

# Linear and k-NN Methods for Classification and Their Extensions

- likelihood example
- logistic regression
  - ext. logistic regression with  $L_1$  penalty, elastic net penalty
- linear and quadratic discriminant analysis
  - ext. regularized discriminant analysis
  - ext. reduced rank discriminant analysis
  - ext. diagonal discriminant analysis
- Nearest-neighbor methods
  - k-NN
  - Local likelihood (local logistic regression)
  - ext. Discriminating Adaptive NN methods (DANN)
- ? Support Vector Machines

# Probability of the data given the model

- Assume we have 15 red balls and 5 blue balls in a bag.
- Repeat 5x:
  - select a ball
  - put it back.
- The probability of the sequence red, blue, blue, red, red is  $\frac{3}{4} \cdot \frac{1}{4} \cdot \frac{1}{4} \cdot \frac{3}{4} \cdot \frac{3}{4}$ .
- The logarithm  $\log_2$  of the probability is  $\approx -0.4 - 2 - 2 - 0.4 - 0.4 = -5.2$

# Likelihood of the model given the data

- Assume we do not know the probabilities, let  $\theta$  be the probability of *red*. We have following probabilities of data for different  $\theta$ .

$\theta$	red	blue	blue	red	red	
$\frac{3}{4}$	$\frac{3}{4}$	$\frac{1}{4}$	$\frac{1}{4}$	$\frac{3}{4}$	$\frac{3}{4}$	$\frac{3^3}{4^5}$

- Take the  $\log_2$  of the probabilities:

$\theta$	red	blue	blue	red	red	
$\frac{1}{2}$	-1	-1	-1	-1	-1	-5
$\frac{3}{5}$	-0.74	-1.32	-1.32	-0.74	-0.74	-4.86
$\frac{3}{4}$	-0.4	-2	-2	-0.4	-0.4	-5.2

- Probability of the data given model is called **likelihood** of the model  $\theta$  given the data.
- Maximum likelihood  $\theta$  estimate is in our case  $\frac{3}{5}$ .
- Predicting probabilities, maximum likelihood estimate is the same as maximum log-likelihood estimate.

# (Log)likelihood

train data		prediction			likelihood	loglik
$x_i$	$g_i$	$P(\text{green} x_i)$	$P(\text{blue} x_i)$	$P(\text{yellow} x_i)$		
1	green	$\frac{1}{2}$	0	$\frac{1}{2}$	$\frac{1}{2}$	-1
1	yellow	$\frac{1}{2}$	0	$\frac{1}{2}$	$\frac{1}{2}$	-1
2	green	$\frac{2}{3}$	$\frac{1}{3}$	0	$\frac{2}{3}$	$\log_2 \frac{2}{3}$
2	green	$\frac{2}{3}$	$\frac{1}{3}$	0	$\frac{2}{3}$	$\log_2 \frac{2}{3}$
2	blue	$\frac{2}{3}$	$\frac{1}{3}$	0	$\frac{1}{3}$	$-\log_2 3$
3	blue	0	1	0	1	0
						$-2 - \log_2 3$ $+2\log_2 \frac{2}{3}$

- **loglik** logarithm with base  $e$  of likelihood function is defined as:

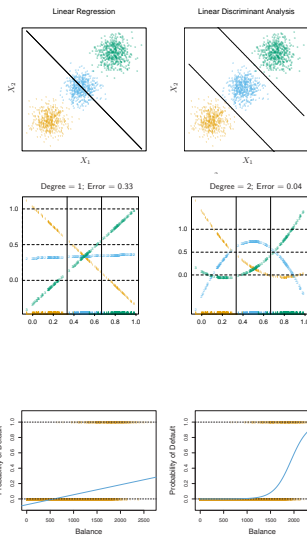
$$\ell(\theta) = \sum_{i=1}^N \log_e(P(G = g_i | x_i, \theta))$$

- Logistic regression uses:

$$P(G = g_k | X = x) = \frac{e^{\beta_k 0 + \beta_k^T x}}{1 + \sum_{l=1, \dots, K-1} e^{\beta_l 0 + \beta_l^T x}}$$

# Logistic Function

- With more than two classes, the linear regression to the class identifier may suffer the **masking problem**.
  - Some class is masked due to the linearity constraint (the blue class in the figure right).
  - Quadratic fit to the identifier function solves this case.
  - Logistic regression, LDA, SVM, k-NN solve this case naturally.
- Probability should be from the interval  $(0, 1)$ .
- Linear prediction is transformed by logistic function (sigmoid) with the maximum  $L$ .
- **logistic**  $\frac{L}{1+e^{-k(x-x_0)}}$ .
- Inverse function is called logit.
- **logit**  $\log \frac{p}{1-p}$ ,



# Logistic Regression

- For  $K$ -class classification we estimate  $(p + 1) \times (K - 1)$  parameters  $\theta = \{\beta_{10}, \beta_1^T, \dots, \beta_{(K-1)0}, \beta_{K-1}^T\}$ .

$$\begin{aligned}\log \frac{P(G = g_1 | X = x)}{P(G = g_K | X = x)} &= \beta_{10} + \beta_1^T x \\ \log \frac{P(G = g_2 | X = x)}{P(G = g_K | X = x)} &= \beta_{20} + \beta_2^T x \\ &\vdots \\ \log \frac{P(G = g_{K-1} | X = x)}{P(G = g_K | X = x)} &= \beta_{(K-1)0} + \beta_{K-1}^T x\end{aligned}$$

that is

$$\begin{aligned}p_k(x; \theta) \leftarrow P(G = g_k | X = x) &= \frac{e^{\beta_{k0} + \beta_k^T x}}{1 + \sum_{l=1, \dots, K-1} e^{\beta_{l0} + \beta_l^T x}} \\ p_K(x; \theta) \leftarrow P(G = g_K | X = x) &= \frac{1}{1 + \sum_{l=1, \dots, K-1} e^{\beta_{l0} + \beta_l^T x}}.\end{aligned}$$

# Fitting Logistic Regression Two class

- This model is estimated iteratively maximizing conditional likelihood of  $G$  given  $X$ .

$$\ell(\theta) = \sum_{i=1}^N \log p_{g_i}(x_i; \theta)$$

- Two class model:  $g_i$  encoded via a 0/1 response  $y_i$ ;  $y_i = 1$  iff  $g_k = g_1$ . Let  $p(x; \theta) = p_1(x; \theta)$ ,  $p_2(x; \theta) = 1 - p(x; \theta)$ . Then:

$$\begin{aligned}\ell(\theta) &= \sum_{i=1}^N (y_i \log p(x_i; \beta) + (1 - y_i) \log(1 - p(x_i; \beta))) \\ &= \sum_{i=1}^N (y_i \beta^T x_i - \log(1 + e^{\beta^T x_i}))\end{aligned}$$

- Set derivatives to zero:

$$\frac{\partial \ell(\beta)}{\partial \beta} = \sum_{i=1}^N x_i (y_i - p(x_i; \beta)) = 0,$$

- which is  $p + 1$  nonlinear equations in  $\beta$ .
- First component:  $x_i \equiv 1$  specifies  $\sum_{i=1}^N y_i = \sum_{i=1}^N p(x_i; \beta)$  the expected

# Newton–Raphson Algorithm

- We use Newton–Raphson Algorithm to solve the system of equations

$$\frac{\partial \ell(\beta)}{\partial \beta} = \sum_{i=1}^N x_i (y_i - p(x_i; \beta)) = 0,$$

- we need the second–derivative or Hessian matrix

$$\frac{\partial^2 \ell(\beta)}{\partial \beta \partial \beta^T} = - \sum_{i=1}^N x_i x_i^T p(x_i; \beta) (1 - p(x_i; \beta)).$$

- Starting with  $\beta^{old}$  a single Newton–Raphson update is

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

- where the derivatives are evaluated at  $\beta^{old}$ .



# Newton–Raphson Algorithm in Matrix Notation

Let us denote:

$\mathbf{y}$  the vector of  $y_i$

$\mathbf{X}$   $N \times (p + 1)$  data matrix  $x_i$

$\mathbf{p}$  the vector of fitted probabilities with  $i$ th element  $p(x_i; \beta^{old})$

$\mathbf{W}$  diagonal matrix with weights  $p(x_i; \beta^{old})(1 - p(x_i; \beta^{old}))$

$$\frac{\partial \ell(\beta)}{\partial \beta} = \mathbf{X}^T (\mathbf{y} - \mathbf{p})$$

$$\frac{\partial^2 \ell(\beta)}{\partial \beta \partial \beta^T} = -\mathbf{X}^T \mathbf{W} \mathbf{X}$$

The Newton–Raphson step is ( $\beta^0 \leftarrow 0$ )

$$\beta^{new} = \beta^{old} + (\mathbf{X}^T \mathbf{W} \mathbf{X})^{-1} \mathbf{X}^T (\mathbf{y} - \mathbf{p})$$

$$= (\mathbf{X}^T \mathbf{W} \mathbf{X})^{-1} \mathbf{X}^T \mathbf{W} (\mathbf{X} \beta^{old} + \mathbf{W}^{-1} (\mathbf{y} - \mathbf{p}))$$

$$= (\mathbf{X}^T \mathbf{W} \mathbf{X})^{-1} \mathbf{X}^T \mathbf{W} \mathbf{z}$$

$$\mathbf{z} = \mathbf{X} \beta^{old} + \mathbf{W}^{-1} (\mathbf{y} - \mathbf{p}) \quad \text{adjusted response}$$

- $p$ ,  $W$ ,  $z$  change each step

This algorithm is referred to as **iteratively reweighted least squares IRLS**

$$\beta^{new} \leftarrow \arg \min_{\beta} (\mathbf{z} - \mathbf{X} \beta)^T \mathbf{W} (\mathbf{z} - \mathbf{X} \beta)$$

# South African Heart Disease

- Analyzing the risk factors of myocardian infarction MI
- (Note: prevalence 5.1%, in the data 160 positive 302 controls, the controls are underrepresented, consider weighting the data.)

TABLE 4.2. Results from a logistic regression fit to the South African heart disease data.

	Coefficient	Std. Error	Z Score
(Intercept)	-4.130	0.964	-4.285
sbp	0.006	0.006	1.023
tobacco	0.080	0.026	3.034
ldl	0.185	0.057	3.219
famhist	0.939	0.225	4.178
obesity	-0.035	0.029	-1.187
alcohol	0.001	0.004	0.136
age	0.043	0.010	4.184

- Wald test: Z score  $|Z| > 2$  is significant at at the 5% level.

TABLE 4.3. Results from stepwise logistic regression fit to South African heart disease data.

	Coefficient	Std. Error	Z score
(Intercept)	-4.204	0.498	-8.45
tobacco	0.081	0.026	3.16
ldl	0.168	0.054	3.09
famhist	0.924	0.223	4.14
age	0.044	0.010	4.52

# South African Heart Disease

- Wald test:  $Z$  score  $|Z| > 2$  is significant at at the 5% level.

TABLE 4.3. Results from stepwise logistic regression fit to South African heart disease data.

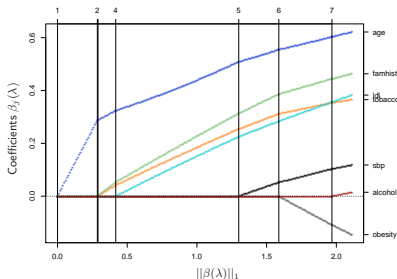
	Coefficient	Std. Error	Z score
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famhist	0.924	0.223	4.14
age	0.044	0.010	4.52

- $P(MI|x_i, \theta) = \frac{e^{-4.204+0.081x_{tobacco}+0.168x_{ldl}+0.924x_{famhist}+0.044x_{age}}}{1+(e^{-4.204+0.081x_{tobacco}+0.168x_{ldl}+0.924x_{famhist}+0.044x_{age}})}$
- Interval estimate  $odds_{tobacco} = e^{0.081 \pm 2 \times 0.026} = (1.03, 1.14)$  increase of odds of  $MI$  based of the increase of  $x_{tobacco}$ .

# $L_1$ regularization 'Lasso'-like

$$\operatorname{argmax}_{\beta_0, \beta} \left( \sum_{i=1}^N (y_i (\beta_0 + \beta^T x_i) - \log(1 + e^{(\beta_0 + \beta^T x_i)})) - \lambda \sum_{j=1}^p |\beta_j| \right)$$

- Newton–Raphson Algorithm or nonlinear programming.
- $\lambda = 0$  standard logistic regression.
- $\lambda \rightarrow \infty$  moves coefficients towards 0.
- $\beta_0$  is not included into the penalty.



# Linear Discriminant Analysis

- LDA assumes multivariate gaussian distribution of each class with a common covariance matrix.

$$\phi(k) = \frac{1}{\sqrt{|2\pi\Sigma|}} e^{-\frac{1}{2}(x-\mu_k)^T \Sigma^{-1}(x-\mu_k)}$$

- Under this assumptions it provides bayes optimal estimate.
- Different covariance matrix for each class leads to Quadratic Discriminant Analysis.
- Let us denote  $N_k$  number of training data in the class  $G_k$ .

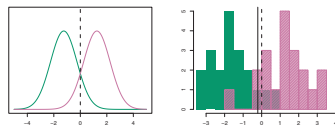
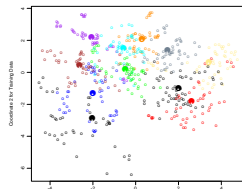


FIGURE 4.4. Left: Two one-dimensional normal density functions are shown.

# Linear Discriminant Analysis

The LDA model parameters: the mean and probability of each class  $\{\mu_i, \pi_i\}_{i=1}^K$  and the common covariance matrix  $\Sigma$  can be evaluated directly.

$$\hat{\pi}_k = \frac{N_k}{N}$$

$$\hat{\mu}_k = \frac{\sum_{\{x_i: G(x_i)=g_k\}} x_i}{N_k}$$

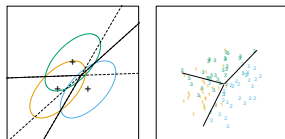
$$\hat{\Sigma} = \sum_{k=1}^K \sum_{\{x_i: G(x_i)=g_k\}} \frac{(x_i - \mu_k)(x_i - \mu_k)^T}{(N - K)}$$

$$\phi_k(x) = N(\mu_k, \Sigma)$$

$$P(G = g_k | X = x) = \frac{\phi_k(x)\pi_k}{\sum_{\ell=1}^K \phi_{\ell}(x)\pi_{\ell}}$$

To classify new instance  $x$  we predict the  $G_k$  with maximal  $\delta_k$ :

$$\delta_k(x) = x^T \Sigma^{-1} \mu_k - \frac{1}{2} \mu_k^T \Sigma^{-1} \mu_k + \log.$$



# Quadratic Discriminant Analysis

Quadratic discriminant analysis estimates the covariance matrix for each class independently. The rest is the same as for the LDA.

$$\begin{aligned}\hat{\pi}_k &= \frac{N_k}{N} \\ \hat{\mu}_k &= \frac{\sum_{\{x_i: G(x_i)=g_k\}} x_i}{N_k} \\ \hat{\Sigma}_k &= \sum_{\{x_i: G(x_i)=g_k\}} \frac{(x_i - \mu_k)(x_i - \mu_k)^T}{(|G_k| - 1)} \\ f_k(x) &= N(\mu_k, \Sigma_k) \\ P(G = g_k | X = x) &= \frac{f_k(x) \pi_k}{\sum_{\ell=1}^K f_\ell(x) \pi_\ell}\end{aligned}$$

To classify new instance  $x$  we predict the  $G_k$  with maximal  $\delta_k$ :

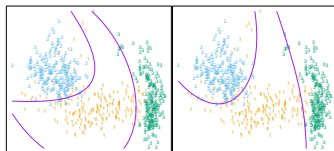
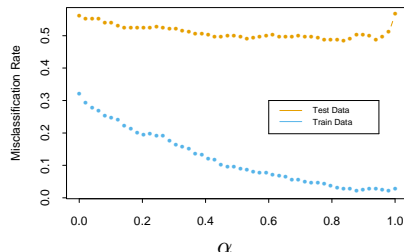
$$\delta_k(x) = -\frac{1}{2}(x - \mu_k)^T \Sigma_k^{-1} (x - \mu_k) - \frac{1}{2} \log |\Sigma_k| + \log \pi_k.$$

# Quadratic and Regularized Discriminant Analysis

- QDA has substantially more parameters. It is questionable whether it is worth to increase the model complexity.
  - LDA parameters:  $(K - 1) \times (p + 1)$
  - QDA parameters:  $(K - 1) \times (\frac{p(p+3)}{2} + 1)$ .
- **Regularized discriminant analysis** takes a weighted average of LDA and QDA to tune the model complexity.

$$\hat{\Sigma}_k(\alpha) = \alpha \hat{\Sigma}_k + (1 - \alpha) \hat{\Sigma}$$

Regularized Discriminant Analysis on the Vowel Data



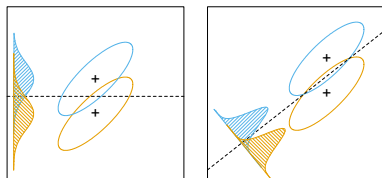
**FIGURE 4.6.** Two methods for fitting quadratic boundaries. The left plot shows the quadratic decision boundaries for the data in Figure 4.1 (obtained using LDA in the five-dimensional space  $X_1, X_2, X_1X_2, X_1^2, X_2^2$ ). The right plot shows the



## Linear and Quadratic Discriminant Analysis

- $O(N^3)$ , often  $O(N^{2.376})$
- QDA and LDA may be computed using matrix decomposition:
  - Compute the eigendecomposition for each  $(x - \hat{\mu}_k)^T \hat{\Sigma}_k^{-1} (x - \hat{\mu}_k) = [\mathbf{U}_k^T (x - \hat{\mu}_k)]^T \mathbf{D}_k^{-1} [\mathbf{U}_k^T (x - \hat{\mu}_k)]$
  - $\log |\hat{\Sigma}_k| = \sum_{\ell} \log d_{k\ell}$ .
- Using this decomposition, LDA classifier can be implemented by the following pair of steps:
  - *Sphere* the data with respect to the common covariance estimate  $\hat{\Sigma}$ :  
 $X^* \leftarrow \mathbf{D}^{-\frac{1}{2}} \mathbf{U}^T X$ , where  $\hat{\Sigma} = \mathbf{U} \mathbf{D} \mathbf{U}^T$ .  
The common covariance estimate of  $X^*$  will now be the identity.
  - Classify to the closest class centroid in the transformed space, modulo the effect of the class prior probabilities  $\pi_k$ .

# Reduced-Rank Linear Discriminant Analysis



$$\max_a \frac{a^T B a}{a^T W a}$$

Finding the sequence of optimal subspaces for LDA:

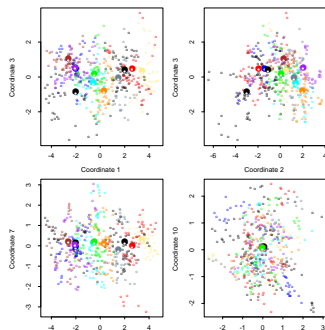
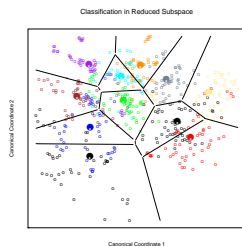
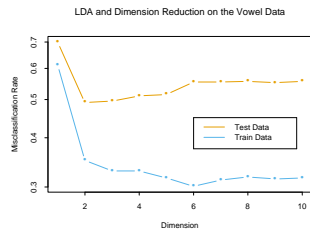
- compute the  $K \times p$  matrix of class centroids  $M$  and the common covariance matrix  $W$  (**within-class** covariance);
- compute  $M^* = MW^{-\frac{1}{2}}$  using the eigen-decomposition of  $W$ ;
- compute  $B^*$  **between-class** covariance, the covariance matrix of  $M^*$  and its eigen-decomposition  $B^* = V^* D_B V^{*T}$ .
  - order  $D_B$  in the decreasing order
  - $v_\ell^*$  of  $V^*$  in sequence define the coordinates of the optimal subspaces
  - $Z_\ell = v_\ell^T X$  with  $v_\ell = W^{-\frac{1}{2}} v_\ell^*$ .

# Vowel Example

Example Vowel data ESL:

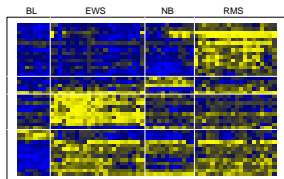
- $X \in \mathbb{R}^{10}$ :
- $k = 11$  classes.

	train	test
Linear regression	0.48	0.67
Linear discriminant analysis	0.32	0.56
Quadratic discriminant analysis	0.01	0.53
Logistic regression	0.22	0.51



# Diagonal Linear Discriminant Analysis

- **With really many dimensions** the even reduced rank does not work.
- Gene expression experiment
  - 2308 genes (columns)
  - 63 samples (rows), from a set of microarray experiments.
  - The samples arose from small, round blue-cell tumors (SRBCT) found in children, and are classified into four major types:
    - BL (Burkitt lymphoma),
    - EWS (Ewing's sarcoma),
    - NB (neuroblastoma),
    - and RMS (rhabdomyosarcoma).
  - There is an additional test data set of 20 observations.



• diagonal-covariance LDA

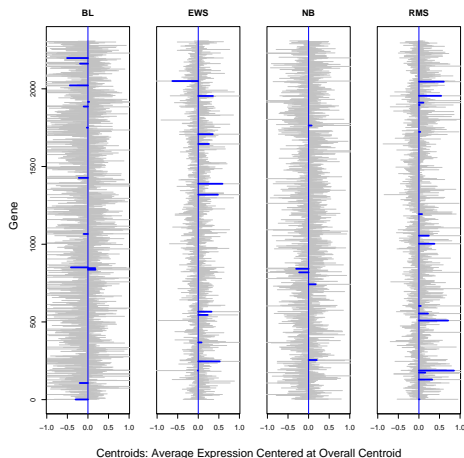
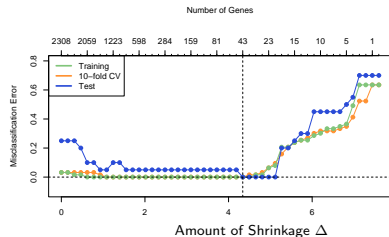
$$\delta_k(x^*) = - \sum_{j=1}^p \frac{(x_j^* - \bar{x}_{jk})^2}{s_j^2} + 2 \log(\pi_k)$$

- $s_j$  is the pooled within-class standard deviation of the  $j$ th gene
- $\bar{x}_{jk} = \sum_{i \in C_k} \frac{x_{ij}}{N_k}$
- $\tilde{x}_k = (\bar{x}_{1k}, \dots, \bar{x}_{jk}, \dots, \bar{x}_{pk})^T$  is the  $k$  class centroid.

# Linear Regression with Elastic Net Penalty

- Elastic net penalty

$$\max_{\{\beta_{0k}, \beta_k \in \mathbb{R}^p\}_1^K} \left[ \sum_{i=1}^N \log P(g_i | x_i) - \lambda \left( \sum_{k=1}^K \sum_{j=1}^p (\alpha |\beta_{kj}| + (1 - \alpha) \beta_{kj}^2) \right) \right]$$

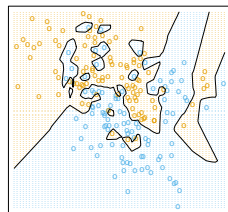
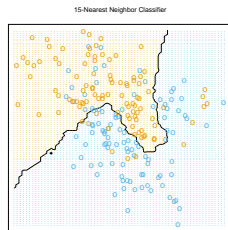
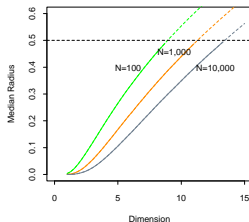


# Nearest-Neighbor Methods

- The **nearest-neighbor methods** use those observations in the training set  $\mathcal{T}$  closest in the input space to  $x$  to form  $\hat{f}$ .

$$\hat{g}(x) = \text{majority}_{x_i \in N_k(x)} g(x_i)$$

- nice, but suffers the curse of dimensionality.



# Discriminang Adaptive Nearest-Neighbor Methods (DANN)

- The metric at a query point is defined by

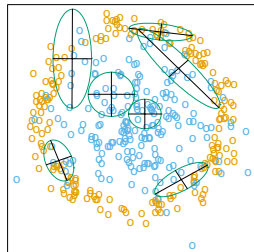
$$D(x, x_0) = (x - x_0)^T \Sigma (x - x_0)$$

- where

$$\Sigma = W^{-\frac{1}{2}} [W^{-\frac{1}{2}} B W^{-\frac{1}{2}} + \epsilon I] W^{-\frac{1}{2}}$$

- where  $\epsilon = 1$  adjusts the neighbourhood and
- $W, B$  are within and between class covariance fitted at the neighbourhood.

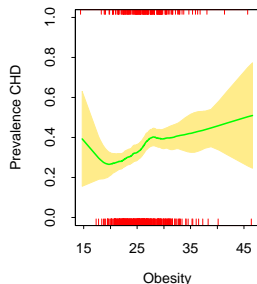
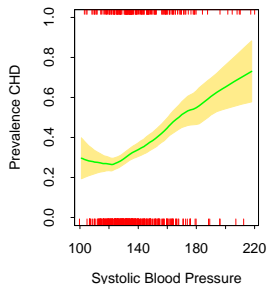
- Close to the class boundary, the  $\Sigma$  shrinks out of the boundary
- in the interior it remains circular.



# Local Likelihood and other methods

- Logistic and log-linear models involve the covariates in a linear fashion.
- We fit the model locally at  $x_0$  and weight the loglik by the kernel  $k_\lambda$
- and center the estimate at  $x_0$ .

$$\begin{aligned}\ell(\beta_{x_0}) &= \sum_{i=1}^N k_\lambda(x_0, x_i) \ell(y_i, (x - x_0)^T \beta_{x_0}) \\ &= \sum_{i=1}^N k_\lambda(x_0, x_i) \left\{ y_i \beta_{x_0}^T (x_i - x_0) - \log(1 + e^{\beta_{x_0}^T (x_i - x_0)}) \right\}\end{aligned}$$



Note: Increased prevalence for small values due to retrospective data: some people with diagnosed CHD started more healthy life.



# Generalized Additive Model (gam)

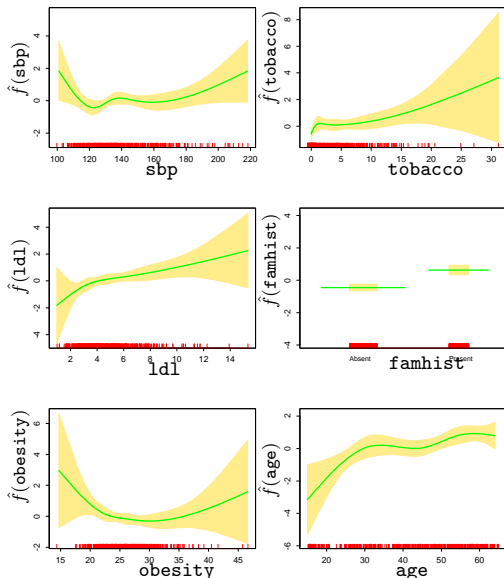
- Each feature  $X_j$  is approximated by a natural spline.
- The overall model is:

$$\text{logit}[P(\text{CHD}|X)] = \theta_0 + h_1(X_1)^T \theta_1 + h_2(X_2)^T \theta_2 + \dots + h_p(X_p)^T \theta_p$$

- $\theta_j$  are vectors of coefficients multiplying their associated vector of natural spline basis functions  $h_j$
- four basis functions (three inner knots) per spline in this example.
- binary *familyhist* with a single coefficient.
- Combine all  $p$  vectors of basic functions into one big vector  $h(X)$ ,  
 $df = 1 + \sum_{j=1}^p df_j$
- each basis function is evaluated at each of the  $N$  samples
- resulting in a  $N \times df$  basis matrix  $\mathbf{H}$ .
- and use 'standard' logistic regression.

# South African Heart Disease continued

- Alcohol not significant by AIC test
- covariance  $Cov(\hat{\theta})$  is estimated by  $\hat{\Sigma} = (\mathbf{H}^T \mathbf{W} \mathbf{H})^{-1}$ 
  - $W$  the diagonal weight matrix
- variance of a single variable  $j$  is:
  - $v_j(X_j) = \text{Var}[f_j(X_j)] = h_j(X_j)^T \hat{\Sigma}_{jj} h_j(X_j)$
- error bounds  $\hat{f}_j(X_j) \pm 2\sqrt{v_j(X_j)}$ .

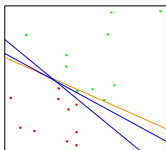


# Summary

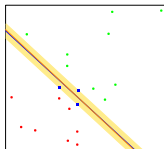
- likelihood example
- logistic regression
  - ext. logistic regression with  $L_1$  penalty, elastic net penalty
- linear and quadratic discriminant analysis
  - ext. regularized discriminant analysis
  - ext. reduced rank discriminant analysis
  - ext. diagonal discriminant analysis
- Nearest-neighbor methods
  - k-NN
  - Local likelihood (local logistic regression)
  - ext. Discriminating Adaptive NN methods (DANN)
- ? Support Vector Machines

# Separating hyperplane, Optimal separating hyperplane

- Classification, we encode the goal class by  $-1$  and  $1$ , respectively.
- separate the space  $X$  by a hyperplane
- **Linear Discriminant Analysis** LDA is not necessary optimal.
- **Logistic regression** finds one if it exists.
- **Perceptron** (a neural network with one neuron) finds separating hyperplane if it exists.
  - The exact position depends on initial parameters.



**FIGURE 4.14.** A toy example with two classes separable by a hyperplane. The orange line is the least squares solution, which misclassifies one of the training points. Also shown are two blue separating hyperplanes found by the perceptron learning algorithm with different random starts.



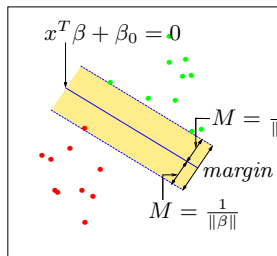
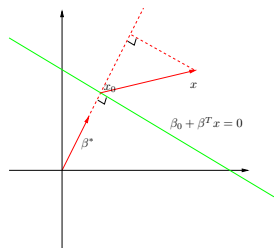
**FIGURE 4.16.** The same data as in Figure 4.14. The shaded region delineates the maximum margin separating the two classes. There are three support points indicated, which lie on the boundary of the margin, and the optimal separating hyperplane (blue line) bisects the slab. Included in the figure is the boundary found using logistic regression (red line), which is very close to the optimal separating hyperplane (see Section 12.3.3).

# Optimal Separating Hyperplane (separable case)

We define **Optimal Separating Hyperplane** as a separating hyperplane with maximal free space  $M$  without any data point around the hyperplane. Formally:

$$\max_{\beta, \beta_0, \|\beta\|=1} M$$

subject to  $y_i(x_i^T \beta + \beta_0) \geq M$  for all  $i = 1, \dots, N$ .



Formally:

$$\max_{\beta, \beta_0, \|\beta\|=1} M$$

subject to  $y_i(x_i^T \beta + \beta_0) \geq M$  for all  $i = 1, \dots, N$ .

We re-define:  $\|\beta\| = 1$  can be moved to the condition (and redefine  $\beta_0$ ):

$$\frac{1}{\|\beta\|} y_i(x_i^T \beta + \beta_0) \geq M$$

Since for any  $\beta$  and  $\beta_0$  satisfying these inequalities, any positively scaled multiple satisfies them too, we can set  $\|\beta\| = \frac{1}{M}$  and we get:

$$\min_{\beta, \beta_0} \frac{1}{2} \|\beta\|^2$$

subject to  $y_i(x_i^T \beta + \beta_0) \geq 1$  pro  $i = 1, \dots, N$ .

This is a convex optimization problem. The Lagrange function, we look for the saddle point w.r.t.  $\beta$  and  $\beta_0$ :

$$L_P = \frac{1}{2} \|\beta\|^2 - \sum_{i=1}^N \alpha_i [y_i(x_i^T \beta + \beta_0) - 1].$$

$$L_P = \frac{1}{2} \|\beta\|^2 - \sum_{i=1}^N \alpha_i [y_i (x_i^T \beta + \beta_0) - 1].$$

Setting the derivatives to zero, we obtain:

$$\beta = \sum_{i=1}^N \alpha_i y_i x_i$$

$$0 = \sum_{i=1}^N \alpha_i y_i$$

Substituting these in  $L_P$  we obtain the so-called Wolfe dual:

$$L_D = \sum_{i=1}^N \alpha_i - \frac{1}{2} \sum_{i=1}^N \sum_{k=1}^N \alpha_i \alpha_k y_i y_k x_i^T x_k$$

subject to  $\alpha_i \geq 0$

The solution is obtained by maximizing  $L_D$  in the positive orthant, for which standard software can be used.

$$L_D = \sum_{i=1}^N \alpha_i - \frac{1}{2} \sum_{i=1}^N \sum_{k=1}^N \alpha_i \alpha_k y_i y_k x_i^T x_k$$

subject to  $\alpha_i \geq 0$ .

In addition the solution must satisfy the Karush–Kuhn–Tucker conditions:

$$\alpha_i [y_i (x_i^T \beta + \beta_0) - 1] = 0$$

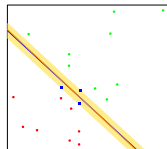
for any  $i$ , therefore for any  $\alpha_i > 0$  must  $[y_i (x_i^T \beta + \beta_0) - 1] = 0$ , that means  $x_i$  is on the boundary and for all  $x_i$  outside the boundary is  $\alpha_i = 0$ .

The boundary is defined by  $x_i$  with  $\alpha_i > 0$  – so called **support vectors**.

We classify new observations

$$\hat{G}(x) = \text{sign}(x^T \beta + \beta_0)$$

- where  $\beta = \sum_{i=1}^N \alpha_i y_i x_i$ ,
- $\beta_0 = y_s - x_s^T \beta$  for any support vector  $\alpha_s > 0$ .





# Optimal Separating Hyperplane (nonseparable case)

- We have to accept incorrectly classified instances in a non-separable case.
- We limit the number of incorrectly classified examples.

We define **slack**  $\xi$  for each data point  $(\xi_1, \dots, \xi_N) = \xi$  as follows:

- $\xi_i$  is the distance of  $x_i$  from the boundary for  $x_i$  at the wrong side of the margin
- and  $\xi_i = 0$ , for  $x_i$  at the correct side.

We require  $\sum_{i=1}^N \xi_i \leq K$ .

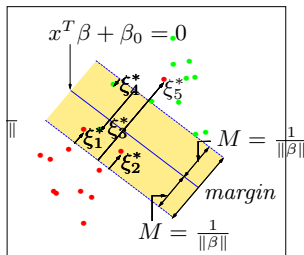
We solve the optimization problem

$$\max_{\beta, \beta_0, \|\beta\|=1} M$$

subject to:

$$y_i(x_i^T \beta + \beta_0) \geq M(1 - \xi_i)$$

where  $\forall i$  is  $\xi_i \geq 0$  a  $\sum_{i=1}^N \xi_i \leq K$ .



# Optimal Separating Hyperplane (nonseparable case)

Again, we omit replace the condition  $\|\beta\|$  by defining  $M = \frac{1}{\|\beta\|}$  and optimize

$$\min \|\beta\| \text{ subject to } \begin{cases} y_i(x^T \beta + \beta_0) \geq (1 - \xi_i) \forall i \\ \xi_i \geq 0, \sum \xi_i \leq \text{constant} \end{cases}$$

We replace the constant by a multiplicative parameter  $\gamma$  and solve

$$\min_{\beta, \beta_0} \frac{1}{2} \|\beta\|^2 + \gamma \sum_{i=1}^N \xi_i$$

subject to  $\xi_i \geq 0$  and  $y_i(x^T \beta + \beta_0) \geq (1 - \xi_i)$ .

- We can set  $\gamma = \infty$  for the separable case.
- Large  $\gamma$ : a complex boundary, fewer support vectors.
- Small  $\gamma$ : a smooth boundary, a robust model, many support vectors.
- $\gamma$  usually set by crossvalidation.

We solve

$$\min_{\beta, \beta_0} \frac{1}{2} \|\beta\|^2 + \gamma \sum_{i=1}^N \xi_i$$

subject to  $\xi_i \geq 0$  and  $y_i(x_i^T \beta + \beta_0) \geq (1 - \xi_i)$ .

Lagrange multipliers again for  $\alpha_i, \mu_i$ :

$$L_P = \frac{1}{2} \|\beta\|^2 + \gamma \sum_{i=1}^N \xi_i - \sum_{i=1}^N \alpha_i [y_i(x_i^T \beta + \beta_0) - (1 - \xi_i)] - \sum_{i=1}^N \mu_i \xi_i$$

Setting the derivative = 0 we get:

$$\beta = \sum_{i=1}^N \alpha_i y_i x_i$$

$$0 = \sum_{i=1}^N \alpha_i y_i$$

$$\alpha_i = \gamma - \mu_i.$$

Substitute to get Wolfe dual:

$$L_D = \sum_{i=1}^N \alpha_i - \frac{1}{2} \sum_{i=1}^N \sum_{k=1}^N \alpha_i \alpha_k y_i y_k x_i^T x_k$$

and maximize  $L_D$  subject to  $0 \leq \alpha_i \leq \gamma$  and  $\sum_{i=1}^N \alpha_i y_i = 0$ .

Solution satisfies:

$$\begin{aligned} \alpha_i [y_i (x_i^T \beta + \beta_0) - (1 - \xi_i)] &= 0 \\ \mu_i \xi_i &= 0 \\ [y_i (x_i^T \beta + \beta_0) - (1 - \xi_i)] &\geq 0 \end{aligned}$$

- The solution is  $\hat{\beta} = \sum_{i=1}^N \hat{\alpha}_i y_i x_i$ .
- **support points** with nonzero coefficients  $\hat{\alpha}_i$  are
  - points at the boundary
    - $\hat{\xi}_i = 0$  (therefore  $0 < \hat{\alpha}_i < \gamma$ ),
  - and points on the wrong side of the margin
    - $\hat{\xi}_i > 0$  (and  $\hat{\alpha}_i = \gamma$ ).
- Any point with  $\hat{\xi}_i = 0$  can be used to calculate  $\hat{\beta}_0$ , typically an average.
  - $\hat{\beta}_0$  for a boundary point  $\alpha_i > 0$ ,  $\xi_i = 0$ :

$$\alpha_i [y_i (x_i^T \hat{\beta} + \hat{\beta}_0) - (1 - 0)] = 0$$

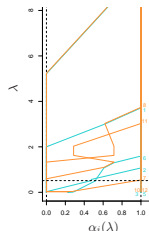
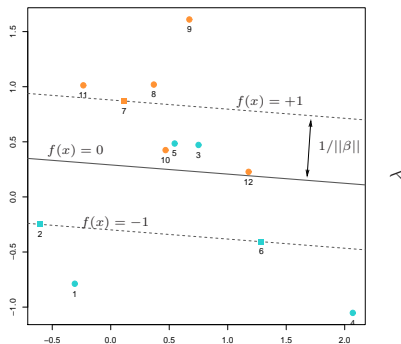
- Parameter  $\alpha$  settled by tuning (crossvalidation)

# SVM Solution

- The solution is  $\hat{\beta} = \sum_{i=1}^N \hat{\alpha}_i y_i x_i$ .
- **support points** with nonzero coefficients  $\hat{\alpha}_i$  are
  - points at the boundary
    - $\hat{\xi}_i = 0$  (therefore  $0 < \hat{\alpha}_i < \gamma$ ),
    - and points on the wrong side of the margin
      - $\hat{\xi}_i > 0$  (and  $\hat{\alpha}_i = \gamma$ ).
- Any point with  $\hat{\xi}_i = 0$  can be used to calculate  $\hat{\beta}_0$ , typically an average.

- $\hat{\beta}_0$  for a boundary point  $\xi_i = 0$ :
 
$$\alpha_i \left[ y_i (x^T \hat{\beta} + \hat{\beta}_0) - (1 - 0) \right] = 0$$

- $\alpha = \xi = 0$  for points 1,4,8,9,11
- $\alpha > 0, \xi = 0$  for points 2,6,8
- misclassified points 3,5.



# Support Vector Machines

Let us have the training data  $(x_i, y_i)_{i=1}^N$ ,  $x_i \in \mathbb{R}^p$ ,  $y_i$  in  $\{-1, 1\}$ . We define a hyperplane

$$\{x : f(x) = x^T \beta + \beta_0 = 0\} \quad (1)$$

where  $\|\beta\| = 1$ .

We classify according to

$$G(x) = \text{sign} [x^T \beta + \beta_0]$$

where  $f(x)$  is a signed distance of  $x$  from the hyperplane.

**Support vector machines replace the scalar product  $\langle x_i, x \rangle$  by a kernel function.**

$$\hat{f}(x) = \beta x + \hat{\beta}_0$$

$$\hat{f}(x) = \sum_{k=1}^N \hat{\alpha}_k y_k x_k^T x + \hat{\beta}_0$$

$$\hat{f}(x) = \sum_{k=1}^N \hat{\alpha}_k y_k \langle x_k, x \rangle + \hat{\beta}_0$$

$$\hat{f}(x) = \sum_{k=1}^N \hat{\alpha}_k y_k K(x_k, x) + \hat{\beta}_0$$

# SVM Example

- **kernel functions** are function to replace scalar product with a scalar product in a transformed space.

<i>d</i> th Degree polynomial:	$K(x, x') = (1 + \langle x, x' \rangle)^d$
Radial basis	$K(x, x') = \exp\left(\frac{-\ x-x'\ ^2}{\ell}\right)$
Neural network	$K(x, x') = \tanh(\kappa_1 \langle x, x' \rangle + \kappa_2)$

- For example a degree 2 with two dimensional input:

$$K(x, x') = (1 + \langle x, x' \rangle)^2 = (1 + 2x_1x'_1 + 2x_2x'_2 + (x_1x'_1)^2 + (x_2x'_2)^2 + 2x_1x'_1x_2x'_2)$$

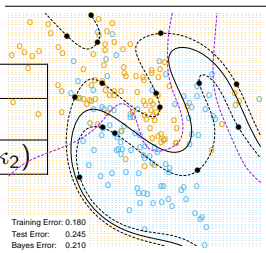
- that is  $M = 6$ ,  $h_1(x) = 1$ ,  $h_2(x) = \sqrt{2}x_1$ ,  
 $h_3(x) = \sqrt{2}x_2$ ,  $h_4(x) = x_1^2$ ,  $h_5(x) = x_2^2$ ,  
 $h_6(x) = \sqrt{2}x_1x_2$ .

The classification function

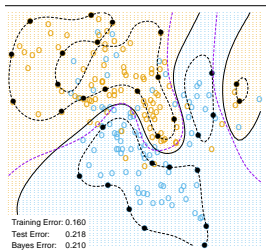
$$\hat{f}(x) = h(x)^T \beta + \beta_0 = \sum_{i=1}^N \alpha_i y_i \langle h(x), h(x_i) \rangle + \beta_0$$

does not need evaluation of  $h(i)$ , only the scalar product  $\langle h(x), h(x_i) \rangle$ .

SVM - Degree-4 Polynomial in Feature Space



SVM - Radial Kernel in Feature Space



# String Kernels and Protein Classification

IPTSALVKETLALLSTHRTLLIANETLRIPVPVHKNHQLCTEEIFQIGITLESQTVQGGTV  
ERLFKNLSLIKKYIDGQKKKCGEERRRVNQFLDY**LQE**FLGVMNTEWI

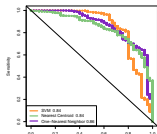
PHRRDLCSRSIWLARKIRSDLTALTESYVKHQGLWSELTEAER**LQEN**LQAYRTFHVLLA  
RLEDQVHFPTPEGDFHQAIHTLLLQVAAFAYQIEELMILLEYKIPRNEADGMLFEKK

- Consider all possible sequences of length  $m$ .
- We define a feature map

$$\Phi_m(x) = \{\phi_a(x)\}_{a \in \mathcal{A}_m}$$

- The kernel function is the inner product:

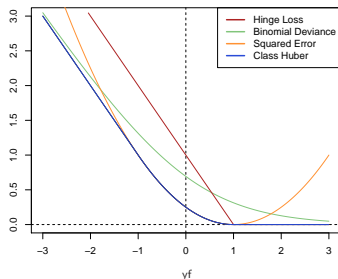
$$K_m(x_1, x_2) = \langle \Phi_m(x_1), \Phi_m(x_2) \rangle.$$





# SVM as a Penalization Method

- We fit a linear function wrt. basis  $\{h_i(x)\}$ :  $f(x) = h^T \beta + \beta_0$ .
- Consider the loss function  $L(y, f) = [1 - yf]_+$ 
  - The optimization problem  $\min_{\beta_0, \beta} \sum_{i=1}^N [1 - yf]_+ + \lambda \|\beta\|^2$
- is equivalent to SVM
  - $\min_{\beta, \beta_0} \frac{1}{2} \|\beta\|^2 + \gamma \sum_{i=1}^N \xi_i$
  - subject to  $\xi_i \geq 0$  and  $y_i(x^T \beta + \beta_0) \geq (1 - \xi_i)$ .
- is similar to smoothing splines penalty:
  - $\min_{\alpha, \alpha_0} \sum_{i=1}^N [1 - yf]_+ + \lambda \alpha^T \mathbf{K} \alpha$
  - where  $\alpha^T \mathbf{K} \alpha = J(f)$  is the smoothing penalty.



Loss Function	$L[y, f(x)]$	Minimizing Function
Binomial Deviance	$\log[1 + e^{-yf(x)}]$	$f(x) = \log \frac{\Pr(Y = +1 x)}{\Pr(Y = -1 x)}$
SVM Hinge Loss	$[1 - yf(x)]_+$	$f(x) = \text{sign}[\Pr(Y = +1 x) - \frac{1}{2}]$
Squared Error	$[y - f(x)]^2 = [1 - yf(x)]^2$	$f(x) = 2\Pr(Y = +1 x) - 1$
"Huberised" Square Hinge Loss	$-4yf(x), \quad yf(x) < -1$ $[1 - yf(x)]_+^2 \quad \text{otherwise}$	$f(x) = 2\Pr(Y = +1 x) - 1$

# SVM and Kernel Dimension

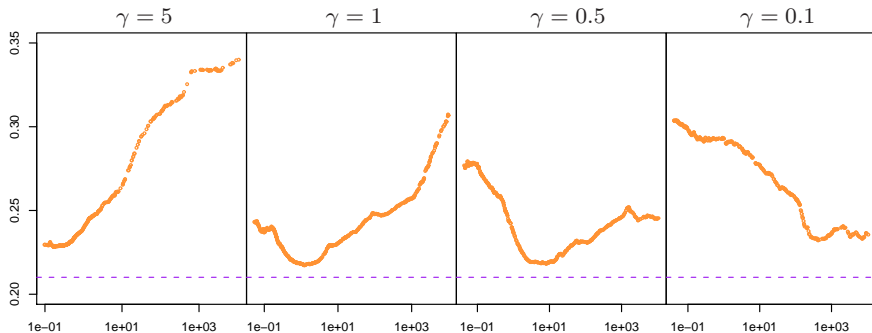
- The first Simulated example
  - 100 observations of each class
  - First class: four standard normal independent features  $X_1, X_2, X_3, X_4$ .
  - Second class conditioned on  $9 \leq \sum X_j^2 \leq 16$ .
- Second example
  - The first one augmented with an additional six standard Gaussian noise features.
- BRUTTO: Additive spline model.
- BRUTTO and MARS has the ability to ignore noisy features.
- We can see the overfitting of SVM. The degree 2 polynomial kernel is the best since the decision boundary is quadratic.

Method	Test Error (SE)	
	No Noise Features	Six Noise Features
SV Classifier	0.450 (0.003)	0.472 (0.003)
SVM/poly 2	0.078 (0.003)	0.152 (0.004)
SVM/poly 5	0.180 (0.004)	0.370 (0.004)
SVM/poly 10	0.230 (0.003)	0.434 (0.002)
BRUTO	0.084 (0.003)	0.090 (0.003)
MARS	0.156 (0.004)	0.173 (0.005)
Bayes	0.029	0.029

## SVM Complexity

The SVM complexity is  $m^3 + mN + mpN$ , where  $m$  is the number of support vectors.

Parameter tuning for different radial basis lengthscale  $\gamma$ .



C

# SVM for Regression

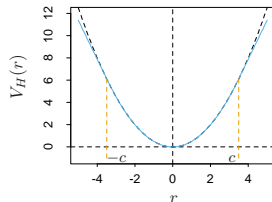
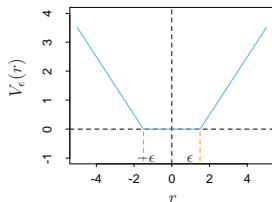
- In regression, we fit a function:  $f(x) = x^T \beta + \beta_0$
- We consider error function  $V_\epsilon$  (left figure)

$$V_\epsilon(r) = \begin{cases} 0 & \text{if } |r| < \epsilon, \\ |r| - \epsilon & \text{otherwise.} \end{cases}$$

- and minimize:

$$H(\beta, \beta_0) = \sum_{i=1}^N V_\epsilon(y_i - f(x_i)) + \frac{\lambda}{2} \|\beta\|^2$$

- 



# SVM for Regression 2

- The solution has the form:  $\hat{\alpha}_i, \hat{\alpha}_i^* \geq 0$

$$\hat{\beta} = \sum_{i=1}^N (\hat{\alpha}_i^* - \hat{\alpha}_i) x_i,$$

$$\hat{f}(x) = \sum_{i=1}^N (\hat{\alpha}_i^* - \hat{\alpha}_i) \langle x, x_i \rangle + \beta_0,$$

- and solve the quadratic programming problem

$$\min_{\alpha_i, \alpha_i^*} \epsilon \sum_{i=1}^N (\alpha_i^* + \alpha_i) - \sum_{i=1}^N y_i (\alpha_i^* - \alpha_i) + \frac{1}{2} \sum_{i,i'=1}^N (\alpha_i^* - \alpha_i) (\alpha_{i'}^* - \alpha_{i'}) \langle x_i, x_{i'} \rangle$$

- subject to the constraints

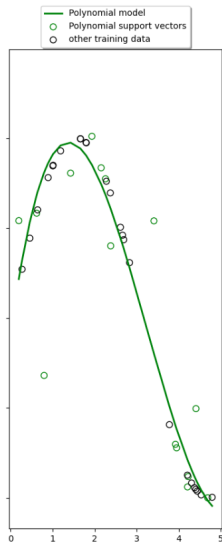
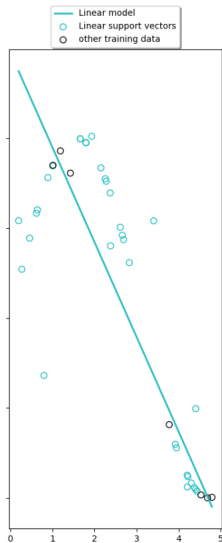
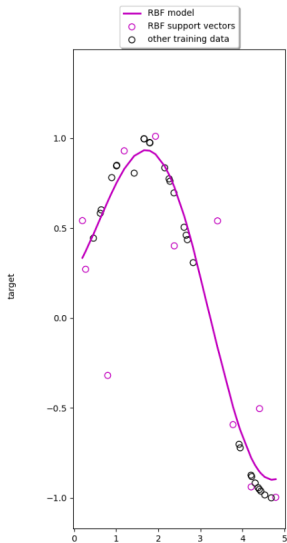
$$0 \leq \alpha_i, \alpha_i^* \leq 1/\lambda,$$

$$\sum_{i=1}^N (\alpha_i^* - \alpha_i) = 0,$$

$$\alpha_i \alpha_i^* = 0.$$

- Support vectors are those with nonzero  $(\hat{\alpha}_i^* - \hat{\alpha}_i)$ .
- With scaled response  $y$ , you may use the default  $\epsilon$ .
- $\lambda$  is tuned by cross-validation.

### Support Vector Regression



# Table of Contents

- 1 Overview of Supervised Learning
- 2 Kernel Methods, Basis Expansion and regularization
- 3 Linear Methods for Classification
- 4 Model Assessment and Selection
- 5 Additive Models, Trees, and Related Methods
- 6 Ensemble Methods
- 7 Clustering
- 8 Bayesian learning, EM algorithm
- 9 Association Rules, Apriori
- 10 Inductive Logic Programming
- 11 Undirected (Pairwise Continuous) Graphical Models
- 12 Gaussian Processes
- 13 PCA Extensions, Independent CA
- 14 Support Vector Machines