

By signing below I pledge that I have not communicated with anyone about this quiz other than the instructor.

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- 1(a). Let $Y_1, \dots, Y_n \sim \text{i.i.d. Poisson}(\theta)$, with density $p(y|\theta) = \theta^y e^{-\theta} / y!$. Write out the log likelihood, find the likelihood equation and the MLE. Show all of your steps.
- 1(b). Identify the asymptotic distribution of the MLE. Specifically, identify the asymptotic distribution of $\sqrt{n}(\hat{\theta}_{MLE} - \theta)$ and justify your result.
- 1(c). Suppose we are interested in estimating the probability that an observation from this population will be zero. In other words, we are interested in $g(\theta) = \Pr(Y = 0|\theta)$, where $Y \sim \text{Poisson}(\theta)$. Find the MLE of $g(\theta)$ based on a sample Y_1, \dots, Y_n , and also specify its asymptotic distribution.
- 1(d). Let X_i be the indicator that $Y_i = 0$, so that $X_i = 1$ if $Y_i = 0$ and $X_i = 0$ otherwise. Find a CAN estimator of $g(\theta)$ based on X_1, \dots, X_n and derive its asymptotic distribution. You may use results from class but explain what results you are using.
- 1(e). Compare the asymptotic variance of the estimator in part (c) to that of the estimator in part (d).

Solution

- 1(a). The log-likelihood is given by

$$\ell_n(\theta) = \sum_{i=1}^n [y_i \log \theta - \theta - \log(y_i!)] = \sum_{i=1}^n y_i \log \theta - n\theta - \sum_{i=1}^n \log(y_i!)$$

Taking the derivative of log-likelihood and setting it equal to 0, we have

$$\frac{\partial \ell_n}{\partial \theta} = \frac{\sum_{i=1}^n y_i}{\theta} - n = 0$$

Therefore, the MLE is $\hat{\theta}_{MLE} = \sum_{i=1}^n Y_i / n = \bar{Y}$.

- 1(b). Since $Y_i \stackrel{\text{iid}}{\sim} \text{Po}(\theta)$, we know that $E[Y_i] = \text{Var}(Y_i) = \theta$. Therefore, using CLT, we have

$$\sqrt{n}(\hat{\theta}_{MLE} - \theta) \xrightarrow{d} N(0, \theta)$$

- 1(c). We have $g(\theta) = \Pr(Y = 0 \mid \theta) = \theta^0 e^{-\theta} / 0! = e^{-\theta}$. Since the MLE for θ is \bar{Y} , the MLE of $g(\theta)$ is $g(\bar{Y}) = \exp(-\bar{Y})$. Since

$$\sqrt{n}(\hat{\theta}_{\text{MLE}} - \theta) \xrightarrow{d} \mathbf{N}(0, \theta)$$

Using Delta's method, we have

$$\sqrt{n} \left(g(\hat{\theta}_{\text{MLE}}) - g(\theta) \right) \xrightarrow{d} \mathbf{N} \left(0, |g'(\theta)|^2 \theta \right) \stackrel{d}{=} \mathbf{N} \left(0, e^{-2\theta} \theta \right)$$

- 1(d). Here $X \sim \text{Ber}(g(\theta))$. A natural choice of estimator of $g(\theta)$ is the sample mean $\widehat{g(\theta)} = \sum_{i=1}^n X_i = \bar{X}$. We know that $\mathbf{E}[X] = g(\theta)$, $\mathbf{Var}(X) = g(\theta)(1 - g(\theta))$. Due to CLT, we have

$$\sqrt{n} \left(\widehat{g(\theta)} - g(\theta) \right) \xrightarrow{d} \mathbf{N} \left(0, e^{-\theta}(1 - e^{-\theta}) \right)$$

Therefore, $\widehat{g(\theta)}$ is a CAN estimator.

- 1(e). It suffices to compare θ and $e^\theta - 1$. Clearly, using Taylor's expansion, we have $\theta < e^\theta - 1$ for any $\theta > 0$. Therefore, the asymptotic variance of $g(\hat{\theta}_{\text{MLE}})$ is smaller than that of $\widehat{g(\theta)}$.

For each $j = 1, \dots, m$, let p_j be a valid p -value for testing a null hypothesis H_j .

- 2(a). Suppose we reject hypothesis j when $p_j < \alpha_E$. In the case that the p -values are mutually independent, derive the value of α_E that makes the global type I error rate (the probability that one or more H_j 's are rejected assuming they are all true) exactly equal to a specified value α_G .
- 2(b). Find a value of α_F such that if we reject hypothesis j when $p_j < \alpha_F$, then the global error rate is less than or equal to α_G , even if the p -values are dependent, and prove your result.
- 2(c). Which threshold, α_E or α_F , is preferable in terms of power if the p -values are truly independent, and why?

Solution

- 2(a). Denote H_0 : All null hypotheses are true. Then the type I error rate is

$$\begin{aligned}\Pr(\text{Reject } H_0 \mid H_0) &= 1 - \Pr(\text{Accept all } H_j, j = 1, \dots, m \mid H_0) \\ &= 1 - (1 - \alpha_E)^m\end{aligned}$$

The second step is due to the facts that (1) all p -values are mutually independent and (2) each p -value follows a uniform distribution under the null hypothesis.

Let $1 - (1 - \alpha_E)^m = \alpha_G$, we have $\alpha_E = 1 - (1 - \alpha_G)^{1/m}$.

- 2(b). Let $\alpha_F = \alpha_G/m$ and we claim that this choice of α_F will lead to a global type I error rate which is less than or equal to α_G . A proof is as follows.

$$\begin{aligned}\Pr(\text{Reject } H_0 \mid H_0) &= \Pr(\text{Reject at least one } H_j, j = 1, \dots, m \mid H_0) \\ &= \Pr\left(\bigcup_{j=1}^m \{\text{Reject } H_j\} \mid H_0\right) \\ &\leq \sum_{j=1}^m \Pr(\text{Reject } H_j \mid H_0) \\ &= \sum_{j=1}^m \alpha_F = m\alpha_F = \alpha_G\end{aligned}$$

The final step is again due to the fact that each p -value follows a uniform distribution under the null hypothesis.

- 2(c). Conceptually, when p -values are truly independent, the threshold α_E is preferable as it is derived under the exactly same assumption and the global type I error rate is exactly α_G . Compared to α_E , α_F is derived under more relaxed assumption and hence will not be as sharp as α_E . Also, the global type I error rate is not exactly equal to α_G . It is more conservative.

Mathematically, let $f(x) = x^d - dx + d - 1$ where $x \in (0, 1)$. It is clear that $f(1) = 0$ for any $d \in (0, 1)$. Since $f'(x) = d(x^{d-1} - 1) < 0$ for any $x \in (0, 1)$ and any $d \in (0, 1)$, $f(x)$ is a monotone decreasing function on $(0, 1)$. Therefore,

$$\begin{aligned} & (1 - \alpha_G)^{1/m} - \frac{1 - \alpha_G}{m} + \frac{1}{m} - 1 < 0 \quad \text{for all } \alpha_G \in (0, 1) \text{ and } m > 1 \\ \implies & \frac{\alpha_G}{m} < 1 - (1 - \alpha_G)^{1/m} \quad \text{for all } \alpha_G \in (0, 1) \text{ and } m > 1 \\ \implies & \alpha_F < \alpha_E \quad \text{for all } \alpha_G \in (0, 1) \text{ and } m > 1 \end{aligned}$$

When $m = 1$, two thresholds are exactly same. When $m > 1$, α_F is smaller than α_E . Therefore, using α_E results in more rejections than using α_F . From the perspective of power comparison, using α_E leads to a more powerful test.

- 3(a). Let $Y_1, \dots, Y_n \sim \text{i.i.d. } P_\theta$ for some unknown $\theta \in \mathbb{R}^+$, where the density for P_θ is $p(y|\theta) = e^{-y/\theta}/\theta$. Write out the log likelihood $l(\theta)$ for θ based on observed values y_1, \dots, y_n , find the MLE $\hat{\theta}$, and write out the formula for $l(\hat{\theta})$, the maximized value of the log likelihood.
- 3(b). Suppose we have two independent observations Y_A and Y_B , one from each of two populations, so that $Y_A \sim P_{\theta_A}$ and $Y_B \sim P_{\theta_B}$. Write out the formula for $l(\theta_A, \theta_B)$, the log likelihood of (θ_A, θ_B) based on observed values y_A and y_B .
- 3(c). We wish to test the hypothesis $H_0 : \theta_A = \theta_B$ versus $H_1 : \theta_A \neq \theta_B$. Compute the minus two log likelihood ratio statistic for this test, which is $t(Y_A, Y_B) = -2 \times (l_0 - l_1)$, where l_1 is the maximum value of $l(\theta_A, \theta_B)$ and l_0 is the maximum subject to the restriction that $\theta_A = \theta_B$. (Hint: Use the result from (a) for some cases where $n = 1$ and $n = 2$.)
- 3(d). Write $t(Y_A, Y_B)$ in terms of the geometric mean $(Y_A Y_B)^{1/2}$ and the arithmetic mean $(Y_A + Y_B)/2$. Provide some intuition as to why this statistic makes sense for evaluating evidence against H_0 .
- 3(e). To actually implement a level- α test of H_0 with this test statistic, you would have to know, approximate, or simulate its null distribution, which is the distribution of $t(Y_A, Y_B)$ when θ_A and θ_B are equal to some common value, say θ . But how could you do this, if you don't know θ ? To solve this problem, show that the distribution of $t(Y_A, Y_B)$ under the null hypothesis doesn't depend on the particular value of θ . (Hint: You may use the fact that if $Y \sim P_\theta$ then $Y \stackrel{d}{=} \theta Z$, where $Z \sim P_1$.)

Solution

- 3(a). The log-likelihood is

$$\ell_n(\theta) = \sum_{i=1}^n (-y_i/\theta - \log \theta) = -\frac{\sum_{i=1}^n y_i}{\theta} - n \log \theta$$

Taking derivative of log-likelihood and setting it equal to 0, we have

$$\frac{\partial \ell_n}{\partial \theta} = \frac{\sum_{i=1}^n y_i}{\theta^2} - \frac{n}{\theta} = 0$$

Therefore, $\hat{\theta}_{\text{MLE}} = \sum_{i=1}^n Y_i/n = \bar{Y}$. The maximized value of the log-likelihood is

$$\ell_n(\hat{\theta}_{\text{MLE}}) = -\frac{n\bar{y}}{\bar{y}} - n \log \bar{y} = -n - n \log \bar{y}.$$

3(b). Since two observations are independent, the joint log-likelihood of θ_A and θ_B is

$$\ell(\theta_A, \theta_B) = -y_A/\theta_A - \log \theta_A - y_B/\theta_B - \log \theta_B$$

3(c). The test statistic is

$$\begin{aligned} t(Y_A, Y_B) &= -2 \log \frac{\max_{\theta} L_0(\theta)}{\max_{\theta_A, \theta_B} L_1(\theta_A, \theta_B)} \\ &= -2 \left(\max_{\theta} \ell_0(\theta) - \max_{\theta_A, \theta_B} \ell_1(\theta_A, \theta_B) \right) \\ &= -2 \left(-2 - 2 \log \frac{y_A + y_B}{2} - (-1 - \log y_A - 1 - \log y_B) \right) \quad (3) \\ &= 4 \left(\log \frac{y_A + y_B}{2} - \log (y_A y_B)^{1/2} \right) \end{aligned}$$

Equation (3) is derived using the result in part (a) with $n = 1$ or 2 .

3(d). Let $M_A = (Y_A + Y_B)/2$ and $M_G = (Y_A Y_B)^{1/2}$. Then

$$t(Y_A, Y_B) = 4 \log \frac{M_A}{M_G}$$

We know that $M_A \geq M_G$ and the equality holds if and only if $Y_A = Y_B$. Let $Y_A/Y_B = m$, then

$$t(Y_A, Y_B) = 4 \log \frac{(m+1)/2}{\sqrt{m}}$$

When the null hypothesis is true, $\theta_A = \theta_B$, we expect to observe that Y_A and Y_B are similar and hence $m \approx 1$. In this case $(m+1)/2 \approx \sqrt{m}$, the value of test statistic is small. However, when the null hypothesis is wrong, m could

be very small or large. In either case, the ratio between $(m+1)/2$ and \sqrt{m} is large and hence the value of test statistic is large.

Therefore, the behavior of the test statistic makes intuitive sense. It evaluates the evidence against the null hypothesis.

- 3(e). By change of variable, it is easy to show that if $Y \sim P_\theta$, then $Y \stackrel{d}{=} \theta Z$ where $Z \sim P_1$. Under the null hypothesis, we have $\theta_A = \theta_B = \theta$ and hence $Y_A \stackrel{d}{=} \theta Z_A$ and $Y_B \stackrel{d}{=} \theta Z_B$ where $Z_A, Z_B \stackrel{\text{iid}}{\sim} P_1$.

Therefore, the test statistic can be rewritten as

$$\begin{aligned} t(Y_A, Y_B) &= 4 \left(\log \frac{y_A + y_B}{2} - \log (y_A y_B)^{1/2} \right) \\ &= 4 \left(\log \frac{\theta Z_A + \theta Z_B}{2} - \log (\theta Z_A \theta Z_B)^{1/2} \right) \\ &= 4 \left(\log \frac{Z_A + Z_B}{2} - \log (Z_A Z_B)^{1/2} \right) \end{aligned}$$

Since $Z_A, Z_B \stackrel{\text{iid}}{\sim} P_1$, the null distribution does not depend on actual value of θ .