# Statistical Learning and Prediction

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#### **Beyond Linearity**

#### Four topics:

- Polynomial regression
- Step functions
- Polynomial regression + step functions = splines:
- Support Vector Classifier
- Support Vector Machine

#### **Polynomial Regression**

- The easiest form of (parametric) nonlinearity is to use a polynomial.
- E.g., a cubic (third order) polynomial:

$$y_i = \alpha + \beta_1 x + \beta_2 x^2 + \beta_3 x^3 + \epsilon_i$$

• More generally, polynomials of order *d*:

$$y_i = \alpha + \sum_{j=1}^d \beta_j x^j + \epsilon_i$$

• Higher order polynomials are very flexible functions (sometimes too flexible)

#### **Polynomial Regression**

• Same trick can be applied in logistic regression (and other generalized linear models, e.g., binomial, count responses):

$$\log\left(\frac{\pi_i}{1-\pi_i}\right) = \alpha + \sum_{j=1}^d \beta_j x^j$$

- No interest in the individual coefficients (difficult to interpret, what is large / small effect?)
- Interest in the shape of the association between predictor and response

#### **Step functions**

• Define cut points  $c_1, c_2, c_3, \ldots, c_K$  and with these functions

$$C_0(X) = I(X < c_1)$$
 $C_1(X) = I(c_1 \le X < c_2)$ 
...
 $C_{K-1} = I(c_{K-1} \le X < c_K)$ 
 $C_K(X) = I(c_K \le X)$ 

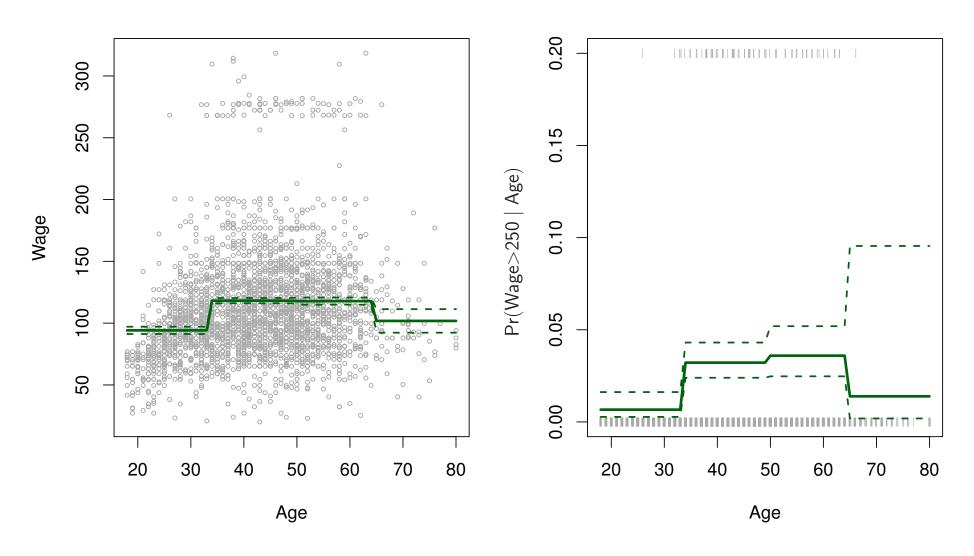
• Use these  $C_k(X)$  as predictors in a linear/logistic regression, e.g.,

$$y_i = \alpha + \sum_{k=1}^K \beta_k C_k(X) + \epsilon_i$$

• Step functions can be used to describe regression and classification trees (where they can be a function of multiple predictor variables; see later)

#### **Example - Step function**

#### **Piecewise Constant**



#### **Basis functions**

- Both polynomials and step functions are special cases of more general *basis functions*:
- Define functions/transformations  $b_1(X), b_2(X), \ldots, b_K(X)$  of the variable X.
- Use these transformations in a linear regression

$$y_i = \alpha + \sum_{k=1}^K \beta_k b_k(X) + \epsilon_i$$

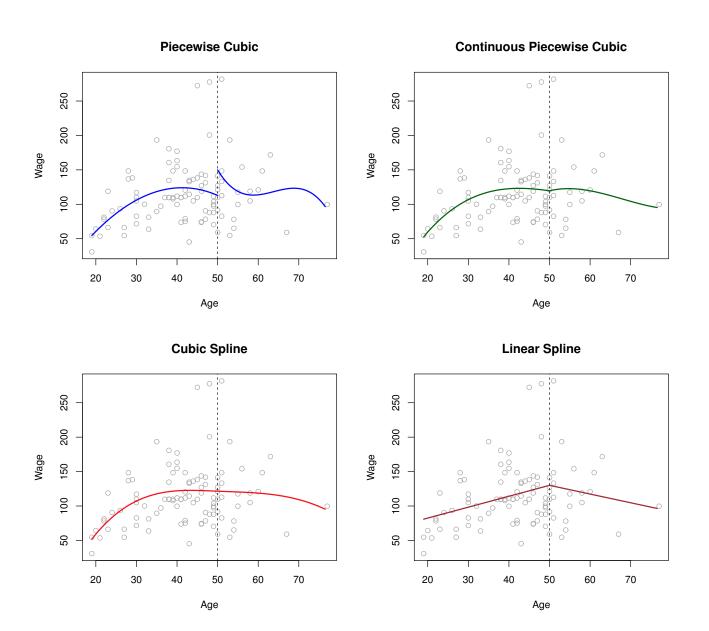
• The functions  $b_k(X)$  are fixed and known.

- Combine the step idea with the polynomial functions
- Create a cutpoint *c*:

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

- This is a piecewise polynomial with 1 knot (cutpoint).
- See next slide: upper left. Can give erratic behavior near the boundaries.

#### **Example - Regression Splines**



- At cutpoint *c* it makes a sudden jump (it is discontinuous).
- We rather have a continuous curve.
- Therefore, we add the constraint that the fitted curve must be continuous (upper right plot).
- Better, but V-shaped join looks unnatural.
- Add two constraints: first- and second-order derivatives of the polynomial curve have to be continuous.
- Thus, we require that the curve is very *smooth*.
- Lower left plot: cubic spline.
- A cubic spline with K knots uses a total of K+4 degrees of freedom.

- Lower right plot: *linear spline* with one knot, K = 1 (a.k.a. hinge function).
- General definition: A degree-d spline is a piecewise degree d polynomial with continuity in the derivatives up to degree d-1.
- In the figures we used a single knot, but we can increase the number of knots, giving more flexible curves.
- Note we have again a bias-variance trade-off: the more knots and the higher the order of the polynomial, the less bias, but the more variance.

• A cubic spline with *K* knots can be modelled as

$$y_i = \beta_0 + \beta_1 b_1(x_i) + \beta_2 b_2(x_i) + \ldots + \beta_{K+3} b_{K+3}(x_i) + \epsilon_i$$

• The basis functions are given by:

$$b_1(x) = x$$
 $b_2(x) = x^2$ 
 $b_3(x) = x^3$ 
 $b_{3+j}(x,\xi_j) = (x-\xi_j)_+^3 = \begin{cases} (x-\xi_j)^3 & \text{if } x>\xi_j \\ 0 & \text{otherwise} \end{cases}$ 

for 
$$j = 1, ..., K$$

• The function has an intercept and K + 3 regression weights, thus using K + 4 degrees of freedom.

- Unfortunately, a cubic spline often still has large variance at the outer ranges of the predictors.
- A *natural spline* is a regression spline with additional boundary contraints: the function is constrained to be linear at the boundaries.
- This uses lower degrees of freedom: quadratic and cubic effects are zero at both boundaries, yielding *K* degrees of freedom.

#### Choosing the number of knots

- Where should we place the knots?
  - Prior knowledge / information.
  - Place the knots in a uniform way, for example based on quantiles.
- How many knots should we use? (or equivalently: How many degrees of freedom?)
  - Prior knowledge / information.
  - Determine by cross validation.

#### **Smoothing Spline**

 Instead of working with a set of basis functions it is also possible to use the fit + penalty approach to fit smooth functions:

$$\sum_{i=1}^{n} (y_i - g(x_i))^2 + \lambda \int g''(t)^2 dt$$

- with  $\lambda$  a non-negative smoothing parameter, which penalizes wiggliness of the fitted function, thereby controlling the bias-variance tradeof.
- The function *g* that minimizes this function is known as a *smoothing spline*.
- $g''(t)^2$  is the second derivative of the function g; the amount by which the slope is changing, i.e. a measure of wigglyness

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- The function *g* that minimizes this function is known as a *smoothing spline*; it consists of basis functions, too.
- $g''(t)^2$  is the second derivative of the function g; the amount by which the slope is changing, i.e. a measure of wigglyness

#### **Generalized Additive Models**

- Often we have multiple predictors:  $x_{i1}, \ldots, x_{ip}$
- We can generalize the ideas:

$$y_i = \beta_0 + \sum_{j=1}^p f_j(x_{ij}) + \epsilon_i$$

- where  $f_j()$  are smooth nonlinear functions (polynomials, cubic splines, natural splines, smoothing splines).
- It is an additive model, the smooth versions of each variable are added.
- Can be generalized to other GLM response variable types (e.g., binomial, count responses).

#### Exercise - splines

- Get the nesda.sav file (Netherlands Study of Depression and Anxiety) from the github repo.
- Read it into R as follows:

```
library("foreign")
read.spss("nesda.sav", use.value.labels=F, to.data.frame=T)
```

- Separate the observations into training (N=400) and test (N=200) parts.
- Use function gam from package mgcv.
- Fit a smoothing spline, predicting mdd based on s (aconscie).
- Set the method argument of function gam () to "REML".
- Use the summary and plot functions to inspect and interpret the result.

#### Exercise - splines

- Type ?s and check out the meaning of the k, bs and sp arguments.
- Repeat the analysis, but now fit an unpenalized smoothing spline: s(aconscie, sp = 0). Also fit a spline using 20 basis functions s(aconscie, k = 20).
- Use the summary and plot functions to inspect and compare the results.
- Fit a GAM containing smoothing splines of both aconscie and neurot as predictors of mdd. Evaluate the misclassification rate on the test dataset, using the predict () function.

#### **Support Vector Machines**

- Maximum Margin Classifier
- Support Vector Classifier
- Support Vector Machine

### Maximum Margin Classifier

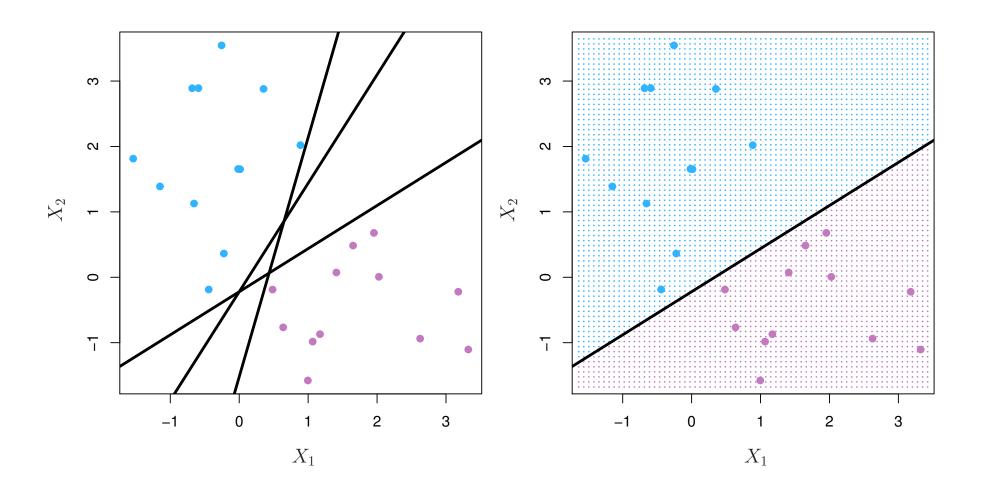
- Classification based on a separating hyperplane.
- That is, finding in the feature space a (hyper)plane that separates the two classes.
- A hyperplane can be seen as an equation of the feature variables:

$$\beta_0 + \beta_1 X_1 + \beta_2 X_2 = 0$$

• If we code  $y \in \{-1,1\}$  then the separating hyperplane has the following property

$$y_i (\beta_0 + \beta_1 x_{i1} + \beta_2 x_{i2}) > 0$$

## **Example - Seperating hyperplane**

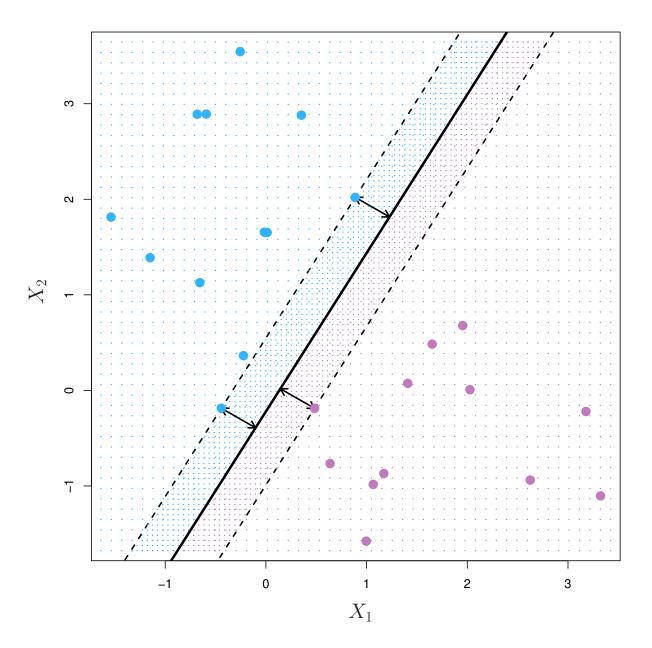


#### Maximum Margin Classifier

- There may be several perfectly separating hyperplanes, then the one with the largest margin is chosen.
- This amounts to:

maximize 
$$M(\beta_0, ..., \beta_p)$$
  
subject to  $\sum_{j=1}^p \beta_j^2 = 1$   
 $y_i (\beta_0 + \beta_1 x_{i1} + \beta_2 x_{i2}) \ge M \ \forall i = 1, ..., n$ 

# Example - Maximum Margin Classifier



- Often no perfectly separating hyperplane exists.
- Generalize the maximum margin classifier idea using a soft margin.
- We allow some observations to be on the wrong side of the margin or even on the wrong side of the hyperplane.
- We introduce *slack variables*:  $\epsilon_1, \dots, \epsilon_n$  and a tuning parameter C:

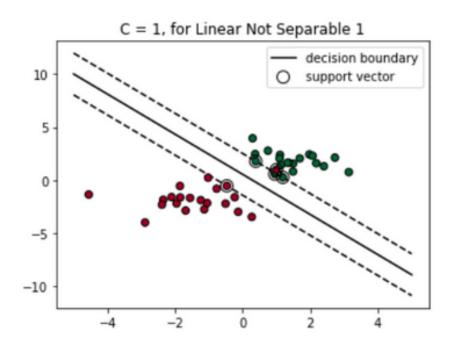
maximize 
$$M(\beta_0, \ldots, \beta_p, \epsilon_1, \ldots, \epsilon_n)$$

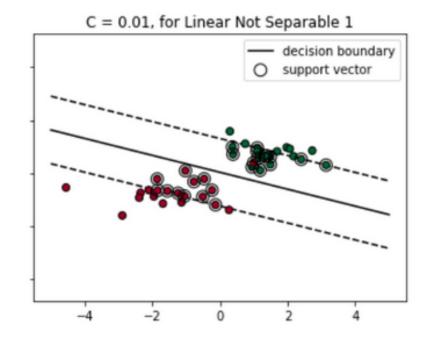
subject to 
$$\sum_{j=1}^{p} \beta_{j}^{2} = 1$$

$$y_{i} (\beta_{0} + \beta_{1}x_{i1} + \beta_{2}x_{i2}) \geq M(1 - \epsilon_{i}) \quad \forall i = 1, ..., n$$

$$\epsilon_{i} \geq 0, \sum_{i=1}^{n} \epsilon_{i} \leq C$$

- The *slack variables*:  $\epsilon_1, \dots, \epsilon_n$  tell us where the *i*-th observation is located, relative to the margin and hyperplane:
  - If  $\epsilon_i = 0$ , the observation is on the correct side of the margin.
  - If  $\epsilon_i > 0$ , the observation is on the incorrect side of the margin.
  - If  $\epsilon_i > 1$ , the observation is on the incorrect side of the hyperplane.
- The tuning parameter *C* functions like a budget:
  - Thus, if C = 0, there is no budget for violations.
  - As C increases, the classifier becomes more tolerant of errors (allows for more *slack*) and the margin becomes wider.





- Observations that are on the margin or on the wrong side of the margin are called *support vectors*
- The decision rule is based only on a small fraction of training observations, which makes it quite robust to behavior of observations far from the hyperplane.
- In contrast, Linear Discriminant Analysis depends on the mean of all observations to construct the discriminant function and is therefore sensitive to points far from the decision line.

#### **Support Vector Machine**

- In practice the decision boundary is often nonlinear
- We can enlarge the feature space to allow for nonlinear boundaries
- Instead of using a support vector classifier on

$$X_1, X_2, \ldots, X_p$$

use

$$X_1, X_1^2, X_2, X_2^2, \ldots, X_p, X_p^2$$

or

$$X_1, X_1^2, X_2, X_2^2, \ldots, X_p, X_p^2, X_1X_2, X_1X_3, \ldots$$

• In the enlarged space the decision boundary is linear, in the original space the decision boundary is nonlinear.

#### **Support Vector Machine**

- There are many ways to enlarge the feature space by using all kinds of functions of the different *X*s.
- The computational burden increases rather fast with increasing number of features.

# **Example Inner Product**

	mdd	aconscie	neurot
1	0	-1.10	-0.22
2	0	-0.09	-1.42
3	0	-1.27	-0.33
4	1	-0.93	0.55
5	0	-0.59	-0.43

#### **Example Inner product**

- $\langle x_i, x_{i'} \rangle$  is the *inner product* defined by  $\langle x_i, x_{i'} \rangle = \sum_{j=1}^p x_{ij} x_{i'j}$
- For person 1 and 2

$$\langle x_1, x_2 \rangle = -1.10 \times -0.09 + -0.22 \times -1.42 = 0.41$$

- It is the 'covariance' between the observations for person 1 and 2
- The inner product is the correlation or more general a measure of similarity
- Important to first standardize the variables, otherwise a variable may be dominant

# **Example Inner product**

	1	2	3	4	5
1	1.25	0.41	1.46	0.90	0.75
2	0.41	2.03	0.58	-0.70	0.67
3	1.46	0.58	1.71	1.00	0.89
4	0.90	-0.70	1.00	1.17	0.31
5	0.75	0.67	0.89	0.31	0.54

#### **Support Vector Machine**

• The linear support vector classifier can be represented as:

$$f(x) = \beta_0 + \sum_{i=1}^n \alpha_i < x, x_i >$$

- where  $\langle x_i, x_i' \rangle$  is the *inner product* defined by  $\langle x_i, x_i' \rangle = \sum_{j=1}^p x_{ij} x_{i'j}$
- The inner product is the correlation or more general a measure of similarity.
- It is possible to define the matrix *K* with all inner products:

$$K(x_i, x_{i'}) = \sum_{j=1}^{p} x_{ij} x_{i'j}$$

• The K is called the kernel, and is an  $n \times n$  matrix.

• It is possible to generalize the kernel, for example:

$$K(x_i, x_{i'}) = (1 + \sum_{j=1}^{p} x_{ij} x_{i'j})^d$$

gives a polynomial kernel of degree d. It is still a  $n \times n$  matrix.

- Using this kernel amounts to fitting a linear support vector classifier in a higher-dimensional space involving polynomials of degree *d*.
- Another choice of kernel is the radial basis kernel which is defined by:

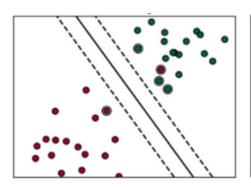
$$K(x_i, x_{i'}) = \exp(-\gamma \sum_{j=1}^{p} (x_{ij} - x_{i'j})^2)$$

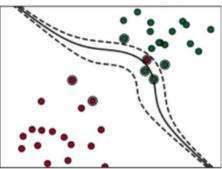
For this kernel the *implicit* feature space is infinite dimensional.

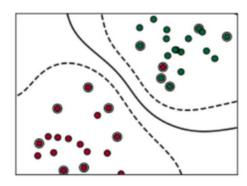
 $\gamma$  is a tuning parameter, with higher values yielding a smoother decision boundary (i.e., higher bias and lower variance).

- In order to use kernels properly it is important to scale the variables first (different scalings give different kernels give different solutions).
- The main advantage of kernels is that they avoid the need to actually transform to an enlarged space; the only thing needed is to compute the kernel and apply the support vector classifier on this kernel.
- This is alo known as the kernel trick.

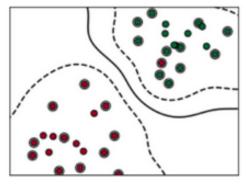
#### **Support Vector Machine**

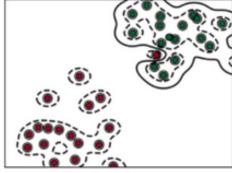


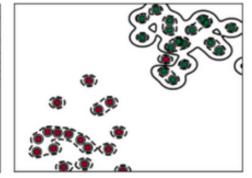




Linear, polynomial, radial basis kernels







Radial basis kernel with  $\gamma = 1$ , 10 or 20.

#### **Exercise - Support Vectors**

- Use aconscie en neurot as predictors
- Use mdd as response (first make it a factor)
- Use the library **e1071**
- To find optimal tuning parameters use the function tune () out=tune (svm, formula, data, kernel="..", ranges=list(cost=c(0.001,..,100)))
- With the optimal budget (cost) fit the model, make a plot and predictions for the test data set:

```
svmfit=svm(formula, data, kernel="..", cost=..,
scale=FALSE)
```

#### **Exercise - Support Vectors**

• Fit a support vector machine with radial basis kernel, tuning the budget and  $\gamma$  parameter::

```
(ranges=list(cost=c(0.1,1,10,100,1000), gamma=c(0.5,1,2,3,4)))
```

ullet With the optimal budget (cost) and  $\gamma$  fit the model again, make a plot and predictions for the test data set

#### **Exercises from book**

• Chapter 7: 1, 4, 7, 8.

For 7 and 8, it is helpful to first make a correlation plot (e.g., plot (Wage)), and then pick a small number of predictors you want to include in your model.

• Chapter 9: 1, 2, 3, 8.