

## The Generalized Likelihood Ratio Test

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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

 $\overline{\theta_0} \in \overline{\Theta_0}$  under  $H_0$ 

and

$$
\theta_1 \in \Theta_1 \text{ under } H_1
$$

given the observed data.

## The Generalized Likelihood Ratio Test (Cont.)

- 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.
- **The many cases where a UMP test does exist,** the GLRT exhibits nearly optimal behavior.

## Generalized Likelihood Ratio Test (GLRT)

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Consider two composite hypotheses  $H_0: \theta \in \Theta_0$  and  $H_1: \theta \in \Theta_1$ . 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):

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

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

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

3. Replace the original problem of testing the composite hypotheses  $H_0$  versus  $H_1$  with the problem of testing the simple hypotheses  $H_0: \Theta_0 = {\theta_0}$ 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  $H_1$  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

$$
L_g(\underline{X}) = \frac{\max_{\underline{\theta} \in \Theta_1} f_{\underline{\theta}}(\underline{X})}{\max_{\underline{\theta} \in \Theta_0} f_{\underline{\theta}}(\underline{X})} \mathop{<}_{H_0} L_0,
$$

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.

# $\boxed{L_0 \text{ and } \gamma}$

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

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

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

We do not use  $E_{\hat{\underline{\theta}}_0}$  [ $\phi(\underline{X})$ ] 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  $H_1: \Theta_0 = \{ \theta \in \mathbb{R} : |\theta| > 1 \}.$ 

We note that in general,

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

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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

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

For this case, we can easily see that

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

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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

$$
L_g(\underline{X}) = \frac{f_{\hat{\theta}_1}(\underline{X})}{f_{\hat{\theta}_0}(\underline{X})} \mathop{>}_{H_0}^{H_1} L_0
$$

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  $F_{\theta}$  (*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 *X* is governed by a distribution  $F_{\theta}(\underline{X})$ ,

but we now assume that  $\theta$  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  $(S, \mathcal{F}, P)$  having two random variables  $\theta$  and  $\overline{X}$  defined on it.

When the random experiment is performed,  $\theta$  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 <u>X</u> takes on a value consistent with the distribution  $F_{\theta}(\underline{x})$ for the value  $\theta$  takes on.

Thus we have a conditional distribution function for *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  $\theta$  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  $\mathbb{R}^n$  is the observation space of the *n*-dimensional observation vector.

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