

Grasslands species diversity mapping from hyperspectral remote sensing

 5^e Colloque Groupe Hyperspectral SFPT-GH

M. Lopes ¹, M. Fauvel ¹, A. Ouin ¹ and S. Girard ²

 1 UMR 1201 DYNAFOR INRA & Institut National Polytechnique de Toulouse 2 Equipe MISTIS-LJK, Universite Grenoble Alpes, INRIA, France

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Grasslands species diversity

- Grasslands represent a significant source of biodiversity in farmed landscapes,
- They provide many ecosystem services (carbon regulation, erosion regulation, pollination...),
- Grasslands surface area and their diversity are declining [OMa12],
- Maps over grassland diversity are required over large area extents.



Spectral Variation Hypothesis

- It assumes that the spectral heterogeneity is correlated with spatial variations and heterogeneity of the habitat [Pal+02]
- Spectral heterogeneity can be used as a proxy for species diversity [Roc+16]
- Several indices have been proposed
 - Standard deviation or coefficient of variations of NDVI
 - PCA
 - Distance to centroids
 - Clustering

Objectives

- Project MUESLI
- Use hyperspectral images to monitor species richness at the parcel level
- Methodological contributions
 - Use of robust high dimensional clustering method
 - Extend conventional heterogeneity/diversity index

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$$H(p) = \frac{1}{n_p} \sum_{i \in p}^{n_p} \|\mathbf{x}_i - \boldsymbol{\mu}_p\|^2$$

where

$$\boldsymbol{\mu}_p = \frac{1}{n_p} \sum_{i \in p}^{n_p} \mathbf{x}_i.$$

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Equivalently, it can be computed as the trace of the empirical covariance matrix of the plot:

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■ Variant: first reduce the dimensionality (PCA, ...)

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Why MDC may not work

The following configurations have the same MDC





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α -diversity

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- Estimated by the Shannon entropy of a given plot

$$E_p = -\sum_{s=1}^{S} p_s \log(p_s)$$

where p is the considered plot, S the total number of species/classes/clusters and p_s is the relative proportion.

α -diversity

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Clusters estimated through the *PCA+Kmeans* pipeline applied on the whole image.

Why Kmeans may not work

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Statistical model

- Mixture model $p(\mathbf{x}) = \sum_{c=1}^{C} \pi_c p(\mathbf{x}|c)$,
- \blacksquare Under Gaussian assumption $p(\mathbf{x}|c)$ is a d-dimensional Gaussian distribution

$$p(\mathbf{x}|c) = \frac{1}{(2\pi)^{d/2} |\mathbf{\Sigma}_c|^{1/2}} \exp\left(-\frac{1}{2} (\mathbf{x} - \boldsymbol{\mu}_c)^\top \mathbf{\Sigma}_c^{-1} (\mathbf{x} - \boldsymbol{\mu}_c)\right)$$

• Curse of dimensionality: special structure for the covariance matrix $\mathbf{\Sigma}_c = \mathbf{Q}_c \mathbf{\Lambda}_c \mathbf{Q}_c^ op$



High dimensional GMM [BGS07]



Under the HDDA model

$$\begin{split} \boldsymbol{\Sigma}_i &= \tilde{\mathbf{Q}}_i \tilde{\mathbf{\Lambda}}_i \tilde{\mathbf{Q}}_i^\top + \lambda_i \mathbf{I}_d \\ \boldsymbol{\Sigma}_i^{-1} &= \tilde{\mathbf{Q}}_i \tilde{\mathbf{V}}_i \tilde{\mathbf{Q}}_i^\top + \lambda_i^{-1} \mathbf{I}_d \end{split}$$

with $\tilde{\mathbf{Q}}_i = \begin{bmatrix} \mathbf{q}_{i1}, \dots, \mathbf{q}_{ip_i} \end{bmatrix}$, $\tilde{\mathbf{\Lambda}}_i = \operatorname{diag} \begin{bmatrix} \lambda_{i1} - \lambda_i, \dots, \lambda_{ip_i} - \lambda_i \end{bmatrix}$, $\tilde{\mathbf{V}}_i = \operatorname{diag} \begin{bmatrix} \frac{1}{\lambda_{i1}} - \frac{1}{\lambda_i}, \dots, \frac{1}{\lambda_{ip_i}} - \frac{1}{\lambda_i} \end{bmatrix}$ and \mathbf{I}_d is the identity matrix of size d.

Spectral heterogeneity revisited 1/2

 \blacksquare Samples covariance matrix for a given plot p

$$\mathbf{\Sigma}_p = \mathbf{B}_p + \mathbf{W}_p$$

where

• \mathbf{B}_p is the between class covariance matrix of plot p

$$\mathbf{B}_p = \sum_{c=1}^{C_p} \pi_{pc} (oldsymbol{\mu}_{pc} - oldsymbol{\mu}_p) (oldsymbol{\mu}_{pc} - oldsymbol{\mu}_p)^ op$$

W_p is the within class covariance matrix of plot p

$$\mathbf{W}_p = \sum_{c=1}^{C_p} \pi_{pc} \mathbf{\Sigma}_{pc}$$

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Spectral heterogeneity revisited 2/2

Trace(
$$\Sigma_p$$
) = Trace(\mathbf{B}_p) + Trace(\mathbf{W}_p)
Trace(\mathbf{B}_p) = $\sum_{c=1}^{C_p} \pi_{pc} \|\boldsymbol{\mu}_{pc} - \boldsymbol{\mu}_p\|^2$
Trace(\mathbf{W}_p) = $\frac{1}{n_p} \sum_{i=1}^{C_p} \sum_{k \in c} \|\mathbf{x}_{pk} - \boldsymbol{\mu}_{pc}\|^2$

	$Trace(\boldsymbol{\Sigma}_p)$	$Trace(\mathbf{B}_p)$	$Trace(\mathbf{W}_p)$
Plot 1	13.63	0	13.63
Plot 2	13.74	12.71	0.973

Improved spectral entropy

For each pixel of the plot, the vector of posterior probabilities is available

$$\left[p(C=1|\mathbf{x}),\ldots,p(C=C_p|\mathbf{x})\right]$$

The relative proportion is then computed as:

$$p_c = \frac{1}{n_p} \sum_{k \in c} p(C = c | \mathbf{x}) = \pi_c$$

It allows to let a pixel belonging to several clusters (not a crisp affectation)

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Data collection





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	Experimental protocol		

Simulations

• Select the number of classes using ICL: stop when $dICL{<}1\%$



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Measure of heterogeneity

ID	С	E	В	W	V	Н	D
6	2	0.68	13.16	11.32	11.17	0.97	0.13
8	1	0.0	inf	11.12	11.12	0.09	3.81
137	4	1.31	10.36	10.97	9.93	0.08	3.97
143	2	0.68	15.02	11.57	11.54	0.04	5.06

B, W and V are in *log* scale

 $\bullet \ E \approx \log(C)$

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Species diversity in semi-natural grasslands

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Conclusions and perspectives

- Species diversity in semi-natural grasslands
- Extension of heterogeneity measures with high dimensional clustering techniques

Conclusions and perspectives

- Species diversity in semi-natural grasslands
- Extension of heterogeneity measures with high dimensional clustering techniques
- Estimated diversity does not correlate (yet!) with field work

Bibliography I

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