

# BIOSTAT/STAT 570: Midterm Take-Home Exam

To be submitted to the course canvas site by 11:59pm Saturday 12th November, 2022. This is an exam, so no collaboration.

In this exam you will investigate different approaches to analyzing a simple prevalence estimation problem, that we looked at briefly in class.

In early April, 2020, 3330 residents of Santa Clara County, California were recruited then and tested for COVID-19 antibodies and 50 people tested positive (Bendavid et al. 2020, MedRxiv).

Suppose a seroprevalence test is carried out with sensitivity  $\delta = \Pr(\text{+ve test} \mid \text{disease})$  and specificity,  $\gamma = \Pr(\text{-ve test} \mid \text{no disease})$ , where  $0 < \gamma \leq 1$ ,  $0 < \delta \leq 1$  and we assume  $\gamma + \delta > 1$ . Let  $\pi$  denote the true prevalence.

We test  $N$  people and  $y$  are recorded as having the disease, and a starting model is

$$y|p \sim \text{Binomial}(N, p)$$

where  $p$  is the probability of a +ve test result, with

$$\begin{aligned} p &= \Pr(\text{+ve test}) \\ &= \Pr(\text{+ve test} \mid \text{disease}) \Pr(\text{disease}) + \Pr(\text{+ve test} \mid \text{no disease}) \Pr(\text{no disease}) \\ &= \pi(\delta + \gamma - 1) + (1 - \gamma) \end{aligned}$$

Suppose, initially, that the sensitivity and specificity are known and we want to estimate  $\pi$ . Assume the sensitivity is 0.8 and the specificity is 0.995.

## 1. Likelihood Analysis

(a) **4 Points** Show that under the binomial model the MLE is

$$\hat{\pi} = \frac{y - N(1 - \gamma)}{N(\delta + \gamma - 1)}.$$

(b) **4 Points** Give an expression for the expected information and hence give the form for a 90% Wald confidence interval.

(c) **4 Points** Obtain the MLE and 90% confidence interval for  $\pi$  the Santa Clara data.

(d) **2 Points** Obtain the MLE and the expected information for  $\theta = \log(\pi/(1 - \pi))$ .

- (e) **2 Points** Suppose one believes the data are overdispersed relative to a binomial. Is it possible to fit an overdispersed model for the data here?

## 2. Bayesian Analysis

We will now carry out a Bayesian analysis with a  $\text{Beta}(1,1)$  prior for  $\pi$ . For the methods listed below, evaluate the normalizing constant, the posterior mean and the posterior variance for  $\theta = \log(\pi/(1 - \pi))$ . Give your results in the form of a table. Also give Monte Carlo confidence intervals on your estimates, for those methods that you can do this for.

- (a) **4 Points** Write down the functional form for the posterior  $p(\theta|y)$ , up to the normalizing constant.
- (b) **4 Points** Gauss-Hermite quadrature using 4 point and 5 point rules. You should decide how to center and scale the integration design points.
- (c) **4 Points** A rejection algorithm, using the prior as proposal. In addition to the normalizing constant, the posterior mean and the posterior variance, give a histogram representation of the posterior distribution for  $\pi$ .
- (d) **4 Points** Importance sampling Monte Carlo. Use a t-distribution with 4 degrees of freedom, with mean and scale based on the MLE and its asymptotic variance.
- (e) **4 Points** A Metropolis algorithm based on a normal proposal distribution centered at the current point with a variance you should choose. In addition to the posterior mean and the posterior variance, give a histogram representation of the posterior distribution for  $\pi$  (For this part, you do not need to evaluate the normalizing constant).
- (f) **6 Points** Suppose that rather than being taken as fixed, beliefs about the sensitivity  $\delta$  and the specificity  $\gamma$  can be represented via beta distributions. Specifically,  $\delta \sim \text{Beta}(160, 40)$  and  $\gamma \sim \text{Beta}(1990, 10)$ . Extend your approach in (b) or (d) to obtain samples from the posterior for  $\pi$ , based on these new beliefs about  $\delta$  and  $\gamma$ . How does the posterior change relative to the point estimates for  $\delta, \gamma$  analysis?
- (g) **BONUS 4 Points** Laplace approximation.

Hint: Section 3.7 of Wakefield (2013) contains many of the details you will need for this work, and the book website contains R code to carry out similar analyses.