

Nonparametric Covariance Estimation for Longitudinal Data via Penalized Tensor Product Splines

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Abstract

With high dimensional longitudinal and functional data becoming much more common, there is a strong need for methods of estimating large covariance matrices. Estimation is made difficult by the instability of sample covariance matrices in high dimensions and a positive-definite constraint we desire to impose on estimates. A Cholesky decomposition of the covariance matrix allows for parameter estimation via unconstrained optimization as well as a statistically meaningful interpretation of the parameter estimates. Regularization improves stability of covariance estimates in high dimensions, as well as in the case where functional data are sparse and individual curves are sampled at different and possibly unequally spaced time points. By viewing the entries of the covariance matrix as the evaluation of a continuous bivariate function at the pairs of observed time points, we treat covariance estimation as bivariate smoothing.

Within regularization framework, we propose novel covariance penalties which are designed to yield natural null models presented in the literature for stationarity or short-term dependence. These penalties are expressed in terms of variation in continuous time lag and its orthogonal complement. We present numerical results and data analysis to illustrate the utility of the proposed method.

keywords: non-parametric, covariance, longitudinal data, functional data, splines, reproducing kernel Hilbert space

1 Introduction

Covariance estimation with high dimensional data has recently gained growing interest; it is generally recognized that there are two primary hurdles responsible for the difficulty in covariance estimation: the instability of sample covariance matrices in high dimensions and a positive-definite constraint we wish estimates to obey. Estimation of population covariance matrices from samples

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of multivariate data has been important for methods in classical multivariate data analysis, time series analysis, spatial statistics and, more recently, the growing field of statistical learning. Covariance estimates play a critical role in establishing independence or conditional independence through graphical models, constructing discriminant functions as in linear discriminant analysis (LDA) and quadratic discriminant analysis (QDA) for the classification of Gaussian data, building confidence intervals for component means and contrasts, and constructing a low-dimensional representation of data via principal components analysis (PCA). One may note that the last two techniques require an estimate of the covariance matrix, and the first two require estimation of the inverse.

Prevalent technological advances in industry and many areas of science make high dimensional longitudinal and functional data a common occurrence, arising in numerous areas including medicine, public health, biology, and environmental science with specific applications including fMRI, spectroscopic imaging, gene microarrays among many others, presenting a need for effective covariance estimation in the challenging situation where parameter dimensionality is possibly much larger than the number of observations. Additional difficulty due to constraints required to yield positive definite estimates make covariance estimation a potentially complex optimization problem. Further, most existing approaches to covariance estimation require data to be sampled at regular grid (time) points, with subjects sharing a set of common observation points. However, in many practical situations, data are irregularly sampled, and subjects may share few common observation times, and methods are needed to accommodate for data collected in this way.

To address the challenge of enforcing positive definiteness, several have considered modeling various matrix decompositions including variance-correlation decomposition, spectral decomposition, and Cholesky decomposition. The Cholesky decomposition has received particular attention, as it which allows for a statistically meaningful interpretation as well as an unconstrained parameterization of elements of the covariance matrix. This parameterization allows for estimation to be accomplished as simply as in least squares regression.

It is well known that the sample covariance matrix is unstable in high dimensions, and there is an extensive existing body of work addressing the issue of high dimensionality in the context of covariance estimation. See ? for a survey of approaches to covariance estimation from the generalized linear modeling and regularization perspectives. However, much of this work addresses high dimensionality arising from functional or times series data sampled on a dense, regular grid. With such data, it is typical that the number of time points is larger than the number of observations. Few have addressed the challenges posed by sparse longitudinal data where measurement times may be almost unique yet sparsely distributed within the observed time range for each individual in the study. In this case, high dimensionality may not be a consequence of having more measurements per subject than the number of subjects themselves, but rather because when pooled across subjects, the total number of unique observed time points is greater than the number of individuals. Incomplete and unbalanced data arise when measurement schedules with targeted time points which are not necessarily equally spaced or if there is missing data. Sparse longitudinal data arise when the measurement schedule has arbitrary or almost unique time points for every individual. A given time point may have very few individuals with corresponding measurements.

We sidestep both issues of high dimensionality and irregularly sampled data by viewing the

response as a stochastic process having continuous covariance function. Recent work outlines the use of function estimation for smoothing elements of the covariance matrix, including [1,2]. To our knowledge, however, no previous work has applied smoothing to both dimensions of the Cholesky factor; we model the generalized autoregressive parameters using tensor product splines. Viewing covariance modeling as bivariate function estimation both accommodates irregularly sampled curved and permits interpolation and extrapolation of the covariance function between two measurements at any pair of time points within the time interval of interest rather than at observed pairs of time points only. The Cholesky decomposition enables covariance estimation through the estimation of a varying coefficient model. A transformation of the design point axes allows for an ANOVA-like decomposition of the coefficient function into two components, corresponding to the lag between time points and an additive component. Through this general framework, we can easily impose penalties on fitted functions to yield natural null models presented in the literature.

2 Cholesky Decomposition of Σ

To present a comprehensive overview our estimation procedure, we begin with the representation of the inverse covariance matrix, $\Omega = \Sigma^{-1}$, in terms of its Cholesky decomposition (see [3] for a detailed discussion.) In the section to follow, we will demonstrate that this parameterization of the precision matrix is particularly attractive due to both the computational advantages as well as the convenient modeling interpretation it permits. For any positive definite matrix Σ , there exists a unique unit lower triangular matrix T with diagonal entries equal to 1 which diagonalizes Σ :

$$T\Sigma T' = D$$

If we assume that the data having covariance matrix Σ follow an autoregressive model, then the entries of the Cholesky factor T and D enjoy a useful interpretation. Let $Y = (Y_1, Y_2, \dots, Y_m)'$ be defined on a probability space with some probability measure \mathcal{P} corresponding to the multivariate Normal distribution with mean 0 and covariance Σ , and let Y_1, Y_2, \dots, Y_m have associated measurement times

$$t_1 < t_2 < \dots < t_m.$$

Let \hat{y}_t denote the linear least-squares predictor of y_t based on its predecessors y_1, \dots, y_{t-1} , and let e_t denote its prediction error with variance $Var(e_t) = \sigma_t^2$, $Cov(e_i, e_j) = 0$ for $i \neq j$. Standard regression theory gives us that we can find coefficients ϕ_{tj} such that

$$y_j = \sum_{k=1}^{j-1} \phi_{jk} y_k + \sigma_j e_j, \quad j = 2, \dots, m, . \quad (1)$$

Let T denote the $m \times m$ unit lower triangular matrix with elements

$$T_{jk} = \begin{cases} -\phi_{jk} & j > k \\ 1 & j = k \\ 0 & otherwise, \end{cases}$$

for $j, k = 1, \dots, m$. In matrix notation, model 1 may then be written

$$e = TY, \quad (2)$$

Taking covariances on both sides of 2, we have

$$D = T\Sigma T' \quad (3)$$

An attractive feature of this reparameterisation is that, regardless of the modelling approach, the estimated covariance matrix is guaranteed to be positive definite. The unconstrained regression coefficients $\{\phi_{jk}\}$ are referred to as the *generalized autoregressive parameters* (GARPs). The $\{\sigma_j^2\}$ are called the *innovation variances* (IVs.) Unconstrained estimation of the $\{\sigma_k^2\}$ is achieved by log transformation; we leave these details for section 2. Expressing the precision matrix in terms of the GARPs and IVs, we have

$$\Omega = \Sigma^{-1} = T'D^{-1}T.$$

Rather than estimating a specific covariance matrix for data observed on a fixed, regular grid, we aim to estimate a smooth covariance function. This accomodates data which may consist of observations on multiple subjects measured at potentially unequally spaced and individual-specific times. In estimation of the means μ of p-vectors of i.i.d. variables, the Gaussian white noise model [9] is the appropriate infinite-dimensional model into which all objects of interest are embedded. In estimation of matrices, a natural analogue is the space $B(l_2, l_2)$, which we write as B , of bounded linear operators from l_2 to l_2 . These can be represented as matrices [cite *Regularized estimation of large covariance matrices by Bickel and Levina - section 4.*]

Rather than m -dimensional vectors, consider Y and e as the values of the stochastic processes $Y(t)$ and $e(t)$ at the set of observation times. We assume that $Y(t)$ is equipped with covariance function $G(s, t)$, and

$$e(s) \sim \mathcal{N}(0, 1)$$

is a zero mean Gaussian process with unit variance. We assume that $G(s, t)$ satisfies some smoothness conditions, where smoothness is defined in terms of square integrability of certain derivatives. [TODO: clean up statement about smoothness of covariance function; integrability of covariance function of a stochastic process?] The entries of Σ , then, correspond to G evaluated at the distinct pairs of observed time points. Similarly, we treat the elements of the precision matrix Ω as the evaluation of some smooth function, $\omega(s, t)$.

Similarly, viewing the elements of D and the elements of the Cholesky factor T as the evaluation of smooth functions $\sigma(\cdot)$ and $\phi^*(\cdot, \cdot)$ leads us to the varying coefficient (VC) models first introduced in the seminal work of Hastie and Tibshirani. The procedures presented by ? and ? utilize varying coefficient models for modeling the mean of longitudinal data; parameterizing the covariance matrix according to 3 allows us to exploit these models in covariance estimation for such data as well. A generalization of traditional linear regression models, varying coefficient models offer more flexibility than their static analogues by allowing the effect of covariates to change smoothly with the value other variables. Both regressors and response variables are assumed to vary according to an *indexing variable*, which permits interpolation of regressors and response variables at values of this indexing variable where there is either missing data or only a

single observation and slope estimation is not feasible. Replacing $\{\phi_{jk}\}$ and $\{\sigma_j\}$ with smooth functions, we model

$$y(t_j) = \sum_{k=1}^{j-1} \phi(t_j, t_k) y(t_k) + \sigma(t_j) \epsilon(t_j) \quad j = 1, \dots, m, \quad (4)$$

for $t_1 < t_2 < \dots < t_m$.

We represent the varying coefficient function and the innovation variances using tensor product smoothing splines and penalized tensor product B-splines alongside penalties to induce simplicity in ϕ and σ^2 to produce final covariance estimates exhibiting the desired null structure. For ease of exposition, we first focus our attention on the estimation of ϕ assume that $\sigma^2(t)$ is fixed and known; we will later propose an iterative procedure for simultaneous estimation of σ^2 and ϕ . Recasting the problem as the estimation of model 4 allows us access to the existing set of tools developed in the bivariate smoothing literature; our approach provides a flexible, comprehensive framework for covariance estimation.

3 Penalized Maximum Likelihood Estimation of ϕ

We employ maximum likelihood for the estimation of the varying coefficient function $\phi(t, s)$ and the innovation variance function $\sigma(t)$, though neither the derivation the form of model 1 nor model 4 via the Cholesky decomposition rely on any assumptions about the distribution of Y . Fixing σ_j^2 , for a sample of N i.i.d. observations Y_1, Y_2, \dots, Y_N from a multivariate Gaussian distribution, the negative log-likelihood as a function of ϕ_{jk} corresponds to the usual error sums of squares and is proportional to

$$-2L(y_1, y_2, \dots, y_N, \Phi) \propto \sum_{i=1}^N \sum_{j=2}^{m_i} \sigma(t_j)^{-2} \left(y_{ij} - \sum_{k=1}^{j-1} \phi(t_{ij}, t_{ik}) y_{ik} \right)^2 \quad (5)$$

where

$$y_i = (y_{i1}, y_{i2}, \dots, y_{i, m_i}), \quad i = 1, \dots, N$$

denotes the vector of observations for subject i with corresponding measurement times

$$t_{i1} < t_{i2} < \dots < t_{i, m_i}.$$

The form of the likelihood of y_1, \dots, y_N indicates that we allow both the number of measurements as well as the observation times to vary across subjects. The $\{t_{ij}\}$ need not be evenly-spaced within or across individuals.

In the case that subjects share a common set of observation times $t_1 < \dots < t_m$, it is well known that the MLE for Σ , $S = \sum_{i=1}^N y_i y_i'$ is highly unstable in high dimensions, a condition that is potentially worsened when one or more subjects has at least one observation time that is unique from the set of observation times common across subjects. To mitigate instability due to high dimensionality and simultaneously permit the estimation of $\phi(\cdot, \cdot)$ as a smooth bivariate function, we obtain a covariance estimator by applying bivariate smoothing of the elements of the Cholesky factor.

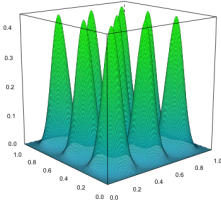


Figure 1: A “thinned” tensor product basis

4 *Smooth via a tensor product B-spline basis*

5 Penalized B-splines for flexible smoothing

Estimating the varying coefficient function ϕ^* is quite different from the usual problem of estimating an arbitrary bivariate function via smoothing. In the case of the latter, we most typically treat both arguments equally in terms of regularization, but in the case of covariance estimation and the generalized coefficient function equal treatment of l and m in terms of penalization perhaps is not the most appropriate approach. The lag component, l , has particularly significant meaning in terms of the covariance function and thus also in terms of ϕ^* and is of considerable more interest than the orthogonal component, m . As discussed in Section 2, we can define an entire class of functional autoregressive models using only the l direction, and additionally, as discussed in Section 3, there is a natural expectation about the functional form of the autoregressive coefficient function (and hence covariance) as a function of l . The use of smoothing splines to estimate ϕ outlined in Section ?? yields smooth null models, but smoothness of the elements of the Cholesky factor alone may not lead to desirable structure in the inverse covariance matrix.

[TODO:] FIX THIS PARAGRAPH

These approaches implicitly adopt different notions of sparsity. Like ? and ?, our aim is to regularize the inverse of the covariance matrix through the Cholesky factor. Expressing the varying coefficient function using a tensor product basis expansion builds the foundation for a flexible estimation framework within which employing multiple notions of smoothness is simple and straightforward.

Equip l and m with

$$\begin{aligned} B_1(l), \dots, B_K(l), \\ B_1(m), \dots, B_L(m) \end{aligned}$$

to build

$$T_{jk}(l, m) = B_j(l) B_k(m)$$

$$\phi^*(l, m) = \sum_{i=1}^K \sum_{j=1}^L \theta_{ij} B_i(l) B_j(m)$$

The truncated power basis, as in their use in defining polynomial regression splines, enjoy a particular ease of interpretation. The coefficient β_{i+k} corresponds to the size of the jump at ℓ_i in the k^{th} derivative of ϕ_1 . This fact is especially useful when tracking change points or in general, any abrupt changes in the regression curve, as the decomposition according to 23 demands. Rather than constructing a basis for \mathcal{H} , we instead consider representing the set of piecewise polynomial functions using B-splines, which exhibit numerical properties which are preferable to those of the TPFs. Let $\ell = \{\ell_i\}$ denote a non-decreasing sequence. The i^{th} B-spline of order k which corresponds to the knot sequence t is defined by

$$B_{i,k,t}(l) = (\ell_{i+k} - \ell_i) [\ell_i, \dots, \ell_{i+k}] (\cdot - l)_+^{k-1}, \quad (6)$$

where $[x_i, \dots, x_{i+k}] f(x)$ denotes the k^{th} order divided difference of the function f at $\{x_i, \dots, x_{i+k}\}$. The placeholder notation, $(\cdot - l)_+^{k-1}$, is used to indicate that the k^{th} divided difference of the function $g(t) = (t - l)_+^{k-1}$ is obtained by fixing l and applying the divided difference to $g(t)$ as a function of t alone. Henceforth, we will write B_i rather than $B_{i,k,t}$ when the spline order and knot sequence can be inferred from surrounding context. Using the properties of the divided difference, one can show that $B_i(x)$ has isolated support:

$$B_i(l) = 0, \quad l \notin [\ell_i, \ell_{i+k}]$$

As a result, for a set of B-splines of order k corresponding to the knot sequence ℓ , only k of them are nonzero on $[\ell_j, \ell_{j+k}]$: $B_{j-k+1}, B_{j-k+2}, \dots, B_j$. See ? for a complete review of B-splines and their properties. A B-spline function is a linear combination of B-spline basis functions:

$$\phi_1(l) = \sum_{i=1}^n \beta_{k+i} B_i(l) \quad (7)$$

5.1 Regularization with difference penalties

Representing the varying coefficient function as a B-spline lays the foundation of a flexible framework for seamlessly instituting multiple types of regularization on the fitted function, and perhaps encouraging more than one notion of structure at once. As in Section ??, one may wish for fitted functions to exhibit smoothness on their support. By using a rich B-spline basis along with a discrete difference penalty on the spline coefficients, we can achieve as much smoothness in the fitted function as desired. ? was the first to propose using a rich B-spline basis and using a penalty to restrict the flexibility of the fitted curve. Like ? in the smoothing spline literature, he applied a penalty to the second derivative of the fitted curve:

$$J = \int_0^1 [\phi_1''(l)]^2 dl.$$

Using the properties of B-splines, for

$$\phi_1(l) = \sum_{j=1}^n \beta_j B_j(l),$$

one can derive a banded matrix P such that

$$J = \beta' P \beta$$

where $\beta = (\beta_1, \dots, \beta_n)$. The i - j^{th} element of P is given by

$$p_{ij} = \int_0^1 B_i''(l) B_j''(l) dl.$$

In some applications, it is useful to work with third and fourth order differences, since for large values of λ , the fitted curve approaches a parametric polynomial model. This may be of particular interest in the context of estimating the elements of the Cholesky factor: many have proposed simple parametric functions of lag only for ϕ^* , such as low order polynomials. See ?. However, with the use of higher order derivatives, the computation of P is nontrivial and becomes very tedious. ? were the first to propose P-splines, or penalized B-splines, as an approach to nonparametric regression which circumvents complexity associated with constructing such penalty matrices by omitting derivatives and integrals altogether, replacing them with finite differences and sums. Instead, flexibility of the fitted function is controlled by using a discrete penalty matrix based on finite difference formulas. Smoothness of the fitted function is achieved by first using a rich B-spline basis with equally spaced knots to purposefully overfit the smooth coefficient vectors. This liberates us from the difficulty of choosing the optimal set of knots. Then by attaching a difference penalty to the goodness of fit measure, one may prevent overfitting and make a potentially ill-conditioned fitting procedure a well-conditioned one.

This is of particular interest within the context of estimating the The finite difference penalty is simple to compute and can be handled mechanically for any order of the differences. Since it is easily introduced into regression equations, it is feasible to evaluate the impact of different orders of the differences on the fitted model. Using the properties of B-splines, it is straightforward to show that the difference penalty is a good discrete approximation to the integrated square of the k^{th} derivative, so little is lost by replacing the derivative-based penalty with the following difference penalty:

$$J_d(\phi_1) = \sum_{j=d}^n (\Delta^d \beta_j)^2 \tag{8}$$

where $\beta = (\beta_1, \dots, \beta_n)$. Let D_d denote the matrix difference operator: $D_d \beta = \Delta^d \beta$, where

$$\Delta \beta_j = \beta_j - \beta_{j-1}, \quad \Delta^2 \beta_j = \Delta(\Delta \beta_j) = \beta_j - 2\beta_{j-1} + \beta_{j-2}$$

In general,

$$\Delta^d \beta_j = \Delta (\Delta^{d-1} \beta_j).$$

Then, 8 can be written in terms of the squared norm of the difference operator applied to the vector of B-spline coefficients:

$$\begin{aligned} J_d(\phi_1) &= \|D_d \beta\|^2 \\ &= \beta' P_d \beta \end{aligned} \quad (9)$$

where $P_d = D_d' D_d$.

By expressing the traditional penalty on the second derivative in terms of the B-spline coefficients, we can examine its connection to a penalty on second-order differences of the third-order B-spline coefficients:

$$\beta' P \beta = \int_0^1 \left\{ \sum_{j=1}^n \beta_j B_{j,3}''(l) \right\}^2 dl. \quad (10)$$

Using the derivative properties of B-splines, this can be written as

$$\beta' P \beta = \int_0^1 \left[\sum_{j=1}^n \sum_{k=1}^n \Delta^2 \beta_j \Delta^2 \beta_k B_{j,1}(l) B_{k,1}(l) \right] dl. \quad (11)$$

Most of the cross products of $B_{j,1}(l)$ and $B_{k,1}(l)$ vanish since B-splines of degree 1 only overlap when j is $k-1$, k , or $k+1$. Thus, we have that

$$\begin{aligned} \beta' P \beta &= \int_0^1 \left[\left\{ \sum_{j=1}^n \Delta^2 \beta_j B_j(l, 1) \right\}^2 + 2 \sum_j \Delta^2 \beta_j \Delta^2 \beta_{j-1} B_j(l, 1) B_{j-1}(l, 1) \right] dl \\ &= \sum_{j=1}^n (\Delta^2 \beta_j)^2 \int_0^1 B_j^2(l, 1) dl \\ &\quad + 2 \sum_{j=1}^n \Delta^2 \beta_j \Delta^2 \beta_{j-1} \int_0^1 B_j(l, 1) B_{j-1}(l, 1) dl \end{aligned} \quad (12)$$

which can be written as

$$\beta' P \beta = c_1 \sum_{j=2}^n (\Delta^2 \beta_j)^2 + c_2 \sum_{j=3}^n \Delta^2 \beta_j \Delta^2 \beta_{j-1} \quad (13)$$

Given a set of equidistant knots, the constants c_1 and c_2 are given by

$$\begin{aligned} c_1 &= \int_0^1 B_{j,1}^2(l) dl \\ c_2 &= \int_0^1 B_{j,1}(l) B_{j-1,1}(l) dl. \end{aligned} \quad (14)$$

This gives us that the traditional smoothness penalty on the squared second derivative can be written as a linear combination of a penalty on the second-order differences of the B-spline coefficients 8 and the sum of the cross products of neighboring second differences. The second term in 13 leads to a complex objective function when minimizing the penalized likelihood, where seven adjacent spline coefficients occur, as opposed to five if only the first term in 13 is used in the penalty. The added complexity is a consequence of overlapping B-splines, which quickly increases when using higher order differences and higher order B-splines. We can seamlessly augment the likelihood with the difference penalty to achieve smooth fitted functions without the complexity posed by the derivative-based penalty.

The following figures display the impact of the squared distance on adjacent basis coefficients on the function itself. When examining the relationship between P-spline curves and their coefficients, it is helpful to consider the coefficients as the skeleton of the function, then draping the B-splines over them to “put the flesh on the bones.” A smoother sequence of coefficients leads to a smoother curve, which is clearly illustrated in Figure ???. As long as the coefficient sequence is smooth, the number of basis functions (and coefficients) is unimportant since the penalty ensures the smoothness of the skeleton.

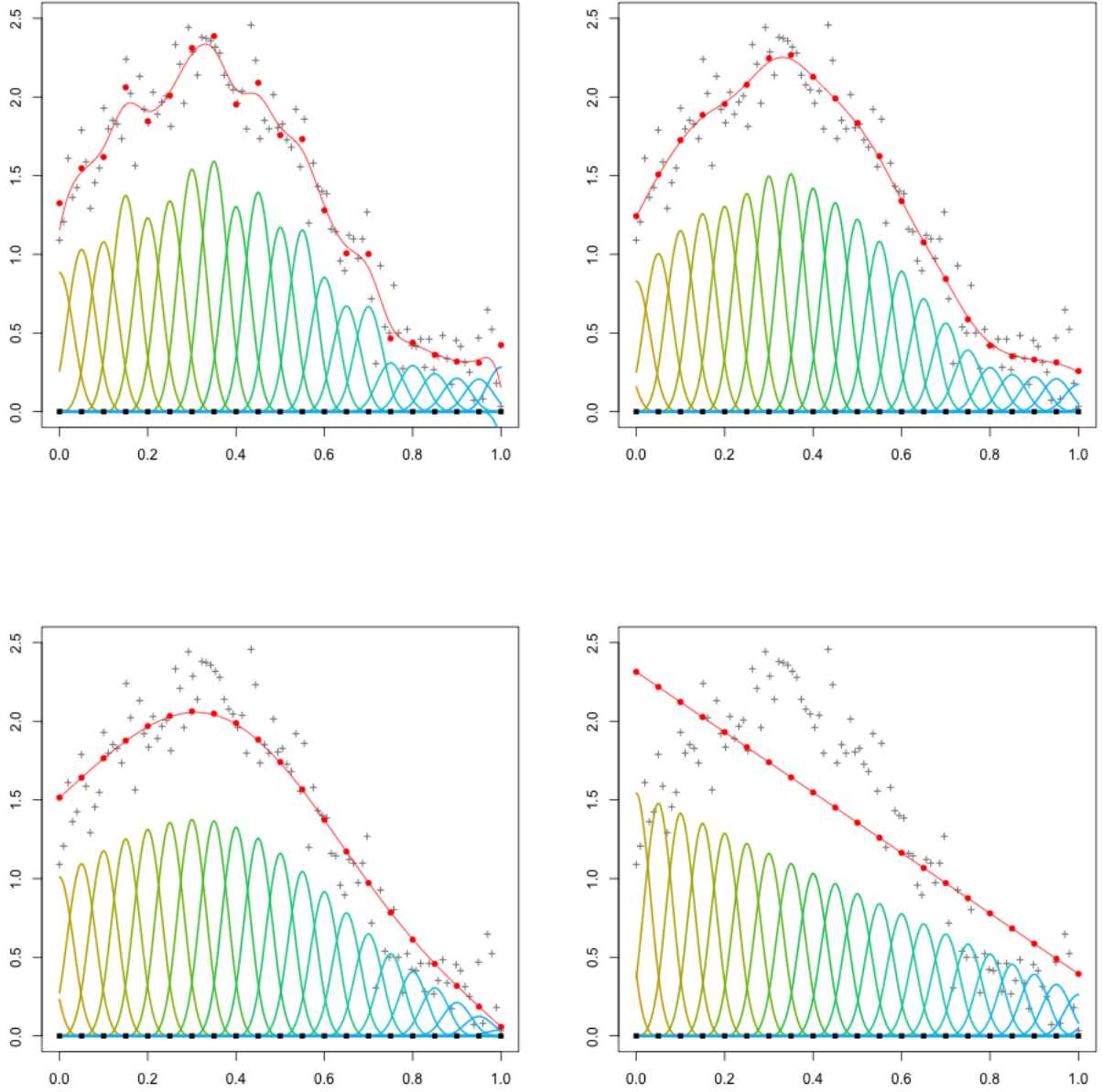


Figure 2: *Illustration of the impact of the second order difference penalty. The number of B-splines used is the same in each plot, with the value of the penalty parameter increasing from left to right and top to bottom across each plot. The fitted curve in the upper left plot is the most “wiggly” of any of the fits, as the penalty plays the weakest roll in the fitted coefficients there. The red circles are the values of each of the B-spline coefficients; as the penalty increases, they form as smoother sequence as we move across the four plots, which results in a smoother fitted function. As the penalty parameter approaches infinity, the fit approaches a linear function as shown in the bottom right plot.*

The number of B-splines can be much larger than the number of observations because penalty ensures that the fitting procedure is well-conditioned. Figure ?? illustrates this utility of the penalty for simulated data; there are $m = 10$ observations and 60 cubic B-splines. This property of P-splines cannot be overly appreciated because it frees us from the concern of choosing the optimal set of knots. Unless computational constraints are of concern, which is possible with large models, it is prudent to use even more B-splines. Figure ?? shows how the fitted function changes as the tuning parameter varies when the data are sparsely sampled. P-splines enjoy a number of additional advantageous properties, many of which are inherited from the attractive properties of B-splines. See ? for a detailed presentation.

5.2 Multidimensional smoothing with B-splines

P-splines can be extended naturally to higher dimensions by constructing a regression basis from the tensor product of the one-dimensional B-splines bases for each dimension. Figure 3 displays the building block of the foundation on which multidimensional P-splines is built: a B-spline tensor product basis $\{T_{kl}\}$ function. If we equip l and m each with a B-spline basis, we can a basis for the varying coefficient function ϕ in ?? by taking the tensor product of the two marginal bases. To regularize the fitted function, the only other modification necessary for the extension into two dimensions is the addition of a difference penalty for each variable l and m . Let $B_1(l), \dots, B_K(l)$ and $B_1(m), \dots, B_L(m)$ denote the B-spline bases for l and m , each having a set of equally spaced knots along their respective domain. It is worth noting that while we have chosen not to distinguish between $\{B_k\}$ and $\{B_l\}$ for the sake of brevity, one is free to specify a different basis for each dimension either by using different order B-spline or, of course, using different numbers of knots, and hence entirely different knot sequences since P-splines rely on bases with equally spaced knots. The tensor product basis functions

$$T_{jk}(l, m) = B_j(l) B_k(m)$$

carve the l - m domain into rectangles. Figure 4 shows a thinned tensor product basis $\{T_{kl}\}$; a portion of the basis was omitted to eliminate overlapping of the basis functions so that the reader can identify individual tensor products. Each “hill” in Figure 4 is associated with an unknown coefficient α_{ij} which determines the height of the hill. For a given knot grid, we can approximate a surface by

$$\phi^*(l, m) = \sum_{i=1}^K \sum_{j=1}^L \alpha_{ij} B_i(l) B_j(m). \quad (15)$$

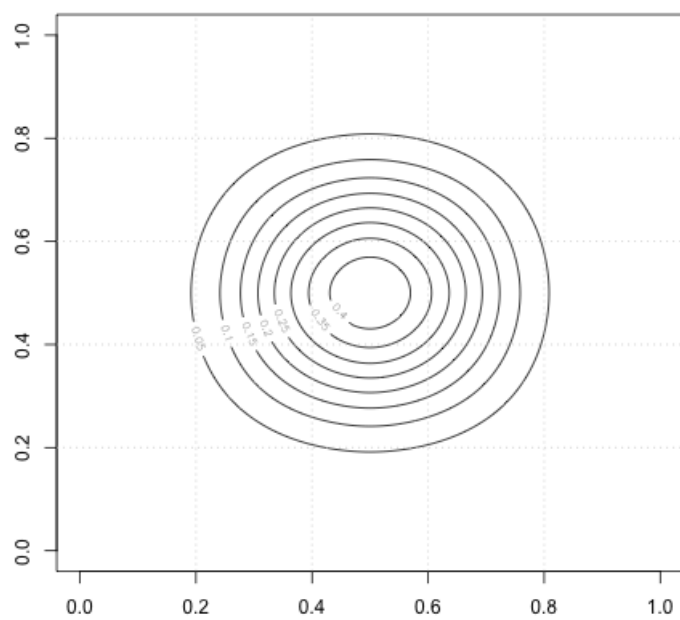
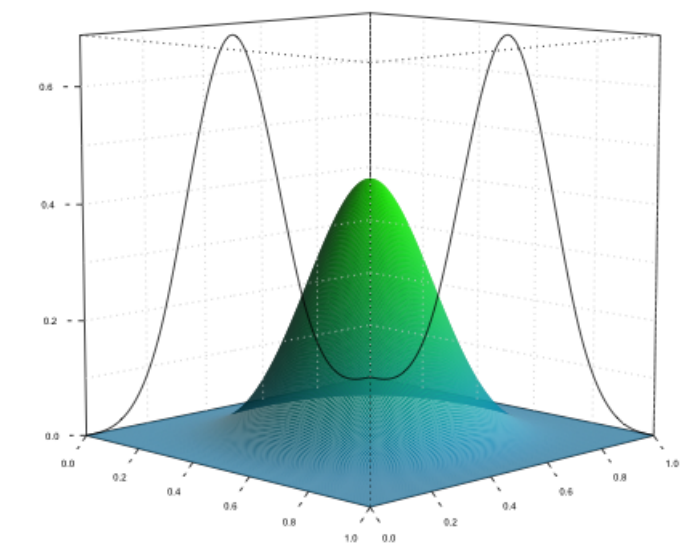


Figure 3: Tensor product of two cubic B-splines

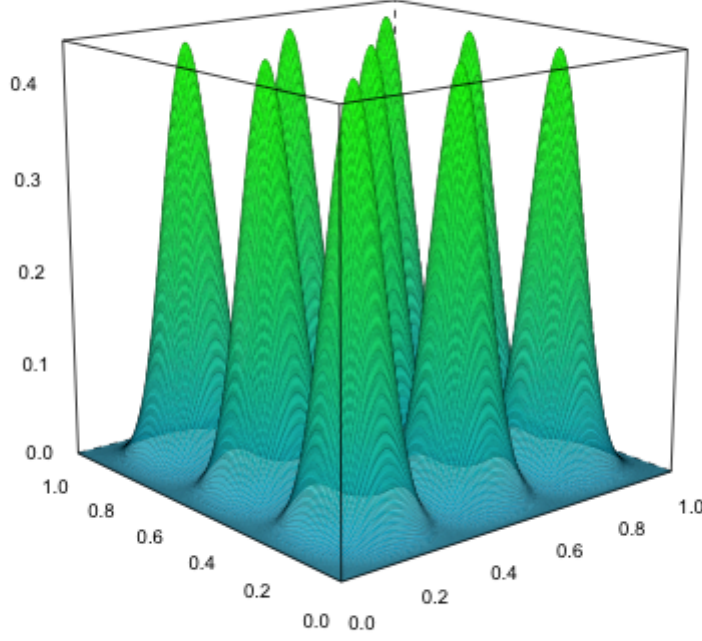


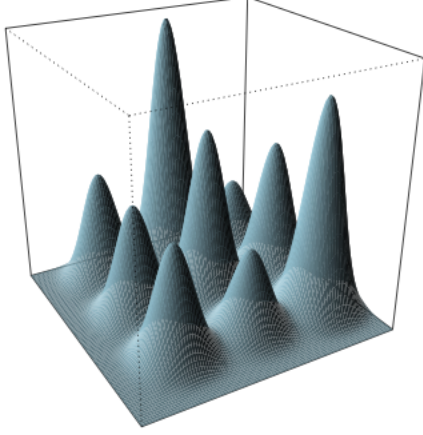
Figure 4: A subset of a full bivariate basis of cubic B-splines

Let p denote the total number of unique within-subject pairs of design points, and index the transformed coordinate pairs $(l, m)_i, i = 1, \dots, p$. Let B_l and B_m denote the B-spline bases along the l and m coordinates of dimension $p \times K$ and $p \times L$, respectively, and let A be the unknown matrix of basis coefficients with elements $[\alpha_{ij}]$. To control the flexibility of the fitted function as with univariate P-splines discussed in Section 5.1, we can apply difference penalties to the rows and columns of A to smooth in the l and m directions respectively, using the penalty

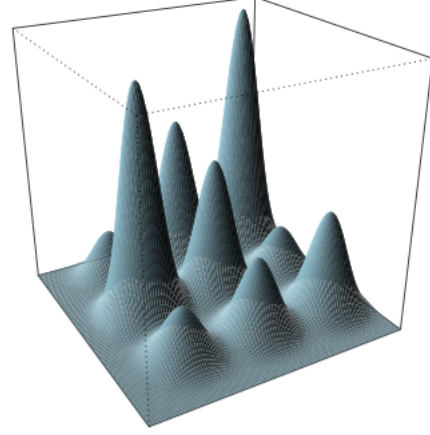
$$\lambda_l \sum_k |D_d \alpha_{k.}|^2 + \lambda_m \sum_l |D_{d_m} \alpha_{.l}|^2. \quad (16)$$

where $\alpha_{k.}$ and $\alpha_{.l}$ denote the k^{th} row and l^{th} column of A , respectively. The first term in 16 imposes a difference penalty of order d_l on the rows of the coefficient matrix while the second term places a difference penalty (of possible different order d_m) on the columns. We give each direction its own smoothing parameter to permit anisotropic smoothing; however, one could opt to use a single smoothing parameter for both directions and dodge the added work of optimizing the amount of smoothing with two separate parameters. Figure ?? shows a potential result of heavy column

penalization (left) and heavy row penalization (right) under a second order difference penalty on each row and each column for large values of λ_l and λ_m . The figure demonstrates that the limiting behaviour of each row and column is linear, but the resulting surface may exhibit slope reversals from one row (column) to the next.



(a) heavy column penalization



(b) heavy row penalization

Figure 5: Nine cubic B-spline tensor products with heavy linear column penalization and heavy linear row penalization

We take the estimator of ϕ^* to be the minimizer of

$$\begin{aligned}
 -2L + J(\phi^*) = & \sum_{i=1}^N \sum_{j=2}^{m_i} \sigma(t_j)^{-2} \left\{ y_{ij} - \sum_{k=1}^{j-1} \left(\sum_{r=1}^L \sum_{s=1}^K \alpha_{rs} B_r(l_{ijk}) B_s(m_{ijk}) \right) y_{ik} \right\}^2 \\
 & + \lambda_l \sum_{r'=1}^K |D_{d_l} \alpha_{r'}|^2 + \lambda_m \sum_{s'=1}^L |D_{d_m} \alpha_{s'}|^2.
 \end{aligned} \tag{17}$$

It is computationally advantageous to express the coefficient matrix in “unfolded” notation, which allows us to express the varying coefficient function at the observed coordinate grid as in the usual multiple regression form:

$$\text{vec} \{ \phi^*(l, m) \} = B\alpha$$

Stacking the columns of A gives the vectorized coefficient matrix $\alpha = \text{vec}(A)$. The $p \times KL$ tensor product basis B is constructed from the tensor product of the marginal B-spline bases defined in ? as the *row-wise Kronecker product* of the individual bases:

$$B = B_l \square B_m = (B_m \otimes 1'_K) \odot (1'_L \otimes B_l). \quad (18)$$

The operator \odot denotes the element-wise matrix product; 1_K (1_L) denotes the column vector of ones having length K (L .) The operations in 18 construct B such that the i^{th} row of $B_m \square B_l$ is the Kronecker product of the corresponding rows of B_m and B_l . The penalty in 16 can also be compactly expressed:

$$\lambda_l |P_l \alpha|^2 + \lambda_m |P_m \alpha|^2 \quad (19)$$

where $P_l = I_L \otimes D'_{d_l} D_{d_l}$ and $P_m = D'_{d_m} D_{d_m} \otimes I_K$. We define the matrix W of historical regressors so that 17 can be written in matrix form as

$$-2L + J(\phi^*) = |Y - WB\alpha|^2 + \lambda_l |P_l \alpha|^2 + \lambda_m |P_m \alpha|^2, \quad (20)$$

with $\hat{\alpha}$ solving the system of equations

$$[(WB)'WB + \lambda_l P_l + \lambda_m P_m] \alpha = WBY \quad (21)$$

From 21, we note that the system of equations depends on basis coefficients remains fixed at KL , even as the number of observations increases. The grid of regression coefficients can be recovered by arranging the elements of $\hat{\alpha}$ into a matrix of L columns having length K . The effective hat matrix is given by

$$H_\lambda = WB(WB'WB + \lambda_l P_l + \lambda_m P_m)^{-1}WB',$$

from which we can easily calculate deletion residuals and obtain a measure of effective model dimension. We will discuss diagnostics and approaches to model selection in detail in Section ??.

6 Recruit the difference penalty to regulate.

For $f(x) = \sum_{i=1}^p B_i(x) \theta_i$, approximate

$$\begin{aligned} \int_0^1 (f''(x))^2 dx &= \int_0^1 \left\{ \sum_{i=1}^p B_i''(x) \theta_i \right\}^2 dx \\ &= k_1 \sum_i (\Delta^2 \theta_i)^2 + k_2, \end{aligned}$$

by

$$\|D_2 \theta\|^2, \quad D_2 \theta = (\Delta^2 \theta_1, \dots, \Delta^2 \theta_{p-2})'$$

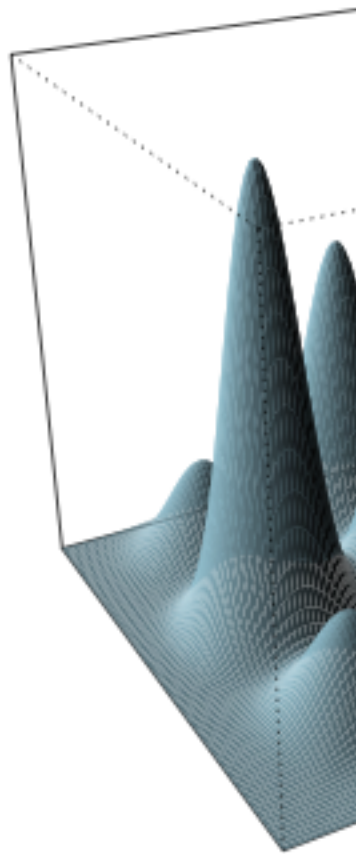
In general, approximate $\int_0^1 (f^{(d)})^2 dx$ with $\|D_d \theta\|^2$

7 *Append the difference penalties to the log likelihood:*

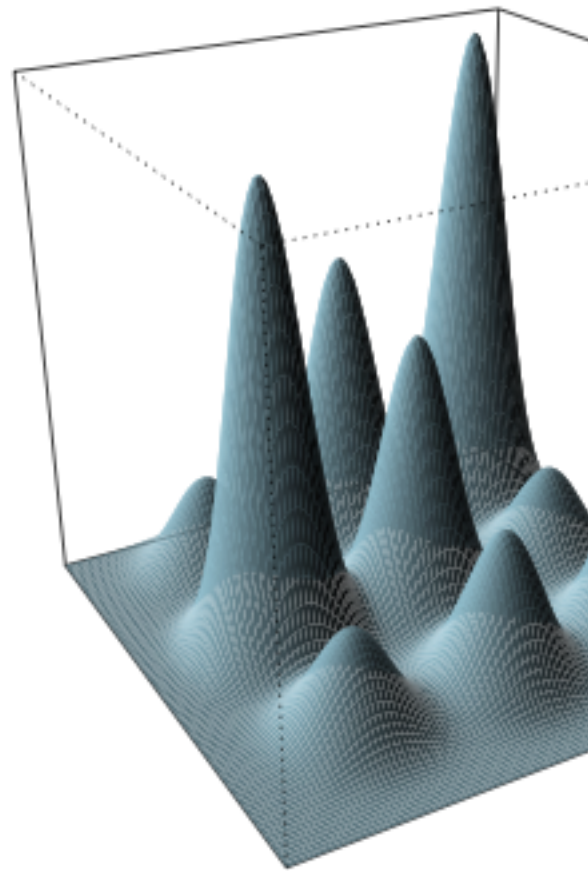
Find θ minimizing

$$\begin{aligned} \mathcal{Q} = \sum_{i=1}^n \sum_{j=2}^{m_i} \sigma_{ij}^{-2} & \left[y_{ij} - \sum_{k=1}^{j-1} \left(\sum_{i=1}^K \sum_{j=1}^L \theta_{ij} B_i(l_{ijk}) B_j(m_{ijk}) \right) y_{ik} \right]^2 \\ & + \lambda_l \sum_k |D_d \theta_{k\cdot}|^2 + \lambda_m \sum_l |D_{d_m} \theta_{\cdot l}|^2 \end{aligned}$$

8 *Differencing in two dimensions*



(a) Strong column penalty



(b) Strong row penalty

Figure 6: *Nine cubic B-spline tensor products with heavy linear column penalty (left) and heavy linear row penalty (right)*

9 “Unfold” the two dimensional surface:

via the kronecker product:

$$\begin{aligned} B &= B_l \square B_m \\ &= (B_m \otimes 1'_K) \odot (1'_L \otimes B_l) \\ P_l &= I_L \otimes D'_{d_l} D_{d_l} \\ P_m &= D'_{d_m} D_{d_m} \otimes I_K \end{aligned}$$

to write

$$\text{vec} \{ \phi^* (l, m) \} = B\theta$$

10 Then the penalized likelihood has convenient form:

$$Q = (Y - WB\theta)' D^{-1} (Y - WB\theta) + \lambda_l \theta' P_l \theta + \lambda_m \theta' P_m \theta$$

$$\hat{\theta}_\lambda = [(WB)' D^{-1} WB + \lambda_l P_l + \lambda_m P_m]^{-1} (WB)' Y$$

Degrees of freedom can be approximated as in the usual smoothing case:

$$\begin{aligned} \text{ED} &= \text{tr} [H_\lambda] \\ &= \text{tr} \left[[(WB)' D^{-1} WB + \lambda_l P_l + \lambda_m P_m]^{-1} (WB)' WB \right] \end{aligned}$$

AIC, GCV, CV accessible for model diagnostics.

10.1 Banding the Cholesky factor

A recent upsurge in the application of regularization to large empirical covariance matrices includes the work ?, ?, ?, ?. ? propose stabilizing the sample covariance matrix by “tapering,” or gradually shrinking the off-diagonal elements to zero. ? propose a sparse estimator by applying an L_1 penalty directly to the elements of the covariance matrix. Instead of regularizing the covariance matrix itself, some have opted to regularize its inverse; ? band the Cholesky decomposition by setting certain diagonals of the Cholesky factor to zero. ? and ? use L_1 penalties to achieve parsimony of the entries of the Cholesky factor; sparsity of the Cholesky factor, however, does not necessarily imply sparsity in the inverse covariance matrix. ? propose banding the Cholesky factor using a nested Lasso penalty which yields sparse estimators of the precision matrix.

? presented a heuristic argument that the GARPs, $\phi_{t,t-l}$ should be monotonically decreasing in absolute value as l increases. The following proposition establishes the relationship between zeros in the Cholesky factor and zeroes in the inverse covariance, connecting the regularization of ϕ^* to the structure in the resulting inverse covariance matrix.

Proposition 10.1. *Let Ω denote a $m \times m$ positive definite matrix with elements ω_{ij} with modified Cholesky decomposition $T'D^{-1}T$, where T is unit lower triangular. Let t_{ij} denote the ij^{th} element of T . For any column j and row $r(j) > j$, $\omega_{mj} = \dots = \omega_{r(j),j} = 0$ if and only if $t_{mj} = \dots = t_{r(j),j} = 0$.*

The proof is left to the appendix. Proposition 10.1 maintains that the modified Cholesky factor T with arbitrary column band lengths corresponds to inverse covariance matrix Ω with the same column band lengths, and hence the inverse covariance matrix is K -banded if and only if its Cholesky factor is K -banded. That is, if ϕ^* is zero or nearly zero for large $|i - j|$, then y_i and y_j are conditionally uncorrelated (or nearly so). This suggests that the effect of y_{t-l} on y_t through model 4 should decrease as the time between the two measurements increases, such that

$$\phi^*(l, m) \approx 0$$

for large l is a reasonable way to institute regularization.

This notion of sparsity is quite different from the notion of smoothness imposed by the usual derivative-based penalties or the analog difference penalties for P-splines; however, the local support of the B-spline basis functions permits the intuitive expression of a penalty to be used for banding the Cholesky factor. The B-spline basis functions are non-negative on their support; therefore, if a B-spline (as in 7) is zero for $l > l_0$, then the coefficients of the B-splines contributing to that region of the domain are also zero. For functions represented using order k B-spline basis functions $\{B_i\}$, $i = 1, \dots, n$, one may truncate the function to zero at some truncation point $l_0 \in (0, 1)$ by penalizing the size of the coefficients corresponding to any basis functions having support on $l > l_0$. This naturally leads to penalties of the form

$$J_B = \sum_{i=i^*-k}^n |\beta_i|^p \quad (22)$$

where

$$i^* = \max_i \{\ell_i : \ell_i \leq l_0\}$$

is the index of the largest knot which is smaller than the truncation point l_0 . Setting $p = 1$ in 22 corresponds to putting a LASSO penalty (see ?) on the coefficients contributing to the function value on $l > l_0$, while setting $p = 2$ corresponds to the usual ridge regression setting.

Several have proposed regularization of the inverse covariance matrix by “banding” the Cholesky factor T : setting all elements of T beyond the K^{th} off-diagonal to zero, i.e. setting $\phi_{t,t-l} = 0$ for $l > K$ for some choice of K . (See ?, ?, ?, and ?.) In terms of model 1, this is equivalent to regressing y_t on only its K immediate predecessors, setting the regression coefficient for y_{t-l} to zero for $l > K$. This notion is instrumental in justifying a family of penalties which induces an alternative decomposition of the function space for which the null space of these penalties comprises functions ϕ^* taking nonzero values on a subset of the domain:

$$\mathcal{H}_0(\mathcal{B}) = \{\phi^* : \phi^*(l, m) = 0 \text{ for all } l > l_0\}, \quad (23)$$

which is equivalent to the set of functions having compact support:

$$\text{supp}(\phi^*) = (0, l_0] .$$

11 Simulations: $\Sigma = 0.3^2 I$
 $N = 30, M = 20, d_l = d_m = 0$

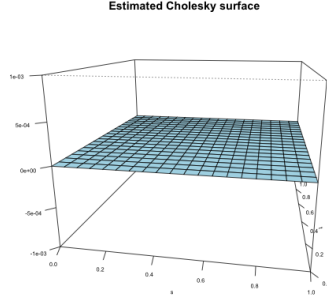
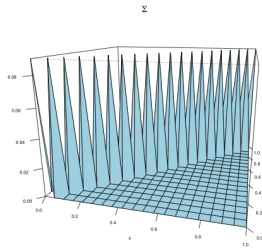
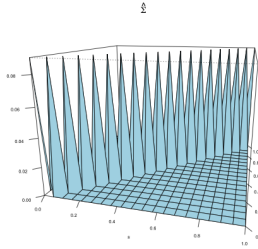


Figure 7: Estimated T

12 Simulations: $\Sigma = 0.3^2 I$
 $N = 30, M = 20, d_l = d_m = 0$

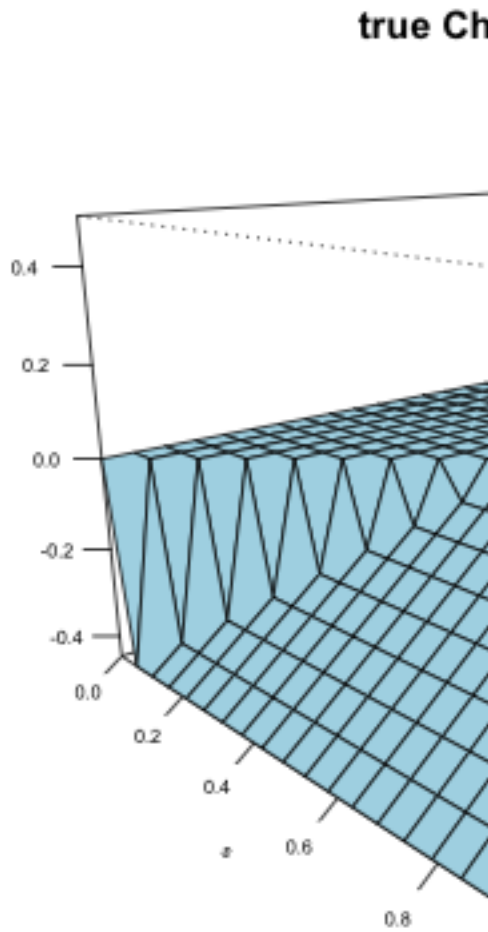


(a) True Σ

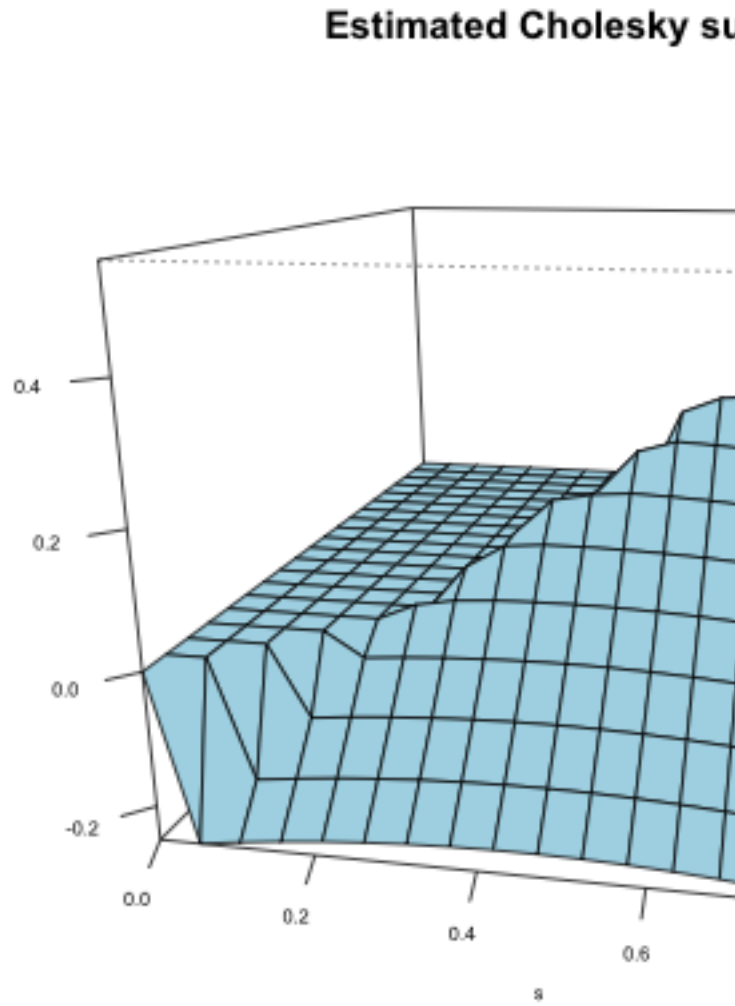


(b) $\hat{\Omega}^{-1}$

13 Simulations: $\phi(s, t) = s - \frac{1}{2}, \sigma^2 = 0.3^2$
 $N = 30, M = 20, d_l = d_m = 2$

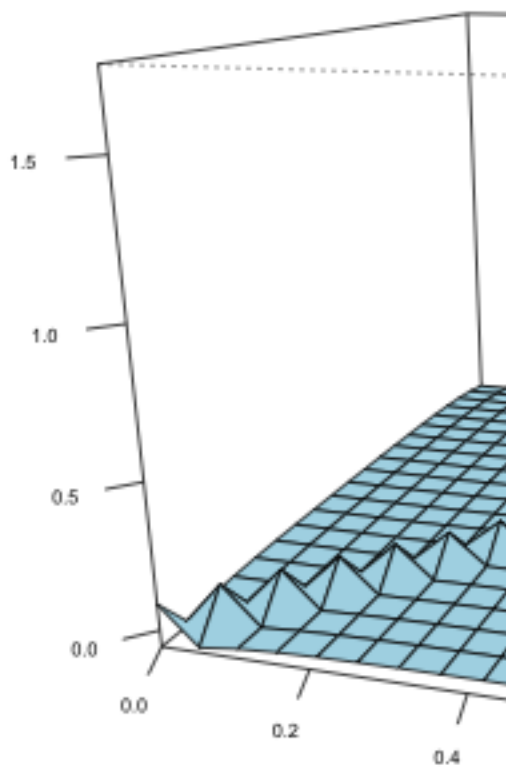


(a) True T

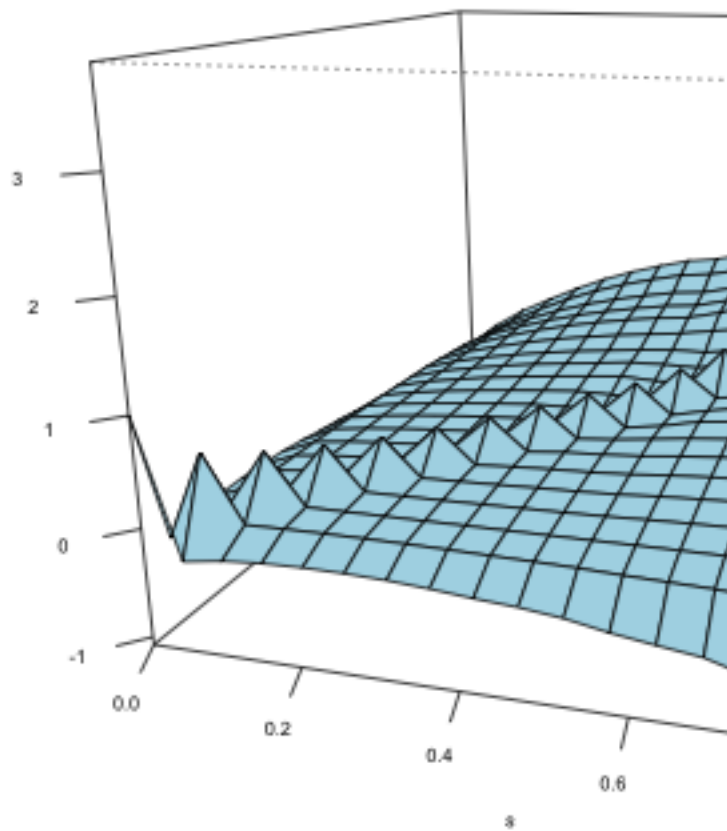


(b) Estimated T

14 Simulations: $\phi(s, t) = s - \frac{1}{2}, \sigma^2 = 0.3^2$
 $N = 30, M = 20, d_l = d_m = 2$

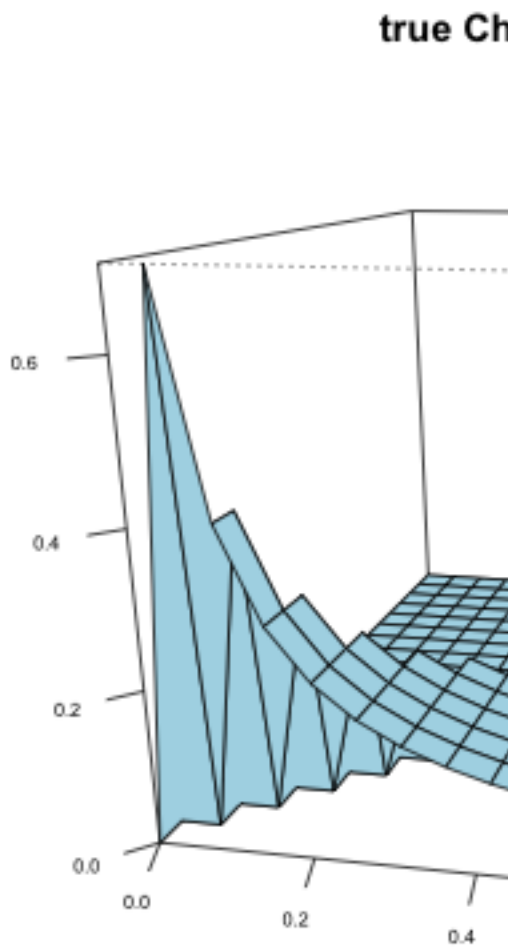


(a) Σ

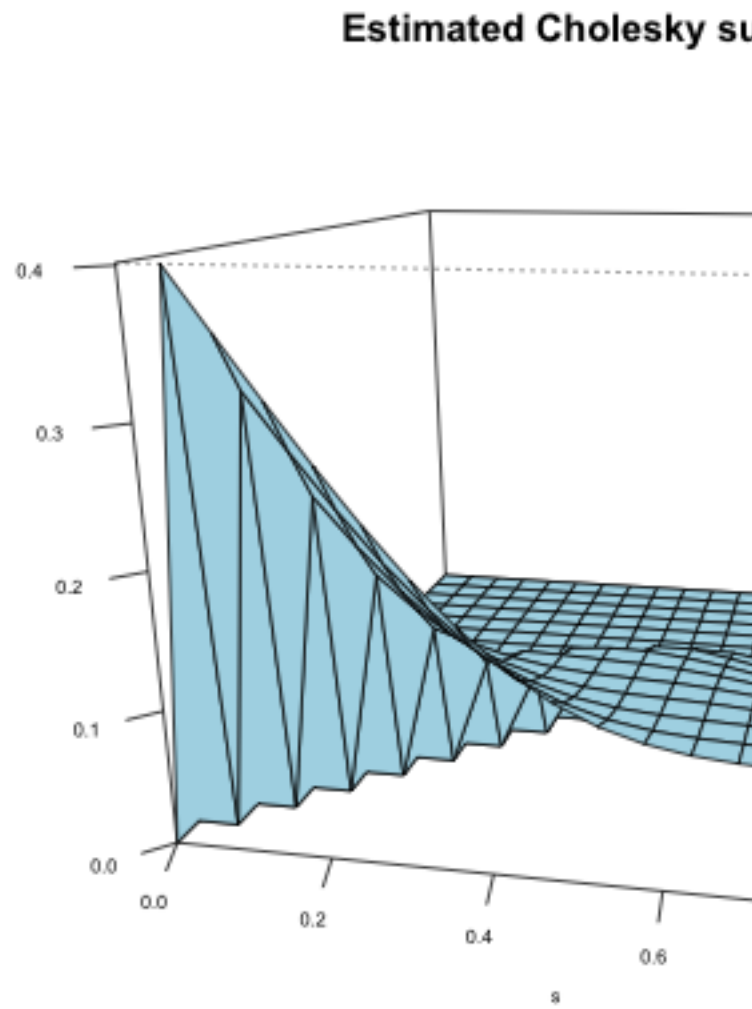


(b) $\hat{\Omega}^{-1}$

15 Simulations: $\Sigma = 0.7J + 0.3I$
 $N = 30, M = 20, d_l = 2, d_m = 1$

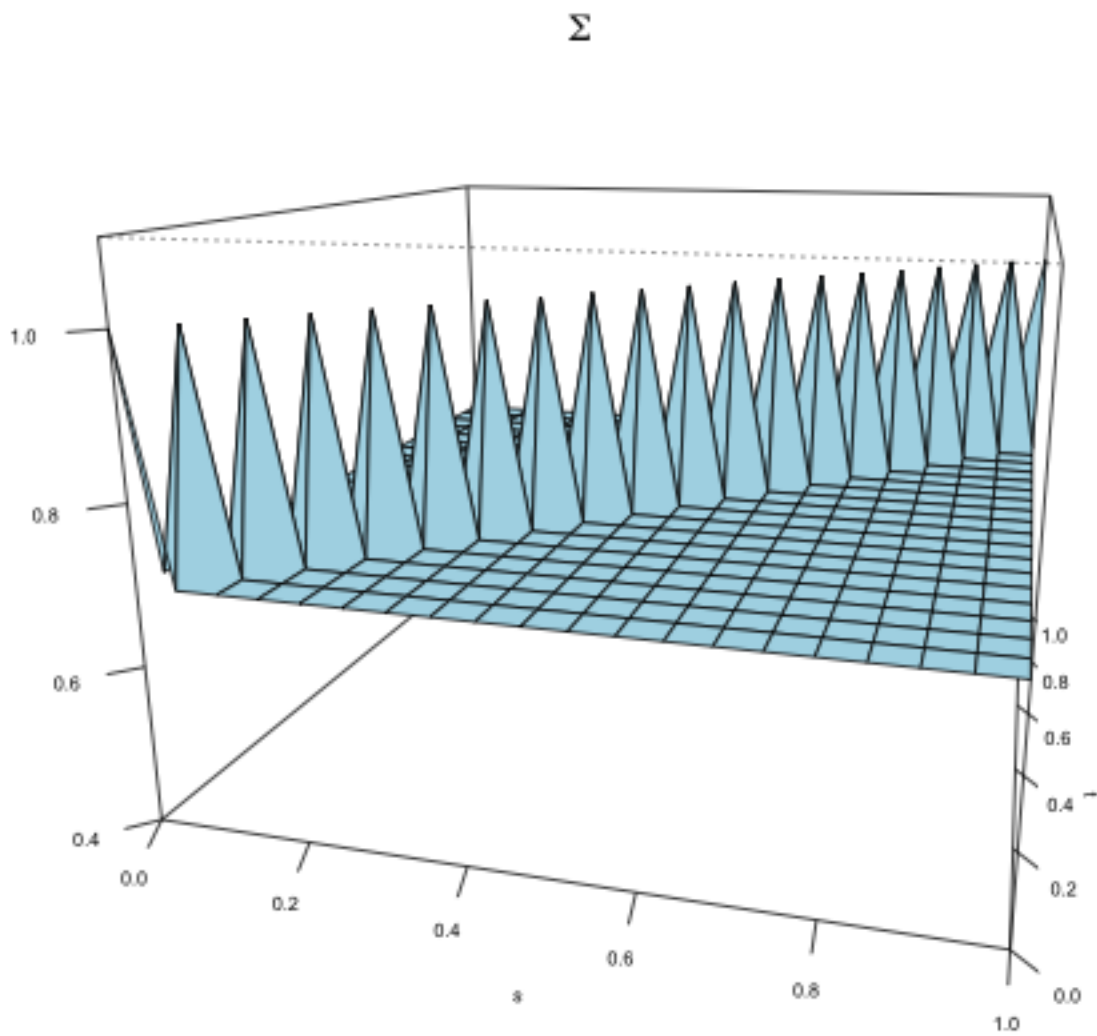


(a) True T



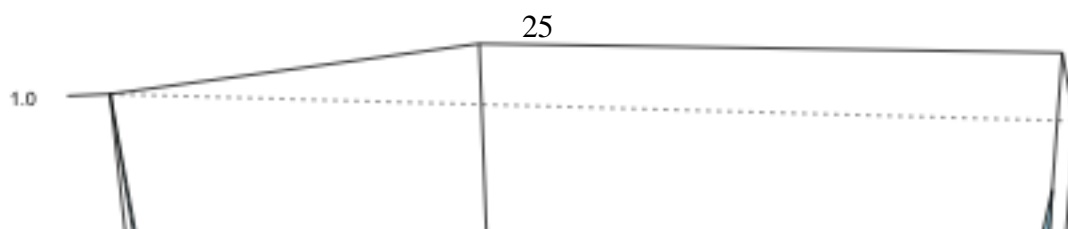
(b) Estimated T

16 Simulations: $\Sigma = 0.7J + 0.3I$
 $N = 30, M = 20, d_l = 2, d_m = 1$



(a) Σ

$\hat{\Sigma}$



17 *Recap: from the top*

- We propose a general framework for unconstrained covariance estimation.
- Flexibility permits imposing various types of regularization with ease.
- Penalty specification is crucial for performance.

18 **What's next?**

- “designer” penalties - impose desirable shape constraints (decay in l)
- Additive, ANOVA models for ϕ^*
- P-spline model reparameterized as mixture models