STAT 457 Homework 05

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Problem 1

Consider two urns each containing an unknown mixture of blue and white marbles. A random sample of size 18 (with replacement) is drawn from urn #1 and a random sample of size 6 (with replacement) is drawn from urn #2. Of the 18 selected marbles from urn #1, 14 are blue. The corresponding number of blue marbles from urn #2 is 2.

$$L(\pi|Y) \propto \pi^{y} (1-\pi)^{n-y} = \pi^{14} (1-\pi)^{4} \sim \text{Binomial}, n_{\pi} = 18, \ y_{\pi} = 14$$

$$\implies p(\pi \mid Y) \sim \text{Beta}(y + \alpha_{0}, n - y + \beta_{0}) \quad \text{where} \quad p(\pi) \sim \text{Beta}(\alpha_{0}, \beta_{0})$$

$$L(\psi|Y) \propto \psi^{y} (1-\psi)^{n-y} = \psi^{2} (1-\psi)^{4} \sim \text{Binomial}, n_{\pi} = 6, \ y_{\pi} = 2$$

$$\implies p(\psi \mid Y) \sim \text{Beta}(y + \alpha_{0}, n - y + \beta_{0}) \quad \text{where} \quad p(\psi) \sim \text{Beta}(\alpha_{0}, \beta_{0})$$

Problem 1a

Let π denote the proportion of blue marbles in urn #1 and let ψ denote the corresponding proportion in urn #2. Under the (i) Haldane, (ii) flat and (iii) non-informative priors, compute $p\left(\ln\left[\frac{\pi}{1-\pi}\right] > \ln\left[\frac{\psi}{1-\psi}\right] \mid \text{data}\right)$ using the normal approximation.

$$p\left(\ln\left[\frac{\pi}{1-\pi}\right] > \ln\left[\frac{\psi}{1-\psi}\right] \mid \text{data}\right) = p\left(\ln\left[\frac{\pi}{1-\pi}\right] - \ln\left[\frac{\psi}{1-\psi}\right] > 0 \mid \text{data}\right)$$

Normal Approx: Probability difference in logodds is greater than 0

| Prior | Haldane - Beta $(0, 0)$ | Flat - $Beta(1, 1)$ | Non-informative - $Beta(.5, .5)$ |
|---------|-------------------------|---------------------|----------------------------------|
| p-value | 0.96533 | 0.97788 | 0.97285 |

Problem 1b

Repeat (1a) by drawing deviates from the appropriate beta distributions. Quantify the Monte Carlo error in your value.

Haldane Prior - Beta(0, 0)

| Iterations | 10000 | 1e + 05 | 1e + 06 |
|----------------|---------|---------|---------|
| p-value | 0.90144 | 0.90114 | 0.90140 |
| Standard Error | 0.00901 | 0.00284 | 0.00090 |

Flat Prior - Beta(1, 1)

| Iterations | 10000 | 1e+05 | 1e+06 |
|----------------|---------|---------|---------|
| p-value | 0.90339 | 0.90072 | 0.90116 |
| Standard Error | 0.00883 | 0.00284 | 0.00090 |

Non-informative Prior - Beta(.5, .5)

| Iterations | 10000 | 1e+05 | 1e+06 |
|------------|---------|---------|---------|
| p-value | 0.89969 | 0.90234 | 0.90110 |

Non-informative Prior - Beta(.5, .5)

| Standard Error | 0.00887 | 0.00284 | 0.00090 |
|----------------|---------|---------|---------|
| Standard Error | 0.00007 | 0.00204 | 0.00090 |

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Problem 1c

Compare your results in (1a) and (1b) to the p-value obtained via Fisher's exact test.

Problem 1d

Add delinquency problem

Problem 2

Suppose a sample of size n is drawn at random and with replacement from some population. For large n the sample proportion (\hat{p}) is normally distributed with mean p and variance $\frac{p(1-p)}{n}$. Find the asymptotic distribution of $2\sin^{-1}\sqrt{\hat{p}}$ using the delta method.

Problem 3

Let x_1, \dots, x_n be an iid sample from $\mathcal{N}(\theta, 1)$ and let y_1, \dots, y_n be an independent iid sample from $\mathcal{N}(\phi, 1)$. Derive the distribution of $\overline{x}/\overline{y}$ (where $\overline{y} \neq 0$) via the delta method.

Problem 4

197 animals are distributed into four categories: $Y = (y_1, y_2, y_3, y_4)$ according to the genetic linkage model $(\frac{2+\theta}{4}, \frac{1-\theta}{4}, \frac{1-\theta}{4}, \frac{1}{4}, \frac{1}{4}, \frac{1}{4}, \frac{1}{4})$. In HW#4 you derived the likelihood for the data Y = (125, 18, 20, 34) and you derived the likelihood for the data Y = (14, 0, 1, 15). In that homework, you also used Newton-Raphson algorithm to obtain the MLE $(\hat{\theta})$ of θ and the standard error of $\hat{\theta}$.

Problem 4a

Plot the normalized likelihood and the associated normal approximation in the same figure for the data Y = (125, 18, 20, 34). Discuss the adequacy of the normal approximation.

Problem 4b

Repeat (4a) for Y = (14, 0, 1, 5)

Problem 5

Use Laplace's method (second order) to compute the posterior mean (under a flat prior) for the genetic linkage model for both data sets.