrix can be interpreted as the score that a particular person belongs to a particular email sending group.

During the inference, the tensor rank estimate gradually reduces to the value 4, indicating that there are four underlying social groups. This is consistent with the results from [2], [3], which are obtained via trial-and-error experiments. After sorting

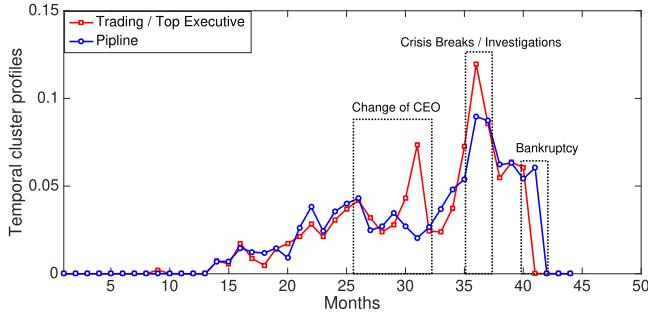


Fig. 9. Temporal cluster profiles (from the third factor matrix) for the ENRON e-mail dataset.

the scores of each column in the first factor matrix, the people with top 10 scores in each social group is shown in Table II. From the information of each person presented in Table II, the clustering results can be clearly interpreted. For instance, the people in the first group are either in legal department or lawyers, thus being clustered together. The clustering results of the proposed algorithms are also consistent with the results from [2], [3], which are obtained via nonlinear programming methods assuming the knowledge of tensor rank. Finally, interesting patterns can be observed from the temporal cluster profiles, which are obtained from the third factor matrix [2], [3], as illustrated in Fig. 9. It is clear that when the company has important events, such as the change of CEO, crisis breaks and bankruptcy, distinct peaks appear. Notice that in this example, all the data entries are nonnegative and the proposed algorithm still works well. This indicates that the tensor CPD with nonnegative factors is a model that matches this social clustering task.

VI. CONCLUSION

In this paper, probabilistic tensor CPD with nonnegative factors has been investigated under unknown tensor rank. In particular, the nonnegative Gaussian-gamma prior, which was shown to have both sparsity-promoting and exponential conjugacy properties, was introduced as the building block of the proposed probabilistic model. Then, an efficient inference algorithm was derived with an integrated feature of automatic rank determination. Extensive numerical results using both synthetic data and real-world data were presented to show the remarkable performance of the proposed algorithm.

APPENDIX

A. Motivation Examples

In this appendix, two motivating examples for probabilistic tensor CPD with nonnegative factors are presented.

1) *Motivating Example 1 (Fluorescence Spectroscopy)*: Fluorescence spectroscopy is a fast, simple and inexpensive method to determine the concentration of any solubilized sample based on its fluorescent properties, and is widely used in chemical, pharmaceutical and biomedical fields [34], [35]. In fluorescence spectroscopy, an excitation beam with a certain wavelength λ_i

passes through a solution in a cuvette. The excited chemical species in the sample will change their electronic states and then emit a beam of light, of which its spectrum is measured at the detector. Mathematically, let the concentration of the r th specie in the sample be c_r , and the excitation value at wavelength λ_i be $a_r(\lambda_i)$. Then, the noise-free measured spectrum intensity at the wavelength λ_j is $a_r(\lambda_i)b_r(\lambda_j)c_r$, where $b_r(\lambda_j)$ is the emission value of the r th species at the wavelength λ_j . If there are R different species in the sample, the noise-free fluorescence excitation-emission measured (EEM) data at λ_j is

$$p_{i,j} = \sum_{r=1}^R a_r(\lambda_i)b_r(\lambda_j)c_r. \quad (27)$$

Assume the excitation beam contains I wavelengths, and the noise-free EEM data is collected at J different wavelengths, an $I \times J$ data matrix is obtained as

$$\mathbf{P} = \sum_{r=1}^R \mathbf{A}_{:,r} \circ \mathbf{B}_{:,r} c_r, \quad (28)$$

where symbol \circ denotes vector outer product, $\mathbf{A}_{:,r} \in \mathbb{R}^{I \times 1}$ is a vector with the i th element being $a_r(\lambda_i)$ and $\mathbf{B}_{:,r} \in \mathbb{R}^{J \times 1}$ is a vector with the j th element being $b_r(\lambda_j)$. Assume $K > 1$ samples with the same chemical species but with different concentration of each specie are measured. Let the concentration of the r th specie in the k th sample be $c_{k,r}$, then after stacking the noise-free EEM data for each sample along a third dimension, a three dimensional (3D) tensor data $\mathcal{P} \in \mathbb{R}^{I \times J \times K}$ can be obtained as

$$\mathcal{P} = \sum_{r=1}^R \mathbf{A}_{:,r} \circ \mathbf{B}_{:,r} \circ \mathbf{C}_{:,r} \triangleq [\mathbf{A}, \mathbf{B}, \mathbf{C}], \quad (29)$$

where $\mathbf{C}_{:,r} \in \mathbb{R}^{K \times 1}$ is a vector with the k th element being $c_{k,r}$; matrices $\mathbf{A} \in \mathbb{R}^{I \times R}$, $\mathbf{B} \in \mathbb{R}^{J \times R}$ and $\mathbf{C} \in \mathbb{R}^{K \times R}$ are matrices with their r th columns being $\mathbf{A}_{:,r}$, $\mathbf{B}_{:,r}$ and $\mathbf{C}_{:,r}$, respectively.

It is easy to see that the noise-free data model in (29) yields exactly the tensor CPD model [1], and that is why CPD algorithms work very well for EEM data analysis [9]. More specifically, accounting for the possible Gaussian noise, the EEM data analysis aims to solve the following problem:

$$\begin{aligned} & \min_{\mathbf{A}, \mathbf{B}, \mathbf{C}} \|\mathcal{P} - [\mathbf{A}, \mathbf{B}, \mathbf{C}]\|_F^2 \\ & \text{s.t. } \mathbf{A} \geq \mathbf{0}_{I \times R}, \mathbf{B} \geq \mathbf{0}_{J \times R}, \mathbf{C} \geq \mathbf{0}_{K \times R}, \end{aligned} \quad (30)$$

where the nonnegative constraints are enforced due to the physical nature of elements in matrices \mathbf{A} , \mathbf{B} and \mathbf{C} as introduced above.

2) *Motivating Example 2 (Social Group Clustering)*: Social group clustering could be benefited by tensor data analysis, by which multiple views of social network are provided [2]–[4]. For example, consider a 3D email data set $\mathcal{Y} \in \mathbb{R}^{I \times J \times K}$ with each element $\mathcal{Y}(i, j, k)$ denoting the number of emails sent from person i to person j at the k th day. Each frontal slice $\mathcal{Y}(:, :, k)$ represents the connection intensity among different pair of peoples in the k th day, while each slice $\mathcal{Y}(:, j, :)$

shows the temporal evolution of the number of received mails for the person j from each of the other person in the data set. Consequently, decomposing the tensor \mathcal{Y} into latent CPD factors $\{\mathbf{A} \in \mathbb{R}^{I \times R}, \mathbf{B} \in \mathbb{R}^{J \times R}, \mathbf{C} \in \mathbb{R}^{K \times R}\}$ reveals different clustering groups from different views (i.e., different tensor dimensions). In particular, using the unfolding property of tensor CPD [1], we have

$$\mathcal{Y}^{(1)} = (\mathbf{C} \diamond \mathbf{B}) \mathbf{A}^T, \quad (31)$$

$$\mathcal{Y}^{(3)} = (\mathbf{B} \diamond \mathbf{A}) \mathbf{C}^T, \quad (32)$$

where $\mathcal{Y}^{(k)}$ is a matrix obtained by unfolding the tensor \mathcal{Y} along its k th dimension [1], and symbol \diamond denotes the Khatri-Rao product (i.e., column-wise Kronecker product). From (31), each column vector $\mathcal{Y}^{(1)}(:, i) \in \mathbb{R}^{JK \times 1}$ can be written as $\mathcal{Y}^{(1)}(:, i) = \sum_{r=1}^R (\mathbf{C} \diamond \mathbf{B})_{:, r} \mathbf{A}_{i, r}$, which is a linear combination of column vectors in matrix $(\mathbf{C} \diamond \mathbf{B}) \in \mathbb{R}^{JK \times R}$ with coefficients $\{\mathbf{A}_{i, r}\}_{r=1}^R$, and it represents the email sending pattern of person i . Thus, each column vector in matrix $\mathbf{C} \diamond \mathbf{B}$ can be interpreted as one of the R underlying email sending patterns, and $\mathbf{A}_{i, r}$ is the linear combining coefficient to generate the person i 's email pattern. Similarly, from (32), each column in $\mathbf{B} \diamond \mathbf{A}$ can be interpreted as a temporal pattern and $\mathbf{C}_{k, r}$ is the coefficient of the r th temporal pattern for generating the k th day's pattern. Obviously, in contrast to the matrix-based model such as k-means or Gaussian mixture model, the tensor CPD model succeeds in mining clustering structures in multidimensional data. To find the latent factor matrices from the social network data \mathcal{Y} , the following problem is usually solved:

$$\begin{aligned} \min_{\mathbf{A}, \mathbf{B}, \mathbf{C}} & \| \mathcal{Y} - [\mathbf{A}, \mathbf{B}, \mathbf{C}] \|_F^2 \\ \text{s.t. } & \mathbf{A} \geq \mathbf{0}_{I \times R}, \mathbf{B} \geq \mathbf{0}_{J \times R}, \mathbf{C} \geq \mathbf{0}_{K \times R}, \end{aligned} \quad (33)$$

where the nonnegative constraints are added to allow only additions among the R latent rank-1 components. This leads to a parts-based representation of the data, in the sense that each rank-1 component is a part of the data, thus further enhancing the model interpretability.

B. Properties on Nonnegative Gaussian-Gamma Prior

Firstly, it is shown that generally a gamma distribution is not conjugate to a truncated Gaussian distribution. Without loss of generality, consider a truncated Gaussian-gamma distribution pair with the following form:

$$\begin{aligned} p(\mathbf{w}, \{\alpha_s\}_{s=1}^S) &= \prod_{s=1}^S p(\mathbf{w}_s | \alpha_s) p(\{\alpha_s\}_{s=1}^S) \\ &= \prod_{s=1}^S \frac{\mathcal{N}(\mathbf{w}_s | \boldsymbol{\mu}_s, \alpha_s^{-1} \mathbf{I}_{M_s})}{\int_{\underline{\mathbf{w}}_s}^{\bar{\mathbf{w}}_s} \mathcal{N}(\mathbf{w}_s | \boldsymbol{\mu}_s, \alpha_s^{-1} \mathbf{I}_{M_s})} \text{U}(\underline{\mathbf{w}}_s, \bar{\mathbf{w}}_s) \\ &\quad \times \frac{b_s^{\alpha_s}}{\Gamma(\alpha_s)} \alpha_s^{\alpha_s-1} \exp(-b_s \alpha_s), \end{aligned} \quad (34)$$

where the truncated Gaussian pdf is with support $[\underline{\mathbf{w}}_s, \bar{\mathbf{w}}_s]$; parameter $\boldsymbol{\mu}_s, \alpha_s \mathbf{I}_{M_s}$ are the mean vector and the precision matrix of the un-rectified Gaussian pdf. In order to compute

the the posterior pdf $p(\alpha_s | \mathbf{w})$, we only keep the terms relevant to α_s and then the following expression is obtained:

$$p(\alpha_s | \mathbf{w}) \propto \frac{\exp\left(-\alpha_s \left(\frac{\|\mathbf{w}_s - \boldsymbol{\mu}_s\|_F^2}{2} + b\right)\right) \alpha_s^{\alpha_s-1+\frac{1}{2}}}{\int_{\underline{\mathbf{w}}_s}^{\bar{\mathbf{w}}_s} \mathcal{N}(\mathbf{w}_s | \boldsymbol{\mu}_s, \alpha_s^{-1} \mathbf{I}_{M_s})}. \quad (35)$$

Obviously, expression (35) does not take the same functional form as that of the gamma distribution, and thus violates the definition of the conjugacy.

But fortunately, when $\boldsymbol{\mu}_s = \mathbf{0}_{M_s \times 1}$, $\mathbf{w}_s = \mathbf{0}_{M_s \times 1}$ and $\bar{\mathbf{w}}_s = \infty$, it is easy to show that $\int_{\mathbf{0}_{M_s \times 1}}^{\infty} \mathcal{N}(\mathbf{w}_s | \mathbf{0}_{M_s \times 1}, \alpha_s^{-1} \mathbf{I}_{M_s}) = \frac{1}{2^{M_s}}$ due to the symmetric property of the Gaussian pdf. Consequently, under this specific setting, equation (35) becomes

$$p(\alpha_s | \mathbf{w}) \propto \exp\left(-\alpha_s \left(\frac{\|\mathbf{w}_s - \boldsymbol{\mu}_s\|_F^2}{2} + b\right)\right) \alpha_s^{\alpha_s-1+\frac{1}{2}}, \quad (36)$$

which takes exactly the same functional form as that of the gamma distribution. As a result, Property 1 is proved, and this important finding motivates the use of the nonnegative Gaussian-gamma prior in the probabilistic modeling. Furthermore, under this parameter setting, the marginal distribution for parameter \mathbf{w} can be computed as follows:

$$\begin{aligned} p^+(\mathbf{w}) &= \int \prod_{s=1}^S p(\mathbf{w}_s | \alpha_s) p(\{\alpha_s\}_{s=1}^S) d\{\alpha_s\} \\ &= \prod_{s=1}^S 2^{M_s} \int \mathcal{N}(\mathbf{w}_s | \boldsymbol{\mu}_s, \alpha_s^{-1} \mathbf{I}_{M_s}) \text{gamma}(\alpha_s | a_s, b_s) d\alpha_s \\ &\quad \times \text{U}(\mathbf{w}_s \geq \mathbf{0}_{M_s \times 1}). \end{aligned} \quad (37)$$

Further using the fact $\int \mathcal{N}(\mathbf{w}_s | \boldsymbol{\mu}_s, \alpha_s^{-1} \mathbf{I}_{M_s}) \text{gamma}(\alpha_s | a_s, b_s) d\alpha_s = (\frac{1}{\pi})^{\frac{M_s}{2}} \frac{\Gamma(a_s + M_s/2)}{(2b_s)^{a_s} \Gamma(a_s)} (2b_s + \mathbf{w}_s^T \mathbf{w}_s)^{-a_s - M_s/2}$ [37], Property 2 can be proved.

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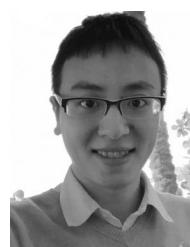
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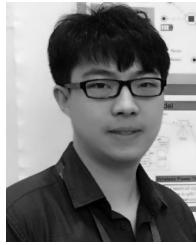
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