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A GEOMETRIC LEARNING APPROACH ON THE SPACE OF COMPLEX COVARIANCE MATRICES

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ABSTRACT

Many signal and image processing applications, including SAR polarimetry and texture analysis, require the classification of complex covariance matrices. The present paper introduces a geometric learning approach on the space of complex covariance matrices based on a new distribution called Riemannian Gaussian distribution. The proposed distribution has two parameters, the centre of mass \bar{Y} and the dispersion parameter σ . After having derived its maximum likelihood estimator and its extension to mixture models, we propose an application to texture recognition on the VisTex database.

Index Terms— Classification, Hermitian matrices, Information geometry, Riemannian centre of mass, EM algorithm.

1. INTRODUCTION

Covariance matrices are used as features for many signal and image processing applications, including biomedical image segmentation [1], radar detection [2,3], texture analysis and scene recognition [4–6], etc. These covariance matrices are used to model different kinds of dependence such as the spatial, spectral and temporal ones to cite a few of them. An interesting space of covariance matrices is the manifold \mathcal{H}_m of $m \times m$ Hermitian positive definite matrices. This latter is involved for example in radar detection, polarimetric SAR image classification [7], texture recognition [8], etc.

Many works have been dedicated in the literature for the statistical modelling of covariance matrices. Due to its mathematical tractability, the Wishart distribution is certainly the most largely used model in the literature [9,10]. Nevertheless, this model assumes Gaussian statistics for the observations which may not be realistic in practice. More advanced models have hence been proposed based on the so-called scalar product model. These compound models include the \mathcal{K} [11], \mathcal{G}^0 [12] and KummerU [13] distributions. They have shown promising results notably for the classification of high resolution polarimetric SAR images. Inspired from clustering approaches on Riemannian manifolds [14, 15], there is another

way to model covariance matrices. By considering Rao's distance on the manifold of covariance matrices, there is a canonical way to define the mean or barycentre of several covariance matrices in this manifold. Based on this concept, the Riemannian Gaussian distribution (RGD) has been introduced to model the statistical variability of real covariance matrices [16]. This distribution is represented by two parameters: its unique mode \bar{Y} (the unique Riemannian centre of mass) and σ its dispersion away from \bar{Y} . An important property of the RGD is that the maximum likelihood estimates of its parameters admit simple expressions and can be computed numerically using different algorithms. However, this RGD has only been proposed on the space \mathcal{P}_m of $m \times m$ real symmetric and strictly positive definite matrices. The main contribution of the paper is to extend this RGD to the space of Hermitian positive definite matrices. Moreover, based on mixture models of RGDs, a new geometric learning approach on \mathcal{H}_m is proposed. This approach is highlighted by an application on the classification of texture images.

The paper is structured as follows. Section 2 recalls some elements on the Riemannian geometry of \mathcal{H}_m . Section 3 introduces the proposed Riemannian Gaussian distribution on \mathcal{H}_m and its extension to mixture models. An expectation-maximization (EM) algorithm is presented to estimate the distribution parameters. Section 4 presents an application to image classification on the VisTex database. Some comparisons with the Wishart model are given to evaluate the potential of the proposed model in the context of texture image classification. Conclusions and future works are finally discussed in Section 5. Note that due to the restriction length, most of the mathematical proofs can not be detailed here and will be given in a forthcoming journal paper.

2. RIEMANNIAN GEOMETRY OF \mathcal{H}_m

This section recalls basic facts on the Riemannian geometry of \mathcal{H}_m , the space of Hermitian positive definite matrices of size $m \times m$. In many applications, \mathcal{H}_m is considered as a Riemannian manifold, equipped with the so-called affine-invariant Riemannian metric. For $Y \in \mathcal{H}_m$, this metric is given by the scalar product

$$g_Y(A, B) = \text{tr}(Y^{-1}AY^{-1}B)$$
 (1)

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where A, B are two Hermitian matrices of size $m \times m$. The Riemannian distance associated to q is the following

$$d(Y,Z) = \sqrt{\operatorname{tr}\left[\log\left(Y^{-1/2}ZY^{-1/2}\right)\right]^2} \tag{2}$$

For more details, the interested reader is referred to [17–19].

The reason for using the Riemannian metric (1) with the resulting Riemannian distance (2) is that these provide particularly suitable mathematical properties [17, 18, 20] which

- (i) invariance by all affine transformations.
- (ii) existence and uniqueness of Riemannian barycentres.

First, let us consider the property of invariance by affine transformations [17,18]. An affine transformation of \mathcal{H}_m is a mapping $Y \mapsto G \cdot Y$, where G is an invertible complex matrix of size $m \times m$. This is given by

$$G \cdot Y = GYG^* \tag{3}$$

where * denotes the conjugate transpose. With regard to the Riemannian distance (2), invariance by affine transformations means that for all $Y, Z \in \mathcal{H}_m$

$$d(Y,Z) = d(G \cdot Y, G \cdot Z) \tag{4}$$

where $G \cdot Y$ and $G \cdot Z$ are defined by (3).

Affine transformations are of special importance in the study of the space \mathcal{H}_m since for any $Y \in \mathcal{H}_m$ there exists G such that $Y = G \cdot I_m$ where $I_m \in \mathcal{H}_m$ is the identity matrix. Replacing this transformation in (4) gives the following identity

$$d(Y,Z) = d(I_m, G^{-1} \cdot Z) \tag{5}$$

This will be used in the next section.

Consider the second property of existence and uniqueness of Riemannian barycentres [20]. The barycentre of N points $Y_1, \cdots, Y_N \in \mathcal{H}_m$ is defined to be

$$\hat{Y}_N = \underset{Y}{\arg\min} \sum_{n=1}^N d^2(Y, Y_n)$$
 (6)

where $d(Y, Y_n)$ is the Riemannian distance (2).

As a Riemannian manifold with non-positive sectional curvature [21], \mathcal{H}_m enjoys the important property of existence and uniqueness of barycentres as defined by (6) [20].

Before going on, let us consider the concept of invariant integration on \mathcal{H}_m . The invariant integral of a function f: $\mathcal{H}_m \to \mathbb{R}$ is given by

$$\mathcal{H}_{m} \to \mathbb{R} \text{ is given by}$$

$$\int_{\mathcal{H}_{m}} f(Y) dv(Y) = \int \dots \int f(Y) \det(Y)^{-m} \prod_{i \leq j} d\operatorname{Re} Y_{ij} \prod_{i < j} d\operatorname{Im} Y_{ij}$$

$$\zeta(\sigma) \text{ can be expressed as}$$

$$\zeta(\sigma) = C \times \int_{\mathbb{R}^{m}} e^{-|r|^{2}/2\sigma^{2}} \prod_{i < j} \sinh^{2}(|r_{i} - r_{j}|/2) dr$$

$$(13)$$

where the subscripts i and j denote matrix elements, while Re and Im denote respectively the real and imaginary parts. In (7), dv(Y) is the Riemannian volume on \mathcal{H}_m defined by

$$dv(Y) = \det(Y)^{-m} \prod_{i \le j} d\operatorname{Re} Y_{ij} \prod_{i < j} d\operatorname{Im} Y_{ij}$$
 (8)

The integral (7) is called an invariant integral because of the following property

$$\int_{\mathcal{H}_m} f(Y)dv(Y) = \int_{\mathcal{H}_m} f(G \cdot Y)dv(Y) \tag{9}$$

where $G \cdot Y$ is the affine transformation given by (3) and G is any invertible complex matrix.

The invariant integral (7) takes on a simplified form when the function f is a central function. This means that f(Y) only depends on the eigenvalues of Y. Precisely, let the spectral decomposition of Y be given by $Y = U \operatorname{diag}(e^{r_1}, \cdots, e^{r_m}) U^*$, where U is a unitary matrix and e^{r_1}, \dots, e^{r_m} are the eigenvalues of Y. The function f is a central function if $f(Y) = f(r_1, \dots, r_m)$.

In this case, the invariant integral (7) reduces to

$$C \times \int_{\mathbb{R}^m} f(r_1, \dots, r_m) \prod_{i < j} \sinh^2 \left(\frac{|r_i - r_j|}{2} \right) dr$$
 (10)

where $dr = dr_1 \cdots dr_m$ and C is a constant [17].

3. RIEMANNIAN GAUSSIAN DISTRIBUTIONS ON

This section contains the definition and the main statistical properties of the Riemannian Gaussian distribution (RGD) on \mathcal{H}_m [17]. This distribution is introduced to give a statistical interpretation of the Riemannian barycentre on \mathcal{H}_m , which is used in many signal and image processing applications [18, 19, 22].

3.1. Definition

The probability density function of the RGD on \mathcal{H}_m , with respect to the Riemannian volume element (8), is given by

$$p(Y|\bar{Y},\sigma) = \frac{1}{\zeta(\sigma)} \exp\left[-\frac{d^2(Y,\bar{Y})}{2\sigma^2}\right]$$
(11)

where $\bar{Y} \in \mathcal{H}_m$ and $\sigma > 0$. In this definition, $\zeta(\sigma)$ is the normalizing constant given by

$$\zeta(\sigma) = \int_{\mathcal{H}_m} \exp\left[-\frac{d^2(Y, \bar{Y})}{2\sigma^2}\right] dv(Y) \tag{12}$$

The fact that it does not depend on \bar{Y} can be derived from (5), by taking $Z = \bar{Y}$, and by replacing it in (9). Moreover, replacing \bar{Y} with the identity matrix I_m in (12) and using (10),

$$\zeta(\sigma) = C \times \int_{\mathbb{R}^m} e^{-|r|^2/2\sigma^2} \prod_{i < j} \sinh^2(|r_i - r_j|/2) dr$$
 (13)

where |r| is the Euclidean norm of r and C is the constant appearing in (10). In practice, the function $\zeta(\sigma)$ can be computed by Monte Carlo integration. Interestingly, for m=2, $\zeta(\sigma)$ admits a closed-form expression which is proportional to $\sigma^2(e^{\sigma^2}-1)$. In the following, the RGD (11) will be denoted by $\mathcal{G}(\overline{Y}, \sigma)$.

Algorithm 1 below describes a simple way to sample from $\mathcal{G}(\bar{Y}, \sigma)$. Note that step 2 can be achieved by a Metropolis-Hastings algorithm.

Algorithm 1 Sampling from $\mathcal{G}(\bar{Y}, \sigma)$

- 1: Generate U from a uniform distribution on the group of unitary matrices of size $m \times m$
- 2: Generate $(r_1, \cdots, r_m) \in \mathbb{R}^m$ from the distribution ∞ $\begin{array}{c} e^{-|r|^2/2\sigma^2}\prod_{i< j}\sinh^2\left(|r_i-r_j|/2\right)\\ \text{3: }X\leftarrow U\cdot\operatorname{diag}(e^{r_1},\cdots,e^{r_m}) \end{array}$
- 4: $Y \leftarrow \bar{Y}^{1/2} \cdot X$

3.2. Maximum likelihood estimates of \bar{Y} and σ

Let Y_1, \ldots, Y_N be N i.i.d samples from $\mathcal{G}(\bar{Y}, \sigma)$ and let \hat{Y}_N be the barycentre of Y_1, \ldots, Y_N . Based on these observations, the maximum likelihood estimate (MLE) of \overline{Y} is the Riemannian barycentre \hat{Y}_N . Moreover, the MLE of σ is the solution $\hat{\sigma}_N$ of

$$\sigma^3 \times \frac{d}{d\sigma} \log \zeta(\sigma) = \frac{1}{N} \sum_{n=1}^{N} d^2(\hat{Y}_N, Y_n)$$

It can be proved that both \hat{Y}_N and $\hat{\sigma}_N$ exist and are unique [17].

For the main application of this paper, numerical computations of \hat{Y}_N and $\hat{\sigma}_N$ are needed. Computation of \hat{Y}_N consists in finding the Riemannian barycentre of Y_1, \ldots, Y_N , defined by (6). This can be done using the Riemannian gradient descent algorithm, which may be found in [23]. The k^{th} iteration of this algorithm produces an approximation \hat{Y}_N^k of \hat{Y}_N in the following way. For $k=1,2,\ldots,$ let Δ_k be the Hermitian matrix $\Delta_k = \sum_{n=1}^N \operatorname{Log}_{\hat{Y}_n^{k-1}}(Y_n)$ where Log denotes the Riemannian logarithm mapping, inverse to the exponential mapping, given by

$$\operatorname{Exp}_{Y}(\Delta) = Y^{1/2} \exp \left(Y^{-1/2} \, \Delta \, Y^{-1/2} \right) \, Y^{1/2}$$

Then, \hat{Y}_N^k is defined to be $\hat{Y}_N^k = \operatorname{Exp}_{\hat{Y}_N^{k-1}} (au_k \, \Delta_k)$ where $\tau_k > 0$ is a step size, which can be determined using a backtracking procedure.

The Riemannian gradient descent algorithm is repeated as long as $\|\Delta_k\| > \epsilon$, where $\|\Delta_k\| = \sqrt{g_{\hat{Y}_N^{k-1}}}(\Delta_k, \Delta_k)$ and ϵ is a precision parameter. In practice ϵ is set to the machine precision. This algorithm is guaranteed to converge, when

a suitable backtracking procedure is used, regardless of the initialization Y_N^0 .

On another hand, computation of $\hat{\sigma}_N$ requires solving a non-linear equation with one variable. This is readily done using Newton's method.

3.3. EM algorithm for mixture estimation

Mixture distributions are effective tools in statistical learning problems as they provide a good approximation to most distributions. The present paper applies this approach to learn distributions on \mathcal{H}_m . A mixture of RGD on \mathcal{H}_m is any probability distribution with density, with respect to the volume element (8), of the form

$$p(Y|(\varpi_{\mu}, \bar{Y}_{\mu}, \sigma_{\mu})_{1 \le \mu \le M}) = \sum_{\mu=1}^{M} \varpi_{\mu} \times p(Y|\bar{Y}_{\mu}, \sigma_{\mu})$$
 (14)

where $\varpi_{\mu} \in (0,1)$ are weights with sum one and where each $p(Y|\bar{Y}_{\mu},\sigma_{\mu})$ is a RGD on \mathcal{H}_m given by (11).

Let Y_1, \ldots, Y_N be independent samples from (14). Based on these samples, the MLE of $\vartheta = \{(\varpi_{\mu}, \tilde{Y}_{\mu}, \sigma_{\mu})\}$ can be computed using an EM algorithm similar to [16, 17]. For this, define for all $\vartheta = \{(\varpi_{\mu}, Y_{\mu}, \sigma_{\mu})\},\$

$$\omega_{\mu}(Y_n, \vartheta) = \frac{\varpi_{\mu} \times p(Y_n | \bar{Y}_{\mu}, \sigma_{\mu})}{\sum_{s=1}^{M} \varpi_s \times p(Y_n | \bar{Y}_s, \sigma_s)}, \quad N_{\mu}(\vartheta) = \sum_{n=1}^{N} \omega_{\mu}(Y_n)$$

The EM algorithm iteratively updates $\hat{\vartheta} = \{(\hat{\varpi}_{\mu}, \hat{Y}_{\mu}, \hat{\sigma}_{\mu})\}$, which is an approximation of the MLE of $\vartheta = (\varpi_{\mu}, \bar{Y}_{\mu}, \sigma_{\mu})$ as follows. Based on the current value of $\hat{\vartheta}$, it updates $\hat{\varpi}_{\mu}, \hat{Y}_{\mu}$ and $\hat{\sigma}_{\mu}$ as follows:

- ► Assign to $\hat{\varpi}_{\mu}$ the value $\hat{\varpi}_{\mu} = N_{\mu}(\hat{\vartheta})/N$
- ightharpoonup Assign to \hat{Y}_{μ} the value

$$\hat{Y}_{\mu} = \underset{Y}{\operatorname{arg\,min}} \sum_{n=1}^{N} \omega_{\mu}(Y_{n}, \hat{\vartheta}) d^{2}(Y, Y_{n})$$

► Assign to $\hat{\sigma}_{\mu}$ the value

$$\hat{\sigma}_{\mu} = \Phi\left(\sqrt{N_{\mu}^{-1}}(\hat{\vartheta}) \times \sum_{n=1}^{N} \omega_{\mu}(Y_{n}, \hat{\vartheta}) \, d^{2}(\hat{Y}_{\mu}, Y_{n}) \right)$$

where the function Φ is the inverse of $\sigma \mapsto \sigma^3 \times$ $\frac{d}{d\sigma}\log\zeta(\sigma)$.

Consider now the problem of classifying new observations modeled as a realization of (14). Following [16, 17], a new observation Y_t is associated to the class C_{μ^*} of the closest cluster, i.e.

$$\mu^* = \operatorname{argmin}_{\mu} \left\{ -\log \, \hat{\varpi}_{\mu} + \log \, \zeta(\hat{\sigma}_{\mu}) + \frac{d^2(Y_t \,, \hat{Y}_{\mu})}{2\hat{\sigma}_{\mu}^2} \right\}$$

This classification rule is optimal in the sense of a Bayesian risk criterion given in [24].

4. APPLICATION TO TEXTURE IMAGE CLASSIFICATION

This section presents an application of the RGD on \mathcal{H}_m for the classification of texture images. The performance of this distribution is first compared with the RGD on \mathcal{P}_m . Second, it is compared with Wishart distribution. Third, the influence of the wavelet decomposition on the classification performance is evaluated. For this purpose, the following numerical experiment on the VisTex database [25] is carried out.

For each texture image of size 512×512 pixels, 196 subimages (128 × 128) are extracted with a 32 pixel overlap. For each subimage, 6 subbands were computed using the complex Gabor filter. Then, two spatial neighborhoods (i.e. horizontal dH and vertical dV) of one pixel are considered. Each subband s of subimage n is composed by a set of two bivariate populations $\Pi_{s,n,dH}$ and $\Pi_{s,n,dV}$, represented respectively by their Hermitian covariance matrices $Y_{s,n,dH}$ and $Y_{s,n,dV} \in \mathcal{H}_2$. These covariance matrices are features which describe the spatial dependence of subband coefficients. The size of the feature space is hence F = 12 (2 spatial neighborhood and 6 subbands). To summarize, each subimage n is represented by a set of F covariance matrices denoted $Y_{f,n}$. Then, the database is equally divided for the classification, where half is used for training and the other half for testing. For each training class, a mixture model is learned by considering the EM algorithm described in Section 3.3. In this experiment, the number of clusters per class (denoted K) is set to 1, 3 or automatically fixed by optimizing the Bayesian information criterion (BIC) [26]. It results that each texture is represented by a set of K clusters containing the maximum likelihood estimates $(\hat{\varpi}_a, Y_{f,a}, \hat{\sigma}_{f,a})$, for $a = 1, \ldots, K$ and $f = 1, \ldots, F$. Then, as the subband are assumed independent, a test subimage t represented by a set of F covariance matrices $Y_{f,t}$ in \mathcal{H}_2 is associated to the class of the closest cluster C_* , i.e. the one realizing the minimum over c of

$$-\log \hat{\varpi}_c + \sum_{f=1}^F \log Z(\hat{\sigma}_{f,c}) + \sum_{f=1}^F \frac{d^2(Y_{f,t}, \hat{Y}_{f,c})}{2\hat{\sigma}_{f,c}^2}$$

Table 1 displays the classification performance in terms of kappa accuracy (mean \pm standard deviation) on the Vis-Tex database. The first line gives the results for the RGD with respectively K=1, K=3 and K fixed by the BIC criterion. The second line shows the corresponding results for the Wishart model. Moreover, the second column in Table 1 shows the results with the Hermitian covariance matrices in \mathcal{H}_m , while the first column exhibits the results obtained for real covariance matrices (obtained by taking the absolute value of the Gabor subbands) in \mathcal{P}_m .

As shown in Table 1, for each model, a significant gain of more than 2% is observed by working on \mathcal{H}_m rather than \mathcal{P}_m . Note also, that the use of the BIC criterion improved the classification performance compared to a fixed number of clusters

model	number of modes (K)	on \mathcal{P}_m	on \mathcal{H}_m
RGD	K=1	78.24 ± 0.36	80.62 ± 0.46
	K=3	90.24 ± 0.40	92.24 ± 0.32
	BIC	91.91 ± 0.45	94.29 ± 0.30
Wishart	K=1	81.93 ± 0.51	82.15 ± 0.50
	K=3	86.07 ± 0.61	87.27 ± 0.71
	BIC	86.14 ± 0.44	87.64 ± 1.06

Table 1: Classification performance (κ accuracy) on the Vis-Tex database.

Wavelet decomposition	RGD	Wishart
Gabor	94.29 ± 0.30	87.64 ± 1.06
CWT	98.03 ± 0.25	97.61 ± 0.26
DT-CWT	98.79 ± 0.17	98.32 ± 0.30

Table 2: Influence of the wavelet decomposition on the classification performance (κ accuracy) on the VisTex database.

(K=1 or 3). Finally, while having the same complexity in terms of number of parameters, the mixture models for the proposed RGD have a significant gain of 4% to 7% compared to the mixture of Wishart model.

A second experiment is carried out on the same database to evaluate the influence of the wavelet decomposition on the classification performance. For that, three decompositions have been considered, which are the Gabor filters, the complex wavelet transform (CWT) and the dual-tree complex wavelet transform (DT-CWT). Here, the number of modes is fixed by optimizing the BIC criterion. As observed in Table 2, the best results are obtained with the DT-CWT. Note also that for each wavelet decomposition, the proposed RGD model leads to better performances compared to the Wishart model. This illustrates the potential of the proposed geometric learning approach on \mathcal{H}_m .

5. CONCLUSION

In this paper, the Riemannian Gaussian distribution (RGD) has been introduced on the space \mathcal{H}_m of Hermitian positive definite matrices. This distribution has a one to one similarity with the RGD on the space \mathcal{P}_m , of $m \times m$ positive definite matrices, previously introduced in [16]. The proposed RGD has two parameters: the unique Riemannian centre of mass \bar{Y} and the dispersion parameter σ . Once having presented the maximum likelihood estimators and the extension of the RGD to mixture models, an experiment on the VisTex database has been carried out to illustrate the potential of the proposed model for texture image classification. A significant gain has been observed compared to the state-of-the-art Wishart model.

Further works will focus on the generalization of the RGD model to structured covariance matrices (e.g. Toeplitz, block Toeplitz matrices).

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