GCVPACK – ROUTINES FOR GENERALIZED CROSS VALIDATION

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Purpose and Description

Purpose

These Fortran-77 subroutines provide building blocks for Generalized Cross-Validation (GCV) (Craven and Wahba, 1979) calculations in data analysis and data smoothing including ridge regression (Golub, Heath, and Wahba, 1979), thin plate smoothing splines (Wahba and Wendelberger, 1980), deconvolution (Wahba, 1982d), smoothing of generalized linear models (O'Sullivan, Yandell and Raynor (1986), Green (1984) and Green and Yandell (1985)), and ill-posed problems (Nychka et al., 1984, O'Sullivan and Wahba, 1985). We present some of the types of problems for which GCV is a useful method of choosing a smoothing or regularization parameter and we describe the structure of the subroutines.

Ridge Regression: A familiar example of a smoothing parameter is the ridge parameter λ in the ridge regression problem which we write as

$$\min_{\mathbf{y}} \quad \frac{1}{n} \| \mathbf{y} - \mathbf{X} \mathbf{y} \|^2 + \lambda \gamma^T \mathbf{y}$$

where γ is a *p*-dimensional parameter vector, **y** is an n-dimensional response vector and **X** is an $n \times p$ design matrix.

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For any positive λ , an optimal γ_{λ} can be easily calculated. Unfortunately, this leaves the question of which value of λ to use. Golub, Heath, and Wahba (1979) demonstrated that minimization of the GCV function $V(\lambda)$ is a powerful criterion for the choice of an optimal λ , where

$$V(\lambda) = \frac{(1/n) \| (\mathbf{I} - \mathbf{A}(\lambda))\mathbf{y} \|^2}{[(1/n) \operatorname{tr}(\mathbf{I} - \mathbf{A}(\lambda))]^2}$$

and $A(\lambda)$ is the $n \times n$ "hat" matrix of the ridge regression

$$\mathbf{A}(\lambda) = \mathbf{X}(\mathbf{X}^{\mathrm{T}}\mathbf{X} + n\,\lambda\mathbf{I})^{-1}\mathbf{X}^{\mathrm{T}}$$

At first glance, optimization of $V(\lambda)$ seems a formidable computational problem since each value of λ has its corresponding $A(\lambda)$. However, Golub, Heath, and Wahba (1979) gave a method of expressing $V(\lambda)$ as an easily-calculated rational function based on the singular value decomposition (SVD) (Dongarra et al., 1979, chapter 10)

$$\mathbf{X} = \mathbf{U}\mathbf{D}\mathbf{V}^{\mathrm{T}}$$

where **U** is $n \times p$ with orthonormal columns, **V** is $p \times p$ and orthogonal, and **D** is $p \times p$ and diagonal with diagonal elements

$$d_1 \ge d_2 \ge \cdots \ge d_p \ge 0$$

which are the nonnegative square roots of the eigenvalues of $\mathbf{X}^{T}\mathbf{X}$. The "hat" matrix can then be written as

$$\mathbf{A}(\lambda) = \mathbf{U}\mathbf{D}^2(\mathbf{D}^2 + n\,\lambda\mathbf{I})^{-1}\mathbf{U}^{\mathrm{T}} ,$$

and using

$$\mathbf{z} = \mathbf{U}^{\mathrm{T}}\mathbf{y}$$

we can write

$$V(\lambda) = \frac{n \left[\left\| \mathbf{y} \right\|^{2} - \left\| \mathbf{z} \right\|^{2} + \sum_{j=1}^{p} \left[\frac{n\lambda}{d_{j}^{2} + n\lambda} \right]^{2} z_{j}^{2} \right]}{\left[n - p + \sum_{j=1}^{p} \frac{n\lambda}{d_{j}^{2} + n\lambda} \right]^{2}} .$$
(1.1)

Once the SVD of X is computed, it is trivial to evaluate $V(\lambda)$ for a wide range of values of λ and determine the optimum value of λ . Equation (1.1) indicates that, for most problems, $d_{\rho}^2 \le n\hat{\lambda} \le d_1^2$. After an optimal λ is chosen, the corresponding γ_{λ} is calculated as

$$\boldsymbol{\gamma}_{\lambda} = \mathbf{V} (\mathbf{D}^2 + n\,\lambda \mathbf{I})^{-1} \mathbf{D} \mathbf{z} \quad . \tag{1.2}$$

Multivariate data smoothing with thin-plate splines: A more important application of GCV is determining smooth representations of an underlying multivariate function from which noisy data is observed. The ridge regression problem serves as an introduction to the idea of GCV

and the computational steps for efficient evaluation of the GCV function but data smoothing using thin-plate smoothing splines (TPSS) is a much more common application of GCV. These methods extend the computational methods derived in Wahba and Wendelberger (1980), Wendelberger (1981), and Wahba (1984a).

For convenience we first describe the calculations for a two-dimensional "independent" variable \mathbf{x} but the software is designed for the general case. The data model for TPSS is

$$y_i = f(\mathbf{x}_i) + \varepsilon_i$$
, $i = 1, \ldots, n$

where the (\mathbf{x}_i, y_i) , i = 1, 2, ..., n, are observed data, f is an unknown function which is assumed to be reasonably smooth, and the ε_i , i = 1, 2, ..., n, are independent, zero-mean random variables.

In general we will measure smoothness of f by the integral over the entire plane of the square of the partial derivatives of f of total order 2. That is,

$$J_2(f) = \int_{-\infty}^{\infty} \int_{-\infty}^{\infty} \left[\left[\frac{\partial^2 f}{\partial x_1^2} \right]^2 + 2 \left[\frac{\partial^2 f}{\partial x_1 \partial x_2} \right]^2 + \left[\frac{\partial^2 f}{\partial x_2^2} \right]^2 \right] dx_1 dx_2 \quad .$$

To allow generalizations, the software uses a smoothness penalty defined by the partial derivatives of total order m as

$$J_m(f) = \int_{-\infty}^{\infty} \int_{i=0}^{\infty} \sum_{j=0}^{m} {m \brack i} \left[\frac{\partial^m f(x_1, x_2)}{\partial x_1^m \partial x_2^{m-i}} \right]^2 dx_1 dx_2$$

In d dimensions,

$$J_m(f) = \underbrace{\int_{-\infty}^{\infty} \cdots \int_{-\infty}^{\infty} \sum \frac{m!}{\alpha_1! \cdots \alpha_d!} \left[\frac{\partial^m f}{\partial x_1^{\alpha_1} \cdots \partial x_2^{\alpha_d}} \right]^2 dx_1 \cdots dx_d$$

with the sum within the integral over $\sum \alpha_i = m$. In general, one must have 2m - d > 0 with d the dimension of **x**. Using this smoothness penalty, the TPSS estimate f_{λ} of f is the minimizer of

$$S_{\lambda}(f) = \frac{1}{n} \sum_{i=1}^{n} (y_i - f(\mathbf{x}_i))^2 + \lambda J_m(f) \quad .$$
(2.1)

From Duchon (1976), the minimizer f_{λ} of (2.1) can be represented as

$$f_{\lambda}(\mathbf{x}) = \sum_{i=1}^{t} \beta_{i} \phi_{i}(\mathbf{x}) + \sum_{i=1}^{n} \delta_{i} E_{m}(\mathbf{x} - \mathbf{x}_{i})$$
(2.2)

where

$$E_m(\mathbf{t}) = (-1)^m \, 2^{1-2m} \, \pi^{-1} \, ((m-1)!)^{-2} \, \| \mathbf{t} \, \|^{(2m-2)} \ln(\| \mathbf{t} \, \|)$$

and t is the dimension of the space of polynomials on two variables of total order at most m-1,

 $t = \begin{pmatrix} m+1\\2 \end{pmatrix} .$

A basis for this space is

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 $\varphi_1(\mathbf{x}) = 1$ $\varphi_2(\mathbf{x}) = x_1$ $\varphi_3(\mathbf{x}) = x_2$ $\varphi_4(\mathbf{x}) = x_1^2$ $\varphi_5(\mathbf{x}) = x_1 x_2$ \dots $\varphi_t(\mathbf{x}) = x_2^{m-1}$

The general definition of E_m , which depends on the dimension, d, of the independent variables **x** is

$$E_{m}(\mathbf{t}) = \begin{cases} (-1)^{1+m+d/2} 2^{1-2m} \pi^{-d/2} ((m-1)! (m-d/2)!)^{-1} \|\mathbf{t}\|^{(2m-d)} \ln(\|\mathbf{t}\|), d \text{ even} \\ \Gamma(\frac{d}{2}-m) 2^{-2m} \pi^{-d/2} ((m-1)!)^{-1} \|\mathbf{t}\|^{(2m-d)}, d \text{ odd} \end{cases}$$
(2.3)

The dimension, t, of the polynomial space is given in general by

$$t = \begin{bmatrix} m+d-1 \\ d \end{bmatrix} .$$

A property of the TPSS representation is that both the function f_{λ} evaluated at the data points and the smoothing penalty J_m can be expressed using the $n \times n$ matrix **K** with entries

$$\{\mathbf{K}\}_{ij} = E_m(\mathbf{x}_i - \mathbf{x}_j) \quad . \tag{2.4}$$

The function f_{λ} also requires the $n \times t$ matrix **T** with entries

$$\{\mathbf{T}\}_{ij} = \phi_j(\mathbf{x}_i) \ .$$

The matrix **T** having full column rank guarantees a unique minimizer of (2.1). Duchon (1976) showed that δ in (2.2) must satisfy

$$\mathbf{T}^{\mathrm{T}}\boldsymbol{\delta} = \mathbf{0} \quad . \tag{2.5}$$

Then β_{λ} and δ_{λ} are the minimizers of (2.1), which can be written as

$$S_{\lambda}(\beta, \delta) = \frac{1}{n} \| \mathbf{y} - \mathbf{T}\beta - \mathbf{K}\delta \|^{2} + \lambda \delta^{\mathrm{T}}\mathbf{K}\delta$$
.

Note that the restriction in (2.5) is important, as **K** will generally have negative eigenvalues, but for any vectors $\boldsymbol{\delta}$ satisfying (2.5) it can be shown that

 $\boldsymbol{\delta}^{\mathrm{T}} \mathbf{K} \boldsymbol{\delta} \ge 0$.

Our objective is to reduce the calculation of the parameters β_{λ} and δ_{λ} , the "hat" matrix $A(\lambda)$, and the GCV function $V(\lambda)$ to a simplified form as was done for the ridge regression case.

No replicates case: When there are no replicates in the x's, we proceed by taking a QR decomposition (Dongarra et al., 1979, chapter 9) of **T** as

$$\mathbf{T} = \mathbf{F}\mathbf{G} = [\mathbf{F}_1:\mathbf{F}_2]\begin{bmatrix}\mathbf{G}_1\\\mathbf{0}\end{bmatrix} = \mathbf{F}_1\mathbf{G}_1$$

where **F** is $n \times n$ and orthogonal while **G** is $n \times t$ and zero below the main diagonal. **F**₁ is the first *t* columns of **F** and **F**₂ is the trailing n - t columns while **G**₁ is the first *t* rows of **G**. The columns of **F**₂ provide a basis for the δ which satisfy

 $\mathbf{T}^{\mathrm{T}} \mathbf{\delta} = \mathbf{0}$

so we can set

$$\delta = \mathbf{F}_2 \boldsymbol{\zeta}$$

where ζ has dimension n-t. Using

$$\mathbf{w}_1 = \mathbf{F}_1^{\mathrm{T}} \mathbf{y}$$
$$\mathbf{w}_2 = \mathbf{F}_2^{\mathrm{T}} \mathbf{y}$$

the objective function of the optimization becomes

$$S_{\lambda}(\boldsymbol{\beta}, \boldsymbol{\delta}) = \frac{1}{n} \| \mathbf{y} - \mathbf{T}\boldsymbol{\beta} - \mathbf{K}\boldsymbol{\delta} \|^{2} + \lambda \boldsymbol{\delta}^{\mathrm{T}} \mathbf{K} \boldsymbol{\delta}$$

$$= \frac{1}{n} \| \mathbf{F}^{\mathrm{T}}(\mathbf{y} - \mathbf{T}\boldsymbol{\beta} - \mathbf{K}\boldsymbol{\delta}) \|^{2} + \lambda \boldsymbol{\delta}^{\mathrm{T}} \mathbf{K} \boldsymbol{\delta}$$

$$= \frac{1}{n} \| \mathbf{w}_{1} - \mathbf{G}_{1} \boldsymbol{\beta} - \mathbf{F}_{1}^{\mathrm{T}} \mathbf{K} \mathbf{F}_{2} \boldsymbol{\zeta} \|^{2} + \frac{1}{n} \| \mathbf{w}_{2} - \mathbf{F}_{2}^{\mathrm{T}} \mathbf{K} \mathbf{F}_{2} \boldsymbol{\zeta} \|^{2} + \lambda \boldsymbol{\zeta}^{\mathrm{T}} \mathbf{F}_{2}^{\mathrm{T}} \mathbf{K} \mathbf{F}_{2} \boldsymbol{\zeta} .$$

(2.6)

Assuming G_1 is non-singular (that is, the points x_i , i = 1, ..., n, are adequately dispersed so that the columns of **T** are linearly independent) the first term in (2.6) can be made zero by solving

$$\mathbf{G}_1 \boldsymbol{\beta}_{\lambda} = \mathbf{w}_1 - \mathbf{F}_1^{\mathrm{T}} \mathbf{K} \mathbf{F}_2 \boldsymbol{\zeta}_{\lambda}$$
$$= \mathbf{w}_1 - \mathbf{F}_1^{\mathrm{T}} \mathbf{K} \boldsymbol{\delta}_{\lambda}$$

for β_{λ} . In practice we check the condition of G_1 and return an error condition if it is computationally singular, indicating that the columns of **T** are strongly correlated. This condition is equivalent to the computational singularity of the problem of least squares regression of the data onto the span $\{\phi_j\}$. Singularity will rarely occur since the column dimension of **T** is small.

We can now reduce the problem to a form like ridge regression by using the fact that $\mathbf{F}_2^T \mathbf{K} \mathbf{F}_2$ is positive definite to form the Cholesky decomposition (Dongarra et al., 1979, chapter 8)

$\mathbf{F}_2^{\mathrm{T}}\mathbf{K}\mathbf{F}_2 = \mathbf{L}^{\mathrm{T}}\mathbf{L}$

where L is $(n-t) \times (n-t)$ and upper triangular. In practice we use a pivoted Cholesky decomposition so we can check the conditioning of $\mathbf{F}_2^T \mathbf{K} \mathbf{F}_2$. If this matrix is computationally singular, which

can occur if $||\mathbf{x}_i - \mathbf{x}_j||$ is very small but non-zero for some $i \neq j$, we return an error condition. A near-singular $\mathbf{F}_2^T \mathbf{K} \mathbf{F}_2$ is usually avoided since, in checking for replicates, we declare \mathbf{x}_i and \mathbf{x}_j to be replicates if the distance between them is very small. See Appendix 1 for more information on the detection of replicates and the computational singularity of **L**.

After ensuring that L is non-singular, we define

and the last two terms of $S_{\lambda}(\beta, \delta)$ in (2.6) can be written as

$$\frac{1}{n} \left\| \mathbf{w}_2 - \mathbf{L}^{\mathrm{T}} \boldsymbol{\gamma} \right\|^2 + \lambda \boldsymbol{\gamma}^{\mathrm{T}} \boldsymbol{\gamma} \ .$$

This has the same form as the ridge regression problem with solution

$$\gamma_{\lambda} = (\mathbf{L}\mathbf{L}^{\mathrm{T}} + n\,\lambda\mathbf{I})^{-1}\mathbf{L}\mathbf{w}_{2}$$
.

We take a SVD of \mathbf{L}^{T} as

$$\mathbf{L}^{\mathrm{T}} = \mathbf{U}\mathbf{D}\mathbf{V}^{\mathrm{T}}$$

and write the estimate as

$$\boldsymbol{\gamma}_{\lambda} = \mathbf{V}(\mathbf{D}^2 + n\,\lambda\mathbf{I})^{-1}\mathbf{D}\mathbf{U}^{\mathrm{T}}\mathbf{w}_2$$

and the "hat" matrix as

$$\mathbf{A}(\lambda) = \mathbf{F}_{1}\mathbf{F}_{1}^{\mathrm{T}} + \mathbf{F}_{2}\mathbf{U}\mathbf{D}^{2}(\mathbf{D}^{2} + n\lambda\mathbf{I})^{-1}\mathbf{U}^{\mathrm{T}}\mathbf{F}_{2}^{\mathrm{T}}$$
$$= \mathbf{F}\begin{bmatrix}\mathbf{I} & \mathbf{0}\\ \mathbf{0} & \mathbf{U}\end{bmatrix}\begin{bmatrix}\mathbf{I} & \mathbf{0}\\ \mathbf{0} & \mathbf{D}^{2}(\mathbf{D}^{2} + n\lambda\mathbf{I})^{-1}\end{bmatrix}\begin{bmatrix}\mathbf{I} & \mathbf{0}\\ \mathbf{0} & \mathbf{U}^{\mathrm{T}}\end{bmatrix}\mathbf{F}^{\mathrm{T}} .$$
(2.7)

As in the ridge regression case, we use

$$\mathbf{z} = \mathbf{U}^{\mathrm{T}}\mathbf{w}_{2}$$

to write

$$V(\lambda) = \frac{n \sum_{j=1}^{n-t} \left(\frac{n\lambda}{d_j^2 + n\lambda}\right)^2 z_j^2}{\left[\sum_{j=1}^{n-t} \frac{n\lambda}{d_j^2 + n\lambda}\right]^2} .$$
(2.8)

The actual calculation of the parameter δ_{λ} corresponding to the E_m 's is performed as

$$\delta_{\lambda} = \mathbf{F}_{2} \mathbf{U} (\mathbf{D}^{2} + n \lambda \mathbf{I})^{-1} \mathbf{z}$$

= $\mathbf{F}_{2} \mathbf{U} (\mathbf{D}^{2} + n \lambda \mathbf{I})^{-1} \mathbf{U}^{\mathrm{T}} \mathbf{F}_{2}^{\mathrm{T}} \mathbf{y} .$ (2.9)

Replicated x values: Replicates of x values introduce some minor complications since we must define only one δ_i corresponding to each unique x position. The best way to handle this is to

pre-process the data by sorting the **x** values to determine the unique **x** values and the number of replicates of each value. Let k be the number of unique **x** values. We can express the objective function optimized by δ_{λ} and β_{λ} as

$$S_{\lambda}(\boldsymbol{\beta},\boldsymbol{\delta}) = \frac{1}{n} \| \mathbf{y} - \mathbf{T}\boldsymbol{\beta} - \mathbf{K}\boldsymbol{\delta} \|^{2} + \lambda \boldsymbol{\delta}^{\mathrm{T}}\mathbf{K}_{U}\boldsymbol{\delta}$$
(3.1)

subject to the condition

 $\mathbf{T}_{II}^{\mathrm{T}} \mathbf{\delta} = \mathbf{0}$

where **T** and **K** are of size $n \times t$ and $n \times k$ respectively, while **T**_U and **K**_U are of size $k \times t$ and $k \times k$ respectively. These matrices are related by

$$T = MT_U$$
$$K = MK_U$$

where **M** is an $n \times k$ indicator matrix (all its entries are ones or zeros and there is only a single one in each row) which, for each row, indicates the unique **x** that corresponds to that observation.

If we take a QR decomposition of M as

$$\mathbf{M} = \mathbf{B}\mathbf{C} = [\mathbf{B}_1:\mathbf{B}_2]\begin{bmatrix}\mathbf{C}_1\\\mathbf{0}\end{bmatrix} = \mathbf{B}_1\mathbf{C}_1$$

and pre-multiply all the vectors in the first term of (3.1) by B, (3.1) divides into

$$S_{\lambda}(\boldsymbol{\beta},\boldsymbol{\delta}) = \frac{1}{n} \| \mathbf{B}_{2}^{\mathsf{T}} \mathbf{y} \|^{2} + \frac{1}{n} \| \mathbf{B}_{1}^{\mathsf{T}} \mathbf{y} - \mathbf{C}_{1} \mathbf{T}_{U} \boldsymbol{\beta} - \mathbf{C}_{1} \mathbf{K}_{U} \boldsymbol{\delta} \|^{2} + \lambda \boldsymbol{\delta}^{\mathsf{T}} \mathbf{K}_{U} \boldsymbol{\delta} .$$
(3.2)

In practice, it is not necessary to explicitly form **M** and take its QR decomposition since C_1 is diagonal with c_{ii} being the square root of the number of replicates of the *i*'th unique **x**. The elements of the vector $C_1^{-1}B_1^Ty$ are the means of the y's at the corresponding unique **x**'s. Further, $||B_2^Ty||^2$ is the sum of squares due to replication.

With this information available we can write

$$\boldsymbol{\omega} = \mathbf{C}_1^{-T} \boldsymbol{\delta}$$

to produce

$$S_{\lambda}(\boldsymbol{\beta},\boldsymbol{\omega}) = \frac{1}{n} \| \mathbf{B}_{2}^{\mathrm{T}} \mathbf{y} \|^{2} + \frac{1}{n} \| \mathbf{B}_{1}^{\mathrm{T}} \mathbf{y} - \mathbf{C}_{1} \mathbf{T}_{U} \boldsymbol{\beta} - \mathbf{C}_{1} \mathbf{K}_{U} \mathbf{C}_{1}^{\mathrm{T}} \boldsymbol{\omega} \|^{2} + \lambda \boldsymbol{\omega}^{\mathrm{T}} \mathbf{C}_{1} \mathbf{K}_{U} \mathbf{C}_{1}^{\mathrm{T}} \boldsymbol{\omega}$$

and proceed as in the case with no replications using C_1T_U in place of **T**, and $C_1K_UC_1^T$ in place of **K**. That is, take a QR decomposition

$$\mathbf{C}_1 \mathbf{T}_U = \mathbf{F} \mathbf{G} = [\mathbf{F}_1 : \mathbf{F}_2] \begin{bmatrix} \mathbf{G}_1 \\ \mathbf{0} \end{bmatrix} = \mathbf{F}_1 \mathbf{G}_1$$

and form $\mathbf{F}_2^{\mathrm{T}} \mathbf{C}_1 \mathbf{K}_U \mathbf{C}_1^{\mathrm{T}} \mathbf{F}_2$ which then determines the Cholesky decomposition

$$\mathbf{F}_2^{\mathrm{T}} \mathbf{C}_1 \mathbf{K}_U \mathbf{C}_1^{\mathrm{T}} \mathbf{F}_2 = \mathbf{L}^{\mathrm{T}} \mathbf{L} \ .$$

A SVD of \mathbf{L}^{T} as

$$\mathbf{L}^{\mathrm{T}} = \mathbf{U}\mathbf{D}\mathbf{V}^{\mathrm{T}}$$

and the product

$$\mathbf{w}_2 = \mathbf{F}_2^{\mathrm{T}} \mathbf{B}_1^{\mathrm{T}} \mathbf{y}$$

allows us to write

$$V(\lambda) = \frac{n\left[\left\| \mathbf{B}_{2}^{\mathrm{T}} \mathbf{y} \right\|^{2} + \sum_{j=1}^{k-t} \left[\frac{n\lambda}{d_{j}^{2} + n\lambda} \right]^{2} z_{j}^{2} \right]}{\left[n - k + \sum_{j=1}^{k-t} \frac{n\lambda}{d_{j}^{2} + n\lambda} \right]^{2}}$$

Given the value of $\hat{\lambda},$ the calculation of δ_λ and β_λ follow as in the no-replicates case. That

$$\delta_{\lambda} = \mathbf{C}_{1}^{\mathrm{T}} \mathbf{F}_{2} \mathbf{U} (\mathbf{D}^{2} + n \lambda \mathbf{I})^{-1} \mathbf{U}^{\mathrm{T}} \mathbf{F}_{2}^{\mathrm{T}} \mathbf{B}_{1}^{\mathrm{T}} \mathbf{y}$$

and β_{λ} is the solution of

$$\mathbf{G}_1 \boldsymbol{\beta}_{\boldsymbol{\lambda}} = \mathbf{F}_1^{\mathrm{T}} (\mathbf{B}_1^{\mathrm{T}} \mathbf{y} - \mathbf{C}_1 \mathbf{K}_U \boldsymbol{\delta}_{\boldsymbol{\lambda}}) \quad .$$

Partial spline models: These are an extension to the thin-plate smoothing spline model in which some of the coordinates of **x**, the "covariates", do not enter into the thin-plate spline. See Wahba (1984b, 1985) and Shiau, Wahba, and Johnson (1985). The model is

$$y_i = f(\mathbf{x}_i) + \sum_{j=1}^{c} \alpha_j \psi_j(\mathbf{x}_i, \mathbf{s}_i) + \varepsilon_i$$
(4.1)

in which \mathbf{s}_i are the "covariates" and $\{\psi_j\}$ are c given functions. For convenience, we will consider these variables as forming another matrix **S** of size $n \times c$. The partial spline estimates of f and α are the minimizers of

$$S_{\lambda}(\boldsymbol{\alpha},\boldsymbol{\beta},\boldsymbol{\delta}) = \frac{1}{n} \sum_{i=1}^{n} \left[y_i - f(\mathbf{x}_i) - \sum_{j=1}^{c} \alpha_j \psi_j(\mathbf{x}_i,\mathbf{s}_i) \right]^2 + \lambda J_m(f) ,$$

and it is known that the minimizing f_{λ} has a representation of the form (2.2). Let **S** be the $n \times c$ matrix with *ij* 'th entry $\psi_j(\mathbf{x}_i, \mathbf{s}_i)$. The matrix [**T**:**S**] must be of full column rank. The objective function for a fixed λ becomes

$$S_{\lambda}(\boldsymbol{\alpha},\boldsymbol{\beta},\boldsymbol{\delta}) = \frac{1}{n} \| \mathbf{y} - \mathbf{S}\boldsymbol{\alpha} - \mathbf{T}\boldsymbol{\beta} - \mathbf{K}\boldsymbol{\delta} \|^{2} + \lambda \, \boldsymbol{\delta}^{\mathrm{T}}\mathbf{K}\boldsymbol{\delta} \ .$$

When determining replicates, we only consider the d variables which determine the spline. When there are no replicates, we proceed as in the basic TPSS case except that we take the initial QR decomposition as

is,

$$[\mathbf{T}:\mathbf{S}] = \mathbf{F}\mathbf{G} = [\mathbf{F}_1:\mathbf{F}_2] \begin{bmatrix} \mathbf{G}_1 \\ \mathbf{0} \end{bmatrix} = \mathbf{F}_1\mathbf{G}_1$$

so $V(\lambda)$ is calculated as in (2.8) with all summations running to n - t - c. That is, after the Cholesky decomposition of $\mathbf{F}_2^T \mathbf{K} \mathbf{F}_2$ and the SVD of the transpose of the Cholesky factor, we have

$$V(\lambda) = \frac{n \sum_{j=1}^{n-t-c} \left[\frac{n\lambda}{d_j^2 + n\lambda}\right]^2 z_j^2}{\left[\sum_{j=1}^{n-t-c} \frac{n\lambda}{d_j^2 + n\lambda}\right]^2}$$

The calculation of ζ_{λ} and δ_{λ} proceeds as in the basic TPSS case. With these available, we solve for α_{λ} and β_{λ} simultaneously. In other words, we have simply replaced T β in (2.6) by

$$[\mathbf{T}:\mathbf{S}]\begin{bmatrix}\boldsymbol{\beta}\\\boldsymbol{\alpha}\end{bmatrix}$$

It can be shown that the implied constraint $S^T \delta = 0$ does not change the solution.

When we have covariates as well as some replications in the d coordinates of the \mathbf{x} 's, we have to distinguish between those columns of \mathbf{S} which follow the replication pattern of the \mathbf{x} 's and those which do not. If all the columns of \mathbf{S} follow the replication pattern, we have an indicator matrix \mathbf{M} for which

$$T = MT_U$$
$$K = MK_U$$
$$S = MS_U$$
.

Taking the QR decomposition of **M** as $\mathbf{M} = \mathbf{BC}$, we then take a QR decomposition of $\mathbf{C}_1[\mathbf{T}_U:\mathbf{S}_U]$ and proceed as above.

If there are columns of **S** which do not follow the replication pattern of the design, we need a more general approach. The covariate matrix is divided into $S = [S_1:S_2]$ in which the columns of S_1 have the same replication structure as the design points \mathbf{x}_i , $i = 1, \dots, n$. We have an indicator matrix **M** for which

$$[\mathbf{T}:\mathbf{S}_{1}:\mathbf{K}]=\mathbf{M}[\mathbf{T}_{U}:\mathbf{S}_{1U}:\mathbf{K}_{U}],$$

and a QR decomposition of $\mathbf{M} = \mathbf{BC}$ as above. However, we cannot easily reduce the objective function to a form such as (3.2) by premultiplying by **B**, as $\mathbf{B}_2^T \mathbf{S}_2$ is not annihilated. Instead we choose to take a QR decomposition of

$$\mathbf{C}_{1}[\mathbf{T}_{U}:\mathbf{S}_{1U}] = \mathbf{F}\mathbf{G} = [\mathbf{F}_{1}:\mathbf{F}_{2}]\begin{bmatrix}\mathbf{G}_{1}\\\mathbf{0}\end{bmatrix} = \mathbf{F}_{1}\mathbf{G}_{2}$$

which is used to reduce the parameter vector and penalty matrix. We proceed as in the case of a general design matrix with a semi-norm penalty as described in the next section by creating the parameter vector

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$$\boldsymbol{\theta} = \begin{bmatrix} \boldsymbol{\beta} \\ \boldsymbol{\alpha} \\ \boldsymbol{\zeta} \end{bmatrix} , \text{ with } \boldsymbol{\delta} = \mathbf{C}_1^{\mathrm{T}} \mathbf{F}_2 \boldsymbol{\zeta} ,$$

and the design matrix

$$\mathbf{X} = [\mathbf{T} : \mathbf{S}_1 : \mathbf{S}_2 : \mathbf{K} \mathbf{C}_1^{\mathrm{T}} \mathbf{F}_2] \quad .$$

The penalty becomes $\theta^{T} \Sigma \theta$, with

$$\boldsymbol{\Sigma} = \begin{bmatrix} \mathbf{0} & \mathbf{0} \\ \mathbf{0} & \mathbf{F}_2^{\mathrm{T}} \mathbf{C}_1 \mathbf{K}_U \mathbf{C}_1^{\mathrm{T}} \mathbf{F}_2 \end{bmatrix}$$

Partial spline models with nodes at selected points, which may not actually correspond to data points, are discussed in Appendix 2.

General design matrix with a semi-norm penalty: The ridge regression case and the TPSS cases which we have considered both have some special structure. In the ridge regression case, the design matrix, **X**, is general but the penalty term, $\gamma^T \gamma$, has a special form so we can streamline the calculations. In the TPSS cases, the penalty term, $\delta^T K \delta$ subject to $T^T \delta = 0$, is more general but the design matrix, [**T**:**K**] is related to the penalty so, again, we can exploit this special structure to provide faster algorithms. Even in the case with both a general design and a general penalty, though, we can still form efficient computational methods for GCV.

The most general GCV calculation we consider is the penalized least squares problem with an objective function

$$S_{\lambda}(\mathbf{\theta}) = \frac{1}{n} \| \mathbf{y} - \mathbf{X}\mathbf{\theta} \|^{2} + \lambda \mathbf{\theta}^{\mathrm{T}} \Sigma \mathbf{\theta}$$
(5.1)

where θ is a *p*-dimensional parameter vector, **y** is an *n*-dimensional response vector, **X** is an $n \times p$ design matrix, and Σ is a $p \times p$ positive semi-definite symmetric matrix defining the smoothness penalty. Note that partial splines can be written in this form as a special case.

A partial spline model with discontinuities in the $\{\psi_j\}$ of (4.1) which fits in the context of (5.1) is described in Shiau, Wahba, and Johnson (1985). Other special cases included splines and vector splines on the sphere (Wahba (1981), Wahba (1982a, 1982b, 1982c)) and remote sensing problems (Wahba (1980a)). Appendix 2 presents some examples and the algebra needed for a partial spline model with basis functions.

The minimization of (5.1) can also be used as a step in the iterative solution of penalized GLIM models (O'Sullivan (1983), O'Sullivan, Yandell and Raynor (1986)), nonlinear regularization problems (O'Sullivan (1983) and O'Sullivan and Wahba (1985)) and iteratively reweighted least squares problems (Green (1984), Green (1985) and Green and Yandell (1985)).

We can find the GCV estimate of λ in the general case by using a series of matrix decompositions to reduce (5.1) to the form of the ridge regression calculation as was done in the TPSS case. First we must isolate the null-space of the semi-norm defined by Σ . That is, we must describe the set of θ 's for which

$$\boldsymbol{\theta}^{\mathrm{T}}\boldsymbol{\Sigma}\,\boldsymbol{\theta} = \boldsymbol{0} \ .$$

We assume the dimension, h, of this space is known and take a pivoted Cholesky decomposition (Dongarra et al., 1979, chapter 8)

$$\mathbf{E}^{\mathrm{T}} \boldsymbol{\Sigma} \, \mathbf{E} = \mathbf{L}^{\mathrm{T}} \, \mathbf{L}$$

where **E** is a $p \times p$ permutation matrix and **L** is $(p-h) \times p$ with zeros below the main diagonal. The conditioning of **L** is evaluated to ensure that **L** actually has computational rank p-h. If **L** is rank deficient, we increase h until the resulting $(p-h) \times p$ matrix **L** is of full row rank and return a non-fatal error code. If the user's value of h was too large, we return a fatal error code as this indicates that the null space of Σ is smaller than expected. As described in Appendix 1, the technique of increasing h until **L** is of full row rank is incompatible with the partial spline code as written here.

A QR decomposition of \mathbf{L}^{T} as

$$\mathbf{L}^{\mathrm{T}} = \mathbf{Q}\mathbf{R} = [\mathbf{Q}_{1}:\mathbf{Q}_{2}]\begin{bmatrix}\mathbf{R}_{1}\\\mathbf{0}\end{bmatrix} = \mathbf{Q}_{1}\mathbf{R}_{1}$$

provides the $h \times p$ matrix \mathbf{Q}_2 which is an orthogonal basis for the null space of the semi-norm defined by Σ . We can now transform to parameters γ and β of dimension p - h and h, respectively, as

$$\begin{pmatrix} \boldsymbol{\gamma} \\ \boldsymbol{\beta} \end{bmatrix} = \begin{bmatrix} \boldsymbol{R}_1^T & \boldsymbol{0} \\ \boldsymbol{0} & \boldsymbol{I} \end{bmatrix} \boldsymbol{Q}^T \boldsymbol{E}^T \boldsymbol{\theta}$$

where β lies in the null space and $S_{\lambda}(\theta)$ from (5.1) can be written

$$S_{\lambda}(\boldsymbol{\beta},\boldsymbol{\gamma}) = \frac{1}{n} \| \mathbf{y} - \mathbf{X}\mathbf{E}\mathbf{Q} \begin{bmatrix} \mathbf{R}_{1}^{-T} & \mathbf{0} \\ \mathbf{0} & \mathbf{I} \end{bmatrix} \begin{bmatrix} \boldsymbol{\gamma} \\ \boldsymbol{\beta} \end{bmatrix} \|^{2} + \lambda \boldsymbol{\gamma}^{T}\boldsymbol{\gamma}$$
$$= \frac{1}{n} \| \mathbf{y} - \mathbf{Z} \begin{bmatrix} \boldsymbol{\gamma} \\ \boldsymbol{\beta} \end{bmatrix} \|^{2} + \lambda \boldsymbol{\gamma}^{T}\boldsymbol{\gamma}$$

with

$$\mathbf{Z} = [\mathbf{Z}_1:\mathbf{Z}_2] = \mathbf{X}\mathbf{E}\mathbf{Q} \begin{bmatrix} \mathbf{R}_1^{-T} & \mathbf{0} \\ \mathbf{0} & \mathbf{I} \end{bmatrix} .$$

This provides the desired form of the penalty term. We must now divide the least squares term into a part that can be made zero by an appropriate choice of β and a part that depends only on γ . Another QR decomposition, this time as

$$\mathbf{Z}_2 = \mathbf{F}\mathbf{G} = [\mathbf{F}_1; \mathbf{F}_2] \begin{bmatrix} \mathbf{G}_1 \\ \mathbf{0} \end{bmatrix}$$

is used to form

$$S_{\lambda}(\boldsymbol{\beta},\boldsymbol{\gamma}) = \frac{1}{n} \| \mathbf{w}_{1} - \mathbf{G}_{1}\boldsymbol{\beta} - \mathbf{J}_{1}\boldsymbol{\gamma} \|^{2} + \frac{1}{n} \| \mathbf{w}_{2} - \mathbf{J}_{2}\boldsymbol{\gamma} \|^{2} + \lambda \boldsymbol{\gamma}^{\mathrm{T}}\boldsymbol{\gamma}$$
(5.2)

where

$$\mathbf{w} = \begin{bmatrix} \mathbf{w}_1 \\ \mathbf{w}_2 \end{bmatrix} = \begin{bmatrix} \mathbf{F}_1^T \\ \mathbf{F}_2^T \end{bmatrix} \mathbf{y} = \mathbf{F}^T \mathbf{y}$$

and

$$\mathbf{J} = \begin{bmatrix} \mathbf{J}_1 \\ \mathbf{J}_2 \end{bmatrix} = \begin{bmatrix} \mathbf{F}_1^T \\ \mathbf{F}_2^T \end{bmatrix} \mathbf{Z}_1 = \mathbf{F}^T \mathbf{Z}_1 \ .$$

After checking that G_1 is non-singular, the first term in (5.2) can be made zero for any value of γ by solving

$$\mathbf{G}_1 \boldsymbol{\beta} = \mathbf{w}_1 - \mathbf{J}_1 \boldsymbol{\gamma} \tag{5.3}$$

for β . This reduces the general penalized least squares to the same form as the ridge regression. A singular value decomposition

$$\mathbf{J}_2 = \mathbf{U}\mathbf{D}\mathbf{V}^{\mathrm{T}} \tag{5.4}$$

produces the representation of the "hat" matrix as

$$\mathbf{A}(\lambda) = \mathbf{F} \begin{bmatrix} \mathbf{I} & \mathbf{0} \\ \mathbf{0} & \mathbf{U} \end{bmatrix} \begin{bmatrix} \mathbf{I} & \mathbf{0} \\ \mathbf{0} & \mathbf{D}^2 (\mathbf{D}^2 + n \lambda \mathbf{I})^{-1} \end{bmatrix} \begin{bmatrix} \mathbf{I} & \mathbf{0} \\ \mathbf{0} & \mathbf{U}^T \end{bmatrix} \mathbf{F}^T \ .$$

The matrix **D** is $a \times a$, with a = min(n, p) - h, and the matrices **U** and **V** are rectangular of sizes $(n-h) \times a$ and $(p-h) \times a$, respectively. Again, using

$$\mathbf{z} = \mathbf{U}^{\mathrm{T}} \mathbf{w}_2 \tag{5.5}$$

the GCV function can be expressed as

$$V(\lambda) = \frac{n \left[\|\mathbf{w}_{2}\|^{2} - \|\mathbf{z}\|^{2} + \sum_{j=1}^{a} \left[\frac{n\lambda}{d_{j}^{2} + n\lambda} \right]^{2} z_{j}^{2} \right]}{\left[n - p + \sum_{j=1}^{a} \frac{n\lambda}{d_{j}^{2} + n\lambda} \right]^{2}} .$$
 (5.6)

.

and the parameters vectors γ_{λ} , β_{λ} , and θ_{λ} are determined in the usual way given $\hat{\lambda}$, with (5.3) and

$$\boldsymbol{\gamma}_{\lambda} = \mathbf{V} (\mathbf{D}^2 + n \,\lambda \mathbf{I})^{-1} \mathbf{D} \mathbf{U}^{\mathrm{T}} \mathbf{w}_2 \ ,$$

yielding

$$\boldsymbol{\theta}_{\lambda} = \mathbf{E} \mathbf{Q} \begin{bmatrix} \mathbf{R}_{1}^{-T} & \mathbf{0} \\ \mathbf{0} & \mathbf{I} \end{bmatrix} \begin{bmatrix} \boldsymbol{\gamma}_{\lambda} \\ \boldsymbol{\beta}_{\lambda} \end{bmatrix} \ .$$

The biggest computational bottleneck is the SVD of J_2 when *n* and *p* are large, particularly since J_2 is often ill-conditioned. We can accelerate the SVD calculation by using a truncated version of the singular value decomposition (Bates and Wahba, 1982). Notice that, in (5.6) and the solution of γ_b , values of d_i such that

$$d_i^2 \ll n\lambda$$

can be set to zero without significantly changing the results. Starting with a tolerance $\tau \rho$, usually a small multiple (τ) of the relative machine precision (ρ), the truncated SVD algorithm finds a matrix \tilde{J}_2 which has $a^* \leq a$ positive singular values and satisfies

$$\frac{\|\|\tilde{\mathbf{J}}_2 - \mathbf{J}_2\|_F^2}{\|\|\mathbf{J}_2\|_F^2} < \tau \rho ,$$

in which $\|\|\bullet\||_F$ is the Frobenius norm. For details of the truncated SVD algorithm, see Appendix 3. We replace \mathbf{J}_2 by $\tilde{\mathbf{J}}_2$ in (5.4), thereby reducing the effective number of parameters to a^* . With the truncation we only calculate an $(n-h) \times a^*$ matrix \mathbf{U} and a $(p-h) \times a^*$ matrix \mathbf{V} so the vector \mathbf{z} defined in (5.5) will be a^* -dimensional, with a replaced by a^* . When \mathbf{J}_2 is ill-conditioned, we get a^* considerably less than a and, since the calculation of the SVD is of order $O(na^2)$, this can create substantial savings in computing time. However, V is sensitive to τ for small λ . To check on the effect of the truncation on the value of $V(\lambda)$ and hence the calculation of $\hat{\lambda}$ we return the diagnostic quantity

$$n\hat{\lambda} / (n\hat{\lambda} + \|\tilde{\mathbf{J}}_2 - \mathbf{J}_2\|_F^2) .$$
(5.7)

This is a lower bound on each of the quantities $n\hat{\lambda}/(d_j^2 + n\hat{\lambda})$ in (5.6) which are replaced by 1 when d_j is set to zero. Preliminary tests indicate that if the diagnostic quantity is above 0.999 then the truncation has negligible effect on V.

Another important method of accelerating the GCV calculations by avoiding the final reduction to diagonal form in the SVD was given by Elden (1984). This involves stopping the evaluation of the singular value decomposition at the intermediate step of the reduction of J_2 to a bidiagonal form, then forming an expression for $V(\lambda)$.

Description

The package has three main subroutine drivers. The first driver, *dtpss* for thin plate smoothing splines, is the most efficient and the most restrictive, allowing covariates only in the case where the replication pattern is the same as that found in the design. The second driver, *dptpss* for partial thin plate smoothing splines, handles general covariates and in turn calls the third driver, *dsnsm* which handles penalized least squares problems with a semi-norm penalty. After a call to *dtpss* or *dptpss* the subroutine *dpred* can be called to evaluate predicted values for additional points not in the design.

Replicates are handled in *dtpss* and *dptpss* using the following routines. The subroutine *dreps* sorts the x vectors and returns C_1 and the information necessary for the routines *duni* and *dsuy* (used only in *dtpss*). Subroutine *duni* reduces a matrix (**T** or **K**) to the corresponding matrix with unique entries (**T**_U or **K**_U). The routine *dsuy* sorts **y** and computes B_1^T **y** and the sum of squares due to replication.

The subroutine *dtpss*, the thin plate spline driver, calls the routine *dsetup* to create the matrices $\mathbf{C}_1 \mathbf{K}_U \mathbf{C}_1^T$ and $\mathbf{C}_1 [\mathbf{T}_U : \mathbf{S}_{1U}]$ from the design points \mathbf{x}_i , i = 1, 2, ..., n using the routines *dmakek* and *dmaket*. The LINPACK routine *dqrdc* is called to decompose $C_1[T_{11}:S_{11}]$ into its QR decomposition **FG**, followed by the routine *dftkf* to calculate $\mathbf{F}^{T}\mathbf{C}_{1}\mathbf{K}_{U}\mathbf{C}_{1}^{T}\mathbf{F}$. *Dsgdc1* does the Cholesky decomposition of $\mathbf{F}_{2}^{T}\mathbf{C}_{1}\mathbf{K}_{U}\mathbf{C}_{1}^{T}\mathbf{F}_{2}$ and the singular value decomposition of the Cholesky factor. Dgcvl uses these results to compute the generalized cross validation estimate of λ and the corresponding estimates of the other parameters. The work in dgcv1 is divided into application of the rotations by \mathbf{F}^{T} in *drsap*, optimization of the $V(\lambda)$ function in *dvlop*, computation of predictive mean square error (if requested) in dpmse, creation of the coefficient vector in dcfcr1, creation of the predicted values in *dpdcr*, and creation of the diagonal of $\mathbf{A}(\hat{\lambda})$ in *ddiag*. Subroutine *dvlop* calls dvmin to minimize $V(\lambda)$ by repeated calls to dvl. The minimization is done by an initial grid search in the $\ln(n\lambda)$ scale followed by a golden ratio search in the neighborhood of the minimizing grid point. The input variable ntbl controls the resolution of the initial grid search. A value for ntbl of 100 or greater is recommended to ensure that the global optimum is located. If a plot of $V(\lambda)$ versus $\ln(n\lambda)$ indicates that a local optimum has been obtained the user may either increase the value of *ntbl* or use the option to specify a reduced range for the search. The grid of $\ln(n\lambda)$ values is returned along with the corresponding $V(\lambda)$ values in the variable *tbl*. The variable auxtbl is returned containing $\hat{\lambda}$, $V(\hat{\lambda})$, V(0) and $V(\infty)$.

The driver dptpss for partial thin plate splines calls routines dreps, dmaket, duni, and dmakek to set up [T:S], $[\mathbf{T}_U:\mathbf{S}_{1U}]$ and \mathbf{K}_U . These are fed to dctsx to create the matrices Σ and \mathbf{X} which are used by the driver dsnsm.

The subroutine dsnsm is a general driver for penalized least squares problems with a seminorm penalty. It calls ddcom which decomposes Σ and X and returns information used by dgcv to find $\hat{\lambda}$, θ_{λ} , and other results. The work in ddcom is split into the decomposition of Σ in a call to dsgdc and the transformation and decomposition of the design in dcrtz and dzdc which in turn calls dtsvdc or dsvdc to perform the singular value decomposition. The work in dgcv is divided into the same subroutines as dgcvI with the exception that dcfcrI is replaced by dcfcr.

In the general case, the driver *dsnsm* allows an option to use a truncation singular value decomposition through the routine *dtsvdc* which preprocesses the design matrix J_2 to reduce the dimensionality before invoking *dsvdc* (see Appendix 3). The truncation tolerance, $\tau \times \rho$ is passed to *dtsvdc* as the parameter *minrat*. The drivers *dtpss* and *dptpss* would not benefit from truncation in the SVD calculation so they use the LINPACK routine *dsvdc*.

Simulation Applications: When GCVPACK is used for simulation studies the option to compute the predictive mean square error should be used. The known "true" response is input in the variable *adiag* and the predictive mean square error, $R(\lambda)$, is returned, along with $V(\lambda)$, in the variable *tbl*. It is recommended that plots of $V(\lambda)$ and $R(\lambda)$ versus $\ln(n\lambda)$ be used to evaluate the success of the GCV function in finding the optimal λ (the λ which minimizes predictive mean square error). The variable *auxtbl* contains $R(\hat{\lambda})$, R(0) and $R(\infty)$.

The decomposition of the X matrix requires the most intensive computation. The subroutines *dtpss* and *dptpss* are both set up to take advantage of the savings in computation that exist for multiple response vectors with the same design. To modify *dtpss* to handle a problem with more than one response vector all code up to and including the call to *dsgdc1* is executed once. A loop can be added to execute the remaining code for each \mathbf{y} vector. In practice this modification would involve adding only a few lines of code.

To modify *dptpss*, or any other driver which calls *dsnsm*, a loop must be added in *dsnsm*. In *dsnsm* there are two subroutines, *ddcom* which needs to be executed once, and *dgcv* which must be executed once for each response vector. In *dptpss*, after the call to *dsnsm*, a transformation is applied to the coefficient vector. This must be done to the coefficient vector corresponding to each **y** vector.

Related Algorithms

The numerical linear algebra in our routines is performed using the LINPACK (Dongarra et al., 1979) routines. The introductory comments of each GCVPACK routine list which LINPACK and BLAS (Basic Linear Algebra Subroutines) routines are called directly or indirectly. There is one machine-dependent constant, the relative machine precision, which is used in these routines to determine error conditions caused by ill-conditioning, but that constant is computed each time it is needed.

The present work generalizes algorithms for ridge regression of Golub, Heath, and Wahba (1979) and Bates and Wahba (1982) which use the singular value decomposition. Elden (1977) gives an algorithm which terminates the singular value decomposition at an intermediate step, reducing X to a bidiagonal form, thereby saving time (see the **Test Results** section). This could be incorporated into GCVPACK but we have not done so yet.

Wendelberger (1981) implemented an algorithm for thin plate splines based on eigenvalueeigenvector decompositions for one-dimensional and multi-dimensional thin plate smoothing splines. Hutchinson (1984) developed an algorithm for thin plate splines with large data sets using the thin plate basis functions of Wahba (1980b); see Appendix 2.

Reinsch (1967) initially proposed a fast algorithm for fixed λ using a Cholesky decomposition (see De Boor (1978)). In the one-dimensional case, the penalty can be written as a product of matrices with only 2m-1 non-zero diagonals. Hutchinson and de Hoog (1985) give an O(n)

algorithm for computing $V(\lambda)$ using a Cholesky decomposition of these matrices. See also O'Sullivan (1985). GCVPACK is not designed to take advantage of the unique structure of one dimensional polynomial smoothing splines, and runs much slower than the code of Hutchinson and de Hoog (1985) in this case.

O'Sullivan, Yandell and Raynor (1986) developed algorithms for smooth generalized linear models based on a Cholesky decomposition of $\mathbf{X}^T \mathbf{X} + n \lambda \mathbf{I}$. Green (1985) and Green and Yandell (1985) presented algorithms for penalized likelihood schemes which include generalized linear models and other iteratively reweighted least squares methods. They present a one-dimensional algorithm based on Reinsch (1967) and a general algorithm based on the Cholesky decomposition. They have also incorporated an iterative algorithm using the SVD to automate the choice of $\hat{\lambda}$, but it needs extensive testing to determine if it is stable. Shiau (1985) developed algorithms for a particular class of partial splines consisting of discontinuities of f or higher order derivatives at known or unknown points. This includes a one-dimensional algorithm based on Hutchinson and de Hoog (1985) and a multidimensional algorithm based on the Cholesky decomposition.

Test Results

The package and drivers have been tested for internal consistency and for accuracy against other known algorithms. Here we present some timing results to show that the methods are feasible for relatively large data sets and to offer insight into which portions of the code should be avoided, if possible. For example, the code allows the computation of the diagonal of $\mathbf{A}(\hat{\lambda})$ for forming diagnostics (Eubank, 1984) but this calculation alone can take 15% or more of the total execution time.

All timing runs were performed on a Vax-11/750 computer with a floating point accelerator and running the 4.2 BSD $UNIX^{TM}$ operating system. We quote two sets of times for the example: one using the driver *dtpss* and the other using *dptpss*. Each of the drivers was timed twice: first using the Fortran version of the Basic Linear Algebra Routines (BLAS) then using Assembler Language BLAS. As explained in Dongarra et al. (1979), the BLAS are a set of low-level routines that perform such elementary tasks as accumulation of dot products and, by replacing them with Assembler language versions, the Linpack routines can be made to run faster.

The design for the example is a 9 by 9 factorial in \mathbf{x}_1 and \mathbf{x}_2 with one covariate, \mathbf{x}_2^2 . Two replicate observations were simulated at each of the 81 design points. Thus n = 162, k = 81, m = 2, d = 2 and c = 1. Our timing results are shown in Tables 1 and 2. The total times are slightly greater than the sum of the times spent in the lower level subroutines since the driver routines have to do some definition of pointers, etc.

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	Fortran BLAS		Assembler BLAS	
Routine	Time (sec.)	Percentage	Time(Sec.)	Percentage
dreps	3.07	3	3.03	4
dsetup	11.85	11	9.42	12
dsgdc1				
Cholesky	3.70	3	2.13	3
bidiag.	31.55	29	15.50	19
diag.	36.10	33	34.65	43
dsuy	0.05	0	0.05	0
dgcv1				
drsap	0.20	0	0.12	0
dvlop	1.70	2	1.68	2
dpmse	1.47	1	1.37	2
dcfcr1	0.28	0	0.15	0
dpdcr	0.25	0	0.12	0
ddiag	18.07	17	11.15	14
Total dtpss	108.92		79.88	

 Table 1: Example 1 using dtpss

	Fortran BLAS		Assembler BLAS	
Routine	Time (sec.)	Percentage	Time (sec.)	Percentage
dreps	3.02	1	3.03	1
make K and T	14.45	4	14.62	6
dctsx	8.38	2	3.98	2
ddcom				
dsgdc				
Cholesky	4.05	1	2.22	1
QR	10.85	3	5.40	2
dcrtz	56.30	16	27.45	12
dzdc				
bidiag.	86.43	24	41.77	18
diag.	101.55	28	82.55	36
dgcv				
drsap	0.42	0	0.23	0
dvlop	1.67	0	1.67	1
dpmse	1.65	0	1.45	1
defer	0.57	0	0.28	0
dpdcr	0.47	0	0.25	0
ddiag	66.02	18	39.65	18
Total dptpss	359.43		226.28	

Table 2: Example 1 using dptpss

The first thing to notice from these tables is that *dtpss* is strongly preferred over *dptpss* for this example since it executes approximately 3 times faster. In general, if *dtpss* can solve the problem, it will do so more quickly. Also, the Assembler BLAS speed things up considerably with most of the gain being in the call to the Linpack SVD routine *dsvdc*.

We have divided the time for dsvdc into two subsections, bidiag and diag. Elden (1984) gave a method of expressing the GCV function $V(\lambda)$ avoiding the diag step. This would result in considerable savings in the dsgdcl or ddcom routines. This savings is offset by the calculations in dgcvl or dgcv becoming more complicated and, possibly, taking longer. However, since those routines take up much less time than diag, we would expect that the overall savings would be worthwhile.

Notice that the calculation of the diagonal of $\mathbf{A}(\hat{\lambda})$ in *ddiag* is comparatively expensive – usually around 15% of the total execution time. If this optional information is not going to be used, it should not be calculated.

In circumstances where there are multiple **y** vectors being analysed for the same design and penalty matrices, such as in Monte-Carlo runs, the decomposition portion, dsgdcl or ddcom, should be called only once while the analysis portion, dgcvl or dgcv, called for each **y**. The analysis portion represents less than 5% of the total time if the calculation of the diagonal of $\mathbf{A}(\hat{\lambda})$ is not undertaken.

The sorting method used in *dreps* is a comparatively primitive sort (a modification of the bubble sort) but, even so, the time taken by *dreps* is a small percentage of the total time. It would be possible to speed up this step by using a more sophisticated sort, but it doesn't appear worthwhile. Also, the evaluation of $V(\lambda)$ after the matrices are decomposed is very quick. In these runs the variable *ntbl* was set to 200 so both $V(\lambda)$ and the mean squared error of prediction (since the data were simulated) were evaluated at 200 different values of λ . Even with 200 evaluations *dvlop* and *dpmse* each represented, at most, 2% of the execution time.

Appendix 1. – replicates and rank-deficient penalty matrices

Because the functions E_m defined in (2.3) are increasing functions of the length of their argument, the matrix **K** defined in (2.4) will be close to singular if $||\mathbf{x}_i - \mathbf{x}_j||$ is very small for some $i \neq j$. To avoid an indeterminacy in the parameters of the thin-plate spline, we determine replicates by comparing $||\mathbf{x}_i - \mathbf{x}_j||$ to a tolerance level rather than checking for $\mathbf{x}_i = \mathbf{x}_j$. The tolerance level is calculated as 100 times the relative machine precision times the length of the diagonal of the smallest rectangle which encloses the $\mathbf{x}_i, i = 1, ..., n$. In all our test cases, this check has been adequate to ensure that the matrix $\mathbf{F}_2^T \mathbf{K} \mathbf{F}_2$ is computationally positive definite.

It is important to note that the determination of replicates involves sorting the \mathbf{x}_i , i = 1, ..., n, in increasing lexicographic order. That is,

th

$$e \text{ rows of} \begin{bmatrix} 1 & 1 \\ 2 & 1 \\ 3 & 1 \\ 4 & 1 \\ 1 & 2 \\ 2 & 2 \\ 3 & 2 \\ 4 & 2 \end{bmatrix} \text{ would be re-ordered as} \begin{bmatrix} 1 & 1 \\ 1 & 2 \\ 2 & 1 \\ 2 & 2 \\ 3 & 1 \\ 3 & 2 \\ 4 & 1 \\ 4 & 2 \end{bmatrix}$$

As mentioned in the **Test Results** section, the sorting algorithm is comparatively primitive (a modification of a bubble sort) and, even though it does not take a substantial percentage of the total execution time, it is to the user's advantage to pass the argument *des* to *dtpss* or *dptpss* with the rows in increasing lexicographic order, if possible, as the sorting time will be minimized.

Replicates are determined in such a way as to avoid a singular penalty matrix because a singular penalty matrix has a different effect for the thin-plate smoothing spline (or partial spline) than it does for the case of a general design matrix with a semi-norm penalty. In the general case, we determine the null space of the penalty so unexpected singularities simply increase the dimension of the null space and that part of the parameter vector is incorporated into the β . Ordinary regression is used to determine β and we assume (and check) that the part of the design matrix corresponding to β is non-singular. Unless the singularity in the penalty corresponds to a singularity in the design, everything works well.

In the case of a thin-plate smoothing spline the least squares part of the objective function (2.6) uses the same matrix $(\mathbf{F}_2^T \mathbf{K} \mathbf{F}_2)$ as the penalty part. Thus, when the penalty is rank-deficient, the "design" matrix (in the regression sense) is also rank deficient and the parameters which lie in the extended null space of the penalty are indeterminant. This can be seen from the form of (2.6). If there are singular values of zero, the corresponding parameters have no effect on the predictions and thus do not enter into the objective function $S_{\lambda}(\beta, \delta)$. There is a parameter vector which can be calculated using (2.9) even with some zero singular values but the part corresponding to the zero singular values can be changed to an arbitrary value without affecting the predictions so, in particular, it could be set to zero. More specifically, consider the last two terms in the last line of (2.6), after the Cholesky decomposition:

$$\frac{1}{n} \|\mathbf{w}_2 - \mathbf{L}^{\mathrm{T}} \mathbf{L} \boldsymbol{\delta} \| + \lambda \boldsymbol{\delta}^{\mathrm{T}} \mathbf{L}^{\mathrm{T}} \mathbf{L} \boldsymbol{\delta} .$$
 (A1.1)

If **L** is not of full row rank, any δ satisfying

$$\mathbf{L}^{\mathrm{T}}\mathbf{L}\mathbf{w}_{2} = [(\mathbf{L}^{\mathrm{T}}\mathbf{L})^{2} + n\,\lambda\,\mathbf{L}^{\mathrm{T}}\mathbf{L}]\,\boldsymbol{\delta}$$

minimizes (A1.1), and in particular we could take

$$\boldsymbol{\delta} = (\mathbf{L}^{\mathrm{T}}\mathbf{L} + n\,\boldsymbol{\lambda}\,\mathbf{I})^{-1}\,\mathbf{w}_2 \ .$$

However, we have chosen not to write the special code that would be required to handle this case.

We have eliminated one source of a computationally singular penalty matrix for the thin plate spline by merging nearly replicated data points. If the computational singularity of $\mathbf{F}_2^T \mathbf{K} \mathbf{F}_2$ is due to other than nearly replicated data points, i.e., due to very large sets of highly irregularly spaced data, the user should consider using thin plate basis functions as described in Appendix 2.

Appendix 2. - partial splines with basis functions

One can use the algorithm for a general design matrix with semi-norm penalty to find partial thin-plate smoothing splines determined by basis functions centered at specified nodes. See Shiau, Wahba, and Johnson (1985). For example, consider the model

 $y_i = \int \cdots \int K(\mathbf{x}_i, \mathbf{x}) f(\mathbf{x}) d\mathbf{x} + \varepsilon_i$ $= L_i f + \varepsilon_i \quad .$

The estimate f_{λ} of f is the minimizer, in an appropriate space, of

$$\frac{1}{n} \sum_{i=1}^{n} (y_i - L_i f)^2 + \lambda J(f)$$
 (A2.1)

where J(f) is an appropriate (quadratic) roughness penalty. If we can approximate f_{λ} by

$$f_{\lambda} = \sum_{i=1}^{b} \theta_i B_i$$

where $\{B_i\}$ are suitably chosen basis functions, then we can define the *ij* 'th entry of **X** as $L_i B_j$ and the matrix Σ by $J(\Sigma \theta_i B_j) = \mathbf{0}^T \Sigma \mathbf{0}$.

The thin plate basis functions were proposed for this purpose by Wahba (1980a). Starting with a set of suitably distributed distinct nodes t_1, t_2, \dots, t_b , the approximation is

$$f_{\lambda}(\mathbf{x}) = \sum_{i=1}^{l} \beta_i \phi_i(\mathbf{x}) + \sum_{i=1}^{b} \delta_i E_m(\mathbf{x} - \mathbf{t}_i)$$
(A2.2)

where $\boldsymbol{\delta} = (\delta_1, \cdots, \delta_b)^T$ must satisfy

$$\sum_{i=1}^{b} \delta_i \phi_j(\mathbf{t}_i) = 0 , j = 1, \cdots t .$$

If f_{λ} is required to be of the form (A2.2), then (A2.1) becomes

$$S_{\lambda}(\boldsymbol{\beta},\boldsymbol{\delta}) = \frac{1}{n} \| \mathbf{y} - \mathbf{T}\boldsymbol{\beta} - \mathbf{K}\boldsymbol{\delta} \|^{2} + \lambda \boldsymbol{\delta}^{\mathrm{T}}\mathbf{K}_{B}\boldsymbol{\delta}$$

subject to $\mathbf{T}_{B}^{\mathrm{T}} \boldsymbol{\delta} = \mathbf{0}$, with

$$\{\mathbf{T}_B\}_{ij} = \phi_j(\mathbf{t}_i) \ .$$

Here, **T** is $n \times t$ and **K** is $n \times b$, with entries

$$\{\mathbf{T}\}_{ij} = L_i \phi_j ,$$

$$\{\mathbf{K}\}_{ij} = L_i E_m (\bullet - \mathbf{t}_j)$$

and \mathbf{K}_{B} is $b \times b$ with entries

$$\{\mathbf{K}_B\}_{ij} = E_m(\mathbf{t}_i - \mathbf{t}_j) \ .$$

If we are interested simply in evaluation functionals, then $L_i f = f(\mathbf{x}_i)$. The matrices \mathbf{T}_B and \mathbf{K}_B remain the same, but the matrices \mathbf{T} and \mathbf{K} have entries

$$\{\mathbf{T}\}_{ij} = \phi_j(\mathbf{x}_i) ,$$

$$\{\mathbf{K}\}_{ij} = E_m(\mathbf{x}_i - \mathbf{t}_j) .$$

We take a QR decomposition

$$\mathbf{T}_{B} = \mathbf{F}\mathbf{G} = [\mathbf{F}_{1};\mathbf{F}_{2}]\begin{bmatrix}\mathbf{G}_{1}\\\mathbf{0}\end{bmatrix} = \mathbf{F}_{1}\mathbf{G}_{1}$$

and use this to construct the parameter vector

$$\boldsymbol{\theta} = \begin{bmatrix} \boldsymbol{\beta} \end{bmatrix}$$
, with $\boldsymbol{\delta} = \mathbf{F}_2 \boldsymbol{\zeta}$,

and to create the design matrix

$$\mathbf{X} = [\mathbf{T} : \mathbf{KF}_2]$$

and penalty matrix

$$\Sigma = \begin{bmatrix} \mathbf{0} & \mathbf{0} \\ \mathbf{0} & \mathbf{F}_2^{\mathrm{T}} \mathbf{K}_B \, \mathbf{F}_2 \end{bmatrix} \quad .$$

We then proceed as in the case of a general design matrix with a semi-norm penalty as described earlier.

Hutchinson's (1984) code implements thin plate basis functions for the case $L_i f = f(\mathbf{x}_i)$, where b is chosen to be much less than n when n is large. Hutchinson's code, or the partial thin plate smoothing spline code described here, should be considered in the case that n is very large or $\mathbf{F}_2^T \mathbf{K} \mathbf{F}_2$ of (2.6) is computationally singular.

Covariates and replicates are handled as before and enter in the same way as for partial spline models. Considering here only the case of no replicates, the model with covariates is

$$f_{\lambda}(\mathbf{x},\mathbf{s}) = \sum_{i=1}^{t} \beta_i \phi_i(\mathbf{x}) + \sum_{i=1}^{b} \delta_i E_m(\mathbf{x}-\mathbf{t}_i) + \sum_{j=1}^{c} \alpha_j \psi_j(\mathbf{x},\mathbf{s}) .$$

The objective function for a fixed λ becomes

$$S_{\lambda}(\boldsymbol{\alpha},\boldsymbol{\beta},\boldsymbol{\delta}) = \frac{1}{n} \| \mathbf{y} - \mathbf{S}\boldsymbol{\alpha} - \mathbf{T}\boldsymbol{\beta} - \mathbf{K}\boldsymbol{\delta} \|^{2} + \lambda \,\boldsymbol{\delta}^{\mathrm{T}}\mathbf{K}_{B}\,\boldsymbol{\delta}$$

subject to $\mathbf{T}_{B}^{T} \boldsymbol{\delta} = \mathbf{0}$, in which **S** is $n \times c$ with entries

$$\{\mathbf{S}\}_{ij} = L_i \psi_j(\bullet, \mathbf{s}_i)$$
,

or, for evaluation functionals,

$$\{\mathbf{S}\}_{ij} = \Psi_j(\mathbf{x}_i, \mathbf{s}_i) \ .$$

The design matrix becomes

$$\mathbf{X} = [\mathbf{T} : \mathbf{S} : \mathbf{KF}_2]$$

with parameter vector

$$\theta = \begin{bmatrix} \beta \\ \alpha \\ \zeta \end{bmatrix}$$

The penalty Σ has the same form, with the addition of rows and columns of zeroes corresponding to α . One would then proceed with the general design matrix with semi-norm penalty.

Appendix 3. - the truncated singular value decomposition

The following theorem of Mirsky (1960) provides a bound for the error in the singular values when using an approximation to a matrix.

Theorem 1: Let **X** and **Y** be $n \times p$ $(n \ge p)$ matrices with singular value decompositions **UDV**^T and **RSW**^T respectively. Denote the ordered singular values of **X** as $\{d_i\}, i=1, \ldots, p$ with . p. Then $d_1 \ge d_2$

$$\geq d_2 \geq \cdots \geq d_p$$
 and the ordered singular values of **x** as $\{s_i\}, i=1, \ldots, p$. Then

$$\sum_{i=1}^{p} (d_i - s_i)^2 \le \left\| \mathbf{X} - \mathbf{Y} \right\|_F^2 = tr \left[(\mathbf{X} - \mathbf{Y})^{\mathrm{T}} (\mathbf{X} - \mathbf{Y}) \right]$$

We will take advantage of this theorem to calculate the SVD of a matrix X_{a^*} which is close to X in the sense that $\|X - X_{a^*}\|$ is small but is better conditioned than is X so the iterative portion of the SVD tends to converge faster and the computational burden is reduced. First, we take a pivoted QR decomposition of X using the pivoting scheme from LINPACK (Dongarra et al., 1979). That is, we determine Q, $n \times n$ orthogonal, R, $n \times p$ and zero below the main diagonal, and **E**, a $p \times p$ permutation matrix, such that

$$\mathbf{X}\mathbf{E} = \mathbf{Q}\mathbf{R} \tag{A3.1}$$

and R has the property that

$$r_{a^*,a^*}^2 \ge \sum_{i=a^*}^j r_{i,j}^2 \quad (j = a^*, a^* + 1, ..., p).$$
(A3.2)

If we take the SVD of \mathbf{R}_p , the triangular matrix composed of the first p rows of \mathbf{R} , as

$$\mathbf{R}_{p} = \mathbf{K} \mathbf{D} \mathbf{L}^{\mathrm{T}} \tag{A3.3}$$

we can produce the SVD of X as

$$\mathbf{X} = \mathbf{Q}_{p} \mathbf{K} \mathbf{D} \mathbf{L}^{\mathrm{T}} \mathbf{E}^{\mathrm{T}} = \mathbf{U} \mathbf{D} \mathbf{V}^{\mathrm{T}}$$
(A3.4)

where \mathbf{Q}_p is the $n \times p$ matrix composed of the first p columns of \mathbf{Q} and $\mathbf{U} = \mathbf{Q}_p \mathbf{K}$ is $n \times p$ while $\mathbf{V} = \mathbf{EL}$ is $p \times p$ and orthogonal. This method would not, however, produce better conditioning for the SVD algorithm since the singular values of \mathbf{R}_p are the same as the singular values of \mathbf{X} .

To provide better conditioning, we truncate the matrix \mathbf{R}_p after the a^* 'th row and take the SVD of the resulting $n \times a^*$ matrix \mathbf{R}_{a^*} ($a^* \le p$) as

$$\mathbf{R}_{a^*} = \mathbf{K}_{a^*} \mathbf{D}_{a^*} \mathbf{L}_{a^*} \tag{A3.5}$$

where \mathbf{K}_{a^*} is $a^* \times a^*$ and \mathbf{L}_{a^*} is $a^* \times p$. The diagonal elements of \mathbf{D}_{a^*} are no longer the singular values of **X** but now represent the singular values of a matrix

$$\mathbf{X}_{a^*} = \mathbf{Q}_{\rho} \begin{bmatrix} \mathbf{R}_{a^*} \\ \mathbf{0} \end{bmatrix} \mathbf{E}^{\mathrm{T}}$$
(A3.6)

which is different from X. However,

$$\|\mathbf{X}-\mathbf{X}_{a^*}\|_F = \left(\sum_{i=a^*+1}^{p} \sum_{j=i}^{p} r_{i,j}^2\right)^{\frac{1}{2}}$$
(A3.7)

so we can choose a^* to be as small as possible subject to the constraint that

$$\frac{\|\mathbf{X} - \mathbf{X}_{a^*}\|_F}{\|\mathbf{X}\|_F} \le \tau \rho \tag{A3.8}$$

where ρ is the relative machine precision (the smallest number such that $1+\rho > 1$ in floating point arithmetic) and τ is a small multiplier.

We initially choose τ as unity but increase it if the LINPACK singular value decomposition routine (*dsvdc*) fails to converge. When such a convergence failure occurs, the user can either increase the number of iterations per singular value allowed in *dsvdc* (we increase this from 30 to 90) or increase τ or both. To increase the maximum allowable number of iterations, change the value of MAXIT in *dsvdc*.

Allowing τ to get too large can result in inaccuracies in the calculation of V. The effect of the truncation is measured by the diagnostic ratio defined in (5.7). In general, values of τ above 100 are not recommended.

The double sum on the right of (A3.7) is easily evaluated a row at a time starting at the p'th row until the constraint (A3.8) is violated and the smallest a^* is determined.

By theorem 1, if $\{d_i\}$, i=1,...,p are the ordered singular values of X and $\{d_{i,a^*}\}$, i=1,...,p are the ordered singular values of X_{a^*} , then

$$\left(\sum_{i=1}^{p} (d_{i} - d_{i,a^{*}})^{2}\right)^{\frac{1}{2}} \le \tau \rho \| \mathbf{X} \| = \tau \rho \left(\sum_{i=1}^{p} d_{i}^{2}\right)^{\frac{1}{2}}$$
(A3.9)

If $n \leq p$, the same procedure is applied to \mathbf{X}^{T} .

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Table 3. GCVPACK notation correspondence			
integer constants			
n	nobs	number of observations	
d	dim	dimension of polynomial space	
m	m	order of derivatives of penalty	
с	ncov	number of covariates	
<i>c</i> ₁	ncov1	number of covariates in S replicating structure of T	
$c - c_1$	ncov2	ncov – ncov1	
a	nuobs	iout[4] = number of unique obs. (dtpss & dptpss)	
a*	npsing	iout[1] = number of positive singular values	
t	mkpoly(m,dim)	dimension of polynomial space	
h=t+c	nnull	$iout[3] = size of null space of \Sigma$	
p = a + t + c	npar	iout[2] = number of parameters	
<i>t</i> + <i>c</i> ₁	ncts1	number of columns in $[\mathbf{T}: \mathbf{S}_1]$	
p−h	pmh	npar – nnull	
n-h	nmh	nobs – nnull	
data and parameter vectors			
У	у	response vector	
β	beta	coefficients for covariates	
α	alpha	coefficients for polynomial	
δ	delta	coefficients for smooth	
γ	coef	coefficients for well-defined smooth basis	
θ	coef	coefficients (in several forms)	

(continued)

		matrices
X	des	design matrix for splined variables
$[\mathbf{T}:\mathbf{S}_1]$	ts1	polynomials and replicated covariates
$[\mathbf{T}_U:\mathbf{S}_{1U}]$	tbsb1	unique polynomials and replicated covariates
\mathbf{S}_2	s2	unreplicated covariates
$\{\mathbf{A}(\hat{\lambda})\}_{ii}$	adiag[i]	diagonal of hat matrix
{ D } _{ii}	svals[i]	singular values
Σ	sigma	penalty matrix
F,G	fg & fgaux	QR decomposition of $[\mathbf{T}:\mathbf{S}_1]$
Е	sgpvt	permutation for pivoted Cholesky of Σ
Q,R	qr & qraux	QR decomposition of Cholesky factor of Σ
{ C ₁ } _{<i>ii</i>}	c1[i]	square root of number of replicates of <i>i</i> 'th unique x
		double precision summaries
Î	lamhat	dout[1] = GCV estimate of lambda
J(f)	penlty	dout[2] = smoothing penalty
$\ \mathbf{I} - \hat{\mathbf{A}}(\hat{\lambda})\ ^2$	rss	dout[3] = residual sum of squares
$tr(\mathbf{I}-\mathbf{A}(\hat{\lambda}))$	_	$dout[4] = trace of \mathbf{I} - \mathbf{A}$
$\ \mathbf{B}_{2}^{\mathrm{T}}\mathbf{y}\ ^{2}$	ssqrep	dout[5] = sum of squares for replication (dtpss)
ρ	machep	relative machine precision
τ	tau	small multiple
τρ	minrat	machine tolerance

Table 3 continued.

Code for Driver Routines

```
.....
dptpss.com
.....
     subroutine dptpss(des,lddes,nobs,dim,m,s,lds,ncov1,ncov2,y,ntbl,
     * adiag, lamlim, dout, iout, coef, svals, tbl, ldtbl, auxtbl, work,
     * lwa,iwork,liwa,job,info)
      integer lddes, nobs, dim, m, lds, ncov1, ncov2, ntbl, iout (4), ldtbl, lwa,
     * liwa,iwork(liwa),job,info
      double precision des (lddes, dim), s (lds, *), y (nobs), adiag (nobs),
     * lamlim(2), dout(4), coef(*), svals(*), tbl(ldtbl, 3),
     * auxtbl(3,3),work(lwa)
С
c Purpose: determine the generalized cross validation estimate of the
        smoothing parameter and fit model parameters for a partial thin
с
С
        plate spline model.
с
c On Entry:
С
    des(lddes,dim)
                        design for the variables to be splined
с
    lddes
                        leading dimension of des as declared in the
                        calling program
с
с
    nobs
                        number of observations
с
    dim
                        number of columns in des
С
                        order of the derivatives in the penalty
    m
с
    s(lds,ncovl+ncov2) design for the covariates
С
                        first ncov1 columns contain covariates which
С
                          duplicate the replication structure of des
с
                        next ncov2 columns contain covariates which
С
                          do not duplicate the replication structure of
с
                          des
```

с	lds	leading dimension of s as declared in the
С		calling program
с	ncovl	number of covariates which duplicate the
¢		replication structure of des
С	ncov2	number of covariates which do not duplicate the
с		replication structure of des
с	y(nobs)	response vector
с	ntbl	number of evenly spaced values for
с		log10(nobs*lambda) to be used in the initial
c		grid search for lambda hat
Ċ		if nth] = 0 only a colden ratio search will be
č		done and this not referenced if n this 0
č		there will be athl rows returned in the
č	adiag (nobe)	"true" y values on entry if predictive man is
2	ading (nobs)	cide y values on encry if predictive mse is
C .	11 d= (2)	lightere lecht het ersch die le 10/sel t
С	LamLim(2)	limits on lambda hat search (in logiU(nobs*
c		lambda) scale) if user input limits are
С		requested. if lamlim(1) = lamlim(2) then lamhat
¢		is set to (10**lamlim(1))/nobs
с	ldtbl	leading dimension of tbl as declared in the
С		calling program
С	job	integer with decimal expansion abc
с		if a is nonzero then predictive mse is computed
с		using adiag as true y
с		if b is nonzero then user input limits on search
с		for lambda hat are used
с		if c is nonzero then adiag will be calculated
c 0	n Exit:	
с	des(lddes,dim)	unique rows of des
с	v(nobs)	predicted values
~	adiac(nobs)	diagonal elements of the hat matrix if requested
2	lamlim(2)	limits on lambda hat search
2	$\operatorname{IamIIIm}(2)$	(in log10 (nobstlambda) scale)
с ~	d_{out} (1)	(in logic (nobb lambad) board,
-	douc (4)	1 lombat generalized gross validation
С		i familat generalized cross varidación
С		2 mestale of the smoothing parameter
с		2 penicy smoothing penalcy
с		3 rss residual sum of squares
с		4 $tr(1-A)$ trace of $1 - A$
с	iout(4)	contains:
С		1 npsing number of positive singular values
с		if info indicates nonzero info
с		from dsvdc then npsing contains
с		info as it was returned from dsvdc
с		2 npar number of parameters
с		(npar = nuobs + nnull)
С		3 nnull size of the null space of sigma
с		(m+dim-1 choose dim)+ncov1+ncov2
с		4 nucbs number of unique rows in des
c	coef(npar)	coefficient estimates [beta':alpha':delta']'
c		coef must have a dimension of at least
c		-
Ŭ		nuobs+nnull
0	svals (nosing)	nuobs+nnull singular values, svals must have a dimension.
c	svals(npsing)	nuobs+nnull singular values, svals must have a dimension, of at least nuobs-nnull.
с с с	svals(npsing)	nuobs+nnull singular values, svals must have a dimension, of at least nuobs-nnull. if info indicates nonzero info in davdo then
000	svals(npsing)	nuobs+nnull singular values, svals must have a dimension, of at least nuobs-nnull. if info indicates nonzero info in dsvdc then suals is as returned from dsvdc
0000	<pre>svals(npsing) tbl(ldtbl_2)</pre>	<pre>nuobs+nnull singular values, svals must have a dimension, of at least nuobs-nnull. if info indicates nonzero info in dsvdc then svals is as returned from dsvdc. column contains</pre>
0 0 0 0	<pre>svals(npsing) tbl(ldtbl,3)</pre>	<pre>nuobs+nnull singular values, svals must have a dimension, of at least nuobs-nnull. if info indicates nonzero info in dsvdc then svals is as returned from dsvdc. column contains arid of log(0(nobstlambds))</pre>
0 0 0 0 0	<pre>svals(npsing) tbl(ldtbl,3)</pre>	<pre>nuobs+nnull singular values, svals must have a dimension, of at least nuobs-nnull. if info indicates nonzero info in dsvdc then svals is as returned from dsvdc. column contains 1 grid of log10(nobs*lambda) 2 U(lertic) </pre>
0 0 0 0 0 0	svals (npsing) tbl (ldtbl, 3)	<pre>nuobs+nnull singular values, svals must have a dimension, of at least nuobs-nnull. if info indicates nonzero info in dsvdc then svals is as returned from dsvdc. column contains 1 grid of log10(nobs*lambda) 2 V(lambda) 2 D(lambda) 3 D(lambda)</pre>
0 0 0 0 0 0 0	<pre>svals(npsing) tbl(ldtbl,3)</pre>	<pre>nuobs+nnull singular values, svals must have a dimension, of at least nuobs-nnull. if info indicates nonzero info in dsvdc then svals is as returned from dsvdc. column contains 1 grid of log10(nobs*lambda) 2 V(lambda) 3 R(lambda) if requested</pre>

```
1st row contains:
с
                             log10(nobs*lamhat), V(lamhat) and
С
                             R(lamhat) if requested
с
                             where lamhat is the gcv estimate of lambda
с
                         2nd row contains:
с
                             0, V(0) and R(0) if requested
с
                         3rd row contains:
\sim
с
                             0, V(infinity) and R(infinity) if requested
    info
                         error indicator
с
                            0 : successful completion
с
                           -1 : log10(nobs*lamhat) <= lamlim(1)
с
                                 (not fatal)
с
                           -2 : log10(nobs*lamhat) >= lamlim(2)
с
                                 (not fatal)
с
с
                            1 : dimension error
с
                            2 : error in dreps, the first ncov1 columns
                                 of s do not duplicate the replication
С
С
                                 structure of des
                            3 : lwa (length of work) is too small
4 : liwa (length of iwork) is too small
С
с
                            5 : error in dmaket
с
                            6 : sigma is rank deficient
с
с
                           1000< info : 1000 + nonzero info returned from
                                         dsnsm
С
c
c Working Storage:
с
    work(lwa)
                         double precision work vector
                         length of work as declared in the calling
    lwa
с
с
                         program
С
                         must be at least 1wa1 + 1wa2 where
                           lwa1 = (nnull-ncov2)*(nobs+nuobs+1)
c
                                   +npar*(nobs+npar)
с
с
                           lwa2 = (npar-nnull)*(npar-2*nnull+2+nobs)
                                   +npar+nobs
с
с
¢
с
    iwork(liwa)
                         integer work vector
                         length of the iwork as declared in the calling
с
    liwa
                         program
С
                         must be at least 3*nobs - (nnull - ncov2)
с
с
c Subprograms Called Directly:
        Gcvpack - dreps dmaket duni dmakek dctsx dsnsm
С
        Linpack - dqrdc dqrsl
С
                - dcopy
С
        Blas
        Other - dprmut dset prmut mkpoly
с
С
c Subprograms Called Indirectly:
        Gevpack - dertz ddcom dgev dsgde dtsvde drsap ddiag
С
                   dvlop dvlop dpmse dcfcr dpdcr dvmin dvl dzdc
Ċ
        Linpack - dchdc dqrdc dqrsl dtrsl dsvdc dtrco
Blas - dcopy ddot dgemv dswap
С
С
        Other - dcpmut dprmut dset dftkf fact mkpoly
Ċ
С
************
dsnsm.com
......
      subroutine dsnsm (x,ldx,y,sigma,ldsigm,nobs,npar,nnull,adiag,
     * tau, lamlim, ntbl, dout, iout, coef, svals, tbl, ldtbl, auxtbl,
     * iwork,liwa,work,lwa,job,info)
      integer ldx,ldsigm,nobs,npar,nnull,ntbl,iout(3),ldtbl,liwa,
     * iwork(liwa),lwa,job,info
      double precision x(ldx,npar),y(nobs),sigma(ldsigm,npar),
     * adiag(nobs),tau,lamlim(2),dout(5),coef(npar),svals(*),
     * tbl(ldtbl,3),auxtbl(3,3),work(lwa)
```

¢		
С	Purpose: determine th	e generalized cross validation estimate of the
c	smoothing param	eter and fit model parameters for a penalized
c	least squares p	problem with a semi-norm smoothing matrix.
С		
С	On Entry:	
с	x(ldx,npar)	design matrix
С	ldx	leading dimension of x as declared in the
с		calling program, must be at least max(nobs,npar)
с	y(nobs)	response vector
с	sigma(ldsigm, npar)	symmetric matrix that defines the semi-norm
с	ldsigm	leading dimension of sigma as declared
ċ		in the calling program
č	nobs	number of observations
õ	ntar	number of parameters
č	npull	dimension of the null snace of sigma
~	adiag (pobe)	"true" y values on entry if computation of
с ~	actag (nobs)	predictive mee is requested
0	1 - 1 + (0)	limite en lambda het goarch (in log10/nobst
c	LamLIM(2)	limits on fambda nat search (in fogio (nobs
C		Tambda) Scale, II user imput Timits are
с		requested if $Iamim(I) = Iamim(Z)$ then $Iamma$
c		is set to (IU^*IamIIm(I))/nobs
c	tau	multiplier controlling the amount of truncation
С		if truncation is requested (try tau = 1
С		to start then try 10 and 100)
с	ntbl	number of evenly spaced values for
С		log10 (nobs*lambda) to be used in the initial
С		grid search for lambda hat
с		<pre>if ntbl = 0 only a golden ratio search will be</pre>
с		done and tbl is not referenced, if $ntbl > 0$
¢		there will be ntbl rows returned in tbl
с	ldtbl	leading dimension of tbl as declared in the
с		calling program
с	job	integer with decimal expansion abcd
с		if a is nonzero then truncation is used
с		if b is nonzero then predictive mse is computed
с		using adiag as true y
с		if c is nonzero then user input limits on search
с		for lambda hat are used
с		if d is nonzero then the diagonal of the hat
с		matrix is calculated
с		
c	On Exit:	
c	x(ldx.npar)	overwritten with many intermediate results
č	v(nobs)	predicted values
č	sigma (ldsigm.npar)	overwritten with the OR decomposition of the
č	erding (rabright, hpur)	Cholesky factor of sigma
č	adiag (nobs)	diagonal elements of the bat matrix if requested
č	lamlim (2)	limits on lambda hat search
~	2000210(2)	(in log10/nobstlambda) scale)
č	dout (5)	(in logic(nobs lambda) scale)
с с	douc (5)	l lambat concerning and charge we lidetion
ن م		I Tanular generalized cross validation
6		estimate of the smoothing parameter
6		2 penity smoothing penalty
6		5 rss residual sum of squares
c		4 $\text{tr}(1-A)$ trace of $1 - A$
c		<pre>> truncation ratio = 1/(1+(normk/(nobs*lamhat)))</pre>
c		where norm $k = norm(R - R sub k) **2$
с	10ut (3)	contains:
C		I npsing number of positive singular
с		values
c		if info indicates nonzero info in

dsvdc then iout(1) contains info as с it was returned from dsvdc С ¢ 2 npar number of parameters 3 nnull size of the null space of sigma С coefficient estimates С coef (npar) svals (npar-nnull) first npsing entries contain singular values С of the matrix j2 с if info indicates nonzero info in dsvdc then с svals is as it was returned from dsvdc С tbl(ldtbl,3) column contains с с 1 grid of log10 (nobs*lambda) 2 V(lambda) С R(lambda) if requested с 3 auxtbl(3,3) auxiliary table С 1st row contains: С log10(nobs*lamhat), V(lamhat) and С R(lamhat) if requested С where lamhat is the gcv estimate of lambda С с 2nd row contains: 0, V(0) and R(0) if requested с с 3rd row contains: 0, V(infinity) and R(infinity) if requested С с info error indicator 0 : successful completion С -3 : nnull is too small (not fatal) с -2 : log10(nobs*lamhat) >= lamlim(2) с (not fatal) С -1 : log10(nobs*lamhat) <= lamlim(1)</pre> с с (not fatal) c 1 : dimension error 2 : 1wa (length of work) is too small С С 3 : liwa (length of iwork) is too small c 4 : error in ntbl or tau 100< info <200 : 100 + nonzero info returned С from ddcom с 200< info <300 : 200 + nonzero info returned с с from dgcv с c Work Arravs: с work(lwa) double precision work vector length of work as declared in the calling С lwa С program с must be at least (npar-nnull) * (npar-2*nnull+2+nobs)+npar+nobs С iwork(liwa) С integer work vector ċ liwa length of iwork as declared in the calling С program С must be at least 2*npar - nnull С c Subprograms Called Directly: Gevpack - ddcom dgev С C c Subprograms Called Indirectly: Gevpack - dertz dsgde defer drsap dvlop dtsvde С с dpmse dvmin dvl dzdc dpdcr ddiag Linpack - dchdc dqrdc dqrsl dtrsl dsvdc dtrco ¢ - dcopy ddot dgemv dswap С Blas С Other - dcpmut dprmut dset ¢ ************ dtpss.com

```
subroutine dtpss(des,lddes,nobs,dim,m,s,lds,ncov,y,ntbl,adiag,
      * lamlim, dout, iout, coef, svals, tbl, ldtbl, auxtbl, work, lwa,
     * iwork,liwa,job,info)
      integer lddes, nobs, dim, m, lds, ncov, ntbl, iout (4), ldtbl, lwa,
     * liwa, iwork (liwa), job, info
      double precision des(lddes,dim),s(lds,*),y(nobs),
      * adiag(nobs),lamlim(2),dout(5),coef(*),svals(*),
     * tbl(ldtbl,3),auxtbl(3,3),work(lwa)
с
c Purpose: determine the generalized cross validation estimate of the
        smoothing parameter and fit model parameters for a thin plate
с
        smoothing spline.
С
С
c On Entry:
    des(lddes,dim)
                         design for the variables to be splined
С
c
    lddes
                         leading dimension of des as declared in calling
с
                         program
                         number of observations
    nobs
с
                         number of columns in des
    dim
С
                         order of the derivatives in the penalty
С
    m
    s(lds,ncov)
                         design for the covariates. The covariates
С
                         must duplicate the replication structure of des.
¢
с
                         See dptpss to handle covariates which do not.
    lds
                         leading dimension of s as declared in calling
С
                         program
с
                         number of covariates
    ncov
С
                         response vector
с
    y(nobs)
                         number of evenly spaced values for
    ntbl
С
                         log10(nobs*lambda) to be used in the initial
с
с
                         grid search for lambda hat
                         if ntbl = 0 only a golden ratio search will be
С
с
                         done and tbl is not referenced, if ntbl > 0
с
                         there will be ntbl rows returned in tbl
С
    adiag(nobs)
                         "true" y values on entry if predictive mse is
С
                         requested
    lamlim(2)
                         limits on lambda hat search (in log10 (nobs*
С
С
                         lambda) scale) if user input limits are
с
                         requested if lamlim(1) = lamlim(2) then lamhat
                         is set to (10**lamlim(1))/nobs
с
с
    ldtbl
                         leading dimension of tbl as declared in the
с
                         calling program
с
    job
                         integer with decimal expansion abdc
с
                         if a is nonzero then predictive mse is computed
С
                            using adiag as true y
                         if b is nonzero then user input limits on search
¢
                            for lambda hat are used
с
с
                         if c is nonzero then adiag will be calculated
с
                         if d is nonzero then there are replicates in the
с
                            design
С
c On Exit:
с
    des(lddes,dim)
                        sorted unique rows of des if job indicates that
                        there are replicates otherwise not changed
с
С
    s(lds,ncov)
                        unique rows of s sorted to correspond to des
с
    v(nobs)
                        predicted values
    adiag(nobs)
                        diagonal elements of the hat matrix if requested
С
c
    lamlim(2)
                        limits on lambda hat search
                        (in log10(nobs*lambda) scale)
¢
    dout (5)
с
                        contains:
¢
                        1 lamhat
                                     generalized cross validation
                                     estimate of the smoothing parameter
С
с
                        2 penlty
                                   smoothing penalty
```

с		3 rss residual sum of squares
c		4 tr(I-A) trace of I - A 5 segren sum of squares for replication
с с	iout (4)	contains:
c	1000 (4)	1 npsing number of positive singular
c		values (npsing = nuobs - ncts).
c		if info indicates nonzero info in
с		dsvdc then npsing contains info as
¢		it was returned from dsvdc.
с		2 npar number of parameters
с		(npar = nuobs + ncts)
С		3 ncts dimension of the polynomial space
с		plus ncov
ų		((m+dim-1 choose dim) + ncov)
c	- · · ·	4 nuobs number of unique rows in des
с	coef (npar)	coefficient estimatés [beta':alpha':delta']
c		coef must have a dimension of at least nuobs+
c		ncts
c	svais (npar-nnuil)	singular values of the matrix j2 if into = 0
c		avala is as it was returned from davdo
с с	+ b1 (1d+b1 3)	column contains
ĉ	CD1 (Ideb1, 5)	1 grid of log10(nobs*lambda)
ĉ		2 V(lambda)
c		3 R(lambda) if requested
c	auxtbl(3.3)	auxiliary table
c		lst row contains:
с		log10(nobs*lamhat), V(lamhat) and
с		R(lamhat) if requested
c		where lamhat is the gcv estimate of lambda
С		2nd row contains:
c		0, V(0) and R(0) if requested
c		3rd row contains:
с		0, V(infinity) and R(infinity) if requested
c	info	error indicator
c		U : Successful completion
c		-1 : logiu(nobs*lamnat) <= lamlim(l)
C		(not istal) -2 : log10(nobstlombat) >= lomlim(2)
0		-2 : IOGIU(NOBS^IAMMAC) \geq IAMIIM(2) (not fatal)
c		1 : dimension error
č		2 : error in dreps, covariates do not
c		duplicate the replication structure of des
c		3 : 1wa (length of work) is too small
с		4 : liwa (length of iwork) is too small
с		10 < info < 20 : 10 + nonzero info returned
с		from dsetup
С		100< info <200 : 100 + nonzero info returned
с		from dsgdc1
c		200< info <300 : 200 + nonzero info returned
с		from dgcvl
°.		
c i	work Arrays:	double presiden work wester
c	WOLK (IWA)	double precision work vector
c	⊥₩a	rengen of work as declared in the calling
c		Must be at least nucls(2+ncts+nucls)+nchs
c	iwork(liwa)	integer work vector
č	liwa	length of iwork as declared in the calling
c		program
c		Must be at least 2*nobs + nuobs - ncts
c		

```
c Subprograms Called Directly:
c Gcvpack - dreps duni dsuy dsetup dsgdcl dgcvl
c Other - dprmut mkpoly
c
c Subprograms Called Indirectly:
c Gcvpack - dcfcrl drsap dvlop dsvtc dpdcr dpmse
c dvmin dvl dmaket dmakek ddiag
c Linpack - dchdc dqrdc dqrsl dtrsl dsvdc
c Blas - ddot dcopy dgemv
c Other - dprmut dset dftkf fact mkpoly
c
```