

STAT 703/J703
February 13th, 2007

-Lecture 9-

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Today

- Asymptotic Normality of the MLE (continued)
- Basics of Hypothesis Testing (Chapter 9)



Theorem B: Under appropriate regularity conditions the MLE is asymptotically normal with mean θ and variance $\frac{1}{nI(\theta)}$.



Sketch of proof: Consider the Taylor series expansion:

$$L'(\hat{\theta}) \approx L'(\theta) + (\hat{\theta} - \theta)L''(\theta)$$



Statistical Inference – Confidence Intervals and Tests of Hypotheses

Test of hypothesis – general method to distinguish between 2 (or more) probability distributions (or models), based on a sample X_1, \dots, X_n assumed to come from one of them.



In particular, based on X_1, \dots, X_n , decide whether $f_1(x)$ or $f_2(x)$ is the pdf (or population) from which the sample came.

More specifically, suppose we think the sample is from a normal population with mean either $\mu=5$ or $\mu=10$ with variance 4. (Or, more generally, $\mu=\mu_1$ vs. $\mu=\mu_2$).



To do this, we follow the “Neyman-Pearson Paradigm” and discuss “significance tests”.

Neyman-Pearson Approach:

Group the possible hypothesized distributions into two categories, the null hypothesis, H_0 , and the alternative hypothesis, H_A (research hypothesis).



E.g. Observe X_1, \dots, X_n .
Either $H_0: N(\mu_1, \sigma^2)$ or $H_A: N(\mu_2, \sigma^2)$
or $H_0: \mu = \mu_1$ vs. $H_A: \mu = \mu_2$,
 μ is the mean of $N(\mu, \sigma^2)$.

Here, if σ^2 is known, each of these hypotheses completely specifies the distribution or population. So, H_0 and H_A are called simple hypotheses.



If $H_0: \mu = 0$ vs. $H_A: \mu > 0$ (μ in $N(\mu, \sigma^2)$, with σ^2 known).

Then H_0 is simple and H_A is a composite hypothesis, i.e. several normal distributions would satisfy it.

H_A is also referred to as a one-sided hypothesis.



If $H_0: \mu=0$ vs. $H_A: \mu \neq 0$ (μ in $N(\mu, \sigma^2)$, σ^2 known), then H_A is a two-sided (composite) hypothesis.

Next, we set up the framework for “testing” H_0 and H_A based on the sample.



9.2 Neyman-Pearson Paradigm

Let $\underline{X} = (X_1, \dots, X_n)$ denote a sample from population $f(x|\theta)$.
Decide on H_0 vs. H_A based on the sample.

A decision on whether or not to reject H_0 in favor of H_A is made on the basis of a statistic

$$T=T(\underline{X})=T(X_1, \dots, X_n).$$



The set of values of T for which H_0 is accepted is called the acceptance region and the set of values of T for which H_0 is rejected is the rejection region of the test.



Two kinds of error may occur:

1. H_0 is rejected when it is true:
Type I error.

$$P(\text{type I error}) = \alpha \\ = P(T \in \text{rejection region} \mid H_0 \text{ true}).$$

If H_0 is simple, α is called the significance level of the test.



2. H_0 is accepted when it is false:
Type II error.

$$P(\text{type II error}) = \beta \\ = P(T \text{ in acceptance region} \mid H_0 \text{ false})$$

If H_A is composite, β depends on which member of H_A is the true pdf.



Power of the test = $P(H_0 \text{ is rejected when false})$
= $1 - P(H_0 \text{ is accepted} \mid H_0 \text{ false})$
= $1 - \beta$.

Ideally, we would want $\alpha = \beta = 0$, but this not possible since the decision is based on data.



Example 1: Consider a sample of size 1 from a normal distribution with variance 1.

Test $H_0: \mu=0$ vs. $H_A: \mu=1$ at $\alpha=0.05$.



Example 2: Consider a sample of size 1 from a normal distribution with variance 1.

Test $H_0: \mu \leq 0$ vs. $H_A: \mu > 0$ at $\alpha=0.05$.



For a composite test the significance level α is the maximum (supremum) of the probabilities of a Type I error over all the possible alternatives.


