

METU Informatics Institute

Min720

Pattern Classification Bio-Medical Applications

Lecture Notes by Neşe Yalabık Spring 2011

Part 3: Estimation of Parameters

Estimation of Parameters

- Most of the time, we have random samples but not the densities given.
- If the parametric form of the densities are given or assumed, then, using the labeled samples, the parameters can be estimated. (supervised learning)

Maximum Likelihood Estimation of Parameters

• Assume we have a sample set:

$$D = \{X_1, X_2, ..., X_n\}$$

- as belonging to a given class. Drawn from $P(X|\omega_j)$

iid (independently drawn from identically distributed r.v.) samples

$$\theta_{j} = [t_{1}, t_{2}, ..., t_{p}]^{T} \text{ (unknown parameter vector)}$$

$$\theta_{j} = (\mu_{j}, \Sigma_{j})^{T} = [\mu_{j1}, \mu_{j2}, ..., \Sigma_{j11}, ...] \text{ for gaussian}$$

The density function $P(X|\omega_j)$ - assumed to be of known form So our problem: estimate θ_j using sample set: $D_i = \{X_{i1}, X_{i2}, ..., X_{in}\}$ *iid*

Now drop j and assume a single density function.

 $\hat{\theta}$: estimate of θ

Anything can be an estimate. What is a good estimate?

- Should converge to actual values
- Unbiased etc

Consider the mixture density $L(\theta) = P(D|\theta) = \prod_{i=1}^{n} P(X_i|\theta)$ (due to statistical independence) $L(\theta)$ is called "likelihood function" $\hat{\theta} - \theta$ that maximizes $L(\theta)$ (Best agrees with drawn samples.) if θ is a singular, Then find θ such that $\frac{dL}{d\theta} = 0$ and for solving for θ . When θ is a vector, then $L = L(t_1, t_2, ..., t_p)$ $\nabla_{\theta} L = 0$

$$\nabla : \text{gradient of} \quad \mathsf{L} \quad \text{wrt} \quad \theta$$

$$Where: \quad \nabla_{\theta} = \begin{bmatrix} \frac{\partial L}{\partial t_1} \\ \frac{\partial L}{\partial t_2} \\ \vdots \\ \frac{\partial L}{\partial t_p} \end{bmatrix} = 0$$

Therefore
$$\hat{\theta} = \arg \max L(\theta)$$

or $\hat{\theta} = \arg \max \ln L(\theta) = \arg \max l(\theta)$ (log-likelihood)

(Be careful not to find the minimum with derivatives)

Example 1: Consider an exponential distribution $f(X;\theta) = \begin{cases} \theta e^{-\theta x} & x \ge 0\\ 0 & \text{otherwise} \end{cases}$ (single feature, single parameter) $\theta |_{f(X|\theta)}$ $L(\theta)$ With a random sample $\{X_1, X_2, ..., X_n\}$ $L(\theta) = f(X_1, X_2, ..., X_n | \theta) = \prod_{i=1}^n \theta \cdot e^{-\theta \cdot x_i}$ valid for $x \ge 0$ valid for $l(\theta) = \ln L(\theta) = \sum_{i=1}^{n} \ln \theta - \theta \sum_{i=1}^{n} x_i = n \ln \theta - \theta \sum_{i=1}^{n} x_i$ $\frac{dl}{d\theta} = \frac{d\ln L(\theta)}{d\theta} = \frac{n}{\theta} - \sum_{i=1}^{n} x_{i} = 0$ $\Rightarrow \frac{n}{\hat{\theta}} = \sum_{i=1}^{n} x_i \Rightarrow \hat{\theta} = \frac{1}{\frac{1}{n} \sum_{i=1}^{n} x_i}$ (inverse of average)

Example 2:

- Multivariate Gaussian with unknown mean vector M. Assume \sum is known.
- k samples from the same distribution:

$$X_{1}, X_{2}, \dots, X_{k}$$
(iid)

$$L(X \mid M) = \prod_{i=1}^{k} \frac{1}{(2\pi)^{n/2} |\Sigma|^{1/2}} e^{-\frac{1}{2}(X_{i} - M)^{T} \Sigma^{-1}(X_{i} - M)}$$

$$\nabla l = \nabla_{M} \log L = \sum_{i=1}^{k} \nabla_{M} \log \frac{1}{(2\pi)^{n/2} |\Sigma|^{1/2}} e^{-\frac{1}{2}(X_{i} - M)^{T} \Sigma^{-1}(X_{i} - M)}$$

$$= \sum_{i=1}^{k} \nabla_{M} (\frac{n}{2} \log(2\pi) - \frac{1}{2} \log|\Sigma| - \frac{1}{2} (X_{i} - M)^{T} \Sigma^{-1}(X_{i} - M))$$

$$= \sum_{i=1}^{k} (\Sigma^{-1}(X_{i} - M))$$
(linear algebra)

$$\Rightarrow 0 = \Sigma^{-1} (\sum_{i=1}^{k} X_i - k \hat{M})$$

$$\hat{M} = \frac{1}{k} \sum_{i=1}^{k} X_i \qquad \text{(sample average or sample mean)}$$

Estimation of \sum when it is unknown. (Do it yourself: not so simple)

$$\hat{\Sigma} = \frac{1}{n} \sum_{k=1}^{n} (X_k - \hat{M}) (X_k - \hat{M})^T \qquad \hat{\Sigma} \quad \text{(sample covariance)}$$

where M is the same as above. Biased estimate : $E(\sigma^2) \neq \sigma^2$

$$=\frac{n-1}{n}\sigma^{2}$$
 use $\frac{1}{n-1}\sum$ for an unbiased estimate.

Example 3:

Binary variables with unknown parameters $p_i, 1 \leq i \leq n$ (n parameters)

$$\begin{split} \log P(X) &= \sum_{i=1}^{n} x_i \log p_i + \sum_{i=1}^{n} (1 - x_i) \log(1 - p_i) \\ \text{So,} \\ l &= \log L = \sum_{j=1}^{k} \log P(X_j) \text{ k samples} \\ &= \sum_{j=1}^{k} (\sum_{i=1}^{n} x_{ij} \log p_i + \sum_{i=1}^{n} (1 - x_{ij}) \log(1 - p_i) \\ \text{here } X_{ij} \text{ is the } i^{th} \text{ element of } j^{th} \text{ sample } X_j \text{ .} \end{split}$$

So,

$$\nabla_{p_{i}} \log L = \begin{bmatrix} \frac{\partial}{\partial p_{1}} \log L \\ \frac{\partial}{\partial p_{2}} \log L \\ \vdots \\ \frac{\partial}{\partial p_{n}} \log L \end{bmatrix}$$

$$\frac{\partial}{\partial p_{i}} \log L = \sum_{j=1}^{k} \left(\frac{x_{ij}}{p_{i}} - ((1 - x_{ij})(1 - p_{i})) \right)$$

$$\Rightarrow 0 = \frac{1}{\hat{p}_{i}} \sum_{j=1}^{k} x_{ij} - \frac{1}{1 - \hat{p}_{i}} \sum_{j=1}^{k} (1 - x_{ij})$$

$$\Rightarrow \hat{p}_{i} = \frac{1}{k} \sum_{j=1}^{k} x_{ij}$$

* \hat{p}_i is the sample average of the feature.

- Since X_i is binary, $\sum_{j=1}^k x_{ij}$ will be the same as counting the occurances of '1'.
- Consider character recognition problem with binary matrices.



- For each pixel, count the number of 1's and this is the estimate of \mathcal{P}_i .



METU Informatics Institute

Min720

Pattern Classification Bio-Medical Applications

Lecture Notes by Neşe Yalabık Spring 2011

Part 4: Features and Feature Extraction

Problems of Dimensionality and Feature Selection

- Are all features independent? Especially in binary features, we might have >100.
- The classification accuracy vs. size of feature set.
- Consider the Gaussian case with same Σ for both categories.

$$P(e) = \frac{1}{\sqrt{2\pi}} \int_{r/2}^{\infty} e^{-u^2/2} du$$

(assuming a priori probabilities are the same) (e:error)

• where r^2 is the square of mahalonobis distance between class means.

$$r^{2} = (\mu_{1} - \mu_{2})^{T} \Sigma^{-1} (\mu_{1} - \mu_{2})$$



• P(e) decreases as r increases (the distance between the means).

If
$$\Sigma = \begin{bmatrix} \sigma_1^2 & & 0 \\ & \sigma_2^2 & & \\ & & \ddots & \\ 0 & & & \sigma_d^2 \end{bmatrix}$$

(all features statistically independent.)

then

$$r^{2} = \sum_{i=1}^{d} \left(\frac{m_{i1} - m_{i2}}{\sigma_{i}}\right)^{2} = \sum_{i=1}^{d} \frac{\left(m_{i1} - m_{i2}\right)^{2}}{\sigma_{i}^{2}}$$

We conclude from here that

- 1-Most useful features are the ones with large distance and small variance.
- 2-Each feature contributes to reduce the probability of error.



- When r increases, probability of error decreases.
- Best features are the ones with distant means and small variances.
- So add new features if the ones we already have are not adequate (more features, decreasing prob. of error.)
- But it was shown that adding new features after some point leads to worse performance.

- ✓ Find statistically independent features
- \checkmark Find discriminating features
- ✓ Computationally feasible features

<u>Principal Component Analysis (PCA) (Karhunen-Loeve</u> <u>Transform)</u>

• Finds (reduces the set) to statistically independent features.



So we either

- □ Throw one away
- \Box Generate a new feature using \mathcal{Y}_1 and \mathcal{Y}_2 (ex:projections of the points to a line)
- $\hfill\square$ Form a linear combination of features.

$$x_{1} = f_{1}(y_{1}, \dots, y_{m})$$

$$x_{2} = f_{2}(y_{1}, \dots, y_{m})$$
Linear
functions
$$x_{d} = f_{d}(y_{1}, \dots, y_{m})$$

-Find a vector X_0 so that sum of the squared distances to X_0 is minimum(Zero degree representation).



$$= \sum \|X_0 - M\|^2 - 2(X_0 - M)^T \sum (X_K - M) + \sum \|X_k - M\|^2$$

$$= \sum \left\| X_0 - M \right\|^2 + \underbrace{\sum \left\| X_k - M \right\|^2}_{\text{Independent of } X_0}$$

Where $X_0 = M$, this expression is minimized.

Consider now 1-d representation from 2-d. -The line should pass through the sample mean.

$$X = M + ae$$
 unit vector in the direction of line



- Now how to find best e that minimizes $J_1 = \sum ||(M + a_k e) X_k||^2$
- It turns out that given the scatter matrix

$$S = \sum_{k=1}^{n} (X_{k} - M)(X_{k} - M)^{T}$$

- e must be the eigenvector of the scatter matrix with the largest eigenvalue lambda λ .

 $Se = \lambda e$

- That is, we project the data onto a line through the sample mean in the direction of <u>the eigenvector of the scatter matrix with</u> <u>largest eigenvalue.</u>
- Now consider d dimensional projection

$$X = M + \sum_{i=1}^{d} a_i e_i$$

 Here e₁,...., e_d are d eigenvectors of the scatter matrix having largest eigenvalues.

<u>Coefficients a_i are called principal components.</u>

• So each m dimensional feature vector is transferred to d dimensional space since the components a_i are given as

$$a_{ki} = e_i^T (X_k - M)$$

<u>Now represent our new feature vector's elements</u>
 So

$$a_{1i} = e_1^T (X_k - M)$$

$$a_{2i} = e_2^T (X_k - M)$$

$$\vdots$$

$$a_{d'i} = e_d^T (X_k - M)$$

FISHER'S LINEAR DISCRIMINANT

- Curse of dimensionality. More features, more samples needed.
- We would like to choose features with more discriminating ability.
- Reduces the dimension of the problem to one in simplest form.
- <u>Seperates samples from different categories.</u>
- <u>Consider samples from 2 different categories now.</u>



-Find a line so that the projection separates the samples best.

Same as:

Apply a transformation to samples X to result with a scalar such that $y = W^T X$ Fisher's criterion function

$$J(W) = \frac{(\mu_1 - \mu_2)^2}{\sigma_1^2 + \sigma_2^2} \quad \text{is maximized, where}$$

$$\mu_i = \frac{1}{n_i} \sum_{y \in C_i} y$$

$$\sigma_i^2 = \frac{1}{n_i} \sum_{y \in C_i} (y - \mu_i)^2$$

- This reduces the problem to <u>1d</u>, by keeping the classes most distant from each other.
- But if we write μ_i and σ_i^2 in terms of M_i and Σ_i

$$M_{i} = \frac{1}{n_{i}} \sum_{X \in C_{i}} X$$

$$\mu_{i} = \frac{1}{n_{i}} \sum_{X \in C_{i}} W^{T} X = W^{T} M$$

$$\sigma_{i}^{2} = \frac{1}{n_{i}} \sum_{X \in C_{i}} (W^{T} X - W^{T} M_{i})^{2} = \frac{1}{n_{i}} \sum_{X \in C_{i}} (W^{T} (X - M_{i}))^{2}$$

$$=\frac{1}{n_i}\sum W^T (X - M_i)(X - M_i)^T W = W^T (\frac{1}{n_i}(\sum (X - M_i)(X - M_i)^T)W)$$

$$= W^{T}S_{i}W$$
Then, $(\mu_{1} - \mu_{2})^{2} = (W^{T}M_{1} - W^{T}M_{2})^{2} = [W^{T}(M_{1} - M_{2})]^{2}$

$$= W^{T}(M_{1} - M_{2})(M_{1} - M_{2})^{T}W = W^{T}S_{B}W$$

$$\sigma_1^2 + \sigma_2^2 = W^T (S_1 + S_2) W = W^T S_W W$$

 S_{B} - within class scatter matrix

 $S_{\scriptscriptstyle W}^{}$ - between class scatter matrix ${\mbox{\cdot}}$ Then , maximize

• It can be shown that W that maximizes J can be found by solving the eigenvalue problem again:

 ${S_W}^{-1}S_BW = \lambda W$ and the solution is given by

 $W = S_W^{-1}(M_1 - M_2)$

 Optimal if the densities are gaussians with equal covariance matrices. That means reducing the dimension does not cause any loss.

Multiple Discriminant Analysis: c category problem. A generalization of 2-category problem.

Non-Parametric Techniques

- Density Estimation
- Use samples directly for classification
 - Nearest Neighbor Rule
 - 1-NN
 - k-NN
- Linear Discriminant Functions: gi(X) is linear.