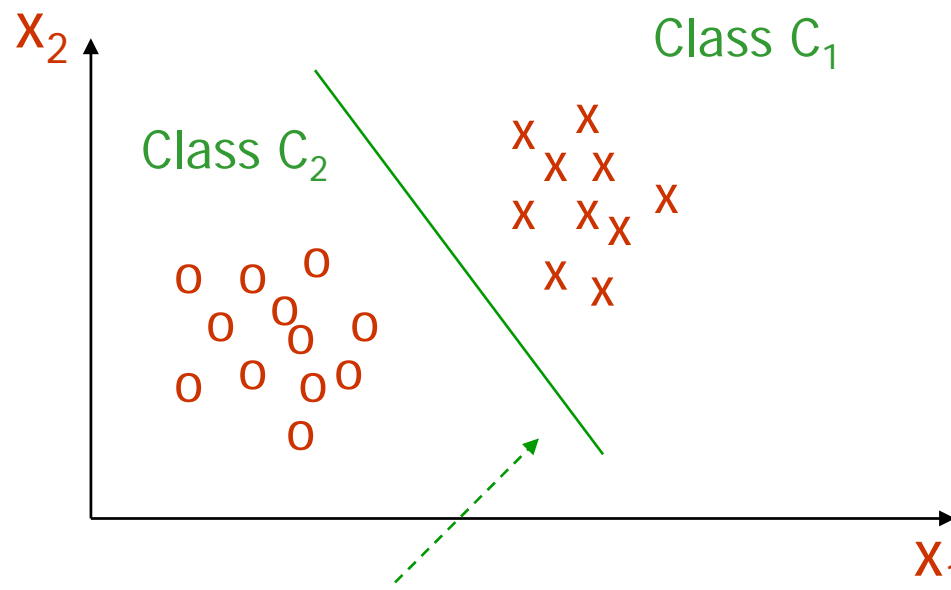




Pattern Classification

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Linear discriminate function



Discriminate function

$$g(\mathbf{x})=0 \quad w_0 + w_1x_1 + w_2x_2 = 0$$

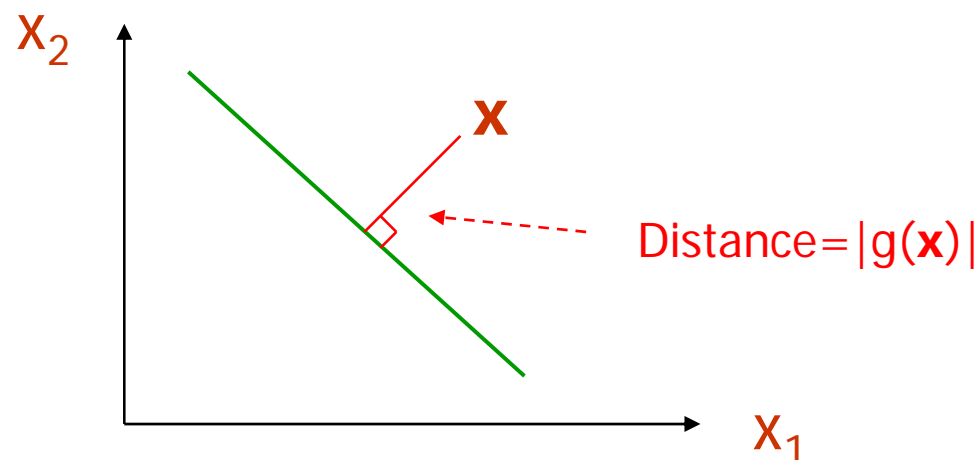
Linear discriminate function (cont.)

A two-class pattern classifier

$$g(\mathbf{x}) = w_0 + w_1x_1 + w_2x_2 + \cdots + w_nx_n = \mathbf{w}^T \mathbf{x}$$

where $\mathbf{w} = [w_0, w_1, w_2, \cdots, w_n]$ and $\mathbf{x} = [1, x_1, x_2, \cdots, x_n]$

$$g(\mathbf{x}) = \begin{cases} \mathbf{w}^T \mathbf{x} > 0 & \text{if } \mathbf{x} \in C_1 \\ \mathbf{w}^T \mathbf{x} \leq 0 & \text{if } \mathbf{x} \in C_2 \end{cases}$$

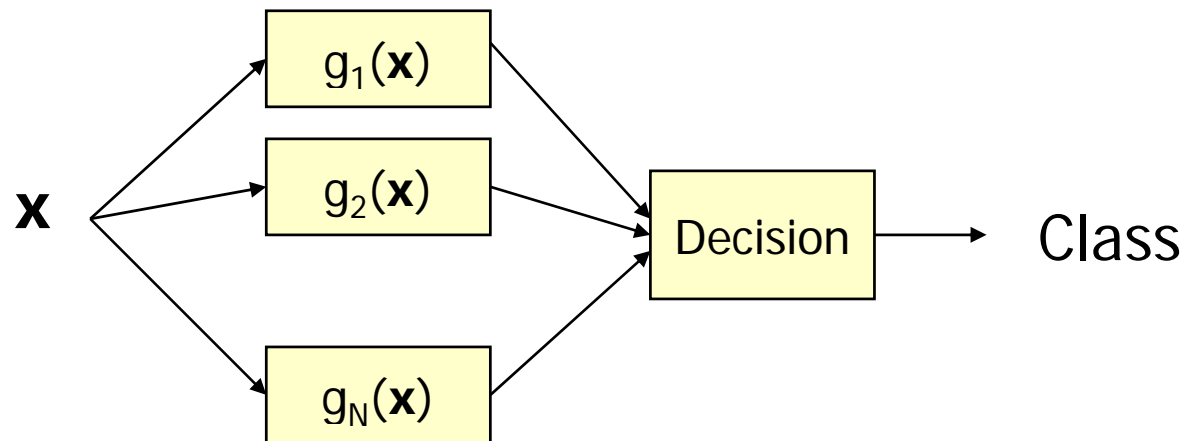


Multi-category classifier

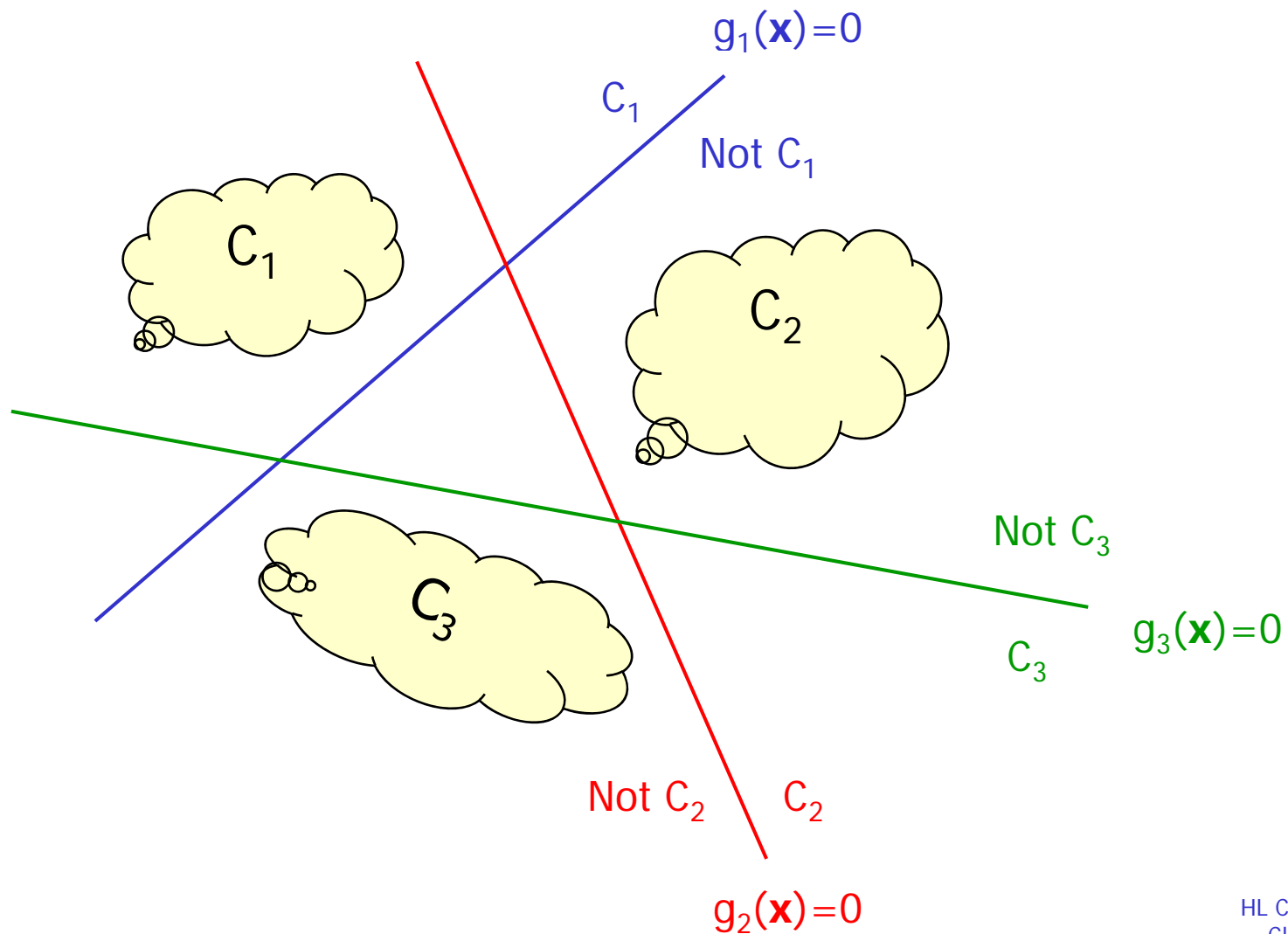
- Build M weight vectors and M discriminate functions

$$g_i(\mathbf{x}) = \begin{cases} \mathbf{w}_i^T \mathbf{x} > 0 & \text{if } \mathbf{x} \in C_i \\ \mathbf{w}_i^T \mathbf{x} \leq 0 & \text{otherwise} \end{cases}, i = 1, 2, \dots, M$$

- Make decision



Build M discriminate functions



Make decisions based on M discriminate functions

- One-to-rest separable

- Each class is separable from the rest by a single discriminate function

if $g_i(\mathbf{x}) > 0$ then $\mathbf{x} \in C_i$

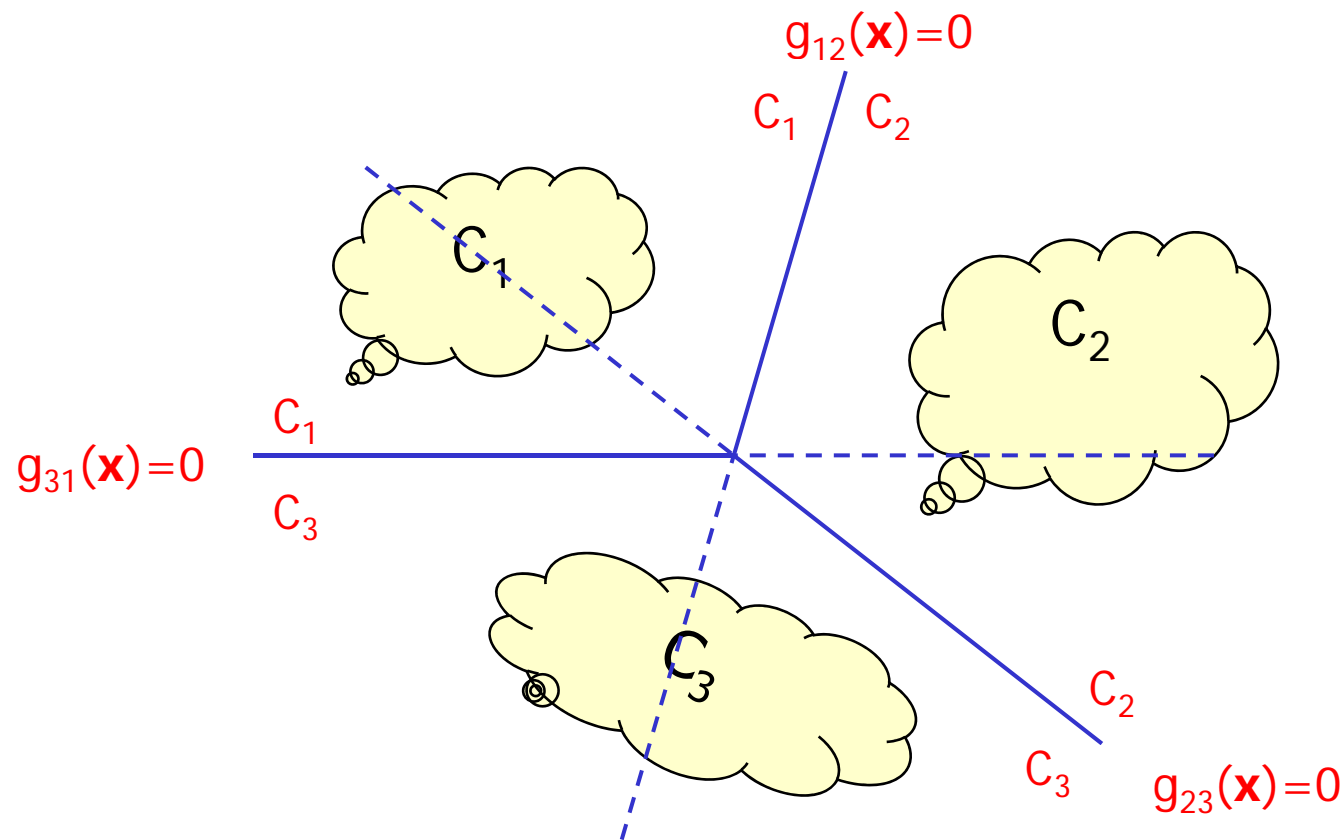
- Pair-wise separable

- There exists M discriminate functions with the property that

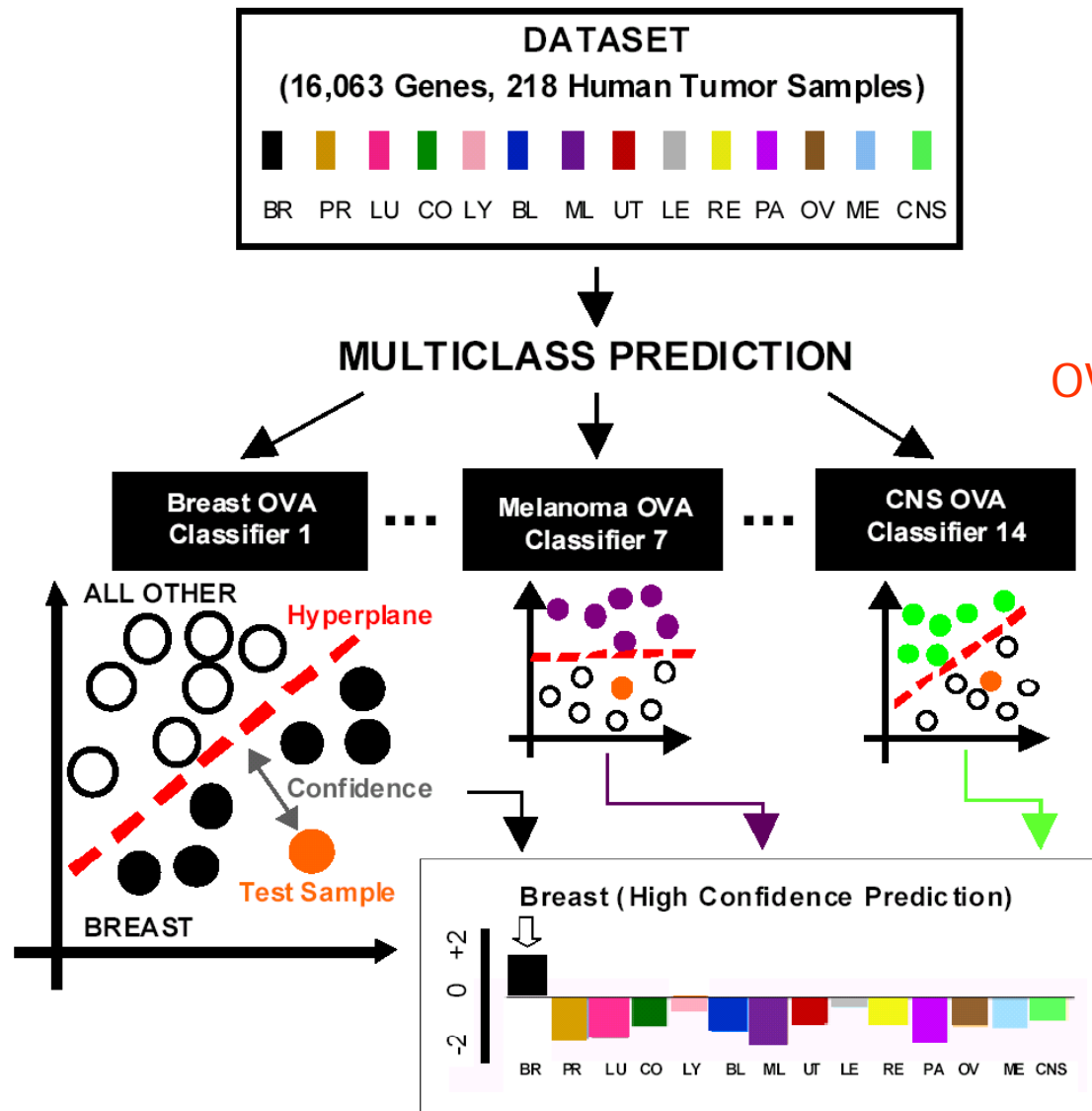
if $g_i(\mathbf{x}) > g_j(\mathbf{x}) \quad \forall j \neq i$ then $\mathbf{x} \in C_i$

$$g_{ij}(\mathbf{x}) = g_i(\mathbf{x}) - g_j(\mathbf{x}) = \mathbf{w}_i \mathbf{x} - \mathbf{w}_j \mathbf{x} = (\mathbf{w}_i - \mathbf{w}_j) \mathbf{x}$$

Pair-wise separable: $M(M-1)/2$ two-class classifier



Minimum distance classifier



OVA (One-versus-all) classifier

Probability model for statistical decision

- Assumes a Gaussian probability density function (PDF) to represent the features for each class
 - Represents the statistical attribute for each class using test samples

Probability model

The a prior probability

$$P(C_i) \quad i = 1, 2, \dots, M$$

The likelihood function of class C_i

$$p(\mathbf{x} | C_i)$$

PDF of \mathbf{x} regardless of class membership

$$p(\mathbf{x}) = \sum_{i=1}^M P(C_i) p(\mathbf{x} | C_i)$$

The a posterior probability

$$P(C_i | \mathbf{x}) = \frac{P(C_i) p(\mathbf{x} | C_i)}{p(\mathbf{x})}$$

The probability that observed
 \mathbf{x} comes from C_i

Bayes classifier for Gaussian PDF

- Gaussian PDF of a single random variable x

$$p(x) = \frac{1}{\sqrt{2\pi}\sigma} \exp\left[-\frac{(x-m)^2}{2\sigma^2}\right]$$

where the mean and the variance

$$m = E[x] = \int_{-\infty}^{\infty} xp(x)dx$$

$$\sigma^2 = E[(x-m)^2] = \int_{-\infty}^{\infty} (x-m)^2 p(x)dx$$

Bayes classifier for Gaussian PDF (cont.)

- In the case of M classes and P pattern vector, $\mathbf{x} = [x_1, x_2, \dots, x_n]$

$$p(x | C_i) = \frac{1}{(2\pi)^{n/2} |\boldsymbol{\Sigma}_i|^{1/2}} \exp \left[-\frac{1}{2} (\mathbf{x} - \mathbf{m}_i)^T \boldsymbol{\Sigma}_i^{-1} (\mathbf{x} - \mathbf{m}_i) \right]$$

where the mean vector and the covariance matrix

$$\mathbf{m}_i = E_i[\mathbf{x}]$$

$$\boldsymbol{\Sigma}_i = E_i[(\mathbf{x} - \mathbf{m}_i)(\mathbf{x} - \mathbf{m}_i)^T]$$

Bayes classifier for Gaussian PDF (cont.)

- Discriminate functions

$$g_i(\mathbf{x}) = p(\mathbf{x} | C_i)P(C_i), \quad i = 1, 2, \dots, M$$

$$g_i(\mathbf{x}) = \ln[p(\mathbf{x} | C_i)P(C_i)] = \ln p(\mathbf{x}/C_i) + \ln P(C_i)$$

$$g_i(\mathbf{x}) = \ln P(C_i) - \frac{1}{2} \ln |\boldsymbol{\Sigma}_i| - \frac{1}{2} [(\mathbf{x} - \mathbf{m}_i)^T \boldsymbol{\Sigma}_i^{-1} (\mathbf{x} - \mathbf{m}_i)]$$

Bayes classifier for Gaussian PDF (cont.)

- If the features are statistically independent $\Sigma_i = \sigma^2 \mathbf{I}$
the bayes classifier is equivalent to linear discriminate function

$$\begin{aligned} g_i(\mathbf{x}) &= \ln P(C_i) - \frac{\|\mathbf{x} - \mathbf{m}_i\|^2}{2\sigma^2} \\ &= \ln P(C_i) - \frac{1}{2\sigma^2} [\mathbf{x}^T \mathbf{x} - 2\mathbf{m}_i^T \mathbf{x} - \mathbf{m}_i^T \mathbf{m}_i] \end{aligned}$$

$$g_i(\mathbf{x}) = \mathbf{w}_i^T \mathbf{x} + \mathbf{w}_{i0}$$

Bayes classifier for Gaussian PDF (cont.)

- If all covariance matrix are equal, the Bayes classifier is also equivalent to linear discriminate functions

$$\begin{aligned} g_i(\mathbf{x}) &= \ln P(C_i) + \mathbf{x}^T \boldsymbol{\Sigma}^{-1} \mathbf{m}_i - \frac{1}{2} \mathbf{m}_i^T \boldsymbol{\Sigma}^{-1} \mathbf{m}_i \\ &= \mathbf{w}_i^T \mathbf{x} + \mathbf{w}_{i0} \end{aligned}$$

Matlab implemenation

- `class = classify(sample,training,group,type)`
 - 'linear' — Fits a multivariate normal density to each group, with a pooled estimate of covariance
 - 'diaglinear' — Similar to 'linear', but with a diagonal covariance matrix estimate
 - 'quadratic' — Fits multivariate normal densities with covariance estimates stratified by group
 - 'diagquadratic' — Similar to 'quadratic',but with a diagonal covariance matrix estimate
 - 'mahalanobis' — Uses Mahalanobis distances with stratified covariance estimates

Fisher's sepal measurements for iris versicolor and virginica

% Training data

```
load fisheriris
```

```
SL = meas(51:end,1);
```

```
SW = meas(51:end,2);
```

```
group = species(51:end);
```

```
h1 = gscatter(SL,SW,group,'rb','v^',[],'off');
```

```
set(h1,'LineWidth',2)
```

```
legend('Fisher versicolor','Fisher virginica', 'Location','NW')
```

% Classify a grid of measurements on the same scale:

```
[X,Y] = meshgrid(linspace(4.5,8),linspace(2,4));
```

```
X = X(:); Y = Y(:);
```

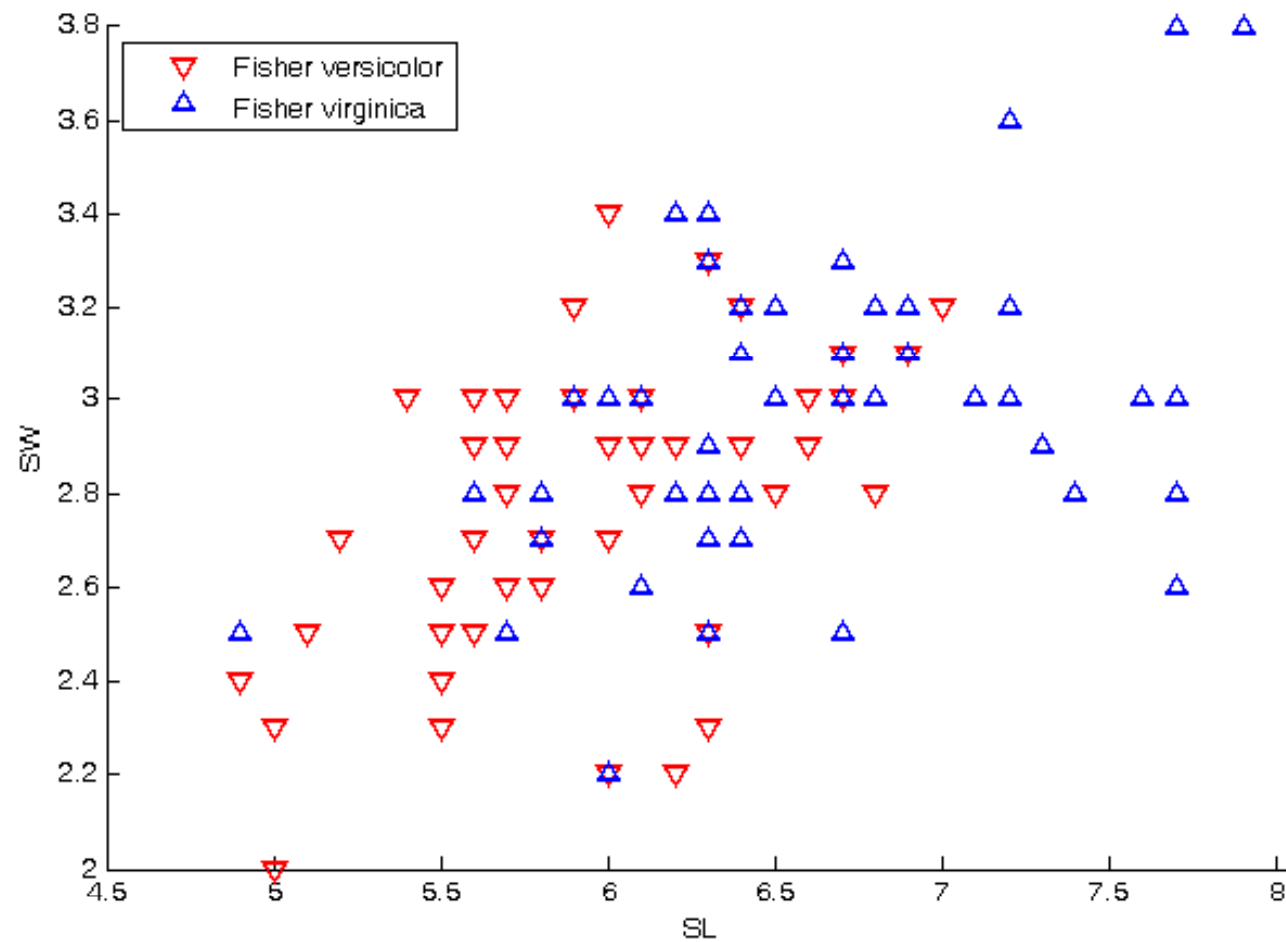
```
[C,err,P,logp,coeff] = classify([X Y],[SL SW],group,'quadratic');
```

Fisher's sepal measurements for iris versicolor and virginica

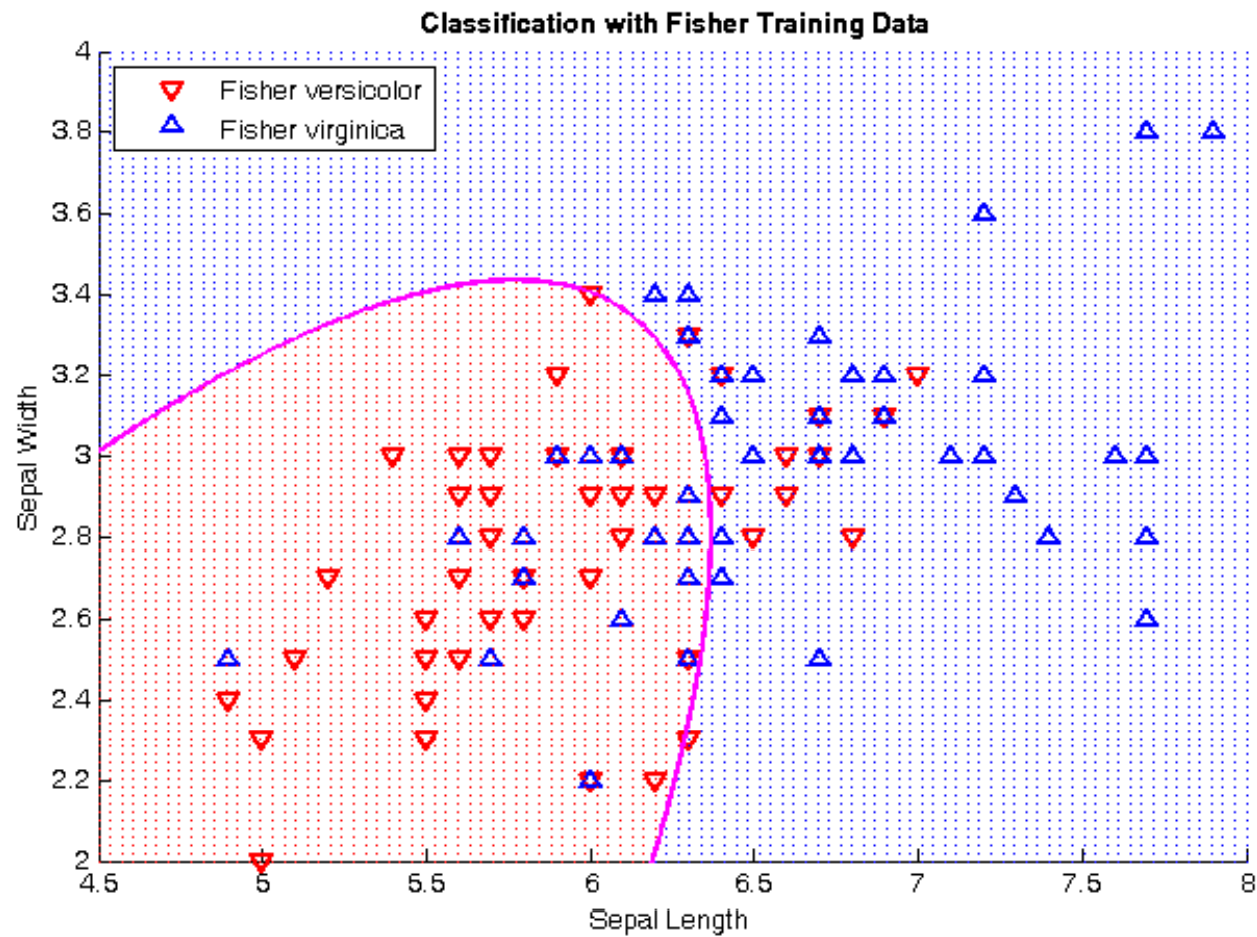
% Visualize the classification

```
hold on;  
gscatter(X,Y,C,'rb','.',1,'off');  
K = coeff(1,2).const;  
L = coeff(1,2).linear;  
Q = coeff(1,2).quadratic;  
f = sprintf('0 = %g+%g*x+%g*y+%g*x^2+%g*x.*y+%g*y.^2',  
    K,L,Q(1,1),Q(1,2)+Q(2,1),Q(2,2));  
h2 = ezplot(f,[4.5 8 2 4]);  
set(h2,'Color','m','LineWidth',2)  
axis([4.5 8 2 4])  
xlabel('Sepal Length')  
ylabel('Sepal Width')  
title('{\bf Classification with Fisher Training Data}')
```

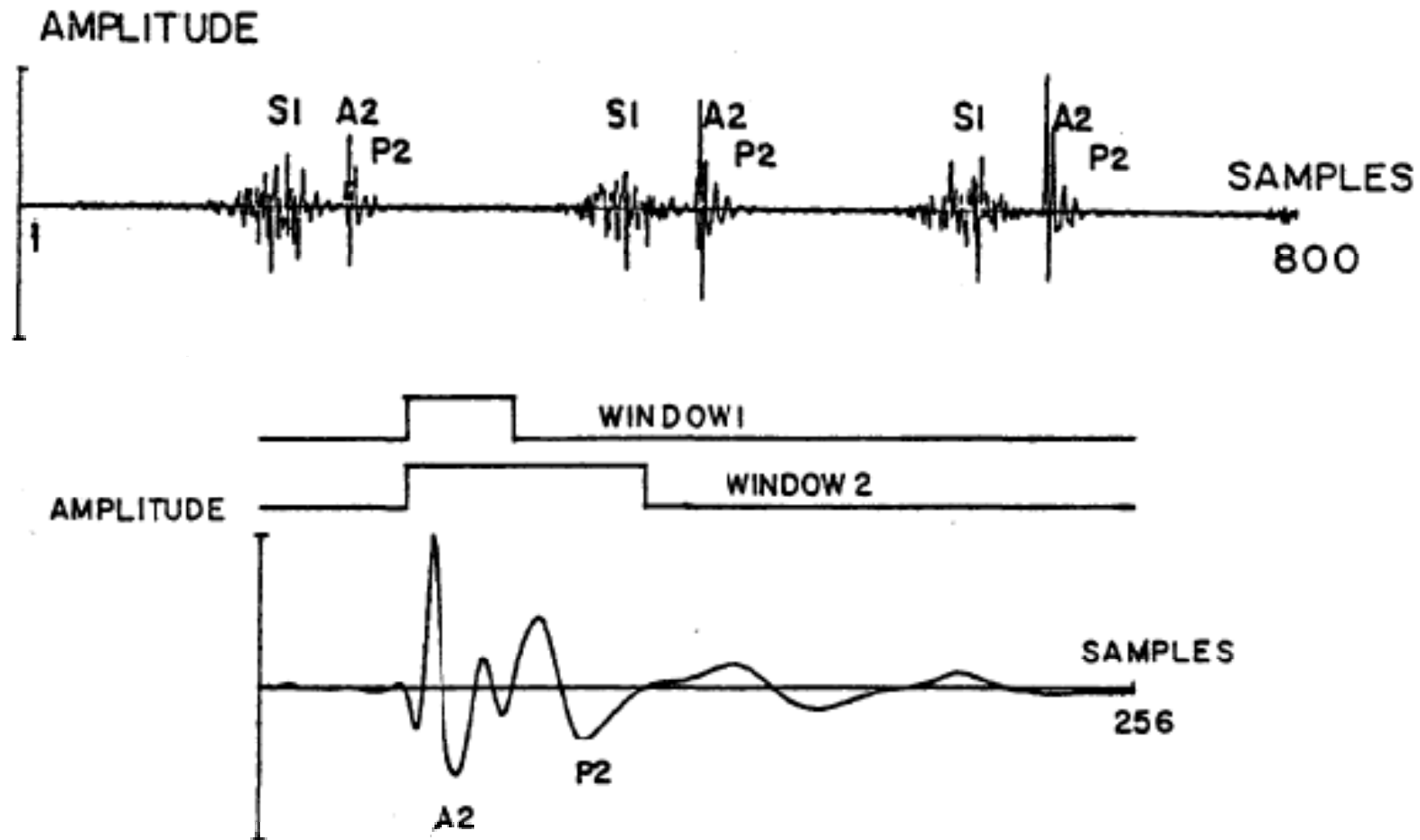
Classifying patterns



Classification



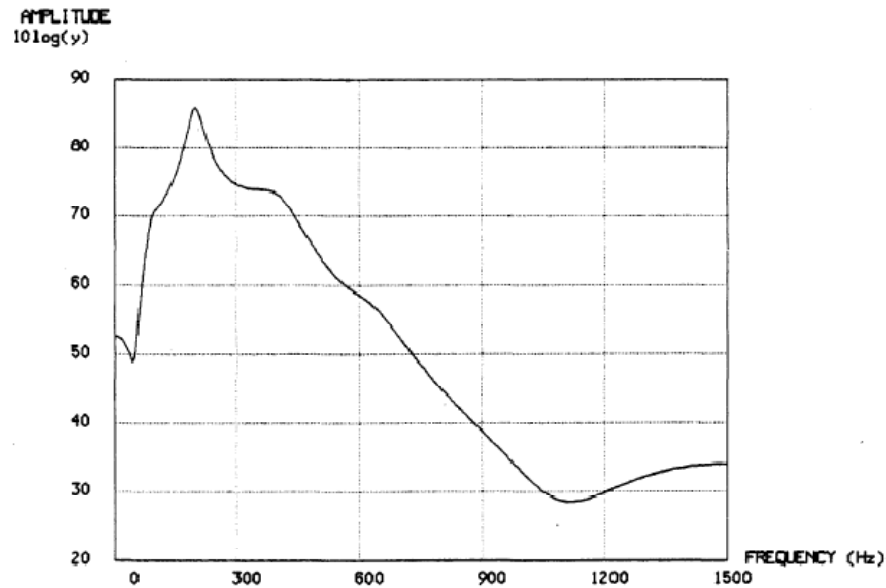
Application in PCG diagnosis



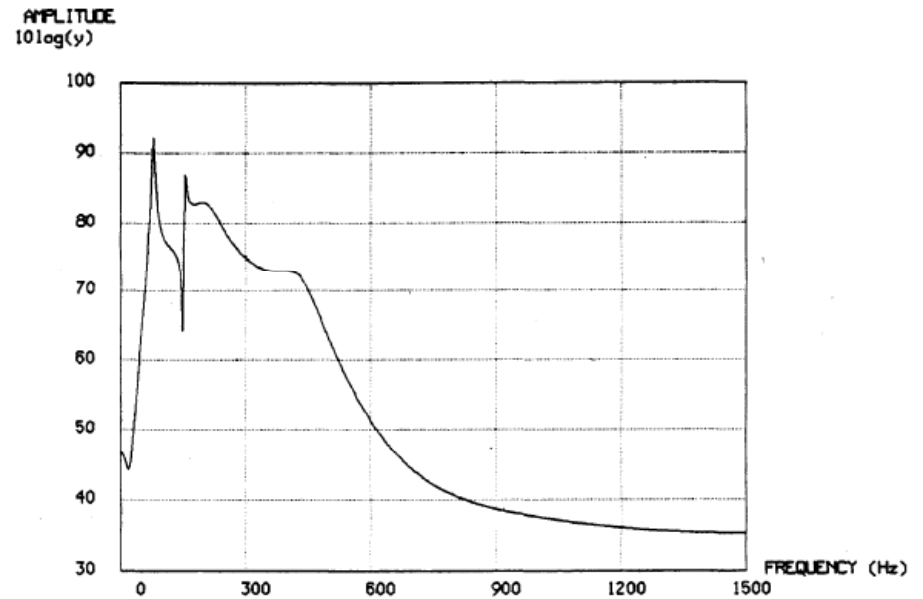
TH. Joo, et al IEEE Trans Biomed Eng 1983

High-resolution spectral analysis by pole-zero modeling

Using window 1



Using window 2



Training sets

TABLE I
THE TRAINING SET USED TO DESIGN THE CLASSIFIER

| Training Set | | |
|--------------|--------|--------|
| Valve | f1(Hz) | f2(Hz) |
| normal#1 | 172. | 79. |
| normal#2 | 140. | 99. |
| normal#3 | 82. | 114. |
| normal#4 | 84. | 123. |
| normal#5 | 123. | 155. |
| normal#6 | 90. | 131. |
| normal#7 | 79. | 137. |
| normal#8 | 134. | 99. |
| normal#9 | 146. | 102. |
| normal#10 | 111. | 73. |
| normal#11 | 93. | 73. |
| normal#12 | 96. | 131. |
| normal#13 | 87. | 143. |
| normal#14 | 125. | 70. |
| average | 121. | 109. |
| abnormal#1 | 184. | 257. |
| abnormal#2 | 158. | 216. |
| abnormal#3 | 213. | 87. |
| abnormal#4 | 193. | 117. |
| abnormal#5 | 178. | 290. |
| abnormal#6 | 131. | 193. |
| average | 176. | 193. |

Classifier design

$$g(x) = g_n(x) - g_a(x)$$

$$g(x) = (x - \mu_a)^T \Sigma_a^{-1} (x - \mu_a) - (x - \mu_n)^T \Sigma_n^{-1} (x - \mu_n) + \ln \frac{|\Sigma_a|}{|\Sigma_n|} - 2 \ln \frac{P(C_a)}{P(C_n)}$$

$$g(x) = \begin{bmatrix} f_1 - 176 \\ f_2 - 193 \end{bmatrix}^T \begin{bmatrix} 1.81e-3 & 2.88e-4 \\ 2.88e-4 & 5.82e-10 \end{bmatrix} \begin{bmatrix} f_1 - 176 \\ f_2 - 193 \end{bmatrix} \\ - \begin{bmatrix} f_1 - 121 \\ f_2 - 109 \end{bmatrix}^T \begin{bmatrix} 1.65e-3 & 7.85e-4 \\ 7.85e-4 & 1.69e-3 \end{bmatrix} \begin{bmatrix} f_1 - 121 \\ f_2 - 109 \end{bmatrix} + 1.82$$

Result

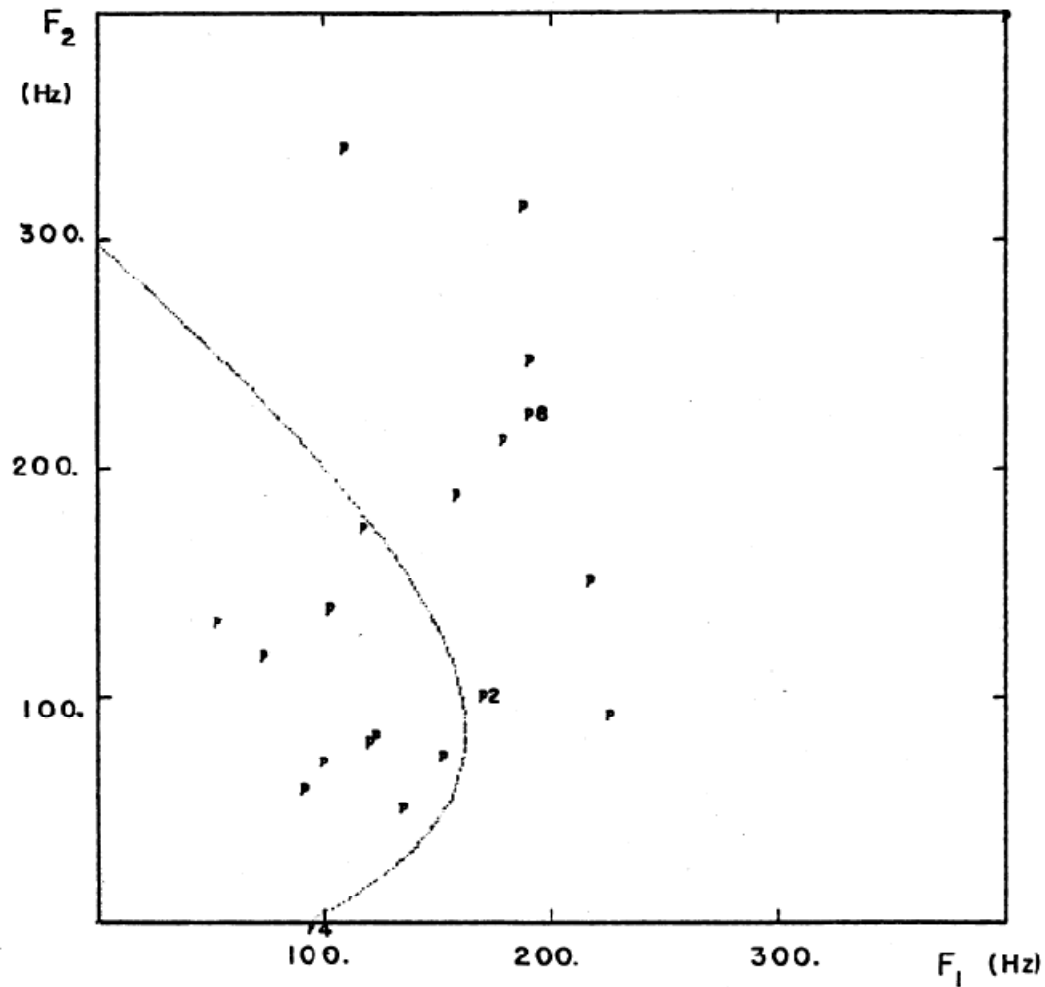


Fig. 6. Test set and the decision boundary in the feature space. Misclassified patients are Patients 2, 4, and 8. Refer to Table II for numerical values.

Test sets

TABLE II
TEST SET - OUTCOME OF THE CLASSIFIER AND THE CLINICAL DIAGNOSIS

| Test Set | | | | | |
|----------|------|------|-------|----------------|-----------|
| patient# | f1 | f2 | g(x) | classification | diagnosis |
| 1 | 73. | 120. | 23.6 | normal | normal |
| 2 | 169. | 102. | -2.68 | abnormal | normal |
| 3 | 120. | 82. | 10.2 | normal | normal |
| 4 | 93. | 0.0 | -0.45 | abnormal | normal |
| 5 | 52. | 134. | 29.4 | normal | normal |
| 6 | 123. | 84. | 9.64 | normal | normal |
| 7 | 134. | 52. | 4.14 | normal | normal |
| 8 | 190. | 225. | -44.7 | abnormal | normal |
| 9 | 99. | 73. | 14.8 | normal | normal |
| 10 | 90. | 61. | 15.5 | normal | normal |
| 11 | 102. | 140. | 12.8 | normal | normal |
| 12 | 117. | 175. | 0.9 | normal | normal |
| 13 | 152. | 76. | 2.1 | normal | normal |
| 14 | 158. | 190. | -17.9 | abnormal | abnormal |
| 15 | 216. | 152. | -24.4 | abnormal | abnormal |
| 16 | 178. | 213. | -34.7 | abnormal | abnormal |
| 17 | 225. | 93. | -15.5 | abnormal | abnormal |
| 18 | 108. | 342. | -85.8 | abnormal | abnormal |
| 19 | 190. | 249. | -57.5 | abnormal | abnormal |
| 20 | 187. | 316. | -103. | abnormal | abnormal |

Measure of decision accuracy

- Let A be the class of abnormal and N the class of normal.
- True positive (TP)
 - The case when test is positive for a subject with disease

$$\text{Sensitivity} = \text{TP fraction} = P(T^+ | A) = \frac{\text{number of TP decision}}{\text{number of subjects with disease}}$$

- True negative (TN)
 - The case when test is negative for a subject without disease

$$\text{Specificity} = \text{TN fraction} = P(T^- | N) = \frac{\text{number of TN decision}}{\text{number of subjects without disease}}$$

Measure of decision accuracy (cont.)

- False negative (FN)

$$\text{FN fraction} = P(T^- | A) = \frac{\text{number of negative decision who has disease}}{\text{number of subjects with disease}}$$

- False positive (FP)

$$\text{FP fraction} = P(T^+ | N) = \frac{\text{number of positive decision who has not disease}}{\text{number of subjects without disease}}$$

Measure of decision accuracy (cont.)

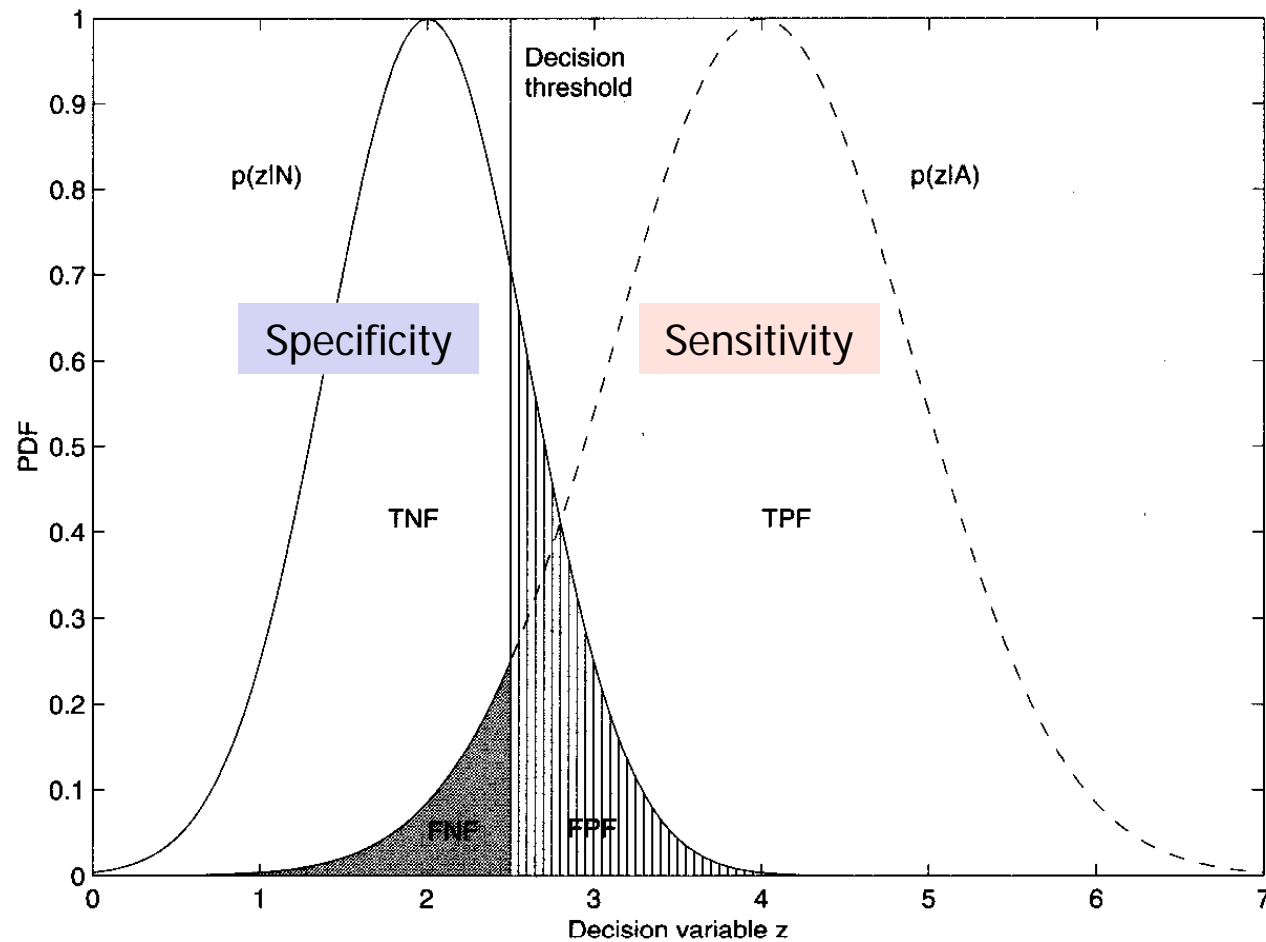
| Actual Group | Predicted Group | |
|--------------|-----------------|----------|
| | Normal | Abnormal |
| Normal | TN | FP |
| Abnormal | FN | TP |

$$TPF + FNF = 1$$

$$TNF + FPF = 1$$

$$\text{Accuracy} = P(A) P(T^+|A) + P(N) P(T^-|N)$$

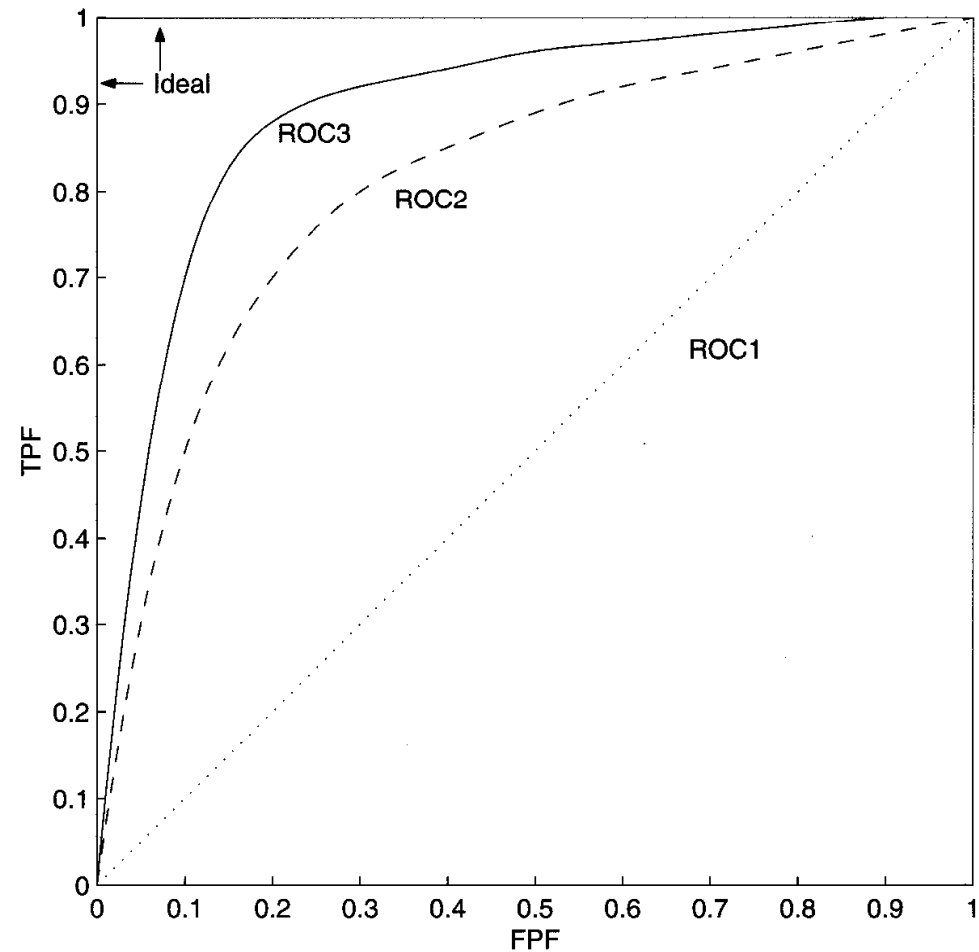
Measure of decision accuracy (cont.)



Decision threshold is the tradeoff between sensitivity and specificity

Receiver operating characteristic (ROC)

- Curve indicate the relationship between sensitivity and (1-specificity)
- A receiver (user) may choose to operate at any point along the curve



Reference

- R. Rangayyan, Biomedical Signal Analysis, John Wiley & Sons, 2002.
- TH. Joo et al, Pole-Zero Modeling and Classification of Phonocardiograms, IEEE Trans. on Biomedical Engineering, 30(2): 110-117, 1983.