## BIOSTAT/STAT 570: Coursework 5

To be submitted to the course canvas site by 11:59pm Monday 4th November, 2022.

1. Consider the data given in Table 1, which are a simplified version of those reported in Breslow and Day (1980). These data arose from a case-control study that was carried out to investigate the relationship between esophageal cancer and various risk factors. Disease status is denoted Y with Y=0/1 corresponding to without/with disease and alcohol consumption is represented by X with X=0/1 denoting  $<80g/\geq80g$  on average per day. Let the probabilities of high alcohol consumption in the cases and controls be denoted

$$p_1 = \Pr(X = 1 \mid Y = 1)$$
 and  $p_2 = \Pr(X = 1 \mid Y = 0)$ ,

respectively. Further, let  $X_1$  be the number exposed from  $n_1$  cases and  $X_2$  be the number exposed from  $n_2$  controls. Suppose  $X_i \mid p_i \sim \mathsf{Binomial}(n_i, p_i)$  in the case (i = 1) and control (i = 2) groups.

$$egin{array}{c|cccc} X = 0 & X = 1 \\ \hline Y = 1 & 104 & 96 & 200 \\ Y = 0 & 666 & 109 & 775 \\ \hline \end{array}$$

Table 1: Case-control data: Y=1 corresponds to the event of esophageal cancer, and X=1 exposure to greater than 80g of alcohol per day. There are 200 cases and 775 controls.

(a) Of particular interest in studies such as this is the odds ratio defined by

$$\theta = \frac{\Pr(Y = 1 \mid X = 1) / \Pr(Y = 0 \mid X = 1)}{\Pr(Y = 1 \mid X = 0) / \Pr(Y = 0 \mid X = 0)}.$$

Show that the odds ratio is equal to

$$\theta = \frac{\Pr(X=1 \mid Y=1) / \Pr(X=0 \mid Y=1)}{\Pr(X=1 \mid Y=0) / \Pr(X=0 \mid Y=0)} = \frac{p_1 / (1-p_1)}{p_2 / (1-p_2)}.$$

- (b) Obtain the MLE and an asymptotic 90% confidence interval for  $\theta$ , for the data of Table 1.
- (c) We now consider a Bayesian analysis. Assume that the prior distribution for  $p_i$  is the beta distribution  $\operatorname{Be}(a,b)$  for i=1,2. Show that the posterior distribution  $\pi(p_1,p_2\mid x_1,x_2)$  is given by the product of the beta distributions  $\operatorname{Be}(a+x_i,b+n_i-x_i)$ , i=1,2.

- (d) Consider the case a=b=1. Obtain expressions for the posterior mean, mode and standard deviation. Evaluate these posterior summaries for the data of Table 1. Report 90% posterior credible intervals for  $p_1$  and  $p_2$ .
- (e) Examine the implied prior distribution for  $\theta$  and give a 90% prior interval.
- (f) Simulate samples  $p_1^{(t)}, p_2^{(t)}, t = 1, \dots, T = 1000$  from the posterior distributions  $p_1 \mid x_1$  and  $p_2 \mid x_2$ . Form histogram representations of the posterior distributions using these samples and obtain sample-based 90% credible intervals.
- (g) Obtain samples from the posterior distribution of  $\theta \mid x_1, x_2$  and form the histogram representation of the posterior. Obtain the posterior median and 90% credible interval for  $\theta \mid x_1, x_2$  and compare with the likelihood analysis.
- (h) Suppose the rate of esophageal cancer is 18 in 100,000. Describe how this information may be used to evaluate

$$q_1 = \Pr(Y = 1 \mid X = 1)$$
 and  $q_0 = \Pr(Y = 1 \mid X = 0)$ .

- (i) Suppose that *a priori* you would like to select a Be(a,b) distribution on the rate of esophageal cancer with 5% of the mass less than 16 in 100,000 and 5% of the mass greater than 20 in 100,000. Find a and b to satisfy these requirements, and hence obtain samples from the posteriors for  $g_1$  and  $g_0$ .
- 2. (a) Consider the "likelihood",  $\widehat{\theta} \mid \theta \sim \mathsf{N}(\theta, V)$  and the prior  $\theta \sim \mathsf{N}(0, W)$  with V and W known. Show that  $\theta \mid \widehat{\theta} \sim \mathsf{N}(r\widehat{\theta}, rV)$  where r = W/(V + W).
  - (b) Suppose we wish to compare the models  $M_0: \theta = 0$  versus  $M_1: \theta \neq 0$ . Show that the Bayes factor is given by

$$\mathsf{BF} = \frac{p(\widehat{\theta}|M_0)}{p(\widehat{\theta}|M_1)} = \frac{1}{\sqrt{1-r}} \exp\left(-\frac{Z^2}{2}r\right)$$

where  $Z = \widehat{\theta}/\sqrt{V}$ .

- (c) Suppose we have a prior probability  $\pi_1 = \Pr(M_1)$  of model  $M_1$  being true. Write down an expression for the posterior probability  $\Pr(M_1|\widehat{\theta}_1)$ , in terms of the BF.
- (d) Now suppose we have summaries from two studies,  $\widehat{\theta}_j, V_j, j = 1, 2$ . Assuming,  $\widehat{\theta}_j \mid \theta \sim \mathsf{N}(\theta, V_j)$  and the prior  $\theta \sim \mathsf{N}(0, W)$ , derive the posterior  $p(\theta | \widehat{\theta}_1, \widehat{\theta}_2)$ .
- (e) Derive the Bayes factor

$$\mathsf{BF} = \frac{p(\widehat{\theta}_1, \widehat{\theta}_2 | M_0)}{p(\widehat{\theta}_1, \widehat{\theta}_2 | M_1)}$$

again comparing the models  $M_0: \theta = 0$  versus  $M_1: \theta \neq 0$ .

We will show these results can be used in the context of a genome-wide association study on Type II diabetes, reported bu Frayling et al. (2007, Science). Two sets of data were independently collected, resulting in two log odds ratios  $\hat{\theta}_j$ , j=1,2, for each SNP. For SNP rs9939609 point estimates of the odds ratio (95% confidence intervals) were 1.27 (1.16, 1.37) and 1.15 (1.09,1.23). Suppose we have a normal prior for the log odds ratio that has a 95% range [log(2/3), log(3/2)].

- (f) Find W from this interval, and then calculate the posterior median and 95% intervals for  $\theta$  based on (i) the first dataset only, (ii) both of the populations.
- (g) Calculate the Bayes factor based on the first dataset only, and then based on both datasets.
- (h) With a prior of  $\pi_1 = 1/5000$ , calculate the probabilities,  $\Pr(M_1|\widehat{\theta}_1)$  and  $\Pr(M_1|\widehat{\theta}_1,\widehat{\theta}_2)$
- 3. We will carry out a Bayesian analysis of the lung cancer and radon data, that were examined in lectures, using INLA. These data are available on the class website.

The likelihood is

$$Y_i \mid \beta \sim_{ind} \mathsf{Poisson} \left[ E_i \exp(\beta_0 + \beta_1 x_i) \right],$$

where  $\beta = [\beta_0, \beta_1]^{\mathsf{T}}$ ,  $Y_i$  and  $E_i$  are observed and expected counts of lung cancer incidence in Minnesota in 1998–2002, and  $x_i$  is a measure of residential radon in county i,  $i = 1, \ldots, n$ .

- (a) Analyze these data using the default prior specifications in INLA. Produce figures of the INLA approximations to the marginal distributions of  $\beta_0$  and  $\beta_1$ , along with the posterior means, posterior standard deviations, and 2.5%, 50%, 97.5% quantiles.
- (b) For a more informative prior specification we may reparameterize the model as

$$Y_i \mid \boldsymbol{\theta} \sim_{ind} \mathsf{Poisson}\left(E_i \theta_0 \theta_1^{x_i - \overline{x}}\right),$$

where  $\boldsymbol{\theta} = [\theta_0, \theta_1]^{\mathsf{T}}$  where

$$\theta_0 = \mathsf{E}[Y/E \mid x = \overline{x}] = \exp(\beta_0 + \beta_1 \overline{x})$$

is the expected standardized mortality ratio in an area with average radon. The parameter  $\theta_1 = \exp(\beta_1)$  is the relative risk associated with a one-unit increase in radon.

For  $\theta_0$  we assume a lognormal prior with 2.5% and 97.5% quantiles of 0.67 and 1.5 to give  $\mu=0,\sigma=0.21$ . For  $\theta_1$  we again take a lognormal prior and assume the relative risk associated with a one-unit increase in radon is between 0.8 and 1.2 with probability 0.95, to give  $\mu=-0.02,\sigma=0.10$ . By converting these into normal priors in INLA, rerun your analysis, and report the same summaries.