## **Decision Estimation**

- Decision estimation and classification are ones of active research areas.
  - Classic measurements of the environment, … , AI (vision, speech recognition.)
    - Systems perform "pattern recognition" or "decision making".
  - Often the information is less than precise, and frequently the decision procedures are statistical in nature.
- Objects of interests are classified into one of classes.
  - These objects are "patterns": printed letters, biological cells,...
  - Systems learn the training data to classify the testing data.
    - Supervised vs. un-supervised.



## **Classification Approach**

- The observation vector x is first transformed into another vector y whose components called features.
  - Feature extraction: the features are intended to be fewer in numbers than the observations.
  - However, they should collectively contain most discernible information for pattern classification.
    - Reduction of the observations to a smaller number of features is anticipated to help design a reliable decision rule.



 Extraction procedures or transformations attempt to compute the components based on intuition or physical considerations of the problem. ⇒ dimensionality reduction.

## Three-class Recognition Example

if  $y_1 + 3y_2 < 9$   $w_1$ : class of 1's, else if  $-y_1 + 3y_2 > 3$   $w_2$ : class of x's, else  $w_3$ : class of 0's.



Discriminant functions:

$$g_1(\mathbf{y}) = -y_1 - 3y_2 + 9,$$
  
 $g_2(\mathbf{y}) = -y_1 + 3y_2 - 3,$   
 $g_3(\mathbf{y}) = g_1(\mathbf{y}) \cdot g_2(\mathbf{y}).$ 

Decision Rule: Choose  $w_i$  where  $g_i(\mathbf{y}) = \max_i [g_i(\mathbf{y})]$ .

- Decision region  $R_i$  is the set  $R_i = \{y : g_i(y) = \max_j [g_j(y)]\}$ .
  - Discriminant functions can be evaluated computationally.
- Decision boundaries are defined by  $g_i(\mathbf{y}) = g_j(\mathbf{y}), i \neq j$ .

## Probability Theory for Random Vectors

- Event A has the associated probability P(A).
  - P(not A) = 1 P(A).
  - The joint probability of two events A & B, denoted as P(AB) or P(A and B) is the probability that A and B both occur simultaneously.
    - P(A or B) = P(A) + P(B) P(A and B).
- Suppose x is a random vector.
  - Its distribution function F(**x**) is defined as

$$F_{\mathbf{x}}(\tilde{\mathbf{x}}) = F_{x_1, x_2, \cdots, x_n}(\tilde{x}_1, \tilde{x}_2, \cdots, \tilde{x}_n) = P(\mathbf{x} \le \tilde{\mathbf{x}}) = P(x_1 \le \tilde{x}_1, x_2 \le \tilde{x}_2, \cdots, x_n \le \tilde{x}_n).$$

•  $F(-\infty)=0, F(+\infty)=1.$ 

- Its density function f(x) is defined as

$$f_{\mathbf{x}}(\tilde{\mathbf{x}}) = \frac{dF_{\mathbf{x}}(\tilde{\mathbf{x}})}{d\mathbf{x}} = \left[\frac{\partial^n F_{\mathbf{x}}(\tilde{\mathbf{x}})}{\partial x_1 \partial x_2 \cdots \partial x_n}\right]_{\mathbf{x} = \tilde{\mathbf{x}}} \Leftrightarrow F_{\mathbf{x}}(\tilde{\mathbf{x}}) = \int_{-\infty}^{\tilde{\mathbf{x}}} f_{\mathbf{x}}(\mathbf{x}) d\mathbf{x} = \int_{-\infty}^{\tilde{x}_1} \int_{-\infty}^{\tilde{x}_2} \cdots \int_{-\infty}^{\tilde{x}_n} f_{\mathbf{x}}(\mathbf{x}) dx_1 dx_2 \cdots dx_n.$$

 $\mathbf{x} = \begin{bmatrix} x_1 \\ x_2 \\ \vdots \\ x_n \end{bmatrix}$ 

Joint Distribution and Density Functions • Suppose **y** is another random vector.  $\mathbf{y} = \begin{pmatrix} y_1 \\ y_2 \\ \vdots \\ \vdots \end{pmatrix}$ • The joint distribution of **x** and **y** is defined by

• The joint distribution of **x** and **y** is defined by  $F_{xy}(\tilde{x}, \tilde{y}) = P(x \le \tilde{x}, y \le \tilde{y}).$ 

The joint density is 
$$f_{\mathbf{x}\mathbf{y}}(\tilde{\mathbf{x}}, \tilde{\mathbf{y}}) = \frac{d^2 F_{\mathbf{x}}(\tilde{\mathbf{x}})}{d\mathbf{x}d\mathbf{y}} = \left[\frac{\partial^m \partial^n F_{\mathbf{x}}(\tilde{\mathbf{x}})}{\partial x_1 \partial x_2 \cdots \partial x_n \partial y_1 \partial y_2 \cdots \partial y_m}\right]_{\mathbf{x}=\tilde{\mathbf{x}}, \mathbf{y}=\tilde{\mathbf{y}}}$$

$$\Rightarrow F_{xy}(-\infty, -\infty) = 0, \quad F_{xy}(\infty, \infty) = 1, \quad F_{xy}(\tilde{\mathbf{x}}, \infty) = F_{x}(\tilde{\mathbf{x}}), \quad F_{xy}(\infty, \tilde{\mathbf{y}}) = F_{y}(\tilde{\mathbf{y}}).$$
  
Marginal p.d.f.

• Example: the joint p.d.f. is  $f_{xy}(\tilde{\mathbf{x}}, \tilde{\mathbf{y}}) = \begin{cases} (\tilde{x}_1 + 3\tilde{x}_2)\tilde{y}_1 & 0 \le \tilde{x}_1, \tilde{x}_2, \tilde{y}_1 \le 1, \\ 0 & \text{otherwise.} \end{cases}$ 

$$\mathbf{x} = \begin{pmatrix} x_1 \\ x_2 \end{pmatrix} \qquad \mathbf{y} = (y_1) \qquad \qquad f_{\mathbf{x}}(\tilde{\mathbf{x}}) = \begin{cases} \int_0^1 (\tilde{x}_1 + 3\tilde{x}_2) y_1 dy_1 = \frac{1}{2} (\tilde{x}_1 + 3\tilde{x}_2) & 0 \le \tilde{x}_1, \tilde{x}_2 \le 1, \\ 0 & \text{otherwise.} \end{cases}$$

Probability Functions jointly with Events
 The joint distribution of a random vector x and an event A is defined by

$$F_{\mathbf{x}A}(\tilde{\mathbf{x}}, A) = P(\mathbf{x} \le \tilde{\mathbf{x}}, A) = \sum_{i=1}^{m} P(\mathbf{x} \le \tilde{\mathbf{x}}, A_i), \quad A = \bigcup_{i=1}^{m} A_i$$

• The conditional probability:  $P(A | B) = \frac{P(A \cap B)}{P(B)} = \frac{P(B | A)P(A)}{P(B)}$ .  $\Rightarrow F_{\mathbf{x}|A}(\tilde{\mathbf{x}} | A) = P(\mathbf{x} \le \tilde{\mathbf{x}} | A) = \frac{P(\mathbf{x} \le \tilde{\mathbf{x}}, A)}{P(A)} = \frac{F_{\mathbf{x}A}(\tilde{\mathbf{x}}, A)}{P(A)}$ . Bayes's Rule  $\Rightarrow f_{\mathbf{x}|\mathbf{y}}(\tilde{\mathbf{x}} | \tilde{\mathbf{y}}) = \frac{f_{\mathbf{xy}}(\tilde{\mathbf{x}}, \tilde{\mathbf{y}})}{f_{\mathbf{y}}(\tilde{\mathbf{y}})}$ .  $\Rightarrow f_{\mathbf{xy}}(\tilde{\mathbf{x}}, \tilde{\mathbf{y}}) = f_{\mathbf{x}}(\tilde{\mathbf{x}}) \cdot f_{\mathbf{y}}(\tilde{\mathbf{y}})$ , if  $\mathbf{x}, \mathbf{y}$  are independent.

$$P(A_{i} | B) = \frac{P(A_{i} \cap B)}{P(B)} = \frac{P(B | A_{i})P(A_{i})}{\sum_{i=1}^{m} P(B | A_{i})}$$
Prior density  

$$\Rightarrow \underbrace{f_{\mathbf{x}|\mathbf{y}}(\tilde{\mathbf{x}} | \tilde{\mathbf{y}})}_{f_{\mathbf{x}}(\tilde{\mathbf{y}})} = \frac{f_{\mathbf{xy}}(\tilde{\mathbf{x}}, \tilde{\mathbf{y}})}{f_{\mathbf{y}}(\tilde{\mathbf{y}})} = \frac{f_{\mathbf{y}|\mathbf{x}}(\tilde{\mathbf{y}} | \tilde{\mathbf{x}})f_{\mathbf{x}}(\tilde{\mathbf{x}})}{\int_{-\infty}^{\infty} f_{\mathbf{y}|\mathbf{x}}(\tilde{\mathbf{y}} | \tilde{\mathbf{x}})f_{\mathbf{x}}(\mathbf{x})d\mathbf{x}}.$$
Prior density  
Posterior density

#### Likelihood Ratio Test

 The hypothesis that a given pattern x belongs one of N<sub>c</sub> classes is tested to minimize the probability of error.

If 
$$f_{w_1|\mathbf{x}}(w_1 \mid \tilde{\mathbf{x}}) = \frac{f_{\mathbf{x}|w_1}(\tilde{\mathbf{x}} \mid w_1)P(w_1)}{f_{\mathbf{x}}(\tilde{\mathbf{x}})} > f_{w_2|\mathbf{x}}(w_2 \mid \tilde{\mathbf{x}}) = \frac{f_{\mathbf{x}|w_2}(\tilde{\mathbf{x}} \mid w_2)P(w_2)}{f_{\mathbf{x}}(\tilde{\mathbf{x}})} \implies w_1 \text{ class,}$$

else  $w_2$  class.

• The decision rule can be 
$$L(\tilde{\mathbf{x}}) \triangleq \frac{f_{\mathbf{x}|w_1}(\tilde{\mathbf{x}} \mid w_1)}{f_{\mathbf{x}|w_2}(\tilde{\mathbf{x}} \mid w_2)} \stackrel{w_1}{\underset{w_2}{>}} \frac{P(w_2)}{P(w_1)}.$$

Likelihood Ratio

Example: the conditional p.d.f. are  $P(w_1) = P(w_2).$   $\Rightarrow L(\tilde{\mathbf{x}}) \triangleq \exp[-\frac{1}{2}(x-4)^2 + \frac{1}{2}(x-10)^2] \stackrel{w_1}{\underset{w_2}{\overset{w_1}{\overset{w_1}{\overset{w_2}{\overset{w_1}{\overset{w_2}{\overset{w_1}{\overset{w_2}{\overset{w_1}{\overset{w_2}{\overset{w_1}{\overset{w_2}{\overset{w_1}{\overset{w_2}{\overset{w_1}{\overset{w_2}{\overset{w_1}{\overset{w_2}{\overset{w_1}{\overset{w_2}{\overset{w_1}{\overset{w_2}{\overset{w_1}{\overset{w_2}{\overset{w_1}{\overset{w_2}{\overset{w_1}{\overset{w_2}{\overset{w_1}{\overset{w_2}{\overset{w_1}{\overset{w_2}{\overset{w_1}{\overset{w_2}{\overset{w_1}{\overset{w_1}{\overset{w_2}{\overset{w_1}{\overset{w_1}{\overset{w_2}{\overset{w_1}{\overset{w_1}{\overset{w_1}{\overset{w_2}{\overset{w_1}}}}{\overset{w_1}{\overset$ 

## Probability of Misclassification

 The probability of error, i.e. Bayes risk, determines the quality of a decision rule.

- A lower value implies a better rule.

$$P(error) = \int_{-\infty}^{\infty} P(error \mid \mathbf{x}) f_{\mathbf{x}}(\mathbf{x}) d\mathbf{x} = P(error \mid w_1) P(w_1) + P(error \mid w_2) P(w_2).$$

$$\varepsilon_1 = P(error \mid w_1) = P(\text{choose } w_2 \mid w_1) = \int_{R_2} f_{\mathbf{x} \mid w_1}(\mathbf{x} \mid w_1) d\mathbf{x}.$$
$$\varepsilon_2 = P(error \mid w_2) = P(\text{choose } w_1 \mid w_2) = \int_{R_1} f_{\mathbf{x} \mid w_2}(\mathbf{x} \mid w_2) d\mathbf{x}.$$

- Bayes risk under Multiple hypotheses can be defined:  $P(error | \mathbf{x}) = \sum_{j=1, j \neq i}^{N_c} P(w_j | \mathbf{x}) = 1 - P(w_i | \mathbf{x}), \text{ if } \mathbf{x} \in R_i.$ 
  - $R_i$  should be defined to be the region where  $P(w_i | \mathbf{x})$  is largest. Choose  $w_i$  where  $P(w_i | \mathbf{x}) = \max_j [P(w_j | \mathbf{x})]$ .

#### **Distance Functions**

- There are several ways to measure the distance d(x,y) between two vectors x & y.
  - Generally, a distance function is any scalar-valued function satisfying the following conditions:
    - d(x,y)>0 for x≠y; d(x,y)=0 if x=y.
    - $\cdot d(\mathbf{x},\mathbf{y}) = d(\mathbf{y},\mathbf{x}).$
    - [Triangular inequality]  $d(\mathbf{x},\mathbf{y})+d(\mathbf{y},\mathbf{z}) \ge d(\mathbf{x},\mathbf{z})$ .
  - Euclidean distance:  $d_E(\mathbf{x}, \mathbf{y}) = |\mathbf{x} \mathbf{y}| = \left(\sum_{i=1}^n (x_i y_i)^2\right)^{\frac{1}{2}}$ .
  - Maximum value distance:  $d_M(\mathbf{x}, \mathbf{y}) = \max_i |x_i y_i|$ .
  - Absolute value distance (city block):  $d_A(\mathbf{x}, \mathbf{y}) = \sum_{i=1}^n |x_i y_i|$ .

#### Linear Transformation

- If x is a vector in X and y is the corresponding (mapped) vector in Y, then y = A x.
- Matrix A is said to be **positive definite**
  - if the quadratic product,  $\mathbf{x}^T A \mathbf{x}$ , is strictly greater than zero for all non-zero vector  $\mathbf{x}$ .  $\mathbf{x}^T A \mathbf{x} = \sum_{i=1}^n \sum_{j=1}^n a_{ij} x_i x_j$ .
- Matrix A is said to be **positive semidefinite** if the quadratic product, **x**<sup>T</sup>A**x**, ≥ 0 for all non-zero vector **x**.

• Example: the positive definite matrix 
$$\mathbf{A} = \begin{pmatrix} 1 & -1 \\ -1 & 2 \end{pmatrix}$$
.  
 $\Leftrightarrow \mathbf{x}^T \mathbf{A} \mathbf{x} = \sum_{i=1}^n \sum_{j=1}^n a_{ij} x_i x_j = x_1^2 - 2x_1 x_2 + 2x_2^2 = (x_1 - x_2)^2 + x_2^2 > 0, \forall \mathbf{x} \neq \mathbf{0}.$   
 $\mathbf{B} = \begin{pmatrix} 2 & 1 & 1 \\ 1 & 2 & -1 \\ 3 & 3 & 0 \end{pmatrix}$  is not a positive definite matrix.

### Differentiation w.r.t. Vectors

- If s is a scalar function of a vector  $\mathbf{x} \in \mathbb{R}^n$ ,
  - the derivative of s w.r.t. x is defined as the vector (gradient)

$$\frac{ds}{d\mathbf{x}} = \begin{bmatrix} \frac{\partial s}{\partial x_1} & \frac{\partial s}{\partial x_2} & \cdots & \frac{\partial s}{\partial x_n} \end{bmatrix}^T$$

If s is a vector ∈ R<sup>m</sup>,
the derivative of s w.r.t. x is the matrix

$$\frac{d\mathbf{s}}{d\mathbf{x}} = \begin{pmatrix} \frac{\partial s_1}{\partial x_1} & \dots & \frac{\partial s_m}{\partial x_1} \\ \vdots & \ddots & \vdots \\ \frac{\partial s_1}{\partial x_n} & \dots & \frac{\partial s_m}{\partial x_n} \end{pmatrix}$$

 $\frac{d\mathbf{r}}{d\mathbf{v}} = \frac{d\mathbf{r}}{d\mathbf{s}}\frac{d\mathbf{s}}{d\mathbf{v}}.$ 

Chained rule

If **B** is symmetric

- If **r** is a linear transformation of **s**, - Namely, **r**=**Bs**, then  $\frac{d\mathbf{r}}{d\mathbf{x}} = \mathbf{B}\frac{d\mathbf{s}}{d\mathbf{x}}$ .
- If r is a nonlinear transformation of s,
- For the quadratic product,  $\frac{d(\mathbf{x}^{\mathrm{T}}\mathbf{B}\mathbf{x})}{d\mathbf{x}} = \cdots = (\mathbf{B} + \mathbf{B}^{\mathrm{T}})\mathbf{x} = 2\mathbf{B}\mathbf{x}.$

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• The correlation matrix **R** of a random vector **x** is  $\mathbf{R} = E(\mathbf{x}\mathbf{x}^{T}) = (r_{ij}), \quad r_{ij} = E(x_{i}x_{j}).$ 

The covariance matrix K of a random vector x is

$$\mathbf{K} = E[(\mathbf{x} - \mathbf{m})(\mathbf{x} - \mathbf{m})^{\mathrm{T}}] = (k_{ij}), \quad k_{ij} = E[(x_i - m_i)(x_j - m_j)].$$

- The diagonal elements  $k_{ii}$  are the variances of the vector components.  $k_{ii} = \sigma_i^2 = Var(x_i) = E[(x_i - m_i)^2].$ 

$$\mathbf{K} = E[(\mathbf{x} - \mathbf{m})(\mathbf{x} - \mathbf{m})^{\mathrm{T}}] = E[\mathbf{x}\mathbf{x}^{\mathrm{T}} - \mathbf{x}\mathbf{m}^{\mathrm{T}} - \mathbf{m}\mathbf{x}^{\mathrm{T}} + \mathbf{m}\mathbf{m}^{\mathrm{T}}]$$
  
=  $E[\mathbf{x}\mathbf{x}^{\mathrm{T}}] - E[\mathbf{x}]\mathbf{m}^{\mathrm{T}} - \mathbf{m}E[\mathbf{x}^{\mathrm{T}}] + \mathbf{m}\mathbf{m}^{\mathrm{T}} = E[\mathbf{x}\mathbf{x}^{\mathrm{T}}] - \mathbf{m}\mathbf{m}^{\mathrm{T}} = \mathbf{R} - \mathbf{m}\mathbf{m}^{\mathrm{T}}.$   $\Rightarrow$   $\mathbf{R} = \mathbf{K} + \mathbf{m}\mathbf{m}^{\mathrm{T}}$ 

- Given a vector  $\mathbf{y} = \mathbf{A} \mathbf{x}$ ,  $\mathbf{m}_{y} = E[\mathbf{A}\mathbf{x}] = \mathbf{A}E[\mathbf{x}] = \mathbf{A}\mathbf{m}_{x}$ .
  - $\mathbf{R}_{\mathbf{y}} = E[(\mathbf{A}\mathbf{x})(\mathbf{A}\mathbf{x})^{T}] = \mathbf{A}E[\mathbf{x}\mathbf{x}^{T}]\mathbf{A}^{T} = \mathbf{A}\mathbf{R}_{\mathbf{x}}\mathbf{A}^{T}. \qquad \mathbf{K}_{\mathbf{y}} = \mathbf{A}\mathbf{K}_{\mathbf{x}}\mathbf{A}^{T}.$

- If **A** is orthogonal, i.e. has orthonormal column vectors,  $|\mathbf{R}_y| = |\mathbf{R}_x|, tr(\mathbf{R}_y) = tr(\mathbf{R}_x), |\mathbf{K}_y| = |\mathbf{K}_x|, tr(\mathbf{K}_y) = tr(\mathbf{K}_x).$ 

## Independent Random Vectors

- If random vectors x and y are independent, they are uncorrelated.
  - The converse is generally not true.
  - Both are uncorrelated if  $\begin{cases} \mathbf{R}_{xy} = E(\mathbf{xy}^{T}) = E(\mathbf{x})E(\mathbf{y}^{T}) = \mathbf{m}_{x}\mathbf{m}_{y}^{T}, \\ \mathbf{K}_{xy} = E[(\mathbf{x} \mathbf{m}_{x})(\mathbf{y} \mathbf{m}_{y})^{T}] = \mathbf{0}. \end{cases}$
- Gaussian Random Vectors:  $f(\mathbf{x}) = \frac{1}{\sigma\sqrt{2\pi}} e^{-\frac{(\mathbf{x}-\mu)^2}{2\sigma^2}} \implies f(\mathbf{x}) = \frac{1}{(2\pi)^{n/2}\sqrt{|\mathbf{K}_{\mathbf{x}}|}} \exp\left[-\frac{(\mathbf{x}-\mathbf{m})\mathbf{K}_{\mathbf{x}}^{-1}(\mathbf{x}-\mathbf{m})^T}{2}\right].$
- Recall: a/(a+c), better for a smaller value of c.
   The ratio of the number of shots detected
   correctly over the actual number of shots.
- Precision: a/(a+b), better for a smaller value of b.
  - ✓ The ratio of the number of shots detected correctly over the total number of shots detected.

detected shots

correct shots

С

Ь

a

### Support Vector Machines (SVM)

#### SVM is a novel kind of Neural Networks.

- Multi-Layer-Perceptron (MLP): Classifier, regressor, etc.
  - Single-layer & Multi-layer with feed-forward connections.
  - Back propagation algorithm, maximum likelihood principle.
  - Training, self-structured: supervised, unsupervised.
  - The performance is justified by a loss function (say, MSE) over unseen samples of the test set.
    - The **expected** risk of the classifier on the test set  $[2] \leq$ The **empirical** risk on the training set [0] + the **estimation** error [1].

Estimation error 
$$\simeq \sqrt{\frac{h}{c}\log(1+2\frac{c}{h})},$$

[0] & [1] should be both min.✓ Minimizing [0] alone do no good!!

c = |Training Set|,

h = VC dimension of the classifer.

(Vapnik-Chervonenkis) = the maximal number of samples correctly classified in the training set.

# **Graphical Illustration**



 In Modeling, the approximation error stems from the model mismatch.

The true f(x) may lie outside the hypothesis space.

- In Learning, the estimation error occurs due to the imperfect learning procedure.
  - The non-optimal model (empirically obtained) may be chosen.
- During the testing (evaluation), the generalization error is met.
- SVM minimizes the Expected risk by controlling VC dimension.
   Learning becomes solving the problem of Quadratic Programming.

## SVM = Optimal Hyperplane Algorithm

- Learning how to classify is estimating a function f:  $R^n \rightarrow \pm 1$  over the training data set = {(x<sub>i</sub>, y<sub>i</sub>)  $\in R^n \times \pm 1$  : i=1...c)}
  - f will correctly classify other unseen example (x, y) under the same unknown probability distribution P(x, y).  $\Rightarrow$  namely, f(x)=y.
  - It is often assumed the data are i.i.d. (identically independent distributed).

Hyperplanes:  $w \cdot x + b = 0$ ,  $w \in \mathbb{R}^n$ ,  $\Leftrightarrow$  Decision Functions:  $f(x) = \operatorname{sgn}(w \cdot x + b)$ .

 $\Rightarrow \exists a unique hyperplane(w,b) \ni \max_{w,b} \min_i \left( \|x - x_i\| : x \in \mathbb{R}^n, w \cdot x + b = 0, i = 1 \cdots c \right).$ Maximize the separation margin.

 $\Rightarrow \text{Optimization problem:} \begin{cases} \min L(w) = \frac{1}{2} \|w\|^2 & \text{Good separation} \\ y_i(w \cdot x_i + b) \ge 1, i = 1 \cdots c. & \text{Correct} \end{cases}$ 

⇒ Solution = the saddle point  $\alpha_i \ge 0$  of the Lagrangian:

$$L(w,b,\alpha) = \frac{1}{2} \|w\|^2 - \sum_{i=1}^{c} \alpha_i [y_i(w \cdot x_i + b) - 1].$$
  
Minimized w.r.t. w & b, maximized w.r.t.  $\alpha_{i \cdot 16}$ 

## Solution

 $\begin{cases} \frac{\partial L(w,b,\alpha)}{\partial b} = 0 \Rightarrow \sum_{i=1}^{c} \alpha_i y_i = 0, & \text{Lying on the margin} \\ \frac{\partial L(w,b,\alpha)}{\partial w} = 0 \Rightarrow w = \sum_{i=1}^{c} \alpha_i y_i x_i. & \text{The solution vector is a linear combination of} \\ a \text{ subset of the training patterns.} \\ \Rightarrow & \text{Support vectors summarize the information.} \end{cases}$ 

 $\Rightarrow b = -\frac{1}{2} w \cdot (x_p + x_q), \alpha_p > 0, \alpha_q > 0, y_p = 1, y_q = -1, \text{ for any SV } x_p, x_q.$ 

However, most classification problems are not linear separable.
 Transform x<sub>i</sub> to a high-dimension space to regain linear separation.

 $x \Rightarrow \Phi(x)$ .  $\Phi(x)$  is hard to compute.

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Decision Functions:  $f(x) = \operatorname{sgn}(w \cdot x + b) = \operatorname{sgn}\left(\sum_{i=1}^{c} \alpha_i y_i(x_i \cdot x) + b\right)$  $\Rightarrow f(x) = \operatorname{sgn}\left(\sum_{i=1}^{c} \alpha_i y_i(\Phi(x_i) \cdot \Phi(x)) + b\right).$ 

The scalar (inner) product,  $\Phi(x_i) \cdot \Phi(x)$ , is easy to compute by a simple kernel. As an example, the polynomial kernel  $k(x,y) = \Phi(x) \cdot \Phi(y) = (x \cdot y)^d$ . ✓ Matrices (K<sub>ii</sub>) are positive definite, where  $K_{ii} = k(x_i, x_i)$ , i, j=1…c.

## Dilemma

- Typically, the data will only be linearly separable in some, possibly very high dimensional space.
  - Separating the data exactly, particularly for a finite amount of data with noise, is favorable. However it will generalize badly.
  - In practice, it may be necessary to employ the non-separable approach (allow some classification error).
- To allow some overlapping between classes, the slack variables  $\tau_i \ge 0$  is introduced.

$$\Rightarrow \text{ Optimization problem:} \begin{cases} \min L(w,\tau) = \frac{1}{2} \|w\|^2 + C \sum_{i=1}^c \tau_i, \quad C \text{ is some constant} \ge 0. \\ y_i(w \cdot x_i + b) \ge 1 - \tau_i, \quad i = 1 \cdots c. \qquad \tau_i \ge 0 \end{cases}$$
$$\Rightarrow L(w,b,\alpha,\tau,\beta) = \frac{1}{2} \|w\|^2 - \sum_{i=1}^c \alpha_i [y_i(w \cdot x_i + b) - 1 + \tau_i] - \sum_{i=1}^c \beta_i \tau_i. \quad \text{Saddle points} \\ C \ge \alpha_i \ge 0 \quad \sum_{i=1}^c \alpha_i y_i = 0, \quad w = \sum_{i=1}^c \alpha_i y_i x_i, \quad \alpha_i + \beta_i = C. \text{ (Any SV } x_i \text{ has } \tau_i = 0.) \end{cases}$$
$$\Rightarrow \max_{\alpha} W(\alpha) = \sum_{i=1}^c \alpha_i - \frac{1}{2} \sum_{i=1}^c \sum_{j=1}^c \alpha_j \alpha_j y_j y_j (x_i \cdot x_j).$$

## **Non-linear Separation**

Using the kernel,

 $\Rightarrow \max_{\alpha} W(\vec{\alpha}) = \sum_{i=1}^{c} \alpha_i - \frac{1}{2} \sum_{i=1}^{c} \sum_{j=1}^{c} \alpha_i \alpha_j y_i y_j k(x_i, x_j).$ 

• Define the matrix Q,  $Q_{ij} = y_i y_j k(\vec{x}_i, \vec{x}_j)$ .  $\Rightarrow W(\vec{\alpha}) = \vec{\alpha}^T \vec{1} - \frac{1}{2} \vec{\alpha}^T Q \vec{\alpha}$ =  $\vec{\alpha}^T (\vec{1} - \frac{1}{2} Q \vec{\alpha})$ .

- Decomposition: break the entire training set into smaller ones.
  - Select the working (active) subset.
    - Other  $\alpha_i$  are fixed in the current iteration.
    - Shrink the problem.
      - There are much less SVs than c.
      - Many SVs have  $\alpha_i = C$ .
      - Caching and incremental updates of the gradient & the termination criteria.

## Generalized Discriminant Analysis (GDA)

GDA is the eigenvalue problem resolution for nonlinear discriminant analysis.

- It is similar in functionality to SVM.
- The input set X has m vectors,  $x_1 \cdots x_m$ , belong to n classes,  $X_1 \cdots X_n$ .

- The cardinality of the subset  $X_i$  is  $m_i$ .  $\Rightarrow X = \bigcup_{i=1}^n X_i$ ,  $\sum_{i=1}^n m_i = m$ .

- The covariance matrix C of all  $x_i$ :  $C = \frac{1}{m} \sum_{i=1}^m x_i x_i^T$ .

Suppose  $x \Rightarrow \Phi(x)$ .  $C \Rightarrow V = \frac{1}{m} \sum_{i=1}^{m} \Phi(x_i) \Phi^T(x_i)$ . Center  $\Phi(x_i)$  in the transform space:  $\tilde{\Phi}(x_i) = \Phi(x_i) - \frac{1}{m} \sum_{k=1}^{m} \Phi(x_k)$ .

- The **inter-class** inertia B is the covariance matrix of the class centers.

$$B = \frac{1}{m} \sum_{i=1}^{n} m_i \overline{\Phi}_i \overline{\Phi}_i^T, \quad \overline{\Phi}_i = \frac{1}{m_i} \sum_{k=1}^{m_i} \Phi(x_{i,k}).$$

The  $k^{\underline{th}}$  vector of Class i is  $x_{i,k}$ .

Likewise,  $V = \frac{1}{m} \sum_{i=1}^{m} \Phi(x_i) \Phi^T(x_i) = \frac{1}{m} \sum_{i=1}^{n} \sum_{k=1}^{m} \Phi(x_{i,k}) \Phi^T(x_{i,k})$ . total inertia

## Formulation

• Using Kernel function:  $k(x_i, x_j) = k_{i,j} = \Phi^T(x_i)\Phi(x_j)$ . For classes p & q,  $(k_{i,j})_{p,q} = \Phi^T(x_{p,i})\Phi(x_{q,j})$ .

• Define a mxm matrix K:  $K = (K_{p,q})_{p,q=1\cdots n}$ ,  $K_{p,q} = (k_{i,j})_{i=1\cdots m_p, j=1\cdots m_q} = K_{q,p}^T$ .

A mxm block diagonal matrix W:  $W = (W_t)_{t=1\cdots n}, \quad W_t = (\frac{1}{m_t})_{m_t \times m_t}.$ 

- The classical criteria for class separability is defined by the quotient between the inter-class inertia and the intra-classes inertia.
  - Its maximization is equivalent to the eigenvalue resolution.
- Assume the classes follow a multivariate Gaussian distribution, and each observation can be assigned to the class having the maximum posterior probability using the <u>Mahalanobis distance</u>.

### **Eigenvalue Resolution**

- Given two symmetric matrices A & B with the same size, and B<sup>-1</sup> exists,  $y^T A y$ 
  - The quotient  $\frac{v^T A v}{v^T B v}$  is maximal for eigenvector v of B<sup>-1</sup>A associated to the large eigenvalue  $\lambda$ .

Since  

$$\frac{(v^T B v)(2Av) - (v^T A v)(2Bv)}{(v^T B v)^2} = 0 \Rightarrow B^{-1}Av = \left(\frac{v^T A v}{v^T B v}\right)v$$

$$\Rightarrow \left(\frac{v^T A v}{v^T B v}\right): \text{ eigenvalue, } v: \text{ eigenvector of } B^{-1}A.$$

Therefore, the quotient  $\frac{B}{V}$  of both inertia's in the problem is maximized:

 $\lambda V v = B v$   $\lambda v = V^{-1} B v \implies \left(\frac{v^T B v}{v^T V v}\right) = \lambda : \text{the largest eigenvalue, } v : \text{eigenvector of } V^{-1} B.$   $B = \frac{1}{m} \sum_{i=1}^{n} m_i \overline{\Phi}_i \overline{\Phi}_i^T, \quad \overline{\Phi}_i = \frac{1}{m_i} \sum_{k=1}^{m_i} \Phi(x_{i,k}).$   $v = \sum_{i=1}^{n} \sum_{k=1}^{m_i} \alpha_{i,k} \Phi(x_{i,k}).$  Linear combination

## Formulation

 $\boldsymbol{\alpha} = \left(\alpha_{i}\right)_{i=1\cdots n}, \quad \alpha_{i} = \left(\alpha_{i,k}\right)_{k=1\cdots m_{i}} \quad \Rightarrow \quad \left(\frac{\boldsymbol{\nu}^{T} \boldsymbol{B} \boldsymbol{\nu}}{\boldsymbol{\nu}^{T} \boldsymbol{V} \boldsymbol{\nu}}\right) = \lambda = \left(\frac{\boldsymbol{\alpha}^{T} \boldsymbol{K} \boldsymbol{W} \boldsymbol{K} \boldsymbol{\alpha}}{\boldsymbol{\alpha}^{T} \boldsymbol{K} \boldsymbol{K} \boldsymbol{\alpha}}\right).$ Proof:  $K = \left(K_{p,q}\right)_{p,q=1\dots,p},$  $\lambda V_V = Bv \implies \lambda \Phi^T(x_{rs}) Vv = \Phi^T(x_{rs}) Bv.$  $K_{p,q} = (k_{i,j})_{i=1\cdots m_{p}, j=1\cdots m_{q}} = K_{q,p}^{T}.$  $Vv = \frac{1}{m} \sum_{p=1}^{n} \sum_{i=1}^{m_p} \Phi(x_{p,i}) \Phi^T(x_{p,i}) \times \sum_{q=1}^{n} \sum_{k=1}^{m_q} \alpha_{q,k} \overline{\Phi(x_{q,k})} = \frac{1}{m} \sum_{q=1}^{n} \sum_{k=1}^{m_q} \alpha_{q,k} \sum_{p=1}^{n} \sum_{i=1}^{m_p} \Phi(x_{p,i}) \Phi^T(x_{p,i}) \Phi(x_{q,k}).$  $\lambda \Phi^{T}(x_{r,s}) V v = \frac{\lambda}{m} \sum_{q=1}^{n} \sum_{k=1}^{m_{q}} \alpha_{q,k} \sum_{p=1}^{n} \sum_{i=1}^{m_{p}} [\Phi^{T}(x_{r,s}) \Phi(x_{p,i})] [\Phi^{T}(x_{p,i}) \Phi(x_{q,k})].$  $\Rightarrow \lambda[\Phi^T(x_{1,m_1}),\cdots,\Phi^T(x_{m_1,m_1}),\cdots,\Phi^T(x_{1,m_n}),\cdots,\Phi^T(x_{m_n,m_n})]Vv = \frac{\lambda}{m}KK\alpha.$  $Bv = \frac{1}{m} \sum_{p=1}^{n} m_p \left[ \frac{1}{m_p} \sum_{i=1}^{m_p} \Phi(x_{p,i}) \right] \left[ \frac{1}{m_p} \sum_{i=1}^{m_p} \Phi(x_{p,i}) \right]^T \times \sum_{q=1}^{n} \sum_{k=1}^{m_q} \alpha_{q,k} \Phi(x_{q,k})$  $= \frac{1}{m} \sum_{q=1}^{n} \sum_{k=1}^{m_q} \alpha_{q,k} \sum_{p=1}^{n} \left[ \sum_{i=1}^{m_p} \Phi(x_{p,i}) \right] \left[ \frac{1}{m_p} \right] \left[ \sum_{i=1}^{m_p} \Phi^T(x_{p,i}) \Phi(x_{q,k}) \right].$  $\Rightarrow [\Phi^{T}(x_{1,m_{1}}), \cdots, \Phi^{T}(x_{m_{1},m_{1}}), \cdots, \Phi^{T}(x_{1,m_{n}}), \cdots, \Phi^{T}(x_{m_{n},m_{n}})]Bv = \frac{1}{m}KWK\alpha.$ 

### **Eigenvalue Resolution**

- By the eigenvectors decomposition of K,  $K = P\Gamma P^T$ 
  - P contains the normalized eigenvectors, say v.
    - P is orthonormal since K is symmetric.

•

 $-\Gamma$  is the diagonal matrix with non-zero eigenvalues.

$$\begin{pmatrix} v^{T}Bv \\ v^{T}Vv \end{pmatrix} = \lambda = \left(\frac{\alpha^{T}KWK\alpha}{\alpha^{T}KK\alpha}\right) = \frac{\alpha^{T}(P\Gamma P^{T})W(P\Gamma P^{T})\alpha}{\alpha^{T}(P\Gamma P^{T})(P\Gamma P^{T})\alpha} = \frac{(\Gamma P^{T}\alpha)^{T}P^{T}WP(\Gamma P^{T}\alpha)}{(\Gamma P^{T}\alpha)^{T}P^{T}P(\Gamma P^{T}\alpha)}.$$

$$\beta = \Gamma P^{T}\alpha \quad \Rightarrow \quad \lambda P^{T}P\beta = \lambda\beta = P^{T}WP\beta \quad \Rightarrow \quad \alpha = P\Gamma^{-1}\beta.$$
Also,  $1 = v^{T}v = \sum_{p=1}^{n}\sum_{k=1}^{m_{p}}\alpha_{p,k}\Phi^{T}(x_{p,k})\sum_{q=1}^{n}\sum_{k=1}^{m_{q}}\alpha_{q,k}\Phi(x_{q,k})$ 

$$= \sum_{p=1}^{n}\sum_{q=1}^{n}\alpha_{p}^{T}K_{p,q}\alpha_{q} = \alpha^{T}K\alpha \quad \Rightarrow \quad \alpha \text{ should be normalized by } \sqrt{\alpha^{T}K\alpha}$$

Given a test vector z, the projections can be computed as

$$v^{T}z = \sum_{p=1}^{n} \sum_{k=1}^{m_{p}} \alpha_{p,k} \Phi^{T}(x_{p,k}) z = \sum_{p=1}^{n} \sum_{k=1}^{m_{p}} \alpha_{p,k} k(x_{p,k},z).$$

## Summary

- **GDA** procedure is summarized in the following steps: **1.** Compute K and W.  $(k_{i,j})_{p,q} = \Phi^T(x_{p,i})\Phi(x_{q,j})$ .  $K = (K_{p,q})_{p,q=1\cdots n}, K_{p,q} = (k_{i,j})_{i=1\cdots m_n, j=1\cdots m_q}, W = (W_t)_{t=1\cdots n}, W_t = (\frac{1}{m_t})_{m \times m}$ .
  - 2. Decompose K using eigenvectors decompositions.  $K = P\Gamma P^{T}$
  - **3.** Compute eigenvectors  $\beta$  and eigenvalues of the system.  $\beta = \Gamma P^T \alpha \implies \lambda P^T P \beta = \underline{\lambda \beta} = P^T W P \beta \implies \alpha = P \Gamma^{-1} \beta.$
  - 4. Compute eigenvectors v using  $\alpha$  and normalize them.

 $v = \sum_{p=1}^{n} \sum_{k=1}^{m_p} \alpha_{p,k} \Phi(x_{p,k}).$   $\alpha$  should be normalized by  $\sqrt{\alpha^T K \alpha}.$ 

5. Compute projections of test points onto the eigenvectors v.

$$\mathbf{v}^{T} z = \sum_{p=1}^{n} \sum_{k=1}^{m_{p}} \alpha_{p,k} \Phi^{T}(x_{p,k}) \ z = \sum_{p=1}^{n} \sum_{k=1}^{m_{p}} \alpha_{p,k} k(x_{p,k},z).$$

## **Kernel Functions**

#### Various kernel functions can be used:

- Gaussian kernel, RBF-kernel:  $k(x, y) = \exp\left(\frac{-\|x-y\|^d}{2\sigma^2}\right)$ .

- Polynomial kernel:  $k(x, y) = (x \cdot y)^d$ .

 $d = 2 \implies (x_1^2, \dots, x_t^2, x_1 x_2, \dots, x_i x_j, \dots): \frac{t(t-1)}{2}$  terms for  $x \in R^t$ .

- Threshold values are learned and chosen.
  - The number of classes minus one is the number of thresholds chosen for classification.

# Biased Discriminant Transform (BDT)

- MM information retrieval relies on the descriptors (or feature vectors), a set of real numbers.
  - Effectiveness of the representation in descriptors.
  - Selection of similarity metric.

- Difference between Traditional and MM DB:
  - Binary "Hit-or-Miss" decision using keywords in traditional DB.
    - The occurrences of the keywords or their synonyms, or
    - Rule-based ranking. etc.
  - In MMDB, the feature space is R<sup>n</sup> (continuous).
    - Inherently, it is a nearest neighbor or a top-k ranking problem.

## Why On-Line Learning?

- "Consensus" interpretation on MM contents:
  - Among all the users
  - Among all the times
    - The correct answer should match the context of conversation.
    - "The bat slipped from his hand." shows different meaning in the context of a baseball game or a cave exploring.
    - Medical image DB may define specific functionalities to perform off-line pre-clustering.

#### On-learning is indispensable.

- The system need to communicate with the user to perceive the specific goal of the queries.
  - In CBIR, a user is required to offer the feature-weighting scheme.
  - In "Relevance Feedback", a user is kept in the loop to tell the relevance of an image or video. (NO R/W of textual description)

## Supervised Classification Problem

One descriptor is assumed to represent the MM object.

- By it, the media type becomes transparent to the system.
- The object can be an whole image, image block, segmented region, shorts, frames, or a key frame.
- $\checkmark$  A point is associated with the descriptor in the feature space.

Relevance feedback: supervised classification problem.

- Learning Speed: the number of iterations.
- Training Size: the number of samples, i.e. their population.
  - Class density, positive/negative samples, etc.
- Top-k returns: not a binary decision.
  - Binary classification (two-class) may not be optimal.
- Initial results are returned; returns/evaluation are iterated.
  - > The goal is to learn the discriminating **subspace**.

## Variants of Relevance Feedback

#### Objectives:

- A user may look for a particular object or a similar one.

#### Feedbacks:

- A user may give back the positive feedback, negative, or both.
- The degree of relevance for each result may be returned.
- Partial likeness: it is like A in color, like B in shape, etc.
- Multiple Descriptors per Sample:
  - A mixed model can be used for refinement (intersection, union) to emphasize the local features.
- Class distribution:
  - Two or more target classes may be assumed.
    - Gaussian: two; Kernel-based: more for non-linearity.
- Data Organization:
  - A hierarchical tree structure may slow learning in real-time.
- Focus:
  - To learn a linear transformation, consider the correlations of feature components, estimate the class density, etc.

## Fisher & Multiple Discriminant Analyses

- The consensus is to find the features to best <u>cluster</u> & <u>separate</u> the positive examples from the negative.
- Traditional approaches:
  - Two-class assumption (FDA): to find a lower dimensional space in which the ratio of between-class scatter over withinclass scatter is maximized.

$$\mathbf{W} = \arg_{\mathbf{W}} \max \frac{\left| \mathbf{W}^{\mathrm{T}} \mathbf{S}_{\mathbf{b}} \mathbf{W} \right|}{\left| \mathbf{W}^{\mathrm{T}} \mathbf{S}_{\mathbf{W}} \mathbf{W} \right|}.$$

$$\mathbf{S}_{\mathbf{b}} = (\mathbf{m}_{\mathbf{x}} - \mathbf{m})(\mathbf{m}_{\mathbf{x}} - \mathbf{m})^{T} + (\mathbf{m}_{\mathbf{y}} - \mathbf{m})(\mathbf{m}_{\mathbf{y}} - \mathbf{m})^{T}.$$

$$\mathbf{S}_{\mathbf{W}} = \sum_{i=1}^{N_{x}} (\mathbf{x}_{i} - \mathbf{m}_{\mathbf{x}})(\mathbf{x}_{i} - \mathbf{m}_{\mathbf{x}})^{T} + \sum_{i=1}^{N_{y}} (\mathbf{y}_{i} - \mathbf{m}_{\mathbf{y}})(\mathbf{y}_{i} - \mathbf{m}_{\mathbf{y}})^{T}.$$
Small intra

- Two-class assumption (MDA):  $\mathbf{S}_{\mathbf{b}} = (\mathbf{m}_{\mathbf{x}} - \mathbf{m})(\mathbf{m}_{\mathbf{x}} - \mathbf{m})^{T} + \sum_{i=1}^{N_{y}} (\mathbf{y}_{i} - \mathbf{m})(\mathbf{y}_{i} - \mathbf{m})^{T}.$   $\mathbf{S}_{\mathbf{W}} = \sum_{i=1}^{N_{x}} (\mathbf{x}_{i} - \mathbf{m}_{\mathbf{x}})(\mathbf{x}_{i} - \mathbf{m}_{\mathbf{x}})^{T}.$ 

## Biased Discriminant Analysis (BDA)

#### • (1+x)-class assumption:

- The user is only interested in one class, while there are an unknown number of other classes.
  - "All happy families are alike, each unhappy family is unhappy in its own fashion" - Leo Tolstoy's Anna Karenina.
  - All positive examples are alike in a way; each negative example is negative in its own way.

$$\mathbf{W} = \arg_{\mathbf{W}} \max \frac{\left| \mathbf{W}^{\mathsf{T}} \mathbf{S}_{\mathbf{y}} \mathbf{W} \right|}{\left| \mathbf{W}^{\mathsf{T}} \mathbf{S}_{\mathbf{x}} \mathbf{W} \right|}.$$
$$\mathbf{S}_{\mathbf{y}} = \sum_{i=1}^{N_{y}} (\mathbf{y}_{i} - \mathbf{m}_{\mathbf{x}}) (\mathbf{y}_{i} - \mathbf{m}_{\mathbf{x}})^{T}.$$
$$\mathbf{S}_{\mathbf{x}} = \sum_{i=1}^{N_{x}} (\mathbf{x}_{i} - \mathbf{m}_{\mathbf{x}}) (\mathbf{x}_{i} - \mathbf{m}_{\mathbf{x}})^{T}.$$

- Regularization and Discounting Factors:
  - Sample-based estimates may be severely biased for small number of training examples.

$$\mathbf{S}_{\mathbf{x}}^{r} = (1 - \mu)\mathbf{S}_{\mathbf{x}} + \frac{\mu}{n}tr[\mathbf{S}_{\mathbf{x}}]\mathbf{I}.$$

$$\mathbf{S}_{\mathbf{y}}^{d} = (1 - \gamma)\mathbf{S}_{\mathbf{y}} + \frac{\gamma}{n}tr[\mathbf{S}_{\mathbf{y}}]\mathbf{I}.$$

$$\mathbf{n} = \text{dim}(\text{original space}).$$

#### Kernel-based BDA (KBDA)

- For non-linearity in the data, a non-linear mapping  $\Phi: \mathbf{x} \rightarrow \Phi(\mathbf{x})$  is used to restore linearity in the transform space.
  - The evaluation of kernel K =  $(k_{ij})$ , where  $k_{ij} = \Phi^{T}(\mathbf{x}_{i})\Phi(\mathbf{x}_{j})$ .

$$\mathbf{W} = \arg_{\mathbf{W}} \max \frac{\left| \mathbf{W}^{\mathsf{T}} \mathbf{S}_{\mathbf{y}}^{\Phi} \mathbf{W} \right|}{\left| \mathbf{W}^{\mathsf{T}} \mathbf{S}_{\mathbf{x}}^{\Phi} \mathbf{W} \right|}.$$
$$\mathbf{S}_{\mathbf{y}}^{\Phi} = \sum_{i=1}^{N_{y}} (\Phi(\mathbf{y}_{i}) - \mathbf{m}_{\mathbf{x}}^{\Phi}) (\Phi(\mathbf{y}_{i}) - \mathbf{m}_{\mathbf{x}}^{\Phi})^{T}.$$
$$\mathbf{S}_{\mathbf{x}}^{\Phi} = \sum_{i=1}^{N_{x}} (\Phi(\mathbf{x}_{i}) - \mathbf{m}_{\mathbf{x}}^{\Phi}) (\Phi(\mathbf{x}_{i}) - \mathbf{m}_{\mathbf{x}}^{\Phi})^{T}.$$

 Let w is the eigenvector associated with the largest eigenvalue for W.

$$\mathbf{w} = \sum_{i=1}^{N_x} \alpha_i \Phi(\mathbf{x}_i) + \sum_{j=1}^{N_y} \alpha_{i+N_x} \Phi(\mathbf{y}_i) = \mathbf{\Phi} \alpha.$$

$$\mathbf{K}_{\mathbf{y}_i} = \mathbf{\Phi}^T \Phi(\mathbf{y}_i) = \left(\mathbf{K}_{\mathbf{y}}\right)_{:,j},$$

$$\mathbf{w}^T \mathbf{S}_{\mathbf{y}}^{\Phi} \mathbf{w} = \alpha^T \mathbf{\Phi}^T \left[\sum_{i=1}^{N_y} (\Phi(\mathbf{y}_i) - \mathbf{m}_{\mathbf{x}}^{\Phi}) (\Phi(\mathbf{y}_i) - \mathbf{m}_{\mathbf{x}}^{\Phi})^T\right] \Phi \alpha$$

$$\mathbf{K}_{\mathbf{mx}} = \mathbf{\Phi}^T \mathbf{m}_{\mathbf{x}}^{\Phi},$$

$$\mathbf{I}_{\mathbf{N}_{\mathbf{x}}} = \frac{1}{\mathbf{N}_{\mathbf{x}}} (\mathbf{1})_{\mathbf{N}_{\mathbf{x}} \times \mathbf{N}_{\mathbf{y}}}.$$

$$\mathbf{W}^T \mathbf{S}_{\mathbf{y}}^{\Phi} \mathbf{w} = \alpha^T \mathbf{\Phi}^T \left[\sum_{i=1}^{N_y} (\Phi(\mathbf{y}_i) - \mathbf{m}_{\mathbf{x}}^{\Phi}) (\Phi(\mathbf{y}_i) - \mathbf{m}_{\mathbf{x}}^{\Phi})^T\right] \alpha$$

$$= \alpha^T \left[\sum_{i=1}^{N_y} (\mathbf{K}_{\mathbf{y}_i} - \mathbf{K}_{\mathbf{mx}}) (\mathbf{K}_{\mathbf{y}_i} - \mathbf{K}_{\mathbf{mx}})^T\right] \alpha.$$

$$= \alpha^T \left[(\mathbf{K}_{\mathbf{y}} - \mathbf{K}_{\mathbf{x}} \mathbf{I}_{\mathbf{N}_{\mathbf{x}}}^{\mathbf{y}}) (\mathbf{K}_{\mathbf{y}} - \mathbf{K}_{\mathbf{x}} \mathbf{I}_{\mathbf{N}_{\mathbf{x}}}^{\mathbf{y}})^T\right] \alpha.$$
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## KBDA (cont'd)



$$\mathbf{w}^{T} \mathbf{S}_{\mathbf{x}}^{\Phi} \mathbf{w} = \mathbf{\alpha}^{T} \left[ (\mathbf{K}_{\mathbf{x}} - \mathbf{K}_{\mathbf{x}} \mathbf{I}_{\mathbf{N}_{\mathbf{x}}}^{\mathbf{x}}) (\mathbf{K}_{\mathbf{x}} - \mathbf{K}_{\mathbf{x}} \mathbf{I}_{\mathbf{N}_{\mathbf{x}}}^{\mathbf{x}})^{T} \right] \mathbf{\alpha}$$
  
=  $\mathbf{\alpha}^{T} \mathbf{K}_{\mathbf{x}} \left[ (\mathbf{I} - \mathbf{I}_{\mathbf{N}_{\mathbf{x}}}^{\mathbf{x}}) (\mathbf{I} - \mathbf{I}_{\mathbf{N}_{\mathbf{x}}}^{\mathbf{x}})^{T} \right] \mathbf{K}_{\mathbf{x}}^{T} \mathbf{\alpha}$   
=  $\mathbf{\alpha}^{T} \mathbf{K}_{\mathbf{x}} (\mathbf{I} - \mathbf{I}_{\mathbf{N}_{\mathbf{x}}}^{\mathbf{x}})^{2} \mathbf{K}_{\mathbf{x}}^{T} \mathbf{\alpha}.$ 

Solve to get α : the eigenvector with the largest eigenvalue.
 Given a new pattern z, find its projection onto w by

$$\mathbf{w}^{T}\Phi(\mathbf{z}) = \sum_{i=1}^{N_{x}} \alpha_{i} k(\mathbf{x}_{i}, \mathbf{z}) + \sum_{j=1}^{N_{y}} \alpha_{i+N_{x}} k(\mathbf{y}_{i}, \mathbf{z}).$$

- In this new space, the nearest neighbors of the positive centroid are returned in each iteration.
  - Combined with the subsequent feedbacks, the new nearest neighbors are output.







RBF-Kernel;

$$k(\mathbf{x}, \mathbf{y}) = \exp\left(-||\mathbf{x} - \mathbf{y}||^2 / (2 \sigma^2)\right),$$

Primal optimization problem:

 $\begin{array}{ll} \text{minimize} & \tau(\mathbf{w}) = \frac{1}{2} ||\mathbf{w}||^2 \\ \text{subject to} & y_i \cdot ((\mathbf{w} \cdot \mathbf{x}_i) + b) \ge 1, \quad i = 1, \dots, \ell. \end{array}$ 

Decision function:



Figure 5 Test results on synthetic training data: six different configurations of non-linearity. The circles are positive examples and the crosses negative. A simulated query process is used for training sample selection, i.e., the 20 nearest neighbors of a randomly selected positive point are used as training samples. The bar diagram shows the averaged hit rate in top 20 returns.