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- Bayes classifier is optimal for minimizing risk. Risk minimization is a very good objective.
- Given class conditional densities we can derive the Bayes classifier for any loss function.
- There are other ways (other than loss function) to trade different errors. For example, NP classifier.
- ROC curve also allows for such trade-off

# Receiver Operating Characteristic (ROC)

- Consider a one dimensional feature space, 2-class problem with a classifier,  $h(X) = 0$  if  $X < \tau$ .
- Consider equal priors, Gaussian class conditional densities with equal variance, 0-1 loss. Now let us write the probability of error as a function of  $\tau$ .

# Receiver Operating Characteristic (ROC)

$$\begin{aligned} P[\text{error}] &= 0.5 \int_{-\infty}^{\tau} f_1(X) dX + 0.5 \int_{\tau}^{\infty} f_0(X) dX \\ &= 0.5 \Phi \left( \frac{\tau - \mu_1}{\sigma} \right) + 0.5 \left( 1 - \Phi \left( \frac{\tau - \mu_0}{\sigma} \right) \right) \end{aligned}$$

- As we vary  $\tau$  we trade one kind of error with another. In Bayes classifier, the loss function determines the ‘exchange rate’.

# ROC curve

- The receiver operating characteristic (ROC) curve is one way to conveniently visualize and exploit this trade off.
- For a two class classifier there are four possible outcomes of a classification decision – two are correct decisions and two are errors.
- Let  $e_i$  denote probability of wrongly assigning class  $i$ ,  $i = 0, 1$ .

# ROC curve

Then we have

$$e_0 = P[X \leq \tau \mid X \in \mathbf{c-1}] \quad (\text{a miss})$$

$$e_1 = P[X > \tau \mid X \in \mathbf{c-0}] \quad (\text{false alarm})$$

$$1 - e_0 = P[X > \tau \mid X \in \mathbf{c-1}] \quad (\text{correct detection})$$

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- For fixed class conditional densities, if we vary  $\tau$  the point  $(e_1, 1 - e_0)$  moves on a smooth curve in  $\mathbb{R}^2$ .
- This is traditionally called the ROC curve. (Choice of coordinates is arbitrary)

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- Hence, varying  $\tau$  we can find ROC and decide which may be the best operating point.
- This can be done for any threshold based classifier irrespective of class conditional densities.
- When the class conditional densities are Gaussian with equal variance, we use this procedure to estimate Bayes error also.

- From our earlier error integral we get

$$\frac{\tau - \mu_0}{\sigma} = \Phi^{-1}(1 - e_1) = a, \text{ say}$$

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- Then,  $|a - b| = \frac{|\mu_1 - \mu_0|}{\sigma} = d$ , the discriminability.
- Knowing  $e_1, (1 - e_0)$ , we can get  $d$  and hence the Bayes error. For our given  $\tau$  we can also get the actual error probability. We can tweak  $\tau$  to match the Bayes error.



- We can in general use the ROC curve in multidimensional cases also. Consider, for example,

$$h(\mathbf{X}) = \text{sgn}(\mathbf{W}^t \mathbf{X} + w_0).$$

We can use ROC to fix  $w_0$  after learning  $\mathbf{W}$ .

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- We need class conditional densities and prior probabilities.
- Prior probabilities can be estimated as fraction of examples from each class.
- Since examples are *iid* and the class labels of examples are known, we have some iid samples from each class conditional distribution.
- The problem: Given  $\{x_1, x_2, \dots, x_n\}$  drawn *iid* according to some distribution, estimate the probability distribution / density.

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- Two main approaches: Parametric and non-parametric.
- Parametric: We assume we have *iid* realizations of a random variable  $X$  whose distribution is known except for values of a parameter vector. We estimate the parameters of the density using the samples available.
- In non-parametric approach we do not assume form of density. It is often modelled as a convex combination of some densities using the samples.



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- Now estimation of density is same as estimation of a parameter vector.

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- $\mathbf{x} = (x_1, \dots, x_n)^T$  – the sample or the data.  
We sometimes use  $\mathcal{D}$  to denote the data.
- It can be thought of as a realization of  $(X_1, \dots, X_n)^T$   
where  $X_i$  are *iid* with density  $f(x | \theta)$ .



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- When we need to remember the sample size, we write  $\hat{\theta}_n$
- For example,

$$\hat{\theta}_n = \frac{1}{n} \sum_{i=1}^n x_i$$

the well-known sample mean.

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- How does one choose estimators



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- In this course, we will consider two methods: Maximum likelihood and Bayesian estimators.
- To begin with, a simple introduction to some general issues in estimation.

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- The  $\hat{\theta}$  is a function of data. Hence the expectation is with respect to the joint density of  $(X_1, \dots, X_n)$ , the *iid* random variables.
- Since  $X_i \sim f(x | \theta)$ , the expectation above needs value of  $\theta$ . So, we write

$$E_{\theta}[\hat{\theta}] = \theta$$

- An unbiased estimator,  $\hat{\theta}$  satisfies

$$E_{\theta}[\hat{\theta}] = \theta$$

- $\hat{\theta}$  is an unbiased estimator, if for every density in the class of densities we are interested in (i.e., every value of the parameter in the parameter space), expected value of the estimator is the true parameter value.



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- So is  $\hat{\theta}'' = x_1$ .
- Unbiasedness alone is not enough

- One possibility: We can say  $\hat{\theta}$  is better than  $\hat{\theta}'$  if,  $\forall \theta$ ,  
$$P_{\theta}[-a \leq (\hat{\theta} - \theta) \leq b] \geq P_{\theta}[-a \leq (\hat{\theta}' - \theta) \leq b] \quad \forall a, b > 0$$
  
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(for any fixed sample size)
- Difficult to get such estimators.



- A weaker method is:  $\hat{\theta}$  is better than  $\hat{\theta}'$  if

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- The mean square error of an estimator is defined by

$$\text{MSE}_{\theta}(\hat{\theta}) = E_{\theta}[(\hat{\theta} - \theta)^2]$$

- Lemma:

$$\text{MSE}_\theta(\hat{\theta}) = V_\theta(\hat{\theta}) + [B_\theta(\hat{\theta})]^2$$

where  $V_\theta(\hat{\theta})$  is the variance given by

$$V_\theta(\hat{\theta}) = E_\theta[(\hat{\theta} - E_\theta[\hat{\theta}])^2]$$

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- For unbiased estimators the variance is the mean square error (because bias is zero).

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and for all  $\hat{\theta}'$  that are unbiased estimators for  $\theta$ .
- If we can get an UMVUE, then it is the 'best' estimator.
- In many cases, it is difficult to get UMVUE.

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- We can also think of asymptotic properties.
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- For example, the sample mean is a consistent estimator of population mean (expectation of the random variable)  
(Law of large numbers)

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- This is not an unbiased estimator.
- But we have the following


$$E[(\hat{\theta}_n - \theta)^2] = E \left[ \left( \frac{1}{n+1} \sum_{i=1}^n (x_i - \theta) - \frac{1}{n+1} \theta \right)^2 \right]$$

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 &= \frac{1}{(n+1)^2} n \sigma^2 + \frac{1}{(n+1)^2} \theta^2 - \\
 &\quad \frac{2\theta}{(n+1)^2} E \left[ \sum (x_i - \theta) \right]
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 &= \frac{1}{(n+1)^2} n \sigma^2 + \frac{1}{(n+1)^2} \theta^2 - \\
 &\quad \frac{2\theta}{(n+1)^2} E \left[ \sum (x_i - \theta) \right] \\
 &= \frac{n}{(n+1)^2} \sigma^2 + \frac{1}{(n+1)^2} \theta^2
 \end{aligned}$$

- Thus,  $E[(\hat{\theta}_n - \theta)^2] \rightarrow 0$  as  $n \rightarrow \infty$ .
- Hence,  $\hat{\theta}$  is consistent (though it is biased).

- Maximum Likelihood (ML) estimation is a general procedure for obtaining consistent estimators.
- It is a parametric method.
- We estimate parameters of a density based on *iid* samples.
- For most densities, ML estimates are consistent.

# Maximum likelihood estimation

- Let  $\mathbf{x} = \{x_1, x_2, \dots, x_n\}$  be the samples.
- Likelihood function is defined by

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- If samples are from a discrete random variable,  $f$  is taken to be the mass function. If samples are from a continuous random variable, then  $f$  is the density function.



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- We essentially look at the likelihood function as a function of  $\theta$  with the  $x_j$  being known values (as given by data).
- To emphasize this we write it as  $L(\theta, \mathbf{x})$  or  $L(\theta \mid \mathbf{x})$  or  $L(\theta \mid \mathcal{D})$ .

Recall that we denote the data samples by  $\mathcal{D}$  also.

# Maximum likelihood estimation contd..

- The maximum likelihood (ML) estimate of  $\theta$  is the value that (globally) maximizes the likelihood function.

# Maximum likelihood estimation contd..

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- Finding MLE is an optimization problem.

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- For many densities we can analytically solve for the maximizer.
- In general we can use numerical optimization techniques.

# Example

- Consider one dimensional case.  
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- Now the likelihood is given by

$$L(\theta \mid \mathbf{x}) = \prod_{j=1}^n \frac{1}{\theta_2 \sqrt{2\pi}} \exp \left( -\frac{(x_j - \theta_1)^2}{2\theta_2^2} \right)$$

- Hence log likelihood would be

$$l(\theta \mid \mathbf{x}) = \sum_{j=1}^n \left[ -\log(\theta_2) - 0.5 \log(2\pi) - \frac{(x_j - \theta_1)^2}{2\theta_2^2} \right]$$

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- To maximize log likelihood we equate the partial derivatives to zero.

- This gives

$$\frac{\partial l}{\partial \theta_1} = \sum_{j=1}^n (x_j - \theta_1) = 0$$

$$\frac{\partial l}{\partial \theta_2} = -\frac{n}{\theta_2} + \frac{1}{\theta_2^3} \sum_{j=1}^n (x_j - \theta_1)^2 = 0$$



- Solving these, we get

$$\hat{\theta}_1 = \frac{1}{n} \sum_{j=1}^n x_j$$

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- ML estimate of variance is **not** unbiased.

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- The mass function has only one parameter, namely,  $p$ .
- Note that we must have  $0 \leq p \leq 1$ .

- The likelihood function is

$$L(p \mid \mathbf{x}) = \prod_{j=1}^n p^{x_j} (1 - p)^{1-x_j} = p^{n\bar{x}} (1 - p)^{n-n\bar{x}}$$

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- The loglikelihood is given by

$$l(p \mid \mathbf{x}) = n\bar{x} \log p + n(1 - \bar{x}) \log(1 - p)$$

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- This is the ML estimate of the parameter  $p$  of a Bernoulli random variable.
- Sample mean is the ML estimator.

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- Often, one maximizes loglikelihood
- For many standard densities we can obtain MLE analytically.