

# Endmember Distinguished Low-Rank and Sparse Representation for Hyperspectral Unmixing

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July 8, 2024



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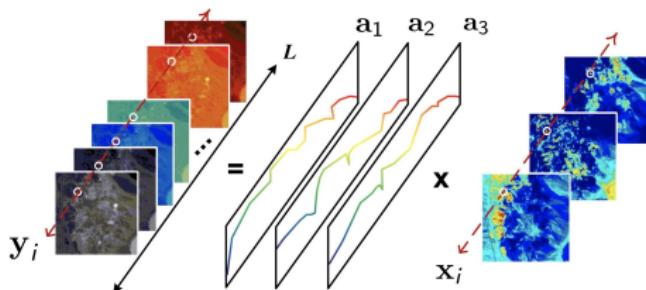
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# Hyperspectral unmixing (HU): Signal model



Courtesy to [Wing-Kin Ma et al. 2014]

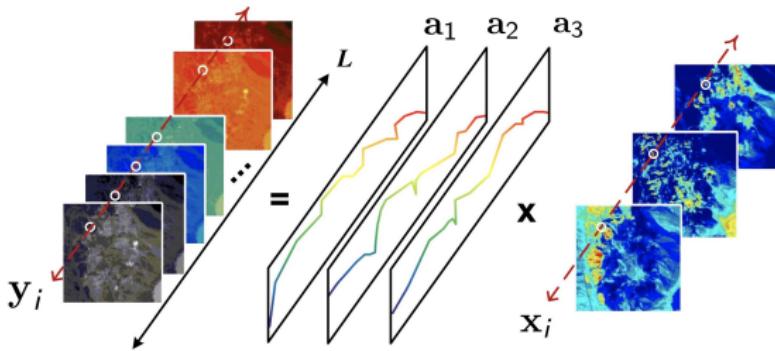
- Linear Mixture Model (LMM):

$$y_i = Ax_i + e_i, \quad i = 1, \dots, k.$$

where

- $A = [a_1, \dots, a_n] \in \mathbb{R}^{m \times n}$ ,  $a_i \in \mathbb{R}^m$  is an **endmember** signature vector;
- $y_i \in \mathbb{R}^m$  is the measured hyperspectral vector at pixel  $i$ ;
- $x_i \in \mathbb{R}^n$  is the **abundance** vector at pixel  $i$ ;  $e_i$  is the noise.





- Let  $Y = [y_1, \dots, y_k]$ ,  $X = [x_1, \dots, x_k]$ ,  $E = [e_1, \dots, e_k]$ . Then

$$Y = AX + E.$$

- **Goal:** recover  $X$  from  $Y$  with knowing  $A$ .

- Assumptions:

- ANC: abundance nonnegative constraint:  $X \geq 0$ , i.e.,  $X_{i,j} \geq 0$ .
  - ASC: abundance sum-to-one constraint:  $\sum_{i=1}^m X_{i,j} = 1$ ,  $\forall j$ .



# Sparse and low-rank regression

## Sparse regression for HU.

- **Idea:** the number of endmembers participating in a mixed pixel is usually very small compared with the (ever-growing) dimensionality (and availability) of spectral libraries.
- The sparse unmixing model [Iordache-Bioucas-Plaza2011] is:

$$\min_{\mathbf{0} \leq \mathbf{X} \in \mathbb{R}^{m \times n}} \frac{1}{2} \|\mathbf{Y} - \mathbf{AX}\|_F^2 + \lambda \|\mathbf{X}\|_1$$

- solve the model by a fast custom-derived solver (e.g., alternating direction method of multipliers (ADMM))
- **SUnSAL:** the sparse unmixing via variable splitting and augmented Lagrangian



## Collaborative sparse regression.

- **Idea:** use a smallest subset of spectral library to represent all mixed pixels in a hyperspectral image.
- The collaborative sparse model [Iordache-Bioucas-Plaza2014] is:

$$\min_{0 \leq X \in \mathbb{R}^{m \times n}} \frac{1}{2} \|Y - AX\|_F^2 + \lambda \|X\|_{2,1}$$

where  $\|X\|_{2,1} = \sum_{i=1}^m \|x^{[i]}\|_2$ , and  $x^{[i]}$  is the  $i$ -th row of  $X$ .

- solve it by ADMM.
- the collaborative SUNSAL (CLSUNSAL) algorithm.



- **SUnSAL**: sparsity on each column vector  $x_j$ , i.e.,  $X$  is sparse.
- **CLSUnSAL**: structural sparsity on  $X$ , i.e.,  $X$  is row-sparse.
  - collaborative sparse on a sliding window [Chen-Nasrabadi-Tran2011], [Qu-Nasrabadi-Tran2014]
  - collaborative sparse with general segments (super-pixels) [Huang-Zhang-Pižurica2017], [Wang-Zhong-Zhang-Xu2017]
- **SUnSAL-TV**: the total variation (TV) regularization is included into the classical sparse regression formulation [Lordache-Bioucas-Plaza2012]

$$\min_{X \geq 0} \frac{1}{2} \|Y - AX\|_F^2 + \lambda \|X\|_1 + \lambda_{TV} TV(X),$$

where  $\lambda \geq 0$  and  $\lambda_{TV} \geq 0$  are regularized parameters.



## Sparse and low-rank unmixing.

- **ADSpLRU**: The spatial correlation among pixels in an HSI translates into a *low-rank* property of the abundance matrix [Giampouras-Themelis-Rontogiannis-Koutroumbas2016].
- The unmixing model is

$$\min_{\mathbf{0} \leq \mathbf{X} \in \mathbb{R}^{m \times K}} \frac{1}{2} \| \mathbf{Y} - \mathbf{AX} \|_F^2 + \lambda \| \mathbf{X} \|_{a,1} + \tau \| \mathbf{X} \|_{f,*}$$

where  $\| \mathbf{X} \|_{a,1} = \sum |a_{i,j} X_{i,j}|$ ,  $\| \mathbf{X} \|_{f,*} = \sum_{i=1}^{\text{rank}(X)} f(\sigma_i) \sigma_i$  is the weighted nuclear norm of  $\mathbf{X}$ , and  $\sigma_i$  is the  $i$ -th singular value of  $\mathbf{X}$ .



- **JSpBLRU**: Partition

$$X = [X_1, \dots, X_s] \in \mathbb{R}^{m \times n},$$

where  $X_j \in \mathbb{R}^{m \times d_j}$ , for  $j = 1, \dots, s$ ,  $\sum_{j=1}^s d_j = n$ , and block number  $s$  is a positive integer for  $1 \leq s \leq n$ . Each  $X_j$  is assumed joint-sparse [Huang-Huang-Deng-Zhao2019].

- Consider the joint-sparse-blocks and low-rank hyperspectral unmixing model

$$\min_{\mathbf{0} \leq X \in \mathbb{R}^{m \times K}} \frac{1}{2} \|Y - AX\|_F^2 + \lambda \sum_{j=1}^s \|X_j\|_{w_j, 2, 1} + \tau \|X\|_{f,*}$$

where the *weighted  $\ell_{2,1}$  norm*, defined as  $\|X_j\|_{w_j, 2, 1} = \sum_{i=1}^m w_{i,j} \|X_j^{[i]}\|_2$ , is used to enhance sparsity along rows in each block in  $X$ ,  $X_j^{[i]}$  is the  $i$ -th row of the  $j$ -th block of  $X$ ,  $w_j = [w_{1,j}, \dots, w_{m,j}]^T$  is a nonnegative vector.

- **BiJSpLRU**: Consider the mode-3 matricization of  $\mathcal{Y}$  along the vertical and horizontal directions simultaneously. [Huang-Di-Wang-Lin-Huang2021]



# Endmember distinguished methods

Idea: divide the endmembers into two groups.

- “Low-rank” endmembers: they participate in the mixing process and are called *active*. The corresponding abundances are gathered in  $X_1 \in \mathbb{R}^{n_1 \times k}$  and we will exploit more spatial information of abundance maps (although sparsity is also important for them).
- “Sparse” endmembers: they are *inactive*, resulting in the sparsity of  $X$ . The corresponding abundances are gathered in  $X_2 \in \mathbb{R}^{n_2 \times k}$ .
- Divide the abundance matrix  $X \in \mathbb{R}^{n \times k}$  into two row blocks according to the activity. That says

$$PX = \begin{bmatrix} X_1 \\ X_2 \end{bmatrix},$$

where  $P = [P_1^T, P_2^T]^T \in \mathbb{R}^{n \times n}$  is a permutation matrix used to reorder the rows of the abundance matrix,  $P_1 X = X_1$  and  $P_2 X = X_2$  are composed of rows corresponding to  $n_1$  low-rank endmembers and  $n_2$  sparse endmembers with  $n_1 + n_2 = n$ , respectively.



# Endmember distinguished methods

## How to distinguish the low-rank and sparse endmembers?

- The key is to determine the permutation matrix  $P$ .
- Step 1. Sort the endmembers in descending order by  $\|X(i, :) \|_2$ , where  $X(i, :)$  denotes the  $i$ -th row of  $X$  for  $i = 1, 2, \dots, n$ .
- Step 2. Use  $\ell_2$ -norm to measure activity. We find the set  $S$  of the top  $K$  endmembers satisfying that

$$\sum_{i \in S} \|X(i, :) \|_2 \geq \rho \sum_{i=1}^n \|X(i, :) \|_2, \quad (1)$$

where  $\rho \in [0, 1]$  is a thresholding parameter.

- Step 3. The endmembers in set  $S$  will be considered as *low-rank* endmembers and the rest will be *sparse* ones.



# Weighted nuclear norm regularization

- We apply the weighted nuclear norm regularization on the **abundance map of each low-rank endmember**, instead of applying on  $X_1$  directly, which can be described as

$$\sum_{i=1}^{n_1} \|Re(X_1(i,:))\|_{g,*}$$

where  $n_1$  is the number of low-rank endmembers,  $Re(\cdot)$  is a function that reshapes the row vector of the  $i$ -th low-rank endmember into abundance map.

- Here the weighted nuclear norm is defined as

$$\|T\|_{g,*} = \sum_i g(\sigma_i) \sigma_i,$$

where  $\sigma_i$  is the  $i$ -th singular value of the matrix  $T$  and  $g$  is a weighting function for singular values defined as  $g(\sigma) = 1/(\sigma + \varepsilon)$ , and  $\varepsilon$  is a small number to avoid singularities [Gu-Xie-Meng-Zuo-Feng-Zhang2017].



# Weighted sparse regularization

- We consider the sparse property for the abundance matrix  $X$ .
- In particular, we introduce a weight factor  $B \in \mathbb{R}^{n \times k}$  into the  $\ell_1$ -norm of  $X$ .
- The weight factor  $B$  consists of a spectral factor  $B_1 \in \mathbb{R}^{n \times k}$  and a spatial factor  $B_2 \in \mathbb{R}^{n \times k}$  to make full use of spectral and spatial information.  
[Zhang-Li-Li-Deng-Plaza2018]
- The weighted  $\ell_1$ -norm regularization on  $X$  is as follows

$$\|B \odot X\|_1, \quad B = \text{sqrt}(B_1 B_2), \quad (2)$$

where the operator  $\odot$  denotes the Hadamard product and  $\text{sqrt}(\cdot)$  is the elementwise square root function.



# Weighted sparse regularization

- The **spectral factor**  $B_1$  is used to enhance the sparsity of the endmembers in the spectral library, which is defined as

$$B_1 = \text{diag} \left[ \frac{k}{\|X(1,:) \|_1 + \varepsilon}, \dots, \frac{k}{\|X(n,:) \|_1 + \varepsilon} \right].$$

- The **spatial factor**  $B_2$  utilizes the spatial correlation between the pixel and its neighbors to promote sparsity, which can be described as

$$B_2(i,j) = \frac{\sum_{k \in \mathcal{N}(j)} w_k}{\sum_{k \in \mathcal{N}(j)} w_k X(i,k) + \varepsilon},$$

where  $\mathcal{N}(j)$  is a collection of the column indices of the  $j$ -th pixel's neighbors (including itself) in the abundance matrix and  $w_k$  is the corresponding weight of neighbors.



# Proposed model

- In conclusion, our proposed unmixing model is given as

$$\begin{aligned} \min_X \frac{1}{2} \|Y - AX\|_F^2 + \lambda \sum_{i=1}^{n_1} \|Re(X_1(i,:))\|_{g,*} + \tau \|B \odot X\|_1 \\ \text{s.t. } X \geq 0, \quad PX = \begin{bmatrix} X_1 \\ X_2 \end{bmatrix}, \end{aligned}$$

- To solve this model under the ADMM framework. We transform this model into a constrained problem as

$$\begin{aligned} \min_X \frac{1}{2} \|Y - AX\|_F^2 + \lambda \sum_{i=1}^{n_1} \|Re(V_1(i,:))\|_{g,*} + \tau \|B \odot V_2\|_1 + \iota_{R_+}(V_3) \\ \text{s.t. } P_1X = V_1, \quad X = V_2, \quad X = V_3, \end{aligned}$$

where  $V_1, V_2, V_3$  are auxiliary variables,  $\iota_{R_+}(x)$  is the indicator function, and  $\tau$  are nonnegative regularization parameters.



# Proposed algorithm

- Let  $\Omega_1 \in \mathbb{R}^{n_1 \times k}$ ,  $\Omega_2 \in \mathbb{R}^{n \times k}$ , and  $\Omega_3 \in \mathbb{R}^{n \times k}$  are Lagrange multipliers.
- Define  $G = [P_1^T, I, I]^T$ ,  $E = \text{diag}(-I, -I, -I)$ ,  $V = [V_1^T, V_2^T, V_3^T]^T$ ,  $\Omega = [\Omega_1^T, \Omega_2^T, \Omega_3^T]^T$ , then the constraints become  $GX + EV = 0$ .
- Define

$$\begin{aligned}\mathcal{L}_\mu(X, V; \Omega) = & \frac{1}{2} \|Y - AX\|_F^2 + \lambda \sum_{i=1}^{n_1} \|Re(V_1(i, :))\|_{g,*} \\ & + \tau \|B \odot V_2\|_1 + \iota_{R_+}(V_3) + \frac{\mu}{2} \|GX + EV - \Omega\|_F^2,\end{aligned}$$

where  $\mu > 0$  is a penalty parameter.

- The ADMM framework is derived

$$\begin{cases} X^{k+1} = \underset{X}{\operatorname{argmin}} \mathcal{L}_\mu(X, V^k; \Omega^k), \\ V^{k+1} = \underset{V}{\operatorname{argmin}} \mathcal{L}_\mu(X^{k+1}, V; \Omega^k), \\ \Omega^{k+1} = \Omega^k - (GX^{k+1} + EV^{k+1}). \end{cases}$$



# Proposed algorithm

- To make the notations clearly, we introduce the soft-thresholding (SHR) operator and singular value thresholding (SVT) operator.
- Let  $X = U\Sigma V^T$  be the singular value decomposition of  $X$  and recall the weighting function. Define

$$\text{SHR}_{g,\alpha}(x) = \text{sign}(x) \max(0, x - \alpha g(x)),$$

$$\text{SVT}_{g,\beta}(X) = U\text{SHR}_{g,\beta}(\Sigma) V^T.$$

- Now we can propose the final algorithm named as Endmember Distinguished Low-rank and Sparse Representation Unmixing (EDLSpRU).



# Proposed algorithm

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**Algorithm 1:** EDLSpRU

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**Input:**  $Y$  and  $A$ .

**Selected parameters:**  $\lambda, \tau, \mu$  and maximum iterations

**Initialization:**  $X^0, V^0, \Omega^0$ , and  $t = 1$

**Repeat:**

$$X^t = (A^T A + 2\mu I + P_1^T P_1)^{-1} [A^T Y + \mu P_1^T (V_1^{t-1} + \Omega_1^{t-1}) + \mu (V_2^{t-1} + \Omega_2^{t-1} + V_3^{t-1} + \Omega_3^{t-1})]$$

for  $i = 1, 2, \dots, n_1$

$$Re(V_1^t(i, :)) = SVT_{g, \frac{\lambda}{\mu}}(Re(P_1 X^t(i, :) - \Omega_1^{t-1}(i, :)))$$

end for

$$V_2^t = SHR_{B, \frac{\tau}{\mu}}(X^t - \Omega_2^{t-1}) \text{ with } B \text{ calculated by (2)}$$

$$V_3^t = \max(X^t - \Omega_3^{t-1}, 0)$$

$$\Omega_1^t = \Omega_1^{t-1} - P_1 X^t + V_1^t$$

$$\Omega_2^t = \Omega_2^{t-1} - X^t + V_2^t$$

$$\Omega_3^t = \Omega_3^{t-1} - X^t + V_3^t$$

Update  $P_1$  and  $n_1$  according to the set  $S$  satisfying (1)

**until** some stopping criterion is satisfied.

**Output:**  $\hat{X} = X^t$

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# Experiments on simulated data

## Experiment setting.

- We will compare EDLSpRU with three state-of-the-art algorithms: ADSpLRU, JSpBLRU and BiJSpLRU.
- RMSE and the signal-to-reconstruction error (SRE) are used to evaluate the performance of unmixing results. They are defined by, respectively,

$$\text{RMSE} = \sqrt{\frac{1}{nk} \sum_{i=1}^k \|\hat{x}_i - x_i\|_2^2}, \quad \text{SRE (dB)} = 10 \log_{10} \left( \frac{\frac{1}{k} \sum_{i=1}^k \|\hat{x}_i\|_2^2}{\frac{1}{k} \sum_{i=1}^k \|\hat{x}_i - x_i\|_2^2} \right),$$

where  $k$  is the number of pixels,  $n$  is the number of endmembers, and  $\hat{x}_i$  and  $x_i$  are estimated and exact abundance vectors of the  $i$ -th pixel, respectively.

- We set the maximum number of iterations to 500.
- We select optimal regularization parameters for EDLSpRU as follows:

$$\lambda, \tau \in \{0.0001, 0.0005, 0.001, 0.005, 0.01, 0.05, 0.1, 0.5, 1, 5\},$$

and select  $\mu \in \{0.1, 0.5, 1, 5\}$ . In addition, we set  $\rho = 0.9$  in all experiments. All possible combinations are considered and the optimal parameters are chosen to get maximum SREs.

- Our test were done by using MATLAB R2021a on a laptop with 2.30 GHz Intel Core i7 and 12 GB memory.



# Experiments on simulated data

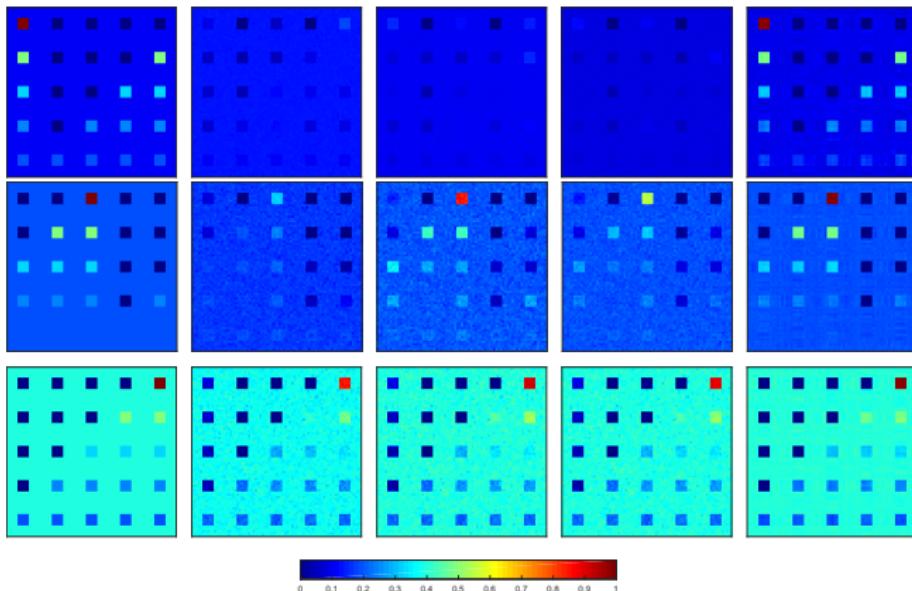
- Data Cube 1 (DC1): The first data cube contains  $75 \times 75$  pixels with 224 spectral bands. The spectral dictionary  $A_1 \in \mathbb{R}^{224 \times 240}$  is extracted from the U.S. Geological Survey (USGS). Five endmembers are randomly chosen from  $A_1$  according to LMM. Finally, the generated data cube is contaminated by white Gaussian i.i.d. noise with signal-to-noise ratio (SNR) of 20, 30, and 40dB respectively.
- Data Cube 2 (DC2): This data cube contains  $100 \times 100$  pixels with 99 spectral bands and the spectral dictionary  $A_2 \in \mathbb{R}^{99 \times 120}$  is from the National Aeronautics and Space Administration Johnson Space Center Spacecraft Materials Spectral Database. Nine endmember signatures are randomly chosen from  $A_2$  and the corresponding abundance maps are used to generate the true data cube, which is also contaminated by Gaussian noise with the same SNR values adopted for DC1.



Table: SRE(dB) and RMSE by different algorithms.

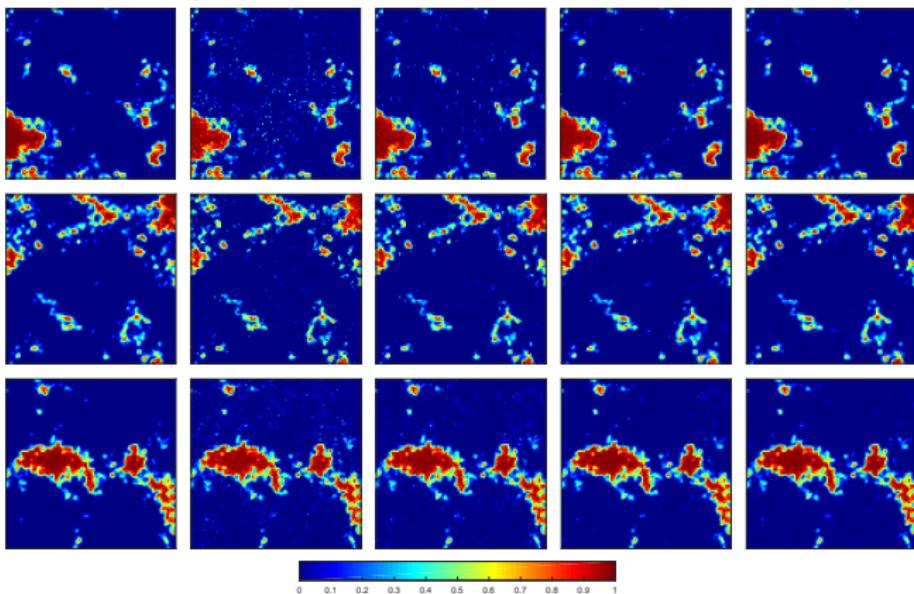
Algorithm	Data Cube 1 (DC1)					
	SNR = 20 dB		SNR = 30 dB		SNR = 40 dB	
	RMSE	SRE	RMSE	SRE	RMSE	SRE
ADSPlRU	0.0168	6.24	0.0079	12.82	0.0014	27.70
JSpBLRU	0.0117	9.42	0.0059	15.40	0.0008	32.50
BiJSpLRU	0.0115	9.53	0.0067	14.30	0.0008	32.82
EDLSpRU	<b>0.0108</b>	<b>10.13</b>	<b>0.0016</b>	<b>26.55</b>	<b>0.0005</b>	<b>36.70</b>
Algorithm	Data Cube 2 (DC2)					
	SNR = 20 dB		SNR = 30 dB		SNR = 40 dB	
	RMSE	SRE	RMSE	SRE	RMSE	SRE
ADSPlRU	0.0375	5.75	0.0114	16.10	0.0034	26.66
JSpBLRU	0.0230	9.98	0.0087	18.42	0.0027	28.44
BiJSpLRU	0.0180	12.12	0.0069	20.48	<b>0.0024</b>	<b>29.55</b>
EDLSpRU	<b>0.0155</b>	<b>13.39</b>	<b>0.0058</b>	<b>22.01</b>	0.0029	28.02





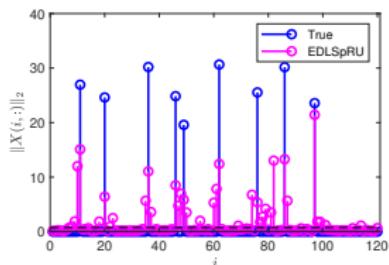
**Figure:** True and estimated abundance maps for endmembers #1, #3 and #5 by different unmixing algorithms for DC1 with SNR = 30 dB. From left to right: True, ADSpLRU, JSpBLRU, BiJSpLRU, and EDLSpRU.



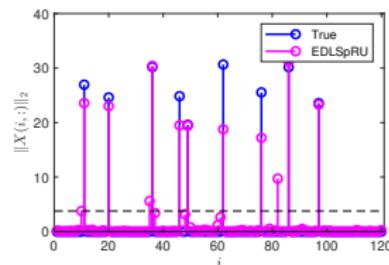


**Figure:** True and estimated abundance maps for endmembers #2 , #3 and #9 by different unmixing algorithms for DC2 with SNR = 30 dB. From left to right: True, ADSpLRU, JSpBLRU, BiJSpLRU, and EDLSpRU.

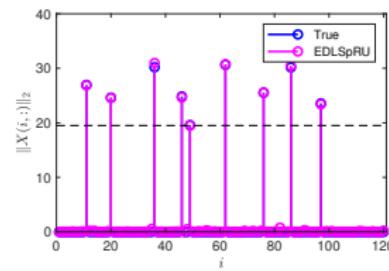




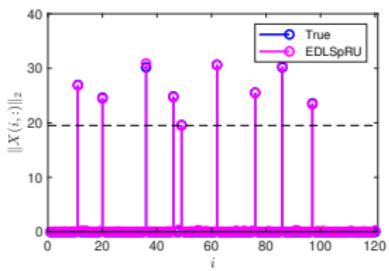
(a) 10 iters



(b) 50 iters



(c) 100 iters



(d) 200 iters

**Figure:** Comparison of each endmember's abundance of the ground truth and the estimations by EDLSprU for DC2 with SNR = 30 dB.



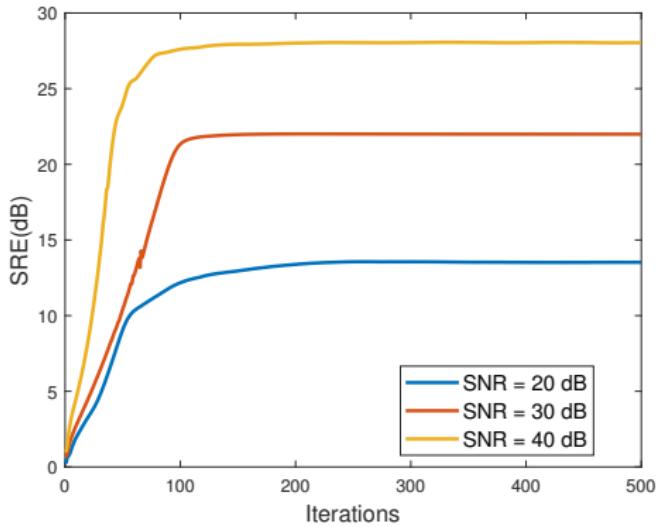


Figure: Plots of SRE (dB) against iterations by EDLSpRU for DC2.



# Experiments on real data

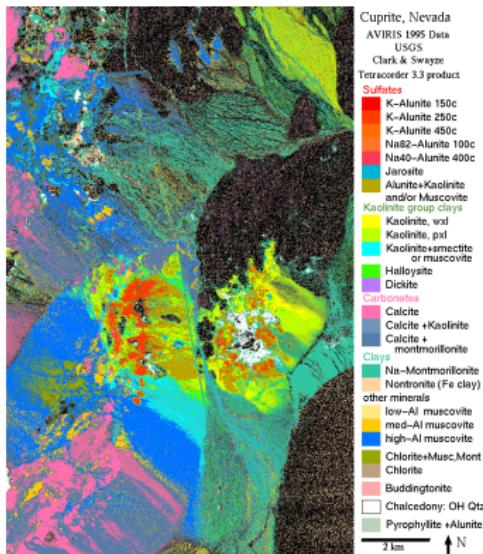
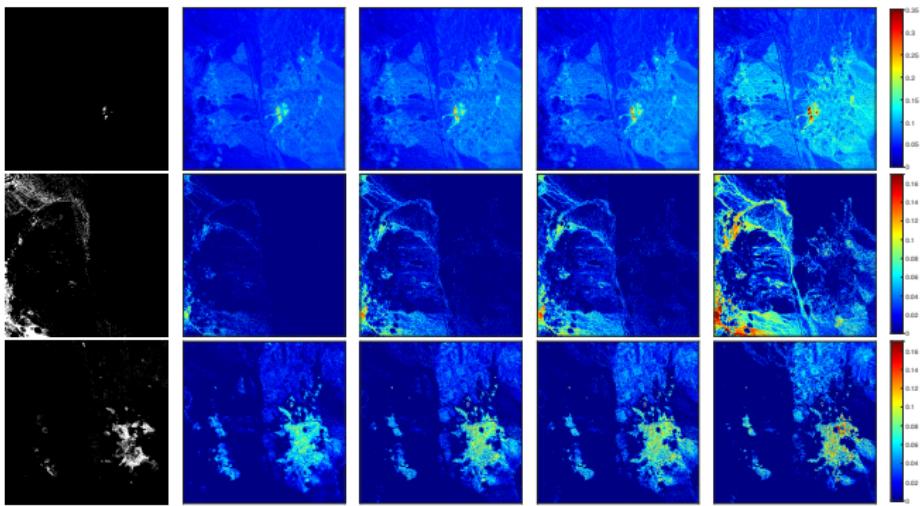


Figure: USGS map showing the location of different minerals in the Cuprite mining district in Nevada.





**Figure:** Estimated abundance maps for minerals buddington, muscovite, and chalcedony by different unmixing algorithms. From left to right: Tetracorder, ADSpLRU, JSpBLRU, BiJSpLRU, and EDLSpRU.



# Conclusion and future work

- We have proposed a new unmixing model which can distinguish endmembers into low-rank ones and sparse ones so that different regularizations can be applied to each type of them. The model aims at considering both low-rank and sparse characteristics for low-rank endmembers, while only considering sparsity for sparse ones.
- We solve our model under the ADMM framework and propose a new algorithm named as EDLSpRU. The experiment results for both simulated and real-data demonstrate the efficacy of the proposed unmixing structure.
- In the future, we will extend the proposed endmember distinguished structure to blind unmixing or its tensor edition for HSI processing.



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# Thank you very much!

