STA 250/MTH 342 – Intro to Mathematical Statistics

Lecture 18

Testing using *p*-value

What is a *p-value*?

The p-value for a test

- ► The *p-value* for a test is a *statistic*, i.e. a function of the data **X** (and therefore is a random variable).
- It is defined to be the smallest α level at which observing **X** will lead to a rejection of the test. (Note the dependence on **X**.)
- In other words, it is the smallest α level for which the corresponding rejection region of the test covers X.

$$p(\mathbf{X}) = \inf\{\alpha : \mathbf{X} \in \mathcal{R}(\alpha)\}.$$

► (Draw a figure.)

- So for sensible tests, the smaller the *p*-value, the *stronger* the evidence *against* H_0 .
- ► For example, for the LR test, what is the meaning of the *p*-value? (Think about the gold miner analogy.)
- ► It is the minimal "budget" you need for considering purchasing the land that covers the observed data.
- ▶ Depending on the context, people use "*p-value*" to refer to either the *p-value statistic* or the observed value of the p-value statistic.

Lot testing example

- We observe X_1, X_2, \dots, X_n i.i.d. Exponential(λ).
- Consider testing

$$H_0: \lambda = 1.0$$
 vs $H_1: \lambda > 1.0$.

The UMP level- α test rejects when

$$\bar{X} < F^{-1}(\alpha)$$

where *F* is the cdf of Gamma $(n, n\lambda)$. So the rejection region is

$$\mathscr{R}(\alpha) = \{(x_1, x_2, \dots, x_n) : \bar{x} < F^{-1}(\alpha).\}$$

What is the *p*-value?

- ▶ If we observe $X_1, X_2, ..., X_n$, what is the *p*-value?
- ▶ By definition of the *p*-value, we know it is

$$p(\mathbf{X}) = \inf\{\alpha : (X_1, X_2, \dots, X_n) \in \mathcal{R}(\alpha)\}$$

$$= \inf\{\alpha : \bar{X} < F^{-1}(\alpha)\}$$

$$= \inf\{\alpha : \alpha > F(\bar{X})\}$$

$$= F(\bar{X}).$$

For example, if $\bar{X} = 0.7$ and n = 20, then

$$p(\mathbf{x}) = F(0.7) \approx 0.077.$$

A heuristic interpretation of the *p*-value

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- ► The *p* value is the "size" of the rejection (as measured by the Type I error) that "barely covers" *X*.
- ► This rejection region contains the data values that provide at least as much evidence in favor of the alternative as the data **X**.
- ▶ For generalized LR tests, this rejection region is the region that contains all data values that correspond to a Λ value no larger than the observed Λ value.
- ► The gold miner analogy.
- ▶ So the *p*-value can be intuitively thought of as the probability to get more "extreme" values of the data under the null.

▶ Back to the lot testing example, the *p*-value is given by

$$p(\mathbf{X}) = F(\bar{X}).$$

▶ This is indeed the chance of observing more "extreme" values under H_0 . (Draw a figure.)

Example: Finding the p-value of a two-sided *t*-test

- Suppose our data are i.i.d. from $N(\mu, \sigma^2)$ where both the mean μ and the variance σ^2 are unknown.
- Consider testing

$$H_0: \mu = \mu_0 \text{ vs } H_1: \mu \neq \mu_1$$

at level α .

We have seen that the level α LR test leads to the t test which rejects when

with
$$T = \frac{\sqrt{n}(\bar{X} - \mu_0)}{s}$$
, and $C = F_{t_{n-1}}^{-1}(1 - \frac{\alpha}{2})$.

In other words, the rejection region, which depends on the level α , is

$$\mathscr{R}(\alpha) = \left\{ (x_1, x_2, \dots, x_n) : \frac{\sqrt{n}|\bar{x} - \mu_0|}{s} > F_{t_{n-1}}^{-1} (1 - \frac{\alpha}{2}) \right\}.$$

► So the p-value is

$$\begin{split} p(\mathbf{X}) &= \inf \left\{ \alpha : \mathbf{X} \in \mathcal{R}(\alpha) \right\} \\ &= \inf \left\{ \alpha : |T| > F_{t_{n-1}}^{-1} \left(1 - \frac{\alpha}{2} \right) \right\} \\ &= \inf \left\{ \alpha : F_{t_{n-1}} (|T|) > 1 - \frac{\alpha}{2} \right\} \\ &= \inf \left\{ \alpha : \alpha > 2 \left(1 - F_{t_{n-1}} (|T|) \right) \right\} \\ &= 2 \left(1 - F_{t_{n-1}} (|T|) \right). \end{split}$$

(Draw a figure.)

► For example, if $\mu_0 = 1$, n = 20, $\bar{X} = 2.2$, $s^2 = 2.3$, then the *p*-value is

$$2\left(1 - F_{t_{n-1}}\left(\frac{\sqrt{10} \times |2.2 - 1|}{\sqrt{2.3}}\right)\right) = 0.0216.$$

▶ This is indeed the probability to observe a value "more extreme" than \mathbf{X} under H_0 . (Draw a figure.)

Testing using *p*-values

- The null is rejected with the test at level α, if and only if the p-value is no larger than α. Why? Let's consider both directions.
 - ▶ Note that $p(\mathbf{X}) = \inf\{\alpha : \mathbf{X} \in \mathcal{R}(\alpha)\}$ means that

$$p(\mathbf{X}) \leq \alpha_1 \Leftrightarrow \alpha_1 \in \{\alpha : \mathbf{X} \in \mathscr{R}(\alpha)\} \Leftrightarrow \mathbf{X} \in \mathscr{R}(\alpha_1).$$

Equivalently,

$$p(\mathbf{X}) > \alpha_1 \Leftrightarrow \alpha_1 \notin \{\alpha : \mathbf{X} \in \mathcal{R}(\alpha)\} \Leftrightarrow \mathbf{X} \notin \mathcal{R}(\alpha_1).$$

► The gold miner analogy.

The *p*-value as a measure of evidence against the null

- ► The *p*-value, however, contains more information than whether the test is rejected or not at a given level, say, $\alpha = .05$.
- ▶ It measures the strength of evidence against the null hypothesis.
- ► E.g., $p \approx .048$ compared to $p \approx .00001$. Both suggest that the null hypothesis is rejected at level α , but $p \approx .00001$ provides much stronger evidence against H_0 .
- So in the previous lot testing example and two-sample *t*-test example, do we reject H_0 at level 0.1, 0.05, 0.01?
- ► The *p*-value gives more information about the evidence from the data than just a reject/accept decision for a particular test level.

A problem with the *p*-value

- ► Note that it says nothing about how much the data supports the alternative in *absolute* terms.
- ▶ In the gold miner analogy: It is the total price for purchasing all the land that are more precious (i.e., with higher gold-to-dollar ratio) than a given point (the observed data), but we have no idea how much gold it contains, which may be very little!

Sampling distribution of the p-value under H_0

- ▶ We have seen that the *p*-value is a statistic, and thus it also has a sampling distribution.
- ▶ Question: If we repeat the experiment many many times, what is the distribution of the p-value that we observe under H_0 ?

- ▶ Let us try to find the cdf of this sampling distribution.
- ▶ For $\alpha \in [0,1]$, we have by the definition of the *p*-value

$$P(p(\mathbf{X}) \le \alpha | H_0) = P(\mathbf{X} \in \mathcal{R}(\alpha) | H_0).$$

- ▶ But $P(\mathbf{X} \in \mathcal{R}(\alpha)|H_0)$ is exactly the Type I error corresponding to the rejection region $R(\alpha)$, which by definition is α .
- Therefore

$$P(p(\mathbf{X}) \le \alpha | H_0) = \alpha$$
 for $\alpha \in [0, 1]$.

What distribution has this cdf?

p-value and the non-reproducibility of published science

- ▶ Under H_0 , $p(\mathbf{X})$ has a standard uniform distribution!
- ▶ So under the null, there is 5% chance that the *p*-value will be no more than 5%!
- ▶ Why are most published science nonreproducible?
- ▶ What does a *p*-value of 0.9999 mean?

The multiple testing problem

- ▶ If we test many null hypotheses, even if all of them are true, by chance we may have some small *p*-values.
- ▶ In such cases, the *p*-values lose their nominal meaning and must be "corrected".

$$P(\min\{U_1, U_2, \dots, U_T\} \le \alpha) = 1 - P(U_1 > \alpha) \times \dots \times P(U_T > \alpha)$$
$$= 1 - (1 - \alpha)^T,$$

which increases to 1 as T goes to infinity!