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.

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

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

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

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

$$\frac{\text{Blue White}}{\text{Urn 1}} \frac{14}{2} \frac{4}{4}$$

$$\text{Urn 2} \frac{14}{2} \frac{4}{4}$$

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.

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

$$p(\pi) \sim \text{Beta}(\alpha_0, \beta_0)$$

$$p(\psi) \sim \text{Beta}(y_\pi, \beta_0)$$

$$p(\pi \mid Y) \sim \text{Beta}(y_\pi + \alpha_0, n_\pi - y_\pi + \beta_0) \stackrel{\text{def}}{=} \text{Beta}(\alpha, \beta)$$

$$p(\pi \mid Y) \sim \text{Beta}(y_\pi + \alpha_0, n_\pi - y_\pi + \beta_0) \stackrel{\text{def}}{=} \text{Beta}(\gamma, \delta)$$

$$\text{Normal Approx Mean} = \ln\left(\frac{\alpha \cdot \delta}{\beta \cdot \gamma}\right)$$

$$\text{Normal Approx Variance} = \frac{1}{\alpha} + \frac{1}{\beta} + \frac{1}{\gamma} + \frac{1}{\delta}$$

$$\text{Normal Approx} \sim \mathcal{N}\left(\ln\left(\frac{\alpha \cdot \delta}{\beta \cdot \gamma}\right), \sqrt{\frac{1}{\alpha} + \frac{1}{\beta} + \frac{1}{\gamma} + \frac{1}{\delta}}\right)$$

Normal Approx: Probability difference in logodds is greater than 0

	Haldane	Flat	Non-informative
Prior p-value			Beta(.5, .5) 0.96706

Problem 1b

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

Probability that the log differences are greater than 0

	Halo	dane Beta(0, 0)	Fl	at Beta(1,	1)	Non-info	rmative B	eta(.5, .5)
Iterations p-value	10000 0.97293	1e+05 0.97254	1e+06 0.97256	10000 0.97303	1e+05 0.97258	1e+06 0.97265	10000 0.97105	1e+05 0.97251	1e+06 0.97281
Standard Error	0.01137	0.00360	0.00114	0.01125	0.00359	0.00113	0.01136	0.00359	0.00113

Problem 1c

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

Odds Ratio > 1
$$\Longrightarrow \frac{\pi}{1-\pi} > \frac{\psi}{1-\psi}$$

```
##
   Fisher's Exact Test for Count Data
##
##
## data: blue
## p-value = 0.06927
## alternative hypothesis: true odds ratio is greater than 1
## 95 percent confidence interval:
   0.8606483
## sample estimates:
## odds ratio
##
     6.334078
## [1] "Probabilitiy that log differences are greater than 0: 0.93073"
```

The Fisher Exact test produces a probability that is lower than the methods in (1a) and (1b).

Problem 1d

Is deliquency releated to birth order?

Row Proportion

	Oldest	In-between	Youngest	Only Child	Total
Most Delinquent			0.25833	0.04722	1.00000
Least Delinquent	0.44402	0.26898	0.20335	0.08366	1.00000

Difference in Proportion for Oldest: Most Delinquent - Least Delinquent

Iterations	10000	100000	1000000
Mean	-0.04753	-0.04558	-0.04565
95% CI	(-0.79796, 0.71730)	(-0.80043, 0.73221)	(-0.79722, 0.73221)
P(diff>0)	0.4313	0.43721	0.436071

Difference in Proportion for In-Between : Most Delinquent - Least Delinquent

Iterations	10000	100000	1000000
Mean	0.04349	0.03782	0.03614
95% CI	(-0.66624, 0.74943)	(-0.67684, 0.73751)	(-0.67381, 0.74165)
P(diff>0)	0.5723	0.56754	0.564099

Difference in Proportion for Youngest : Most Delinquent - Least Delinquent

Iterations	10000	100000	1000000
Mean	0.02591	0.02642	0.02761
95% CI	(-0.64223, 0.68651)	(-0.61413, 0.67724)	(-0.61398, 0.68130)
P(diff>0)	0.558	0.56111	0.563085

Difference in Proportion for Only: Most Delinquent - Least Delinquent

Iterations	10000	100000	1000000
Mean	-0.01886	-0.01806	-0.01811
95% CI	(-0.44356, 0.29082)	(-0.43012, 0.28790)	(-0.42690, 0.28280)
P(diff>0)	0.3603	0.36121	0.359778

Combined p-values: $X = -2\sum_{i=1}^4 (p_i) \sim \chi_{df=8}^2$

Iterations	10000	1e+05	1e+06
Combined p-value	0.02310	0.02168	0.02265

There is evidence that birth order has an effect on deliquency rates.

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.

$$\begin{split} \hat{p} &\sim \mathcal{N}\left(p, \ \frac{p(1-p)}{n}\right) \\ \text{Let } g(\hat{p}) &= 2 \sin^{-1} \sqrt{\hat{p}} \\ \sqrt{n} \left(g(\hat{p}) - g(p)\right) \overset{\mathcal{D}}{\rightarrow} \mathcal{N}\left(0, \ \sigma^2[g'(p)]^2\right) \\ \sigma^2 &= \frac{p(1-p)}{n} \\ g'(p) &= \frac{1}{\sqrt{1-p}\sqrt{p}} \\ [g'(p)]^2 &= \frac{1}{(1-p)p} \\ \sigma^2[g'(p)]^2 &= \frac{p(1-p)}{n} \cdot \frac{1}{(1-p)p} = \frac{1}{n} \\ \sqrt{n} \left(g(\hat{p}) - g(p)\right) \overset{\mathcal{D}}{\rightarrow} \mathcal{N}\left(0, \ \frac{1}{n}\right) \\ g(\hat{p}) \overset{\mathcal{D}}{\rightarrow} \mathcal{N}\left(g(\hat{p}), \ 1\right) \\ 2 \sin^{-1} \sqrt{\hat{p}} \overset{\mathcal{D}}{\rightarrow} \mathcal{N}\left(2 \sin^{-1} \sqrt{p} \ , \ 1\right) \end{split}$$

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.

$$\overline{x} \sim \mathcal{N}(\theta, 1/n)$$

$$\overline{y} \sim \mathcal{N}(\phi, 1/n)$$
Let $h(x, y) = x/y$
Let $h(B) = \overline{x}/\overline{y}$
Let $h(\beta) = \theta/\phi$

$$\sqrt{n} (h(B) - h(\beta)) \stackrel{\mathcal{D}}{\to} \mathcal{N} \left(0, \nabla h(\beta)^T \cdot \Sigma \cdot \nabla h(\beta)\right)$$

$$\Sigma = \begin{bmatrix} 1/n & 0 \\ 0 & 1/n \end{bmatrix}$$

$$\nabla h(\beta)^T = \begin{bmatrix} \frac{\partial h}{\partial x} & \frac{\partial h}{\partial y} \\ \frac{\partial h}{\partial y} \end{bmatrix}_{\theta, \phi}$$

$$= \begin{bmatrix} \frac{1}{y} & -\frac{x}{y^2} \end{bmatrix}_{\theta, \phi}$$

$$= \begin{bmatrix} \frac{1}{\phi} & -\frac{\theta}{\phi^2} \end{bmatrix} \begin{bmatrix} 1/n & 0 \\ 0 & 1/n \end{bmatrix} \begin{bmatrix} \frac{1}{\phi} \\ -\frac{\theta}{\phi^2} \end{bmatrix}$$

$$\nabla h(\beta)^T \cdot \Sigma \cdot \nabla = \begin{bmatrix} \frac{1}{\phi} & -\frac{\theta}{\phi^2} \end{bmatrix} \begin{bmatrix} 1/n & 0 \\ 0 & 1/n \end{bmatrix} \begin{bmatrix} \frac{1}{\phi} \\ -\frac{\theta}{\phi^2} \end{bmatrix}$$

$$= \begin{bmatrix} \frac{1}{n\phi} & -\frac{\theta}{n\phi^2} \end{bmatrix} \begin{bmatrix} \frac{1}{\phi} \\ -\frac{\theta}{\phi^2} \end{bmatrix}$$

$$= \frac{1}{n} \left(\frac{1}{\phi^2} - \frac{\sigma^2}{\phi^4} \right)$$

$$\sqrt{n} (\overline{x}/\overline{y} - \theta/\phi) \stackrel{\mathcal{D}}{\to} \mathcal{N} \left(0, \frac{1}{n} \left(\frac{1}{\phi^2} - \frac{\sigma^2}{\phi^4} \right) \right)$$

$$\frac{\overline{x}}{\overline{y}} \stackrel{\mathcal{D}}{\to} \mathcal{N} \left(\frac{\theta}{\phi}, \frac{1}{\phi^2} - \frac{\sigma^2}{\phi^4} \right)$$

197 animals are distributed into four categories: $Y = (y_1, y_2, y_3, y_4)$ according to the genetic linkage model $(\frac{2+\theta}{4}, \frac{1-\theta}{4}, \frac{1-\theta}{4}, \frac{\theta}{4})$. 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 $\hat{\theta}$.

$$L(\theta \mid \mathbf{Y}) = \frac{(y_1 + y_2 + y_3 + y_4)!}{y_1! y_2! y_2! y_4!} \left(\frac{2 + \theta}{4}\right)^{y_1} \left(\frac{1 - \theta}{4}\right)^{y_2} \left(\frac{1 - \theta}{4}\right)^{y_3} \left(\frac{\theta}{4}\right)^{y_4}$$

$$\propto (2 + \theta)^{y_1} \cdot (1 - \theta)^{y_2 + y_3} \cdot (\theta)^{y_4}$$

$$\ell(\theta \mid \mathbf{Y}) \propto y_1 \log(2 + \theta) + (y_2 + y_3) \log(1 - \theta) + y_4 \log(\theta)$$

$$\frac{\partial \ell}{\partial \theta} = \frac{y_1}{2 + \theta} - \frac{y_2 + y_3}{1 - \theta} + \frac{y_4}{\theta}$$

$$\frac{\partial^2 \ell}{\partial \theta^2} = -\frac{y_1}{(2 + \theta)^2} - \frac{y_2 + y_3}{(1 - \theta)^2} - \frac{y_4}{\theta^2}$$

$$\theta^{(i+1)} = \theta^{(i)} - \frac{\frac{y_1}{2 + \theta^{(i)}} - \frac{y_2 + y_3}{1 - \theta^{(i)}} + \frac{y_4}{\theta^{(i)}}}{-\frac{y_1}{(2 + \theta^{(i)})^2} - \frac{y_2 + y_3}{(1 - \theta^{(i)})^2} - \frac{y_4}{(\theta^{(i)})^2}} \xrightarrow{\text{Newton-Raphson } \hat{\theta}}$$

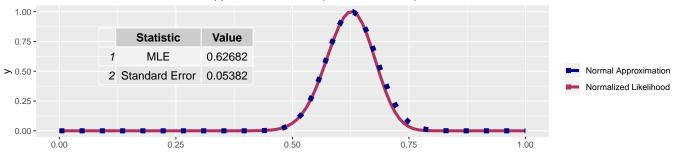
$$s.e.(\hat{\theta}) = \sqrt{1/\mathcal{I}(\theta)}$$

$$\mathcal{I}(\theta) = \left[\frac{\partial^2 \ell}{\partial \theta^2}\right]_{\hat{\theta}} = -\frac{y_1}{(2 + \hat{\theta})^2} - \frac{y_2 + y_3}{(1 - \hat{\theta})^2} - \frac{y_4}{\hat{\theta}^2}$$

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.

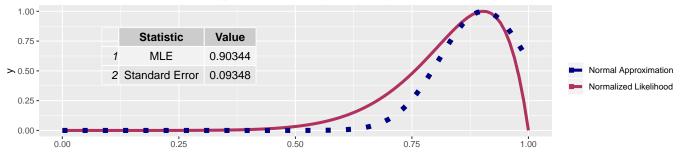




Problem 4b

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

Normal Likelihood and Normal Approximation for Y=(14, 0, 1, 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.

$$L(\theta \mid Y) \propto (2 + \theta)^{y_1} (1 - \theta)^{y_2 + y_3} (\theta)^{y_4}$$

$$\ell(\theta \mid Y) \propto y_1 \ln(2 + \theta) + (y_2 + y_3) \ln(1 - \theta) + y_4 \ln(\theta)$$

$$p(\theta) = \theta^{\alpha - 1} (1 - \theta)^{\beta - 1}$$
Flat Prior $p(\theta) = \theta^{1 - 1} (1 - \theta)^{1 - 1} = 1$

$$-nh(\theta) = \ell(\theta \mid Y) + \ln(p(\theta))$$

$$= y_1 \ln(2 + \theta) + (y_2 + y_3) \ln(1 - \theta) + y_4 \ln(\theta) + \ln(1)$$

$$= y_1 \ln(2 + \theta) + (y_2 + y_3) \ln(1 - \theta) + y_4 \ln(\theta)$$

$$-nh^*(\theta) = \ell(\theta \mid Y) + \ln(p(\theta)) + \ln(g(\theta))$$

$$= y_1 \ln(2 + \theta) + (y_2 + y_3) \ln(1 - \theta) + (y_4 + 1) \ln(\theta)$$

$$\hat{\theta} = \text{Newton-Raphson Result}$$

$$\theta^* = \text{Newton-Raphson Result}$$

$$\hat{\sigma} = [h''(\theta)]_{\hat{\theta}}^{-1/2} \qquad h(\theta) = \frac{1}{-n} \cdot -nh(\theta)$$

$$\sigma^* = [(h^*)''(\theta)]_{\theta^*}^{-1/2} \qquad h^*(\theta) = \frac{1}{-n} \cdot -nh^*(\theta)$$

$$\mathbb{E}_{\theta} [\theta] = \frac{\sigma^*}{\hat{\sigma}} \cdot \frac{\exp\{-nh^*(\theta^*)\}}{\exp\{-nh(\hat{\theta})\}}$$

Laplace's Method (Second Order) of Posterior Mean for Y=(125, 18, 20, 34)

Statistic	Value
theta.hat	0.62682
theta.star	0.63099
sigma.hat	0.72238
sigma.star	0.71529
Posterior Mean	0.62275

Laplace's Method (Second Order) of Posterior Mean for Y=(14, 0, 1, 5)

Statistic	Value
theta.hat	0.90344
theta.star	0.91217
sigma.hat	0.41696
sigma.star	0.38003
Posterior Mean	0.82752