Doubly Nonparametric Sparse Nonnegative Matrix Factorization based on Dependent Indian Buffet Processes

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Abstract—Sparse Nonnegative Matrix Factorization (SNMF) aims to factorize a data matrix into two optimized nonnegative sparse factor matrices, which could benefit many tasks, such as document-word co-clustering. However, traditional SNMF typically assumes the number of latent factors (i.e., dimensionality of the factor matrices) to be fixed. This assumption makes it inflexible in practice. In this paper, we propose a doubly sparse nonparametric NMF framework to mitigate this issue by using dependent Indian Buffet Processes (dIBP). We apply a correlation function for the generation of two stick weights associated with each column pair of factor matrices, while still maintaining their respective marginal distribution specified by IBP. As a consequence, the generation of two factor matrices will be column-wise correlated. Under this framework, two classes of correlation function are proposed: (1) using bivariate Beta distribution and (2) using Copula function. Compared to single IBP-based NMF, our work jointly makes two factor matrices nonparametric and sparse, which could be applied to broader scenarios such as co-clustering. Our work is seen to be more flexible than Gaussian Process (GP)-based and Hierarchial Beta Process (HBP)-based dIBPs in terms of allowing the two corresponding binary matrix columns to have greater variations in their non-zero entries. Our experiments on synthetic data show the merits of our work compared to state-of-the-art models in respect of factorization efficiency, sparsity, and flexibility. Experiments on real-world datasets demonstrate the efficiency of our work in document-word co-clustering tasks.

Index Terms—Text mining, co-clustering, nonnegative matrix factorization, probability graphical model

I. INTRODUCTION

S PARSE Nonnegative Matrix Factorization (SNMF) [1] aims to factorize a matrix into two optimized sparse nonnegative factor matrices, which is recognized as an efficient tool for unsupervised learning in many research areas such as image processing, gene analysis, and recommender systems [2]. The improvement of SNMF over classical NMF is due to its sparse data representation [3] which is generally desirable because it assists human understanding (e.g., with gene expression data), reduces computational costs, and obtains better generalization in

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learning algorithms [4]. When two factor matrices from NMF both have the sparsity constraint, they are referred to as *doubly sparse NMF*.

A motivating example that could benefit from *doubly sparse NMF* is the document-word co-clustering task [5], [6], which clusters documents and words simultaneously given a document-word count matrix. Document clustering is useful for organizational purposes or browsing, and word clustering is useful for the automatic construction of a statistical thesaurus or the enhancement of queries [7]. Traditional separative clustering methods do not utilize the relationship between documents and words well, whereas co-clustering achieves much better performance in terms of discovering the structure of data [8]. With the new sparse representations from *doubly sparse NMF*, documents and words could be simultaneously and accurately clustered by considering their interior relations.

The assumption that the dimensionality of factor matrices from doubly sparse NMF needs to be predefined, however, prevents its use in many real-world applications. Normally, this dimensionality is assigned by experts with domain knowledge, but inaccurate assignment will impact the performance of doubly sparse NMF on such applied tasks as document-word co-clustering. Furthermore, the increase in the amount of data and the complexity of the tasks means that even experts are inadequate for this job. Therefore, it is more reasonable and practical to automatically learn the dimensionality from the data (which is known as *nonparametricity*). An intuitive solution from Bayesian nonparametric learning [9], which utilizes stochastic processes [10] as the tools for data analysis, is to use the Indian buffet process (IBP) [11] as the prior for the factor matrices. Unfortunately, IBP can only enable one factor matrix *sparsity* and *nonparametricity* (called Single Nonparametric Sparse NMF [sIBP-NMF] in this paper). Despite its success on some tasks (e.g., factor analysis), sIBP-NMF fails in a number of broader tasks, such as the aforementioned document-word co-clustering. Therefore, our idea is to jointly assign two factor matrices IBP priors (named doubly Nonparametric Sparse NMF), thus endowing both factor matrices with sparsity and nonparametricity.

In this paper, we propose a doubly nonparametric sparse NMF framework to enable traditional NMF with both *double sparsity* and *nonparametricity*. As the core of the proposed framework, two new dependent IBPs (dIBPs) are

innovatively designed as the joint prior for the two factor matrices from NMF based on bivariate Beta distribution and copula. It is dIBP that helps us endow the NMF with both *double sparsity* and *nonparametricity*. Two closely related approaches use GP-based dIBP [12] and HBP-based dIBP [13]. By comparison, the proposed dIBPs allow both factor matrices to have much greater flexibility and variation in terms of their non-zero entries. Furthermore, instead of correlating two IBPs at the binary matrices level, the two proposed dIBPs correlate with the two IBPs at the very beginning (at the Beta random variable level). This strategy results in the implementation of the doubly sparse nonparametric NMF framework based on new dIBPs with a simpler model structure than GP-based dIBP. Nevertheless, introducing bivariate Beta distribution and copula presents a challenge for the model inference. We have designed four inference algorithms for four implementations of the doubly nonparametric sparse NMF framework: GP-based dIBP model, HBP-based dIBP model, bivariate Beta distribution-based dIBP model, and copula-based dIBP model. The experiments on the synthetic data show the merits of our work compared to traditional NMF, single IBP-based NMF, GP-based NMF, and HBP-based NMF on factorization efficiency, sparsity, and correlation flexibility. The experiments on real-world datasets show that the proposed models perform well in the document-word co-clustering task without explicitly predefining the dimensionality of the factor matrices.

The contributions of this paper are summarized as:

- Two new dIBPs (i.e., bivariate Beta distribution-based and copula-based) with simpler model structure and larger correlation flexibility are proposed as alternatives to the existing GP-based and HBP-based dIBPs;
- Four dIBP-based nonparametric doubly sparse non-negative matrix factorization models are proposed to endow the traditional NMF with both *double sparsity* and *nonparametricity*.

The rest of this paper is organized as follows. Preliminary details of NMF and IBP are briefly introduced in Section II. Section III reviews related work. Our dIBP-based doubly sparse nonparametric NMF framework is proposed with four implementations in Section IV, and Gibbs samplers are designed for the four models in Section V. A set of experiments on synthetic data and real-world tasks are conducted in Section VI. Lastly, Section VII concludes the study and discusses further work.

II. PRELIMINARY KNOWLEDGE

A. Sparse nonnegative matrix factorization

Given a nonnegative matrix $Y_{M \times N}$, sparse nonnegative matrix factorization (SNMF) aims to find two matrices $A_{M \times K}$ and $X_{N \times K}$ to minimize the following cost function,

$$J = \|Y - AX^T\|_F^2 + \|A\|_1 + \|X\|_1 \qquad (1)$$

where $\| \cdot \|_F$ is the Frobenius norm, $\| \cdot \|_1$ is the ℓ_1 norm and the elements of A and X have nonnegative constraint.

TABLE I
NOTATIONS IN THIS PAPER

Symbol	meaning in this paper
M	the row number of Y
N	the column number of Y
K	the latent factor number
$K^\dagger$	the truncation level
Y	data matrix with size $M \times N$
$y_{m,n}$	the element of Y at m row and n column
A	factor matrix with size $M \times K$
X	factor matrix with size $N \times K$
$a_{m,k}$	the element of A at m row and k column
$x_{n,k}$	the element of X at n row and k column
$Z^{(1)}$	mask matrix for A with size $M \times K$
$Z^{(2)}$	mask matrix for X with size $N \times K$
$z^{(1)}_{m,k}$	the mask binary variable for the element of A at m row and k column
$z^{(2)}_{n,k}$	the mask binary variable for the element of X at n row and k column
$V^{(1)}$	loading matrix for A with size $M \times K$
$V^{(2)}$	loading matrix for X with size $N \times K$
$v^{(1)}_{m,k}$	the loading variable for the element of A at m row and k column
$v^{(2)}_{n,k}$	the loading variable for the element of X at n row and k column
$\mu^{(1)}_k$	k-th stick weight of IBP for A
$\mu^{(2)}_k$	k-th stick weight of IBP for X
θ	parameters for a bivariate Beta distribution or a copula
$corr$	correlation value

The ℓ_1 norm in the cost function is used for the sparsity constraint.

Several important notations used throughout this paper are summarized in Table I.

Take the document-word co-clustering task as an example. The input $Y_{M \times N}$ denotes the frequency of N words in M documents. The $A_{M \times K}$ denotes the documents' interests in K factors (i.e., topics), and $X_{N \times K}$ denotes the words' interests in K factors (i.e., topics). All documents and words are projected into the same K-dimensional space by NMF. Based on $A_{M \times K}$ and $X_{N \times K}$, the document-word co-clustering task can be accomplished.

One problem of NMF is that the dimensionality of the factor matrices K needs to be predefined. Normally, this variable is experimentally adjusted within a range. This paper will resolve this problem using Bayesian nonparametric learning.

B. Indian buffet process

The Indian Buffet Process (IBP) [14] is defined as a prior for the binary matrices with an infinite number of columns. A stick-breaking construction for IBP [15] is proposed as follows:

$$\nu_j \sim Beta(\alpha, 1), \mu_k = \prod_{j=1}^{k} \nu_j, z_{n,k} \sim Ber(\mu_k) \qquad (2)$$

where $Ber()$ denotes Bernoulli distribution, $z_{n,k}$ forms a matrix $Z_{N \times K}$, $\{\nu_j\}$ is a set of variables with a Beta distribution, α is the parameter of the Beta distribution, and μ_k is the stick weight of column k. The bigger μ_k is,

the more 'ones' appear in column k of the binary matrix $Z_{N \times K}$. The K is determined by the data and the parameter α of IBP.

III. RELATED WORK

The related work mainly falls into two categories: one concerns research on Nonparametric Nonnegative Matrix Factorization and the other concerns the Indian Buffet Process. These state-of-the-art researches inspire our idea of using a dependent Indian Buffet Processes for Bayesian nonparametric nonnegative matrix factorization.

A. Bayesian nonparametric NMF

The Bayesian nonparametric extension of Nonnegative Matrix Factorization mainly relies on the machinery of stochastic processes, and there are two categories that principally use stochastic processes to build Bayesian nonparametric NMF.

One category of models decomposes the data matrix $Y = Z \odot A$ into a binary (mask) matrix Z and a factor matrix A, and binary (mask) matrix is given an infinite prior [16], i.e., IBP [11]. The binary matrix Z functions as the feature selection matrix and the other factor matrix A functions as a feature matrix. The dimensionality of A will change with the change of Z. Another similar approach is to use Beta process [17], in which the data matrix $Y = (S \odot Z)A$ is decomposed into three matrices where S is an ordinary nonnegative matrix and Z is modeled by a Bernoulli process which is in turn given a Beta process prior [18]. Due to the relationship between IBP and the Beta-bernoulli process [17], this approach is similar to the model using IBP. However, there is only one (mask) matrix controlled to have nonparametricity and sparsity properties, which is more like a factor analysis than NMF.

The other category of research assigns the data matrix a Poisson (likelihood) distribution as $Y_{m,n} \sim Poi(\sum_k r_k \phi_{m,k} \varphi_{n,k})$ (named Gamma-Poisson NMF [19]), where r is given an infinite prior, i.e., Gamma process [20]. ϕ and φ, which account for two factor matrices from NMF respectively, are normal nonnegative matrices. The nonparametricity of the model mainly depends on the Gamma process r. The innovative idea of this model is to control the two factor matrices through the coefficient of Poisson data likelihood. This idea is extended by replacing the Gamma-Poisson process with the Beta-Negative binomial process, which allows overdispersion for the count data (matrix) [21]. Since the Beta-Negative binomial process is (in distribution) equal to the Beta-Gamma-Gamma-Poisson process, the idea in [21] about nonparametricity control is same. This kind of method has been applied for assortative network modeling [22], joint document and network modeling [23], and discovering words in spoken utterances [24]. However, although the sparsity of observation is considered [19], the sparsity of the factor matrices, i.e., ϕ and φ, is overlooked, so this idea cannot be used for sparse NMF. Apart from model design, there are also researchers who are focusing on the

inference methods for nonparametric NMF models, such as Power-EP [11], stochastic structured mean-field variational inference [25], and particle filtering [26]. Both of these are specially designed for the first category of nonparametric NMF.

To summarize, although there are a number of preliminary studies on the nonparametric NMF problem, the current state-of-the-art works cannot be directly used for our target: NMF with both *double sparsity* and *nonparametricity*.

B. IBP and dIBP

IBP is proposed in [14], and is a marginalization of the Beta-Bernoulli process. Its widespread popularity is due to its power to generate a binary matrix with infinite columns. Restricted IBP [27] is proposed to allow an arbitrary prior distribution rather than a fixed Poisson distribution form for the number of features in each observation; Integrative IBP [28] is developed for integrating multimodal data in a latent space; Gibbs-type IBP [29] is a generalization of IBP with two-parameter IBP and three-parameter IBP [30] as special cases. All these models provide extensions for the original IBP to endow it with new meaningful features, e.g., power-law behavior. This paper focuses on the dIBP for NMF problem, so the original IBP is used here; the abovementioned works could be considered as interesting extensions of our work in the future.

The dIBP was first proposed in [12] based on Gaussian process, and can be used for the nonparametric NMF after being embedded in our proposed framework, discussed in Section IV. GP-based dIBP models the dependence between IBP with different covariants, such as the time tags of documents, geographic locations of people or GDPs of countries. The idea of dependent nonparametric processes was first proposed by MacEachern [31], and seven different classes of dependence are summarized [32]. The hierarchial Beta process is proposed so that the different Beta processes share a common base discrete measure [13]; it can also be seen as an implicit realization of dIBP which is used for nonparametric NMF in Section IV. The Phylogenetic IBP [33] considers the tree structure of the data points, which can be seen as a supervised IBP. A coupled IBP [34] is proposed for collaborative filtering, which links two IBPs through a factor matrix. However, this coupling does not guarantee that the factor matrices have the same dimensionality which is important for the NMF. To summarize, there is no existing work on using or constructing dIBP for nonparametric NMF.

IV. DOUBLY SPARSE NONPARAMETRIC NMF FRAMEWORK AND ITS THREE IMPLEMENTATIONS

In this section, we propose a Doubly Sparse Nonparametric NMF Framework followed by its four implementations, and we then discuss the differences between these implementations and analyze the properties of models.

A. Doubly sparse nonparametric NMF framework

In our Doubly Sparse Nonparametric NMF Framework, the data matrix is modeled as,

$$Y = AX^T = (V^{(1)} \odot Z^{(1)})(V^{(2)} \odot Z^{(2)})^T \quad (3)$$

where $Z^{(1)}$ and $Z^{(2)}$ are two binary matrices, $V^{(1)}$ and $V^{(2)}$ are loading matrices, and $\odot$ denotes the Hadamard product. Next, we assign these latent variable probability distributions as follows,

$$\begin{aligned} a_{m,k} = v_{m,k}^{(1)} z_{m,k}^{(1)}, & \ v_{m,k}^{(1)} \sim Gam(1, \tau_1) \\ x_{n,k} = v_{n,k}^{(2)} z_{n,k}^{(2)}, & \ v_{n,k}^{(2)} \sim Gam(1, \tau_2) \end{aligned} \quad (4)$$

where $Gam(\cdot, \cdot)$ denotes the Gamma distribution and τ_1 and τ_2 are two parameters. It is evident that $a_{m,k} \geq 0$ and $x_{n,k} \geq 0$ are always satisfied under the above probability distributions. With A and X determined, the likelihood of the model is defined as,

$$y_{m,n}|a_{m,k}, x_{n,k} \sim Exp(y_{m,n}; \sum_k a_{m,k} x_{n,k} + \epsilon) \quad (5)$$

where $Exp(\cdot)$ denotes the Exponential distribution and ϵ is a very small positive number to make the parameter of Exponential distribution greater than zero. The selection of the Exponential distribution is used to guarantee each element of Y with support $[0, +\infty)$. The above parameter setting is used to retain the desired expectations of these distributions, i.e., the expectation of $y_{m,n}$ is $\sum_k a_{m,k} x_{n,k} + \epsilon$.

At the point, we have completed the construction of the doubly sparse nonparametric NMF framework except for an appropriate prior for $Z^{(1)}$ and $Z^{(2)}$. In the following subsections, we will introduce four implementations of this framework using four kinds of dIBP.
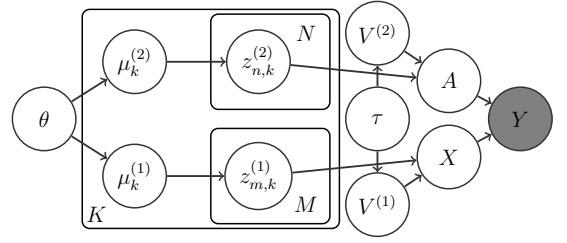
B. Implementation by bivariate Beta distribution-based dIBP

There are different ways to link two IBPs. Here, we propose a method that links the initials of two IBPs, i.e., ν. In the original IBP, ν satisfies a Beta distribution with parameter $(\alpha, 1)$ as in Eq. (2). Therefore, our innovative idea is to find a joint distribution $(\nu^{(1)}, \nu^{(2)})$ with Beta distributions $Beta(\alpha_1, 1)$ and $Beta(\alpha_2, 1)$ as marginal distributions. Following this idea, an intuitive candidate would be Dirichlet distribution. There is a strictly negative relation, i.e., $\nu^{(1)} + \nu^{(2)} = 1$, between the samples of the Dirichlet distribution, but we hope to preserve the freedom of two $(\nu^{(1)}, \nu^{(2)})$.

Instead of the Dirichlet distribution, a bivariate Beta distribution [35] is adopted here whose probability density function is,

$$p(r_1, r_2) = \frac{r_1^{a^0-1} r_2^{b^0-1} (1-r_1)^{b^0+c^0-1} (1-r_2)^{a^0+c^0-1}}{B(a^0, b^0, c^0)(1-r_1 r_2)^{a^0+b^0+c^0}}$$
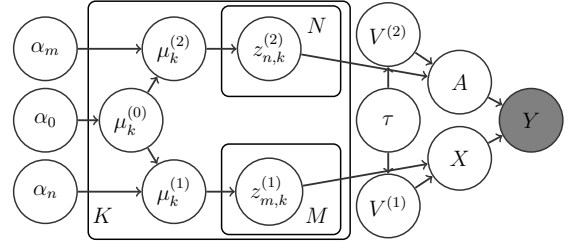$$s.t., \quad 0 \leq r_1, r_2 \leq 1, \quad a^0, b^0, c^0 > 0 \quad (6)$$

where a^0, b^0, c^0 are three parameters of this distribution and $B(a^0, b^0, c^0)$ is the normalization constant that is difficult

(a) Implementations by Bivariate or Copula-based dependent IBP
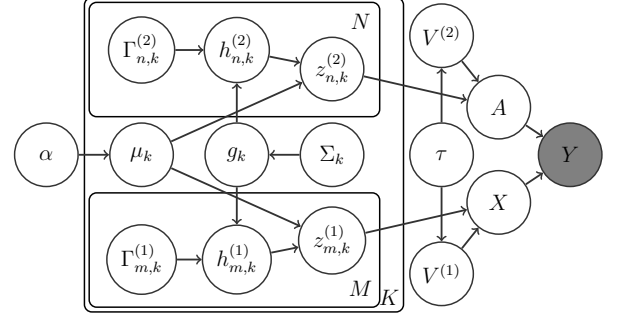
(b) Implementation by GP-based dependent IBP

(c) Implementations by HBP-based dependent IBP

Fig. 1.   Graphical Models for (a) Bivariate Beta distribution-based or Copula-based dependent Indian Buffet Processes, (b) GP-based dependent Indian Buffet Processes, and (c) HBP-based dependent Indian Buffet Processes.

to evaluate. One merit of this bivariate Beta distribution is that two marginal distributions are,

$$r_1 \sim Beta(a^0, c^0), \quad r_2 \sim Beta(b^0, c^0) \quad (7)$$

which satisfy our requirement concerning Beta marginal distributions of $\nu^{(1)}$ and $\nu^{(2)}$. Another merit is that this distribution can model positive correlation between (r_1, r_2) with the range $[0, 1]$ adjusted by the three parameters (a^0, b^0, c^0) compared to the strictly negative correlation from the Dirichlet distribution. Here, we set c^0 of the bivariate Beta distribution to 1 because we must ensure that the marginal distribution of each ν is a Beta distribution with parameter form $(\alpha, 1)$. This condition is necessary for the distribution of the generated binary matrices that satisfy the IBPs. Considering Eq. (7), we give c^0 a fixed value. Even with a fixed value for c^0, the bivariate Beta distribution is still able to model different correlations of two variables. The correlation can be expressed as a function of a^0 and b^0 when $c^0 = 1$,

$$\begin{aligned} corr = & \sqrt{a^0 b^0 (a^0 + 2)(b^0 + 2)} \\ & \left[\frac{\Gamma(a+2)\Gamma(b+2)}{(a+b+1)\Gamma(a+b+2)} \mathbb{F} - 1 \right] \end{aligned} \quad (8)$$

where

$$\mathbb{F} =_3F_2(a^0 + 1, b^0 + 1, a^0 + b^0 + 1; \\ a^0 + b^0 + 2, a^0 + b^0 + 2; 1) \quad (9)$$

is a Hypergeometric function[1] that can be evaluated given the parameters. For example, if $a^0 = 2.5$ and $b^0 = 4$, the correlation between r_1 and r_2 is 0.995; if $a^0 = 0.05$ and $b^0 = 0.1$, the correlation between r_1 and r_2 is 0.080.

With the desired bivariate Beta distribution in hand, we build the BiBeta-based dIBP as follows,

$$(\nu_k^{(1)}, \nu_k^{(2)}) \sim biBeta(\theta), \theta : \{a^0, b^0, c^0 = 1\} > 0$$
$$z_{n,k}^{(1)} \stackrel{\text{i.i.d}}{\sim} Ber(\mu_k^{(1)}), \mu_k^{(1)} = \prod_{j=1}^{k} \nu_j^{(1)}$$
$$z_{n,k}^{(2)} \stackrel{\text{i.i.d}}{\sim} Ber(\mu_k^{(2)}), \mu_k^{(2)} = \prod_{j=1}^{k} \nu_j^{(2)} \quad (10)$$

where $biBeta(\theta)$ denotes the bivariate Beta distribution in Eq. (6) and θ denotes the parameters of the distribution. The graphical model is shown in Fig. 1(a).

Although the bivariate Beta distribution-based dIBP has extended the freedom of the relation between $\nu^{(1)}$ and $\nu^{(2)}$, the relation is restricted to positive relations in bivariate Beta distribution. Next, we use the copula to capture more freedom of their relations.

C. Implementation by copula-based dIBP

Copula links two variables with given marginal distributions, and is used to define a joint distribution for variables with known marginal distributions in statistics. Here, we use the Farlie-Gumbel-Morgenstern (FGM) Copula [36] as an example. The definition of the FGM Copula is,

$$C_\rho(o_1, o_2) = o_1 o_2 + \rho o_1 o_2 (1 - o_1)(1 - o_2)$$
$$c_\rho(o_1, o_2) = 1 + \rho(2o_1 - 1)(2o_2 - 1) \quad (11)$$

where $C_\rho(o_1, o_2)$ is the joint cumulative distribution function, $c_\rho(o_1, o_2)$ is the joint probability density function, $\rho \in [-1, 1]$ is the parameter of the FGM copula, and o_1 and o_2 are the cumulative distribution function values of two marginal distributions that are known in advance.

For our problem, the two variables are $\nu^{(1)}$ and $\nu^{(2)}$ and their marginal distributions are Beta distributions with parameters $(\alpha_1, 1)$ and $(\alpha_2, 1)$. Therefore, o_1 and o_2 are defined as,

$$o_1 = F^{beta}(\nu^{(1)}) \sim (\nu^{(1)})^{\alpha_1}, f^{beta}(\nu^{(1)}) \sim \alpha_1(\nu^{(1)})^{\alpha_1 - 1}$$
$$o_2 = F^{beta}(\nu^{(2)}) \sim (\nu^{(2)})^{\alpha_2}, f^{beta}(\nu^{(2)}) \sim \alpha_2(\nu^{(2)})^{\alpha_2 - 1}$$

where $F^{beta}(\nu^{(1)})$ and $f^{beta}(\nu^{(1)})$ represent the cumulative distribution function and probabilistic density function of $\nu^{(1)}$.

In Copula, the correlation between $\nu^{(1)}$ and $\nu^{(2)}$ is modeled or reflected by the value of ρ. Various correlations (i.e., positive, null or negative) can be captured by the different ρ. The Spearman correlation can be evaluated by

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[1]https://en.wikipedia.org/wiki/Generalized_hypergeometric_function

$corr = \frac{\rho}{3}$. Since the support of ρ is $[-1, 1]$, the correlation range that can be modeled by the FGM Copula is $[-\frac{1}{3}, \frac{1}{3}]$. In particular, there is no correlation if $\rho = 0$.

Copula-based dIBP is defined by replacing the bivariate Beta distribution in Eq. (10) with joint distribution defined by the FGM copula,

$$p(\nu_k^{(1)}, \nu_k^{(2)}) = c_\rho(\theta), \\ \theta : \{\rho \in [-1, 1], \alpha_1 > 0, \alpha_2 > 0\} \quad (12)$$

The graphical model is the same as the BiBeta-based dIBP in Fig. 1(a) but with different parameters θ.

There are a number of works concerning the constructing of dependent random measures through the Levy Copula [37], [38], such as Levy Copula-based dependent Dirichlet process [39] and Levy Copula-based dependent Poisson-Dirichlet process [40]. Levy Copula-based dependent random measures have an advantage that the marginal random measures do not have to be of the same type of process [37]. For example, a Beta process could be linked with a Gamma process. The Levy copula is mainly used to link two Levy processes, but IBP is not a Levy process although its de Finetti mixing distribution (i.e., Beta process) is. The Copula used in this work is an ordinary one that is used for linking two random variables with fixed marginal distributions (Beta distributions in this work).

D. Implementation by GP-based dIBP

The first dIBP is proposed based on Gaussian Process (GP) [12]. In this GP-based dIBP, each stick weight μ_k is used to generated a different number of columns of binary matrices. In other words, the GP-based dIBP uses the same set of sticks for different binary matrices. We can use this GP-based dIBP for the NMF as the prior for the matrices $Z^{(1)}$ and $Z^{(2)}$. The graphical model is shown in Fig. 1(b), and the generative process is as follows,

$$\nu_j \sim Beta(\alpha, 1), \ \mu_k = \prod_{j=1}^{k} \nu_j \quad (13)$$

where the $\{\mu_k\}$ are IBP sticks as in Eq. (2). This set of sticks is shared by two binary matrices through

$$g_k \sim GP(0, \Sigma_k)$$
$$h_{m,k}^{(1)} \sim GP(g_k, \Gamma_{m,k}^{(1)})$$
$$h_{n,k}^{(2)} \sim GP(g_k, \Gamma_{n,k}^{(2)})$$
$$\Sigma_k(t, t') = \sigma^2 \exp(-\frac{(t - t')^2}{s^2})$$
$$\Gamma^{(1)} = \Gamma^{(2)} = \eta^2 I$$
$$z_{m,k}^{(1)} = \delta\{h_{m,k}^{(1)} < F^{-1}(\mu_k | 0, (\Sigma_k)^{(1,1)} + (\Gamma_{m,k}^{(1)})^{(1,1)})\}$$
$$z_{n,k}^{(2)} = \delta\{h_{n,k}^{(2)} < F^{-1}(\mu_k | 0, (\Sigma_k)^{(2,2)} + (\Gamma_{n,k}^{(2)})^{(2,2)})\}$$

where $GP(0, \Sigma_k)$ denotes a Gaussian process parameterized by a mean function 0 and a kernel function Σ_k. g_k is a random draw from $GP(0, \Sigma_k)$ which is in turn set as the mean function of $GP(g_k, \Gamma_{m,k}^{(1)})$ and $GP(g_k, \Gamma_{n,k}^{(2)})$, so there is a hierarchy between different GPs where the

correlation is captured. $\Gamma_{m,k}^{(1)}$ and $\Gamma_{n,k}^{(2)}$ are identity matrices with a parameter η. $h_{m,k}^{(1)}$ and $h_{n,k}^{(2)}$ are two random draws from the corresponding GPs. $F^{-1}(\cdot)$ is the inverse normal cumulative distribution function, and $\delta\{\cdot\}$ is the indicator function. More details can be found in [12]. Since there are only two binary matrices, the GP is degenerated to a two-dimensional Gaussian distribution, and Σ_k is equal to a 2×2 matrix.

E. Implementation by HBP-based dIBP

Another possible candidate for the dIBP construction is the Hierarchial Beta process (HBP) [13]. The graphical model of HBP-dIBP is shown in Fig. 1(c) and the generative process is as follows,

$$B = \sum_{k=1}^{\infty} \mu_k^{(0)} \delta_{\theta_k} \sim BP(\alpha_0, B_0)$$

$$B^{(1)} = \sum_{k=1}^{\infty} \mu_k^{(1)} \delta_{\theta_k} \sim BP(\alpha_m, B), Z^{(1)} \sim BeP(B^{(1)})$$

$$B^{(2)} = \sum_{k=1}^{\infty} \mu_k^{(2)} \delta_{\theta_k} \sim BP(\alpha_n, B), Z^{(2)} \sim BeP(B^{(2)})$$

where $BP(\alpha, B_0)$ denotes a Beta process parameterized by a base measure B_0 and a concentration parameter α_0; $BeP(B^{(1)})$ denotes a Bernoulli process parameterized by a base measure $B^{(1)}$. In HBP, B, which is a random draw from $BP(\alpha_0, B_0)$, is set as the base measure of $BP(\alpha_m, B)$ and $BP(\alpha_n, B)$. Strictly speaking, the above model is not dependent on a dIBP because no explicit dIBP is built in this model. As proven in [13], when $\widetilde{Z} \sim BeP(\widetilde{B})$ and $\widetilde{B} \sim BP(1, \widetilde{B_0})$ where the total mass of $\widetilde{B_0}$ is $\widetilde{\alpha}$, the marginal distribution of $\widetilde{Z}$ with $\widetilde{B}$ integrated out is $IBP(\widetilde{\alpha})$. However, the marginal distribution of $B^{(1)}$ or $B^{(2)}$ with B integrated out is not necessarily a BP (not to mention with the specific parameters, i.e., 1 and $\widetilde{B_0}$), so we cannot draw the conclusion that $Z^{(1)}$ and $Z^{(2)}$ are with IBP marginal distributions. Although there is no dIBP explicitly built in this model, it can still be seen as a prior for the $Z^{(1)}$ and $Z^{(2)}$. Note that the stick weights $\mu_k^{(1)}$ and $\mu_k^{(2)}$ for two IBPs both center on $\mu_k^{(0)}$. This means that there are the same expected non-zero items in $Z^{(1)}$ and $Z^{(2)}$ across all columns, which is too restrictive. Take document-word co-clustering as an example. This restriction means that the non-zero items of documents and words tend to have a similar number on all factors/topics. It is more reasonable to let documents and words have flexible behavior/'interests' on the different factors/topics.

F. Discussion on models

One apparent advantage of bivariate Beta distribution-based and copula-based dIBP compared with GP-based dIBP is that fewer latent variables are involved. This can easily be observed in the graphical models in Fig. 1. More latent variables tend to slow down the convergence of the model inference. The advantage of Copula compared to
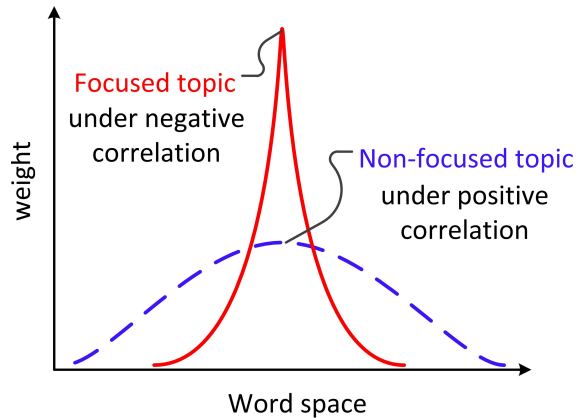

Fig. 2. Illustration of the meaning of learned correlation. The red/solid curve denotes a focused distribution on words from a large negative correlation; the blue/dashed curve denotes a non-focused distribution on words from a large positive correlation.

bivariate Beta distribution is that we can easily obtain both the cumulative distribution function and probability density function of $(\nu_k^{(1)}, \nu_k^{(2)})$. This will impact on the model inference, which will be discussed later.

Another advantage of the proposed dIBPs is their greater flexibility. GP-based dIBP works in the following fashion: Let $Z_j^{(1)}$ and $Z_j^{(2)}$ be the j-th corresponding columns of two identical sized binary matrices $Z^{(1)}$ and $Z^{(2)}$. The model assumes that $Z_j^{(1)}$ and $Z_j^{(2)}$ share the same stick weight μ_j, making the expectation of the number of non-zero entries identical for both $Z_j^{(1)}$ and $Z_j^{(2)}$. The GP is merely used to control the correlations between individual entry pairs WITHIN $Z_j^{(1)}$ and $Z_j^{(2)}$, by thresholding a Gaussian cumulative distribution function (CDF) to make the entry either one or zero. HBP-based dIBP is similar to GP-based dIBP, which also makes the expectation of the number of non-zero entries identical for both $Z_j^{(1)}$ and $Z_j^{(2)}$ through a hierarchal structure. The two parameters α_m and α_n are instead used to control the respective variance from the expectation for two matrices. Although both models could be used to implement a doubly sparse nonparametric NMF, it nonetheless is inadequate in many matrix factorisation scenarios as the assumption that the total number of non-zero entries are (in expectation) distributed identically does not hold universally. Take document-word co-clustering as an example. The number of non-zero factors of a Document column may be drastically different from the number of non-zero factors of a corresponding Word column in a document-word matrix. Contrarily, there is no restriction on the expectations of μ_1 and μ_2 for two binary matrices $Z^{(1)}$ and $Z^{(2)}$ in the proposed bivariate Beta distribution-based dIBP and Copula-based dIBP, which allows both $Z_j^{(1)}$ and $Z_j^{(2)}$ to have much greater flexibility and variation in terms of their non-zero entries.

What does the correlation between $\mu^{(1)}$ and $\mu^{(2)}$ reveal from the data? We explain its meaning using the document-word matrix as an example again. We obtain two factor matrices $A_{M \times K}$ and $X_{N \times K}$ after factorization. The K

hidden factors are also named *topics* in some literatures. We know that $\mu_k^{(1)}$ and $\mu_k^{(2)}$ account for the generation of $Z_k^{(1)}$ and $Z_k^{(2)}$ which are the k-th column of $Z_{M \times K}^{(1)}$ and $Z_{N \times K}^{(2)}$. If $\mu_k^{(1)}$ is larger, the number of 'ones' in $Z_k^{(1)}$ is greater; the same holds for $\mu_k^{(2)}$ and $Z_k^{(2)}$. When there is a large negative correlation between $\mu_k^{(1)}$ and $\mu_k^{(2)}$, the number of 'ones' in $Z_k^{(2)}$ will be smaller if the number of 'ones' in $Z_k^{(1)}$ is larger. This means that the commonly used topics in documents (the number of 'ones' in $Z_k^{(1)}$ is large) tend to be *focused topics* which only focus on infrequent words (the number of 'ones' in $Z_k^{(2)}$ is small), as illustrated by the red/solid curve/distribution on words in Fig. 2. In contrast, when there is a large positive correlation between $\mu_k^{(1)}$ and $\mu_k^{(2)}$, the number of 'ones' in $Z_k^{(2)}$ will be larger if the number of 'ones' in $Z_k^{(1)}$ is also larger. This means that the commonly used topics in documents (the number of 'ones' in $Z_k^{(1)}$ is large) tend to be *non-focused topics* which have many words (the number of 'ones' in $Z_k^{(2)}$ is also large), as illustrated by the blue/dashed curve/distribution on words in Fig. 2. Therefore, we can conclude that the correlation between $\mu^{(1)}$ and $\mu^{(2)}$ can be seen as the measurement of the *focus degree* of commonly-used hidden factors/topics in the dataset. Note that *focused* here is different from the one used in [41] where *focused* means each document could flexibly select their own topics rather than rigidly following the correlation defined in HDP [42], but it is here to highlight the topic with limited number of words with dominant weights.

V. Model Inference

The objective of this section with data matrix Y is to estimate the hidden variables by a properly designed MCMC inference algorithm for their posterior distribution, $p(\mu, Z, V, \theta | Y)$. It is difficult to perform posterior inference under infinite mixtures, thus a common work-around solution in Bayesian nonparametric learning is to use a truncation method. The truncation method, which uses a relatively big $K^\dagger$ as the (potential) maximum number of factors, is widely accepted. We want to highlight that GP-based and HBP-based dIBPs are not our contribution. Our contribution is to link the GP-based and HBP-based dIBPs with the nonnegative matrix factorization likelihood. The inference of these models is given in the Appendix for the self-contained purpose.

A. Update stick weights μ

When updating μ_k, we need to find its conditional distribution given μ_{k-1} and μ_{k+1}, because the order of μ must be maintained to make the marginal distributions of Z satisfy two IBPs. Based on the M-H sampler, the acceptance ratio of a new sample for $\mu_k = [\mu_k^{(1)}, \mu_k^{(2)}]$ is,

$$\min \left(1, \frac{p(Z|\mu_k^*) p(\mu_k^* | \mu_{k+1}, \mu_{k-1})}{p(Z|\mu_k) p(\mu_k | \mu_{k+1}, \mu_{k-1})} \times \frac{q(\mu_k)}{q(\mu_k^*)} \right) \quad (14)$$

where $q(\cdot)$ is a proposal distribution which will be explained later; μ_k^* is a new sample drawn from the proposal

distribution $q(\cdot)$; $p(Z|\mu_k)$ is the likelihood of μ_k to generate the kth column of binary matrix Z as in Eq. (10) and the $p(\mu_k^* | \mu_{k+1}, \mu_{k-1})$ is the conditional probability density function of μ_k^* within the range of $[\mu_{k+1}, \mu_{k-1}]$, which will be different when different strategies in Section IV are used to link $\mu_k^{(1)}$ and $\mu_k^{(2)}$.

Next, we derive the conditional probability density function of $p(\mu_k | \mu_{k+1}, \mu_{k-1})$ as follows: for the first column,

$$\mu_1^{(1)} = \nu_1^{(1)}, \quad \mu_1^{(2)} = \nu_1^{(2)}$$

and

$$p(\mu_1^{(1)}, \mu_1^{(2)}) = p(\nu_1^{(1)}, \nu_1^{(2)})$$

where $p(\cdot, \cdot)$ is the joint probability density function given by Eq. (6) or Eq. (11) from two proposed dIBPs. For the k-th column ($k > 1$),

$$\mu_k^{(1)} = \nu_k^{(1)} \mu_{k-1}^{(1)}, \quad \mu_k^{(2)} = \nu_k^{(2)} \mu_{k-1}^{(2)}$$

and

$$p(\mu_k^{(1)}, \mu_k^{(2)}) = \frac{p\left(\mu_k^{(1)} / \mu_{k-1}^{(1)}, \mu_k^{(2)} / \mu_{k-1}^{(2)} \right)}{\mu_{k-1}^{(1)} \mu_{k-1}^{(2)}}$$

To summarize, the conditional density of μ_k is

$$p(\mu_k | \mu_{k-1}, \mu_{k+1})$$
$$\propto \frac{p\left(\mu_k^{(1)} / \mu_{k-1}^{(1)}, \mu_k^{(2)} / \mu_{k-1}^{(2)} \right)}{\mu_{k-1}^{(1)} \mu_{k-1}^{(2)}} \frac{p\left(\mu_{k+1}^{(1)} / \mu_k^{(1)}, \mu_{k+1}^{(2)} / \mu_k^{(2)} \right)}{\mu_k^{(1)} \mu_k^{(2)}}$$

Considering the support of μ_k, the proposal distribution, $q(\cdot)$, is set as the product of two independent truncated Beta distributions: $\mu_k^{(1)} \sim Beta\left(\frac{\alpha_1}{K^\dagger}, 1 \right), \mu_k^{(1)} \in [\mu_{k+1}^{(1)}, \mu_{k-1}^{(1)}]$ and $\mu_k^{(2)} \sim Beta\left(\frac{\alpha_2}{K^\dagger}, 1 \right), \mu_k^{(2)} \in [\mu_{k+1}^{(2)}, \mu_{k-1}^{(2)}]$ It is quite easy to sample the truncated Beta distribution because Beta distribution is a standard distribution.

B. Update binary matrices Z

Two binary matrices, $Z : \{Z^{(1)}, Z^{(2)}\}$, can be updated separately. Each element in the two matrices satisfies a Bernoulli distribution with the following conditional posterior probabilities:

$$p(z_{m,k}^{(1)} = 1) \propto \mu_k^{(1)} \prod_n e^{-y_{m,n}(\sum_l v_{m,l}^{(1)} z_{m,l}^{(1)} v_{n,l}^{(2)} z_{n,l}^{(2)} + \epsilon)}$$

$$p(z_{m,k}^{(1)} = 0) \propto (1 - \mu_k^{(1)}) \prod_n e^{-y_{m,n}(\sum_l v_{m,l}^{(1)} z_{m,l}^{(1)} v_{n,l}^{(2)} z_{n,l}^{(2)} + \epsilon)}$$

$$(15)$$

and

$$p(z_{n,k}^{(2)} = 1) \propto \mu_k^{(2)} \prod_m e^{-y_{m,n}(\sum_l v_{m,l}^{(1)} z_{m,l}^{(1)} v_{n,l}^{(2)} z_{n,l}^{(2)} + \epsilon)}$$

$$p(z_{n,k}^{(2)} = 0) \propto (1 - \mu_k^{(2)}) \prod_m e^{-y_{m,n}(\sum_l v_{m,l}^{(1)} z_{m,l}^{(1)} v_{n,l}^{(2)} z_{n,l}^{(2)} + \epsilon)}$$

$$(16)$$

where ϵ is a small positive number. Since elements of Z having discrete distribution, it is easy to obtain samples.

C. Update loading matrices V

Since the prior for V is Gamma distribution and the likelihood is exponential distribution, the conditional distribution $p(v_{m,k}^{(1)}|\cdots)$ is proportional to,

$$e^{v_{m,k}^{(1)}\tau_1} \prod_n e^{-y_{m,n}(\sum_k v_{m,k}^{(1)} z_{m,k}^{(1)} v_{n,k}^{(2)} z_{n,k}^{(2)} + \epsilon)} \quad (17)$$

and $p(v_{n,k}^{(2)}|\cdots)$ is proportional to,

$$e^{v_{n,k}^{(2)}\tau_2} \prod_m e^{-y_{m,n}(\sum_l v_{m,l}^{(1)} z_{m,l}^{(1)} v_{n,l}^{(2)} z_{n,l}^{(2)} + \epsilon)} \quad (18)$$

Due to the existence of the ϵ, the posterior of V is not a Gamma distribution, so we have to use the M-H sampler to obtain its samples.

D. Update model parameter θ

The graphical model in Fig. 1(c) has the parameter θ. For the different strategies to link $(\nu^{(1)}, \nu^{(2)})$, the parameters must be different. Therefore, we design corresponding update methods for the proposed two strategies: bivariate Beta distribution and copula.

1) Bivariate Beta distribution: The parameters of bivariate Beta distribution, $\theta : \{a^0, b^0\}$, are given two Gamma priors. The conditional distributions are,

$$p([a^0 \; b^0]) \propto Gam([a^0 \; b^0]; hp) \prod_{k=1}^{K} p(\mu_k^{(1)}, \mu_k^{(2)} | a^0, b^0) \quad (19)$$

where hp is the hyper-parameter of the prior for a^0 and b^0 and K is the number of active columns of Z. The 'active' column means that there is at least one element with 1 in that column.

2) Copula: There are three parameters for each copula, $\theta : \{\rho, \alpha_1, \alpha_2\}$. Their conditional distributions are,

$$p([\alpha_1 \; \alpha_2]) \propto Gam([\alpha_1 \; \alpha_2]; hp) \prod_{k=1}^{K} c(\mu_k^{(1)}, \mu_k^{(2)}) \quad (20)$$

where $c(\cdot, \cdot)$ is the copula density in Eq. (11). We give the ρ of the FGM copula a uniform distribution on its support $[-1, 1]$, and its posterior is

$$p(\rho|\cdots) \propto \prod_{k=1}^{K} c(\mu_k^{(1)}, \mu_k^{(2)} | \rho) \quad (21)$$

After introducing update methods for all the latent variables, we summarize the inference (i.e., Gibbs sampler) for the four models in Algorithm 1 for the bivariate Beta distribution-based dIBP NMF (BB-dIBP-NMF) model, the Copula-based dIBP NMF (C-dIBP-NMF) model, and HBP-based dIBP NMF (HBP-dIBP-NMF) model. Algorithm 2 is for the GP-based dIBP NMF (GP-dIBP-NMF) model.

Algorithm 1: Gibbs Sampler for BB-dIBP-NMF, C-dIBP-NMF, and HBP-dIBP-NMF

Input: Y
Output: A, X
initialization;
while $i \le max_{iter}$ **do**
    // latent variables of dIBP
    //for BB-dIBP-NMF
    Update μ by Eq. (14);
    //for C-dIBP-NMF
    Update μ by Eq. (14);
    //for HBP-dIBP-NMF
    Update μ by Eqs. (32), (30) and (31);
    // binary matrix
    Update Z by Eq. (15) and (16);
    // latent variables of NMF
    Update V by Eq. (17) and (18);
    $i + +$;
return A and X;

Algorithm 2: Gibbs Sampler for GP-dIBP-NMF

Input: Y
Output: A, X
initialization;
while $i \le max_{iter}$ **do**
    // latent variables of dIBP
    Update μ by Eq. (22);
    Update Z by Eq. (28) and (29);
    Update g by Eq. (24);
    Update h by Eq. (25);
    Update s by Eq. (27);
    // latent variables of NMF
    Update V by Eq. (17) and (18);
    $i + +$;
return A and X;

E. Computational complexity analysis

The Gibbs sampling updates of BB-dIBP-NMF can be calculated in $O((M + N)(K^\dagger)^2 + (M + N + 3)K^\dagger)$ time, where $K^\dagger$ is the truncation level. Note that $O((M+N)K^\dagger)$ accounts for the latent variable of NMF that is shared across four implementations, and $O(3K^\dagger + (M+N)(K^\dagger)^2)$ accounts for BB-dIBP. When $K^\dagger$ is set relatively large in comparison to the scale of data matrix (i.e., M and N), the cost of BB-dIBP is primarily of the order of $O((M + N)(K^\dagger)^2)$. We do not consider the cost from the latent variable of NMF for other implementations. The cost of the posterior inference for the C-dIBP update is $O(2K^\dagger + (M + N)(K^\dagger)^2)$ which is smaller than BB-dIBP with $K^\dagger$. Under relatively large $K^\dagger$, they are almost equal. The update cost for GP-dIBP is $O(MNK^\dagger + (M + N)(K^\dagger)^2)$ which is higher than BB-dIBP and C-dIBP. Lastly, the inference of BB-dIBP can be calculated in $O((M+N+3)K^\dagger + (M+N)(K^\dagger)^2)$ time, which is lower than GP-dIBP but a little higher than BB-dIBP and C-dIBP.
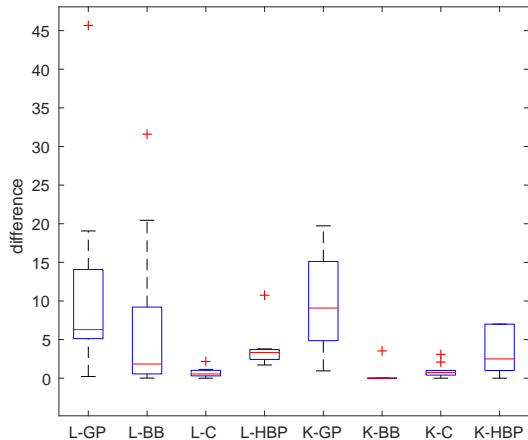
Fig. 3. Comparison of convergence of different models. For instance, $L-GP$ denotes the difference between the data likelihoods from GP-dIBP-NMF; $K-GP$ denotes the difference between effective factor number from GP-dIBP-NMF.

VI. EXPERIMENTS

In this section, we first take a series of experiments on the synthetic data to show the merits of our work by comparing the traditional NMF, single IBP-based NMF, and GP-based NMF on sparsity and nonparametricity (Section VI.A) and flexibility (Section VI.B). A real-world task, i.e., document-word co-clustering, is conducted to show the usefulness of our work compared to other models (Section VI.C).

A. Evaluation of convergence

This section checks and compares the convergence of the proposed models. First, we randomly generate a data matrix with size $M = 10, N = 20$. The Geweke test [43] is then used for the sampling convergence check. This test splits the samples into two parts (after removing a burn-in period, i.e., the first 1,000 iterations): the first 1,000 and the last 2,000. If the chain is at stationarity, the means of the two parts should be equal. A smaller difference between them therefore indicates better convergence. Here, we use two latent variables: one is the latent factor number and the other is the data likelihood. Fig. 3 shows the statistical results of the models, i.e., GP-dIBP-NMF, BB-dIBP-NMF, C-dIBP-NMF, and HBP-dIBP-NMF, after 10 independent runs on the same synthetic dataset. From this figure, we can see that the proposed models (BB-dIBP and C-dIBP) have better convergence than GP-dIBP. Of the proposed models, the Copula-based model achieves the best performance on convergence.

B. Evaluation of sparsity and nonparametricity

We randomly generate a matrix $Y_{20 \times 30}$ using the following procedure: 1) give a vector $[0.5, 0.4, 0.3, 0.2, 0.1]$ as the parameters of five Bernoulli distributions; 2) randomly generate the elements of i-th column (i.e., $A(:, i)$ and $X(:, i)$) of both matrices $A_{20 \times 5}$ and $X_{30 \times 5}$ using
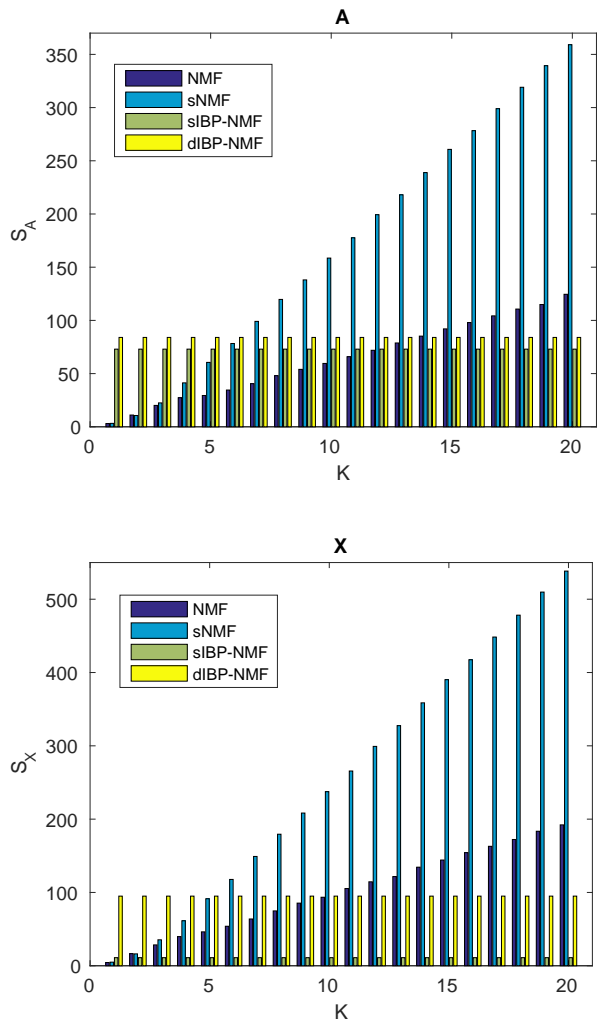

Fig. 4. Comparison of the sparsity on synthetic dataset between traditional NMF (NMF), traditional sparse NMF (sNMF), single IBP-based sparse NMF (sIBP-NMF), and doubly IBP-based sparse NMF (dIBP-NMF). Note that dIBP-NMF here is based on bivariate Beta distribution.

the Bernoulli distribution with i-th value of the above vector; 3) generate Y as the product of A and X as $Y = AX^T$. Since the values of the parameters of Bernoulli distributions are small, the generated factor matrices A and X tend to be sparse. Here, the matrix Y is used as the input data for different algorithms (i.e., traditional sparse NMF (sNMF) in Eq. (1), single nonparametric sparse NMF (sIBP-NMF), and the proposed dIBP-NMF), and the sparsity of the learned factor matrices from the different algorithms are evaluated and compared. We design the following metric to quantitatively compare the sparsity from different algorithms: $S_A = \sum_m \sum_k \mathbb{1}(A(m, k) = 0)$ and $S_X = \sum_n \sum_k \mathbb{1}(X(n, k) = 0)$, where $\mathbb{1}(\cdot)$ is an indicator function parameterized by a condition which equals 1 if condition is satisfied; 0, otherwise. Here, we separately evaluate the sparsity of A and X considering the sIBP-NMF. Note that the 0.00000001 acts as a relaxation of the sparsity in the implementation. In the experiments, we randomly generate a number 100 of $Y_{20 \times 30}$ using the above procedure. The results are shown in Fig. 4, which compares
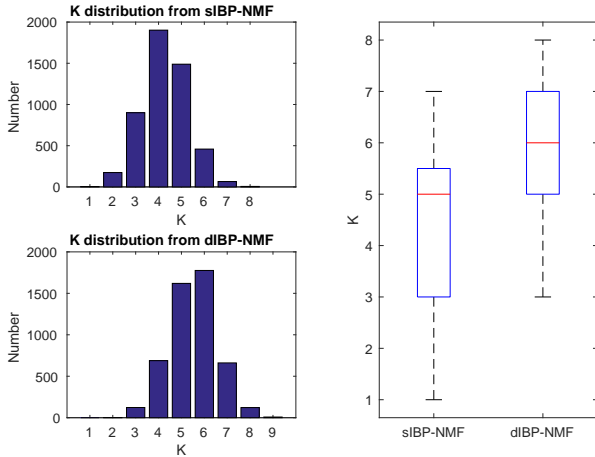
Fig. 5. Comparison of the learned topic number distribution on synthetic dataset between single IBP-based sparse NMF (sIBP-NMF) and doubly IBP-based sparse NMF (dIBP-NMF). Note that dIBP-NMF here is based on bivariate Beta distribution.
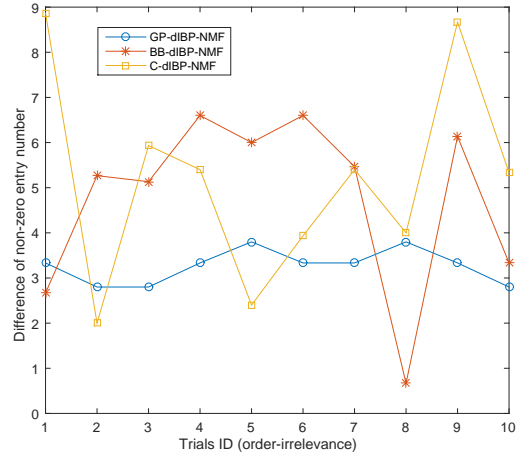

Fig. 6. Results on synthetic data to show the flexibility of the different models. The x-axis denotes the trial IDs (order is irrelevant).

the sparsity of the learned factor matrices A and X from different algorithms. Note that the results are an average of 100 trials. The x-axis denotes the number of factors for NMF and sNMF since they need this as input, but sIBP-NMF and dIBP-NMF do not. We can see that sNMF has greater sparsity than NMF due to the sparse constraint. It appears that the sparsity increases as the number of factors also increases. Since sIBP-NMF and dIBP-NMF do not need the factor number as input, the values of S_A and S_X from two algorithms are equal. For each trial, we can obtain a distribution on factor number, as shown by two examples in Fig. 5 for sIBP-NMF and dIBP-NMF with 5,000 Gibbs samples. The peak of distribution is seen as the final learned factor number from the algorithm. The statistics on the learned factor numbers from 100 trials are shown in the right subfigure in Fig. 5. The averages are around the benchmark (i.e., 5 in the generative procedure), which denotes the relative accuracy of the factor learning (i.e., nonparametric property). Not surprisingly, sIBP-NMF has sparse A but not sparse X (the value of S_A is large even compared to the value in sNMF but S_X is small). Its sparsity of A is due to its IBP prior. Since there are two IBP priors for A and X in dIBP-NMF, the resultant A and X are both sparse. Therefore, we can draw the conclusion that the proposed dIBP-NMF could obtain two sparse factor matrices but sIBP-NMF could only obtain one, which will impact on the ability to conduct the co-clustering task that will be demonstrated in Section VI.D.

C. Evaluation of correlation flexibility

As claimed, our proposed models have greater correlation flexibility for allowing the numbers of non-zero entries in factor matrices more different from each other compared to GP-based dIBP. To show this flexibility, we first design a metric to measure the flexibility by comparing the number of non-zero entries of two factor matrices (A and X) from

the models. The mean of the differences between the corresponding columns of A and X is $\frac{1}{K}\sum_{k=1}^{K}|N_k^{(1)} - N_k^{(2)}|$, where K is the number of columns of both matrices, $N_k^{(1)}$ is the number of non-zero entries of k-th column of A, and $N_k^{(2)}$ is the number of non-zero entries of k-th column of X. It appears that the larger this metric is, the more flexibility a model has. We have ten randomly generated matrices of the same size: 20×30, and we run three models on ten matrices. The designed metric has been evaluated on the learned factor matrices of different models. As shown in Fig. 6, we can see that the metrics on BB-dIBP-NMF and C-dIBP-NMF are larger than those on GP-dIBP-NMF in most trials and are also with more fluctuations and larger non-zero entry number differences compared to GP-dIBP-NMF. We conclude that the proposed dIBPs are more flexible than GP-based dIBP.

D. Real-world task: Document-word co-clustering

In this subsection, we apply the proposed algorithms to a real-world task: document-word co-clustering. The real-world datasets[2] used for this task are:

- **Cora Dataset** The Cora dataset consists of 2708 scientific publications classified into seven classes. The dictionary consists of 1433 unique words.
- **Citeseer Dataset** The CiteSeer dataset consists of 3312 scientific publications. The dictionary consists of 3703 unique words. The labels of these papers are set as their research areas.

The above datasets already have benchmarks for the document clusters but do not have benchmarks for the word clusters. We use the co-occurrence relations between words to generate a distance matrix through which we can obtain a number (10 being the best) of word clusters by spectral clustering algorithm. These word clusters are seen as benchmarks. The evaluation metrics (bigger means

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[2] http://linqs.cs.umd.edu/projects/projects/lbc/

TABLE II
NUMERICAL RESULTS OF DOCUMENT-WORD CO-CLUSTERING ON *Citeseer* DATASET

Evaluation Metric		Models or Algorithms							
		Bayesian Nonparametric NMF Models					Parametric Models or Algorithms		
		sIBP	BB-dIBP	C-dIBP	GP-dIBP	HBP-dIBP	NMF	SNMF	Spectral Clustering
doc	JC	0.2390	0.2483	0.2411	0.2323	0.2406	0.2050±0.0321	0.2318±0.0312	0.1784
	FM	0.3860	0.4132	0.3876	0.3902	0.3906	0.3420±0.0440	0.3764±0.0404	0.4219
	F1	0.3510	0.3417	0.3729	0.3851	0.3697	0.3398±0.0447	0.3753±0.0415	0.3029
word	JC	0.6521	0.6923	0.6793	0.6762	0.6627	0.6412±0.0425	0.6940±0.0467	N/A
	FM	0.7798	0.8417	0.8301	0.8361	0.8154	0.7809±0.0242	0.8325±0.0263	N/A
	F1	0.7973	0.8223	0.8119	0.8182	0.8063	0.7667±0.0273	0.8186±0.0294	N/A

TABLE III
NUMERICAL RESULTS OF DOCUMENT-WORD CO-CLUSTERING ON *Cora* DATASET

Evaluation Metric		Models or Algorithms							
		Bayesian Nonparametric Models					Parametric Models or Algorithms		
		sIBP	BB-dIBP	C-dIBP	GP-dIBP	HBP-dIBP	NMF	SNMF	Spectral Clustering
doc	JC	0.1680	0.1720	0.1700	0.1650	0.1683	0.1524±0.0239	0.1678±0.0170	0.1691
	FM	0.2770	0.2790	0.2610	0.2630	0.2700	0.2729±0.0366	0.2882±0.0263	0.3123
	F1	0.2890	0.2930	0.3090	0.2610	0.2633	0.2688±0.0354	0.2870±0.0254	0.2737
word	JC	0.4380	0.5320	0.5190	0.5170	0.5210	0.4328±0.0545	0.5098±0.0464	N/A
	FM	0.6780	0.7320	0.7190	0.7300	0.6809	0.5946±0.0314	0.7129±0.0292	N/A
	F1	0.6420	0.6770	0.6650	0.6650	0.6532	0.5739±0.0361	0.6742±0.0366	N/A

better) for clustering are Jaccard Coefficient: $JC = \frac{a}{a+b+c}$, Folkes&Mallows: $FM = \left(\frac{a}{a+b}\frac{a}{a+c}\right)^{1/2}$ and F1 measure: $F1 = \frac{2a^2}{2a^2+ac+ab}$, where a is the number of two points that are in the same cluster of both benchmark result and clustering result; b is the number of two points that are in the same cluster of benchmark result but in different clusters of clustering result; c is the number of two points that are not in the same cluster of the two benchmark result but are in the same cluster of clustering result.

We compare the following models on the document-word co-clustering task: three parametric models or algorithms (classical NMF, sparse NMF (SNMF) in Eq. (1), and spectral clustering) and five Bayesian nonparametric models (single IBP-based sparse NMF (sIBP-NMF) [11], GP-dIBP-NMF, BB-dIBP-NMF, C-dIBP-NMF, and HBP-dIBP-NMF). When applied to document-word co-clustering, the output A and X from each of the above models can be considered as new representations of documents and words on latent factors. Based on these new representations, we can use a clustering algorithm (*K-means*, in this section, and K is set as the number of benchmark clusters) to conduct document and word clustering. Since a common algorithm (i.e., *K-means*) is adopted for all models, the performance of models will be only determined by the learned new data representation A and X. We also compare the performance of above models with spectral clustering on document clustering. Note that since the benchmark of word clustering is built based on spectral clustering, we do not compare spectral clustering with other models or algorithms on word clustering.

The results on *Citeseer* are all listed in Table II. Since NMF and SNMF require the factor number as input, we adjust this parameter for them from 1 to 100; thus, there are fluctuations in the results from NMF and SNMF in Table II where the standard deviations are given after the mean value. The other models are all nonparametric

models which do not need the factor number as input, so the results of these models are listed in the table with only one value. The learned factor numbers are 9 (from sIBP-NMF), 10 (from BB-dIBP-NMF), 12 (from C-dIBP-NMF), 12 (from GP-dIBP-NMF) and 10 (from HBP-dIBP-NMF). The results on *Cora* are all listed in Table III. The learned factor numbers are 10 (from sIBP-NMF), 17 (from BB-dIBP-NMF), 15 (from C-dIBP-NMF), 22 (from GP-dIBP-NMF) and 12 (from HBP-dIBP-NMF). From these results, we can see that: 1) There are fluctuations in the performance of SNMF and NMF on document and word clustering due to the factor number parameter. Note that spectral clustering needs the cluster number to be fixed in advance, like traditional clustering methods. For the sake of comparison, we simply feed the benchmark cluster number of the datasets to algorithms, so there is no fluctuation in its results in the table. 2) Although sIBP-NMF has relatively good performance on document clustering, the performance on word clustering is poor, which may be due to its single side sparse and nonparametric control. 3) Five Bayesian nonparametric models achieve comparable or better (in most cases) performance than three parametric models or algorithms. Considering the release of the factor number assumption, these Bayesian nonparametric models are an improvement on traditional NMF. 4) Comparing sIBP-NMF, the other Bayesian nonparametric models all perform better on both document and word clustering with a weak exception by HBP-dIBP on document clustering on *Cora*. To summarize, without prior knowledge of the number of factors, the proposed algorithms achieve relatively good performance on the document clustering task. Of the three algorithms, BB-dIBP-NMF achieves the best performance overall.

We also check how the learned models perform on the prediction of held-out data by 5-fold cross validation. The setting is as follows: 1) train the models on the training data ($Y_{doc \times words}^{training}$) to obtain new representations for

TABLE IV
NUMERICAL RESULTS OF HELD-OUT DATA PREDICTION ON *Citeseer* DATASET

Prediction precision		Models or Algorithms						
		Bayesian Nonparametric NMF Models					Parametric Models or Algorithms	
		sIBP	BB-dIBP	C-dIBP	GP-dIBP	HBP-dIBP	NMF	SNMF
doc	training	0.6981	0.7165	0.7293	0.7097	0.7121	0.6613±0.0659	0.6792±0.0953
	test	0.6977	0.7063	0.7087	0.6944	0.7002	0.6386±0.0753	0.6973±0.0891
word	training	0.8638	0.8810	0.8677	0.8598	0.8701	0.8655±0.0159	0.8712±0.0155
	test	0.8203	0.8332	0.8307	0.8267	0.8312	0.8193±0.0269	0.8170±0.0288

TABLE V
NUMERICAL RESULTS OF HELD-OUT DATA PREDICTION ON *Cora* DATASET

Prediction precision		Models or Algorithms						
		Bayesian Nonparametric NMF Models					Parametric Models or Algorithms	
		sIBP	BB-dIBP	C-dIBP	GP-dIBP	HBP-dIBP	NMF	SNMF
doc	training	0.6790	0.7212	0.7197	0.7112	0.7081	0.6362±0.0832	0.6487±0.0877
	test	0.6298	0.6765	0.6904	0.6600	0.6556	0.6117±0.0762	0.6231±0.0965
word	training	0.8952	0.9091	0.9126	0.8889	0.9001	0.8960±0.0157	0.8940±0.0262
	test	0.8643	0.8673	0.8799	0.8620	0.8692	0.8759±0.0280	0.8431±0.0301

the training data, i.e., new representations of documents $A_{doc \times factor}^{training}$; 2) use the new representation of data to train a classifier (i.e., KNN here); 3) predict the labels of the test data using the new representations of test data $A_{doc \times factor}^{test} = Y^{test}(X_{factor \times words}^{test})^{-1}$ and the trained classifier. The procedure for words is similar to the above. The prediction metric is the ratio of the correctly predicted data number to the number of all data. The results are listed in Tables IV and V. Note that the standard deviations in these tables are from the different factor numbers, not the cross validation. We can see from these results that: 1) The models are not overfitting because there is not a big difference between the prediction on the training and test data; 2) The Bayesian nonparametric models have relatively better performance compared to NMF and SNMF; 3) BB-dIBP-NMF and C-dIBP-NMF achieve the best prediction on *Citeseer* and *Cora*, respectively.

VII. CONCLUSION AND FURTHER STUDY

Nonnegative matrix factorization is advantageous for many machine learning tasks (e.g., co-clustering), but the assumption that the dimension of the factors is known in advance makes NMF impractical for many applications. To resolve this issue, we have proposed a doubly sparse nonparametric NMF framework based on dIBP to remove the assumption. First, two models were built by implementing this framework using GP-based dIBP and HBP-based dIBP, which successfully remove the assumption but suffers from larger model complexity or less flexibility. Then, we proposed two new dIBPs through bivariate Beta distribution and a copula. The advantages of the models based on the new dIBPs is that 1) they have simpler model structures than models with GP-based dIBP, and 2) the correlation in data can be directly learned out, which can be seen as a measurement of the focus degree of hidden factors/topics. Lastly, four inference algorithms have been designed for the proposed models, respectively. The experiments on synthetic and real-world datasets demonstrates the capability of the proposed models to perform NMF without predefining the dimensionality and more correlation flexibility compared to GP-based dIBP and HBP-based dIBP.

One possible future area of study for this work is the aspect of efficiency. Current Gibbs sampling inference is not efficient enough for big data. Our future study will focus on the efficiency of the inference of the proposed models using the variational inference strategy.

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APPENDIX A
THE CONDITIONAL DISTRIBUTIONS FOR THE GP-BASED DIBP NMF

The conditional distributions are:

Sampling μ

$$p(\mu_k | \cdots) \propto \frac{\mu_K^\alpha}{\mu_k} \prod_{t=1}^{2} \prod_n^{N_t} (\gamma_k^t)^{z_{n,k}^t} (1 - \gamma_k^t)^{1 - z_{n,k}^t} \quad (22)$$

where

$$\gamma_k^t = F(F^{-1}(\mu_k | 0, \Sigma_k^{(t,t)} + \eta^2) - g_k^t | 0, \eta^2) \quad (23)$$

where $F()$ is a normal cumulative distribution function.

Sampling g

$$p(g_k | \cdots) \propto \mathcal{N}(g_k | 0, \Sigma_k) \cdot \prod_t \prod_n^{N_t} \mathcal{N}(h_{n,k}^t | g_k^t, \eta^2) \quad (24)$$

where $\mathcal{N}()$ denotes normal distribution.

Sampling h

$$p(h_{n,k}^t|\cdots) \propto \mathcal{N}(g_k^t, \eta^2), \begin{cases} h_{n,k}^t \in (-\infty, \widetilde{\mu}_k^t] & \text{if } z_{n,k}^t = 1 \\ h_{n,k}^t \in [\widetilde{\mu}_k^t, +\infty) & \text{if } z_{n,k}^t = 0 \end{cases} \tag{25}$$

where

$$\widetilde{\mu}_k^t = F^{-1}(\mu_k|0, \Sigma_k^{(t,t)} + \eta^2) \tag{26}$$

Sampling s

$$p(s|\cdots) \propto Gam(s; hs, 1) \prod_k^K \mathcal{N}(g_k|0, \Sigma_k) \tag{27}$$

Sampling Z

$$p(z_{m,k}^{(1)} = 1) \propto \gamma_k^1 \prod_n e^{-y_{m,n}(\sum_l v_{m,l}^{(1)} z_{m,l}^{(1)} v_{n,l}^{(2)} z_{n,l}^{(2)} + \epsilon)}$$

$$p(z_{m,k}^{(1)} = 0) \propto (1-\gamma_k^1) \prod_n e^{-y_{m,n}(\sum_l v_{m,l}^{(1)} z_{m,l}^{(1)} v_{n,l}^{(2)} z_{n,l}^{(2)} + \epsilon)}$$

$$\tag{28}$$

and

$$p(z_{n,k}^{(2)} = 1) \propto \gamma_k^2 \prod_m e^{-y_{m,n}(\sum_l v_{m,l}^{(1)} z_{m,l}^{(1)} v_{n,l}^{(2)} z_{n,l}^{(2)} + \epsilon)}$$

$$p(z_{n,k}^{(2)} = 0) \propto (1-\gamma_k^2) \prod_m e^{-y_{m,n}(\sum_l v_{m,l}^{(1)} z_{m,l}^{(1)} v_{n,l}^{(2)} z_{n,l}^{(2)} + \epsilon)}$$

$$\tag{29}$$

where γ_k^t is the same as in Eq. (23).

APPENDIX B
THE CONDITIONAL DISTRIBUTIONS FOR THE HBP-BASED dIBP NMF

When using truncation level $K^\dagger$, the approximation of the original HBP-based dIBP [44] is as $\mu_k^{(0)} \sim Beta\left(\frac{\alpha_0}{K^\dagger}, \alpha_0\left(1 - \frac{1}{K^\dagger}\right)\right)$ where $\mu_k^{(1)} \sim Beta\left(\alpha_m \mu_k^{(0)}, \alpha_m(1 - \mu_k^{(0)})\right)$ and $\mu_k^{(2)} sim Beta\left(\alpha_n \mu_k^{(0)}, \alpha_n(1 - \mu_k^{(0)})\right)$ The conditional distributions of the hidden variables are:

Sampling μ_k^1

$$\mu_k^{(1)} \sim Beta(\alpha_m \mu_k^{(0)} + \sum_m z_{m,k}^{(1)},$$
$$\alpha_m(1 - \mu_k^{(0)}) + M - \sum_m z_{m,k}^{(1)}) \tag{30}$$

Sampling μ_k^2

$$\mu_k^{(2)}|\cdots \sim Beta(\alpha_n \mu_k^{(0)} + \sum_n z_{n,k}^{(2)},$$
$$\alpha_n(1 - \mu_k^{(0)}) + N - \sum_n z_{n,k}^{(2)}) \tag{31}$$

Sampling μ_k^0 The log probabilistic density of μ_k^0 is proportional to

$$\begin{aligned} &\alpha_0(\frac{1}{K^\dagger} - 1) \log \mu_k^{(0)} + \frac{\alpha_0}{K^\dagger} \log(\mu_k^{(0)}) \\ &+ \alpha_m \mu_k^{(0)} \log \mu_k^{(1)} + \alpha_m(1 - \mu_k^{(0)}) \log(1 - \mu_k^{(1)}) \\ &+ (\alpha_n \mu_k^{(0)}) \log \mu_k^{(1)} + \alpha_n(1 - \mu_k^{(0)}) \log(1 - \mu_k^{(2)}) \\ &- \log \Gamma(\alpha_m \mu_k^{(0)}) - \log \Gamma(\alpha_m(1 - \mu_k^{(0)})) \\ &- \log \Gamma(\alpha_n \mu_k^{(0)}) - \log \Gamma(\alpha_n(1 - \mu_k^{(0)})) \end{aligned} \tag{32}$$

Finally, the parameter α_m could also join the sampler with the density $Gam(\alpha_m; hp) \prod_{k=1}^K Beta(\alpha_m \mu_k^{(0)}, \alpha_m(1 - \mu_k^{(0)}))$
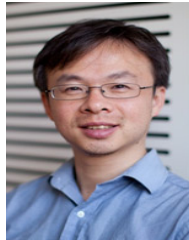
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