INTERPRETABLE MULTIPLE LOSS FUNCTIONS IN A LOW-RANK DEEP IMAGE PRIOR BASED METHOD FOR SINGLE HYPERSPECTRAL IMAGE SUPER-RESOLUTION

*Tatiana Gelvez*¹ *Jorge Bacca*² *Henry Arguello*²

¹Department of Electrical Engineering, Universidad Industrial de Santander, Bucaramanga, Colombia ²Department of Computer Science, Universidad Industrial de Santander, Bucaramanga, Colombia

ABSTRACT

This paper presents a single hyperspectral image superresolution (HSI-SR) method based on the non data-driven deep image prior approach, where the prior information of the image is intrinsically learned through the weights and the structure of a neural network. Precisely, we propose a scheme composed of a sequence of independent and interpretable deep-blocks, whose outputs are affected in the back-propagation step not only by the general loss-function, but also by a single loss-function, resulting in a scheme using multiple loss-functions. The internal architecture of each deep-block consists of a low-rank decomposition with two block-layers inspired by the linear mixture model, where the spectral image is formed as the product between an abundance and an endmember matrix, such that, the features of the first block-layer and the weights of the second block-layer can be interpreted as the abundances and the endmembers, respectively. In the experimental section, we showed the remarkable performance of the proposed HSI-SR method concerning the extensive reviewed state-of-the-art approaches.

Index Terms— Neural networks, Deep image prior, Low-rank approximation, Hyperspectral image super-resolution.

1. INTRODUCTION

Single hyperspectral image super-resolution (HSI-SR) recovery technique addresses the spatial-spectral tradeoff of current optical devices, that acquire a limited amount of incident energy when sensing HSIs with dozens or hundreds of narrow spectral bands, sacrificing spatial resolution. For this, HSI-SR methods aim to infer a high-spatial-resolution HSI (HR-HSI) from the acquired low-spatial-resolution HSI (LR-HSI) with no additional auxiliary images by exploiting the high spatialspectral correlations as prior information. Model-based optimization methods incorporate such prior information through hand-crafted regularizers as total-variation [1], sparse representation [2], and low-rankness [3]. In contrast, deep learning (DL)-based methods learn the prior information or the entire decoder operator to model the relationship between the LR-HSI and the HR-HSI through the training of a huge amount of datasets [4, 5]. For instance, authors in [6, 7, 8] introduce convolutional neural networks (CNNs) based on 3D filters to better exploit the spatial-spectral correlations. Still, DLbased methods can become impractical given the high cost of acquiring several HSI training datasets [9, 10]. The recent so-called deep image prior (DIP) approach showed that the prior information can be embodied in the weights of the neural network allowing to solve various inverse problems, such as HSI-SR, in a non data-driven manner [11]. For instance, [12] integrated DIP with 3D convolutions containing higher low-level image information, [13] boosted DIP strategy by adding explicit regularization, and [14] provided an evolutionary strategy to automatically search for the optimal hyper-parameters of the DIP architecture.

Previous DIP-based methods for HSI-SR commonly employ a standard loss-function of the output of the model. Contrary, we propose a DIP-based HSI-SR method that incorporates the idea of using multiple loss-functions associated with multiple deep-blocks forming the deep architecture. This multiple deep-blocks scheme considers that the deeper the network, the finer the learned features estimating the spatialspectral correlations of HSIs. Besides, we propose each deepblock to be interpretable in the sense that its internal structure consists of two block-layers inspired by the linear mixture model (LMM), where the outputs can be interpreted as a lowrank approximation of the HR-HSI, decomposed as a matrix product between, an endmember matrix containing the spectral response of few independent materials in the scene, and an abundance matrix including the fractional proportions of the endmembers at each spatial location [15]. The proposal then recovers the HR-HSI from the LR-HSI in a non datadriven manner by including the prior information on the architecture, where neither the abundances nor the endmembers are known. Instead, at each deep-block, the features of the first block-layer can be interpreted as the abundances, and the weights of the second block-layer as the endmembers, where some included regularizers consider the LMM constraints.

Simulations over datasets showed the effectiveness of the proposed interpretable architecture for HSI-SR, even when compared against data-driven methods. In particular, the improvement is obtained when using the sum of the single loss-functions affecting the multiple deep-blocks, and the intrinsic regularization by the neural network architecture.

2. THEORETICAL BACKGROUND

2.1. Single Hyperspectral Image Super-resolution

Let $\mathbf{f} \in \mathbb{R}^{N_h^2 L}$ denote the vector form of an HR-HSI with $(N_h \times N_h)$ spatial pixels and L spectral bands. The LR-HSI acquisition is given by the following forward linear model

$$\mathbf{f}_{low} = \mathbf{D}\mathbf{B}\mathbf{f} + \boldsymbol{\eta},\tag{1}$$

where $\mathbf{f}_{low} \in \mathbb{R}^{N_l^2 L}$ stands for the observed LR-HSI, with $N_l \times N_l$ spatial pixels, and *L* spectral bands; $\mathbf{D} \in \mathbb{R}^{N_l^2 L \times N_h^2 L}$ denotes a spatial downsampling operator, where $N_l = (1/d)N_h$, with $d \in \mathbb{Z}_+$ being a downsampling factor; $\mathbf{B} \in \mathbb{R}^{N_h^2 L \times N_h^2 L}$ denotes a spatial blurring operator, and $\boldsymbol{\eta} \in \mathbb{R}^{N_l^2 L}$ stands for added additive Gaussian noise.

2.2. Linear Mixture Model for Hyperspectral Images

The linear mixture model (LMM) states that natural scenes contain a few number $r \ll L$ of materials defined uniquely by its spectral response. Then, the i^{th} spatial pixel $\mathbf{f}_i \in \mathbb{R}^L$ can be modeled as $\mathbf{f}_i = \mathbf{E}\mathbf{a}_i$, where $\mathbf{E} \in \mathbb{R}^{L \times r}$ is an endmember matrix, whose columns contain a unique spectral response, and $\mathbf{a}_i \in \mathbb{R}^r$ is an abundance vector, containing the fractional proportions of each endmember at the i^{th} spatial pixel, for $i = 1, \ldots, N_h^2$. Then, \mathbf{f} can be low-rank decomposed as

$$\mathbf{f} = (\mathbf{E} \otimes \mathbf{I}_{N_h^2}) \bar{\mathbf{a}} = \bar{\mathbf{E}} \bar{\mathbf{a}},\tag{2}$$

where $\bar{\mathbf{a}} \in \mathbb{R}^{N_h^2 r}_+ = [\mathbf{a}_1^T \dots \mathbf{a}_i^T \dots \mathbf{a}_{N_h^2}^T]^T$ stacks the abundances; $\bar{\mathbf{E}} \in \mathbb{R}^{N_h^2 L \times N_h^2 r}_+$ encompasses the endmembers; $\mathbf{I}_{N_h^2}$ refers to an identity matrix of size N_h^2 ; and \otimes denotes the Kronecker product, introduced to apply the endmembers along the spatial pixels, maintaining a vector notation.

The LMM entails the physical constraints that a mixed pixel has an entire composition, such that, the sum of non-negative fractional proportions must be one, i.e. $\mathbf{a}_i[a] \ge 0$, and $\sum_{a=1}^{r} \mathbf{a}_i[a] = 1, \forall i$. Also, since the endmembers connote light intensities they are non-negative [16].

3. PROPOSED SINGLE HYPERSPECTRAL SUPER-RESOLUTION METHOD

The proposed idea is to find a deep model based on the lowrank decomposition $\overline{\mathbf{E}}\overline{\mathbf{a}}$, such that, a super-resolved image can be estimated. Then, the proposed HSI-SR scheme solves the following optimization problem based on the DIP approach that considers the image prior information implicitly through fitting the weights of a deep model given by

$$\hat{\theta} \in \underset{\theta}{\operatorname{argmin}} \quad \left\| \mathbf{f}_{low} - \mathbf{DB} \mathcal{M}_{\theta}(\mathbf{f}^{0}) \right\|_{2}^{2}, \tag{3}$$

where $\hat{\mathbf{f}} := \mathcal{M}_{\hat{\theta}}(\mathbf{f}^0)$ denotes the estimated HR-HSI, $\mathbf{f}^0 \in \mathbb{R}^{N_h^2 L}$ denotes a bicubic interpolation taken from \mathbf{f}_{low} , given

as the input of the deep generator model $\mathcal{M}_{\theta}(\cdot) : \mathbb{R}^{N_h^2 L} \to \mathbb{R}^{N_h^2 L}$, with θ as the adjustable deep parameters. Notice that, although problem in (3) is based on a deep model, it only requires the LR-HSI, f_{low} , and the forward operators **DB**, i.e, it is non data-driven. Precisely, we aim to capture the prior information implicitly by proposing an adequate design of the deep generative model, $\mathcal{M}_{\theta}(\cdot)$. For this, we determine the architecture of the deep generative model to be composed by a sequence of multiple interpretable deep-blocks, containing two block-layers inspired by the LMM, so that, the k^{th} deep-block can be mathematically expressed as

$$\mathcal{M}_{\theta^k}(\mathbf{f}^{k-1}) = \mathbf{f}^k = \mathcal{E}_{\mathbf{E}^k}(\mathcal{A}_{\theta}^k(\mathbf{f}^{k-1})), \tag{4}$$

where $\mathcal{A}_{\theta}^{k}(\cdot) : \mathbb{R}^{N_{h}^{2}L} \to \mathbb{R}^{N_{h}^{2}r}$ models the first block-layer, referred to as the abundance block-layer, consisting of a CNN that receives and properly filters the input to obtain an output whose properties and dimensions should match for being interpreted as the abundances. $\mathcal{E}_{\mathbf{E}^{k}}(\cdot) : \mathbb{R}^{N_{h}^{2}r} \to \mathbb{R}^{N_{h}^{2}L}$ models the fully connected second block-layer, referred to as the endmember block-layer, consisting of an operator that performs the matrix multiplication between the learned features in the abundance block-layer $\mathbf{\bar{E}}$, whose dimensions should match to be interpreted as the end-member shock-layer $\mathbf{\bar{E}}$, whose dimensions should match to be interpreted as the end-members according to the LMM in (2).

Then, the proposed deep model can be expressed as

$$\mathcal{M}_{\theta}(\mathbf{f}^{0}) = \hat{\mathbf{f}} = \mathcal{M}_{\theta^{K}}(\mathcal{M}_{\theta^{K-1}}(\cdots \mathcal{M}_{\theta^{1}}(\mathbf{f}^{0}))), \quad (5)$$

where $\boldsymbol{\theta} = [\boldsymbol{\theta}^1, \dots, \boldsymbol{\theta}^k, \dots \boldsymbol{\theta}^K]$ stacks the adjustable weights of each deep-block, $\mathbf{f}^k := \mathcal{M}_{\theta^k}(\mathbf{f}^{k-1})$, with each \mathbf{f}^k using a single loss-function as detailed in following sub-sections.

3.1. Abundance Block-layer

The abundance block-layer involves a CNN that filters and operates the spatial dimension of the previous deep-block output. This, given that the abundances entail the spatial correlations of the underlying HSI. Mathematically, the output of the abundance block-layer at the k^{th} deep-block, interpreted as the estimated abundances $\bar{\mathbf{a}}^k \in \mathbb{R}^{N_h^2 r}$ can be expressed as

$$\bar{\mathbf{a}}^k = \mathcal{A}_{\theta^k}(\mathbf{f}^{k-1}). \tag{6}$$

Notice that r denotes a tunable hyper-parameter related to the amount of endmembers present in the scene. The abundance block-layer is based on an autoencoder network, with six 2D convolutional layers with kernel sizes of 3×3 , where the number of features increases symmetrically from L to 3L, and then decreases to L. A value of L = 32 was chosen for this work, where the spatial dimensions are preserved using a padding strategy. The abundance block-layer also adds the LMM physical constraints described in Section 2.2 by using the *sigmoid* function as the activation of the network output $\bar{\mathbf{a}}^k$, and including an explicit regularization term in the loss-function, $R(\bar{\mathbf{a}}^k) = \sum_{a=1}^r \mathbf{a}_i^k[a] = 1$, with $\mathbf{a}_i[a] \ge 0, \forall i$.

3.2. Endmember Block-layer

The endmember block-layer consists of a trainable layer that follows the LMM to adjust the underlying spectral responses by performing a matrix multiplication between the output of the previous abundance block-layer $\bar{\mathbf{a}}^k$, and the learned weights, in such a manner that these weights can be interpreted as the estimated endmembers at the k^{th} deep-block. Mathematically, the endmember block-layer is expressed as

$$\mathbf{f}^{k} = \mathcal{E}_{\mathbf{E}^{k}}\left(\bar{\mathbf{a}}^{k}\right) = \left(\mathbf{E}^{k} \otimes \mathbf{I}_{N_{k}^{2}}\right)\bar{\mathbf{a}}^{k},\tag{7}$$

This block-layer also includes the non-negative constraint described in Section 2.2 by projecting the estimated endmembers into \mathbb{R}_+ at each gradient step.

3.3. Multiple Loss-functions

In (3) the DIP-based approach applies the forward operators **DB** at the output of the deep model to estimate the LR-HSI. Then, the proposal employs such forward operators at the output of each one of the multiple deep-blocks. Furthermore, we propose adjusting each internal architecture weights to refine the learned estimation as the block is deeper by minimizing a single loss-function containing the loss from the difference between the observed LR-HSI and the predicted LR-HSI of each deep-block and the LMM physical constraints. The proposal then solves the optimization problem given by

$$\{\boldsymbol{\theta}^*\} \in \operatorname*{arg\,min}_{\boldsymbol{\theta}} \sum_{k} \tau_k \, \mathcal{L}^k \left(\mathcal{M}_{\theta^k}(\mathbf{f}^{k-1}) | \, \mathbf{f}_{low}, \mathbf{DB} \right), \qquad (8)$$

$$\mathcal{L}^{k}(\mathbf{f}^{k} | \mathbf{f}_{low}, \mathbf{DB}) = \left\| \mathbf{f}_{low} - \mathbf{DB}\mathbf{f}^{k} \right\|_{2}^{2} + \gamma^{k} \sum_{a=1}^{r} \mathbf{a}_{i}^{k}[a] = 1,$$

where, $\tau_k > 0$ index the weights of the single loss-function of each deep-block, and $\gamma_k > 0$ denote the regularization parameters that control the trade-off between the fidelity-data and the abundance constraint. The output at each deep-block can be interpreted as the estimated HR-HSI; nonetheless, the average between the outputs of the last two deep-blocks were chosen as the estimated HR-HSI to better consider the learned features in the final estimations as, $\hat{\mathbf{f}} = (\mathbf{f}^K + \mathbf{f}^{K-1})/2$.

Figure 1 outlines our DIP-based HSI-SR method, where the HR-HSI is estimated trough multiple interpretable deepblocks inspired by the LMM with two block-layers, each one referring to the abundance and endmembers, respectively.

4. SIMULATIONS AND RESULTS

The conducted experiments employ two public available datasets, the hyperspectral Pavia Center¹, and the Stuff_toys image taken from the spectral CAVE dataset [17] for two downsampling factors, d = 4 and d = 8. The Pavia Center



Fig. 1: Proposed DIP-based single HSI-SR method. A bicubic interpolation of the LR-HSI is used as the input \mathbf{f}_0 of a sequence of deep-blocks shown at the top. The zoomed version of the k^{th} deep-block depicted at the bottom, where the architecture is composed of the abundance block-layer, consisting in a CNN, and the end-member block-layer performing the matrix product according to the LMM. Finally, the general loss-function considers the sum of the single losses employed by the multiple deep-blocks.

was acquired with the ROSIS sensor in 2001. It contains 102 spectral bands and 1096×1096 spatial pixels, where the water vapor absorption and noisy spectral bands were removed from the original 115 observed spectral bands. We used one sub-region of 224×224 spatial pixels for testing in concordance to experiments in [6]. The CAVE dataset consists of 32 spectral images with 512×512 spatial pixels, and 31 spectral bands ranging from 400nm to 700nm at 10nm steps.

The performance of the proposal was compared against several state-of-the-art methods, including the bicubic interpolation; four deep sinlge gray/RGB image SR methods, EDSR [18], RCAN [19], and SAN [20]; four data-driven DLbased single HSI-SR methods, 3DCNN [8], GDRRN [21], and SSPSR [6]; and a non data-driven, DL-based method, DIP [12]. Remark that the proposed method follows a non data-driven framework. The optimization was solved using the the Adam algorithm [22] with a learning rate of $1e^{-3}$, all hyper-parameters γ^k set to 0.5, and a different value for the rank, r in (6), and number of deep-blocks, K, for each dataset through cross-validation. Precisely, we established r = 6 and K = 3, and r = 12 and K = 4, for Pavia Center and Stuff_toys, respectively. All experiments were run on an Intel Xeon W-3223, 64GB of memory, and a NVIDIA RTX 3090 GPU with 24GB of memory².

The quality improvement is quantified through the spectral angle mapper (SAM), the root mean squared error (RMSE), the dimensionless global relative error of synthesis (ERGAS), the peak signal-to-noise ratio (PSNR), and the

¹Available in http://www.ehu.eus/ccwintco/. Accessed: 17-Feb-2021.

²The source code is publicly available in https://github.com/ TatianaGelvez/Interpretable_HSI_SR

structural similarity (SSIM) metrics calculated as in [23, 24].

Tables 1 and 2 compare the quantitative results for the single HSI-SR along the evaluated methods for the Pavia Center and Stuff_toys datasets, respectively. There, it can be seen that our proposed non data-driven method outperforms or achieves

Table 1: Quantitative Results for the Pavia Center Dataset

Method	d	$SAM{\downarrow}$	RMSE↓	ERGAS↓	$\text{PSNR}\uparrow$	SSIM \uparrow
Bicubic	4	6.1399	0.0437	6.8814	27.5874	0.6961
EDSR RCAN SAN	4 4 4	5.8657 5.9785 5.9590	0.0379 0.0376 0.0374	6.0199 6.0485 5.9903	28.7981 28.8165 28.8554	$\begin{array}{c} 0.7722 \\ 0.7719 \\ 0.7740 \end{array}$
3DCNN GDRRN SSPSR	4 4 4	5.8669 5.4750 <u>5.4612</u>	$\begin{array}{c} 0.0396 \\ 0.0393 \\ \underline{0.0362} \end{array}$	6.2665 6.2264 5.8014	28.4114 28.4726 <u>29.1581</u>	$\begin{array}{c} 0.7501 \\ 0.7530 \\ \underline{0.7903} \end{array}$
DIP Proposed	4 4	6.2665 4.2120	0.0410	6.4845 5 8084	28.1061 29 914	0.7365 0.8396
roposea	•	1.2120	0.0002	5.0004	27,717	0.0070
Bicubic	8	7.8478	0.0630	4.8280	24.5972	0.4725
Bicubic EDSR RCAN SAN	8 8 8 8	7.8478 7.8594 7.9992 8.0371	0.0630 0.05983 0.0604 0.0609	4.8280 4.6359 4.6930 4.7646	24.5972 25.0041 24.9183 24.8485	0.4725 0.5130 0.5086 0.5054
Bicubic EDSR RCAN SAN 3DCNN GDRRN SSPSR	8 8 8 8 8 8 8 8 8 8 8 8	7.8478 7.8594 7.9992 8.0371 7.6878 7.3531 7.3312	0.0630 0.05983 0.0604 0.0609 0.0605 0.0607 0.0586	$ \begin{array}{r} \frac{5.6004}{4.8280} \\ 4.8280 \\ 4.6359 \\ 4.6930 \\ 4.7646 \\ 4.6469 \\ 4.6220 \\ 4.5266 \\ \end{array} $	24.5972 25.0041 24.9183 24.8485 24.9336 24.8648 25.1985	0.4725 0.5130 0.5086 0.5054 0.5038 0.5014 0.5365

Table 2: Quantitative Results for the Stuff_toys Dataset

Method	d	$SAM{\downarrow}$	RMSE↓	$ERGAS{\downarrow}$	PSNR↑	SSIM \uparrow
Bicubic	4	4.1759	0.0212	5.2719	34.7214	0.9277
EDSR RCAN SAN	4 4 4	3.5499 3.6050 3.5951	$\begin{array}{c} 0.0149 \\ 0.0142 \\ 0.0143 \end{array}$	3.5921 3.4178 3.4200	38.1575 38.7585 38.7188	0.9522 0.9530 0.9531
3DCNN GDRRN SSPSR	4 4 4	3.3463 <u>3.4143</u> 3.1846	$\begin{array}{c} 0.0154 \\ 0.0145 \\ 0.0138 \end{array}$	3.7042 3.5086 3.3384	37.9759 38.4507 <u>39.0892</u>	0.9522 0.9538 0.9553
DIP Proposed	4 4	8.4935 5.3285	0.0124 0.0110	2.5358 2.1997	38.1329 39.1640	0.9631 0.9821
r		0.0200			0712010	000021
Bicubic	8	5.8962	0.0346	4.2175	30.2056	0.8526
Bicubic EDSR RCAN SAN	8 8 8 8	5.8962 5.6865 5.9771 5.8683	0.0346 0.0279 0.0268 0.0267	4.2175 3.3903 3.1781 3.1437	30.2056 32.4072 32.9544 33.0012	0.8526 0.8842 0.8884 0.8888
Bicubic EDSR RCAN SAN 3DCNN GDRRN SSPSR	8 8 8 8 8 8 8 8	5.8962 5.6865 5.9771 5.8683 5.0948 5.3597 4.4874	0.0346 0.0279 0.0268 0.0267 0.0292 0.0280 0.0257	4.2175 3.3903 3.1781 3.1437 3.5536 3.3460 3.0419	30.2056 32.4072 32.9544 33.0012 31.9691 32.5763 33.4340	0.8526 0.8526 0.8842 0.8884 0.8888 0.8888 0.8863 0.8890 0.9010



Fig. 2: Visualization of some learned features and adjusted weights for d = 4. The output of each deep-block can be effectively interpreted as the endmembers **E** and the abundances \bar{a} .

competitive quality even against the data-driven methods such as the SSPSR for both datasets, both downsampling factors, and all evaluated metrics. Further, we remark that the obtained spectral quality measured with the SAM metric results particularly improved for the hyperspectral Pavia Center dataset. This, because of the low-rank prior can be better exploited when there is a high number of spectral bands that cause higher spectral correlations, which are taken into account within our interpretable deep-blocks. In order to see the additional advange of the proposed inspired network, Fig. 2 depicts the obtained adjusted weights of the second blocklayer **E**, and the features of the first block-layer $\bar{\mathbf{a}}$. There, it can be observed that such outputs can effectively be interpreted as a set of endmembers and abundances, what becomes in additional information that can be useful for further highlevel applications. Finally, Fig. 3 shows a visual comparison of the RGB mapping of some super-resolved images for the Pavia dataset, where the proposed method shows a better spatial quality, particularly for high down-sampling factors.

5. CONCLUSIONS

This paper presented a single HSI-SR method based on the non data-driven DIP approach, where the proposed deep generative model is composed of multiple interpretable deepblocks. The internal architecture is further inspired by the low-rank decomposition given by the LMM, such that, the first block-layer involves a CNN whose output dimensions match the abundances, and the second block-layer involves a matrix product whose adjusted weights match the endmembers. In addition, the general loss-function is composed by the sum of multiple single losses affecting independently each deep-block. The experiments showed that the proposal outperforms state-of-the-art methods, even those data-driven. A better performance was evidenced in images with many spectral bands, where the low-rankness can be fully exploited.



Fig. 3: RGB representation of the reconstructed composite images of Pavia datatest with upsampling factor d = 4 (*top*) and d = 8 (*bottom*). Notice that, the spatial quality is especially improved for a factor d = 8.

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