

Bayesian Wavelet-Based Functional Mixed Models

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Functional Data Analysis

- **Functional Data:**
 - Ideal units of observation: **curves**
 - Observed data: **curves sampled on fine grid**
- Increasingly encountered in scientific research
- FDA (Ramsay & Silverman, 1997)

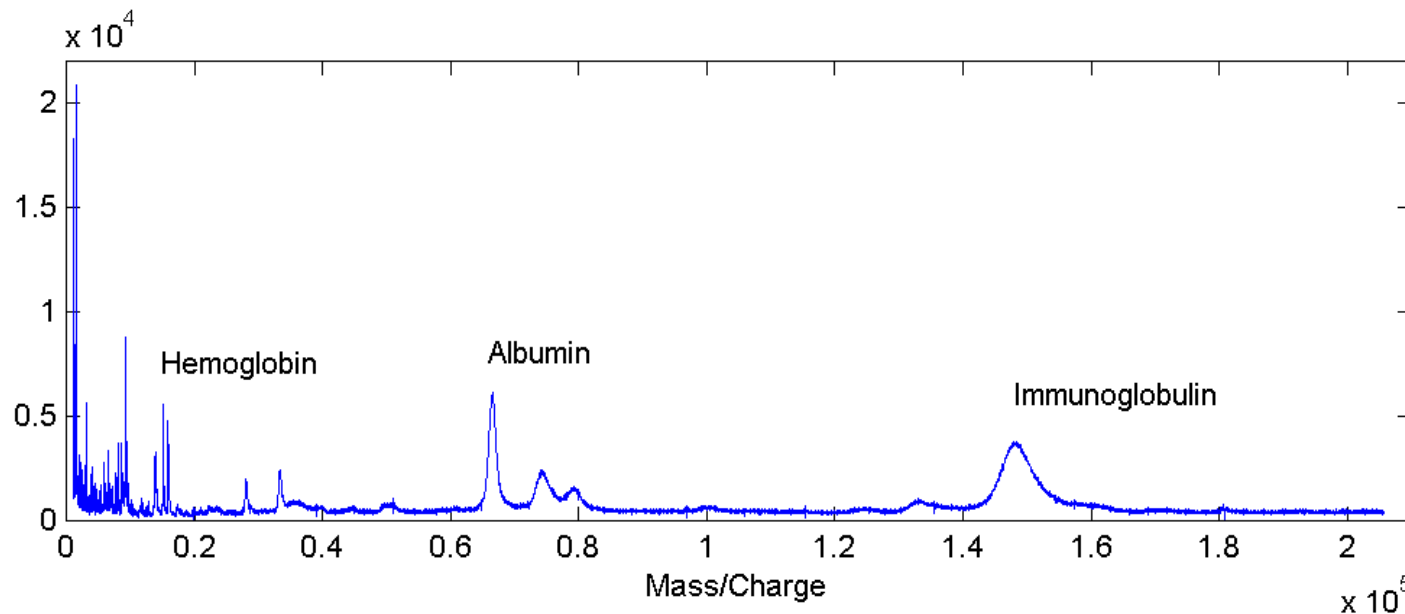
2 issues with functional data

1. **Regularization:** exploiting the assumed smoothness or regularity between measurements within a curve
2. **Replication:** combining information across N curves

Example: Mass Spectrometry Proteomics

- **Central dogma:** DNA \rightarrow mRNA \rightarrow protein
- **Microarrays:** measure expression levels of 10,000s of genes in sample (amount of mRNA)
- **Proteomics:** look at proteins in sample.
 - Gaining increased attention in research
 - Proteins more biologically relevant than mRNA
 - Can use readily available fluids (e.g. blood, urine)
 - **MALDI-TOF:** mass spectrometry instrument that can see 100s or 1000s of proteins in sample

Sample MALDI-TOF Spectrum



- **MALDI-TOF Spectrum: observed function**
- $g(t)$ = intensity of spectrum at m/z value t
- Intensity at peak (roughly) estimates the abundance of some protein with molecular weight of t Daltons

Example: Mouse proteomics study

- 16 nude mice had 1 of 2 cancer **cell lines** injected into 1 of 2 **organs** (**lung** or **brain**)
- **Cell lines:**
 - **A375P:** human melanoma, low metastatic potential
 - **PC3MM2:** human prostate, highly metastatic
- Blood Serum extracted from each mouse – placed on 2 MALDI plates
- Samples run at 2 different **laser intensities** (**low/ high**)
- Total of 32 spectra (observed functions), 2 per mouse
- Sampled on equally-spaced grid of roughly 24,000
 - Downsampled to grid of size 2000

Example: Mouse proteomics study

- **Goal:**

Find proteins differentially expressed by:

- Host organ site (lung/brain)
- Donor cell line (A375P/PC3MM2)
- Organ-by-cell line interaction

- **Combine information across laser intensities:**

Requires us to include in modeling:

- Functional **laser intensity effect**
- **Random effect functions** to account for correlation between spectra from same mouse

Linear Mixed Models

Linear Mixed Model (Laird and Ware, 1982):

$$\underbrace{Y}_{N \times 1} = \underbrace{X}_{N \times p} \underbrace{\beta}_{p \times 1} + \underbrace{Z}_{N \times m} \underbrace{u}_{m \times 1} + \underbrace{e}_{N \times 1}$$

$$\begin{aligned} u &\sim N(0, \overbrace{P}^{m \times m}) \\ e &\sim N(0, \underbrace{R}_{N \times N}) \end{aligned}$$

- **Fixed effects** part, $X\beta$, accommodate a broad class of mean structures, including main effects, interactions, and linear coefficients.
- **Random effects** part, Zu , provide a convenient mechanism for modeling correlation among the N observations.
- Marginally, $Y \sim N(X\beta, ZPZ' + R)$

Functional Mixed Model (FMM)

Suppose we observe a sample of N curves,
 $Y_i(t)$, $i=1, \dots, N$, on a compact set \mathcal{T}

$$\underbrace{Y(t)}_{N \text{ functions}} = \underbrace{\overbrace{X}^{N \times p}}_{p \text{ functions}} \underbrace{B(t)}_{p \text{ functions}} + \underbrace{\overbrace{Z}^{N \times m}}_{m \text{ functions}} \underbrace{U(t)}_{m \text{ functions}} + \underbrace{E(t)}_{N \text{ functions}}$$

$$\begin{aligned} U(t) &\sim \text{MGP}(P, Q) \\ E(t) &\sim \text{MGP}(R, S) \end{aligned}$$

- **DEFN:** $U(t) \sim \text{MGP}(P, Q)$ implies the rows of $P^{-1/2}U(t)$ are ind. mean zero Gaussian Processes with covariance surface $Q(t_1, t_2)$.
 - Functional generalization of **Matrix Normal** (Dawid, 1981).
 - Implies $\text{Cov}\{U_i(t_1), U_j(t_2)\} = P_{ij} * Q(t_1, t_2)$
- **P** and **R** are covariance matrices (between-curve)
- $Q(t_1, t_2)$ and $S(t_1, t_2)$ are covariance surfaces on $\mathcal{T} \times \mathcal{T}$

Discrete Version of FMM

Suppose each observed curve is sampled on a common equally-spaced grid of length T .

$$\underbrace{Y}_{N \times T} = \underbrace{X}_{N \times p} \underbrace{B}_{p \times T} + \underbrace{Z}_{N \times m} \underbrace{U}_{m \times T} + \underbrace{E}_{N \times T}$$

$$\begin{aligned} U &\sim MN(P, Q) \\ E &\sim MN(R, S) \end{aligned}$$

- U and E follow the **Matrix Normal distn.**
 - $U \sim MN(P, Q)$ implies $\text{Cov}\{U_{ij}, U_{i'j'}\} = P_{ii'} * Q_{jj'}$
- P and R are covariance matrices ($m \times m$ & $N \times N$)
- Q and S are within-curve covariance matrices ($T \times T$)

Functional Mixed Models

- **Key feature of FMM:** Does not require specification of parametric form for curves
- Most existing literature for nonparametrically modeling functional data is based on kernels or splines.
- Kernels/splines may not work well for spatially heterogeneous data

Introduction to Wavelets

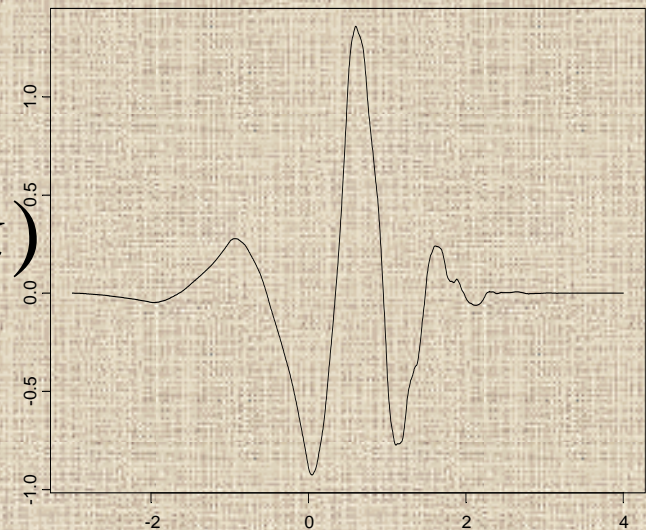
- Wavelets: families of orthonormal basis functions

$$g(t) = \sum_{j,k \in \mathfrak{T}} d_{jk} \psi_{jk}(t)$$

$$\psi_{jk}(t) = 2^{-j/2} \psi(2^{-j/2} t - k)$$

$$d_{jk} = \int g(t) \psi_{jk}(t) dt$$

Daubechies Basis Function



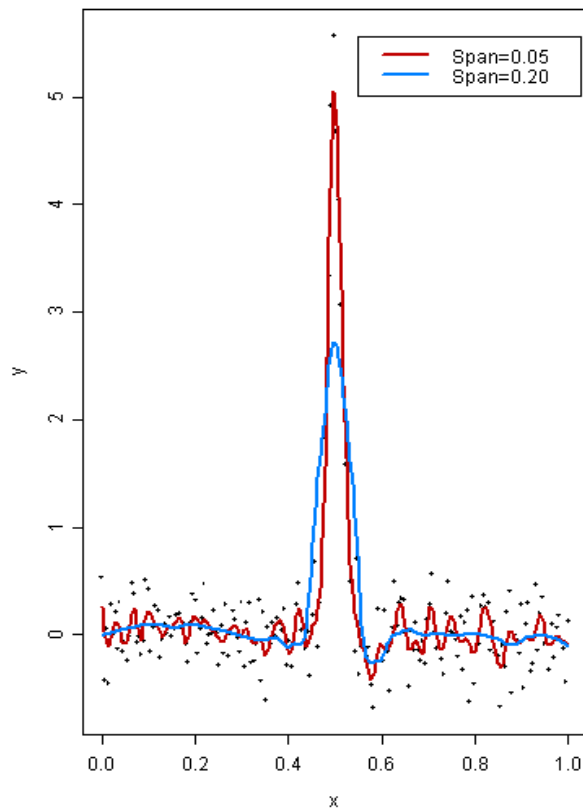
- Discrete Wavelet Transform (DWT): fast algorithm $\{O(T)\}$ for obtaining empirical wavelet coefficients for curves sampled on equally-spaced grid of length T .

Wavelet Regression

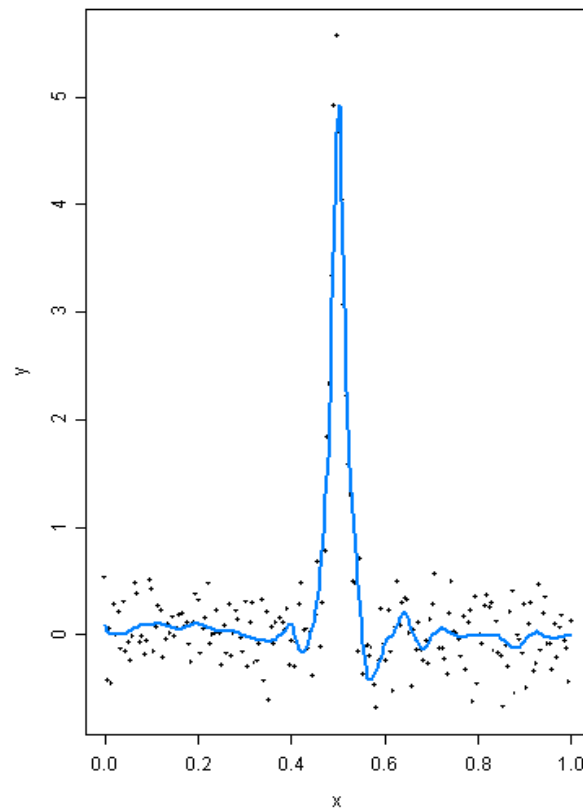
- Useful properties of wavelets:
 - Whitening property
 - Compact support
 - Parsimonious representation
- **Wavelet Regression** – 3 step process
 1. Project data into wavelet space
 2. Threshold/shrink coefficients
 3. Project back to data space
- Yields *adaptively regularized* nonparametric estimates

Adaptive Regularization

Regularization by Local Linear Smoothing



Adaptive Regularization by Wavelet Shrinkage



Wavelet-Based Hierarchical Functional Models

- Most existing wavelet regression methods for single function case
- **Morris, Vannucci, Brown, and Carroll (2003)**
 - Bayesian wavelet-based method for estimating mean function for functional data from nested design.
 - Extends wavelet regression to hierarchical functional context.
- **Goal:** Develop Bayesian wavelet-based methodology for much more general setting of functional mixed models.

Wavelet-Based FMM:

General Approach

1. **Project** observed functions Y **into wavelet space.**
2. **Fit FMM** in wavelet space.
(Use MCMC to get posterior samples)
3. **Project** wavelet-space estimates
(posterior samples) **back to data space.**

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Wavelet-Based FMM

1. Project observed functions Y to wavelet space

- Apply DWT to rows of Y to get wavelet coefficients corresponding to each observed function

$$\underbrace{D}_{N \times T} = \underbrace{Y}_{N \times T} \underbrace{W'}_{T \times T}$$

- Projects the observed curves into the space spanned by the wavelet bases.

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Projecting FMM to Wavelet Space

$$\underbrace{Y}_{N \times T} = \underbrace{X}_{N \times p} \underbrace{B}_{p \times T} + \underbrace{Z}_{N \times m} \underbrace{U}_{m \times T} + \underbrace{E}_{N \times T}$$

$$U \sim MN(P, Q)$$
$$E \sim MN(R, S)$$

Projecting FMM to Wavelet Space

$$\underbrace{Y}_{N \times T} \underbrace{\mathbf{W}'}_{T \times T} = \underbrace{X}_{N \times p} \underbrace{B}_{p \times T} \underbrace{\mathbf{W}'}_{T \times T} + \underbrace{Z}_{N \times m} \underbrace{U}_{m \times T} \underbrace{\mathbf{W}'}_{T \times T} + \underbrace{E}_{N \times T} \underbrace{\mathbf{W}'}_{T \times T}$$

$$U\mathbf{W}' \sim MN(P, \mathbf{W}Q\mathbf{W}')$$

$$E\mathbf{W}' \sim MN(R, \mathbf{W}S\mathbf{W}')$$

Projecting FMM to Wavelet Space

$$\underbrace{D}_{N \times T} = \underbrace{X}_{N \times p} \underbrace{B^*}_{p \times T} + \underbrace{Z}_{N \times m} \underbrace{U^*}_{m \times T} + \underbrace{E^*}_{N \times T}$$

$$U^* \sim MN(P, Q^*)$$
$$E^* \sim MN(R, S^*)$$

Wavelet Space FMM

D : empirical wavelet coefficients for observed curves

Row i contains wavelet coefficients for observed curve i

Each column **double-indexed** by wavelet scale j and location k

$$\underbrace{D}_{N \times T} = \underbrace{X}_{N \times p} \underbrace{B^*}_{p \times T} + \underbrace{Z}_{N \times m} \underbrace{U^*}_{m \times T} + \underbrace{E^*}_{N \times T}$$

$$\begin{aligned} U^* &\sim MN(P, Q^*) \\ E^* &\sim MN(R, S^*) \end{aligned}$$

- $B^*=BW'$ & $U^*=UW'$: Rows contain wavelet coefficients for the fixed and random effect functions,
- $E^*=EW'$ is the matrix of residuals
- $Q^*=WQW'$ and $S^*=WSW'$ model the covariance structure between wavelet coefficients for a given function.
- P, Q^*, R and S^* are typically too large to estimate in an unstructured fashion.

Covariance Assumptions

- We choose **parametric structures** for P and R to model the covariance structure between the curves.
 - Based on the experimental design
 - As in linear mixed models.
- We assume the between-wavelet covariance matrices Q^* and S^* are **diagonal**.
 - Assume wavelet coefficients within given function independent
 - Heuristically justified by whitening property of DWT
 - Common assumption in wavelet regression
 - Is parsimonious in wavelet space (T parameters), yet leads to flexible class of covariance structures in data space

Wavelet Space Model

$$\underbrace{D}_{N \times T} = \underbrace{X}_{N \times p} \underbrace{B^*}_{p \times T} + \underbrace{Z}_{N \times m} \underbrace{U^*}_{m \times T} + \underbrace{E^*}_{N \times T}$$

$$U^* \sim MN(P, Q^*)$$
$$E^* \sim MN(R, S^*)$$

Model Each Column Separately

$$\underbrace{d_{jk}}_{N \times 1} = \underbrace{X}_{N \times p} \underbrace{B_{jk}^*}_{p \times 1} + \underbrace{Z}_{N \times m} \underbrace{U_{jk}^*}_{m \times 1} + \underbrace{E_{jk}^*}_{N \times 1}$$

$$U_{jk}^* \sim N(0, P \cdot Q_{jk}^*)$$
$$E_{jk}^* \sim N(0, R \cdot S_{jk}^*)$$

Single Wavelet Coefficient Model

- Independence assumption allows us to fit wavelet-space model **one column at a time**.
 - i.e., we have a series of T (scalar) mixed models, with the only shared parameters being the between-curve covariance parameters in P and R .
- In principal, we could fit this model using standard mixed models software.
- However, fitting this model without additional mechanism for regularization would result in **rough, noisy estimates** of the p fixed effects functions $B_i(t)$

Prior Assumptions

Mixture prior on β_{ijk}^* :

$$\beta_{ijk}^* = \gamma_{ijk}^* N(0, \tau_{ij}) + (1 - \gamma_{ijk}^*) \delta_0$$

$$\gamma_{ijk}^* = \text{Bernoulli}(\pi_{ij})$$

- Nonlinearly shrinks β_{ijk}^* towards 0, leading to **adaptively regularized** estimates of $B_i(t)$.
- τ_{ij} & π_{ij} are **regularization parameters**
 - Can be estimated from the data using **empirical Bayes**
 - Extend Clyde&George (1999) to functional mixed model

Model Fitting

- **MCMC** to obtain posterior samples of model quantities
 - Work with marginal likelihood; U^* integrated out;
- Let Ω be a vector containing ALL covariance parameters (i.e. for P , Q^* , R , and S^*).

MCMC Steps

1. Sample from $f(B^*/D, \Omega)$:

Mixture of normals and point masses at 0 for each i, j, k .

2. Sample from $f(\Omega/D, B^*)$:

Metropolis-Hastings steps for each j, k

3. If desired, sample from $f(U^*/D, B^*, \Omega)$:

Multivariate normal

Wavelet-Based FMM:

General Approach

1. **Project** observed functions Y **into wavelet space.**
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Wavelet-Based FMM

3. **Project** wavelet-space estimates (posterior samples) **back to data space**.
 - Apply IDWT to posterior samples of B^* to get posterior samples of fixed effect functions $B_i(t)$ for $i=1, \dots, p$, on grid t .
 - **$B=B^*W$**
 - Posterior samples of $U(t)$, P , Q , R , and S are also available, if desired.
 - Can be used for Bayesian inference/prediction

Example: Model

Let $Y(t)$ be the $N=32$ MALDI spectra, preprocessed.

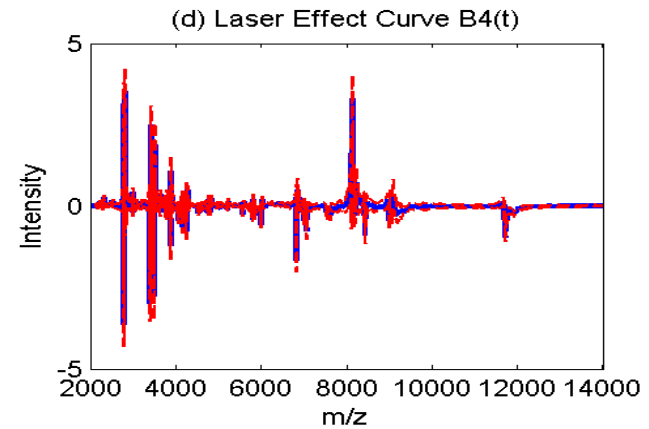
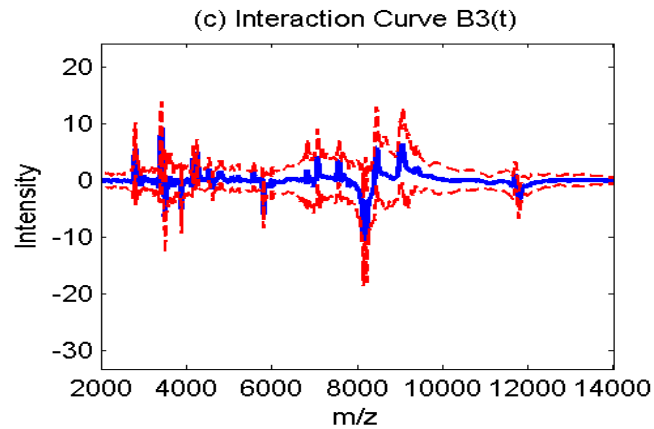
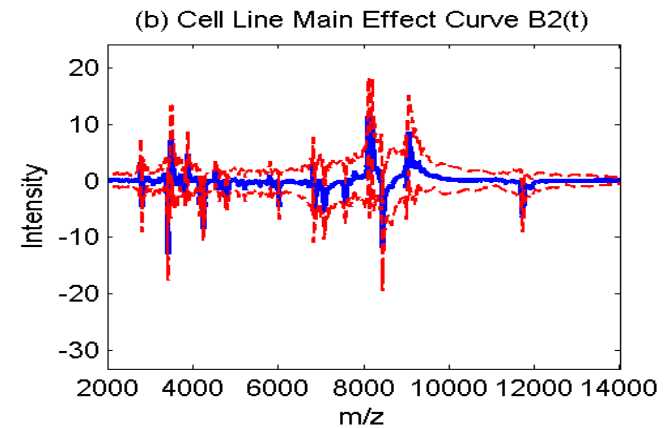
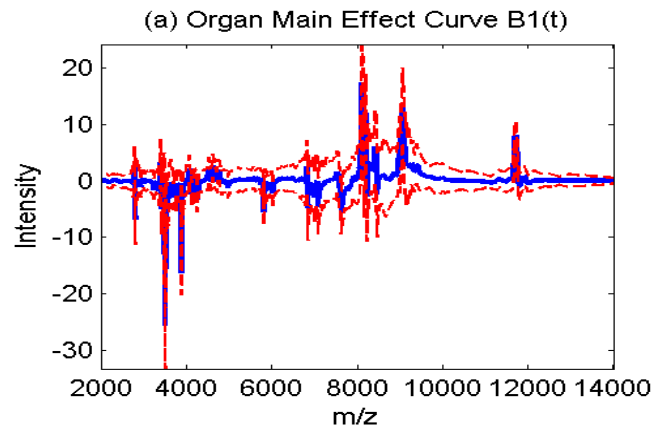
$$Y(t) = XB(t) + ZU(t) + E$$

- $X = \{1, X_{organ}, X_{cell-line}, X_{int}, X_{laser}\}$, where
 $X_{organ} = 1$ for lung, -1 brain. $X_{cell-line} = 1$ for A375P, -1 for PC3MM2
 $X_{int} = X_{organ} * X_{cell-line}$ $X_{laser} = 1$ for low laser intensity, -1 high.
- $B(t) = \{B_0(t), B_1(t), B_2(t), B_3(t), B_4(t)\}$, where
 $B_0(t)$ = overall mean spectrum $B_1(t)$ = organ main effect function
 $B_2(t)$ = cell-line main effect $B_3(t)$ = org x cell-line int function
 $B_4(t)$ = laser intensity effect function
- $Z = 1_2 \otimes I_{16}$, and $U(t) = \{U_1(t), \dots, U_{16}(t)\}$ are mouse random effect functions.

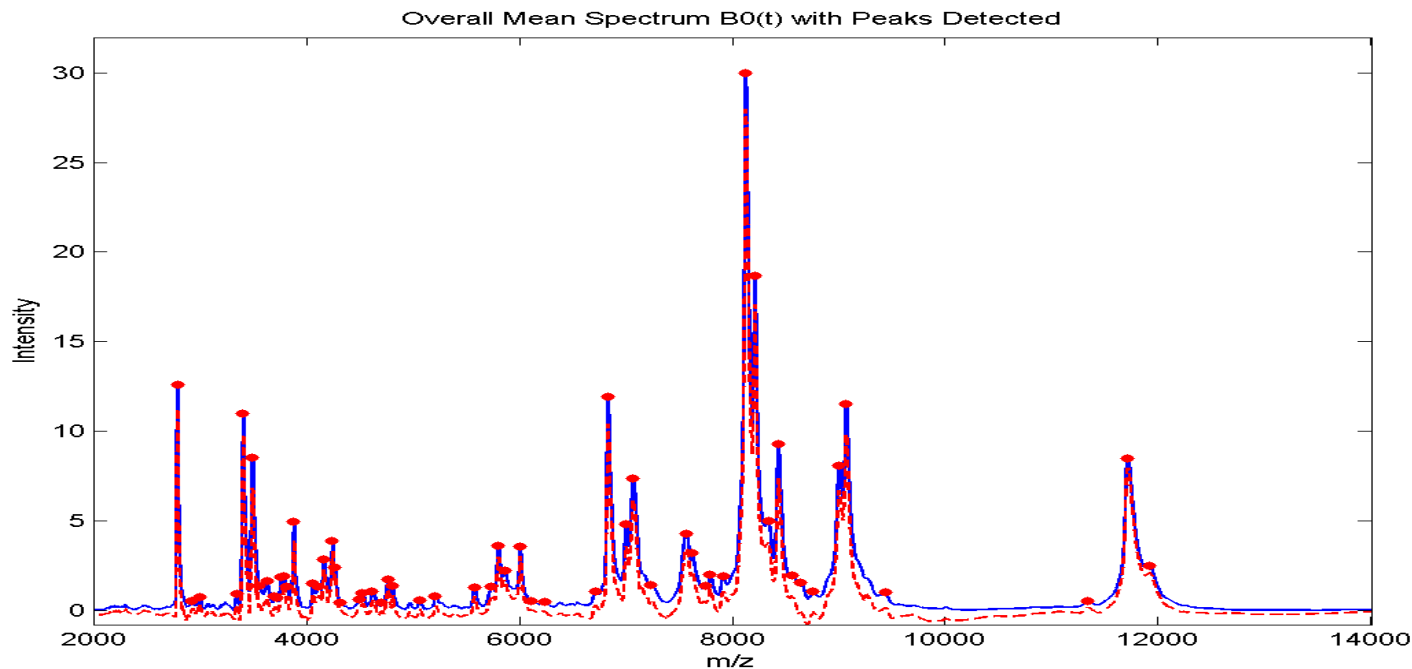
Example: Model Fitting

- Daubechies 8 wavelet basis, $J=11$ levels
- **Empirical Bayes** procedure used to estimate regularization parameters π_{ij} and τ_{ij} from data.
- Burn-in 1000; 20,000 MCMC samples; thin=10
- Took **7hr 53min** on Win2000 P-IV 2.8GHz 2GB RAM
 - That is Matlab code; C++ code takes ~2 hours.
- Trace plots indicated good convergence properties
- Metropolis Hastings acceptance probabilities good:
 - Range of (0.04, 0.53)
 - (10th, 50th, 90th) percentiles of (0.20, 0.29, 0.50)

Example: Results



Example: Peak detection



- We first did “**peak detection**”
Local maxima in posterior mean (denoised) estimate of $B_0(t)$ with High posterior probability of nonzero mean; $\Pr\{B_0(t) > 0/Y\} \geq 0.95$
- Using this criterion, we found 58 peaks
- We restrict inference to values of t at peaks

Example: Flagged peaks

Detecting ‘significant’ peaks: (assoc. w/ organ,cl,int)

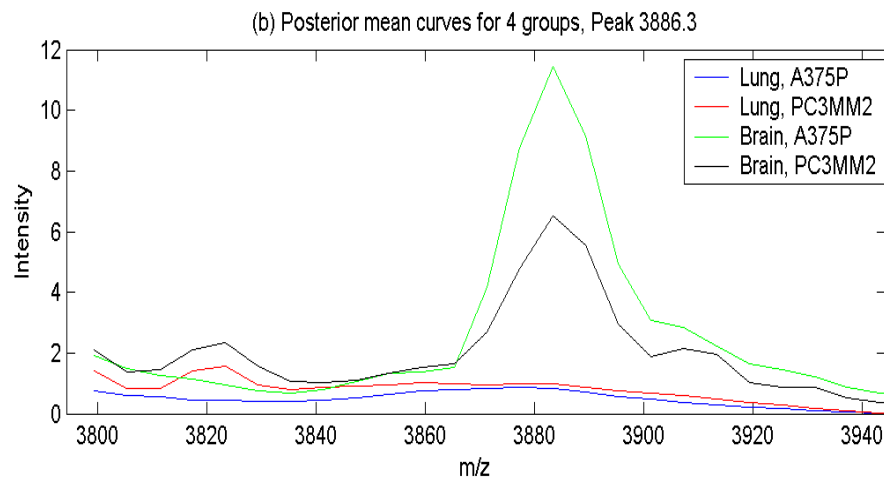
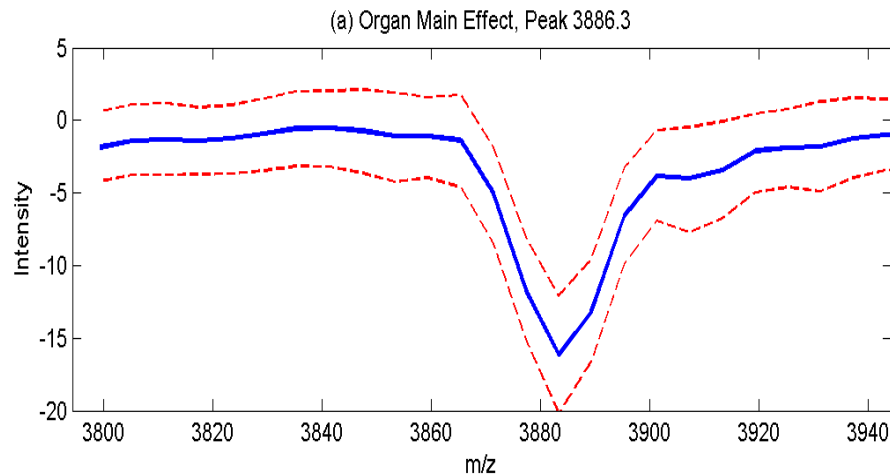
For each t at a peak, compute $p_i(t) = \min[\Pr\{B_i(t) > 0\}, \Pr\{B_i(t) < 0\}]$

Flag any peak for which $p_i(t)$ is very small. ($< 0.05/58 = 0.00086$)

Using this criterion, we flagged **9 peaks** as interesting

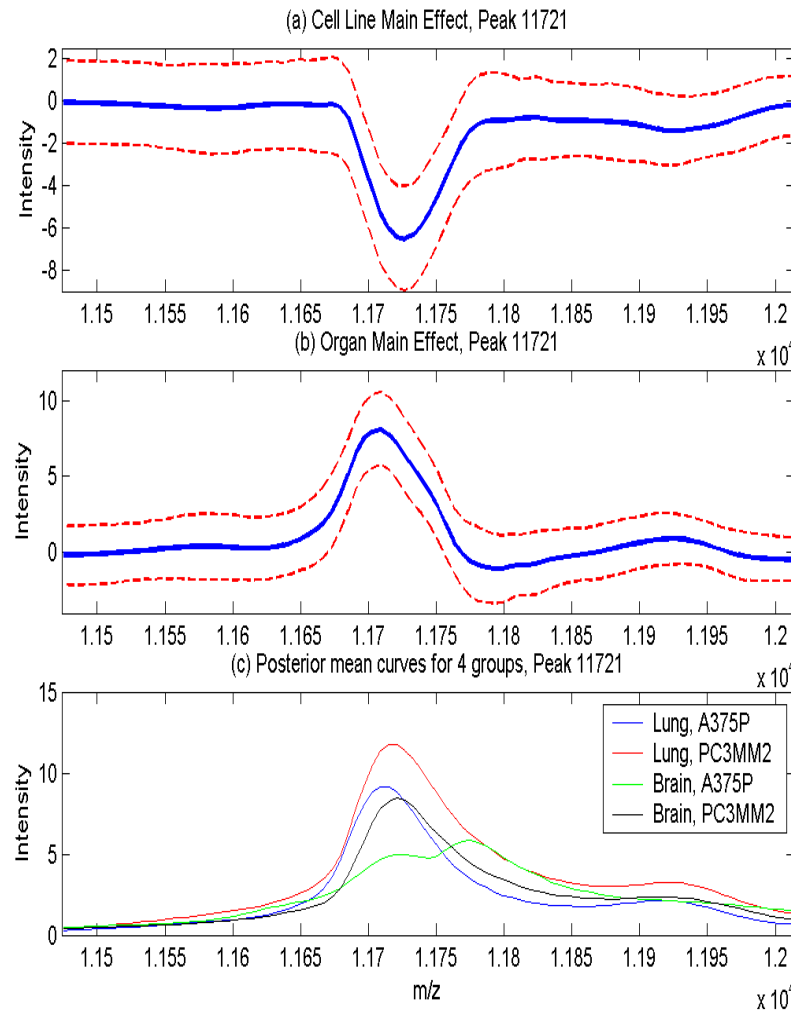
m/z	Effect	p	Comment
3412.6	int.	<0.0005	PC3MM2>A375P for brain-injected only
3496.6	organ	<0.0005	Only expressed in brain-injected mice
3886.3	organ	<0.0005	Only expressed in brain-injected mice
4168.2	int.	0.0005	PC3MM2>A375P in brain-injected only
4252.1	int.	<0.0005	PC3MM2>A375P in brain-injected only
4270.1	cell line	<0.0005	PC3MM2>A375P
5805.3	int.	<0.0005	brain>lung only for mice given A375P cell-line
6015.2	cell line	<0.0005	PC3MM2>A375P
11721	cell line	<0.0005	PC3MM2>A375P
11721	organ	<0.0005	lung>brain

Example: Results



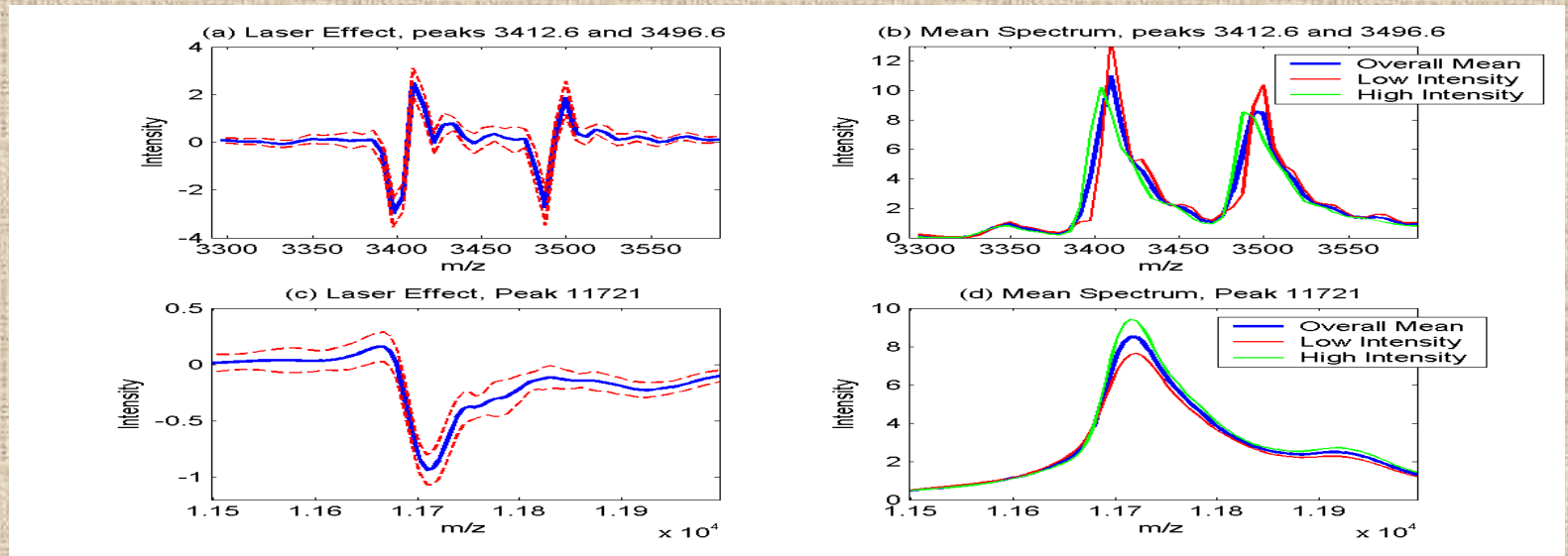
- Specific to brain-injected mice
- May be **CGRP-II** (3882.34 Dal), peptide in mouse proteome that dilates blood vessels in brain
- Host response to tumor implanted in brain?

Example: Results



- **Higher in mice injected with metastatic (PC3-MM2) cell line**
- **May be *MTS1* (11721.43 Dalt), metastatic cell protein in mouse proteome.**
- **Also higher in lung-injected mice than brain-injected mice**

Example: Results



- **Laser intensity effect adjusts for:**
 - **Offsets** in m/z scale
 - **Shifts** in intensities
- Important proof of principle that “linear” functional term can be used to adjust for functional effects of nuisance factors

Discussion

- Introduced unified modeling approach for FDA
 - Applied here to MALDI-TOF, but method is general.
- Method based on mixed models; is **FLEXIBLE**
 - Accommodates a **wide range of experimental designs**
 - Addresses **large number of research questions**
- Posterior samples allow **Bayesian inference and prediction**
 - **Posterior credible intervals**; pointwise or joint
 - **Predictive distributions** for future sampled curves
 - **Predictive probabilities** for group membership of new curves
 - Bayesian functional inference can be done via **Bayes Factors**
- Since a unified modeling approach is used, all **sources of variability** in the model **propagated throughout inference**.

Discussion

- Since functions adaptively regularized using wavelet shrinkage, the method is **appropriate for spatially heterogeneous functional data**.
- Approach is Bayesian. The **only informative priors to elicit are regularization parameters**, which can be estimated from data using empirical Bayes.
- Method **generalizes to higher dimensional functions**, e.g. image data, space/time (fixed domain) data.
- We used wavelet bases, but approach can be generalized to **other orthogonal basis functions**.
- Difficult to develop unified statistical modeling approach for replicated functional data, but worth the effort.

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