>>> Spectral Unmixing >>> GRSS Summer School

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Date: [2017-04-26 Wed 13:30]-[2017-04-26 Wed 16:30]

# 1. Motivations

# 2. Spectral mixture models

Linear models

Non linear model

# 3. Subspace and Endmember extraction

Subspace extraction
Endmember extraction
Endmember extraction in action

#### 4. Linear unmixing

Abundance estimation
Advanced abundance estimation

5. References

[~]\$ \_

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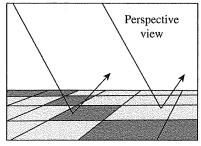
## 4. Linear unmixing

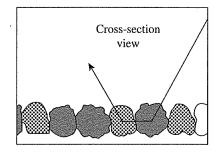
Abundance estimation
Advanced abundance estimation

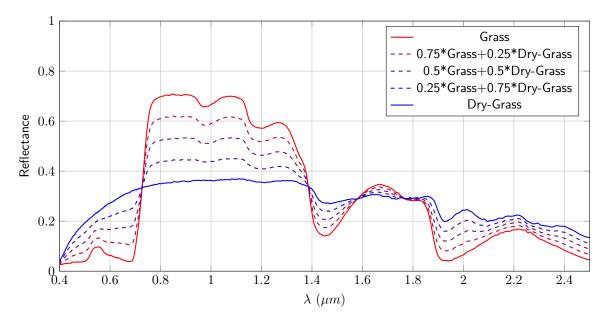
## >>> Spectral mixing

- \* When a pixel contains several materials: all these materials contribute to the collected reflectance [MLC16]
- \* It is called a *mixel* or *mixed spectra* and pure contributions are termed *endmembers*









# >>> Applications

- ★ Conventional mapping applications [Bio+12]:
  - ★ Mineral proportions
  - ⋆ Vegetation covers
- \* Multisource fusion:
  - ★ Sources of various spatial resolution
  - \* Sources of various spectral resolution
- ⋆ Misc:
  - \* Post-improvement of classification maps
  - \* Clouds reconstruction of multi-temporal HI

[1. Motivations]\$ \_ [6/43]

>>> Unmixing processing chain

- 1. Athmospheric correction
- 2. Dimensionality reduction
- 3. Find endmembers
  - \* Hyperspectral library
  - \* Unsupervised
- 4. Find abundances

# >>> Outline

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\* Each pixel is a convex linear combination of endmenbers (pure spectra):

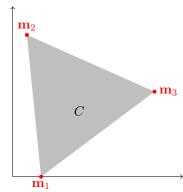
$$\mathbf{x} = \sum_{j=1}^{p} \alpha_j \mathbf{m}_j + \mathbf{e}$$

- $\star$   $\mathbf{m}_j$  is the  $j^{\text{th}}$  endmember,
- $\star \ \alpha_j$  is the abundance of endmember j,
- \* e is the modelling error.
- \* The abundances are subject to the following constraints:
  - ★ Non negativity:  $\alpha_j \ge 0, \forall j = 1, \ldots, p$
  - \* Sum to one:  $\sum_{j=1}^{p} \alpha_j = 1$ .
- \* Matricial formulation:

$$x = M\alpha + e$$

where 
$$\mathbf{M} = \left[\mathbf{m}_1, \dots, \mathbf{m}_p\right]$$
.

- $\star$  If endmembers are affinely independent, hyperspectral pixels live in the convex hull C generated by the endmembers
- \*  $C := \left\{ \mathbf{x} = \mathbf{M} \boldsymbol{\alpha} | \sum_{j=1}^{p} \alpha_j = 1, \ \alpha_j \ge 0, \forall j = 1, \dots, p \right\}$



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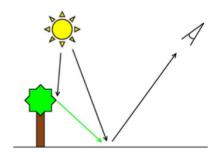
Endmember extraction in action

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★ Consider multiple reflections [HPG14]



★ The model

$$\mathbf{x} = \sum_{j=1}^{p} \alpha_j \mathbf{m}_j + \sum_{j,k=1}^{p} \beta_{j,k} \mathbf{m}_j \odot \mathbf{m}_j$$

 $\star$  Several possible constraints on  $\alpha$  and  $\beta$ 

>>> Other models

- ⋆ Physics based model: Intimate mixture (e.g., Hapke model)
- ⋆ Computer graphics: ray tracing
- ★ Kernel methods:
  - ★ "kernelized" linear models
  - ★ Construct "smart" kernels

[2. Spectral mixture models]\$ \_ [14/43]

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Endmember extraction
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>>> Conventional methods

Find the subspace signal in order to search the endmembers:

- \* Principal component analysis: find the subspace that minimizes reconstruction error
- \* Minimum noise fraction: find the subspace that maximizes the SNR
- ★ HySime (hyperspectral subspace identification)
  - \* Estimation of the signal and noise subspaces
  - \* Find the subpace that best represents the signal subspace

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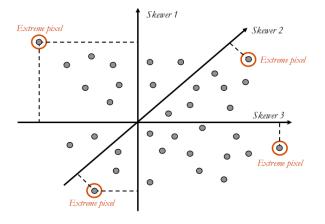
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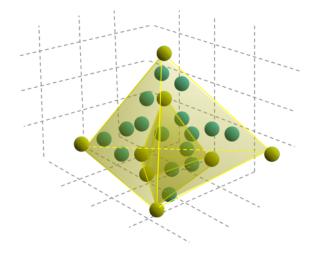
# >>> Pixel Purity Index [BKG95]

- 1. Apply MNF
- 2. Project pixels onto random vector and find extreme projected pixels and store them
- 3. Pixels with the highest score are identified as the *purest one*



# >>> N-FINDR [Win99]

- $\star$  Assumes pure pixels are present
- ★ Find the pixels that maximizes the volume of the simplex



>>> Vertex component analysis [ND05]

- \* Iteratively project pixels on *orthogonal direction* to the subspace spanned by previously selected endmembers
- ★ New endmember correpond to the extreme of the projection
- \* Works similarly to *orthogonal subspace projection* but accounting for the noise

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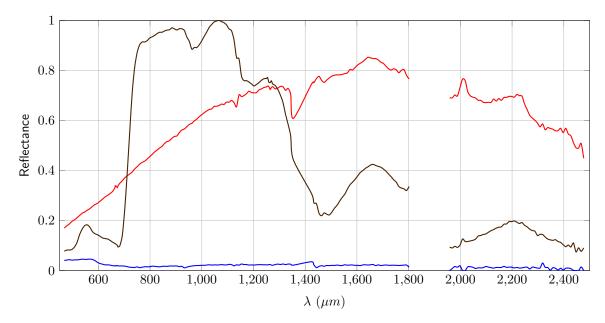
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Subspace extraction
Endmember extraction in action

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#### 5. References

[4. Linear unmixing]\$ \_ [25/43]

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 $\star$  One the endmembers are estimated, abundances can be estimated by minimizing a reconstruction error Re:

$$\hat{\boldsymbol{\alpha}} = \min_{\boldsymbol{\alpha}} \left\{ Re(\mathbf{x}, \mathbf{M}\boldsymbol{\alpha}) \right\}$$

 $\star$  Conventionally Re is chosen as the mean square square error:

$$\hat{\boldsymbol{\alpha}} = \min_{\boldsymbol{\alpha}} \|\mathbf{x} - \mathbf{M}\boldsymbol{\alpha}\|^2$$

★ Without any constraint the solution is given by

$$\hat{\boldsymbol{\alpha}} = (\mathbf{M}^{\top}\mathbf{M})^{-1}\mathbf{M}^{\top}\mathbf{x}$$

★ Positive constraints:

$$\hat{m{lpha}} = \min_{m{lpha}} \| \mathbf{x} - \mathbf{M} m{lpha} \|^2$$
 Subject to  $lpha_j \geq 0, orall j = 1, \dots, p$ 

★ Full set of constraints

$$\hat{\pmb{\alpha}} = \min_{\pmb{\alpha}} \| \mathbf{x} - \mathbf{M} \pmb{\alpha} \|^2$$
 Subject to  $\sum_{j=1}^p \alpha_j = 1, \alpha_j \geq 0, \forall j=1,\dots,p$ 

- \* Not explicit solution! Needs to optimize a quadratic function under convex constraints.
- \* This is known as non negative least squares unmixing or fully constraints least square unmixing
- \* Easy trick: use only positive constraints that add manually the sum to one constraint!

[4. Linear unmixing]\$ \_ [28/43]

#### >>> NNLS

```
from scipy import optimize
import scipy as sp

# Load endmembers
M = sp.loadtxt(".../Unmixing/figures/endmembers.csv",delimiter=',')[:,1:]
x = 0.2*M[:,0] + 0.7*M[:,1] + 0.1*M[:,2]

# NNLS
res = optimize.nnls(M,x)
print res[0]

[ 0.2  0.7  0.1]
```

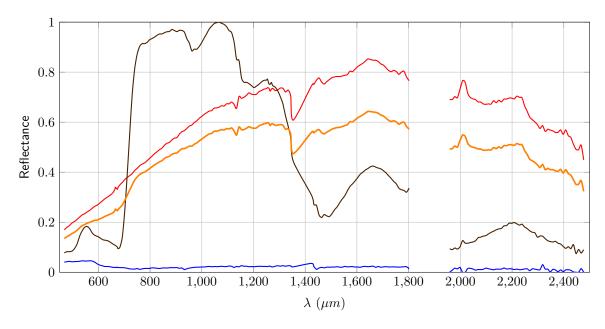
[4. Linear unmixing]\$ \_ [29/43]

```
>>> FCLS 1/2
from scipy import optimize
import scipy as sp
# Loss
def loss(alpha,x,M):
    e = x-sp.dot(M,alpha)
    return (e**2).sum()
def jac(alpha,x,M):
    e = x-sp.dot(M,alpha)
    return -2*sp.dot(M.T,e)
cons = {'type':'eq','fun':lambda alpha: 1-alpha.sum(),'jac':lambda alpha: -alpha}
bnds = ((0, None), (0, None), (0, None,))
# Load endmembers
M = sp.loadtxt("../Unmixing/figures/endmembers.csv",delimiter=',')[:,1:]
x = 0.2*M[:.0] + 0.7*M[:.1] + 0.1*M[:.2]
# Optimize
alpha0 = sp.ones((3,))/3.0
res = optimize.minimize(loss, alpha0, args=(x,M,), jac=jac,constraints=cons, method='SLSQP',
                        bounds=bnds.)
print res['x']
[ 0.16441937  0.67116125  0.16441937]
```

[4. Linear unmixing]\$ \_ [30/43]

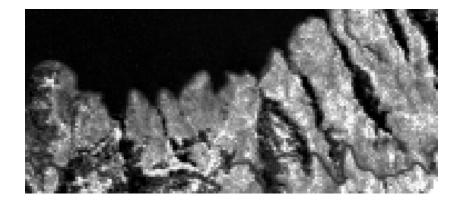


[4. Linear unmixing]\$ \_



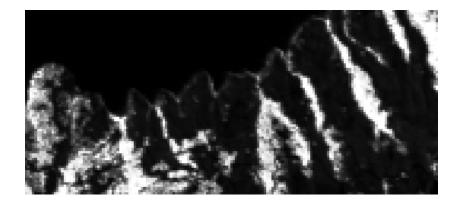
```
import scipy as sp
import rasterTools as rt
import pysptools.eea as eea
# Number of endmembers
NE = 3
# Load images
im,GeoT,Proj = rt.open_data('../Data/Moffett_full.tif')
 [h,w,b]=im.shape
wave = sp.loadtxt('../Data/wave_moffett.csv',delimiter=',')
# Compute endmenbers
nfindr = eea.NFINDR()
M = nfindr.extract(im.astype(float), NE, normalize=False).T
abundances = sp.empty((h,w,NE))
for h in xrange(h):
    for w_ in xrange(w):
        x = im[h, w,:]
        res = optimize.nnls(M,x)
        a = res[0]
         abundances[h ,w ,:] = (a/a.sum())
# Write the image
rt.write_data("../Data/Moffett_abundances.tif",abundances,GeoT,Proj)
[4. Linear unmixing]$
```

>>> Abundance maps 1/4
from scipy import optimize





[4. Linear unmixing]\$ \_ [34/43]



[4. Linear unmixing]\$ \_ [35/43]

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- \* When the number of endmembers are large (e.g. selected from spectral library), the observed spectral vector is usually a combination of few ones.
- \* Add *sparsity* constraints in the optimization problem such as

$$\min_{\boldsymbol{\alpha}} \|\boldsymbol{\alpha}\|_0$$

Subject to 
$$\|\mathbf{x} - \mathbf{M}\boldsymbol{\alpha}\|^2 \le \delta, \alpha_j \ge 0, \forall j = 1, \dots, p$$

- \* However, this problem is NP-hard: no straightforward solution
- ★ Use convex formulation from sparse regression (LASSO):

$$\min_{\alpha} \|\mathbf{x} - \mathbf{M}\alpha\|^2 + \lambda \|\alpha\|_1$$

Subject to  $\alpha_i \geq 0, \forall j = 1, \dots, p$ 

[4. Linear unmixing]\$ \_\_ [37/43]

- ★ Add spatial constraints during the optimization
- $\star$  Total variation [IBP12]: measure the spatial variation of abundances for neighboring pixels

$$TV(\alpha_i) = \sum_{j \in \mathcal{N}_i} \|\alpha_i - \alpha_j\|_1$$

★ Global optimization rather than pixel wise optimization:

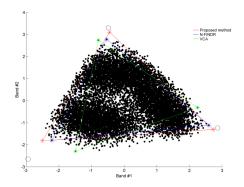
$$\sum_{i=1}^{n} \|\mathbf{x}_i - \mathbf{M}\boldsymbol{\alpha}_i\|^2 + \lambda \sum_{i=1}^{n} \|\boldsymbol{\alpha}_i\|_1 + \lambda_{TV} \sum_{i=1}^{n} TV(\boldsymbol{\alpha}_i)$$
Subject to  $\alpha_{i,j} \ge 0, \forall j = 1, \dots, p$  and  $i = 1, \dots, n$ 

[4. Linear unmixing]\$ \_ [38/43]

\* Likelihood formulation

$$f(\mathbf{x}_i|\mathbf{M}, \boldsymbol{\alpha}_i, \sigma^2) = \left(\frac{1}{2\pi\sigma^2}\right)^{d/2} \exp\left(-\frac{\|\mathbf{x}_i - \mathbf{M}\boldsymbol{\alpha}_i\|^2}{2\sigma^2}\right)$$

- ★ Conjoint estimation of endmembers and abundances [Dob+09]
- ★ Rely in MCMC algorithm
  - ⋆ Long processing time
  - \* Provide usually better results than geometric methods in critical situation (no pure pixels...)
- ⋆ Possible to add spatial dependencies [EDT11]



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[6. References]\$ \_ [42/43]

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