

## The Generalized Likelihood Ratio Test

The generalized likelihood ratio test (GLRT) gets around the composite hypothesis testing problem by effectively turning it into a test between two simple hypotheses.

These simple hypotheses are selected to be the most likely value of

 $\theta_0 \in \Theta_0$  under  $H_0$ 

and

$$\theta_1 \in \Theta_1$$
 under  $H_1$ 

given the observed data.

# The Generalized Likelihood Ratio Test (Cont.)

- O Using these two simple hypotheses, a likelihood ratio test is implemented to test between them.
- The composite hypothesis corresponding to the simple hypothesis declared by the simple likelihood test is the composite hypothesis declared by the GLRT.
- While the GLRT is not optimal in any particular sense, it seems like a reasonable approach to dealing with the composite hypothesis testing problem.
- In many cases where a UMP test does exist, the GLRT exhibits nearly optimal behavior.

## Generalized Likelihood Ratio Test (GLRT)

Consider two composite hypotheses  $H_0: \underline{\theta} \in \Theta_0$  and  $H_1: \underline{\theta} \in \Theta_1$ . The Generalized Likelihood Ratio Test (GLRT) consists of the following procedure:

1. Assume  $H_0$  is true and estimate the value of  $\theta$  from the observed data using a maximum likelihood estimate (MLE):

$$\underline{\hat{\theta}}_0 = \arg\max_{\theta\in\Theta_0} f_{\underline{\theta}}(\underline{X}).$$

2. Assume  $H_1$  is true and estimate the value of  $\theta$  from the observed data using a (MLE):

$$\hat{\theta}_1 = \arg \max_{\theta \in \Theta_1} f_{\underline{\theta}}(\underline{X}).$$

3. Replace the original problem of testing the composite hypotheses  $H_0$  versus  $H_1$  with the problem of testing the simple hypotheses  $\hat{H}_0 : \hat{\Theta}_0 = \{\hat{\theta}_0\}$  versus  $\hat{H}_1 : \hat{\Theta}_1 = \{\hat{\theta}_1\}$ . If  $\hat{H}_0$  is decided in the simple hypothesis problem, then  $H_0$  is decided as the composite hypothesis. If  $\hat{H}_1$  is decided in the simple hypothesis problem, then  $H_1$  is decided as the composite hypothesis.

### **GLRT-continued**

When we carry out this procedure, we get a GLRT of the form

$$L_g(\underline{X}) = \frac{\max_{\underline{\theta}\in\Theta_1} f_{\underline{\theta}}(\underline{X})}{\max_{\underline{\theta}\in\Theta_0} f_{\underline{\theta}}(\underline{X})} \stackrel{H_1}{\underset{H_0}{\gtrsim}} L_0,$$

which yields a statistical test of the form

$$\phi(\underline{X}) = \begin{cases} 1, & \text{for } L_g(\underline{X}) > L_0, \\ \gamma, & \text{for } L_g(\underline{X}) = L_0, \\ 0, & \text{for } L_g(\underline{X}) < L_0, \end{cases}$$

where, in principle,  $L_0$  and  $\gamma$  are selected to yield a size  $\alpha$  test.

# $L_0$ and $\gamma$

$$\phi(\underline{X}) = \begin{cases} 1, & \text{for } L_g(\underline{X}) > L_0, \\ \gamma, & \text{for } L_g(\underline{X}) = L_0, \\ 0, & \text{for } L_g(\underline{X}) < L_0. \end{cases}$$

Selecting  $L_0$  and  $\gamma$  to yield a size  $\alpha$  test is difficult.

The reason is that the size of the test is still defined as

 $\overline{\alpha} = \sup_{\underline{\theta} \in \Theta_0} \mathrm{E}_{\underline{\theta}} \left[ \phi(\underline{X}) \right].$ 

We do not use  $E_{\underline{\hat{\theta}}_0}[\phi(\underline{X})]$  as the size of the test.

**Example:** Suppose we wish to test the hypotheses  $H_0$  versus  $H_1$  that the random sample  $\underline{X} = (X_1, \ldots, X_N)$  comes from a density

$$f_{\theta}(x_n) = \frac{1}{\sqrt{2\pi}} \exp\left\{\frac{-(x_n - \theta)^2}{2}\right\}$$

where under  $H_0$ :  $\Theta_0 = [-1, 1]$ , and under  $H_1$ :  $\Theta_0 = \{\theta \in \mathbf{R} : |\theta| > 1\}.$ 

We note that in general,

$$f_{\theta}(\underline{X}) = \frac{1}{(2\pi)^{N/2}} \exp\left\{-\frac{1}{2}\sum_{n=1}^{N} (x_n - \theta)^2\right\},\tag{1}$$

from which it follows that the (unconstrained) maximum likelihood estimate  $\hat{\theta}_{ML}$  of  $\theta$  can be found by solving

$$\frac{\partial}{\partial \theta} f_{\theta}(\underline{X}) = 0$$

for  $\theta$ , yielding

$$\hat{\theta}_{ML} = \frac{1}{N} \sum_{n=1}^{N} X_n$$

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For this case, we can easily see that

$$\hat{\theta}_0 = \arg\max_{\underline{\theta}\in\Theta_0} f_{\underline{\theta}}(\underline{X}) = \begin{cases} \bar{\theta}_{ML}, & \text{for } \bar{\theta}_{ML} \in [-1,1], \\ -1, & \text{for } \hat{\theta}_{ML} < -1, \\ 1, & \text{for } \hat{\theta}_{ML} > 1, \end{cases}$$

and

$$\hat{\theta}_1 = \arg\max_{\underline{\theta}\in\Theta_1} f_{\underline{\theta}}(\underline{X}) = \begin{cases} \hat{\theta}_{ML}, & \text{for } \hat{\theta}_{ML} \notin [-1,1] \\ -1, & \text{for } \hat{\theta}_{ML} \in [-1,0] \\ 1, & \text{for } \hat{\theta}_{ML} \in (0,1]. \end{cases}$$

Using this  $\hat{\theta}_0$  and  $\hat{\theta}_1$  we can now construct the GLRT

$$L_g(\underline{X}) = \frac{f_{\hat{\theta}_1}(\underline{X})}{f_{\hat{\theta}_0}(\underline{X})} \stackrel{>}{\underset{H_0}{\overset{>}{\atop}}} L_0$$

with an appropriately chosen threshold  $L_0$ .

Recall: 
$$\alpha = \sup_{\underline{\theta} \in \Theta_0} E_{\underline{\theta}} [\phi(\underline{X})]; \quad \alpha \neq E_{\underline{\hat{\theta}}_0} [\phi(\underline{X})]$$

#### **Bayesian Detection Theory**

In classical detection, we decide between  $H_0: \underline{\theta} \in \Theta_0$  versus  $H_1: \underline{\theta} \in \Theta_1$  based on an observation <u>X</u> governed by the parameterized cdf  $F_{\theta}(\underline{x})$ 

Here  $\underline{\theta}$  was assumed to be an *unknown* but fixed parameter. The parameter  $\underline{\theta}$  was *not* assumed to be random, just unknown.

In the Bayesian detection framework, we again assume that our observation  $\underline{X}$  is governed by a distribution  $F_{\theta}(\underline{X})$ ,

but we now assume that  $\underline{\theta}$  is a random vector taking on one of two possible values:  $\underline{\theta}_0$  or  $\underline{\theta}_1$ . These values are taken on with probabilities

$$p_0 = P(\{\underline{\theta} = \underline{\theta}_0\}),$$
  

$$p_1 = P(\{\underline{\theta} = \underline{\theta}_1\}) = 1 - p_0,$$

where  $p_0 \in [0, 1]$ .

We have a random experiment with probability space  $(\mathcal{S}, \mathcal{F}, P)$  having two random variables  $\underline{\theta}$  and  $\underline{X}$  defined on it.

When the random experiment is performed,  $\underline{\theta}$  takes on a value from the set  $\{\underline{\theta}_0, \underline{\theta}_1\}$  with probabilities  $p_0$  and  $p_1$ , respectively.

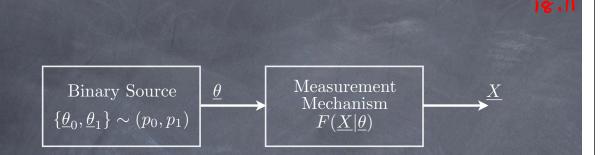
The observed value of  $\underline{X}$  takes on a value consistent with the distribution  $F_{\underline{\theta}}(\underline{x})$  for the value  $\underline{\theta}$  takes on.

Thus we have a conditional distribution function for  $\underline{X}$ :

$$F(\underline{x}|\underline{\theta}_{j}) = P(\{\underline{X} \le \underline{x}\}|\{\underline{\theta} = \underline{\theta}_{j}\}) = F_{\underline{\theta}_{j}}(\underline{x}), \quad j = 0, 1$$

The joint distribution of  $\underline{\theta}$  and  $\underline{X}$  is given by

$$F(\underline{\theta},\underline{x}) = F(\underline{x}|\underline{\theta})P(\underline{\theta}) = \begin{cases} p_0 F(\underline{x}|\underline{\theta}_0), & \text{for } \underline{\theta} = \underline{\theta}_0, \\ p_1 F(\underline{x}|\underline{\theta}_1), & \text{for } \underline{\theta} = \underline{\theta}_1. \end{cases}$$



It follows that

$$P(\underline{\theta}) = \int_{\mathbf{R}^n} dF(\underline{\theta}, \underline{x}) = \int_{\mathbf{R}^n} f(\underline{\theta}, \underline{x}) d\underline{x} = \begin{cases} p_0, & \text{for } \underline{\theta} = \underline{\theta}_0, \\ p_1, & \text{for } \underline{\theta} = \underline{\theta}_1. \end{cases}$$

and

$$F(\underline{x}) = F(\underline{x}|\underline{\theta}_0)P(\underline{\theta}_0) + F(\underline{x}|\underline{\theta}_1)P(\underline{\theta}_1)$$
$$= p_0F(\underline{x}|\underline{\theta}_0) + p_1F(\underline{x}|\underline{\theta}_1),$$

where  $\mathbf{R}^n$  is the observation space of the *n*-dimensional observation vector.

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