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# Affine-transformation invariant clustering models

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## Abstract

We develop a cluster process which is invariant with respect to unknown affine transformations of the feature space without knowing the number of clusters in advance. Specifically, our proposed method can identify clusters invariant under (I) orthogonal transformations, (II) scaling-coordinate orthogonal transformations, and (III) arbitrary nonsingular linear transformations corresponding to models I, II, and III, respectively and represent clusters with the proposed heatmap of the similarity matrix. The proposed Metropolis-Hastings algorithm leads to an irreducible and aperiodic Markov chain, which is also efficient at identifying clusters reasonably well for various applications. Both the synthetic and real data examples show that the proposed method could be widely applied in many fields, especially for finding the number of clusters and identifying clusters of samples of interest in aerial photography and genomic data.

**Keywords:** Dirichlet process, Ewens process, Metropolis-Hastings algorithm, Markov chain Monte Carlo sampling, Unsupervised learning

## 1 Introduction

Clustering of objects invariant with respect to affine transformations of feature vectors is an important research topic since objects may be recorded via different angles and positions so that their coordinates may vary and their nearest neighbors may belong to other clusters. For example, the longitude, latitude, and altitude coordinates of an object which are recorded by devices equipped in aircrafts or satellites change across different observational time. In this situation, distance-based clustering method including  $k$ -means (MacQueen 1967), hierarchical clustering (Ward 1963), clustering based on principal components, spectral clustering (Ng et al. 2001), and others (Jain and Dubes 1988; Ozawa 1985) may fail to identify the correct clusters by grouping nearest points. Another category is distribution-based clustering methods (Banfield and Raftery 1993; Fraley and Raftery 1998; Fraley and Raftery 2002; Fraley and Raftery 2007; McCullagh and Yang 2008; Vogt et al. 2010) which may specify a partition as a parameter in a likelihood function and estimate it under a Bayesian framework.

In certain areas of application, the goal is to cluster objects  $i = 1, \dots, n$  into disjoint subsets based on their feature vectors  $Y_i \in \mathbb{R}^d$ . In this paper, we propose group

invariance by considering three cases of a cluster process that are invariant with respect to three groups of affine transformations  $g: \mathbb{R}^d \rightarrow \mathbb{R}^d$  acting on the feature space. The group invariance implies that the feature configurations  $Y$  and  $Y'$  in  $\mathbb{R}^{n \times d}$  determine the same clustering, or probability distribution on clusterings, if they belong to the same group orbit that is an equivalence class. For example, if the feature space is Euclidean and  $\mathcal{G}$  is the group of Euclidean isometries or congruences, the clustering is a function only of the maximal invariant, which is the array of Euclidean distances  $D_{ij} = \|Y_i - Y_j\|$ . For example, image data such as the aerial photography and three-dimensional protein structures are two motivating examples. The shape and relative locations of data may vary due to the change of the viewer's angle and location.

Our goal is to develop a novel clustering method which can identify clusters of  $Y = (Y_1, \dots, Y_n)$  even when all  $Y_i$ 's are mapped by an unknown affine transformation  $Y'_i = \mathbf{a} + AY_i$ , where  $\mathbf{a} = (a_1, \dots, a_d) \in \mathbb{R}^d$  and  $A \in \mathbb{R}^{d \times d}$  is nonsingular. Affine-invariant clustering is important when the clusters are not well-separated in the observational space. Although there are previous work on affine-invariant clustering methods (Fitzgibbon and Zisserman 2002; Begelfor and Werman 2006; Shioda R. and Tunçel 2007; Brubaker. S.C. and Vempala 2008; Kumar and Orlin 2008; García-Escudero et al. 2010; Lee et al. 2014), these existing methods handle different problems from ours. These methods aim to cluster the same item observed in different angles or mapped by different unknown affine transformations. Instead, in our problem setting we consider only one unknown affine transformation that is applied to all objects.

The affine transformations consist of three types: (1) index permutations, rotation, one-scaling on all variables, and location-translation transformations that are under the first type of covariance structures and named model I whose transformation and covariance structure  $\sigma^2 I_d$  were also adopted by Vogt et al. (2010); (2) each variable may have different scaling transformations that are under the second type of covariance structures and named model II; (3) the variables are transformed by a nonsingular matrix that is named model III, where the observed variables may be linear combinations of some latent variables in model I. These models cover fairly general situations of clustering in nature.

McCullagh and Yang (2008) constructed a Dirichlet cluster process together with a random partition representing the clustering. In this paper, we follow their setup and extend their framework. We assume that the random partition of objects follows Ewens distribution (Ewens 1972), and we propose a likelihood of the responses which is invariant respect to affine transformations.

## 2 Cluster process and prior distributions

In this paper, an  $\mathbb{R}^d$ -valued *cluster process*  $(Y, B)$  means a random partition  $B$  of the natural numbers, together with an infinite sequence  $Y_1, Y_2, \dots$  of random vectors in the state space  $\mathbb{R}^d$ . The restriction of such a process to a finite sample  $[n] = \{1, \dots, n\}$  of units or specimens consists of the restricted partition  $B[n]$  accompanied by the finite sequence  $Y[n] = (Y_1, \dots, Y_n)$ . A partition  $B[n]: [n] \times [n] \rightarrow \{0, 1\}$  is the partition of the sample units expressed as a binary cluster-factor matrix of  $B_{ij} = 1$  if  $Y_i$  and  $Y_j$  are of the same cluster (denoted as  $i \sim j$ ), and  $B_{ij} = 0$  otherwise (McCullagh and Yang 2008). For example, when  $n = 3$ , the partition  $\{\{1, 2\}, 3\}$  and the cluster labels 112 correspond to an equivalence relation

$$B = \begin{pmatrix} 1 & 1 & 0 \\ 1 & 1 & 0 \\ 0 & 0 & 1 \end{pmatrix}.$$

Notice that the elements of  $B$  are transitional. i.e., if individuals  $i, j, k$  belong to the same cluster, then  $B_{i,j} = 1$  and  $B_{j,k} = 1$  imply  $B_{i,k} = 1$ .

The term *cluster process* implies infinite exchangeability, which means that the joint distribution  $p_n$  of  $(Y[n], B[n])$  is symmetric (McCullagh and Yang 2006) or invariant under permutations of indices (Pitman 2002), and  $p_n$  is the marginal distribution of  $p_{n+1}$  under deletion of the  $(n+1)$ th unit from the sample.

Similar to (McCullagh and Yang 2008), we construct an exchangeable Gaussian mixture as a simple example of clustering processes. First,  $B \sim p$  is some infinitely exchangeable random partition. Secondly, the conditional distribution of the samples  $Y$ , which is regarded as a matrix  $(Y_{i,r})$  of order  $n \times d$  given  $B$  (say the cluster label  $cl(Y_i) = l$ ) and  $\theta$ , is Gaussian with mean and variance as follows

$$E(Y_i | B, \mu_l) = \mu_l, \quad \text{Cov}(Y_{i,r}, Y_{j,s} | B, \theta) = (\delta_{i,j} + \theta B_{i,j}) \Sigma_{r,s},$$

where  $\mu_l = (\mu_{l1}, \dots, \mu_{ld}) \in \mathbb{R}^d$  is the centroid of cluster  $k$ ,  $\delta$  is Kronecker's delta, that is,  $\delta_{i,j} = 1$  if  $i = j$  and 0 if  $i \neq j$ ,  $\theta > 0$  is a ratio parameter connecting the within- and between-cluster covariance matrices, and  $\Sigma = (\Sigma_{r,s})$  is a positive definite matrix of order  $d \times d$ , known as the within-cluster covariance matrix. In our settings, the between-cluster covariance matrix is simply  $\theta \Sigma$ , the cluster centroids  $\mu_1, \dots, \mu_k$  are iid from  $N(\mu, \theta \Sigma)$ , and the mean of  $Y$  given  $B$  and  $\mu_1, \dots, \mu_k$  is

$$E(Y | B, \mu_1, \dots, \mu_k) = (\mu_{cl(Y_1)}, \dots, \mu_{cl(Y_n)})$$

and the covariance of  $Y$  given  $B$  can also be represented by the covariance of its vector form  $\text{Vec}(Y) = (Y_{11}, \dots, Y_{1d}, \dots, Y_{n1}, \dots, Y_{nd})^\top$  as

$$\text{Cov}(\text{Vec}(Y) | B, \theta) = (I_n + \theta B) \otimes \Sigma$$

which is an  $nd \times nd$  matrix with “ $\otimes$ ” indicating the Kronecker product.  $\Sigma$ , the column covariance of  $Y$ , is assumed identical for all clusters,  $I_n + \theta B$  is assumed an exchangeable structure for the row covariance of  $Y$ , and  $\theta$  is the product of the standard deviations of two rows. There exist competing algorithms that are affine-equivariant and do not impose this requirement (Shioda R. and Tunçel 2007; Kumar and Orlin 2008; García-Escudero et al. 2010; Lee et al. 2014). The identity matrix itself is also a partition in which each cluster consists of one element.

Given the number of clusters  $k$ , the cluster sizes  $(n_1, \dots, n_k)$  may follow a multinomial distribution with category probabilities  $\pi = (\pi_1, \dots, \pi_k)$ , where  $\pi$  follows an exchangeable Dirichlet distribution  $\text{Dir}(\lambda/k, \dots, \lambda/k)$ . After integrating out  $\pi$ , the partition  $B$  follows a Dirichlet-multinomial prior

$$p_n(B | \lambda, k) = \frac{k!}{(k - \#B)!} \frac{\Gamma(\lambda) \prod_{b \in B} \Gamma(n_b + \lambda/k)}{\Gamma(n + \lambda) [\Gamma(\lambda/k)]^{\#B}}$$

where  $\#B \leq k$  denotes the number of clusters presented in the partition  $B$  and  $n_b$  is the size of cluster  $b$  (MacEachern 1994; Dahl 2005; McCullagh and Yang 2008). The limit as  $k \rightarrow \infty$  is well defined and known as the Ewens's sampling formula (ESF) with parameter  $\lambda > 0$

$$p_n(n_1, \dots, n_k | \lambda) = \frac{\Gamma(\lambda) \lambda^{\#B}}{\Gamma(n + \lambda)} \prod_{b \in B} \Gamma(n_b),$$

which is also known as Chinese restaurant process (CRP) (Ewens 1972; Neal 2000; Blei and Jordan 2006; Crane 2016). McCullagh and Yang (2008) provided a framework with a finite number of clusters and general covariance structures. In this paper, we adopt the CRP prior for partition  $B$  which implies  $k = \infty$  in the population with the proposed Gaussian likelihood to get the affine-invariant clusters. Note that  $\#B \leq n$  for any given sample size  $n$ .

We choose a proper prior distribution for the variance ratio  $\theta$ , the symmetric  $F$ -family

$$p(\theta) \propto \frac{\theta^{\alpha-1}}{(1+\theta)^{2\alpha}}$$

with  $\alpha > 0$  allowing a range of reasonable choices (Chaloner 1987).

We propose a sampling procedure to estimate the partition  $B$  and the parameter  $\theta$  from conditional probabilities. Since the conditional distribution of  $\theta$  does not have a recognized form, we propose to use a discrete version  $\{p(\theta_j)\}_{j=1}^J$ , where  $J$  is a predetermined moderately large integer. Based on our experience,  $J = 100$  works reasonably well for the real data examples that we have examined.

### 3 Affine-transformation invariant clustering

The affine-transformation invariant clustering identified in this manuscript is invariant even when the objects are mapped by an unknown affine transformation. The conditional distribution on partitions of  $[n] = \{1, \dots, n\}$  is determined by the finite sequence  $Y = (Y_1, \dots, Y_n)$  regarded as a configuration of  $n$  labeled points in  $\mathbb{R}^d$ . The exchangeability condition implies that any permutation  $\pi$  of the sequence induces a corresponding permutation in  $B$ , i.e.  $p_n(B^\pi | Y = y^\pi) = p_n(B | Y = y)$ , where  $y_i^\pi = y_{\pi(i)}$  and  $B_{i,j}^\pi = B_{\pi(i),\pi(j)}$ . In many cases, it is reasonable to assume additional symmetries involving transformations in  $\mathbb{R}^d$ , for example  $p_n(B | Y) = p_n(B | -Y)$ . We are asking, in effect, whether two labeled configurations  $Y$  and  $Y'$  which are *geometrically equivalent* in  $\mathbb{R}^d$  should determine the same conditional distribution on sample partitions.

If the state space  $\mathbb{R}^d$  is regarded as a  $d$ -dimensional Euclidean space with the standard Euclidean inner product and Euclidean metric, the configurations  $Y$  and  $Y'$  are *congruent* if there exists a vector  $\mathbf{a} = (a_1, \dots, a_d) \in \mathbb{R}^d$  and an orthogonal matrix  $A \in \mathbb{R}^{d \times d}$  such that  $Y'_i = \mathbf{a} + AY_i$  for each  $i$ . Equivalently, the  $n \times n$  arrays of squared Euclidean distances  $D_{ij} = \|Y_i - Y_j\|^2$  and  $D'_{ij} = \|Y'_i - Y'_j\|^2$  are equal. The configurations are geometrically *similar* if  $Y'_i = \mathbf{a} + bY_i$  for  $b \in \mathbb{R}$  and  $b \neq 0$ , implying that the arrays of distances are proportional  $D' = b^2 D$ .

The geometric equivalence is defined by regarding the observation  $Y$  as a group orbit rather than a point. In general, the group is the affine group  $GA(\mathbb{R}^d)$ ,  $\mathcal{G} = \mathbb{R}^d \times L$  and  $L$  is the collection of all  $d \times d$  nonsingular matrices, with the operation  $(\mathbf{a}_1, A_1) \circ (\mathbf{a}_2, A_2) = (\mathbf{a}_1 + A_1 \mathbf{a}_2, A_1 A_2)$  for  $\mathbf{a}_i \in \mathbb{R}^d, A_i \in L$  with  $i = 1, 2$ , which is consistent with compositions of affine transformations. The orbit of an element  $Y = (Y_1, \dots, Y_n)^\top \in \mathbb{R}^{n \times d}$  is defined as

$$\text{Orb}(Y) = \left\{ X \in \mathbb{R}^{n \times d} : \exists g \in \mathcal{G} \text{ s.t. } X = g \star Y \right\}, \quad (1)$$

where the group action is that  $\mathcal{G}$  acts on  $\mathbb{R}^{n \times d}$  as

$$(\mathbf{a}, A) \star Y = (\mathbf{a} + AY_1, \dots, \mathbf{a} + AY_n)^\top = \mathbf{1}_n \mathbf{a}^\top + YA^\top \quad (2)$$

where  $\mathbf{1}_n$  is a length- $n$  vector of 1's. It can be verified that its vector form  $\text{Vec}((\mathbf{a}, A) \star Y) = \mathbf{1}_n \otimes \mathbf{a} + (I_n \otimes A)\text{Vec}(Y)$ . If

$$\text{Vec}(Y) \sim N(\mathbf{1}_n \otimes \boldsymbol{\mu}, (I_n + \theta B) \otimes \boldsymbol{\Sigma}),$$

then an element in the same orbit

$$\text{Vec}((\mathbf{a}, A) \star Y) \sim N(\mathbf{1}_n \otimes (\mathbf{a} + A\boldsymbol{\mu}), (I_n + \theta B) \otimes (A\boldsymbol{\Sigma}A^\top))$$

More specifically,

$$\begin{aligned} \text{Vec}((-\boldsymbol{\mu}, I_d) \star Y) &\sim N(\mathbf{0}, (I_n + \theta B) \otimes \boldsymbol{\Sigma}) \\ \text{Vec}((-T^{-1}\boldsymbol{\mu}, T^{-1}) \star Y) &\sim N(\mathbf{0}, (I_n + \theta B) \otimes I_d) \end{aligned}$$

where  $T$  is a  $d \times d$  nonsingular matrix satisfying  $\boldsymbol{\Sigma} = TT^\top$ .

**Theorem 1** If  $n \leq d$ , then all  $Y \in \mathbb{R}^{n \times d}$  of full rank  $n$  belong to the same orbit. If  $n = d + 1$ , then all  $Y \in \mathbb{R}^{n \times d}$  satisfying  $\text{rank}(Y) = \text{rank}(Y - \mathbf{1}_n \mathbf{1}_n^\top Y/n) = d$  belong to the same orbit.

The proof of Theorem 1 is relegated to the Appendix A. According to the proof, if  $n = d + 1$ , then  $\text{rank}(Y) = d$  implies that  $\text{rank}(Y - \mathbf{1}_n \mathbf{1}_n^\top Y/n)$  is either  $d$  or  $d - 1$ . The case of  $d - 1$  only occupies a lower-dimensional subspace.

According to Theorem 1, for  $n \leq d + 1$ , the action is essentially transitive in the sense that all configurations of  $n$  distinct points in  $\mathbb{R}^d$  belong to the same orbit: all other orbits are negligible in that they have Lebesgue measure zero. As a result, the observation  $Y$  regarded as a group orbit  $\mathcal{G}Y$  is uninformative for clustering unless  $n > d + 1$ . We name the orbit and group action defined above as model III.

In model I, which is the case considered in Vogt et al. (2010), the covariance between features are proportional to an identity matrix. The group is  $\mathcal{G} = \mathbb{R}^d \times \mathbb{R} \setminus \{0\}$  with the operation  $(\mathbf{a}_1, b_1) \circ (\mathbf{a}_2, b_2) = (\mathbf{a}_1 + b_1 \mathbf{a}_2, b_1 b_2)$  for  $\mathbf{a}_i \in \mathbb{R}^d, b_i \in \mathbb{R} \setminus \{0\}, i = 1, 2$ . The orbit of an element  $Y \in \mathbb{R}^{n \times d}$  and the group action are defined similarly as in (1) and (2) with  $A$  replaced by  $b$ . Then  $(\mathbf{a}, b) \star Y = \mathbf{1}_n \mathbf{a}^\top + bY$  and  $\text{Vec}((\mathbf{a}, b) \star Y) = \mathbf{1}_n \otimes \mathbf{a} + b\text{Vec}(Y)$ . If

$$\text{Vec}(Y) \sim N(\mathbf{1}_n \otimes \boldsymbol{\mu}, (I_n + \theta B) \otimes \sigma^2 I_d),$$

then  $\text{Vec}((-\boldsymbol{\mu}, 1) \star Y) \sim N(\mathbf{0}, (I_n + \theta B) \otimes \sigma^2 I_d)$  and  $\text{Vec}((-\boldsymbol{\mu}/\sigma, 1/\sigma) \star Y) \sim N(\mathbf{0}, (I_n + \theta B) \otimes I_d)$ , which correspond to elements in  $\text{Orb}(Y)$ .

In essence, the observation is not regarded as a point in  $\mathbb{R}^{n \times d}$  but is treated as a *group orbit* generated by the group of rigid transformations, or similarity transformations if scalar multiples are permitted. In statistical terms, this approach meshes with the submodel in which the matrix  $\boldsymbol{\Sigma}$  in model I is a scaled identity matrix  $I_d$ . An equivalent way of saying the same thing for  $n > d$  is that the column-centered sample matrix  $\tilde{Y} = Y - \mathbf{1}_n \mathbf{1}_n^\top Y/n$  determines the sample covariance matrix  $S = (\tilde{Y}^\top \tilde{Y})/(n - 1)$  and hence the Mahalanobis metric  $\|x - x^*\|^2 = (x - x^*)^\top S^{-1}(x - x^*)$  in the state space (Mahalanobis 1936; Gnanadesikan and Kettenring 1972). One implication is that the  $n \times n$  matrix

$D = (D_{ij}) = \left( \|Y_i - Y_j\|^2 \right)$  of standardized inter-point Mahalanobis distances is maximal invariant, and the conditional distribution on sample partitions depends on  $Y$  only through this matrix.

In practice, the  $d$  variables are sometimes measured on scales that are not commensurate with one another, so the state space seldom has a natural metric. In this case, we assume that  $Y$  and  $Y'$  as equivalent configurations for each feature  $Y_{\cdot j}$  if there are  $a_j \in \mathbb{R}$  and  $b_j \in \mathbb{R} \setminus \{0\}$ , such that  $Y'_{\cdot j} = a_j + b_j Y_{\cdot j}$ . In model II, the group is the affine group  $GA(\mathbb{R})^d, \mathcal{G} = \mathbb{R}^d \times D$  and  $D = \{\text{diag}\{b_1, \dots, b_d\} \mid b_i \neq 0, i = 1, \dots, d\}$  with the operation  $(\boldsymbol{a}_1, A_1) \circ (\boldsymbol{a}_2, A_2) = (\boldsymbol{a}_1 + A_1 \boldsymbol{a}_2, A_1 A_2)$  for  $\boldsymbol{a}_i \in \mathbb{R}^d, A_i \in D$  with  $i = 1, 2$ . The orbit of an element  $Y \in \mathbb{R}^{n \times d}$  and the group action are defined in (1) and (2) with  $A \in D$ . If

$$\text{Vec}(Y) \sim N(\mathbf{1}_n \otimes \boldsymbol{\mu}, (I_n + \theta B) \otimes \text{diag}\{\sigma_1^2, \dots, \sigma_d^2\})$$

then  $\text{Vec}((- \boldsymbol{\mu}, I_d) \star Y) \sim N(\mathbf{0}, (I_n + \theta B) \otimes \text{diag}\{\sigma_1^2, \dots, \sigma_d^2\})$ , and furthermore  $\text{Vec}((\boldsymbol{a}, A) \star Y) \sim N(\mathbf{0}, (I_n + \theta B) \otimes I_d)$  with  $\boldsymbol{a} = -(\mu_1/\sigma_1, \dots, \mu_d/\sigma_d)^\top$  and  $A = \text{diag}\{\sigma_1^{-1}, \dots, \sigma_d^{-1}\}$ , which correspond to elements of the group orbit. No linear combinations are permitted here, so that the integrity of the variables is preserved.

Moreover, in some cases, the location information or shapes of objects from aerial photography applications may be distorted by the viewer's angle or position so that the variables may be strongly correlated. A more extreme approach avoids the metric assumption by regarding  $Y$  and  $Y'$  as equivalent configurations if there exists a vector  $\boldsymbol{a} \in \mathbb{R}^d$  and a non-singular matrix  $A \in \mathbb{R}^{d \times d}$  such that  $Y'_i = \boldsymbol{a} + AY_i$  with  $A^\top A$  is a positive definite matrix for all  $i$ . Consequently, models I, II, III specify the structures of the covariance matrix between features, and the partition  $B$  of  $Y$  is affine invariant and the same as the partition  $B$  of the group orbit  $\mathcal{G}Y \subset \mathbb{R}^{n \times d}$ , which is independent of the mean.

### 3.1 Gaussian marginal probabilities

The distribution of the column-centered group orbit,  $\mathcal{G}Y$ , is assumed to be a Gaussian distribution

$$N(\mathbf{0}, (I_n + \theta B) \otimes A^\top A)$$

which depends only on  $I_n + \theta B$  and  $A^\top A$ . Actually, it can be verified that for any  $(\boldsymbol{a}, A) \in \mathcal{G}$ , the two distributions of group orbits induced by  $N(\mathbf{1}_n \otimes \boldsymbol{\mu}, (I_n + \theta B) \otimes \Sigma)$  and  $N(\mathbf{1}_n \otimes (\boldsymbol{a} + A\boldsymbol{\mu}), (I_n + \theta B) \otimes (A\Sigma A^\top))$  respectively are the same.

McCullagh (2008) studied the  $d$  time series with an autocorrelation  $\Gamma$  and  $n$  observations in time or space following three Gaussian distribution models  $N(\mathbf{0}, \Gamma \otimes \Sigma)$  under different assumptions of  $\Sigma$  as follows :

$$\text{Model I: } \Sigma = \sigma^2 I_d \tag{3}$$

$$\text{Model II: } \Sigma = \text{diag}\{\sigma_1^2, \dots, \sigma_d^2\} \tag{4}$$

$$\text{Model III: } \Sigma \in PD_d \tag{5}$$

where  $PD_d$  is the collection of  $d \times d$  symmetric positive definite matrices. These three models correspond to our three models of affine transformed equivalence classes which

we discussed in the previous section. In this paper, we set  $(I_n + \theta B)$  as  $\Gamma$  and  $A^\top A$  as  $\Sigma$ . Following (McCullagh 2008), the log-likelihood based on  $Y$  for all three models is:

$$\begin{aligned} l(\Gamma, \Sigma | Y) &= -\frac{1}{2} \log \det(\Gamma \otimes \Sigma) - \frac{1}{2} \text{tr}(Y^\top \Gamma^{-1} Y \Sigma^{-1}) \\ &= -\frac{d}{2} \log \det(\Gamma) - \frac{n}{2} \log \det(\Sigma) - \frac{1}{2} \text{tr}(Y^\top \Gamma^{-1} Y \Sigma^{-1}), \end{aligned}$$

**Lemma 1**  $(I_n + \theta B)^{-1} = I_n - \theta WB$ , where  $W = \text{diag}\{(1 + \theta N_1)^{-1}, \dots, (1 + \theta N_n)^{-1}\}$  and  $N_i$  is the  $i$ th diagonal element of  $N = B\mathbf{1}_n$ .

According to Lemma 1 and its proof, which is relegated to the Appendix A,  $\Gamma = I_n + \theta B$  is always nonsingular for  $\theta > 0$  and its inverse  $\Gamma^{-1} = (I_n + \theta B)^{-1} = I_n - \theta WB$  can be obtained explicitly. To ensure that  $Y^\top \Gamma^{-1} Y$  is positive definite with probability 1 (McCullagh 2008), as well as informative group orbits (see Theorem 1 and its subsequent discussion), we assume  $n > d + 1$ .

After plugging in the maximum likelihood estimator of  $\Sigma$  which for model III is  $\hat{\Sigma}_\Gamma = Y^\top \Gamma^{-1} Y / n$ , for model II is  $\text{diag}(\hat{\Sigma}_\Gamma)$ , and for model I is  $\text{tr}(\hat{\Sigma}_\Gamma) I_d / d$  (McCullagh 2008), the profile likelihood of  $\Gamma$  is

$$L_p(\Gamma^{-1} | \mathcal{G}Y) = \begin{cases} \det(\Gamma^{-1})^{d/2} / \text{tr}(Y^\top \Gamma^{-1} Y)^{nd/2} & (\text{I}) \\ \det(\Gamma^{-1})^{d/2} / \prod_{r=1}^d \left(Y_{(r)}^\top \Gamma^{-1} Y_{(r)}\right)^{n/2} & (\text{II}) \\ \det(\Gamma^{-1})^{d/2} / \det(Y^\top \Gamma^{-1} Y)^{n/2} & (\text{III}) \end{cases}$$

where  $Y_{(r)} \in \mathbb{R}^n$  is the  $r$ th column of  $Y$ ,  $r = 1, \dots, d$ .

The conditional distribution on partitions of  $[n]$  depends on the group orbit and the assumptions made regarding  $\Sigma$ . For group I, with  $\Sigma \propto I_d$  in the Gaussian model, the likelihood depends only on the distance matrix  $D$ , so the likelihood is constant on the orbits associated with the larger group of Euclidean similarities. Therefore, for model I, the similarity transformation can be generalized as if  $Y'_i = \mathbf{a} + AY_i$  for  $A^\top A = \sigma^2 I_d$  and  $\sigma \neq 0$ , implying that the arrays of distances are proportional  $D' = \sigma^2 D$ . Consequently, there is a representative element of the group orbit with feature mean vector  $\mathbf{0}$ , so that  $\text{Vec}(Y) \sim N(\mathbf{0}, (I_n + \theta B) \otimes \sigma^2 I_d)$ .

For model II, the affine transformation can be generalized as  $Y'_i = \mathbf{a} + AY_i$ , where  $\mathbf{a} \in \mathbb{R}^d$  and  $A \in \mathbb{R}^{d \times d}$  with  $A^\top A$  as a diagonal matrix with positive diagonal entries for all  $i$ . As a result, there is a representative element of the group orbit with feature mean vector  $\mathbf{0}$ , so that  $\text{Vec}(Y) \sim N(\mathbf{0}, (I_n + \theta B) \otimes \text{diag}\{\sigma_1^2, \dots, \sigma_d^2\})$ . This is to work with  $GA(\mathbb{R}^d)$  which is the general affine group acting independently on the  $d$  columns of  $Y$ . For model III,  $\Sigma$  is an arbitrary matrix in  $PD_d$ . The group is  $GA(\mathbb{R}^d)$  and  $n > d + 1$ . These three models are nested by model I  $\subset$  model II  $\subset$  model III.

Affine invariance in  $\mathbb{R}^d$  is a strong requirement, which comes at a small cost for moderate  $d$  provided that  $d/n$  is small. When  $d/n \leq 1$ ,  $Y^\top \Gamma^{-1} Y$  is positive definite with probability one (McCullagh 2008), then model III works. However, while  $d/n < 1$  is not small, model III may be inefficient due to some eigenvalues of  $Y^\top \Gamma^{-1} Y$  and  $\det(Y^\top \Gamma^{-1} Y)$  close to zero (Dempster 1972; Stein 1975). As a result, the profile likelihood of  $\Gamma$  becomes unstable. In contrast, model II is less computationally expensive than model III, and model I is the most efficient one.

#### 4 Markov chain Monte Carlo algorithm for sampling partitions

We use the prior and posterior distributions of  $\theta$  and  $B$  discussed in Section 2 through a Markov chain Monte Carlo (MCMC) algorithm for sampling partitions. The iterative  $\theta$  is obtained by Gibbs sampling (Geman and Geman 1984) according to the conditional distribution  $p_n(\theta_j|B, \mathcal{G}Y) \propto p(\theta_j) \times L_p(\Gamma^{-1}|\mathcal{G}Y)$ , where  $p(\theta_j) \propto \theta_j^{\alpha-1} / (1 + \theta_j)^{2\alpha}$  for  $j = 1, \dots, J$ . For instance,  $\alpha = 1$  and the discrete set  $\{2^{-3}, 2^{-2}, \dots, 2^{10}\}$  for the range of  $\theta$  are used as the default setting in our experiments. For updating  $B$ , the conditional distribution on partitions is

$$p_n(B|\theta, \mathcal{G}Y) \propto p_n(B|\lambda) \times L_p(\Gamma^{-1}|\mathcal{G}Y),$$

where  $p_n(B|\lambda)$  is the Ewens distribution, and a Metropolis-Hastings algorithm (Metropolis et al. 1953; Hastings 1970) is used to choose the iterative  $B$ .  $\lambda$  is set as 1 in the following applications. After burning in a certain number of the resulting Markov chain, we use the average of the partition matrix as the similarity matrix to make inference on partition. The proposal distribution  $q(B^{(i+1)}|B^{(i)}, \mathcal{G}Y)$  is proportional to  $\exp(-\alpha \times d_{xc})$ , where  $d_{xc}$  is the distance between each point and the corresponding centroid of the clusters and  $\alpha$  is a scale hyperparameter which was set as 2 in our experiments. More specifically, a partition candidate  $B^*$  is generated by re-assigning the label of each point with the probability proportional to the reciprocal of the distance between each point and the corresponding centroid.

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#### Algorithm 1 Metropolis-Hastings algorithm

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- 1: Initialize  $B$  and  $\theta$ .
- 2: **for**  $i = 1 : N$  **do**  $N$  is the number of total iterations. Suppose that the current values are  $\theta^{(i)}$  and  $B^{(i)}$ .
- 3:     Randomly sample  $\theta^{(i+1)}$  from  $p_n(\theta_j|B^{(i)}, \mathcal{G}Y), j = 1, \dots, J$ .
- 4:     Propose  $B^* \sim q(B^{(i+1)}|B^{(i)}, \mathcal{G}Y)$ .
- 5:     Calculate
$$R = \frac{p_n(B^*|\lambda) L_p(\Gamma(\theta^{(i+1)}, B^*)^{-1}|\mathcal{G}Y) q(B^{(i)}|B^*, \mathcal{G}Y)}{p_n(B^{(i)}|\lambda) L_p(\Gamma(\theta^{(i+1)}, B^{(i)})^{-1}|\mathcal{G}Y) q(B^*|B^{(i)}, \mathcal{G}Y)}.$$
- 6:     Accept  $B^{(i+1)} = B^*$  with probability  $\min\{1, R\}$
- 7:     Keep  $B^{(i+1)} = B^{(i)}$  with probability  $1 - \min\{1, R\}$
- 8: **end for**
- 9: **return** all the  $B^{(i)}$ 's and  $\theta^{(i)}$ 's.

---

Since Algorithm 1 is a Metropolis-Hastings algorithm, it satisfies the detailed balance condition, and therefore the generated Markov chain has a stationary distribution (Chib and Greenberg 1995; Gamerman 1997; Robert and Casella 2010). Since we leave a small but positive probability that the partition stays the same in the Gibbs sampling and the discrete posterior of  $\theta$  stays positive always, then the transition probability

$$p_n(\theta^{(k+1)}, B^{(k+1)}|\theta^{(k)}, B^{(k)}) > 0$$

where  $\theta^{(k+1)} = \theta^{(k)}$  and  $B^{(k+1)} = B^{(k)}$ , and then the  $(\theta, B)$ -valued Markov chain constructed by Algorithm 1 is aperiodic.

**Lemma 2** *If  $n > d + 1$ , the  $(\theta, B)$ -valued Markov chain constructed by Algorithm 1 is aperiodic.*

Since there is always a positive chance that the partition can be split further into the simplest partition in which each element is a cluster, then all possible partitions communicate with each other, so that the  $(\theta, B)$ -valued Markov chain constructed by Algorithm 1 is irreducible. Given the sample size  $n$ , the size of the state space of  $B$  known as the Bell number (Bell 1934), and the size of the state space of  $\theta$  are all finite, then the irreducibility also implies positive recurrence. Consequently, the  $(\theta, B)$ -valued Markov chain constructed by Algorithm 1 is ergodic (Isaacson and Madsen 1976; Gilks et al. 1996). The properties are summarized as the following lemma and theorem, whose proofs are relegated to the Appendix A.

**Lemma 3** *If  $n > d + 1$ , the  $(\theta, B)$ -valued Markov chain constructed by Algorithm 1 is irreducible, and thus is positive recurrent.*

**Theorem 2 (Ergodic theorem)** *If  $n > d + 1$ , the  $(\theta, B)$ -valued Markov chain constructed by Algorithm 1 converges to its stationary distribution  $p_n(\theta, B | \mathcal{G}Y) \propto p(\theta) \times p_n(B | \lambda) \times L_p(\Gamma^{-1} | \mathcal{G}Y)$ . More specifically, for any real-valued function  $f$  satisfying  $\sum_{(\theta, B)} |f(\theta, B)| p_n(\theta, B | \mathcal{G}Y) < \infty$ , we have*

$$\frac{1}{n+1} \sum_{i=0}^n f(\theta^{(i)}, B^{(i)}) \longrightarrow \sum_{(\theta, B)} f(\theta, B) p_n(\theta, B | \mathcal{G}Y)$$

*almost surely for all initial value  $(\theta^{(0)}, B^{(0)})$ .*

## 5 Analysis of simulated and real data

We test the proposed Bayesian cluster process with Algorithm 1 on both synthetic and real data. Algorithm 1 with model I and point-wise updating is equivalent to the method of (Vogt et al. 2010). If there is no prior information of the number of clusters, users can set the initial partition  $B$  as  $I_n$  in which each observation is a block. In practice, we use a randomly sampled clusters from a discrete uniform distribution of a range chosen by users. The clustering result is represented by the average of the estimated similarity matrix

$$S = \sum_{k=n_0+1}^N \frac{B^{(k)}}{N - n_0},$$

where  $n_0$  is the number of burn-in iterations. Furthermore, we also define a dissimilarity matrix  $D$  as  $\mathbf{1}_n \mathbf{1}_n^\top - S$ . The dissimilarity matrix,  $D$ , can be expressed by a heatmap which represents a matrix with grayscale colors with white as 1, black as 0, and the spectrum of gray as values between 0 and 1. The heatmap of the original similarity matrix cannot be recognized with the naked eye and equivalence relation needs to be decoded from the matrix  $B$ . However, in practice, users can identify clusters through including the names of rows and columns of the similarity matrix to find which individuals are clustered together. Additionally, the heatmap function of the stats R package can permute the order of individuals to have cluster blocks with hierarchical dendrograms. It is challenging to monitor convergence of the Markov chain because the sampled clusters are random and may vary

in each iteration. To determine convergence, we run Algorithm 1 ten times for each data set and stop the chain when we observe the number of clusters remain the same in the given chain length (Chang and Fisher 2013).

### 5.1 Illustrative simulated data

**Four clusters on the vertices of a unit square data** Three simulated data sets are generated for illustration. In the simulation study, 1000 initial burn-in iterations were discarded, and 2000 Markov chains of  $B$  samples based on each model were used to calculate  $D$ . We first applied the proposed cluster process with model I on the synthetic data for four clusters centered at the four vertices of a unit square. For each vertex  $\mu_k$ , we generate 20 points from  $N(\mu_k, (1/4)I_2)$  for  $k = 1, \dots, 4$  (see Fig. 1, the left panel). We call the data  $X_I$ , and then apply model I to cluster  $X_I$  with the average within- and between-cluster distances. The resulting heatmap successfully reveals the true clusters for most of the points (not shown here).

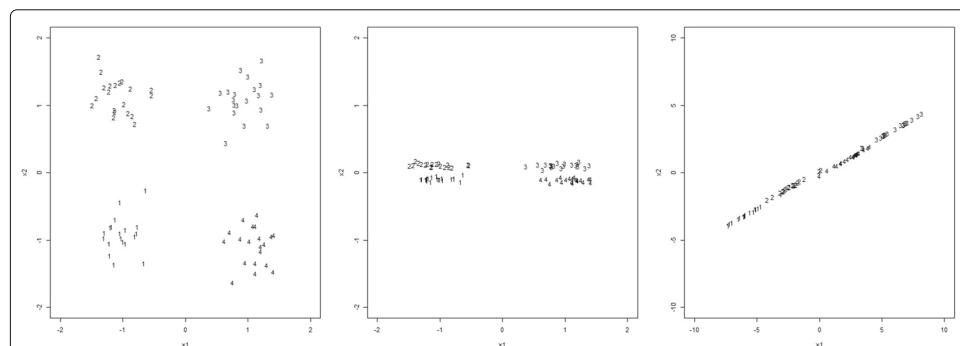
Then we transform the data by  $X_{II} = X_I \times \begin{pmatrix} 3 & 0 \\ 0 & 1/3 \end{pmatrix}$ . The transformed features seem to have two groups (see Fig. 1, the middle panel), clusters (1, 2) and clusters (3, 4). The cluster process with model I does not work well for this case, while the heatmap based on model II without knowing the transformation can reveal the true clusters for most of the points (not shown here).

Furthermore we transform the data by  $X_{III} = X_I \times \begin{pmatrix} 4.1 & 2.1 \\ 1.9 & 1.1 \end{pmatrix}$ . The transformed features are aligned in a straight line (see Fig. 1, the right panel). The transformed data  $X_{III}$  is more difficult to cluster than  $X_I$  and  $X_{II}$ , since the original four clusters are transformed to be not well separated.

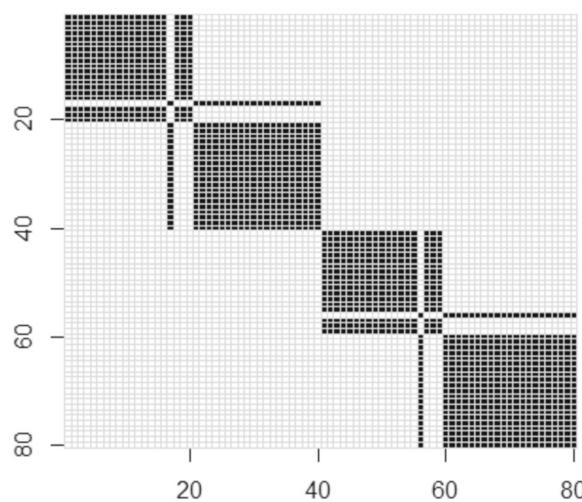
The resulting heatmap using model III with the initial clusters assigned randomly and uniformly from  $\{1, 2, 3, 4\}$  reveals the true four clusters for most of the points (see Fig. 2).

### 5.2 Applications to real data

Besides the synthetic data, we also evaluate the performance of the proposed approach by using real data. We run 3000 MCMC iterations and burn in the first 1000 iterations,



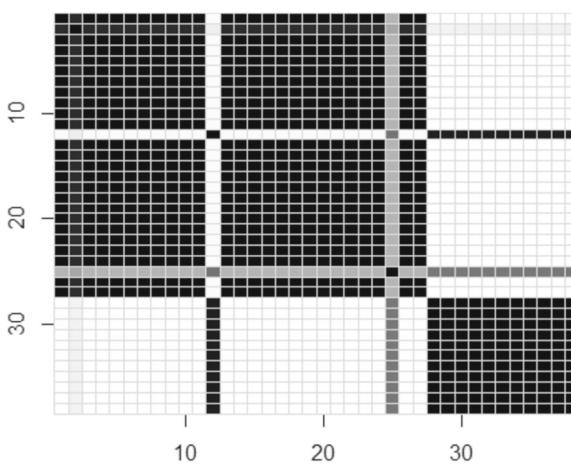
**Fig. 1** The scatter plots for  $X_I$ ,  $X_{II}$ , and  $X_{III}$  of the unit square synthetic data from the left to the right. The most left panel is the original features which have four clusters at the vertices of the unit square with equal size 20; the middle panel is the features which are transformed by scaling each dimension differently, clusters 1 and 2 are grouped as well as clusters 3 and 4 are grouped; the right panel shows the transformed features are aligned as a straight line



**Fig. 2** The heatmap of the similarity matrix using model III reveals the true four clusters for most of the transformed data  $X_{\parallel\parallel}$

and use the heatmap of matrix  $S$  to visualize the clusters. The accuracy rate is based on the average proportion of identical elements of matrix  $B$  of the cluster and the true matrix  $B$ , and compared the accuracy rates with  $k$ -means (MacQueen 1967) and Mclust using R package ‘mclust’ with its default setting (Fraley and Raftery 2002). The reason why we chose R package ‘mclust’ is that Mclust is a model-based clustering approach using the Gaussian mixture model, which assumes a Gaussian distribution for each component under one of the three types covariance structures (the argument of Mclust: modelName) 1. Spherical (EII), 2. Diagonal (VVI), and 3. General (VVV) for comparing with our proposed model I, II, III, correspondingly. The main difference is that the Mclust obtains clusters with an expectation–maximization (EM) algorithm (Dempster 1972; McLachlan and Peel 2000), but our method uses a Metropolis-Hastings algorithm with the profile likelihood of  $\Gamma$  to sample clusters.

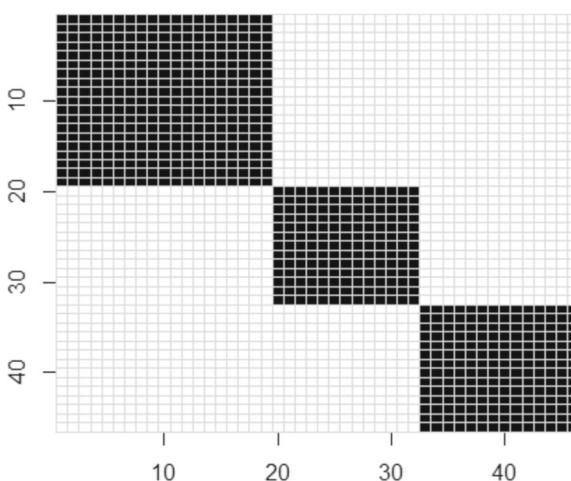
**Model I: Gene expression data of Leukemia patients** The gene expression microarray data (Dua and Graff 2019) has been used to study genetic disorder such as identifying diagnostic or prognostic biomarkers or clustering and classifying diseases (Dudoit et al. 2002). For example, (Golub et al. 1999) classified patients of acute leukemia into two sub types, Acute Lymphoblastic Leukemia (ALL) and Acute Myeloid Leukemia (AML). For illustration purpose, we use the training set of the leukemia data which consists of 3051 genes and 38 tumor mRNA samples. Pretending we do not know the label information, we would like to cluster the 38 samples according to their 3051 features (gene expression levels). The two clusters comprise 27 ALL cases and 11 AML cases. Since the number of features is larger than the sample size, our approach is not applicable to this dataset directly. Therefore, we first reduce the dimension by projecting the data on the subspace which consists of the first twenty principal components (PC) (Jolliffe 1986). Note that these PC scores are orthonormal which satisfies the assumption of model I. The resulting heatmap based on model I (Fig. 3) reveal the cluster of the 11 AML cases. The accuracy rate using the proposed model I with the initial clusters assigned randomly and uniformly from  $\{1, 2\}$  is 0.9164, while the accuracy rates of  $k$ -means and Mclust are 0.6994 and 0.5886, respectively. We noticed that Mclust resulted in only one cluster.



**Fig. 3** The heatmap of the similarity matrix using model I identifies the ALL group in the left upper corner and the AML group in the right bottom corner

#### Model II: Geographic coordinate data of Denmark's 3D Road Network

This three-dimensional road network dataset of geographic coordinates includes the altitude, latitude, and longitude degrees of each road segments in North Jutland in northern Denmark, which is publicly available at the UC Irvine Machine Learning Repository (Kaul 2013; Dua and Graff 2019). Since three spatial dimensional features are orthogonal, it satisfies the assumption of model II so that we use this dataset to demonstrate model II. Three subjects with the road maps OSM ID 144552912 (19 observations), 125829151 (13 observations), 145752974 (14 observations) are used for the clustering analysis. Note that each objects may have several observations measured from different angles, and the altitude values are extracted from NASA's Shuttle Radar Topography Mission (SRTM) data (Jarvis et al. 2008). The accuracy rate using model II with the initial clusters assigned randomly and uniformly from  $\{1, 2, 3, 4, 5\}$  is 1, while the accuracy rates of  $k$ -means with  $k = 3$  and Mclust are 0.7486 and 0.9490, respectively. The resulting heatmap using model II (Fig. 4) reveals 3 clusters correctly.



**Fig. 4** The heatmap of the similarity matrix using model II correctly reveals three clusters corresponding to the three buildings in the Denmark 3-D road map data

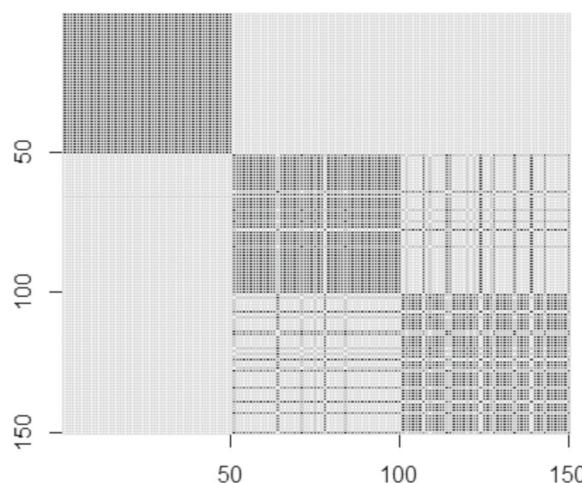
### Model III: Iris data

This iris dataset (Fisher 1936) contain three species—Setosa, Versicolor, and Virginica with four features which are the measurements of the variables sepal length and width and petal length and width in centimeters, respectively. Each species consists of 50 iris flowers. The data points are clustered by their four features. Here,  $d = 4$ ,  $n = 150$ ,  $k = 3$ . The heatmap of the similarity matrix using model III correctly reflects three clusters corresponding to the three species of iris for most points (Fig. 5). The accuracy rate using the proposed model III with the initial clusters assigned randomly and uniformly from  $\{1, 2, 3\}$  is 0.9087, while the accuracy rates of  $k$ -means with  $k = 3$  and Mclust are 0.7740 and 0.7763, respectively. We noticed that both the  $k$ -means and Mclust result in two clusters by grouping Versicolor and Virginica as a cluster.

## 6 Concluding discussion

The proposed clustering method is invariant under different groups of affine transformations and computationally efficient. It identifies clusters for most samples without knowing the number of clusters in advance, and it may group a big cluster as several small clusters. These problems are dealt with an exchangeable partition prior which avoids label-switching problems and the partition valued in the MCMC algorithm is invariant under linear transformations under three types of covariance structures. The advantage of replacing the Dirichlet-multinomial prior with its limiting process is that we do not need to know the number of clusters in advance. The disadvantage is that it may be less efficient computationally if the number of clusters is known. Note that the proposed approach does not target the partition maximizing the posterior distribution. Instead, it estimates the expected partition or the similarity matrix.

The three clustering models are based on the covariance matrix between variables. There are guidelines of telling which model work best in practice by the experimental design or testing its sample covarinace matrix. If the features are othornormal or orthogonal, then model I and model II are applicable, respectively. Models I and II run faster than model III due to the structure of the covariance matrix. Otherwise, model III can be used in general. It works reasonably well across various applications.



**Fig. 5** The heatmap of the similarity matrix using model III correctly reveals three clusters corresponding to the three species of iris for most points

Since we use the profile likelihood of  $\Gamma$  in our model, we do not sample the covariance matrix directly, and Lemma 1 and Theorem 1 implies as  $n > d + 1$ , the proposed Metropolis-Hastings algorithm can work. However, the maximum likelihood estimator (MLE) of the general unstructured covariance matrix will be less efficient if the diagonal covariance structure is actually correct because it will tend to have small eigenvalues and a large determinant of the inverse covariance matrix (i.e.  $\Gamma^{-1}$ ). Indeed, when using model III and even if  $n > d + 1$ ,  $\Gamma$  may be near singular. This may make the sampling less efficient. i.e. the acceptance rate may become small (Roberts and Rosenthal 2001). Although the stationary distribution of the sampled clusters' Markov chain using Algorithm 1 is independent of the initial clusters according to Theorem (2), we practically suggest to set the initial clusters sampled from a discrete uniform distribution of a range given by users instead of setting each individual as a cluster in order to obtain convergent sampled clusters without using a long Markov chain. This makes the proposed Algorithm 1 sample more efficiently from a smaller collection of partition candidates.

The proposed clustering algorithm produces the desired clusters with 2000 iterations after 1000 burn-in iterations in our experiments. The main contributions of our work include: 1) The proposed three clustering models with three types of covariance structures can handle general cases of affine transformations. In contrast, (Vogt et al. 2010) only dealt with the case of model I. 2) Algorithm 1 is efficient, since it updates all individuals' clusters instead of a single individual's cluster per iteration. It also ensures that the resulting partition-valued Markov chain is ergodic and convergent in distribution. 3) The experiments show the advantages of our cluster process which successfully identifies the true clusters using the proposed distance matrix. In particular if the clusters are not well separated, the similarity matrix with probabilistic nature can still reveal the relationships through hierarchical approaches. The proposed method could be used to extract interesting information from aerial photography, genomic data, and data with attributes under different scales, especially when the nearest neighbors may belong to different clusters in the feature space. The proposed method can be improved in the further work by modeling the mean of each cluster with regression on covariates or non-Gaussian distributions.

## Appendix A

**Proof of Theorem 1:** For any  $Y \in \mathbb{R}^{n \times d}$ , denote  $\tilde{Y} = Y - \mathbf{1}_n \mathbf{1}_n^\top Y/n$ . Let  $\tilde{Y}_{(n-1)}$  be the  $(n-1) \times d$  matrix consisting of the first  $n-1$  rows of  $\tilde{Y}$ . Since  $\mathbf{1}_n^\top \tilde{Y} = 0$ , then  $\text{rank}(\tilde{Y}) = \text{rank}(\tilde{Y}_{(n-1)})$ .

If  $n \leq d+1$  and  $\text{rank}(\tilde{Y}_{(n-1)}) = n-1$ , that is,  $\tilde{Y}_{(n-1)}$  is of full row rank, then there exists an orthogonal matrix  $O \in \mathbb{R}^{d \times d}$  (column permutations), such that,  $\tilde{Y}_{(n-1)} O = (U, V)$ , where  $U \in \mathbb{R}^{(n-1) \times (n-1)}$  is of full rank, and  $V \in \mathbb{R}^{(n-1) \times (d+1-n)}$ . We let

$$\boldsymbol{\alpha} = \frac{1}{n} Y^\top \mathbf{1}_n, \quad A = O \begin{pmatrix} U^\top & \mathbf{0} \\ V^\top & I_{d+1-n} \end{pmatrix}, \quad Z = \begin{pmatrix} I_{n-1} & \mathbf{0} \\ -\mathbf{1}_{n-1}^\top & \mathbf{0} \end{pmatrix}.$$

It can be verified that  $Y = (\boldsymbol{\alpha}, A) \star Z$ . That is,  $Y \in \text{Orb}(Z)$ , where  $Z$  is a constant matrix.

If  $n \leq d$  and  $\text{rank}(Y) = n$ , then  $\text{rank}(\tilde{Y}_{(n-1)}) = n-1$ , since  $\tilde{Y}_{(n-1)} = WY$ , where  $W = (I_{n-1} - \mathbf{1}_{n-1} \mathbf{1}_{n-1}^\top/n, -\mathbf{1}_{n-1}/n)$  is of full row rank  $n-1$ .

Suppose  $n = d+1$  and  $\text{rank}(Y) = d = n-1$ . Without any loss of generality, we assume  $\text{rank}(\tilde{Y}_{(n-1)}) = n-1$ , where  $\tilde{Y}_{(n-1)}$  consists of the first  $n-1$  rows of  $Y$ . Then  $Y_n = c_1 Y_1 + \cdots + c_{n-1} Y_{n-1}$  for some  $c_1, \dots, c_{n-1} \in \mathbb{R}$  and  $\tilde{Y}_{(n-1)} = DY_{(n-1)}$ , where

$D = I_{n-1} - \mathbf{1}_{n-1}\mathbf{1}_{n-1}^\top/n - \mathbf{1}_{n-1}\mathbf{c}^\top/n$ , and  $\mathbf{c}^\top = (c_1, \dots, c_{n-1})$ . It can be verified that if  $c_1 + \dots + c_{n-1} \neq 1$ , then  $\text{rank}(D) = n-1$  and  $\text{rank}(\tilde{Y}_{(n-1)}) = n-1$ ; if  $c_1 + \dots + c_{n-1} = 1$ , then  $\text{rank}(D) = n-2$  and  $\text{rank}(\tilde{Y}_{(n-1)}) = n-2$ . Note that if  $n = d+1$  but  $\text{rank}(\tilde{Y}_{(n-1)}) = n-2$ , then  $Y \notin \text{Orb}(Z)$ .  $\square$

**Proof of Lemma 1:** Suppose the partition matrix  $B$  consists of  $k$  blocks with block sizes  $n_1, \dots, n_k$ , where  $k \geq 1$ ,  $n_i > 0$  for  $i = 1, \dots, k$ , and  $n_1 + \dots + n_k = n$ .

We first assume that  $B = \text{diag}\{\mathbf{1}_{n_1}\mathbf{1}_{n_1}^\top, \dots, \mathbf{1}_{n_k}\mathbf{1}_{n_k}^\top\}$ , which is in its standard form. Then  $B = LL^\top$  with  $L = \text{diag}\{\mathbf{1}_{n_1}, \dots, \mathbf{1}_{n_k}\} \in \mathbb{R}^{n \times k}$  and  $I_n + \theta B = I_n + EE^\top$  with  $E = \sqrt{\theta}L$ .

According to the Sherman-Morrison-Woodbury formula (see, for example, Section 2.1.4 in Golub and Van Loan (2013)), for matrices  $A \in \mathbb{R}^{n \times n}$  and  $U, V \in \mathbb{R}^{n \times k}$ ,  $(A + UV^\top)^{-1} = A^{-1} - A^{-1}U(I + V^\top A^{-1}U)^{-1}V^\top A^{-1}$  if both  $A$  and  $I + V^\top A^{-1}U$  are nonsingular. In our case,  $A = I_n$  is nonsingular,  $U = V = E$ , and  $I + V^\top A^{-1}U = I_k + E^\top E = \text{diag}\{1 + \theta n_1, \dots, 1 + \theta n_k\}$  is also nonsingular. Thus

$$\begin{aligned} & (I_n + EE^\top)^{-1} \\ &= I_n - E(I_k + E^\top E)^{-1}E^\top \\ &= I_n - \theta L(I_k + \theta L^\top L)^{-1}L^\top \\ &= I_n - \theta \cdot \text{diag}\left\{\frac{1}{1 + \theta n_1}\mathbf{1}_{n_1}\mathbf{1}_{n_1}^\top, \dots, \frac{1}{1 + \theta n_k}\mathbf{1}_{n_k}\mathbf{1}_{n_k}^\top\right\} \\ &= I_n - \theta \cdot \text{diag}\left\{\frac{1}{1 + \theta n_1}I_{n_1}, \dots, \frac{1}{1 + \theta n_k}I_{n_k}\right\} \cdot \text{diag}\left\{\mathbf{1}_{n_1}\mathbf{1}_{n_1}^\top, \dots, \mathbf{1}_{n_k}\mathbf{1}_{n_k}^\top\right\} \\ &= I_n - \theta WB. \end{aligned}$$

In general, by row-switching and column-switching transformations, we can always transform  $B$  into its standard form. That is, there exists an orthogonal matrix  $O$  such that  $B_r = OBO^\top$  is in standard form. Let  $W_r = OWOT$ . Then  $(I_n + \theta B)^{-1} = O^\top(I_n + \theta B_r)^{-1}O = O^\top(I_n - \theta W_r B_r)O = I_n - \theta \cdot O^\top W_r O \cdot O^\top B_r O = I_n - \theta WB$ .  $\square$

**Proof of Lemma 3:** In our case, the Markov chain built by Algorithm 1 is actually a discrete chain. It is irreducible since  $p_n(\theta^{(k+1)}, B^{(k+1)} | \theta^{(k)}, B^{(k)}) > 0$  for each pair of states. As a direct conclusion of Theorem 4.1 in Gilks et al. (1996), our Markov chain is positive recurrent.  $\square$

**Proof of Theorem 2:** Algorithm 1 is a Gibbs sampler plus a Metropolis-Hastings component for sampling  $B^{(i+1)}$ . Given  $B^{(i)}$  and  $\theta^{(i+1)}$ , the Metropolis-Hastings ratio with proposal distribution  $q(B|B^{(i)}, \mathcal{G}Y)$  and target distribution  $p_n(\theta, B|\mathcal{G}Y)$  is

$$\begin{aligned} R(B^{(i)}, B^*) &= \frac{p_n(\theta^{(i+1)}, B^* | \mathcal{G}Y) \cdot q(B^{(i)} | B^*, \mathcal{G}Y)}{p_n(\theta^{(i+1)}, B^{(i)} | \mathcal{G}Y) \cdot q(B^* | B^{(i)}, \mathcal{G}Y)} \\ &= \frac{p(\theta^{(i+1)}) \cdot p_n(B^* | \lambda) \cdot L_p(\Gamma(\theta^{(i+1)}, B^*)^{-1} | \mathcal{G}Y) \cdot q(B^{(i)} | B^*, \mathcal{G}Y)}{p(\theta^{(i+1)}) \cdot p_n(B^{(i)} | \lambda) \cdot L_p(\Gamma(\theta^{(i+1)}, B^{(i)})^{-1} | \mathcal{G}Y) \cdot q(B^* | B^{(i)}, \mathcal{G}Y)} \\ &= \frac{p_n(B^* | \lambda) \cdot L_p(\Gamma(\theta^{(i+1)}, B^*)^{-1} | \mathcal{G}Y) \cdot q(B^{(i)} | B^*, \mathcal{G}Y)}{p_n(B^{(i)} | \lambda) \cdot L_p(\Gamma(\theta^{(i+1)}, B^{(i)})^{-1} | \mathcal{G}Y) \cdot q(B^* | B^{(i)}, \mathcal{G}Y)} \end{aligned}$$

which is exactly  $R$  in Algorithm 1. Since Metropolis-Hastings algorithms satisfy detailed balance condition, the target distribution  $p_n(\theta, B|\mathcal{G}Y)$  is a stationary distribution. By Lemmas 2 and 3, the convergence statements follow as a direct conclusion of Theorems 4.3 and 4.4 in Gilks et al. (1996).  $\square$

**Appendix B journal name abbreviations for use in *Boundary-Layer meteorology***

Journal Name	Abbreviation used in BLM
ACM Transactions of Mathematical Software	ACM Trans Math Soft
Acoustics Australia	Acoust Aust
Acta Geophysica	Acta Geophys
Acta Mechanica Sinica	Acta Mech Sinica
Acta Mechanica Supplement	Acta Mech Suppl
Advances in Atmospheric Science	Adv Atmos Sci
Advances in Ecological Research	Adv Ecol Res
Advances in Meteorology	Adv Meteorol
Advances in Science and Research	Adv Sci Res
Advances in Water Resources	Adv Water Resour
Aeolian Research	Aeolian Res
Aerospace Science and Technology	Aerosp Sci Technol
Agricultural Meteorology	Agric Meteorol
Agricultural and Forest Meteorology	Agric For Meteorol
Agricultural Water Management	Agric Water Manag
American Institute of Aeronautics and Astronautics	Am Inst Aeronaut Astronaut
Annals of Glaciology	Ann Glaciol
Annalen der Meteorologie	Ann Meteorol
Annals of Statistics	Ann Stat
Antarctic Science	Antarct Sci
Annual Review of Fluid Mechanics	Annu Rev Fluid Mech
Applied Energy	Appl Energy
Applied Mechanics Review	Appl Mech Rev
Applied Numerical Mathematics	Appl Numer Math
Applied Physics B	Appl Phys B
Applied Optics	Appl Opt
Aquatic Botany	Aquat Bot
Archiv fur Meteorologie Geophysik und Bioklimatologie Serie A-Meteorologie und Geophysik	Arch Meteorol Geophys Bioklim Ser A
Archiv fur Hydrobiologie	Arch Hydrobiol
Artificial Intelligence	Artif Intell
Astronomy & Astrophysics	Astron Astrophys
Atmospheric Measurement Techniques	Atmos Meas Tech
Atmosphere-Ocean	Atmos-Ocean

Journal Name	Abbreviation used in BLM
Atmospheric Research	Atmos Res
Atmospheric Science Letters	Atmos Sci Lett
Australian Journal of Physics	Aust J Phys
Australian Journal of Botany	Aust J Bot
Beitrag zur Physik der Atmosphäre	Beitr Phys Atmos
Biogeosciences	Biogeosciences
Biometrika	Biometrika
Biosystems Engineering	Biosyst Eng
Boreal Environment Research	Boreal Environ Res
Boundary-Layer Meteorology	Boundary-Layer Meteorol
Building and Environment	Build Environ
Bulletin of the American Meteorological Society	Bull Am Meteorol Soc
Climate Research	Clim Res
Cold Regions Science and Technology	Cold Reg Sci Technol
Communications in Agricultural and Applied Biological Sciences	Commun Agric Appl Biol Sci
Communications in Mathematical Physics	Commun Math Phys
Communications on Pure and Applied Mathematics	Commun Pure Appl Math
Comptes Rendus Physique	C R Phys
Computers and Electronics in Agriculture	Comput Electron Agric
Computing and Informatics	Comput Inf
Computer Methods in Applied Mechanical Engineering	Comput Methods Appl Mech Eng
Computational Statistics and Data Analysis	Comput Stat Data Anal
Contributions to Atmospheric Physics	Contr Atmos Phys
Crop Protection	Crop Prot
Deep Sea Research Part II	Deep Sea Res II
Dynamics of Atmospheres and Oceans	Dyn Atmos Oceans
Earth System Science Data Discussions	Earth Syst Sci Data Discuss
Earth Surface Processes and Landforms	Earth Surf Process Landf
Ecological Applications	Ecol Appl
Ecological Indicators	Ecol Indic
Ecological Modelling	Ecol Model
Ecology	Ecology

Journal Name	Abbreviation used in BLM
Electronic Journal of Operational Meteorology	Electron J Oper Meteorol
Energies	Energies
Energy and Buildings	Energy Buil
Energy Conversion and Management	Energy Convers Manag
Environmental Fluid Mechanics	Environ Fluid Mech
Environmental Modeling and Software	Environ Modell Softw
Environmental Pollution	Environ Pollut
Environmental Research Letters	Environ Res Lett
Environmental Science and Technology	Environ Sci Technol
Environmental Software	Environ Softw
Eos, Transactions, American Geophysical Union	Eos Trans AGU
European Journal of Forest Research	Eur J For Res
Experiments in Fluids	Exp Fluids
Fisheries Research	Fish Res
Flow Turbulence and Combustion	Flow Turbul Combust
Forestry	Forestry
Freshwater Biology	Freshwater Biol
Functional Ecology	Funct Ecol
Acta Geodaetica et Geophysica Hungarica	Geod Geophys
Geografiska Annaler Series A	Geogr Ann Ser A
Geography Compass	Geogr Compass
Geomorphology	Geomorphology
Geophysical Research Letters	Geophys Res Lett
Geoscientific Instrumentation, Methods and Data Systems	Geosci Instrum Method Data Syst
Geoscientific Model Development	Geosci Model Dev
Global Biogeochemical Sciences	Glob Biogeochem Cycles
Glocal Change Biology	Glob Change Biol
Hydrology and Earth System Sciences	Hydrol Earth Syst Sci
Hydrological Processes	Hydrol Proc
IEEE Journal of Ocean Engineering	IEEE J Ocean Eng
IEEE Transactions on Geoscience and Remote Sensing	IEEE Trans Geosci Remote
International Journal of Climatology	Int J Climatol
International Journal of Wildland Fire	Int J Wildland Fire
International Journal of Heat and Fluid Flow	Int J Heat Fluid Flow

Journal Name	Abbreviation used in BLM
International Journal of Numerical Methods for Fluids	Int J Numer Methods Fluids
International Journal of Remote Sensing	Int J Remote Sens
Izvestiya, Atmospheric and Oceanic Physics	Izv Atmos Ocean Phys
Journal of Advances in Modeling Earth Systems	J Adv Model Earth Syst
Journal of Aerosol Science	J Aerosol Sci
Journal of Agricultural Engineering Research	J Agric Eng Res
Journal of the Air Pollution Control Association	J Air Pollut Control Assoc
Journal of Aircraft	J Aircr
Journal of Applied Meteorology and Climatology	J Appl Meteorol Clim
Journal of Applied Meteorology	J Appl Meteorol
Journal of Aquatic Plant Management	J Aquat Plant Manag
Journal of Arid Environments	J Arid Environ
Journal of Atmospheric and Oceanic Technology	J Atmos OceanTechnol
Journal of Atmospheric Science	J Atmos Sci
Journal of Climate	J Clim
Journal of Computational Physics	J Comput Phys
Journal of Earth Simulation	J Earth Simul
Journal of Earth System Science	J Earth Syst Sci
Journal of Environmental Engineering	J Environ Eng
Journal of Experimental Botany	J Exp Bot
Journal of the Faculty of Science Hokkaido University	J Fac Sci Hokkaido Univ
Journal of Field Robotics	J Field Robot
Journal of Fluid Mechanics	J Fluid Mech
Journal of Geophysical Research	J Geophys Res
Journal of Geophysical Research-Atmospheres	J Geophys Res Atmos
Journal of Glaciology	J Glaciol
Journal of Great Lakes Research	J Great Lakes Res
Journal of Hazardous Materials	J Hazard Mater A
Journal of Heat Transfer	J Heat Transf
Journal of Hydraulic Engineering	J Hydraul Eng
Journal of Hydrology	J Hydrol
Journal of Hydrometeorology	J Hydrometeorol
Journal of Marine Research	J Mar Res
Journal of Marine Systems	J Mar Syst
Journal de Mathematiques Pures et Appliquees	J Math Pures Appl

Journal Name	Abbreviation used in BLM
Journal of Meteorology	J Meteorol
Journal of the Meteorological Society of Japan	J Meteorol Soc Jpn
Journal of Oceanography	J Oceanogr
Journal of Operational Oceanography	J Oper Oceanogr
Journal of the operational Research Society	J Oper Res Soc
Journal of the Optical Society of America	J Opt Soc Am
Journal of Plankton Research	J Plankton Res
Journal of Solar Energy Engineering	J Sol Energy Eng
Journal of Quantitative Spectroscopy and Radiative Transfer	J Quant Spectrosc Radiat Transf
Journal of Renewable and Sustainable Energy	J Renew Sust Energy
Journal of Scientific Statistical Computing	J Sci Stat Comput
Journal of Statistical Physics	J Stat Phys
Journal of Thermophysics and Heat Transfer	J Thermophys Heat Transf
Journal of Tropical Ecology	J Trop Ecol
Journal of Turbulence	J Turbul
Journal of Wind Engineering and Industrial Aerodynamics	J Wind Eng Ind Aerodyn
Landscape and Urban Planning	Landsc Urban Plan
Limnology and Oceanography	Limnol Oceanogr
Low Temperature Science	Low Temp Sci
Machine Learning	Mach Learn
Marine Chemistry	Mar Chem
Mathematische Annalen	Math Ann
Meteorological Applications	Meteorol Appl
Meteorology and Atmospheric Physics	Meteorol Atmos Phys
Meteorologische Zeitschrift	Meteorol Z
Monthly Weather Review	Mon Weather Rev
Natural hazards and Earth System Sciences	Nat Hazards Earth Syst Sci
Nature Climate Change	Nat Clim Change
Nature Letters	Nat Clim Change
Nature Geoscience	Nat Geosci
Neural Computation	Neural Comput
Nonlinear Processes in Geophysics	Nonlin Process Geophys
New Zealand Journal of Science	N Z J Sci
Oceanography	Oceanography
Ocean Dynamics	Ocean Dyn
Ocean Engineering Science	Ocean Eng Sci

Journal Name	Abbreviation used in BLM
Ocean Modeling	Ocean Model
Papers in Physical Oceanography and Meteorology	Pap Phys Oceanogr Meteorol
Particle & Particle Systems Characterization	Part Syst Charact
Particuology	Particuology
Philosophical Transactions of the Royal Society of London	Philos Trans R Soc
Photogrammetric Engineering and Remote Sensing	Photogramm Eng Remote Sens
Physical Review Letters	Phys Rev Lett
Physical Review E	Phys Rev E
Physics and Chemistry of the Earth	Phys Chem Earth
Physics of Fluids	Phys Fluids
Physics A - Statistical Mechanics and its Applications	Physica A Stat Mech Appl
Physica D	Physica D
Plant Biosystems	Plant Biosyst
PLOS One	PLOS One
Powder technology	Powder Technol
Proceedings of the Royal Society	Proc Roy Soc
Progress in Aerospace Science	Prog Aerosp Sci
Progress in Heat and Mass Transfer	Prog Heat Mass Transf
Progress in Physical Geography	Prog Phys Geogr
Pure and Applied Geophysics	Pure Appl Geophys
Quarterly Journal of the Royal Meteorological Society	Q J R Meteorol Soc
Remote Sensing	Remote Sens
Remote Sensing of Environment	Remote Sens Environ
Renewable Energy	Renew Energy
Reviews of Geophysics	Rev Geophys
Reviews of Geophysics and Space Physics	Rev Geophys Space Phys
Review of Scientific Instruments	Rev Sci Inst
Science	Science
Science of the Total Environment	Sci Tot Environ
Sedimentology	Sedimentol
Siam Journal on Applied Mathematics	SIAM J Appl Math
Tellus	Tellus
Tellus Series B - Chemical and Physical Meteorology	Tellus Ser B Chem Phys Meteorol

Journal Name	Abbreviation used in BLM
Theoretical and Applied Climatology	Theor Appl Climatol
Theoretical and Computational Fluid Dynamics	Theor Comput Fluid Dyn
Theoretical Computational Fluid Dynamics	Theor Comput Fluid Mech
Thermal Science Engineering	Therm Sci Eng
Transactions of the American Society of Agricultural Engineers	Trans ASAE
Tree Physiology	Tree Physiol
Trudy Geofizicheskogo Instituta, Akademiya Nauk SSSR	Trudy Geofiz Inst AN SSSR
Urban Climate	Urban Clim
Water Air and Soil Pollution	Water Air Soil Pollut
Water Resources Research	Water Resour Res
Waterway Port Coastal and Ocean Engineering	Waterw Port Coast Ocean Eng
Weather	Weather
Weather and Forecasting	Weather Forecast
Wind Energy	Wind Energy
Wind Engineering	Wind Eng
Zeitschrift für Angewandte Mathematik und Mechanik	Z Agnew Math Mech

#### Abbreviations

ALL: Acute lymphoblastic leukemia; AML: Acute myeloid leukemia; MCMC: Markov chain Monte Carlo; PC: Principal components PC; SRTM: NASA's shuttle radar topography mission

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#### Authors' contributions

Hsin-Hsiung Huang wrote the draft of the manuscript, developed the algorithms, and conducted the experiments. Jie Yang proposed the methods and the initial algorithms. Both authors read and approved the final manuscript.

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#### Competing interests

The authors declare that they have no competing interests.

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