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Unmixing Prior to Supervised Classification of Remotely Sensed Hyperspectral Images

Inmaculada Dópido, Maciel Zortea, Alberto Villa, Student Member, IEEE, Antonio Plaza, Senior Member, IEEE, and Paolo Gamba, Senior Member, IEEE

Abstract—Supervised classification of hyperspectral images is a very challenging task due to the generally unfavorable ratio between the number of spectral bands and the number of training samples available a priori, which results in the Hughes phenomenon. For this purpose, several feature extraction methods have been investigated in order to reduce the dimensionality of the data to the right subspace without significant loss of the original information that allows for the separation of classes. In this letter, we explore the use of spectral unmixing for feature extraction prior to supervised classification of hyperspectral data using support vector machines. The proposed feature extraction strategy has been implemented in the form of four different unmixing chains and evaluated using two different scenes collected by National Aeronautics and Space Administration Jet Propulsion Laboratory’s Airborne Visible/Infrared Imaging Spectrometer. The experiments suggest competitive results but also show that the definition of the unmixing chains plays an important role in the final classification accuracy. Moreover, differently from most feature extraction techniques available in the literature, the features obtained using linear spectral unmixing are potentially easier to interpret due to their physical meaning.

Index Terms—Endmember extraction, feature extraction, hyperspectral image classification, spectral unmixing, support vector machines (SVMs).

I. INTRODUCTION

In MANY studies, hyperspectral analysis techniques are divided into full- and mixed-pixel classification techniques [1]–[3], where each pixel vector defines a spectral signature or fingerprint that uniquely characterizes the underlying materials at each site in a scene. Full-pixel classification techniques assume that each spectral signature comprises the response of one single underlying material. Often, however, this is not a realistic assumption. If the spatial resolution of the sensor is not fine enough to separate different pure signature classes at a macroscopic level, these can jointly occupy a single pixel, and the resulting spectral signature will be a composite of the individual pure spectra, often called endmembers in hyperspectral imaging terminology [4]. Let us denote a remotely sensed hyperspectral scene with \( n \) bands by \( \mathbf{I} \), in which each pixel is represented by a vector \( \mathbf{X} = [x_1, x_2, \ldots, x_n] \in \mathbb{R}^n \), where \( \mathbb{R} \) denotes the set of real numbers in which the pixel’s spectral response \( x_k \) at sensor channels \( k = 1, \ldots, n \) is included. Under the linear mixture model assumption, each pixel vector can be modeled using

\[
\mathbf{X} = \sum_{z=1}^{p} \Phi_z \cdot \mathbf{E}_z + \mathbf{n}
\]

where \( \mathbf{E}_z \) denotes the spectral response of endmember \( z \), \( \Phi_z \) is a scalar value designating the fractional abundance of the endmember \( z \) at the pixel \( \mathbf{X} \), \( p \) is the total number of endmembers, and \( \mathbf{n} \) is a noise vector. Two physical constraints can be imposed into the model described in (1): abundance nonnegativity constraint, i.e., \( \Phi_z \geq 0 \), and abundance sum-to-one constraint, i.e., \( \sum_{z=1}^{p} \Phi_z = 1 \) [5].

Several machine learning techniques have been applied, under the full-pixel assumption, to extract relevant information from hyperspectral data. The good classification performance exhibited by the support vector machine (SVM) [1], [6], [7] using spectral signatures as input features can be improved by applying suitable feature extraction strategies that are able to reduce the dimensionality of the data to a subspace without losing the original information [8]. Techniques used for this purpose include principal component analysis (PCA) [3], minimum noise fraction (MNF) [9], or independent component analysis (ICA) [10]. PCA and MNF maximize the amount of data variance and signal-to-noise ratio (SNR), respectively, yielding a transformed data set in a new uncorrelated coordinate system, while ICA tries to find components as statistically independent as possible. However, all these methods maximize the information contained in the first transformed components, relegating variations of less significant size to low-order components. If such low-order components are not preserved, small classes may be affected. The inclusion of spatial features such as morphological profiles can be used to address this issue [8], [11], [12].

In this letter, we explore an alternative strategy focused on the use of spectral unmixing for feature extraction prior to classification. Previous efforts in this direction were presented in [13] and [14], but the analysis of whether spectral unmixing can replace standard feature extraction transformations remains an unexplored topic. Although classification techniques often neglect the impact of mixed pixels in the provision of a set of final class labels, widely used benchmark data sets in the literature—e.g., the Airborne Visible/Infrared Imaging Spectrometer (AVIRIS) Indian Pines scene—are known to be
dominated by mixed pixels, even if the associated ground-truth information is only available in full-pixel form. Hence, the use of spectral unmixing presents distinctive features with regard to other approaches such as PCA, MNF, or ICA. First, it provides additional information for classification in hyperspectral analysis scenarios with moderate spatial resolution, since the subpixel composition of training samples can be used as part of the learning process of the classifier. Second, the components estimated by spectral unmixing can be physically explained as the abundances of spectral endmembers. Third, spectral unmixing does not penalize classes which are not relevant in terms of variance or SNR. Here, we design different unmixing processing chains with the goal of addressing three specific research questions.

1) Is spectral unmixing a feasible strategy for feature extraction prior to classification?
2) Does the inclusion of spatial information at the endmember extraction stage lead to better classification results?
3) Is it really necessary to estimate pure spectral endmembers for classification purposes?

We have structured the remainder of this letter as follows. Section II describes the considered spectral unmixing chains. Section III presents different experiments specifically designed to address the aforementioned research questions and provide a comparison between the proposed unmixing-based strategy and other feature extraction approaches in the literature. Section IV concludes with some remarks and future research avenues.

II. UNMIXING-BASED FEATURE EXTRACTION

A. Unmixing Chain #1

In this section, we describe our first approach to design an unmixing-based feature extraction chain which can be summarized by the flowchart in Fig. 1. First, we estimate the number of endmembers \( p \) directly from the original \( n \)-dimensional hyperspectral image \( I \). For this purpose, we use in this letter two standard techniques widely used in the literature such as the HySime method [15] and the virtual dimensionality (VD) concept [16]. Once the number of endmembers \( p \) has been estimated, we apply an automatic algorithm to extract a set of endmembers from the original hyperspectral image [17]. Here, we use an orthogonal subspace projection technique [18] which has been shown in previous work to provide a very good tradeoff between the signature purity of the extracted endmembers and the computational time to obtain them. Preliminary experiments conducted with other endmember extraction techniques, such as vertex component analysis [19] and N-FINDR [20], have shown very similar results in terms of classification accuracy. Finally, linear spectral unmixing (either unconstrained or constrained) can be used to estimate the abundance of each endmember in each pixel of the scene, providing a set of \( p \) abundance maps. Then, standard SVM classification is performed on the stack of abundance fractions using randomly selected training samples.

B. Unmixing Chain #2

In this section, we introduce a variation of the unmixing-based feature extraction chain which includes spatial preprocessing prior to endmember extraction in order to guide the endmember searching process to those areas which are more spatially homogeneous. This approach is shown in Fig. 1. The spatial preprocessing strategy adopted in this letter is described in detail in [21]. As in the previous chain, the features resulting from the proposed (spatially enhanced) unmixing process are used to train an SVM classifier with a few randomly selected labeled samples. The classifier is then tested using the remaining labeled samples.

C. Unmixing Chain #3

Our main motivation for introducing a third unmixing-based feature extraction chain is the fact that the estimation of the number of endmembers \( p \) in the original image is a very challenging issue. Fig. 2 shows a new chain in which the endmembers are extracted from the set of available (labeled) training samples instead of from the original image. This chain introduces two important variations: 1) As a simplification to the challenging estimation problem, the number of endmembers to be extracted is set as the total number of different classes \( c \) in the training set, and 2) the endmember searching process is conducted only on the training set, which reduces computational complexity. However, the number of endmembers in the original image \( p \) is probably different than \( c \), the number of labeled classes. Therefore, in order to unmix the original image, we need to address a partial unmixing problem (in which not all endmembers may be available a priori). A successful
technique for this purpose is mixture-tuned matched filtering (MTMF) [22], also known as constrained energy minimization [23], which combines linear spectral unmixing and statistical matched filtering. From matched filtering, it inherits the ability to map a single known target without knowing the other background endmember signatures. From spectral mixture modeling, it inherits the leverage arising from the mixed pixel model and the constraints on feasibility.

D. Unmixing Chain #4

The fourth unmixing chain tested in our experiments represents a slight variation of unmixing chain #3 in which the spectral signatures used for unmixing purposes are not obtained via endmember extraction but through averaging of the spectral signatures associated to each labeled class in the training set. To keep the number of estimated components low, only one component is allowed for each class. This averaging strategy produces \( c \) signatures, each representative of a labeled class, which are then used to partially unmix the original hyperspectral scene using MTMF.

III. EXPERIMENTAL RESULTS

A. Hyperspectral Data

The first data set used in our experiments was collected by the AVIRIS sensor over the Indian Pines region. The scene comprises 145 lines by 145 samples and 220 spectral channels with the wavelength range from 0.4 to 2.5 \( \mu \text{m} \) nominal spectral resolution of 10 nm, and spatial resolution of 20 m by pixel. After removing noisy and water absorption bands, 202 channels were left. The ground truth contains 16 land cover classes. The number of pixels in the smallest class is 20, while the number of pixels in the largest class is 2468.

The second data set was collected by the AVIRIS sensor over the Kennedy Space Center, Florida, in March 1996. The portion of this scene used in our experiments has dimensions of 292 \( \times 383 \) pixels. After removing water absorption and low-SNR bands, 176 bands were used for the analysis. The spatial resolution is 20 m by pixel. Twelve ground-truth classes were available, where the number of pixels in the smallest class is 134 while the number of pixels in the largest class is 761.

B. Experiments

1) Experiment 1. Use of Unmixing as a Feature Extraction Strategy: In this experiment, we use the AVIRIS Indian Pines and Kennedy Space Center data sets to analyze the impact of imposing nonnegativity and sum-to-one constraints in abundance estimation prior to classification. For the AVIRIS Indian Pines image, we construct ten small training sets by randomly selecting 5\%, 10\%, and 15\% of the ground-truth pixels. For the AVIRIS Kennedy Space Center, since the size of the smaller classes is bigger, we decided to reduce the training sets even more and selected 1\%, 3\%, and 5\% of the available ground-truth pixels. Then, the three considered types of input features (original, reduced, and unmixing based) are built for the selected training samples and used to train an SVM classifier in which two types of kernels, i.e., polynomial and Gaussian, are used. The SVM was trained with each of these training subsets and then evaluated with the remaining test set. Each experiment was repeated ten times, and the mean and standard deviation accuracy values were reported. Kernel parameters were optimized by a grid search procedure, and the optimal parameters were selected using tenfold cross validation. The LIBSVM library was used for the experiments.

Table I summarizes the overall classification accuracies obtained after applying the considered SVM classification system (with polynomial and Gaussian kernels) to the features extracted after applying unmixing chain #1 (see Fig. 1) to the AVIRIS scenes.

Table: Classification Accuracies (Percentage) and Standard Deviation Obtained After Applying the Considered SVM Classification System (With Gaussian and Polynomial Kernels) to Three Different Types of Features (Original, Reduced, and Unmixing Based) Extracted From the AVIRIS Indian Pines and Kennedy Space Center Scenes (Ten Randomly Chosen Training Sets)

<table>
<thead>
<tr>
<th>Image</th>
<th>Type of feature</th>
<th># of features</th>
<th>Polynomial kernel</th>
<th>Gaussian kernel</th>
</tr>
</thead>
<tbody>
<tr>
<td></td>
<td></td>
<td></td>
<td>5%</td>
<td>10%</td>
</tr>
<tr>
<td></td>
<td></td>
<td></td>
<td>5%</td>
<td>10%</td>
</tr>
<tr>
<td></td>
<td></td>
<td></td>
<td>5%</td>
<td>10%</td>
</tr>
<tr>
<td>Original spectral information</td>
<td>202</td>
<td>75.78±0.123</td>
<td>81.55±0.086</td>
<td>83.58±0.078</td>
</tr>
<tr>
<td>PCA</td>
<td>18</td>
<td>77.07±0.146</td>
<td>81.66±0.088</td>
<td>83.11±0.052</td>
</tr>
<tr>
<td>MNF</td>
<td>18</td>
<td>82.97±0.93</td>
<td>87.41±0.31</td>
<td>88.38±0.57</td>
</tr>
<tr>
<td>ICA</td>
<td>18</td>
<td>76.63±1.27</td>
<td>81.00±0.21</td>
<td>82.94±0.36</td>
</tr>
<tr>
<td>Chain #1</td>
<td>18</td>
<td>74.56±1.04</td>
<td>79.20±1.12</td>
<td>80.97±0.50</td>
</tr>
<tr>
<td>Chain #2</td>
<td>18</td>
<td>71.93±0.96</td>
<td>77.58±0.92</td>
<td>79.31±0.33</td>
</tr>
<tr>
<td>Chain #3</td>
<td>16</td>
<td>81.32±0.84</td>
<td>85.56±0.84</td>
<td>86.83±0.57</td>
</tr>
<tr>
<td>Chain #4</td>
<td>16</td>
<td>82.36±1.09</td>
<td>86.87±0.59</td>
<td>87.97±0.57</td>
</tr>
</tbody>
</table>

1Available online: http://dynamo.ecn.purdue.edu/~biehl/MultiSpec
2Available online: http://www.csr.utexas.edu/hyperspectral/data/KSC/
3http://www.csie.ntu.edu.tw/~cjlin/libsvm/
It is better to have a representative endmember selected per class. In order to address this issue, a possible solution is endmember searching process to spatially homogeneous areas provided by PCA, MNF, or ICA, or the original spectral information. This is because endmember extraction is generally sensitive to outliers and anomalies; hence, a strategy for directing the endmember searching process to spatially homogeneous areas could improve the final classification results.

2) Experiment 2. Impact of Including Spatial Information at the Endmember Extraction Stage: In this experiment, we apply unmixing chain #2 for feature extraction prior to classification. As shown in Table I, spatial preprocessing prior to endmember extraction cannot lead to improved classification results with regard to chain #1 and the original spectral information. This is due to the spectral similarity of the most spatially representative classes in our considered scenes. For instance, in the AVIRIS Indian Pines scene, the corns and soybeans were very early in their growth cycle at the time of data collection, which resulted in low coverage of the soil (≈5%) [24]. Given this low canopy ground cover, the variation in spectral response among different classes is very low, and spatial information cannot significantly increase discrimination between different classes. In order to address this issue, a possible solution is to conduct the endmember extraction process in a supervised fashion, taking advantage of the information contained in the available labeled samples in order to guarantee that a highly representative endmember is selected per class.

3) Experiment 3. Impact of Endmember Purity on the Final Classification Results: In a supervised endmember extraction framework, our first experiment is based on applying unmixing chain #3 to select endmembers only from the available training samples. Apart from reducing computational complexity (which, in this case, involves a search for c endmembers in the pixels belonging to the training set), Table I reveals that this strategy improves the classification results reported for chains #1 and #2. However, in order to make sure that only one endmember per labeled class is used for unmixing purposes, we also apply unmixing chain #4 in which spectral averaging of the available training pixels in each class is conducted in order to produce a final set of c spectral signatures. Although averaging of endmembers can lead to degradation of spectral purity, it can also reduce the effects of noise and/or average out the subtle spectral variability of a given class, thus obtaining a more representative endmember for the class as a whole. This is illustrated by the classification results for unmixing chain #4 in Table I, which outperform those reported for most other tested methods except the MNF. This indicates that, in a supervised unmixing scenario, the use of spectrally pure signatures is not as important as the choice of signatures which are representative of the available training samples.

Table II shows the statistical differences (average value of ten comparisons) between traditional dimensionality reduction methods and unmixing chains #3 and #4, computed using McNemar’s test [25] for the case of the polynomial kernel. The differences are statistically significant at a confidence level of 95% if |Z| > 1.96. For each couple of compared feature extraction chains, we report also how many times each chain wins/ties/loses after comparing the thematic maps obtained using the same training set. If the value of Z reported for each entry in Table II is positive and larger than 1.96, the first compared chain wins. By convention, the comparison is always performed with the first chain in a line in Table II and the second chain in a column in Table II. It can be noticed that using chains #3 and #4 always perform significantly better than PCA and ICA. MNF performs better than chain #3, while the differences with chain #4 are in general not significant.

To conclude this section, Fig. 3 shows the best classification results (out of ten runs) obtained after applying the SVM—trained with 10% of the available training samples—to each feature extraction strategy considered for the AVIRIS Indian Pines scene. As shown in Fig. 3, both the MNF in Fig. 3(d) and chain #4 in Fig. 3(h) provide the best classification scores, with less confusion in heavily mixed classes.
samples during the learning stage, with the potential advantage of improved interpretability of features due to the physical nature of the extracted abundance maps. Although final classification accuracies are likely to be dependent on the particular data set considered, the chains tested suggest higher accuracies with respect to traditional methods, such as PCA and ICA, and comparable accuracies related to MNF.

Further research is needed to define an optimality criterion to design unmixing chains as a feature reduction tool for classification purposes. A start point might be chain #4 which indicates that, in the context of a supervised unmixing scenario, the use of spectrally pure signatures is not as important as the choice of signatures which are highly representative of the available training samples.

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Fig. 3. Best classification results for AVIRIS Indian Pines (using SVM classifier with Gaussian kernel, trained with 10% of the available samples per class). (a) Ground-truth. (b) Original image (84.27%). (c) PCA (83.33%). (d) MNF (89.41%). (e) Chain #1 (81.29%). (f) Chain #2 (79.64%). (g) Chain #3 (87.99%). (h) Chain #4 (89.26%). (i) Class accuracies (MNF versus Chain #4).