- degrees of freedom for cubic splines, 1 covariate
 - K knots, K + 1 intervals, 4(K + 1) parameters
 - 3K restrictions
 - leaves K + 4 parameters, omit constant term, leaves K + 3
- degrees of freedom for natural splines
 - linear on $(-\infty, \xi_1]$ and $[\xi_K, \infty)$: 4 pars
 - cubic in interior intervals 4(K-1)
 - 3K restrictions, leaves K parameters, but
 - 2 knots added at x₍₁₎, x_(n): K + 2
 - omit constant term get K + 1

what does omit constant term mean? Example

$$\begin{array}{rcl} S_0(x) &=& a_0+b_0x+c_0x^2+d_0x^3, \quad 0\leq x\leq 1\\ S_1(x) &=& a_1+b_1(x-1)+c_1(x-1)^2+d_1(x-1)^3, \quad 1\leq x\leq \end{array}$$

• e.g.
$$a_0 = b_0 = c_0 = d_0 = 1$$
; fixes $a_1 = 4$, $b_1 = 6$, $c_1 = 8$

or, force a₀=0, add it in later

degrees of freedom for smoothing splines

$$\min_{f} \Sigma \{ y_i - f(x_i) \}^2 + \lambda \int \{ f''(t) \}^2 dt$$

solution is natural cubic splines with knots at unique x_i

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

$$\blacktriangleright \min_{\theta} \Sigma \{ y_i - \Sigma_j N_j(x_i) \theta_j \}^2 + \lambda \Sigma_{jk} \theta_j \theta_k \Omega_{jk}$$

•
$$\min_{\theta} (\mathbf{y} - N\theta)^T (\mathbf{y} - N\theta) + \lambda \theta^T \Omega_N \theta$$

•
$$\Omega_{jk} = \int N_j''(t) N_k''(t) dt$$

► just like ridge regression: $\hat{\theta} = (N^T N + \lambda \Omega_N)^{-1} N^T y$

•
$$\hat{f} = N\hat{\theta} = n(N^T N + \lambda \Omega_N)^{-1} N^T y = S_\lambda y$$
, say

- degrees of freedom defined to be trace S_λ by analogy
- Same formula works for regression splines (actually easier) cf (5.15)

- Suppose we have X_1 , X_2 , and $E(y \mid X) = f(X_1, X_2)$
- one solution is to combine separate spline bases for X₁ and X₂
- e.g. additively: $f(X_1, X_2) = f_1(X_1) + f_2(X_2)$ (this is what was done for heart data)
- doesn't permit interactions
- alternative is to use all possible cross products: called tensor products
- $f(X_1, X_2) = \sum_{j=1}^{M_1} \sum_{k=1}^{M_2} \theta_{jk} h_{1j}(X_1) h_{2k}(X_2)$
- ► analogous to forming quadratic functions in regression using, e.g., x₁², x₁x₂, x₂²

alternative to derive smoothing splines in two dimensions:

$$\min_{f} \sum_{i=1}^{N} \{ y_i - f(\underline{x}_i) \}^2 + \lambda J(|f|)$$

►
$$J(|f|) = \int \int (\partial_1^2 f + \partial_2^2 f + 2\partial_{12} f)^2 dx dy$$

- as in univariate case, solution exists in a spline basis similar to natural splines
- ► (5.39): $f(\underline{x}) = \beta_0 + \beta^T \underline{x} + \sum_{j=1}^N \alpha_j h_j(\underline{x})$

$$h_j(\underline{x}) = \eta(||\underline{x} - \underline{x}_j||), \quad \eta(z) = z^2 \log z$$

- called radial basis functions: take this form because of symmetry of penalty
- note uses N knots; reduced in implementation by regularization

Kernel methods for regression: univariate

• model: E(Y | x) = f(x) ("smooth")

• data:
$$y_i = f(x_i) + \epsilon_i$$

- ▶ running mean smoother: $\hat{f}(x_0) = \operatorname{ave}(y_i \mid x_i \in N_k(x_0))$
- ► N_k(x₀) set of k "nearest neighbours": k smallest values of |x_i x₀|
- smoother estimate using kernel weighted average

$$\hat{f}(\mathbf{x}_0) = \frac{\sum_{i=1}^N K_{\lambda}(\mathbf{x}_0, \mathbf{y}_i) \mathbf{y}_i}{\sum_{i=1}^N K_{\lambda}(\mathbf{x}_0, \mathbf{x}_i)}$$

[Figure 6.1]

kernel

$$\mathcal{K}_{\lambda}(\mathbf{x}_0, \mathbf{x}) = D\left(rac{|\mathbf{x} - \mathbf{x}_0|}{\lambda}
ight) ext{ or } D\left(rac{|\mathbf{x} - \mathbf{x}_0|}{h_{\lambda}(\mathbf{x}_0)}
ight)$$

kernel methods for regression

- ► λ determines the width of the neighbourhood, hence smoothness
- increasing \(\lambda\) gives smoother function (higher bias, lower variance)
- metric window width $(h_{\lambda}(x_0) = \lambda)$ constant bias, variance $\propto 1/\text{local density}$
- ► nearest neighbour window width (*h*_λ(*x*₀) depends on *x*₀) constant variance, bias ∝ 1/local density
- Choice of kernel:

$$D(t) = \begin{cases} \frac{3}{4}(1-t^2), |t| \le 1 & \text{Epanichakov} \\ 0 & \\ = \begin{cases} (1-|t|^3)^3, |t| \le 1 & \text{tri-cube} \\ 0 & \\ = & \phi(t) = \frac{1}{\sqrt{2\pi}}\exp(-t^2/2) & \text{Gaussian} \end{cases}$$

R or Splus:

ksmooth(x,y,kernel=c("box","normal"),bandwidth=0.5,range

loess(formula)

more later

- > eps<-rnorm(100,0,1/3)</pre>
- > x<-runif(100)</pre>
- > $sin4x <- function(x) {sin(4*x)}$
- > y<-sin4(x)+eps</pre>
- > plot(sin4,0,1,type="l",ylim=c(-1.0,1.5),xlim=c(0,1))
- > points(x,y)
- > lines(ksmooth(x,y,"box",bandwidth=.2),col="blue")
- > lines(ksmooth(x,y,"normal",bandwidth=.2),col="green")
- > plot(sin4,0,1,type="l",ylim=c(-1.0,1.5),xlim=c(0,1))
- > lines(ksmooth(x,y,"normal",bandwidth=.2),col="green")
- > lines(ksmooth(x,y,"normal",bandwidth=0.4),col="blue")
- > lines(ksmooth(x,y,"normal",bandwidth=0.6),col="red")

(Figure 6.1)

Local linear regression

replace weighted average of x_i's with weighted linear (or polynomial) regression: better endpoint behaviour

$$\begin{split} \min_{\alpha(x_0),\beta(x_0)} \sum K_{\lambda}(x_0,x_i) \{y_i - \alpha(x_0) - \beta(x_0)x_i\}^2 \\ \hat{f}(x_0) &= (1,x_0)(X^T W(x_0)X)^{-1} X^T W(x_0)y \\ X &= \begin{pmatrix} 1 & x_1 \\ 1 & x_2 \\ \vdots & \vdots \\ 1 & x_n \end{pmatrix} = B \\ W(x_0) &= \text{diag } K_{\lambda}(x_0,x_i) \end{split}$$

Notes

Recall weighted least squares:

$$\min_{\beta} \sum w_i (y_i - \beta_0 - \beta_1 x_i)^2 \text{ or } \min_{\beta} (y - X\beta)^T W(y - X\beta)$$

$$\hat{\boldsymbol{\beta}} = (\boldsymbol{X}^T \boldsymbol{W} \boldsymbol{X})^{-1} \boldsymbol{X}^T \boldsymbol{W} \boldsymbol{y}$$

- can combine the least squares weights with the kernel weights; see Figure 6.4 and pp. 169, 170.
- can also do local quadratic regression (and higher) but increases bias at endpoints
- for extrapolation book recommends local linear fits; for good fits in middle local quadratic
- In R there are several smoothers: ksmooth and loess are built in
- The first uses kernel smoothing, the second uses local linear regression (robustified)

- scatter.smooth fits a loess curve to a scatter plot
- Loess takes a family argument : family = gaussian gives weighted least squares using K_λ as weights and family=symmetric gives a robust version using Tukey's biweight
- supsmu implements "Friedman's super smoother": a running lines smoother with elaborate adaptive choice of bandwidth
- Library KernSmooth has locpoly for local polynomial fits, and by setting degree = 0 gives a kernel smooth

```
> lo1 <- loess(y^x, degree=1, span=0.75)
> attributes(lo1)
$names
                "fitted" "residuals" "enp"
 [1] "n"
                                                       "s"
                                                                   "one.
[7] "two.delta" "trace.hat" "divisor"
                                          "pars"
                                                       "kd"
                                                                   "call
[13] "terms"
              "xnames"
                              " x "
                                          "v"
                                                       "weights"
$class
[1] "loess"
> plot(sin4,0,1,tvpe="l",vlim=c(-1.0,1.5),xlim=c(0,1))
> points(x,lo1$fitted,pch=".",col="red")
> plot(x,lol$fitted,pch=".",col="red",ylim=c(-1.0,1.5),xlim=c(0,1))
> lines(ksmooth(x,y,"normal",bandwidth=0.4),col="blue")
> plot(x,lo1$fitted,pch=".",col="red",ylim=c(-1.0,1.5),xlim=c(0,1))
> lo2 < -loess(v^x, degree=1, span=0.4)
> points(x,lo2$fitted,pch=".",col="green")
> points(x,loess(v<sup>x</sup>x,degree=2,span=0.4)$fitted,pch=".",col="purple")
```

Notes

- $\hat{f} = S_{\lambda} y$ and df=trace(S_{λ}), as in smoothing splines
- X can have up to 4 numerical predictors
- ▶ while possible to fit these models in R^p, (see §6.3, 6.4), doesn't seem so useful
- §6.4 describes ways to impose some structure to get a more interpretable model
- can use the same idea for likelihood functions and maximum likelihood estimates:

$$\max_{\beta} \sum \ell(\beta; y_i)$$

replaced by

$$\max_{\beta} \sum K_{\lambda}(x_0, x_i) \ell(\beta; y_i)$$

called local likelihood and described in $\S6.5$

Kernel methods for classification

- model: $X \sim f(\cdot)$
- training data (x_1, \ldots, x_N)
- $\hat{f}(x_0) = \frac{\#\{x_i \in n_\lambda(x_0)\}}{N\lambda}$ (a histogram)
- $\hat{f}(x_0) = \frac{1}{N\lambda} \sum K_{\lambda}(x_0, x_i)$: smooth density estimate
- implemented in R as density(x, ...) with a large choice of kernels; default is Gaussian, see (6.23)
- ▶ for classification: compute $\hat{f}_j(X)$ for each class

$$\hat{\mathrm{pr}}(\mathsf{Y}=j \mid \mathsf{X}=\mathsf{x}_0) = \hat{\pi}_j \hat{f}_j(\mathsf{x}_0) / \sum \hat{\pi}_k \hat{f}_j(\mathsf{x}_0)$$

with *p* inputs (§6.6.3); treat the inputs as independent

$$\hat{f}_j(\underline{X}) = \prod_{k=1}^p \hat{f}_{jk}(X_k)$$

the Naive Bayes classifier:

$$\hat{\mathrm{pr}}(\mathsf{Y} = j \mid \underline{X} = \underline{x}_0) = \hat{\pi}_j \hat{f}_j(\underline{(x_0)} / \Sigma \hat{\pi}_j \hat{f}_j(\underline{x}_0))$$