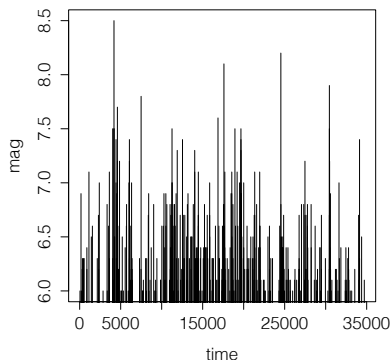


Today

- ▶ HW 3 due April 1
- ▶ Project due April 15
- ▶ nonparametric and semi-parametric regression
- ▶ In the News:
 - Statschat
 - New Yorker
- ▶ recap: kernel regression – summarize; plus where it is implemented
- ▶ honey

... example 10.31

SM



```
library(SMPracticals); data(quake)
with(quake, plot(time, mag, type="h"))
```

Recap: nonparametric Regression

ELM, Ch. 11; SM, §10.7

- ▶ model $y_i = f(x_i) + \epsilon_i$, $i = 1, \dots, n$ x_i scalar
- ▶ mean function $f(\cdot)$ assumed to be “smooth”
- ▶ introduce a **kernel function** $K(u)$ and define a set of weights

$$w_i = \frac{1}{\lambda} K\left(\frac{x_i - x_0}{\lambda}\right)$$

- ▶ estimate of $f(x)$, at $x = x_0$:

$$\hat{f}_\lambda(x_0) = \frac{\sum_{i=1}^n w_i y_i}{\sum_{i=1}^n w_i}$$

- ▶ Nadaraya-Watson estimator – local averaging

Local Polynomials

- ▶ better estimates can be obtained using local regression at point x_0 instead of local averaging



$$\begin{pmatrix} y_1 \\ y_2 \\ \vdots \\ y_n \end{pmatrix} = \begin{pmatrix} 1 & (x_1 - x_0) & \cdots & (x_1 - x_0)^k \\ \vdots & \vdots & & \vdots \\ 1 & (x_n - x_0) & \cdots & (x_n - x_0)^k \end{pmatrix} \begin{pmatrix} \beta_0 \\ \beta_1 \\ \vdots \\ \beta_k \end{pmatrix} + \begin{pmatrix} \varepsilon_1 \\ \varepsilon_2 \\ \vdots \\ \varepsilon_n \end{pmatrix},$$



$$\hat{\beta} = (X^T W X)^{-1} X^T W y, \quad w_i = \frac{1}{\lambda} K \left(\frac{x_i - x_0}{\lambda} \right)$$



$$\hat{f}_\lambda(x_0) = \hat{\beta}_0$$

... local polynomials

- ▶ odd-order polynomials work better than even; usually local linear fits are used
- ▶ kernel function is often a Gaussian density, or the tricube function

$$K(u) = (1 - |u|^3)^3, \quad |u| < 1$$

- ▶ as with Nadaraya-Watson estimators, choice of **bandwidth**, λ controls smoothness of function
- ▶ `loess` computes local linear regression, and includes a robust option

```
loess(formula, data, weights, subset, na.action, model = FALSE,  
      span = 0.75, enp.target, degree = 2,  
      parametric = FALSE, drop.square = FALSE, normalize = TRUE,  
      family = c("gaussian", "symmetric"),  
      method = c("loess", "model.frame"),  
      control = loess.control(...), ...)
```

- ▶ `"symmetric"` implements robust version

... cross-validation



$$CV(\lambda) = \sum_{i=1}^n \{y_i - \hat{f}_{-i}(x_i)\}^2$$

- ▶ for local polynomials

$$CV(\lambda) = \sum_{i=1}^n \left\{ \frac{y_i - \hat{f}_{\lambda}(x_i)}{1 - S_{ii}(\lambda)} \right\}^2$$

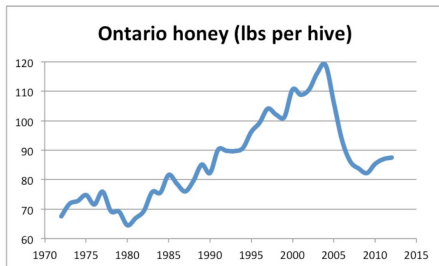
- ▶ simpler

$$GCV(\lambda) = \sum_{i=1}^n \left\{ \frac{y_i - \hat{f}_{\lambda}(x_i)}{1 - \text{tr}(S_{\lambda})/n} \right\}^2$$

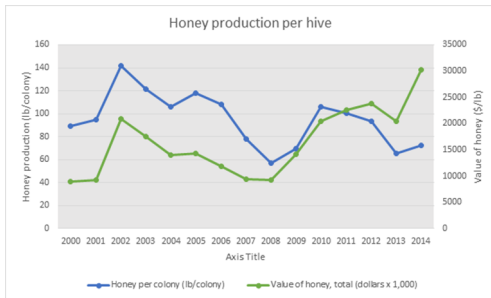
- ▶ library `sm` does local linear regression, computes optimal bandwidth via `hcv`

Example: honey production

Stat Can

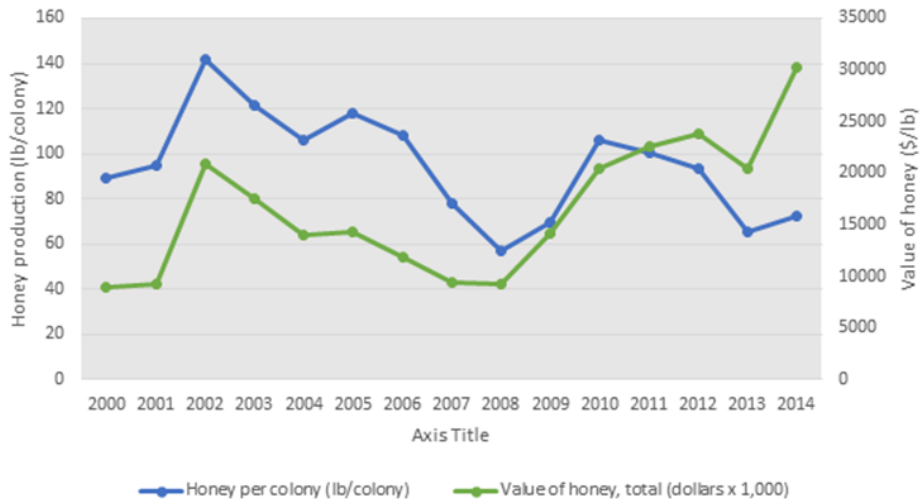


5-year running mean

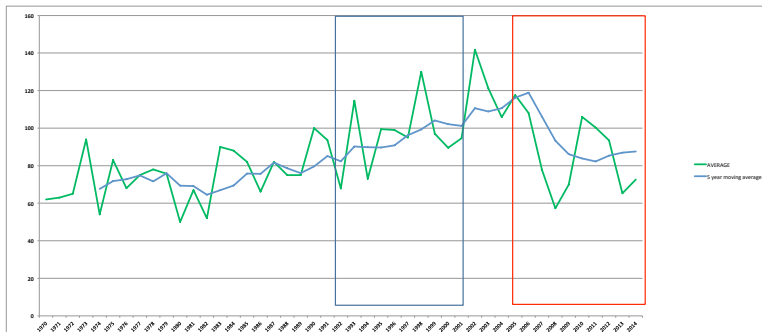


annual averages

Honey production per hive



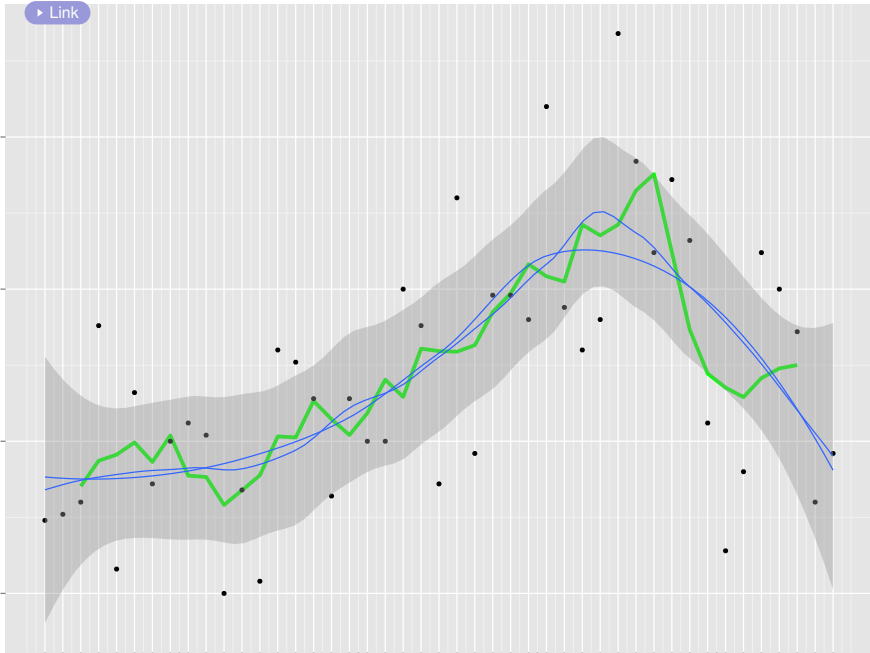
YEAR	1970	1971	1972	1973	1974	1975	1976	1977	1978	1979	1980	1981	1982	1983	1984	1985	1986	1987	1988	1989	1990	1991	1992
AVERAGE	62	63	65	94	54	83	68	75	78	76	50	67	52	90	88	82	66	82	75	75	100	94	68
5 year moving average					68	72	73	75	72	76	69	69	65	67	69	76	76	82	79	76	80	85	82



[▶ Link](#)

average

year



- ▶ $\hat{\beta} = (X^T W X)^{-1} X^T W y$
- ▶ $W = \text{diag}(w_1, \dots, w_n)$
- ▶ $\hat{f}_\lambda(x_0) = \hat{\beta}_0 = \sum_{i=1}^n S(x_0; x_i, \lambda) y_i$; $\hat{f} = (\hat{f}_\lambda(x_1), \dots, \hat{f}_\lambda(x_n)) = S_\lambda y$
- ▶ $S(x_0; x_1, \lambda), \dots, S(x_0; x_n, \lambda)$ first row of “hat” matrix
 $(X^T W X)^{-1} X^T W$
- ▶ $E\{\hat{f}_\lambda(x_0)\} = \sum_{i=1}^n S(x_0; x_i, \lambda) f_\lambda(x_i)$
- ▶ $\text{var}\{\hat{f}_\lambda(x_0)\} = \sigma^2 \sum_{i=1}^n S^2(x_0; x_i, \lambda)$
- ▶ $\tilde{\sigma}^2 = \frac{1}{n-2\nu_1+\nu_2} \sum_{i=1}^n \{y_i - \hat{f}_\lambda(x_i)\}^2$
- ▶ $\nu_1 = \text{tr}(S_\lambda)$, $\nu_2 = \text{tr}(S_\lambda^T S_\lambda)$

potential estimates of ‘degrees of freedom’

Flexible modelling using basis expansions

ELM §11.2; SM §10.7.2

- ▶ $y_i = f(x_i) + \epsilon_i$

- ▶ Flexible linear modelling

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

- ▶ This is called a **linear basis expansion**, and h_m is the m th basis function
- ▶ For example: $f(x) = \beta_0 + \beta_1 x + \beta_2 x^2$, or $f(x) = \beta_0 + \beta_1 \sin(x) + \beta_2 \cos(x)$, etc.
- ▶ Simple linear regression has $h_1(x) = 1$, $h_2(x) = x$

Piecewise polynomials

- ▶ piecewise constant basis functions
$$h_1(x) = I(x < \xi_1), \quad h_2(x) = I(\xi_1 \leq x < \xi_2),$$
$$h_3(x) = I(\xi_2 \leq x)$$
- ▶ equivalent to fitting by local averaging
- ▶ piecewise linear basis functions , with constraints
$$h_1(x) = 1, \quad h_2(x) = x$$
$$h_3(x) = (x - \xi_1)_+, \quad h_4(x) = (x - \xi_2)_+$$
- ▶ windows defined by **knots** $\xi_1, \xi_2, \dots$
- ▶ **piecewise cubic basis functions**
$$h_1(x) = 1, h_2(x) = x, h_3(x) = x^2, h_4(x) = x^3$$
- ▶ continuity $h_5(x) = (x - \xi_1)_+^3, \quad h_6(x) = (x - \xi_2)_+^3$
- ▶ continuous function, continuous first and second derivs

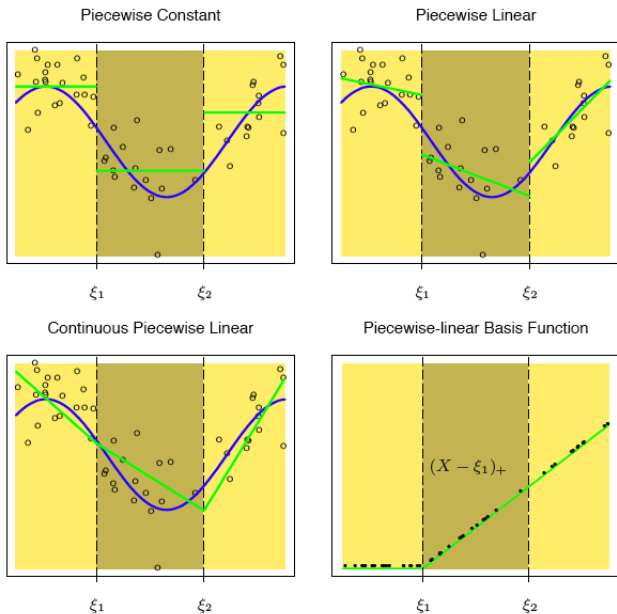


FIGURE 5.1. The top left panel shows a piecewise constant function fit to some artificial data. The broken vertical lines indicate the positions of the two knots ξ_1 and ξ_2 . The blue curve represents the true function, from which the data were

Piecewise Cubic Polynomials

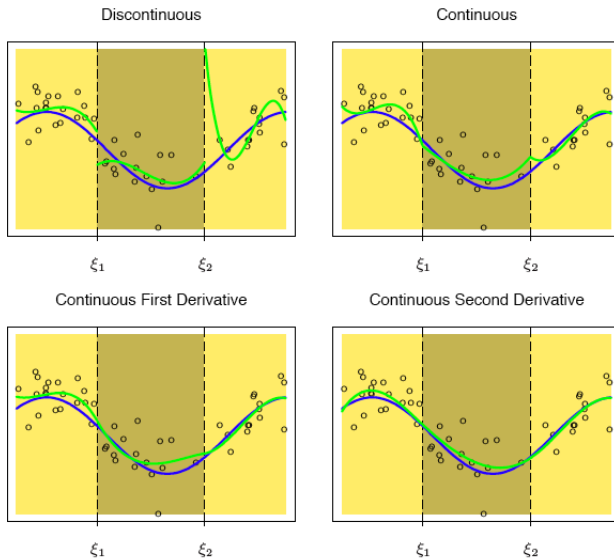
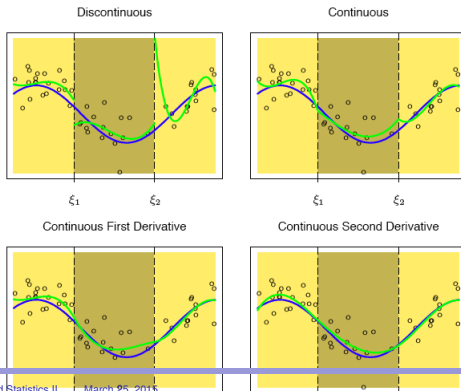


FIGURE 5.2. A series of piecewise-cubic polynomials, with increasing orders of continuity.

Cubic splines

- ▶ **truncated power basis** of degree 3
- ▶ need to choose number of knots K and placement of knots $\xi_1, \dots, \xi_K$
- ▶ construct features matrix using truncated power basis set
- ▶ use constructed matrix as set of predictors

Piecewise Cubic Polynomials



... cubic splines

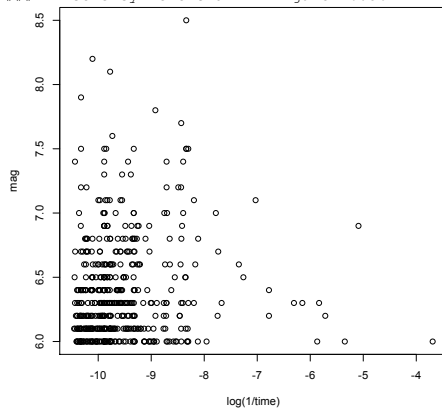
```
> with(quake, bs(log(1/time))[1:10,])
#bs(x) with no other arguments just gives a
# single cubic polynomial
      1      2      3
[1,] 0.0000000 0.0000000 1.0000000
[2,] 0.1018013 0.3903714 0.4989780
[3,] 0.1359705 0.4189773 0.4303434
[4,] 0.1884790 0.4408886 0.3437743
[5,] 0.2056632 0.4436068 0.3189471
...
attr(,"degree")
[1] 3
attr(,"knots")
numeric(0)
attr(,"Boundary.knots")
[1] -10.454784 -3.690961
attr(,"intercept")
[1] FALSE
attr(,"class")
[1] "bs"      "basis"   "matrix"
```

... cubic splines

```
> with(quake, bs(log(1/time), df=5)[1:10,])
# gives a proper cubic spline basis, here with 5 df
      1          2          3          4          5
[1,] 0 0.00000000 0.0000000 0.0000000 1.0000000
[2,] 0 0.01110655 0.1250814 0.4247847 0.4390274
[3,] 0 0.01846075 0.1661869 0.4486889 0.3666635
[4,] 0 0.03370916 0.2283997 0.4600092 0.2778819
[5,] 0 0.03989014 0.2484715 0.4585984 0.2530400
...
attr(,"degree")
[1] 3
attr(,"knots")
33.33333% 66.66667%
-9.943294 -9.520987
attr(,"Boundary.knots")
[1] -10.454784 -3.690961
```

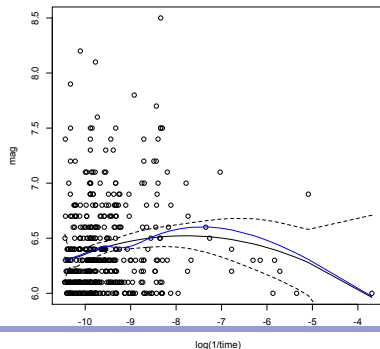
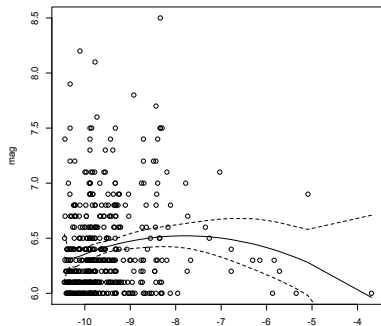
Example: earthquake data

```
> data(quake, package="SMPracticals")
> head(quake)
      time mag
1  40.08333 6.0
2 162.38889 6.9
3 210.22917 6.0
> with(quake, plot(log(1/time), mag))## using a different measure of
### intensity here than in Figure 10.36
```



... earthquake data

```
> quake.bs = lm(mag ~ bs(log(1/time),df=5),data = quake)
> quake.pred = predict(quake.bs, se.fit = TRUE, interval = "confidence")
> quake.pred
$fit
      fit      lwr      upr
1  5.962665 5.216283 6.709047
2  6.279641 5.979190 6.580092
3  6.323859 6.042772 6.604946
> lines(log(1/quake$time),quake.pred[[1]])
> lines(log(1/quake$time),quake.pred[[1]]+quake.pred$se.fit, lty=2)
> lines(log(1/quake$time),quake.pred[[1]]-quake.pred$se.fit, lty=2)
> quake.lo = loess(mag ~ log(1/time), data = quake)
> lines(quake.lo$x,quake.lo$fitted,col="blue")
```



Regression splines

- ▶ regression splines are 'hand-crafted': $\text{bs}(x, \text{df} = ?)$
- ▶ once crafted, model is parametric

- ▶
$$f(x) = \sum_{m=1}^M \beta_m h_m(x) \quad \text{df} = M$$

- ▶ degrees of freedom controls how many **knots** are placed
- ▶ equivalent to how many parameters are fit
- ▶ ELM, §11.2 constructs linear splines 'by hand' Figure 11.7
- ▶ a different approach is **smoothing splines** – put a LOT of knots, and regularize

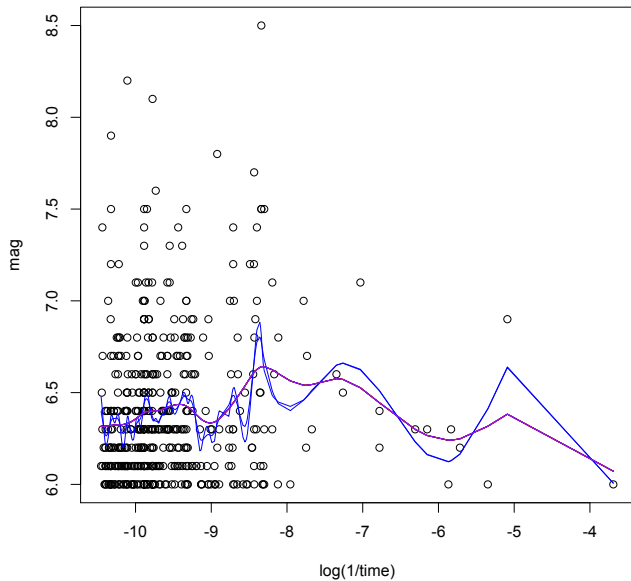
- ▶ $y_i = f(x_i) + \epsilon_i, \quad i = 1, \dots, n$
- ▶ choose $f(\cdot)$ to solve

$$\min_f \sum_{i=1}^n \frac{\{y - f(x_i)\}^2}{2\sigma^2} - \frac{\lambda}{2\sigma^2} \int_a^b \{f''(t)\}^2 dt, \quad \lambda > 0$$

- ▶ solution is a cubic spline, with knots at each observed x_i value

see SM Figure 10.18 for a non-regularized solution

- ▶ has an explicit, finite dimensional solution
- ▶ $\hat{f} = \{\hat{f}(x_1), \dots, \hat{f}(x_n)\} = (I + \lambda K)^{-1} y$
- ▶ K is a symmetric $n \times n$ matrix of rank $n - 2$



... smoothing splines

```
> quake$int = log(1/quake$time)
> quake[1:4,]
      time mag      int
1  40.08333 6.0 -3.690961
2 162.38889 6.9 -5.089994
3 210.22917 6.0 -5.348198
4 303.85417 6.2 -5.716548

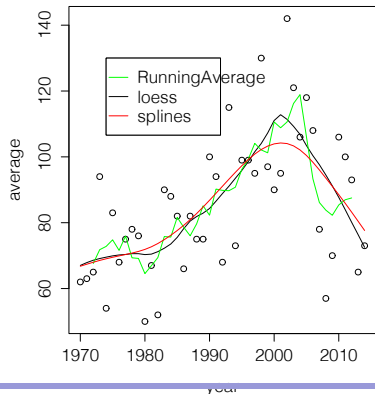
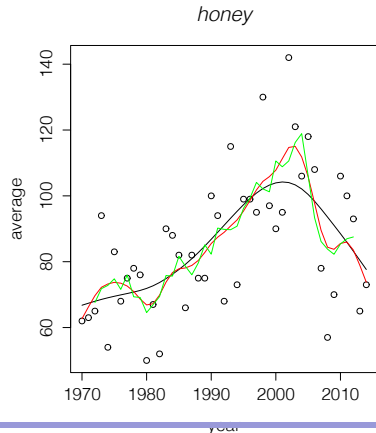
> attach(quake)
> plot(int,mag)
> quake.ss2 = smooth.spline(x = int, y = mag, df = 5)
> lines(quake.ss2, col="red")
> quake.ss3
Call:
smooth.spline(x = int, y = mag, cv = TRUE)

Smoothing Parameter spar= 1.499945 lambda= 0.0001340604 (25 iterations)
Equivalent Degrees of Freedom (Df): 11.35023
Penalized Criterion: 64.57512
PRESS: 0.1730025
> lines(quake.ss3, col="blue")
```

... smoothing splines

```
> attach(honey)

> plot(year, average)
> plot(year, average, main = "honey")
> lines(smooth.spline(honey$year, honey$average))
> lines(smooth.spline(honey$year, honey$average, df = 11), col="red")
> lines(honey$year, honey$runmean, col="green")
```



Multidimensional splines

- ▶ so far we are considering just 1 X at a time
- ▶ for regression splines we replace each X by the new columns of the basis matrix
- ▶ for smoothing splines we get a univariate regression
- ▶ it is possible to construct smoothing splines for two or more inputs simultaneously, but computational difficulty increases rapidly
- ▶ these are called thin plate splines
- ▶ alternative:

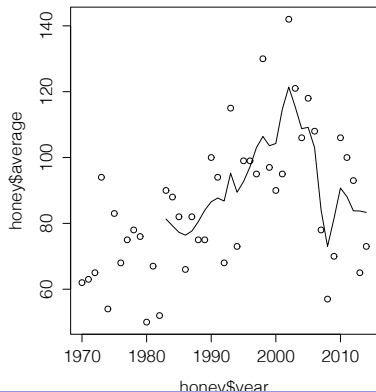
$$E(Y \mid X_1, \dots, X_p) = f_1(X_1) + f_2(X_2) + \dots + f_p(X_p)$$

additive models

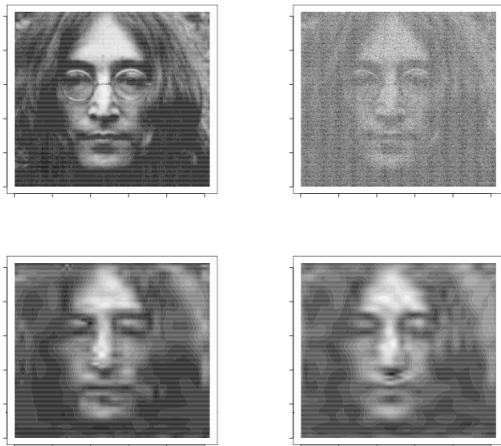
ELM Ch. 12

- ▶ basis functions with very 'local' support
- ▶ useful for capturing regions of high variation
- ▶ useful for data compression – transmit only local components

Fig 11.11



... wavelets



Vidaković and Mueller, "Wavelets for kids (Part I)" 1994.

Which smoothing method?

ELM §11.6

- ▶ basis functions: natural splines, Fourier, wavelet bases
- ▶ regularization via cubic smoothing splines
- ▶ kernel smoothers: locally constant/linear/polynomial
- ▶ adaptive bandwidth, running medians, running M -estimates
- ▶ with very little noise, a small amount of local smoothing (e.g. nearest neighbours)
- ▶ with moderate amounts of noise, kernel and spline methods are effective
- ▶ with large amounts of noise, parametric methods are more attractive
- ▶ “It is not reasonable to claim that any one smoother is better than the rest”
 - ▶ `loess` is robust to outliers, and provides smooth fits
 - ▶ spline smoothers are more efficient, but potentially sensitive to outliers
 - ▶ kernel smoothers are very sensitive to bandwidth

- ▶ Ridge regression



$$\begin{aligned}\hat{\beta}_{LS} &= (X^T X)^{-1} X^T y \\ \hat{\beta}_{ridge} &= (X^T X + \lambda I)^{-1} X^T y\end{aligned}$$

- ▶ can show that $\hat{\beta}_{ridge}$ satisfies

$$\begin{aligned}\min_{\beta} & \left(\sum \{y_i - \beta_0 - \sum_{j=1}^p x_{ij} \beta_j\}^2 + \lambda \sum_{j=1}^p \beta_j^2 \right) \\ \min_{\beta} & \sum \{y_i - \beta_0 - \sum_{j=1}^p x_{ij} \beta_j\}^2 \quad \text{s.t. } \sum \beta_j^2 \leq t\end{aligned}$$

- ▶ Assume x_j 's are centered and put these in matrix X (with no column of 1's):

$$\min_{\beta} (y - X\beta)^T (y - X\beta) \quad \text{s.t. } \|\beta\|^2 \leq t$$

... ridge regression



$$\min_{\beta} \{ (y - X\beta)^T (y - X\beta) + \lambda \|\beta\|^2 \}$$

- ▶ λ is a tuning parameter: $\lambda = 0$ gives $\hat{\beta}_{LS}$, $\lambda \rightarrow \infty$
- ▶ in R the `library MASS library(MASS)` has a ridge regression version of `lm` called `lm.ridge`
- ▶ if columns of X are nearly linearly dependent (multicollinearity), $\hat{\beta}$'s for these columns should be shrunk towards 0.
- ▶ essential that the predictors are all scaled to the same units
- ▶ this is difficult for interpretation of the coefficients
- ▶

$$df(\lambda) = \text{tr}[X(X^T X + \lambda I)^{-1} X^T] = \sum_{j=1}^p \frac{d_j^2}{d_j^2 + \lambda}$$

effective number of parameters

Lasso



$$\min_{\beta} \left(\sum \{y_i - \beta_0 - \sum_{j=1}^p x_{ij} \beta_j\}^2 + \lambda \sum_{j=1}^p |\beta_j| \right)$$



$$\min_{\beta} \sum \{y_i - \beta_0 - \sum_{j=1}^p x_{ij} \beta_j\}^2 \quad \text{s.t.} \quad \sum |\beta_j| \leq t$$

- ▶ quadratic programming problem
- ▶ $\hat{\beta}^{lasso}$ is nonlinear function of y
- ▶ Tibshirani (1996), JRSS B and (2011), JRSS B
- ▶ <http://http://www-stat.stanford.edu/~tibs/lasso.html>

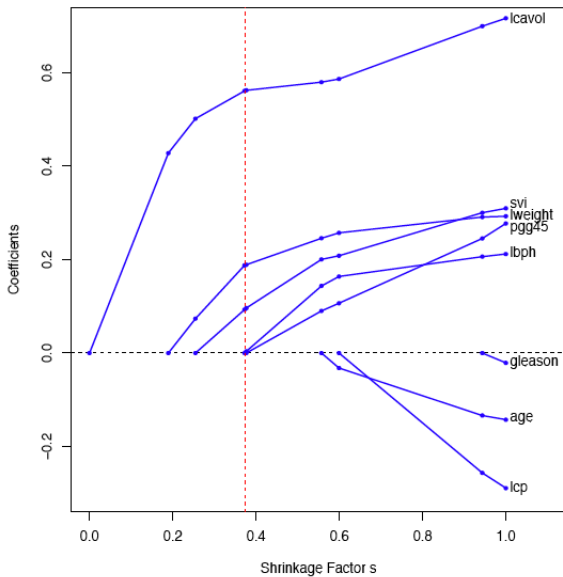


FIGURE 3.10. Profiles of lasso coefficients, as the tuning parameter t is varied. Coefficients are plotted versus $s = t / \sum_1^p |\hat{\beta}_j|$. A vertical line is drawn at $s = 0.36$, the value chosen by cross-validation. Compare Figure 3.8 on page 65; the lasso

... shrinkage

- ▶ ridge regression gives “proportional shrinkage”
- ▶ subset selection gives “hard thresholding” (some $\beta_j \rightarrow 0$)
- ▶ lasso gives “soft thresholding”: blend of shrinkage and zeroing
- ▶ **elastic net** combines lasso and ridge regression

$$\min_{\beta} \left(\sum \{y_i - \beta_0 - \sum_{j=1}^p x_{ij}\beta_j\}^2 + \lambda_1 \sum_{j=1}^p |\beta_j| + \lambda_2 \sum_{j=1}^p \beta_j^2 \right)$$

- ▶ implemented in R in `library(glmnet)`
- ▶ estimates of coefficients are biased (but may have small mean-squared error)
- ▶ Lasso is now used as a variable selection method
- ▶ improvements in algorithms allow fast computation even for $p > n$

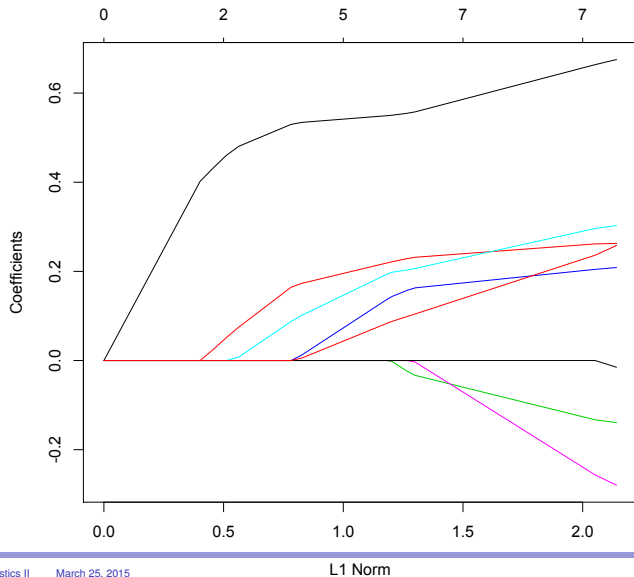
```
> prostate <- read.csv(file="prostate.data",sep="\t")
## data is at http://statweb.stanford.edu/~tibs/ElemStatLearn/datasets/prostate.data

> head(prostate)
      lcavol  lweight age      lbph svi      lcp gleason  pgg45
1 -0.5798185 2.769459 50 -1.386294 0 -1.386294      6      0
2 -0.9942523 3.319626 58 -1.386294 0 -1.386294      6      0
3 -0.5108256 2.691243 74 -1.386294 0 -1.386294      7     20
4 -1.2039728 3.282789 58 -1.386294 0 -1.386294      6      0
5  0.7514161 3.432373 62 -1.386294 0 -1.386294      6      0
6 -1.0498221 3.228826 50 -1.386294 0 -1.386294      6      0

      lpsa train
1 -0.4307829 TRUE
2 -0.1625189 TRUE
3 -0.1625189 TRUE
4 -0.1625189 TRUE
5  0.3715636 TRUE
6  0.7654678 TRUE

> xp <- scale(prostate[,1:8])
> y <- prostate[,9]
> train <- prostate[,10]
## standardize data; y is the response (log psa); extract training data
##
> library(glmnet)
> pr.lasso <- glmnet(xp[train,],y[train]) #Lasso
> pr.lasso2 <- glmnet(xp[train,],y[train], alpha=0) # ridge
> plot(pr.lasso); plot(pr.lasso2)
```

... prostate data



... prostate data

