

IRAP PhD Day 2019

Taking in account panchromatic data in hyperspectral image unmixing

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An hyperspectral image

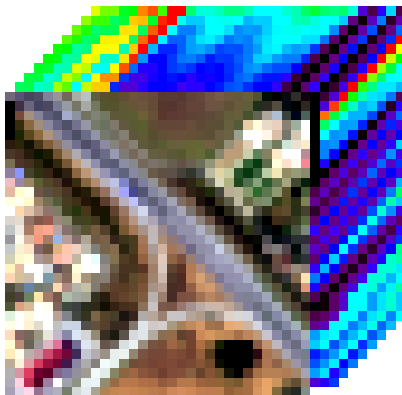
Acquiring a scene with several hundred of spectral bands

Expectation:



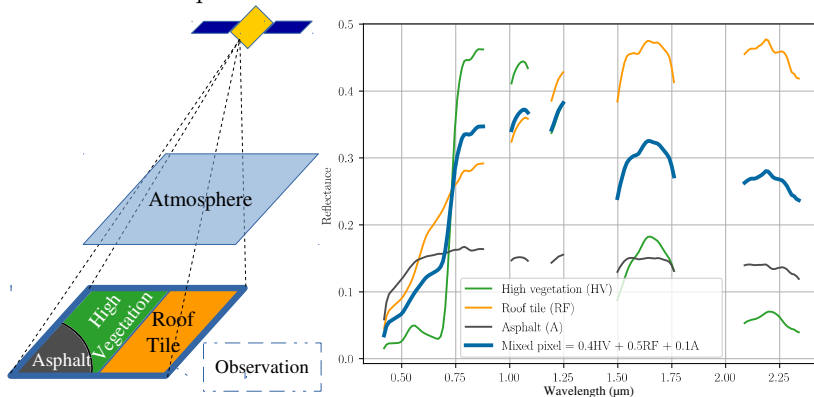
An hyperspectral image

Acquiring a scene with several hundred of spectral bands
Reality:



Problem statement

Due to the spatial sampling, a hyper-spectral photosite acquires a mix of the material spectra in the covered area.



Problem statement

Example: linear mixing model

$$\vec{y}_i = \sum_{k=1}^P x_{ki} \vec{s}_k + \vec{w}$$

- ✏ The pixel i $\vec{y} \in \mathbb{R}_+^L$
- ✏ The abundance fractions $\vec{x}_k \in \mathbb{R}_+^P$
- ✏ The endmembers $\vec{s}_k \in \mathbb{R}_+^L$
- ✏ A noise $\vec{w}$

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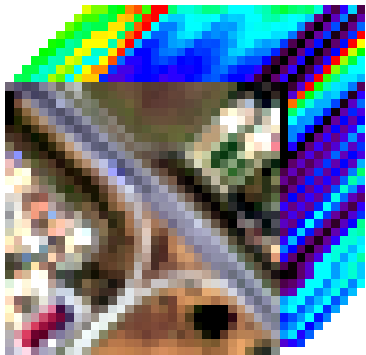
Unmixing goal

Blindly extract or estimate the endmembers $\vec{s}_k$ as well as their abundance fractions $\vec{x}_{k,i}$

Adding a panchromatic image

Since 2011 the CNES studies the possible outcomes of shipping two co-registered instrument:

- ➔ an hyperspectral camera covering the $[0.4 - 2.5]$ μm range with 10 nm spectral step and a 8 m GSD.
- ➔ a 2 m GSD panchromatic camera.



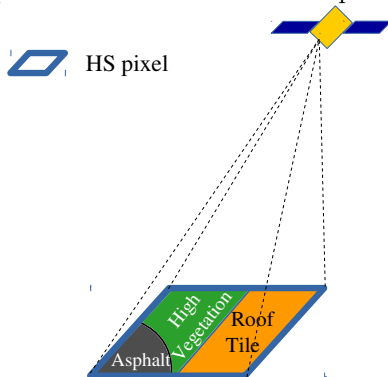
Hyperspectral image (HSI) - 8 m GSD



Panchromatic image (Pan) - 2 m GSD

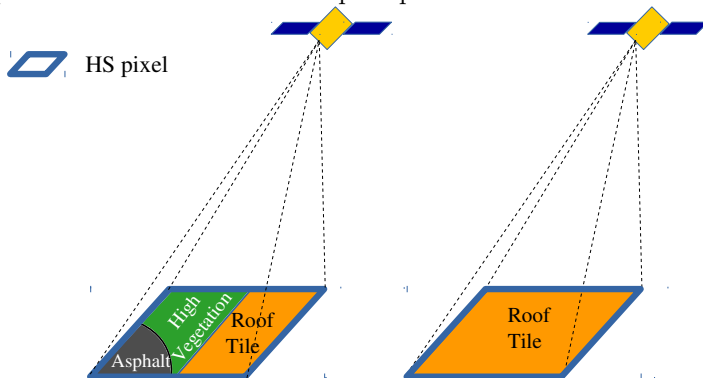
Proposed approach

Homogeneity-Based Endmember Extraction (HBEE): A homogeneous panchromatic area leads to a pure pixel



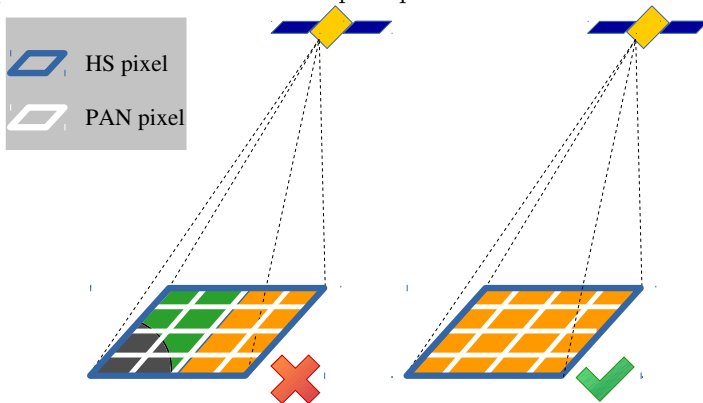
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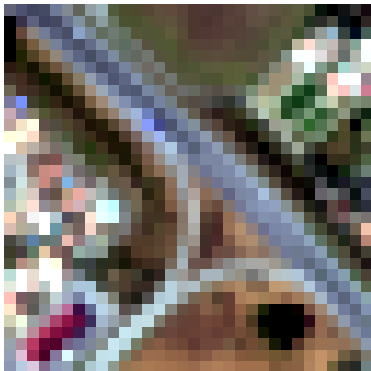


Proposed approach

Homogeneity-Based Endmember Extraction (HBEE): A homogeneous panchromatic area leads to a pure pixel



Homogeneous pixel detection

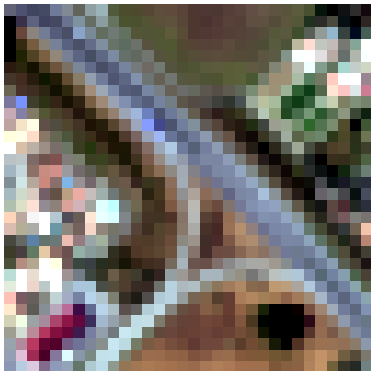


HS image with an 8m GSD

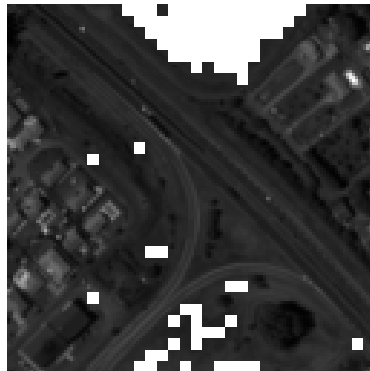


Pan image with a 2m GSD

Homogeneous pixel detection

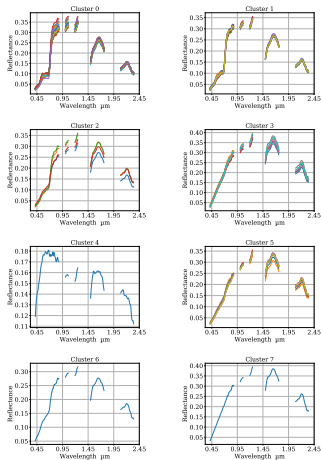


HS image with an 8m GSD

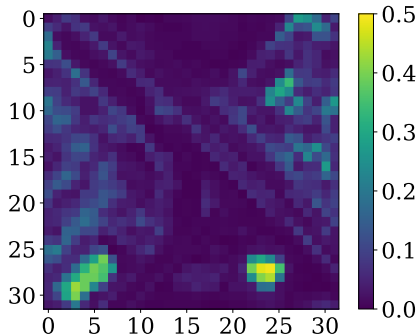


Detected homogeneous area

HBEE results



Pure spectra clusters



Reconstruction Error (RE) map after the HBEE stage:

$$RE = \frac{\| \hat{y}_{ij} - y_{ij} \|}{\| \hat{y}_{ij} \|}$$

Proposed approach - LSNMF stage

A poorly reconstructed area contains a yet unextracted endmember.

Locally-Semi-supervised Non-negative Matrix Factorisation (LSNMF)

- 1 Calculation of the RE map.
- 2 Selection of the most poorly reconstructed area.
- 3 Initialisation of the endmember and abundance fraction matrices.
- 4 Estimation of an unknown endmember by applying an NMF onto the area.
- 5 Iterate until all the previously detected areas are correctly rebuilt.

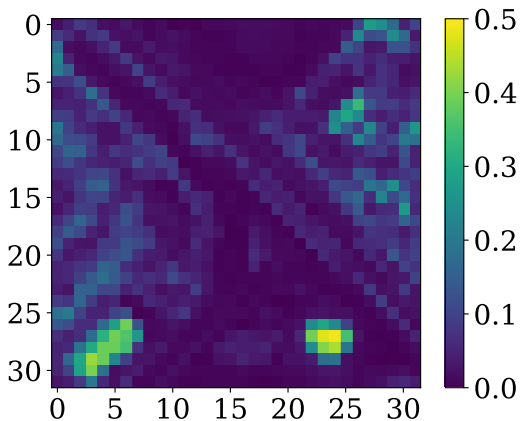
NMF

To estimate the endmember, the multiplicative update strategy is considered here :

$$X_{k+1} \leftarrow X_k \frac{S_k^T Y}{S_k^T S_k X_k + \epsilon} \quad S_{k+1} \leftarrow S_k \frac{Y X_k^T}{S_k X_k X_k^T + \epsilon}$$

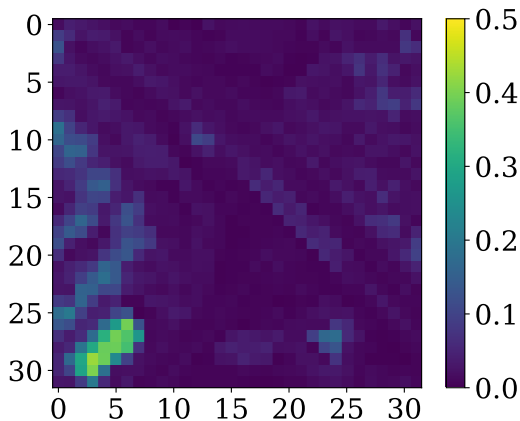
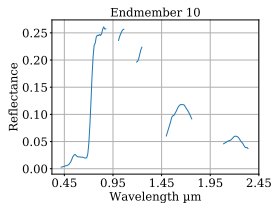
LSNMF results

Iteration 0



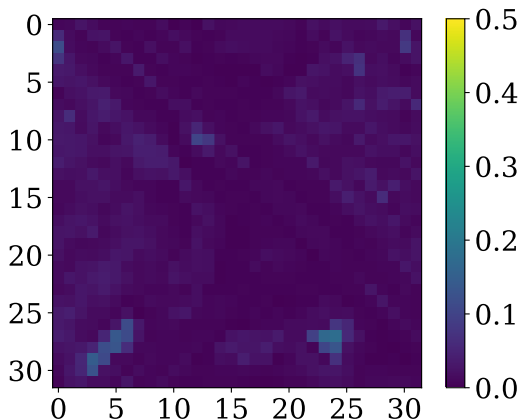
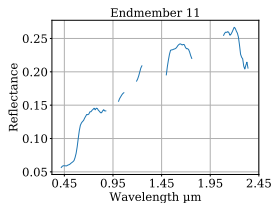
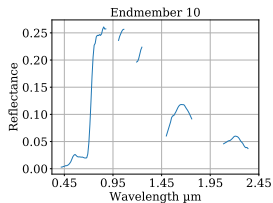
LSNMF results

Iteration 1



LSNMF results

Iteration 2



Conclusion

Limitations and possible enhancements

- The parameters depend on the image

Contribution

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- HBEE extracts truly pure spectra as endmembers.

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- HBEE extracts truly pure spectra as endmembers.
- LSNMF estimates satisfactory unextracted endmembers.

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- The overall performances are equivalent to others state of the art methods in a spectral sense.

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Contribution

- HBEE extracts truly pure spectra as endmembers.
- LSNMF estimates satisfactory unextracted endmembers.
- The overall performances are equivalent to others state of the art methods in a spectral sense.
- Naturally estimates a satisfactory number of endmember



IRAP, Université de Toulouse, CNRS, UPS, CNES, Toulouse || ONERA The French Aerospace Lab, Toulouse || *CNES, Toulouse

4. Andrew Veith and Barbara Kuster, "Quick shift and leaved methods for multi-unfolding," in *Computer Vision - ECCV 2004*, David Forsyth, Philip Torr, and Andrew Zisserman, E. Sarti, Heidelberg, 2004, pp. 705–718, Springer Berlin.

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Main unmixing methods

Several strategies :

Endmember estimation

- ➔ Finding pure pixels : VCA, ATGP, NFINDR.
- ➔ Minimizing the volume of the simplex containing the data and induced by the endmembers : MVSA, SISAL, MVC-NMF.
- ➔ Estimating the endmembers by a Bayesian approach.
- ➔ The Non-negative Matrix Factorization (NMF).

Abundance fraction estimation knowing the endmembers

- ➔ Least-Square
- ➔ Non-negatively constrained Least-Square (NCLS)
- ➔ Fully-Constrained Least-Square (FCLS)

Spectral clustering and selection

Clustering of the extracted spectra

A simple unsupervised algorithm will iteratively build p clusters of the extracted pure spectra using a Spectral Angle Measurement (SAM) criterion :

$$\theta = \left(\frac{\langle s_k, s_l \rangle}{\|s_k\| \cdot \|s_l\|} \right)$$

Endmembers selection

The chosen endmembers s_k are the ones with the best homogeneity score for each class.

Abundance fraction estimation

The abundance fractions matrix X is estimated using a Non-negatively Constrained Least-Square algorithm (NCLS)

Annexes

Clustering spectra

- 1 The pure spectrum list is sorted by their homogeneity level
- 2 Création d'une classe C_k à partir du premier spectre de la liste.
- 3 Agrégation à C_k tous les spectres de la listes présentant une valeur de SAM avec le premier spectre inférieure à ϵ_{SAM} et suppression de ces spectres de la liste.
- 4 Recommencer à l'étape 1 jusqu'à ce que la liste soit vide.

Annexes

Reconstruction error map criterion

$$RE = \frac{||\hat{y}_{ij} - y_{ij}||}{||\hat{y}_{ij}||} \quad (1)$$

Annexes

