

How to mix spatial and spectral information when processing hyperspectral images

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Presentation summary

- ⇒ Introduction: Hyperspectral images and spatial information
- ⇒ Markov Random Fields
- ⇒ Split-and-merge and Unin-Find strategies
- ⇒ Anisotropic diffusion
- ⇒ Conclusion

Hyperspectral imaging at Cemagref

Spectrometry & chemometrics (J-M Roger et al)

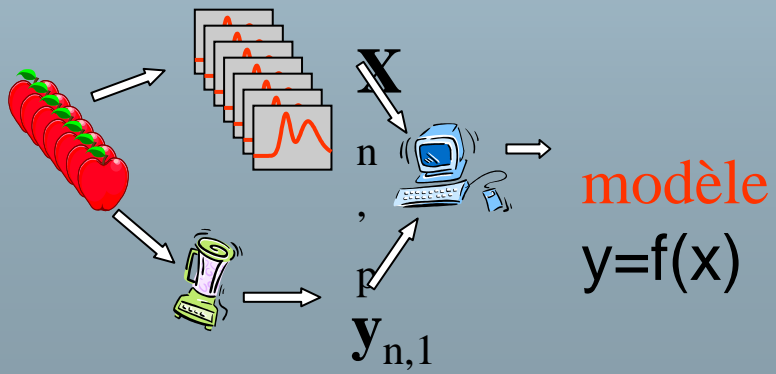
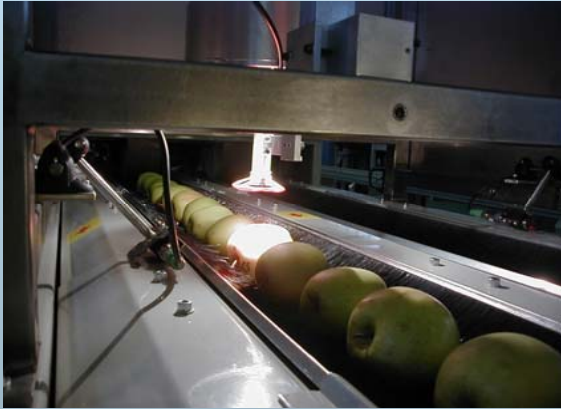
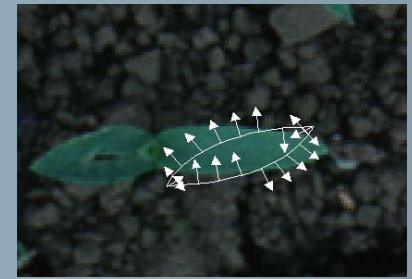
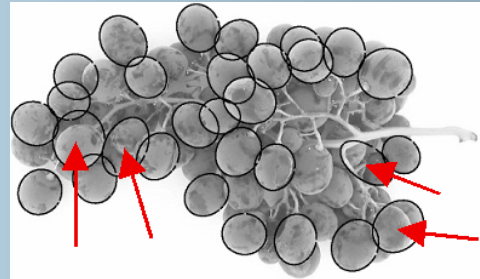


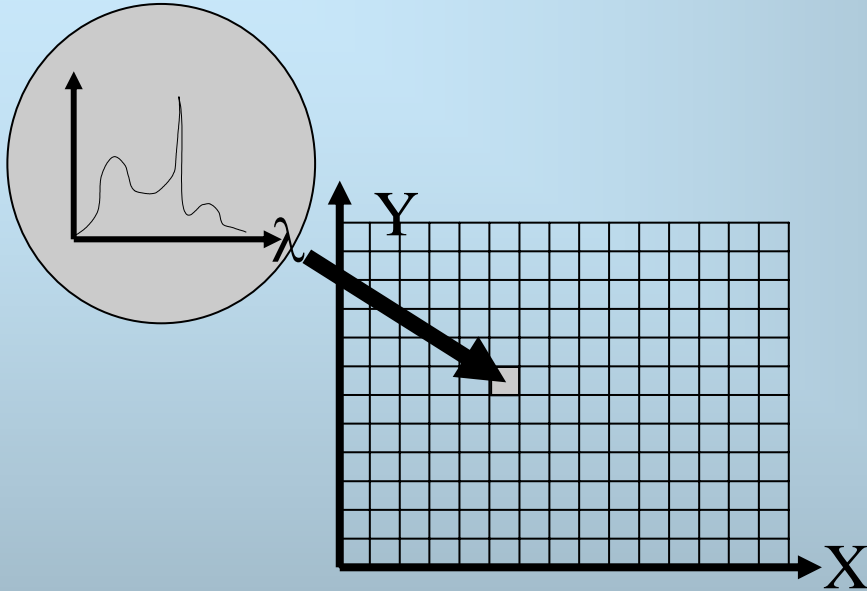
Image processing (G. Rabatel et al)



« Proximity » hyperspectral imaging



Hyperspectral imaging in a few words



⇒ Compared to standard imagery:
➤ More pixel information → better classification potential

⇒ Compared to spectrometry:
➤ Spatial mapping of the data

⇒ Some applications:

- Image segmentation (urban areas, forests, etc.)
- Object detection (e.g. military applications)
- Mapping of chemical components (pharmacy, biology, geology...)

The basics of hyperspectral processing

⇒ Two main approaches

- Supervised learning:
output/spectra relationship learned from examples
- Unsupervised processing:
clusterisation, pure spectra extraction...

⇒ A common requirement: model parcimony

- Looking for a data subspace that keeps class separability
PLS (supervised) ; PCA, projection pursuit (unsupervised)

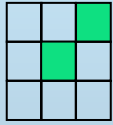
The present situation

- ⇒ In almost cases, **pixel-based** processing
 - Pixels are just considered as a large set of samples
 - Their neighbourhood relations are not taken into account:
 - neither in the calibration step
 - nor in the classification step
 - A spatial mixing of the pixels would lead to the same processing and the same results

- ⇒ A simple question: can we do better by considering pixels as « picture elements », not only as samples ?

Spatial relationships

Basic notion:
neighbourhood



Texture



Homogeneity



Region:
set of connected pixels
with the same attributes
(level or texture)

Shape



Homogeneity constraint

- ⇒ Every pixel in the image have a high *a priori* probability to have spectral attributes close to its' neighbour ones

- ⇒ Can be used to:
 - overcome classification ambiguities
 - remove erroneous pixels (outsiders)
 - Improve pixel clusterisation

Example 1: Markov Random Fields (MRF)

Markov Random Field: definition

- ⇒ Let us consider a 2D image $I(x,y)$
(I : attribute vector, attribute scalar, or label)
- ⇒ Let us consider $p(I(i,j) = x_s \ / \ I(k,l) \{k \neq i, l \neq j\})$
Probability for the pixel (i,j) to have a value x_s , knowing every other values in the image

If $I(x,y)$ is a Markov Random Field, this probability only depends on (i,j) neighbourhood, and can be written as:

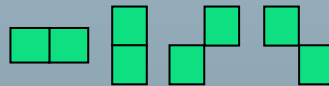
$$p(x_s) = \frac{1}{Z} \exp(-\sum_C V(C))$$

- ⇒ Z : normalisation constant
- ⇒ $V(C)$: potential of the clique C
- ⇒ C : clique or neighbourhood configuration

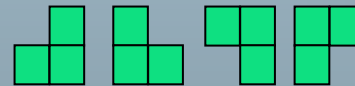
Cliques:



1st order



2nd order



3rd order



4th order

Can be used for texture modelisation

MRF for homogeneity regularisation

The Potts model: limited to cliques of 1st and 2nd orders

⇒ First order potential:

$$V(C1) = \text{Log} [\text{Pr}(x_s / S)]$$

$\text{Pr}(x_s / S)$: probability to have a label x_s for a spectrum S

Gaussienne hypothesis:
$$V(C1) = (S - \mu_{x_s})^T \cdot \text{COV}^{-1} \cdot (S - \mu_{x_s})$$

⇒ Second order potential:

$$\begin{aligned} V(C2) = V(x_s, x_t) &= -\beta \text{ if } x_s = x_t \\ &= +\beta \text{ if } x_s \neq x_t \end{aligned}$$

β parameter: tuning of the regularisation effect

MRF implementation


⇒ The problem: to find the output image corresponding to the maximal probability $\text{pr}(x_s)$ for every pixel

➤ Simulated annealing (optimal, heavy computation)

- Introduction of a temperature parameter
- Slow decreasing of T to zero.

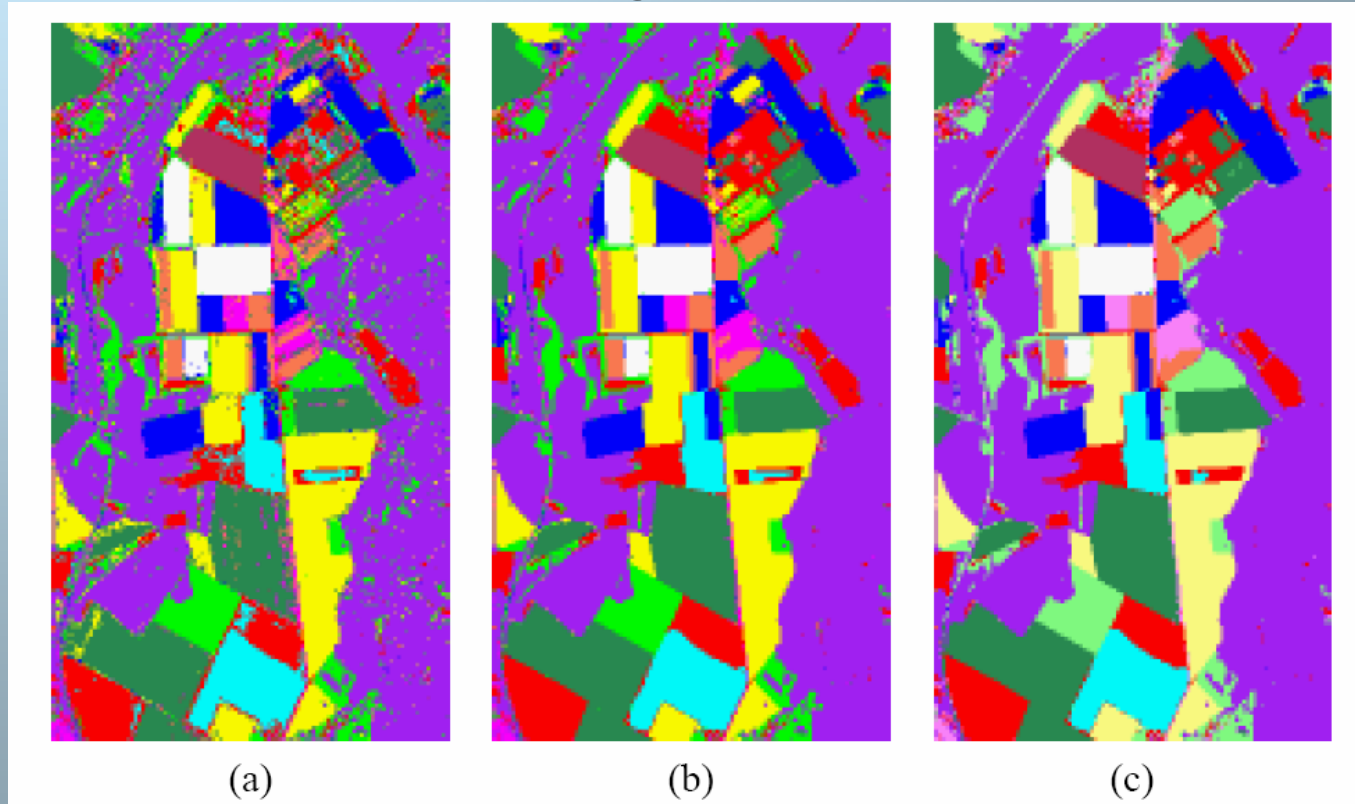
$$p(x_s, T) = \frac{1}{Z(T)} \exp\left(-\frac{\sum_C V(C)}{T}\right)$$

➤ Algorithm ICM (sub-optimal, faster)

- 
- Image scanning
 - For each pixel, setting of the x_s value with the higher probability $\text{pr}(x_s)$
 - Re-iteration until a stationary state is reached

Hyperspectral example (Prony et al, 2000)

ICM regularisation



a) No regulation

b) $\beta = 3$

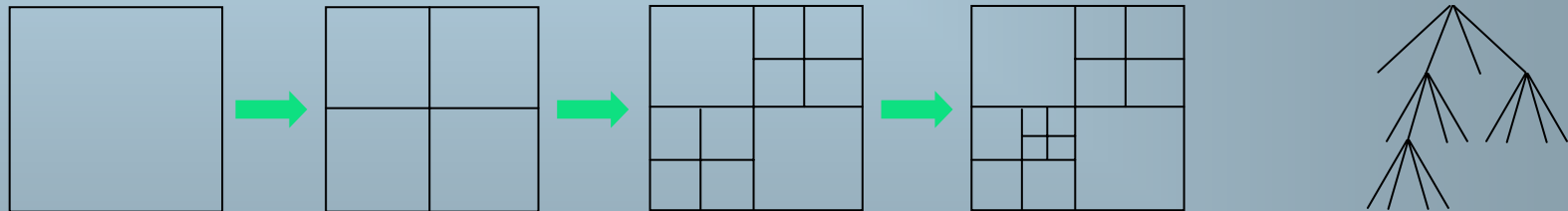
c) $\beta = 5$

Unsupervised segmentation based on split-and-merge

N. Gorretta, Cemagref
(under development)

Basic principles

- ⇒ Objective: to partition the hyperspectral image in a set of homogeneous regions
- ⇒ Splitting process:
Building of a quad-tree
 - Root = initial image
 - For each tree node: children nodes are created as long as the subimage variance is too high



- ⇒ Merging process:
Adjacent regions are merged if the distance between their average pixel values is below a threshold

Implementation

⇒ Before the split-and-merge segmentation

- a PCA is made on the spectral data of the subregion
- Score images are builded using a grey-level renormalisation
- Subregion variances (split) and Euclidian distances (merge) are computed using the multi-dimensional set of scores

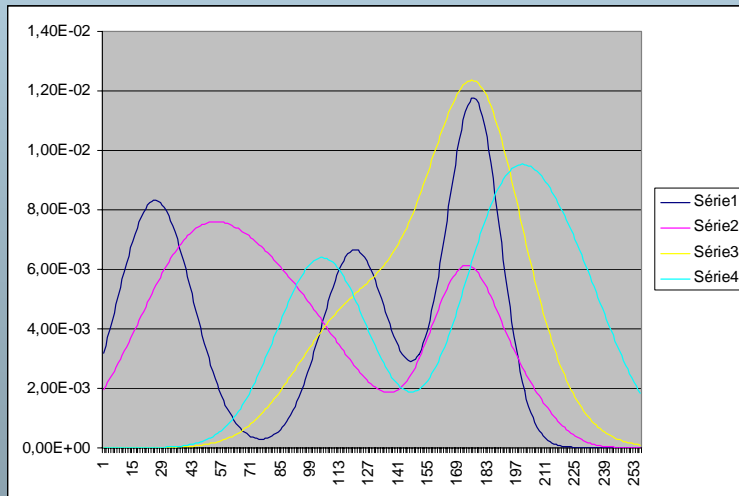
⇒ Local segmentation refinement

- For each resulting sub-region, the entire process is launched again (PCA computation + split-and-merge process)
- Refinement is stopped when no more subregions are created
- This allows a hierachical image segmentation, where the user can tune the level of segmentation detail required.

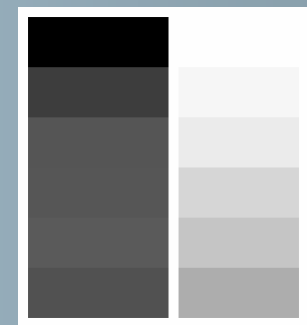
Results: synthetic test image

⇒ The test image

- Four different spectra have been generated using a random process
- A synthetic image has been built, using a particular concentration pattern



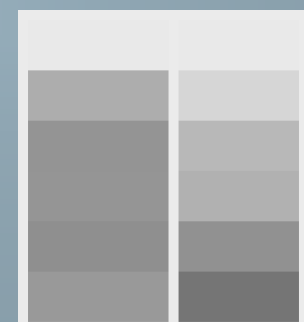
C1



C2

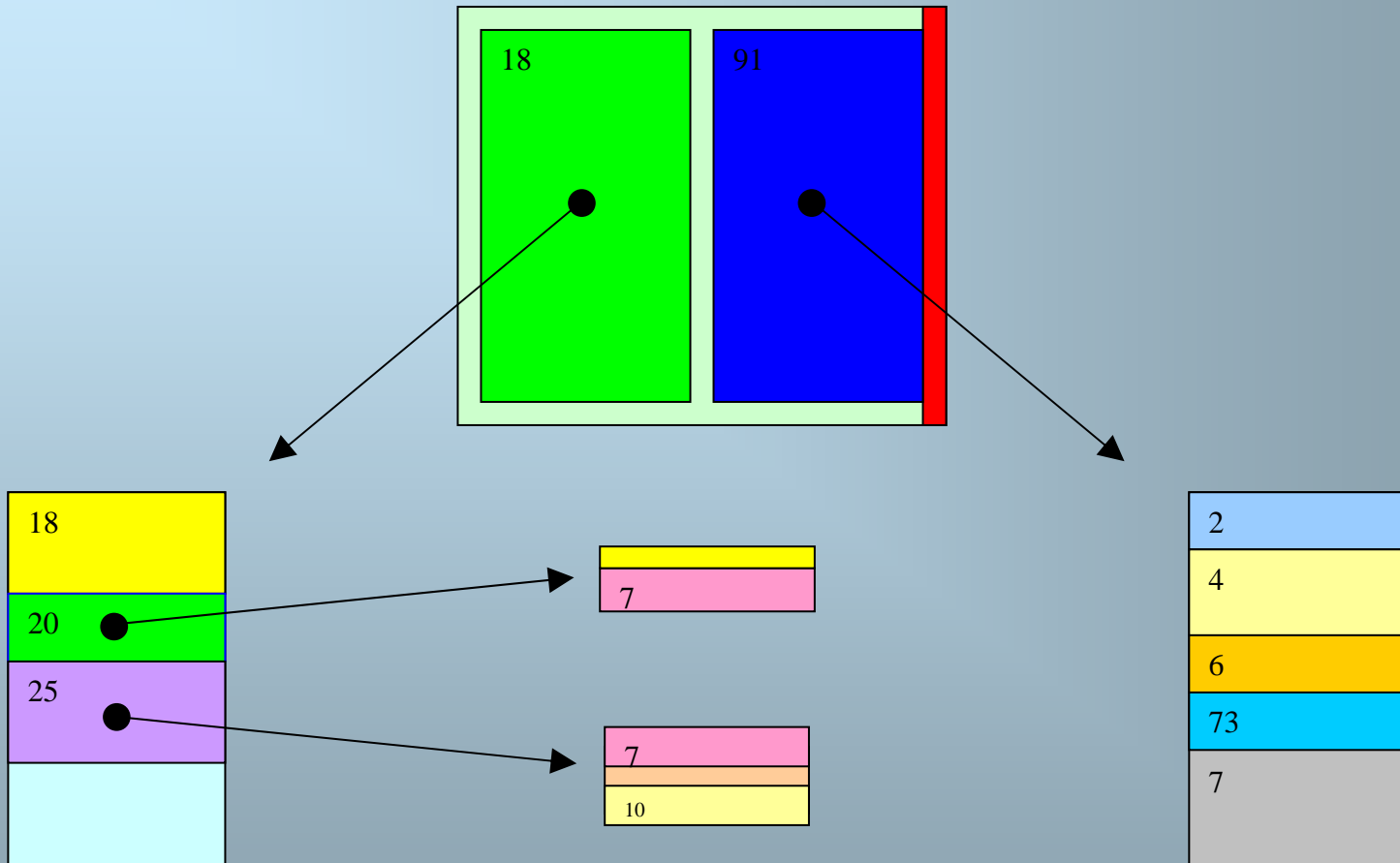


C3



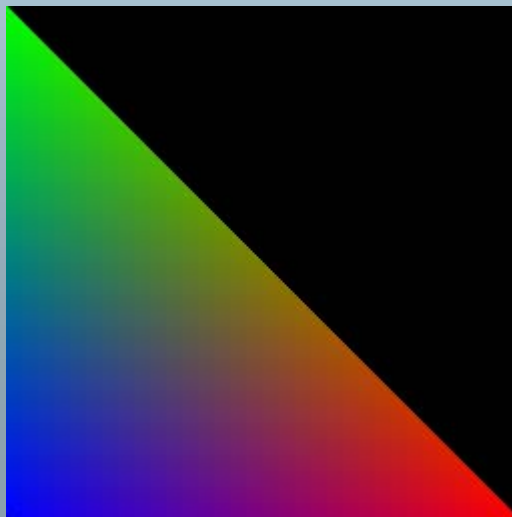
C4

Results: image segmentation

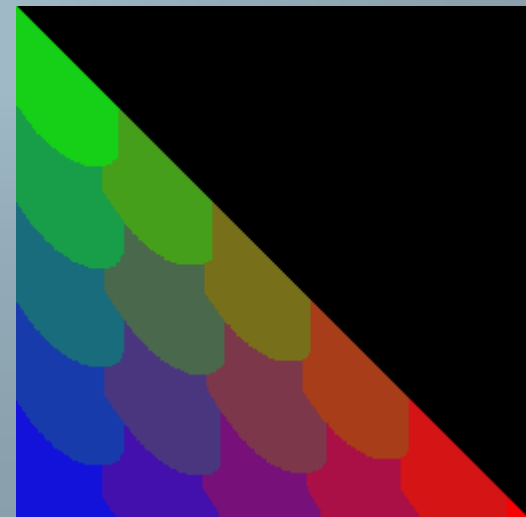
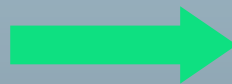


Alternative: Union-Find algorithm (Fiorio, 1996)

- ⇒ Starting with a region associated to each pixel
- ⇒ Image scanning, and comparison of each pixel (region) with the pixel (region) above and on the right.
- ⇒ Merging condition: euclidian distance $< S$

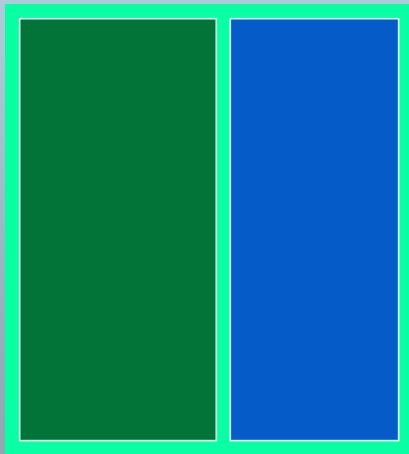


$S = 40$

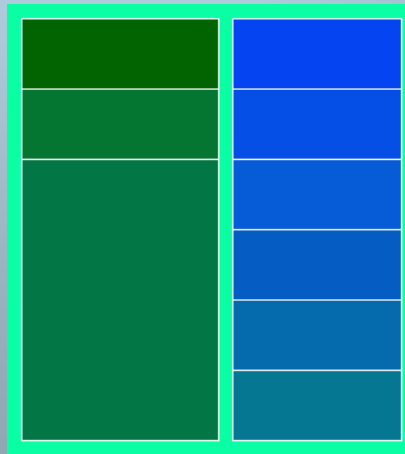


Union-Find result

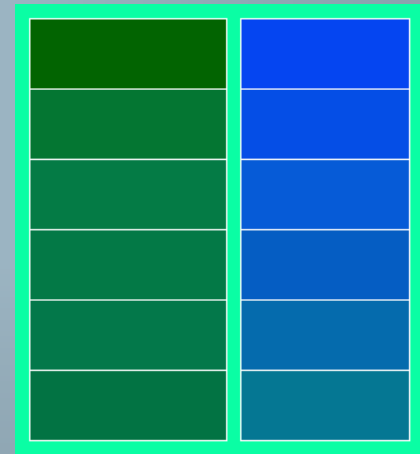
Application to the PC score vector
(after normalisation 0-255)



$S = 100$



$S = 10$



$S = 2$

Anisotropic diffusion

Basic principle (Perona & Malik, 1990)

- ⇒ Objective: to smooth an image (noise reduction) while preserving edges

- ⇒ Method:
 - Implementation of the Gaussian smoothing as an iterative process, using PDE formalism (PDE: Partial Derivative Equation)
 - Local attenuation of the iterative smoothing in high gradient areas

Isotropic diffusion

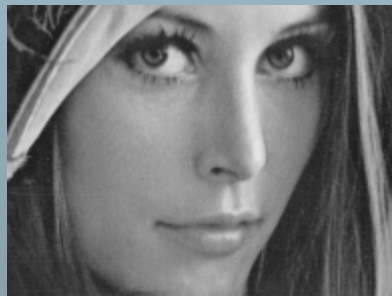
⇒ Let us consider the temporal equation for the image $I(x,y)$:

$$\frac{\partial I(x,y)}{\partial t} = \text{div}(\nabla I) = \Delta I \quad \ll \text{Heat equation} \gg$$

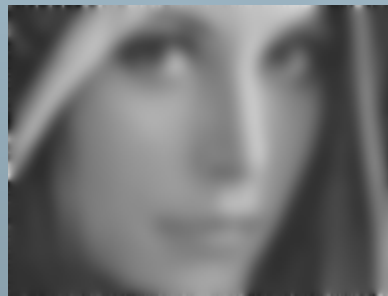
⇒ The equation solution is a temporal gaussian filtering:

$$I(x,y,t) = I(x,y,t_0) * G(x,y,t) \quad \text{where:} \quad G(x,y,t) = \frac{1}{4\pi t} \exp\left(-\frac{(x^2+y^2)}{4t}\right)$$

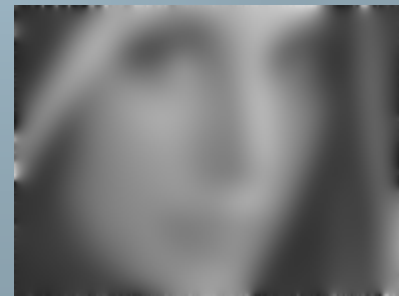
Gaussian with a variance $\sigma^2 = 2 t$



Original image



30 iterations



100 iterations

Anisotropic diffusion

⇒ Introduction of a function $g(\nabla I)$ so that:

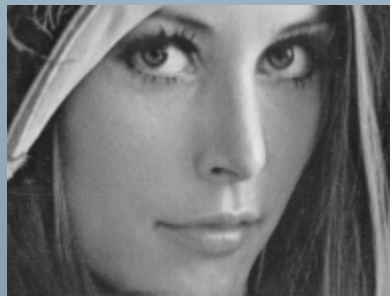
➤ $g(\nabla I) \approx 0$ if ∇I high

➤ $g(\nabla I) \approx 1$ if ∇I low

e.g:

$$g(\nabla I) = \exp(-\nabla I^2)$$

$$\frac{\partial I(x, y)}{\partial t} = \text{div}(g(\nabla I) \cdot \nabla I)$$



Original image



30 iterations



100 iterations

Extension to vectoriel data

⇒ The notion of gradient ∇I has to be redefined

Di Zenzo analysis (Di Zenzo, 1986)

$$\|dI\|^2 = \begin{bmatrix} dx \\ dy \end{bmatrix}^T \begin{bmatrix} g_{11} & g_{12} \\ g_{21} & g_{22} \end{bmatrix} \begin{bmatrix} dx \\ dy \end{bmatrix} \quad \text{where} \quad g_{ij} = \left\langle \frac{\partial I}{\partial x_i}, \frac{\partial I}{\partial x_j} \right\rangle$$

Maximum and minimum vectorial variations (eigen values of $[g_{ij}]$) :

$$\begin{cases} \lambda_+ = g_{11} + g_{22} + \sqrt{(g_{11} - g_{22})^2 + 4g_{12}^2} \\ \lambda_- = g_{11} + g_{22} - \sqrt{(g_{11} - g_{22})^2 + 4g_{12}^2} \end{cases}$$

⇒ Then:

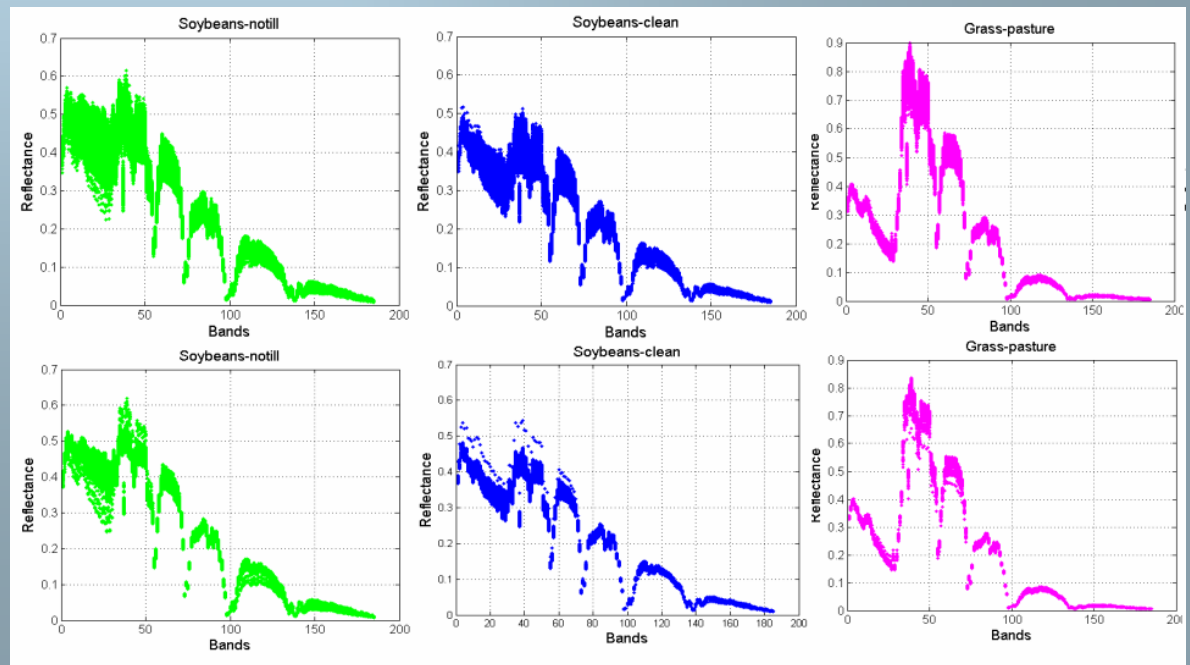
- $|\nabla I| = \sqrt{\lambda_+}$ (Di Zenzo)
- $|\nabla I| = \sqrt{(\lambda_+ - \lambda_-)}$ (Sapiro)

Hyperspectral example (Velez-Reyes, 2006)

Anisotropic
diffusion



Reduction of the
spectra deviation



Conclusion

⇒ Taking into account the spatial relationships between hyperspectral imaging pixels can help in :

- Reduction of classification errors
- Unsupervised segmentation
- Spectral noise reduction

⇒ An apparent paradox

- « Classical » hyperspectral processing requires a very accurate instrumental calibration
- Image processing tools have been developed for years to overcome image signal inaccuracy (8 bits signal level, lighting variations, etc.)

Can spatial information reduce hyperspectral calibration requirements ?

- Unsupervised clusterisation merging smooth spectral variations
- Chemometric modelisation adapted to these «deviating » clusters (e.g. EPO (J-M Roger, 2003)

References

- ⇒ Duarte-Carvajalino, Julio M.; Vélez-Reyes, Miguel; Castillo, Paul. *Scale-space in hyperspectral image analysis*. Algorithms and Technologies for Multispectral, Hyperspectral, and Ultraspectral Imagery XII. Edited by Shen, Sylvia S.; Lewis, Paul E.. Proceedings of the SPIE, Volume 6233, pp. 623315 (2006).
- ⇒ Fiorio, C. and J. Gustedt, *Two linear time Union-Find strategies for image processing*. Theoretical Computer Science, 1996. **154**(1996): p. 165-181.
- ⇒ Perona, P. and J. Malik, *Scale-space and edge detection using anisotropic diffusion*. IEEE Transactions on Pattern Analysis and Machine Intelligence, 1990. **12**(7): p. 629-639.
- ⇒ Prony, O., Descombes, X., Zerubia, J. Classification d'images satellitaires hyperspectrales en zone rurale et périurbaine. Rapport de recherche INRIA N°4008, septembre 2000.
- ⇒ Roger, J-M, Chauchard, F., Bellon-Maurel V. *EPO-PLS external parameter orthogonalisation of pls : Application to temperature- independent measurement of sugar content of intact fruits*. Chemometrics and Intelligent Laboratory Systems, 66-2 :191-204, 2003.