



Statistical learning for biological data

David Causeur

Department of Statistics and Computer Science

Agrocampus Ouest

IRMAR CNRS UMR 6625

<http://www.agrocampus-ouest.fr/math/causeur/>



Outline

- 1 Regression modeling
 - Why 'regression'?
 - Fitting linear regression models
 - Feature selection for prediction
- 2 Penalized regression
 - Sparse regression modeling
 - Penalized estimation of classification models
- 3 Latent variable models for prediction
 - Partial Least Squares
 - Linear Discriminant Analysis
- 4 Nonparametric regression
 - Smooth effect curve
 - Generalized Additive Modeling



Outline

- 1 Regression modeling
 - Why 'regression'?
 - Fitting linear regression models
 - Feature selection for prediction
- 2 Penalized regression
 - Sparse regression modeling
 - Penalized estimation of classification models
- 3 Latent variable models for prediction
 - Partial Least Squares
 - Linear Discriminant Analysis
- 4 Nonparametric regression
 - Smooth effect curve
 - Generalized Additive Modeling



Outline

- 1 Regression modeling
 - Why 'regression'?
 - Fitting linear regression models
 - Feature selection for prediction
- 2 Penalized regression
 - Sparse regression modeling
 - Penalized estimation of classification models
- 3 Latent variable models for prediction
 - Partial Least Squares
 - Linear Discriminant Analysis
- 4 Nonparametric regression
 - Smooth effect curve
 - Generalized Additive Modeling



Outline

- 1 Regression modeling
 - Why 'regression'?
 - Fitting linear regression models
 - Feature selection for prediction
- 2 Penalized regression
 - Sparse regression modeling
 - Penalized estimation of classification models
- 3 Latent variable models for prediction
 - Partial Least Squares
 - Linear Discriminant Analysis
- 4 Nonparametric regression
 - Smooth effect curve
 - Generalized Additive Modeling



Nonlinear regression function

In some situations, the regression function $x \mapsto f(x)$ is obviously **nonlinear**

... but no biological theory can help in setting a parametric framework:

$$Y = f(x) + \varepsilon, \varepsilon \sim \mathcal{N}(0; \sigma)$$

Still, f is usually assumed to be *regular*: continuous, differentiable, twice differentiable, ...

► Predicting the daily maximum ozone concentration using \mathbb{R}



Nonlinear regression function

How to draw a regression curve without introducing any biological knowledge?



Local polynomial approximation: the *loess* method

Let x_0 be any value of X in $[\min(x_i); \max(x_i)]$:

$$f(x_0) = \mathbb{E}(Y \mid X = x_0)$$

How to estimate $f(x_0)$?

In the *rare* situations where replications of response values are observed for items with $X = x_0$

a natural estimate of $f(x_0)$ is just the average of those response values.

Most often, we do not even have any observation at $X = x_0$.



Local polynomial approximation: the *loess* method

More generally, change " $X = x_0$ " into " X close to x_0 "

Let $d_i = |x_0 - x_i|$ with $d_{(1)} \leq d_{(2)} \leq \dots \leq d_{(n)}$

For $1 \leq k \leq n$, the **k -neighborhood** of x_0 is defined as follows:

$$N_k(x_0) = \{i = 1, \dots, n, d_i \leq d_{(k)}\}$$

► Illustration using R



Local polynomial approximation: the *loess* method

How to aggregate the response values of items within $N_k(x_0)$ to form an estimate of $f(x_0)$?

The *loess* answer: using a *weighted* local polynomial fit of the regression function.



Local polynomial approximation: the *loess* method

Why weighting the sampling items within $N_k(x_0)$?

... in order to estimate $f(x_0)$, the closest data points should be favored.

A possible weighting function:

$$\omega(x_j) = (1 - u_j^3)^3, \text{ where } u_j = \frac{d_j}{\max_{i \in N_k(x_0)} d_i}.$$

► Displaying the tricube function in \mathbb{R}



Local polynomial approximation: the *loess* method

Local weighted least-squares fit of a polynomial

$$\hat{D}(x) = \hat{a} + \hat{b}x + \hat{c}x^2, \text{ where}$$
$$(\hat{a}, \hat{b}, \hat{c}) \text{ minimizes } \sum_{i \in N_k(x_0)} \omega(x_i) [Y_i - a - bx_i - cx_i^2]^2.$$

Finally, $\hat{f}(x_0) = \hat{D}(x_0)$.

- ▶ Local polynomial approximation using package `gam` in R



Local polynomial approximation: the *loess* method

What is the best value for k (or span)?

In a prediction accuracy perspective, k can be chosen so as to minimize the PRESS

► Optimal span using R



Spline smoothing

Let us assume that f is a spline function of degree d on $[a, b]$

There exists a partition $a = t_0 < t_1 < t_2 < \dots < t_L < b = t_{L+1}$ of $[a, b]$ such that:

- f is a piecewise polynomial of degree d on the partition;
- f is $d - 1$ times continuously differentiable on $[a, b]$.

... Spline($t_1, t_2, \dots, t_L; d$) is a $(L + D + 1)$ -dimensional linear space

Note: L is the number of interior nodes.

Note also: usually, $D = 3$ (*cubic splines*)



Spline smoothing

For $D = 0$, $L + 1$ basis functions $B_{i,0}$, $i = 0, \dots, L$:

$$B_{i,0}(x) = \begin{cases} 1 & \text{if } x \in [t_i, t_{i+1}[\\ 0 & \text{otherwise} \end{cases}$$

For $D = 1$, $L + 2$ basis functions $B_{i,1}$, $i = -1, 0, \dots, L$ with support $[t_i, t_{i+2}]$:

$$B_{i,1}(x) = \frac{x - t_j}{t_{i+1} - t_j} B_{i,0}(x) + \frac{t_{i+2} - x}{t_{i+2} - t_{i+1}} B_{i+1,0}(x),$$

where, for all x , $B_{-1,0}(x) = 0$ and $B_{L+1,0}(x) = 0$.

For $D = 2, \dots$

► Display B-splines using R



Spline smoothing

Since $\text{Spline}(t_1, t_2, \dots, t_L; d)$ is a linear space:

$$Y = b_{-3}B_{-3}(x) + b_{-2}B_{-2}(x) + \dots + b_L B_L(x) + \varepsilon$$

Estimation of $b = (b_{-3}, \dots, b_L)$ is just a **linear** least-squares minimization issue.

► Estimation of B-spline coefficients using \mathbb{R}



Spline smoothing

loess and *spline* are both **linear smoothers**:

$$\hat{Y} = S_{\lambda} Y,$$

where λ is a generic hyper-parameter for tuning regularity:

- the number of classes in the partition for *loess*
- the dimension of the B-spline basis for *spline*

e.g for spline smoothing: if B_{λ} stands for the matrix of B-spline functions, then

$$S_{\lambda} = B_{\lambda}(B'_{\lambda}B_{\lambda})^{-1}B_{\lambda}$$

What distinguishes the smoothing matrices of a **smooth** fit and a **non-smooth** fit?



Spline smoothing

Two extreme smoothing matrices:

- The averaging matrix (the smoothest possible):

$$\hat{Y}_i = \bar{Y}, \text{ for all } i = 1, \dots, n,$$

where all elements in S_λ equal $\frac{1}{n}$

- The identity matrix (the least smooth):

$$\hat{Y}_i = Y_i, \text{ for all } i = 1, \dots, n, \text{ where } S_\lambda = I_n$$

Nonparametric degree of freedom: $df_\lambda = \text{trace}(S_\lambda)$ is interpreted as an *equivalent number of parameters*.

► Nonparametric degree of freedom using \mathbb{R}



Spline smoothing

Is the effect of temperature on Ozone concentration significantly nonlinear?



Nonparametric vs parametric model

Suppose \mathcal{M}_0 is a parametric submodel of the regression model $\mathcal{M}: Y = f(x) + \varepsilon$.

Let d_0 and d denote the residual degrees of freedom of \mathcal{M}_0 and \mathcal{M} respectively.

Then, for the test of $H_0: \mathcal{M}$ is not better than \mathcal{M}_0 , the test statistics is the nonlinear F-test:

$$F = \frac{\frac{\text{RSS}_0 - \text{RSS}}{d_0 - d}}{\frac{\text{RSS}}{d}}$$

and the null distribution of F is the Fisher distribution with $d_0 - d$ and d degrees of freedom.



Nonparametric vs parametric model

Is there a gain in prediction accuracy of the present model
w.r.t the linear model?

► Prediction performance of a nonparametric model using \mathbb{R}



Nonparametric vs parametric model

How to improve the prediction accuracy by completing the profile of explanatory variables?



Additive regression models

Suppose $x = (x_1, \dots, x_p)$ is a profile of explanatory variables:

$$Y = \beta_0 + f_1(x_1) + \dots + f_p(x_p) + \varepsilon, \quad \varepsilon \sim \mathcal{N}(0; \sigma)$$

Estimation using a *Backfitting* algorithm :

- Initialization: $\hat{\beta}_0 = \bar{Y}$, $\hat{f}_j = \hat{f}_j^{(0)}$
- Cycling over the marginal effects: if \hat{f}_j , $j \neq k$, are the current estimates, update f_k :

$$\left[Y - \hat{\beta}_0 - \sum_{j=1, j \neq k}^p \hat{f}_j(x_j) \right] = f_k(x^{(k)}) + \varepsilon,$$

- Iteration until convergence.

► Fitting full GAM using R



Additive regression models

Is it relevant to consider all the explanatory variables in the model?

▶ Nonparametric ANOVA using R



Nonparametric model selection

The **Akaike Information Criterion** for the following additive model

$$Y = \beta_0 + f_1(x_1) + \dots + f_p(x_p) + \varepsilon, \varepsilon \sim \mathcal{N}(0; \sigma)$$

with k nonparametric degrees of freedom is:

$$\text{AIC} \propto n \log\left(\frac{\text{RSS}}{n}\right) + 2k$$

Stepwise model selection for gam is implemented in R package

gam: `step.Gam`.

► Stepwise nonparametric model selection using R