

BEADS : filtrage asymétrique de ligne de base (tendance) et débruitage pour des signaux positifs avec parcimonie des dérivées

Séminaire ICube

L. DUVAL, A. PIRAYRE

IFP Energies nouvelles

1 et 4 av. de Bois-Préau, 92852 Rueil-Malmaison - France

X. NING, I. W. SELESNICK

Polytechnic School of Engineering

New York University

19 juin 2015

The fast way

- ▶ Question: where is the string behind the bead?
- ▶ Smoothness, sparsity, asymmetry



Outline

INTRODUCTION

OUTLINE

BACKGROUND

MODELING

NOTATIONS

COMPOUND SPARSE DERIVATIVE MODELING

BEADS ALGORITHM

MAJORIZE-MINIMIZE

EVALUATION AND RESULTS

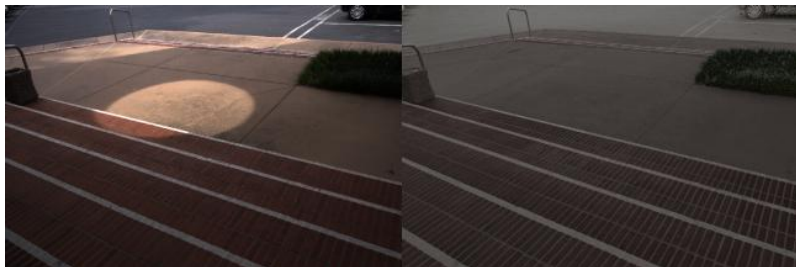
SIMULATED BASELINE AND NOISE

POISSON NOISE

GC×GC

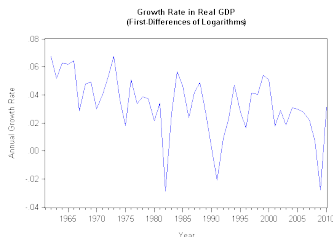
CONCLUSIONS

Background on background



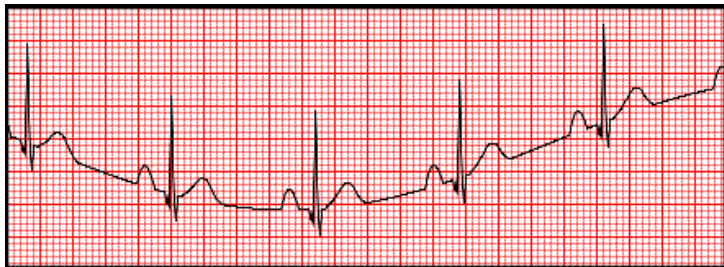
- ▶ Background affects quantitative evaluation/comparison
- ▶ In some domains: (instrumental) bias, (seasonal) trend
- ▶ In analytical chemistry: drift, continuum, wander, *baseline*
- ▶ Rare cases of parametric modeling

Background on background



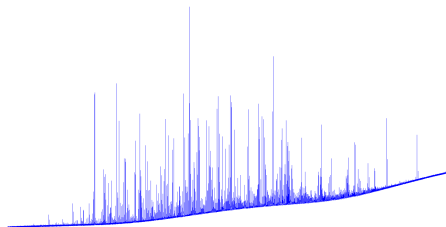
- ▶ Background affects quantitative evaluation/comparison
- ▶ In some domains: (instrumental) bias, (seasonal) trend
- ▶ In analytical chemistry: drift, continuum, wander, *baseline*
- ▶ Rare cases of parametric modeling

Background on background



- ▶ Background affects quantitative evaluation/comparison
- ▶ In some domains: (instrumental) bias, (seasonal) trend
- ▶ In analytical chemistry: drift, continuum, wander, *baseline*
- ▶ Rare cases of parametric modeling

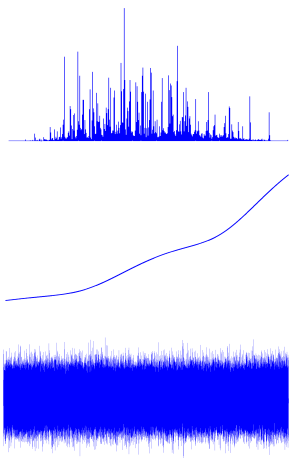
Background on background



- ▶ Background affects quantitative evaluation/comparison
- ▶ In some domains: (instrumental) bias, (seasonal) trend
- ▶ In analytical chemistry: drift, continuum, wander, *baseline*
- ▶ Rare cases of parametric modeling

Background on background

For analytical chemistry data:



Notations

Morphological decomposition:

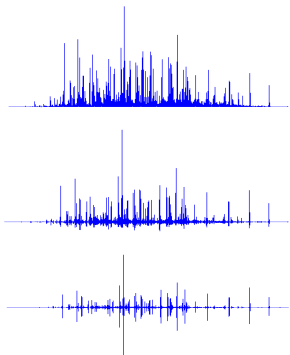
$$\mathbf{y} = \mathbf{x} + \mathbf{f} + \mathbf{w}, \quad (\mathbf{y}, \mathbf{x}, \mathbf{f}, \mathbf{w}) \in (\mathbb{R}^N)^4.$$

- ▶ \mathbf{y} : observation
- ▶ \mathbf{x} : clean series of peaks
- ▶ \mathbf{f} : baseline
- ▶ \mathbf{w} : noise

Assumption: in the absence of peaks, the baseline can be approximately recovered from a noise-corrupted observation by low-pass filtering

- ▶ $\hat{\mathbf{f}} = \mathbf{L}(\mathbf{y} - \hat{\mathbf{x}})$ (\mathbf{L} : low-pass filter)
- ▶ formulated as $\|\mathbf{y} - \hat{\mathbf{s}}\|_2^2 = \|\mathbf{H}(\mathbf{y} - \hat{\mathbf{x}})\|_2^2$
- ▶ $\mathbf{H} = \mathbf{I} - \mathbf{L}$: high-pass filter

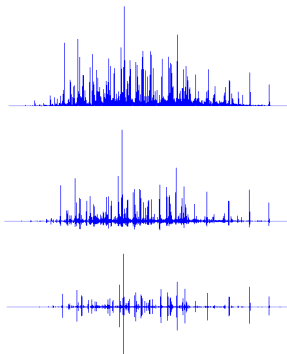
Compound sparse derivative modeling



An estimate $\hat{\mathbf{x}}$ can be obtained (with \mathbf{D}_i diff. operators) via:

$$\hat{\mathbf{x}} = \arg \min_{\mathbf{x}} \left\{ F(\mathbf{x}) = \frac{1}{2} \|\mathbf{H}(\mathbf{y} - \mathbf{x})\|_2^2 + \sum_{i=0}^M \lambda_i R_i(\mathbf{D}_i \mathbf{x}) \right\}.$$

Compound sparse derivative modeling



Examples of (smooth) sparsity promoting functions for R_i

- ▶ $\phi_i^A = |x|$
- ▶ $\phi_i^B = \sqrt{|x|^2 + \epsilon}$
- ▶ $\phi_i^C = |x| - \epsilon \log(|x| + \epsilon)$

Compound sparse derivative modeling

Take the positivity of chromatogram peaks into account:

$$\hat{\mathbf{x}} = \arg \min_{\mathbf{x}} \left\{ F(\mathbf{x}) = \frac{1}{2} \|\mathbf{H}(\mathbf{y} - \mathbf{x})\|_2^2 + \lambda_0 \sum_{n=0}^{N-1} \theta_\epsilon(x_n; r) + \sum_{i=1}^M \lambda_i \sum_{n=0}^{N_i-1} \phi([\mathbf{D}_i \mathbf{x}]_n) \right\}.$$

Start from:

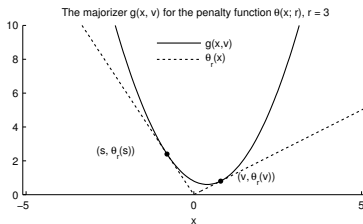
$$\theta(x; r) = \begin{cases} x, & x \geq 0 \\ -rx, & x < 0 \end{cases}$$

Compound sparse derivative modeling

Take the positivity of chromatogram peaks into account:

$$\hat{\mathbf{x}} = \arg \min_{\mathbf{x}} \left\{ F(\mathbf{x}) = \frac{1}{2} \|\mathbf{H}(\mathbf{y} - \mathbf{x})\|_2^2 + \lambda_0 \sum_{n=0}^{N-1} \theta_{\epsilon}(x_n; r) + \sum_{i=1}^M \lambda_i \sum_{n=0}^{N_i-1} \phi([\mathbf{D}_i \mathbf{x}]_n) \right\}.$$

and majorize it

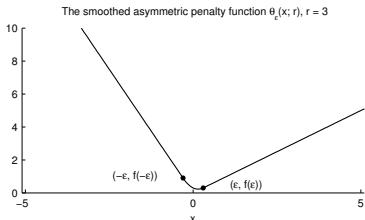


Compound sparse derivative modeling

Take the positivity of chromatogram peaks into account:

$$\hat{\mathbf{x}} = \arg \min_{\mathbf{x}} \left\{ F(\mathbf{x}) = \frac{1}{2} \|\mathbf{H}(\mathbf{y} - \mathbf{x})\|_2^2 + \lambda_0 \sum_{n=0}^{N-1} \theta_{\epsilon}(x_n; r) + \sum_{i=1}^M \lambda_i \sum_{n=0}^{N_i-1} \phi([\mathbf{D}_i \mathbf{x}]_n) \right\}.$$

then smooth it:



Compound sparse derivative modeling

Take the positivity of chromatogram peaks into account:

$$\hat{\mathbf{x}} = \arg \min_{\mathbf{x}} \left\{ F(\mathbf{x}) = \frac{1}{2} \|\mathbf{H}(\mathbf{y} - \mathbf{x})\|_2^2 + \lambda_0 \sum_{n=0}^{N-1} \theta_{\epsilon}(x_n; r) + \sum_{i=1}^M \lambda_i \sum_{n=0}^{N_i-1} \phi([\mathbf{D}_i \mathbf{x}]_n) \right\}.$$

then majorize it:

$$g_0(x, v) = \begin{cases} \frac{1+r}{4|v|} x^2 + \frac{1-r}{2} x + |v| \frac{1+r}{4}, & |v| > \epsilon \\ \frac{1+r}{4\epsilon} x^2 + \frac{1-r}{2} x + \epsilon \frac{1+r}{4}, & |v| \leq \epsilon. \end{cases}$$

BEADS Algorithm

We now have a majorizer for F

$$G(\mathbf{x}, \mathbf{v}) = \frac{1}{2} \|\mathbf{H}(\mathbf{y} - \mathbf{x})\|_2^2 + \lambda_0 \mathbf{x}^\top [\boldsymbol{\Gamma}(\mathbf{v})] \mathbf{x} \\ + \lambda_0 \mathbf{b}^\top \mathbf{x} + \sum_{i=1}^M \left[\frac{\lambda_i}{2} (\mathbf{D}_i \mathbf{x})^\top [\Lambda(\mathbf{D}_i \mathbf{v})] (\mathbf{D}_i \mathbf{x}) \right] + c(\mathbf{v}).$$

Minimizing $G(\mathbf{x}, \mathbf{v})$ with respect to \mathbf{x} yields

$$\mathbf{x} = \left[\mathbf{H}^\top \mathbf{H} + 2\lambda_0 \boldsymbol{\Gamma}(\mathbf{v}) + \sum_{i=1}^M \lambda_i \mathbf{D}_i^\top [\Lambda(\mathbf{D}_i \mathbf{v})] \mathbf{D}_i \right]^{-1} (\mathbf{H}^\top \mathbf{H} \mathbf{y} - \lambda_0 \mathbf{b}).$$

with notations

$$c(\mathbf{v}) = \sum_n \left[\phi(v_n) - \frac{v_n}{2} \phi'(v_n) \right].$$

BEADS Algorithm

We now have a majorizer for F

$$G(\mathbf{x}, \mathbf{v}) = \frac{1}{2} \|\mathbf{H}(\mathbf{y} - \mathbf{x})\|_2^2 + \lambda_0 \mathbf{x}^\top [\boldsymbol{\Gamma}(\mathbf{v})] \mathbf{x} \\ + \lambda_0 \mathbf{b}^\top \mathbf{x} + \sum_{i=1}^M \left[\frac{\lambda_i}{2} (\mathbf{D}_i \mathbf{x})^\top [\Lambda(\mathbf{D}_i \mathbf{v})] (\mathbf{D}_i \mathbf{x}) \right] + c(\mathbf{v}).$$

Minimizing $G(\mathbf{x}, \mathbf{v})$ with respect to \mathbf{x} yields

$$\mathbf{x} = \left[\mathbf{H}^\top \mathbf{H} + 2\lambda_0 \boldsymbol{\Gamma}(\mathbf{v}) + \sum_{i=1}^M \lambda_i \mathbf{D}_i^\top [\Lambda(\mathbf{D}_i \mathbf{v})] \mathbf{D}_i \right]^{-1} (\mathbf{H}^\top \mathbf{H} \mathbf{y} - \lambda_0 \mathbf{b}).$$

with notations

$$[\boldsymbol{\Gamma}(\mathbf{v})]_{n,n} = \begin{cases} \frac{1+r}{4|v_n|}, & |v_n| \geq \epsilon \\ \frac{1+r}{4\epsilon}, & |v_n| \leq \epsilon \end{cases}$$

BEADS Algorithm

We now have a majorizer for F

$$G(\mathbf{x}, \mathbf{v}) = \frac{1}{2} \|\mathbf{H}(\mathbf{y} - \mathbf{x})\|_2^2 + \lambda_0 \mathbf{x}^\top [\boldsymbol{\Gamma}(\mathbf{v})] \mathbf{x} \\ + \lambda_0 \mathbf{b}^\top \mathbf{x} + \sum_{i=1}^M \left[\frac{\lambda_i}{2} (\mathbf{D}_i \mathbf{x})^\top [\boldsymbol{\Lambda}(\mathbf{D}_i \mathbf{v})] (\mathbf{D}_i \mathbf{x}) \right] + c(\mathbf{v}).$$

Minimizing $G(\mathbf{x}, \mathbf{v})$ with respect to \mathbf{x} yields

$$\mathbf{x} = \left[\mathbf{H}^\top \mathbf{H} + 2\lambda_0 \boldsymbol{\Gamma}(\mathbf{v}) + \sum_{i=1}^M \lambda_i \mathbf{D}_i^\top [\boldsymbol{\Lambda}(\mathbf{D}_i \mathbf{v})] \mathbf{D}_i \right]^{-1} (\mathbf{H}^\top \mathbf{H} \mathbf{y} - \lambda_0 \mathbf{b}).$$

with notations

$$[\boldsymbol{\Lambda}(\mathbf{v})]_{n,n} = \frac{\phi'(v_n)}{v_n}$$

BEADS Algorithm

We now have a majorizer for F

$$G(\mathbf{x}, \mathbf{v}) = \frac{1}{2} \|\mathbf{H}(\mathbf{y} - \mathbf{x})\|_2^2 + \lambda_0 \mathbf{x}^\top [\boldsymbol{\Gamma}(\mathbf{v})] \mathbf{x} \\ + \lambda_0 \mathbf{b}^\top \mathbf{x} + \sum_{i=1}^M \left[\frac{\lambda_i}{2} (\mathbf{D}_i \mathbf{x})^\top [\boldsymbol{\Lambda}(\mathbf{D}_i \mathbf{v})] (\mathbf{D}_i \mathbf{x}) \right] + c(\mathbf{v}).$$

Minimizing $G(\mathbf{x}, \mathbf{v})$ with respect to \mathbf{x} yields

$$\mathbf{x} = \left[\mathbf{H}^\top \mathbf{H} + 2\lambda_0 \boldsymbol{\Gamma}(\mathbf{v}) + \sum_{i=1}^M \lambda_i \mathbf{D}_i^\top [\boldsymbol{\Lambda}(\mathbf{D}_i \mathbf{v})] \mathbf{D}_i \right]^{-1} \left(\mathbf{H}^\top \mathbf{H} \mathbf{y} - \lambda_0 \mathbf{b} \right).$$

with notations

$$[\mathbf{b}]_n = \frac{1-r}{2}$$

BEADS Algorithm

Writing filter $\mathbf{H} = \mathbf{A}^{-1}\mathbf{B} \approx \mathbf{B}\mathbf{A}^{-1}$ (banded matrices) we have

$$\mathbf{x} = \mathbf{A}\mathbf{Q}^{-1} \left(\mathbf{B}^T\mathbf{B}\mathbf{A}^{-1}\mathbf{y} - \lambda_0\mathbf{A}^T\mathbf{b} \right)$$

where \mathbf{Q} is the banded matrix,

$$\mathbf{Q} = \mathbf{B}^T\mathbf{B} + \mathbf{A}^T\mathbf{M}\mathbf{A},$$

and \mathbf{M} is the banded matrix,

$$\mathbf{M} = 2\lambda_0\mathbf{\Gamma}(\mathbf{v}) + \sum_{i=1}^M \lambda_i \mathbf{D}_i^T [\Lambda(\mathbf{D}_i\mathbf{v})] \mathbf{D}_i.$$

BEADS Algorithm

Using previous equations, the MM iteration takes the form:

$$\mathbf{M}^{(k)} = 2\lambda_0\mathbf{\Gamma}(\mathbf{x}^{(k)}) + \sum_{i=1}^M \lambda_i \mathbf{D}_i^T [\Lambda(\mathbf{D}_i \mathbf{x}^{(k)})] \mathbf{D}_i.$$

$$\mathbf{Q}^{(k)} = \mathbf{B}^T \mathbf{B} + \mathbf{A}^T \mathbf{M}^{(k)} \mathbf{A}$$

$$\mathbf{x}^{(k+1)} = \mathbf{A}[\mathbf{Q}^{(k)}]^{-1} \left(\mathbf{B}^T \mathbf{B} \mathbf{A}^{-1} \mathbf{y} - \lambda_0 \mathbf{A}^T \mathbf{b} \right)$$

BEADS Algorithm

Input: \mathbf{y} , \mathbf{A} , \mathbf{B} , λ_i , $i = 0, \dots, M$

1. $\mathbf{b} = \mathbf{B}^T \mathbf{B} \mathbf{A}^{-1} \mathbf{y}$
2. $\mathbf{x} = \mathbf{y}$ (Initialization)

Repeat

3.
$$[\Lambda_i]_{n,n} = \frac{\phi'([\mathbf{D}_i \mathbf{x}]_n)}{[\mathbf{D}_i \mathbf{x}]_n}, \quad i = 0, \dots, M,$$

4.
$$\mathbf{M} = \sum_{i=0}^M \lambda_i \mathbf{D}_i^T \Lambda_i \mathbf{D}_i$$

5.
$$\mathbf{Q} = \mathbf{B}^T \mathbf{B} + \mathbf{A}^T \mathbf{M} \mathbf{A}$$

6.
$$\mathbf{x} = \mathbf{A} \mathbf{Q}^{-1} \mathbf{b}$$

Until converged

8.
$$\mathbf{f} = \mathbf{y} - \mathbf{x} - \mathbf{B} \mathbf{A}^{-1} (\mathbf{y} - \mathbf{x})$$

Output: \mathbf{x} , \mathbf{f}

Evaluation 1

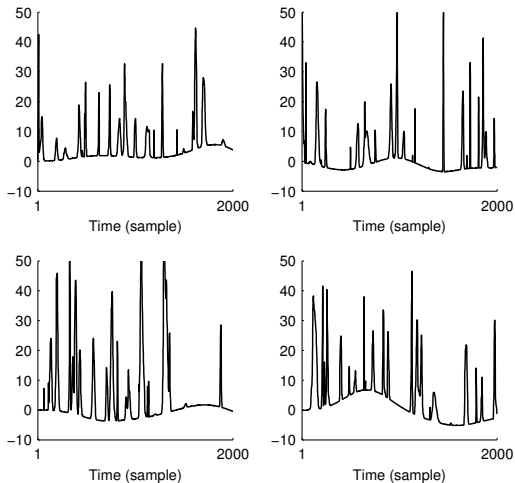
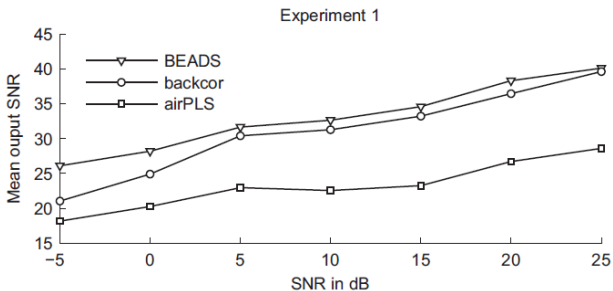


Figure : Simulated chromatograms w/ polynomial+sine baseline.

Evaluation 1 with Gaussian noise



	0 dB		10 dB		20 dB	
	Mean	Std	Mean	Std	Mean	Std
BEADS	28.1	8.52	32.64	8.02	38.33	6.74
backcor	24.91	9.75	31.27	8.33	36.47	6.53
airLPS	20.26	9.65	22.54	10.15	26.71	7.76

Evaluation 2

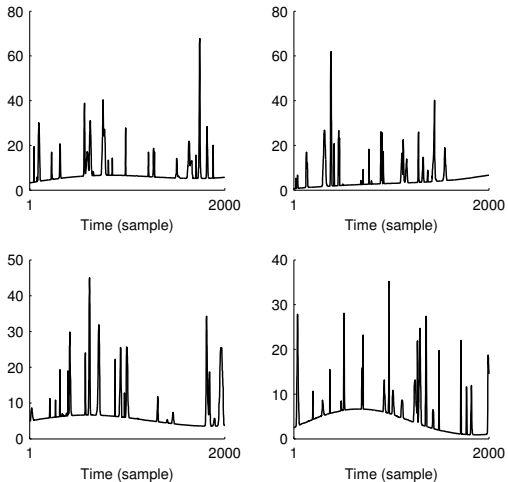
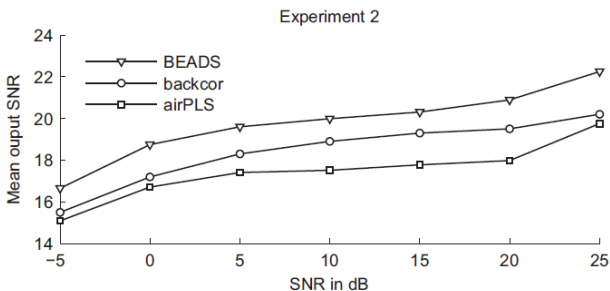


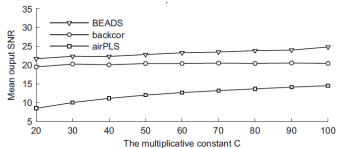
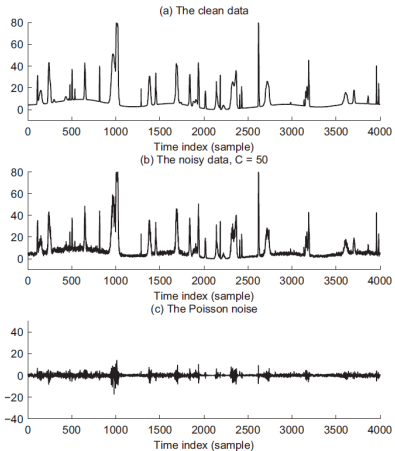
Figure : Simulated chromatograms w/ limited power spectrum noise.

Evaluation 2 with Gaussian noise



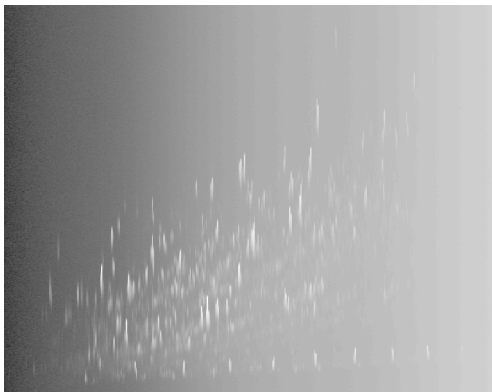
	0 dB		10 dB		20 dB	
	Mean	Std	Mean	Std	Mean	Std
BEADS	18.75	3.71	19.99	3.17	20.89	3.32
backcor	17.20	4.57	18.93	3.74	19.54	3.18
airLPS	16.71	4.80	17.52	5.54	17.98	4.82

Evaluation 3 with Poisson noise



Two-dimensional chromatography data 1

Hyphenated, two-dimensional gas chromatography data



Original data

Two-dimensional chromatography data 1

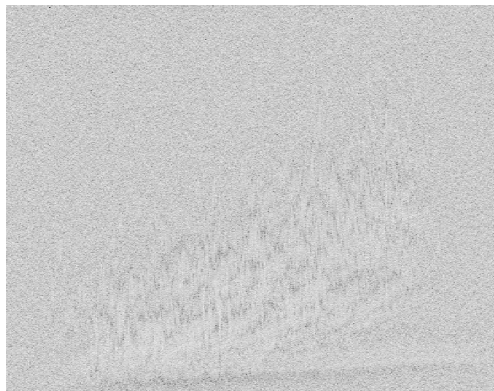
Hyphenated, two-dimensional gas chromatography data



2D background

Two-dimensional chromatography data 1

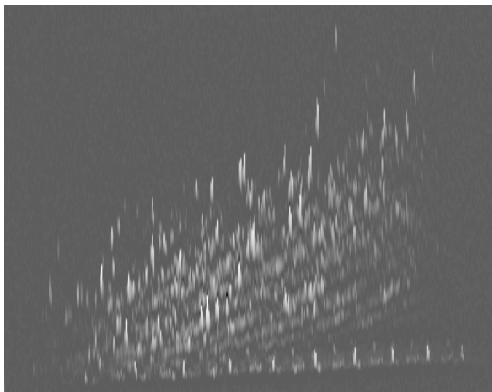
Hyphenated, two-dimensional gas chromatography data



Noise

Two-dimensional chromatography data 1

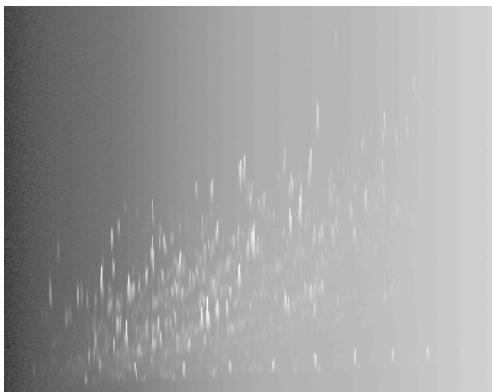
Hyphenated, two-dimensional gas chromatography data



Corrected

Two-dimensional chromatography data 1

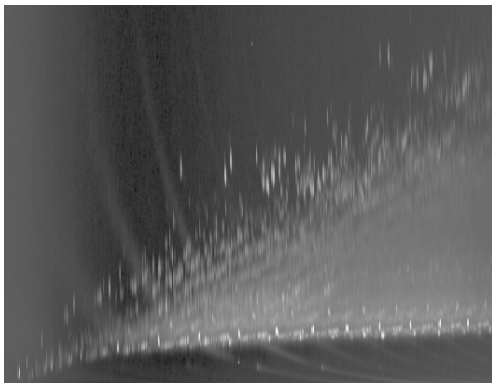
Hyphenated, two-dimensional gas chromatography data



Original data (again!)

Two-dimensional chromatography data 2

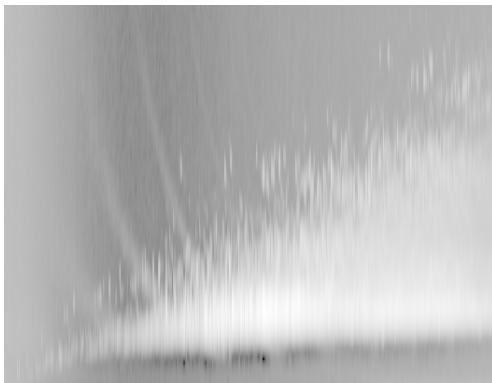
Hyphenated, two-dimensional gas chromatography data



Original data

Two-dimensional chromatography data 2

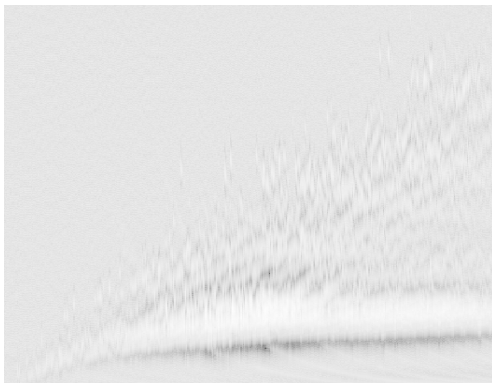
Hyphenated, two-dimensional gas chromatography data



2D background

Two-dimensional chromatography data 2

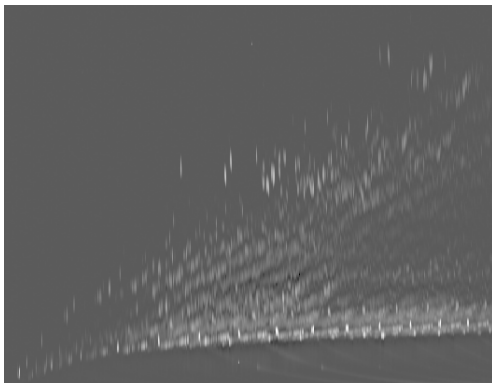
Hyphenated, two-dimensional gas chromatography data



Noise

Two-dimensional chromatography data 2

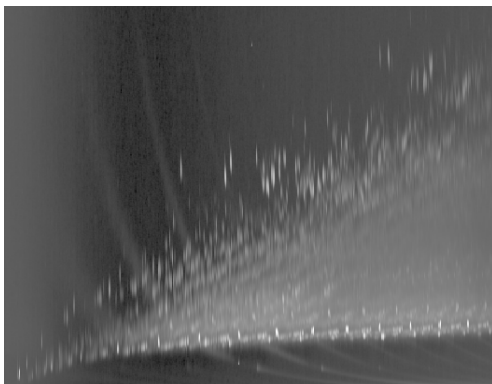
Hyphenated, two-dimensional gas chromatography data



Corrected

Two-dimensional chromatography data 2

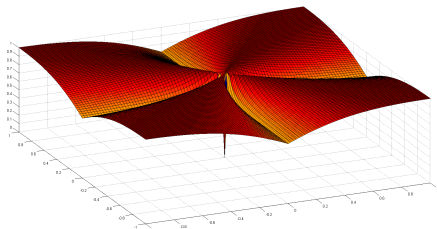
Hyphenated, two-dimensional gas chromatography data



Original data (again!)

Conclusions and work to come

- ▶ BEADS: Baseline Estimation And Denoising... Sparsely
- ▶ Asymmetric penalties with Majorization-Minimization
- ▶ Matlab toolbox: <http://lc.cx/beads>
- ▶ Important pre-processing for image alignment
- ▶ Further tests on other analytical chemistry signals for routine analysis
 - ▶ gas, liquid or ion chromatography; infrared, Raman, Nuclear Magnetic Resonance (NMR) spectroscopy; mass spectrometry
- ▶ Use closer to ℓ_0 sparse penalties: SOOT or smoothed ℓ_1/ℓ_2



More for free: additional references



V. Mazet, C. Carteret, D. Brie, J. Idier, and B. Humbert.

Background removal from spectra by designing and minimising a non-quadratic cost function.
Chemometr. Intell. Lab. Syst., 76(2):121–133, 2005.



C. Vendevre, R. Ruiz-Guerrero, F. Bertoncini, L. Duval, D. Thiébaud, and M.-C. Hennion.

Characterisation of middle-distillates by comprehensive two-dimensional gas chromatography (GC × GC): A powerful alternative for performing various standard analysis of middle-distillates.
J. Chrom. A, 1086(1-2):21–28, 2005.



C. Vendevre, R. Ruiz-Guerrero, F. Bertoncini, L. Duval, and D. Thiébaud.

Comprehensive two-dimensional gas chromatography for detailed characterisation of petroleum products.
Oil Gas Sci. Tech., 62(1):43–55, 2007.



X. Ning, I. W. Selesnick, and L. Duval.

Chromatogram baseline estimation and denoising using sparsity (BEADS).
Chemometr. Intell. Lab. Syst., 139:156–167, Dec. 2014.



A. Repetti, M. Q. Pham, L. Duval, E. Chouzenoux, and J.-C. Pesquet.

Euclid in a taxicab: Sparse blind deconvolution with smoothed ℓ_1/ℓ_2 regularization.
IEEE Signal Process. Lett., 22(5):539–543, May 2015.



C. Couprie, M. Moreaud, L. Duval, S. Henon and V. Souchon,

BARCHAN: Blob Alignment for Robust CHromatographic ANalysis.
J. Chrom. A, 2016...