

# Lecture 7: Splines and Generalized Additive Models

## Computational Statistics

Thierry Denœux

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# Overview

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# Moving beyond linearity

- Linear models are widely used in econometrics.
- In particular, linear regression, linear discriminant analysis, logistic regression all rely on a linear model.
- It is extremely unlikely that the true function  $f(X)$  is actually linear in  $X$ . In regression problems,  $f(X) = \mathbb{E}(Y|X)$  will typically be nonlinear and nonadditive in  $X$ , and representing  $f(X)$  by a linear model is usually a convenient, and sometimes a necessary, approximation.
  - Convenient because a linear model is easy to interpret, and is the first-order Taylor approximation to  $f(X)$ .
  - Sometimes necessary, because with  $N$  small and/or  $p$  large, a linear model might be all we are able to fit to the data without overfitting.
- Likewise in classification, it is usually assumed that some monotone transformation of  $\mathbb{P}(Y = 1|X)$  is linear in  $X$ . This is inevitably an approximation.



# Linear basis expansion

- The core idea in this chapter is to augment/replace the vector of inputs  $X$  with additional variables, which are transformations of  $X$ , and then use linear models in this new space of derived input features.
- Denote by  $h_m(X) : \mathbb{R}^p \rightarrow \mathbb{R}$  the  $m$ -th transformation of  $X$ ,  $m = 1, \dots, M$ . We then model

$$f(X) = \sum_{m=1}^M \beta_m h_m(X)$$

a linear basis expansion in  $X$ .

- The beauty of this approach is that once the basis functions  $h_m$  have been determined, the models are linear in these new variables, and the fitting proceeds as for linear models.



## Popular choices for basis functions $h_m$

Some simple and widely used examples of the  $h_m$  are the following:

- $h_m(X) = X_m$ ,  $m = 1, \dots, p$  recovers the original linear model.
- $h_m(X) = X_j^2$  or  $h_m(X) = X_j X_k$  allows us to augment the inputs with polynomial terms to achieve higher-order Taylor expansions. Note, however, that the number of variables grows exponentially in the degree of the polynomial. A full quadratic model in  $p$  variables requires  $O(p^2)$  square and cross-product terms, or more generally  $O(p^d)$  for a degree- $d$  polynomial.
- $h_m(X) = \log(X_j)$ ,  $\sqrt{X_j}$ , ... permits other nonlinear transformations of single inputs. More generally one can use similar functions involving several inputs, such as  $h_m(X) = \|X\|$ .
- $h_m(X) = I(L_m \leq X_k < U_m)$ , an indicator for a region of  $X_k$ . By breaking the range of  $X_k$  up into  $M_k$  such nonoverlapping regions results in a model with a piecewise constant contribution for  $X_k$ .



# Discussion

- Sometimes the problem at hand will call for particular basis functions  $h_m$ , such as logarithms or power functions.
- More often, however, we use the basis expansions as a device to achieve more flexible representations for  $f(X)$ .
- Polynomials are an example of the latter, although they are limited by their global nature – tweaking the coefficients to achieve a functional form in one region can cause the function to flap about madly in remote regions.



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# Fitting polynomials

- In most of this lecture, we assume  $p = 1$ .
- Create new variables  $h_1(X) = X$ ,  $h_2(X) = X^2$ ,  $h_3(X) = X^3$ , etc. and then do multiple linear regression on the transformed variables.
- We either fix the degree  $d$  at some reasonably low value, else use cross-validation to choose  $d$ .
- Polynomials have unpredictable tail behavior – very bad for extrapolation.



# Example in R

```

x=seq(0,10,0.5)
n<-length(x)
y1=x[1:10]+2*cos(x[1:10])+2*rnorm(10)
xtest<- seq(-2,12,0.01)
ftest<- xtest+2*cos(xtest)
d<-3

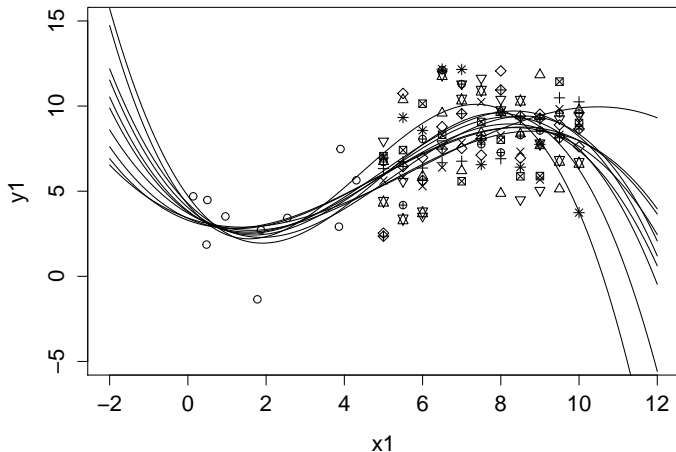
plot(x1,y1,xlim=c(-2,12),ylim=c(-5,15),
      main=paste('degree = ',as.character(d)))
for(i in 1:10){
  y2=x[11:21]+2*cos(x[11:21])+2*rnorm(11)
  points(x[11:21],y2,pch=i+1)
  y<-c(y1,y2)
  reg<-lm(y ~ poly(x,degree=d))
  ypred<-predict(reg,newdata=data.frame(x=xtest),interval="c")
  lines(xtest,ypred[, "fit"],lty=1)
}

```



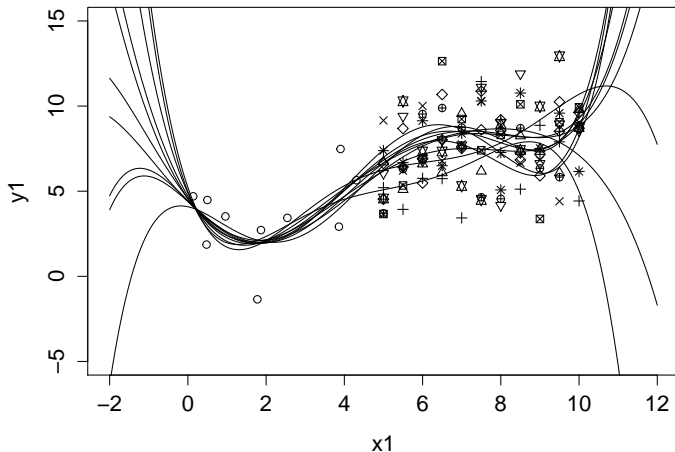
Result,  $d = 2$ 

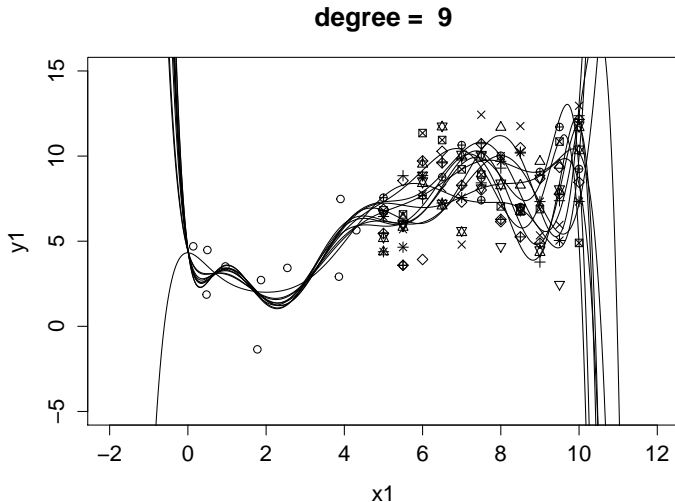
degree = 3



Result,  $d = 3$ 

degree = 5



Result,  $d = 4$ 

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# Step Functions

- Another way of creating transformations of a variable is to cut the variable into distinct regions.

$$h_1(X) = I(X < \xi_1), h_2(X) = I(\xi_1 \leq X < \xi_2), \dots, \\ h_M(X) = I(X \geq \xi_{M-1})$$

- Since the basis functions are positive over disjoint regions, the least squares estimate of the model  $f(X) = \sum_{m=1}^M \beta_m h_m(X)$  is  $\hat{\beta}_m = \bar{Y}_m$ , the mean of  $Y$  in the  $m$ -th region.



# Example in R

```
library("ISLR")

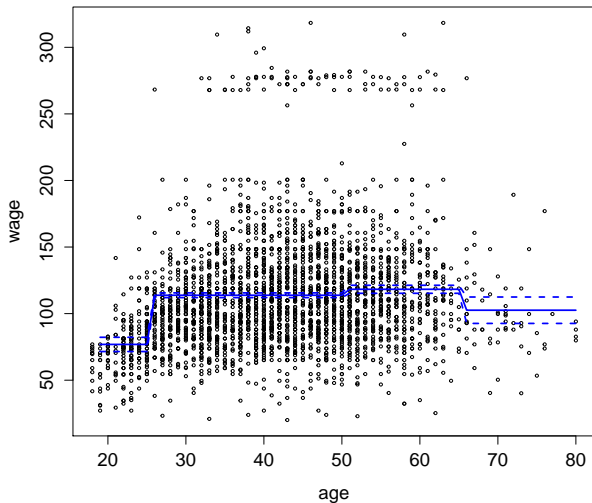
reg<-lm(wage ~ cut(age, c(18, 25, 50, 65, 90)),data=Wage)
ypred<-predict(reg,newdata=data.frame(age=18:80),interval="c")

plot(Wage$age,Wage$wage,cex=0.5,xlab="age",ylab="wage")
lines(18:80,ypred[, "fit"],lty=1,col="blue",lwd=2)
lines(18:80,ypred[, "lwr"],lty=2,col="blue",lwd=2)
lines(18:80,ypred[, "upr"],lty=2,col="blue",lwd=2)
```





## Result



## Step functions – continued

- Easy to work with. Creates a series of dummy variables representing each group.
- Useful way of creating interactions that are easy to interpret. For example, interaction effect of Year and Age:

$$I(\text{Year} < 2005) \cdot \text{Age}, I(\text{Year} \geq 2005) \cdot \text{Age}$$

would allow for different linear functions in each age category.

- Choice of cutpoints or knots can be problematic. For creating nonlinearities, smoother alternatives such as splines are available.



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# Piecewise Polynomials

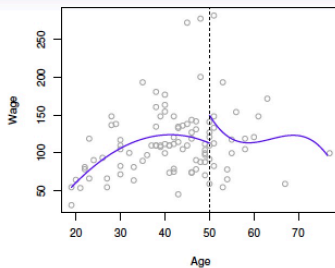
- Instead of a single polynomial in  $X$  over its whole domain, we can rather use different polynomials in regions defined by knots. E.g. (see figure)

$$y_i = \begin{cases} \beta_{01} + \beta_{11}x_i + \beta_{21}x_i^2 + \beta_{31}x_i^3 + \epsilon_i & \text{if } x_i < \xi, \\ \beta_{02} + \beta_{12}x_i + \beta_{22}x_i^2 + \beta_{32}x_i^3 + \epsilon_i & \text{if } x_i \geq \xi, \end{cases}$$

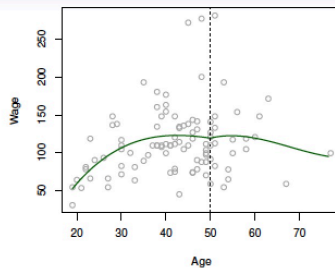
- Better to add constraints to the polynomials, e.g. continuity.
- Splines have the “maximum” amount of continuity.



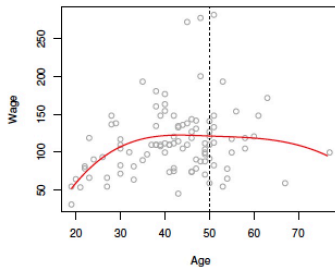
Piecewise Cubic



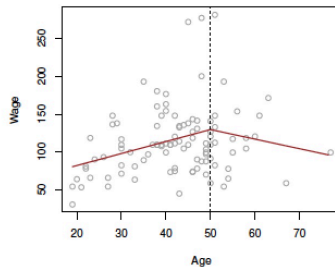
Continuous Piecewise Cubic



Cubic Spline



Linear Spline



# Linear Splines

- A linear spline with knots at  $\xi_k$ ,  $k = 1, \dots, K$  is a piecewise linear polynomial continuous at each knot.
- The set of linear splines with fixed knots is a vector space.
- The number of degrees of freedom is  $2(K + 1) - K = K + 2$ . We can thus decompose linear splines on a basis of  $K + 2$  basis functions,

$$y = \sum_{m=1}^{K+2} \beta_m h_m(x) + \epsilon.$$

- The basis functions can be chosen as

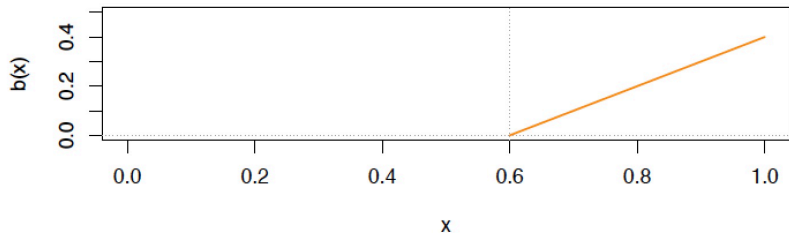
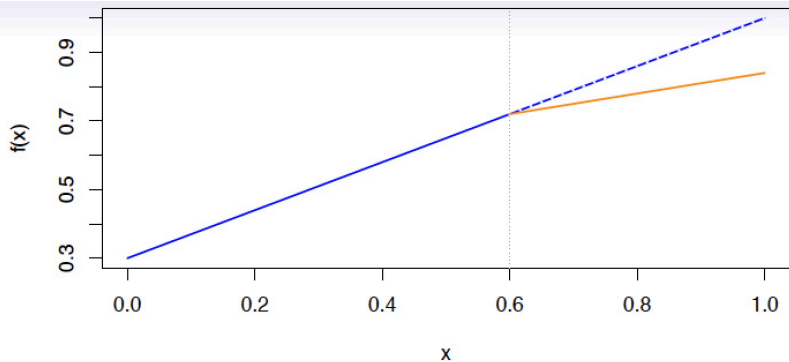
$$h_1(x) = 1$$

$$h_2(x) = x$$

$$h_{k+2}(x) = (x - \xi_k)_+, \quad k = 1, \dots, K,$$

where  $(\cdot)_+$  denotes the positive part, i.e.,  $(x - \xi_k)_+ = x - \xi_k$  if  $x > \xi_k$  and  $(x - \xi_k)_+ = 0$  otherwise.







# Cubic Splines

- A cubic spline with knots at  $\xi_k$ ,  $k = 1, \dots, K$  is a piecewise cubic polynomial with continuous derivatives up to order 2 at each knot.
- Enforcing one more order of continuity would lead to a global cubic polynomial.
- Again, the set of cubic splines with fixed knots is a vector space, and the number of degrees of freedom is  $4(K + 1) - 3K = K + 4$ . We can thus decompose cubic splines on a basis of  $K + 4$  basis functions,

$$y = \sum_{m=1}^{K+4} \beta_m h_m(x) + \epsilon.$$

- We can choose truncated power basis functions,

$$\begin{aligned} h_k(x) &= x^{k-1}, \quad k = 1, \dots, 4, \\ h_{k+4}(x) &= (x - \xi_k)_+^3, \quad k = 1, \dots, K. \end{aligned}$$



## order- $M$ splines

- More generally, an order- $M$  spline with knots  $\xi_k$ ,  $k = 1, \dots, K$  is a piecewise-polynomial of order  $M - 1$ , which has continuous derivatives up to order  $M - 2$ .
- A cubic spline has  $M = 4$ . A piecewise-constant function is an order-1 spline, while a continuous piecewise linear function is an order-2 spline.
- The general form for the truncated-power basis set is

$$h_k(x) = x^{k-1}, \quad k = 1, \dots, M,$$
$$h_{k+M}(x) = (x - \xi_k)_+^{M-1}, \quad k = 1, \dots, K.$$

- It is claimed that cubic splines are the lowest-order spline for which the knot-discontinuity is not visible to the human eye. There is seldom any good reason to go beyond cubic-splines.
- In practice the most widely used orders are  $M = 1, 2$  and  $4$ .



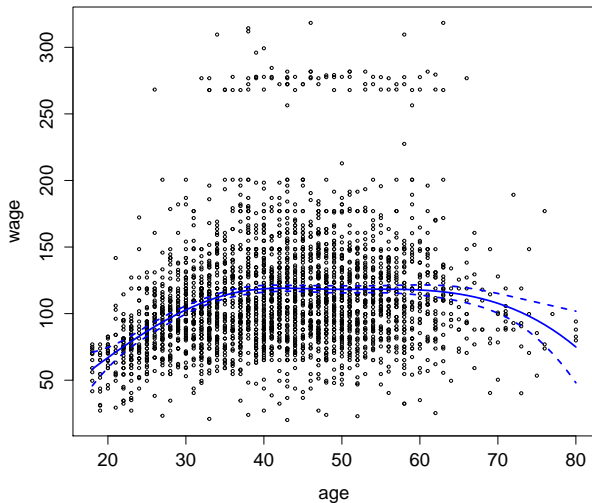
# Splines in R

```
library('splines')  
fit<-lm(wage~bs(age,5),data=Wage)  
  
ypred<-predict(fit,newdata=data.frame(age=18:80),interval="c")  
  
plot(Wage$age,Wage$wage,cex=0.5,xlab="age",ylab="wage")  
lines(18:80,ypred[, "fit"],lty=1,col="blue",lwd=2)  
lines(18:80,ypred[, "lwr"],lty=2,col="blue",lwd=2)  
lines(18:80,ypred[, "upr"],lty=2,col="blue",lwd=2)
```

- By default, degree=3, and the intercept is not included in the basis functions.
- The number of knots is df-degree. If not specified, the knots are placed at quantiles.



## Result



# B-spline basis

- Since the space of spline functions of a particular order and knot sequence is a vector space, there are many equivalent bases for representing them (just as there are for ordinary polynomials.)
- While the truncated power basis is conceptually simple, it is not too attractive numerically: powers of large numbers can lead to severe rounding problems.
- In practice, we often use another basis: the B-spline basis, which allows for efficient computations even when the number of knots  $K$  is large (each basis function has a local support).



# B-spline basis

## Construction

- Before we can get started, we need to augment the knot sequence.
- Let  $\xi_0 < \xi_1$  and  $\xi_K < \xi_{K+1}$  be two boundary knots, which typically define the domain over which we wish to evaluate our spline. We now define the augmented knot sequence  $\tau$  such that
  - $\tau_1 \leq \tau_2 \leq \dots \leq \tau_M \leq \xi_0$
  - $\tau_{j+M} = \xi_j, j = 1, \dots, K$
  - $\xi_{K+1} \leq \tau_{K+M+1} \leq \tau_{K+M+2} \leq \dots \leq \tau_{K+2M}$ .
- The actual values of these additional knots beyond the boundary are arbitrary, and it is customary to make them all the same and equal to  $\xi_0$  and  $\xi_{K+1}$ , respectively.



## B-spline basis

## Construction – Continued

- Denote by  $B_{i,m}(x)$  the  $i$ th B-spline basis function of order  $m$  for the knot-sequence  $\tau$ ,  $m \leq M$ . They are defined recursively in terms of divided differences as follows:

$$B_{i,1}(x) = \begin{cases} 1 & \text{if } \tau_i \leq x < \tau_{i+1} \\ 0 & \text{otherwise} \end{cases}$$

for  $i = 1, \dots, K + 2M - 1$ . (By convention,  $B_{i,1} = 0$  if  $\tau_i = \tau_{i+1}$ ).

$$B_{i,m} = \frac{x - \tau_i}{\tau_{i+m-1} - \tau_i} B_{i,m-1}(x) + \frac{\tau_{i+m} - x}{\tau_{i+m} - \tau_{i+1}} B_{i+1,m-1}(x)$$

for  $i = 1, \dots, K + 2M - m$ .

- Thus with  $M = 4$ ,  $B_{i,4}$ ,  $i = 1, \dots, K + 4$  are the  $K + 4$  cubic B-spline basis functions for the knot sequence  $\xi$ .



# B-spline basis

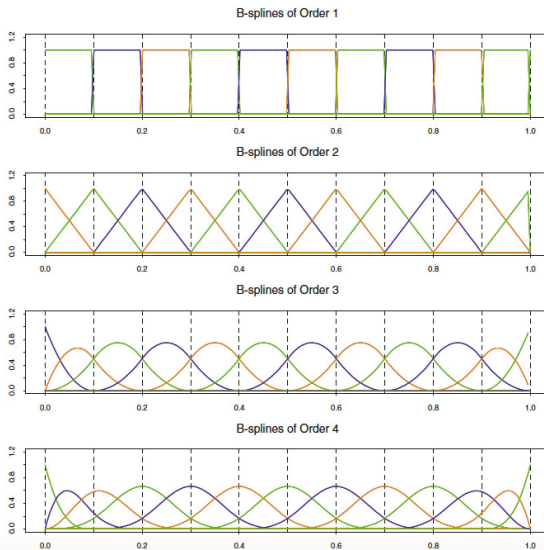
## Properties

- The B-splines span the space of cubic splines for the knot sequence  $\xi$ .
- They have local support and they are nonzero on an interval spanned by  $M + 1$  knots (see next slide).





## Sequence of B-splines up to order 4 with 10 knots evenly spaced from 0 to 1



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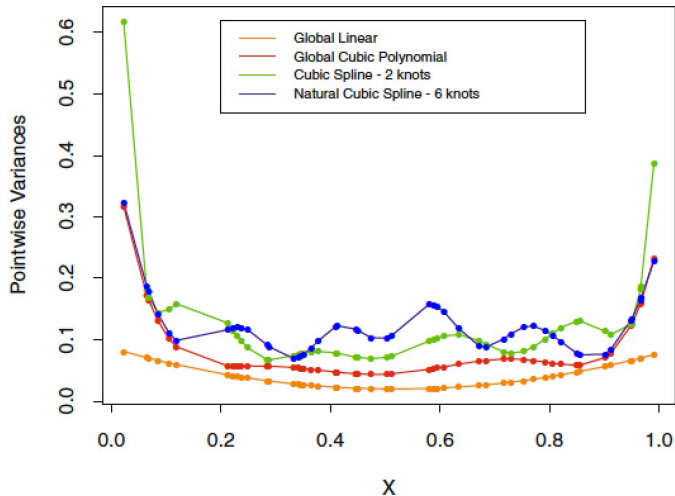


# Variance of splines beyond the boundary knots

- We know that the behavior of polynomials fit to data tends to be erratic near the boundaries, and extrapolation can be dangerous.
- These problems are exacerbated with splines. The polynomials fit beyond the boundary knots behave even more wildly than the corresponding global polynomials in that region.



## Example



# Explanation of the previous figure

- Pointwise variance curves for four different models, with  $X$  consisting of 50 points drawn at random from  $U[0, 1]$ , and an assumed error model with constant variance.
- The linear and cubic polynomial fits have 2 and 4 df, respectively, while the cubic spline and natural cubic spline each have 6 df.
- The cubic spline has two knots at 0.33 and 0.66, while the natural spline has boundary knots at 0.1 and 0.9, and four interior knots uniformly spaced between them.

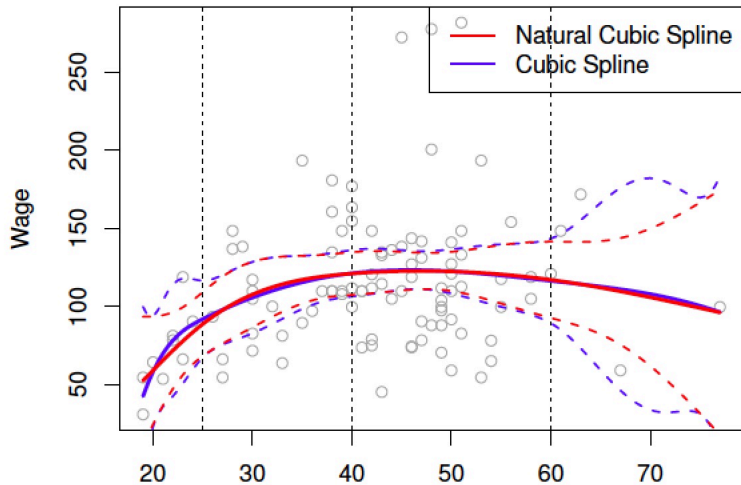


# Natural cubic spline

- A natural cubic spline adds additional constraints, namely that the function is linear beyond the boundary knots.
- This frees up four degrees of freedom (two constraints each in both boundary regions), which can be spent more profitably by putting more knots in the interior region.
- There will be a price paid in bias near the boundaries, but assuming the function is linear near the boundaries (where we have less information anyway) is often considered reasonable.



# Example



# Natural cubic spline basis

- A natural cubic spline with  $K$  knots has  $K$  degrees of freedom: it can be represented by  $K$  basis functions.
- One can start from a basis for cubic splines, and derive the reduced basis by imposing the boundary constraints. For example, starting from the truncated power series basis,

$$f(X) = \sum_{j=0}^3 \beta_j X^j + \sum_{k=1}^K \theta_k (X - \xi_k)_+^3,$$

the constraints  $f''(X) = 0$  and  $f^{(3)}(X) = 0$  for  $X < \xi_1$  and  $X > \xi_K$  lead to the conditions

$$\beta_2 = \beta_3 = 0, \quad \sum_{k=1}^K \theta_k = 0, \quad \sum_{k=1}^K \xi_k \theta_k = 0$$





# Natural cubic spline basis – continued

- These conditions are automatically satisfied by choosing the following basis,

$$N_1(X) = 1, \quad N_2(X) = X,$$

$$N_{k+2}(X) = d_k(X) - d_{K-1}(X), \quad k = 1, \dots, K-2$$

with

$$d_k = \frac{(X - \xi_k)_+^3 - (X - \xi_K)_+^3}{\xi_K - \xi_k}$$



# Example in R

```
fit1<-lm(y ~ ns(x,df=5))
```

```
fit2<-lm(y ~ bs(x,df=5))
```

```
ypred1<-predict(fit1,newdata=data.frame(x=xtest),interval="c")
```

```
ypred2<-predict(fit2,newdata=data.frame(x=xtest),interval="c")
```

```
plot(x,y,xlim=range(xtest))
```

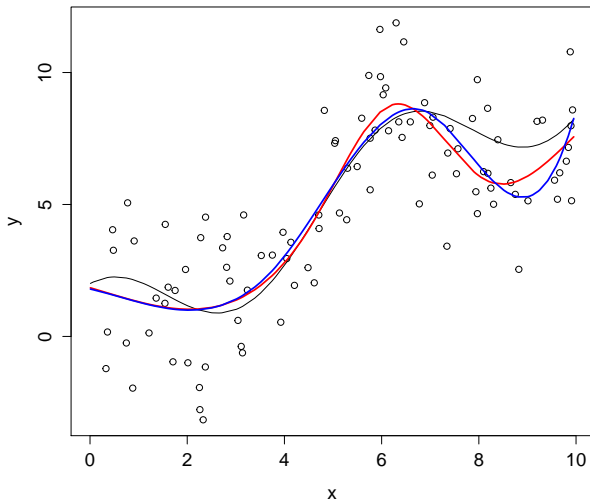
```
lines(xtest,ftest)
```

```
lines(xtest,ypred1[, "fit"],lty=1,col="red",lwd=2)
```

```
lines(xtest,ypred2[, "fit"],lty=1,col="blue",lwd=2)
```



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# Using splines with logistic regression

- Until now, we have discussed regression problems. However, splines can also be used when the response variable is qualitative.
- Consider, for instance, natural splines with  $K$  knots. For binary classification, we can fit the logistic regression model,

$$\log \frac{\mathbb{P}(Y = 1 | X = x)}{\mathbb{P}(Y = 0 | X = x)} = f(x)$$

with  $f(x) = \sum_{k=1}^K \beta_k N_k(x)$ .

- Once the basis functions have been defined, we just need to estimate coefficients  $\beta_k$  using a standard logistic regression procedure.
- A smooth estimate of the conditional probability  $\mathbb{P}(Y = 1|x)$  can then be used for classification or risk scoring.



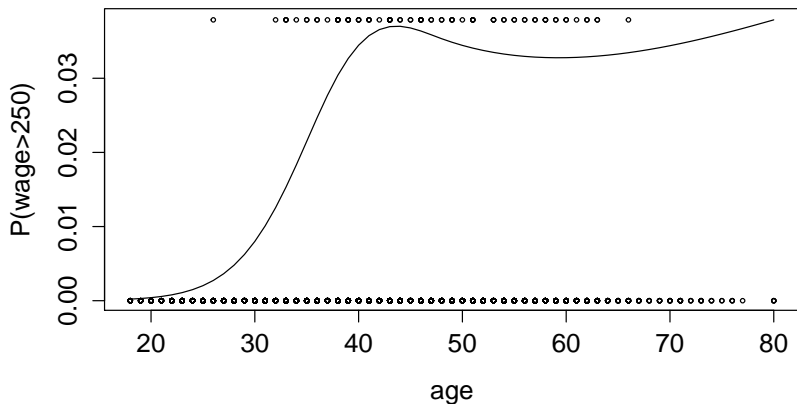
# Example in R

```
class<-glm(I(wage>250) ~ ns(age,3),data=Wage,family='binomial')
proba<-predict(class,newdata=data.frame(age=18:80),type='response')

plot(18:80,proba,xlab="age",ylab="P(wage>250)",type="l")
ii<-which(Wage$age>250)
points(Wage$age[ii],rep(max(proba),length(ii)),cex=0.5)
points(Wage$age[-ii],rep(0,nrow(Wage)-length(ii)),cex=0.5)
```



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# Smoothing splines

## Problem formulation

- Here we discuss a spline basis method that avoids the knot selection problem completely by using a maximal set of knots. The complexity of the fit is controlled by regularization.
- Problem: among all functions  $f(x)$  with two continuous derivatives, find one that minimizes the penalized residual sum of squares

$$RSS(f, \lambda) = \sum_{i=1}^N (y_i - f(x_i))^2 + \lambda \int [f''(t)]^2 dt,$$

where  $\lambda$  is a fixed smoothing parameter.

- The first term measures closeness to the data, while the second term penalizes curvature in the function, and  $\lambda$  establishes a tradeoff between the two. Special cases:  $\lambda = 0$  (no constraint on  $f$ ) and  $\lambda = \infty$  ( $f$  has to be linear).



# Smoothing splines

## Solution

- It can be shown that this problem has an explicit, finite-dimensional, unique minimizer which is a natural cubic spline with knots at the unique values of the  $x_i, i = 1, \dots, N$ .
- At face value it seems that the family is still over-parametrized, since there are as many as  $N$  knots, which implies  $N$  degrees of freedom. However, the penalty term translates to a penalty on the spline coefficients, which are shrunk some of the way toward the linear fit.
- The solution is thus of the form

$$f(x) = \sum_{j=1}^N N_j(x) \theta_j,$$

where the  $N_j(x)$  are an  $N$ -dimensional set of basis functions for representing this family of natural splines.



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# Computation

- The criterion can be written as

$$RSS(\theta, \lambda) = (\mathbf{y} - \mathbf{N}\theta)^T(\mathbf{y} - \mathbf{N}\theta) + \lambda\theta^T\mathbf{\Omega}_N\theta,$$

where  $\{\mathbf{N}\}_{ij} = N_j(x_i)$  and  $\{\mathbf{\Omega}_N\}_{jk} = \int N_j''(t)N_k''(t)dt$ .

- The solution is

$$\hat{\theta} = (\mathbf{N}^T\mathbf{N} + \lambda\mathbf{\Omega}_N)^{-1}\mathbf{N}^T\mathbf{y},$$

a generalized ridge regression.

- The fitted smoothing spline is given by

$$\hat{f}(x) = \sum_{j=1}^N N_j(x)\hat{\theta}_j.$$

- In practice, when  $N$  is large, we can use only a subset of the  $N$  interior knots (rule of thumb: number of knots proportional to  $\log N$ ).



# Degrees of freedom

- Denote by  $\hat{\mathbf{f}}$  the  $N$ -vector of fitted values  $f(x_i)$  at the training predictors  $x_i$ . Then,

$$\hat{\mathbf{f}} = \mathbf{N}\hat{\boldsymbol{\theta}} = (\mathbf{N}^T \mathbf{N} + \lambda \boldsymbol{\Omega}_N)^{-1} \mathbf{N}^T \mathbf{y} = \mathbf{S}_\lambda \mathbf{y}$$

- As matrix  $\mathbf{S}_\lambda$  does not depend on  $\mathbf{y}$ , the smoothing spline is a linear smoother.
- In the case of cubic spline with knot sequence  $\xi$  and, we have

$$\hat{\mathbf{f}} = \mathbf{B}_\xi \hat{\boldsymbol{\theta}} = (\mathbf{B}_\xi^T \mathbf{B}_\xi)^{-1} \mathbf{B}_\xi^T \mathbf{y} = \mathbf{H}_\xi \mathbf{y},$$

where  $\mathbf{B}_\xi$  is the  $N \times M$  matrix of basis functions. The degrees of freedom is  $M = \text{trace}(\mathbf{H}_\xi)$ .

- By analogy, the effective degrees of freedom of a smoothing spline is defined as

$$\text{df}_\lambda = \text{trace}(\mathbf{S}_\lambda)$$



# Selection of smoothing parameters

- As  $\lambda \rightarrow 0$ ,  $\text{df}_\lambda \rightarrow N$  and  $\mathbf{S}_\lambda \rightarrow \mathbf{I}$ . As  $\lambda \rightarrow \infty$ ,  $\text{df}_\lambda \rightarrow 2$  and  $\mathbf{S}_\lambda \rightarrow \mathbf{H}$ , the hat matrix for linear regression on  $\mathbf{x}$ .
- Since  $\text{df}_\lambda$  is monotone in  $\lambda$ , we can invert the relationship and specify  $\lambda$  by fixing  $\text{df}_\lambda$  (this can be achieved by simple numerical methods). Using  $\text{df}$  in this way provides a uniform approach to compare many different smoothing methods.
- The leave-one-out (LOO) cross-validated error is given by

$$RSS_{cv}(\lambda) = \sum_{i=1}^N (y_i - \hat{f}_\lambda^{(-i)}(x_i))^2 = \sum_{i=1}^N \left[ \frac{y_i - \hat{f}_\lambda(x_i)}{1 - \{\mathbf{S}_\lambda\}_{ii}} \right]^2$$

# Smoothing splines in R

```
ss1<-smooth.spline(x,y,df=3)
ss2<-smooth.spline(x,y,df=15)
ss<-smooth.spline(x,y)

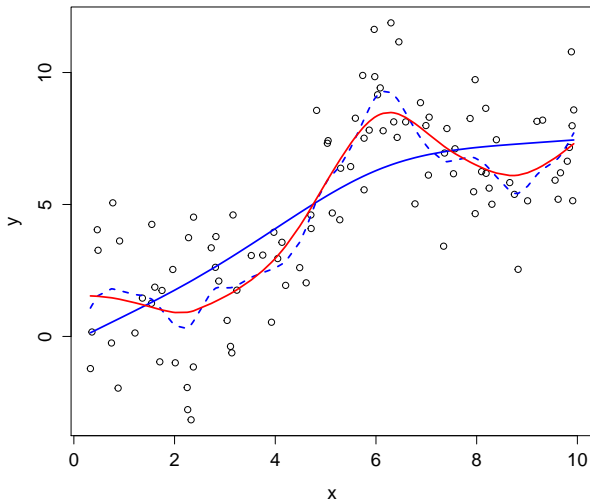
plot(x,y)
lines(x,ss1$y,col="blue",lwd=2)
lines(x,ss2$y,col="blue",lwd=2,lty=2)
lines(x,ss$y,col="red",lwd=2)

> ss$df
7.459728
```





## Result



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# Application to logistic regression

- The smoothing spline problem has been posed in a regression setting. It is typically straightforward to transfer this technology to other domains.
- Here we consider logistic regression with a single quantitative input  $X$ . The model is

$$\log \frac{\mathbb{P}(Y = 1|X = x)}{\mathbb{P}(Y = 0|X = x)} = f(x),$$

which implies

$$\mathbb{P}(Y = 1|X = x) = \frac{e^{f(x)}}{1 + e^{f(x)}} = p(x).$$



# Penalized log-likelihood

- We construct the penalized log-likelihood criterion

$$\begin{aligned}\ell(f; \lambda) &= \sum_{i=1}^N [y_i \log p(x_i) + (1 - y_i) \log(1 - p(x_i))] - \frac{1}{2} \lambda \int \{f''(t)\}^2 dt \\ &= \sum_{i=1}^N [y_i f(x_i) - \log(1 + e^{f(x)})] - \frac{1}{2} \lambda \int \{f''(t)\}^2 dt\end{aligned}$$

- As before, the optimal  $f$  is a finite-dimensional natural spline with knots at the unique values of  $x$ . We can represent  $f$  as

$$f(x) = \sum_{j=1}^N N_j(x) \theta_j.$$



# Optimization

- We compute the first and second derivatives

$$\begin{aligned}\frac{\partial \ell(\theta)}{\partial \theta} &= \mathbf{N}^T (\mathbf{y} - \mathbf{p}) - \lambda \mathbf{\Omega} \theta \\ \frac{\partial^2 \ell(\theta)}{\partial \theta \partial \theta^T} &= -\mathbf{N}^T \mathbf{W} \mathbf{N} - \lambda \mathbf{\Omega},\end{aligned}$$

where  $\mathbf{p}$  is the  $N$ -vector with elements  $p(x_i)$ , and  $\mathbf{W}$  is a diagonal matrix of weights  $p(x_i)(1 - p(x_i))$ .

- Parameters  $\theta_j$  can be estimated using the Newton method,

$$\theta^{new} = \theta^{old} - \left( \frac{\partial^2 \ell(\theta^{old})}{\partial \theta \partial \theta^T} \right)^{-1} \frac{\partial \ell(\theta^{old})}{\partial \theta}$$



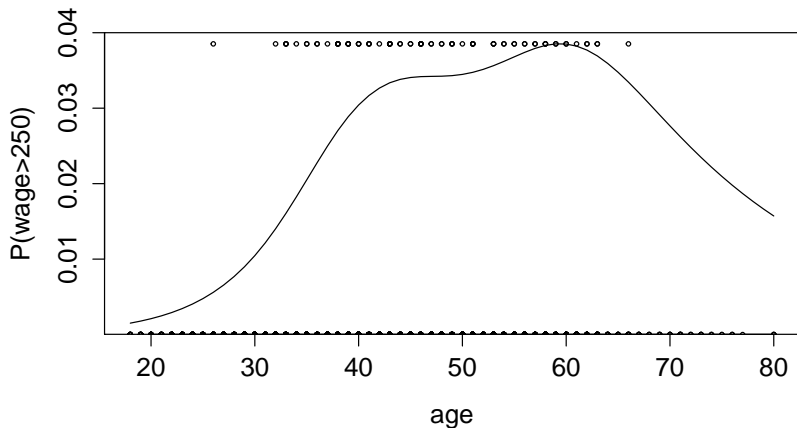
# Nonparametric logistic regression in R

```
library(gam)
class<-gam(I(wage>250) ~ s(age,df=3),data=Wage,family='binomial')
proba<-predict(class,newdata=data.frame(age=18:80),type='response')

plot(18:80,proba,xlab="age",ylab="P(wage>250)",type="l")
ii<-which(Wage$wage>250)
points(Wage$age[ii],rep(max(proba),length(ii)),cex=0.5)
points(Wage$age[-ii],rep(0,nrow(Wage)-length(ii)),cex=0.5)
```



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# Motivation

- Regression models play an important role in many data analyses, providing prediction and classification rules, and data analytic tools for understanding the importance of different inputs.
- Although attractively simple, the traditional linear model often fails in these situations: in real life, effects are often not linear.
- Here, we describe more automatic flexible statistical methods that may be used to identify and characterize nonlinear regression effects. These methods are called **generalized additive models** (GAMs).



# GAM for regression

- In the regression setting, a generalized additive model has the form

$$\mathbb{E}(Y|X_1, X_2, \dots, X_p) = \alpha + f_1(X_1) + f_2(X_2) + \dots + f_p(X_p)$$

- As usual  $X_1, X_2, \dots, X_p$  represent predictors and  $Y$  is the outcome.
- The  $f_j$ 's are unspecified smooth (nonparametric) functions.



# GAM for binary classification

- For two-class classification, recall the logistic regression model for binary data discussed previously. We relate the mean of the binary response  $\mu(X) = \mathbb{P}(Y = 1|X)$  to the predictors via a linear regression model and the logit link function:

$$\log \frac{\mu(X)}{1 - \mu(X)} = \alpha + \beta_1 X_1 + \dots + \beta_p X_p$$

- The additive logistic regression model replaces each linear term by a more general functional form

$$\log \frac{\mu(X)}{1 - \mu(X)} = \alpha + f_1(X_1) + \dots + f_p(X_p)$$

where again each  $f_j$  is an unspecified smooth function.

- While the nonparametric form for the functions  $f_j$  makes the model more flexible, the additivity is retained and allows us to interpret the model in much the same way as before.

# GAM: general form

- In general, the conditional mean  $\mu(X)$  of a response  $Y$  is related to an additive function of the predictors via a link function  $g$ :

$$g[\mu(X)] = \alpha + f_1(X_1) + \dots + f_p(X_p)$$

- Examples of classical link functions are the following:
  - $g(\mu) = \mu$  is the identity link, used for linear and additive models for Gaussian response data.
  - $g(\mu) = \text{logit}(\mu)$  as above, or  $g(\mu) = \text{probit}(\mu)$ , the probit link function, for modeling binomial probabilities. The probit function is the inverse Gaussian cumulative distribution function:  
 $\text{probit}(\mu) = \Phi^{-1}(\mu)$ .
  - $g(\mu) = \log(\mu)$  for log-linear or log-additive models for Poisson count data.



# Mixing linear and nonlinear effects, interactions

- We can easily mix in linear and other parametric forms with the nonlinear terms, a necessity when some of the inputs are qualitative variables (factors).
- The nonlinear terms are not restricted to main effects either; we can have nonlinear components in two or more variables, or separate curves in  $X_j$  for each level of the factor  $X_k$ , e.g.,
  - $g(\mu) = X^T \beta + \sum_k \alpha_k I(V = k) + f(Z)$  – a semiparametric model, where  $X$  is a vector of predictors to be modeled linearly,  $\alpha_k$  the effect for the  $k$ th level of a qualitative input  $V$ , and the effect of predictor  $Z$  is modeled nonparametrically.
  - $g(\mu) = f(X) + \sum_k g_k(Z) I(V = k)$  – again  $k$  indexes the levels of a qualitative input  $V$ , and thus creates an interaction term for the effect of  $V$  and  $Z$ ,
  - etc...



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# GAMs with natural splines

- If we model each function  $f_j$  as a natural spline, then we can fit the resulting model using simple least square (regression) or likelihood maximization algorithm (classification).
- For instance, with natural cubic splines, we have the following GAM:

$$g(\mu) = \sum_{j=1}^p \sum_{k=1}^{K(j)} \beta_{jk} N_k(X_j) + \epsilon,$$

where  $K(j)$  is the number of knots for variable  $j$ .





# Example in R

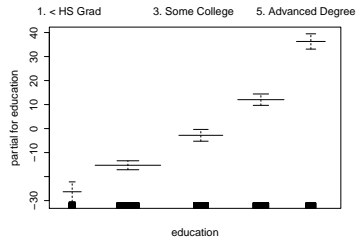
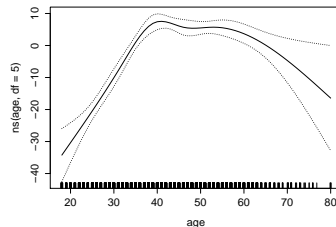
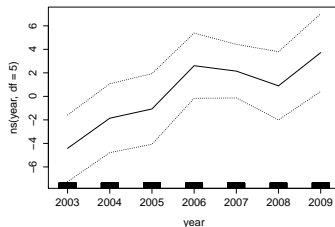
```
library("ISLR") # For the Wage data
library("splines")

fit1<-lm(wage ~ ns(year,df=5)+ns(age,df=5)+education,data=Wage)

library("gam")
fit2<-gam(wage ~ ns(year,df=5)+ns(age,df=5)+education,data=Wage)
plot(fit2,se=TRUE)
```



## Result



# GAMs with smoothing splines

- Consider an additive model of the form

$$Y = \alpha + f_1(X_1) + f_2(X_2) + \dots + f_p(X_p) + \epsilon,$$

where the error term  $\epsilon$  has mean zero.

- We can specify a penalized sum of squares for this problem,

$$SS(\alpha, f_1, \dots, f_p) = \sum_{i=1}^N \left( y_i - \alpha - \sum_{j=1}^p f_j(x_{ij}) \right)^2 + \sum_{j=1}^p \lambda_j \int f_j''(t_j)^2 dt_j,$$

where the  $\lambda_j \geq 0$  are tuning parameters.

- It can be shown that the minimizer of  $SS$  is an additive cubic spline model; each of the functions  $f_j$  is a cubic spline in the component  $X_j$  with knots at each of the unique values of  $x_{ij}$ ,  $i = 1, \dots, N$ .



# Unicity of the solution

- Without further restrictions on the model, the solution is not unique.
- The constant  $\alpha$  is not identifiable, since we can add or subtract any constants to each of the functions  $f_j$ , and adjust  $\alpha$  accordingly.
- The standard convention is to assume that  $\sum_{i=1}^N f_j(x_{ij}) = 0$  for all  $j$  – the functions average zero over the data.
- It is easily seen that  $\hat{\alpha} = \text{ave}(y_i)$  in this case.
- If in addition to this restriction, the matrix of input values (having  $ij$ th entry  $x_{ij}$ ) has full column rank, then  $SS$  is a strictly convex criterion and the minimizer is unique.



# Backfitting algorithm

- A simple iterative procedure exists for finding the solution.
- We set  $\hat{\alpha} = \text{ave}(y_i)$ , and it never changes.
- We apply a cubic smoothing spline  $S_j$  to the targets  $\{y_i - \hat{\alpha} - \hat{f}(x_{ik})\}_{i=1}^N$ , as a function of  $x_{ij}$  to obtain a new estimate  $\hat{f}_j$ .
- This is done for each predictor in turn, using the current estimates of the other functions  $\hat{f}_k$  when computing  $y - \hat{\alpha} - \sum_{k \neq j} \hat{f}_k(x_{ik})$ .
- The process is continued until the estimates  $\hat{f}_j$  stabilize.
- This procedure (known as **backfitting**) is grouped cyclic coordinate descent algorithm.



# Backfitting algorithm

- 1 Initialize:  $\hat{\alpha} = \text{ave}(y_i)$ ,  $\hat{f}_j = 0$ ,  $\forall i, j$ .
- 2 Cycle:  $j = 1, 2, \dots, p, 1, 2, \dots, p, \dots$ ,

$$\hat{f}_j \leftarrow \mathcal{S}_j \left[ \{y_i - \hat{\alpha} - \sum_{k \neq j} \hat{f}_k(x_{ik})\}_{i=1}^N \right]$$

$$\hat{f}_j \leftarrow \hat{f}_j - \frac{1}{N} \sum_{i=1}^N \hat{f}_j(x_{ij})$$

until the functions  $\hat{f}_j$  change less than a prespecified threshold.



# Example in R

```
library("gam")  
fit3<-gam(wage ~ s(year,df=5)+s(age,df=5)+education,data=Wage)  
plot(fit3,se=TRUE)
```



## Result

