

1. A study conducted in the Maternal and Child Health Department asks each of n randomly selected mothers of infants less than 6 months of age a set of k “yes” or “no” questions. Assuming the probability of answering “yes” in each question is denoted by π , the researcher is interested in estimating the probability of “perfect care”, which is defined by answering “yes” in all of the k questions, i.e., $\theta = \pi^k$.

- (a) Let X_i denote the number of questions mother i answering “yes” among those k questions. One can assume that X_i follows a binomial distribution with pdf

$$f(x|\pi) = \binom{k}{x} \pi^x (1 - \pi)^{k-x}.$$

Assuming X_1, \dots, X_n is a random sample of size n , derive the maximum likelihood estimator (MLE) of θ .

- (b) Derive Cramér-Rao Lower Bound (CRLB) of the variance of any unbiased estimator of θ .
- (c) Show that $T = \sum_{i=1}^n X_i$ is a complete and sufficient statistic for π (as well as for θ), and that

$$W = \begin{cases} 1 & \text{if } X_1 = k \\ 0 & \text{otherwise,} \end{cases}$$

is an unbiased estimator of θ .

- (d) By Lehmann-Sheffe Theorem, the estimator $\phi(T) = E(W|T)$ is the uniformly minimum variance unbiased estimator (UMVUE). Show that

$$\phi\left(\sum_{i=1}^n X_i\right) = \frac{\binom{(n-1)k}{\sum_{i=1}^n X_i - k}}{\binom{nk}{\sum_{i=1}^n X_i}}$$

if $\sum_{i=1}^n X_i = k, k+1, \dots, nk$, and $\phi(\sum_{i=1}^n X_i) = 0$ otherwise. Here you may use the fact that $\sum_{i=1}^n X_i$ follows a binomial distribution with parameters nk and π , denoted by $\text{Bin}(nk, \pi)$.

- (e) Without using $E(E(W|T)) = E(W)$, show that $\phi(\sum_{i=1}^n X_i)$ is indeed an unbiased estimator of θ .
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- (f) One may see that the variance of $\phi(\sum_{i=1}^n X_i)$ is quite complicated and difficult to derive, which makes the estimator less useful because one cannot make any inference on θ (e.g., does a mother with private insurance more likely have a “perfect care”?) A biostatistician suggests using the MLE in (a) to make inference on θ since the large sample property of the estimator can be easily derived. Comment on the approach of the biostatistician and derive the large sample distribution of the MLE of θ in (a). [Hint: Use Central Limit Theorem (CLT) and Delta method].
2. In malaria study, human-to-mosquito transmission is mediated by sexual stage parasites called gametocytes. *P. falciparum* gametocytes often represent only about 1% of the total parasite load in blood, which makes the detection of the gametocytes difficult. Suppose the amount of the gametocytes detected by a current technology, e.g., Pfs25 RT-PCR, distribute like a location-shifted exponential distribution with pdf

$$f(x|\theta) = \frac{1}{\beta} \exp\left(-\frac{x-\theta}{\beta}\right),$$

where $x > \theta$ and $\theta > 0$ is so-called detection limit. Assuming β is *known*, a researcher is eager to know how low is the amount of gametocytes the current technology can detect.

- (a) If the research collects a random sample X_1, \dots, X_n of size n from an essay, derive the maximum likelihood estimator $\hat{\theta}$ of θ .
- (b) To test a null hypothesis $H_0 : \theta \leq \theta_0$ versus the alternative hypothesis $H_0 : \theta \geq \theta_0$, one can derive a likelihood ratio test (LRT) statistic $\lambda(\mathbf{x})$ and claim that the null hypothesis is rejected if $\lambda(\mathbf{x}) \leq c$ for some cutoff c . Derive the LRT statistic $\lambda(\mathbf{x})$.
- (c) One may find the cutoff c is difficult to find since one may not know the distribution of $\lambda(\mathbf{X})$. Instead, we can find an equivalent rejection region using the MLE in (a). Show that the equivalent region is $\{\mathbf{x}; \hat{\theta} \geq c^*\}$.
- (d) Find the cutoff c^* such that the test has a size α , i.e., $\alpha = \sup_{\theta \in \Theta_0} P(\hat{\theta} \geq c^*)$.
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