Solution of BIOSTAT 802 HW3

Solution of Problem 1:

Let $\Theta_H = \{P_0\}$ and $\Theta_K = \{P_1, P_2\}$. The hypothesis of interest is

$$H: \Theta_H \text{ versus } K: \Theta_K.$$

To begin we calculate likelihood ratios of $P_i(x)/P_0(x)$, i = 1, 2, that are listed in the following table:

From this table, it is clear that x = 5 leads to the action of rejection as the corresponding LRs is ∞ .

(i) Consider the case of $\alpha = 0.01$. According to the **Neyman-Pearson Lemma**, we can construct an MP test $\phi_1(x)$ for $H': P_0$ versus $K': P_1$. That is, there exists a constant k_1 such that

$$\phi_1(x) = \begin{cases} 1, & P_1(x)/P_0(x) > k_1; \\ \gamma_1(x), & P_1(x)/P_0(x) = k_1; \\ 0, & P_1(x)/P_0(x) < k_1, \end{cases}$$

where the cutoff k_1 is so chosen that its size $E_0\phi_1(X) = 0.01$. Here we choose $k_1 = 4$, and must take $\gamma_1(x) = 1/2$ (at x = 3), so x = 3 leads to the 50% chance of rejection! In this case, the size (or the probability of type I error) is, under P_0 , 0.00 + (1/2) (0.02) = 0.01, as desired. In summary, the critical function of the MP test is given by

$$\phi_1(x) = \phi_2(x) = 1(x=5) + (1/2)1(x=3),$$

where 1(.) is the indicator function.

Following the above steps, similarly we can construct an MP test for $H'': P_0$ against $K'': P_2$. The N-P Lemma ensures that there exists a constant k_2 such that the test $\phi_2(x)$ takes the following form:

$$\phi_2(x) = \begin{cases} 1, & P_2(x)/P_0(x) > k_2; \\ \gamma_2(x), & P_2(x)/P_0(x) = k_2; \\ 0, & P_2(x)/P_0(x) < k_2, \end{cases}$$

where the cutoff k_2 is so chosen that its size $E_0\phi_2(X) = 0.01$. Here we choose $k_2 = 6$, and must take $\gamma_2(x) = 1/2$ (at x = 3), so x = 3 leads to the 50% chance of rejection! In this case, the size (or the probability of type I error) is 0.00 + (1/2) (0.02) = 0.01, as desired. In summary, the critical function of the MP test $\phi_2(x)$ is given by

$$\phi_2(x) = 1(x=5) + (1/2)1(x=3).$$

Although k_1 and k_2 are different, the rejection rule is the same $\phi_1(x) = \phi_2(x)$ on any data x. In other words, the rejection rule is uniform over the sample space regardless of alternative P_1 or P_2 . Thus, the UMP test of P_0 against P_1 and P_2 is of the form

$$\phi(x) = 1(x = 5) + (1/2)1(x = 3).$$

(ii) Similarly, for $\alpha = 0.05$, the most powerful tests of P_0 against P_1 and P_2 take the following forms, respectively,

$$\phi_1(x) = 1(x \in \{2, 3, 5\}) + \gamma_1(x)1(x \in \{1, 4\}),$$

$$\phi_2(x) = 1(x \in \{2, 3, 5\}),$$

where $\gamma_1(x) \neq 0$ in order to satisfy the size condition $E_0\phi_1(X) = 0.05$. Because $\phi_1(x) \neq \phi_2(x)$ on the sample space, these two MP tests cannot be unified, and thus there is no UMP test for the hypothesis H versus K.

(iii) For $\alpha = 0.07$, the most powerful tests of P_0 against P_1 and P_2 take the following forms, respectively,

$$\phi_1(x) = 1(x \in \{2, 3, 5\}) + \gamma(x)1(x \in \{1, 4\}),$$

$$\phi_2(x) = 1(x \in \{1, 2, 3, 5\}),$$

where $\gamma(x) = 1(x = 1)$. Although $\phi_1(x)$ and $\phi_2(x)$ have different analytic forms, they are effectively the same critical function defining the same rejection region. Thus, there is a UMP test for the hypothesis H versus K, that is, $\phi(x) = 1(x \in \{1, 2, 3, 5\})$.

Solution of Problem 2

(h) let To=T and T1 = 1-TT. The above prob. of error may be expressed as follows.

$$Prob(enin) = \pi, \int_{x} (1 - \psi(x)) dP_{1}(x) + \pi_{0} \int_{x} \psi(x) dP_{0}(x)$$

$$= \pi_{1} - \pi_{1} \int_{x} \psi(x) P_{1}(x) dx + \pi_{0} \int_{x} \psi(x) P_{0}(x) dx$$

$$= \pi_{1} + \int_{x} \psi(x) \{ \pi_{0} P_{0}(x) - \pi_{1} P_{1}(x) \} dx,$$

where Ti, is a constant independent data & and decision rule U(1).

Set
$$\mathfrak{X}_1 = \left\{ \chi: \Pi_0 \mathfrak{P}_0(x) - \Pi_1 \mathfrak{P}_1(x) > 0 \right\} = \left\{ \chi: \frac{\Pi_0}{\Pi_1} \mathfrak{P}_0(x) > \mathfrak{P}_1(x) \right\}$$

$$\mathfrak{X}_2 = \left\{ \chi: \Pi_0 \mathfrak{P}_0(x) - \Pi_1 \mathfrak{P}_1(x) = 0 \right\} = \left\{ \chi: \mathfrak{P}_1 = \frac{\Pi_0}{\Pi_1} \mathfrak{P}_0(x) \right\}$$

$$\mathfrak{X}_3 = \left\{ \chi: \Pi_0 \mathfrak{P}_0(x) - \Pi_1 \mathfrak{P}_1(x) < 0 \right\} = \left\{ \chi: \mathfrak{P}_1 \nearrow \frac{\Pi_0}{\Pi_1} \mathfrak{P}_0(x) \right\},$$

and X = X, U.X, U Xz, and they are disjoint sets. Since O = \(\partial (x) \le 1,

$$\psi(x) = \underset{0 \le \psi \le 1}{\operatorname{arg min}} \int \psi(x) \left\{ \pi_{\sigma} \, \mathcal{P}_{\sigma}(x) - \pi_{\sigma} \, \mathcal{P}_{\sigma}(x) \right\} dx = \left\{ \begin{array}{c} 1, & x \in \mathcal{X}_{3}, \\ 0, & x \in \mathcal{X}_{1}, \end{array} \right.$$

wanch, $\psi(x) = \begin{cases} 1, & \text{if } \gamma_i(x) \nearrow \frac{\pi_0}{\pi_i} p_i(x) \nearrow k p_0(x), & \text{if } \frac{\pi_0}{\pi_i} p_0(x) \nearrow k p_0(x), \end{cases}$

(c)
$$p(H_0|X=x) = \frac{p(X=x|H_0) p(H_0)}{p(X=x)} = \frac{p_0(X) \pi_0}{\pi_0 p_0(X) + \pi_1 p_1(X)}$$
, by the Bayes formula.

(i. helvise $p(H_1|X=x) = \frac{\gamma_1(x) \pi_1}{\pi_0 p_0(x) + \pi_1 p_1(x)}$

Give. X=x, when if p(x) > To p(x) <=> p(H) | X=x) > p(H) | X=x) while accept Ho of p(x) < To po(x) <=> p(H) | X=x) > p(H) | X=x).

This means that this Baye let under decision accordis to lenger posterior protein protein. Since T(x) is a sufficient statistic, the conditional distribution of X/T is independent of parameter $\theta \in \Theta$. That is, $\phi(t) = E[\psi(x)/t]$ is parameter θ free. $A(so, 0 \le \psi(x) \le 1$, so,

 $0 \le \phi(t) \le 1$.

Therefore, $\phi(t)$ defines a critical function, or a test. Now, look at the power function of $\phi(t)$, because of the property of conditional expectation

 $E_{\theta} \Phi(T) = E_{\theta} \{ E[\Psi(x) | T] \} = E_{\theta} \Psi(x),$ i.e. the two tests have the name power function.