

**QUALIFYING EXAM SOLUTIONS**  
**Statistical Methods**  
**Fall, 2011**

1. (a) The model is

$$Y \sim \text{Poisson}(\alpha + \beta X).$$

Suppose  $(Y_i, X_i)$  for  $i = 1, \dots, n$  are observed. The loglikelihood function is

$$\ell(\alpha, \beta) = -\sum_{i=1}^n Y_i! + \sum_{i=1}^n Y_i \log(\alpha + \beta X_i) - \sum_{i=1}^n (\alpha + \beta X_i).$$

Then, the likelihood equations are

$$\begin{aligned} \frac{\partial \ell(\alpha, \beta)}{\partial \alpha} &= \sum_{i=1}^n \frac{Y_i}{\alpha + \beta X_i} - n \\ \frac{\partial \ell(\alpha, \beta)}{\partial \beta} &= \sum_{i=1}^n \frac{Y_i X_i}{\alpha + \beta X_i} - \sum_{i=1}^n X_i. \end{aligned}$$

It is clear that those are different from the least square equations. Therefore, the MLE is not equal to the LSE.

- (b) Let  $\pi_i$  be the probability. Then the model is

$$\log \frac{\pi_j}{\pi_1} = \beta_{j0} + \beta_{j1} X.$$

Then, the model is equivalent to

$$\pi_j = \pi_1 e^{\beta_{j0} + \beta_{j1} X}$$

and

$$\sum_{j=1}^J \pi_j = 1.$$

If we fit  $J - 1$  separate logistic regression model for  $(1, j)$ , we also have

$$\pi_j = \pi_1 e^{\beta_{j0} + \beta_{j1} X}$$

which is the same as the formula from the previous model. However, the constraint is  $\pi_j + \pi_1 = 1$ , which is different from the previous one. Therefore, the two models are different.

- (c) Let  $\pi_j$  be the probability of the  $j$ -th category. The model assumption is

$$\log \frac{\pi_1 + \dots + \pi_j}{\pi_{j+1} + \dots + \pi_5} = \theta_j - \beta_2 I_{X=2} - \beta_3 I_{X=3} - \beta_4 I_{X=4},$$

for  $j = 1, 2, 3, 4$ . Then, we have  $\hat{\theta}_1 = -0.9188$ ,  $\hat{\theta}_2 = -0.5183$ ,  $\hat{\theta}_3 = 0.4922$ ,  $\hat{\theta}_4 = 1.8579$ ,  $\hat{\beta}_2 = 0.1176$ ,  $\hat{\beta}_3 = 0.3174$ ,  $\hat{\beta}_4 = 0.5208$ . If the last category is used as a baseline, then, we have the model

$$\log \frac{\pi_1 + \dots + \pi_j}{\pi_{j+1} + \dots + \pi_5} = \theta'_j - \beta'_1 I_{X=1} - \beta'_2 I_{X=2} - \beta'_3 I_{X=3}.$$

Compare it with the previous model, we have

$$\theta'_j = \theta_j - \beta_4$$

and

$$\beta'_i = \beta_i + \beta_4.$$

Therefore, we have  $\hat{\theta}'_1 = -1.4396$ ,  $\hat{\theta}'_2 = -1.0391$ ,  $\hat{\theta}'_3 = -0.0286$ ,  $\hat{\theta}'_4 = 1.3371$ ,  $\hat{\beta}'_1 = -0.5208$ ,  $\hat{\beta}'_2 = -0.4032$ , and  $\hat{\beta}'_3 = -0.2034$ .

2. (a) Because the model does not contain any interaction effect, it assumes all the interaction effects are zero:

$$\log \lambda_{ijkl} = \mu + \alpha_i + \beta_j + \gamma_k + \delta_l,$$

where  $\lambda_{ijkl}$  is the expected value of the response,  $\alpha_i$ ,  $\beta_j$ ,  $\gamma_k$  and  $\delta_l$ , represents Heart, Comps, Smoke, and BW, respectively. It assumes all the independent variables are independent.

- (b) The expected count is

$$e^{5.15385-1.55769-0.04539-0.99056} = 12.94.$$

- (c) The residual degree of freedom is  $df = 16 - 5 = 11$ . Since the residual deviance is  $164.7 > \chi^2_{0.05,11} = 19.67$ , we reject the null hypothesis and conclude that at least one of the interaction effects are not zero. Therefore, the four independent variables are not independent.

- (d) The model is

$$\log \lambda_{ijkl} = \mu + \alpha_i + \beta_j + \delta_l + (\alpha\beta)_{ij} + (\alpha\delta)_{il} + (\beta\delta)_{jl}.$$

Try to interpret it by yourself. Keep in mind that if the interaction between two independent variable is not in the model, then they are independent.

- (e) The residual degree of freedom is 9. Since  $G^2 = 9.561 < \chi^2_{0.05,9} = 16.92$ , we accept the model.
3. (a) Let  $n$  be the total number of the subjects. We can choose a random order 1 to  $n$  and assign the  $i$ -th number to the  $i$ -th subject. If the  $i$ -th number is less than or equal to  $n/2$ , then for (D1) we assign one treatment otherwise we assign the second treatment, for (D2) we assign A first then B otherwise B first and A.
- (b) For (D1) treatment A and treatment B effect are independent measure, but they are dependent in (D2). However, the treatment effect is nested in subject in (D1) but not in (d2).
- (c) For (D1), let  $\bar{Y}_1$  be the average of treatment 1 and  $\bar{Y}_2$  be the average of the treatment 2. Then

$$V(\bar{Y}_1 - \bar{Y}_2) = 2 \times 10^2 / 50 = 4.$$

For (D2), let  $y_{i1}$  be the measurement of treatment A and  $y_{i2}$  be the measurement of treatment B. Let  $\delta_i = y_{i1} - y_{i2}$ , and  $\hat{\delta} = \sum_{i=1}^{100} \delta_i$ .

$$V(\delta_i) = 2 \times 10^2 - 2 \times 0.82 \times 10^2 = 36.$$

Then

$$V(\bar{\delta}) = \frac{36}{100} = 0.36.$$

Therefore, the second measure has lower variance, which is more powerful if the treatment mean is not affected.

- (d) We need to include: (i) the nested effect in (D1) and the dependence in (d2); (ii) the power function comparison.

4. (a) An exponential family PMF (or PDF) has the form of

$$f(y, \theta, \phi) = \exp \frac{y\theta - b(\theta)}{a(\phi)} + c(\theta, \phi).$$

The PMF of the negative binomial is

$$f(y; k, \mu) = \exp \left\{ y \log \frac{\mu}{\mu + k} + k \log \frac{k}{\mu + k} - \log \frac{\Gamma(y + k)}{\Gamma(k)\Gamma(y + 1)} \right\}.$$

Because  $k$  is known, the above is an exponential family if we use  $\theta = \log \frac{\mu}{\mu + k}$ . In this case, we have  $b(\theta) = -k \log(1 - e^\theta)$  and  $\phi = 1$ . Thus,

$$E(y) = b'(\theta) = \mu.$$

- (b) The canonical link is  $g(\mu) = \theta$  which is

$$g(\mu) = \log \frac{\mu}{\mu + k}.$$

- (c) We consider the model

$$\log \frac{\mu_i}{\mu_i + k} = \alpha + \beta x_i$$

where  $x = 1$  if the  $i$ -th person is a patient, and  $x = 0$  if not. Let  $x_i = 1$  for  $i = 1, 2, 3$  and  $x_i = 0$  for  $i = 4, 5, 6$ . Then  $\mu_i = k/(1 - e^\alpha)$  for  $i = 1, 2, 3$  and  $\mu_i = k/(1 - e^{\alpha+\beta})$  for  $i = 4, 5, 6$ . The loglikelihood function is

$$\begin{aligned} \ell(\alpha, \beta) &= \sum_{i=1}^6 f(y_i; k, \mu_i) \\ &= \sum_{i=1}^6 \log \frac{\Gamma(y_i + k)}{\Gamma(k)\Gamma(y_i + 1)} + 3k \log(1 - e^\alpha) + 3k \log(1 - e^{\alpha+\beta}) \\ &\quad + \alpha(y_1 + y_2 + y_3) + (\alpha + \beta)(y_4 + y_5 + y_6). \end{aligned}$$

5. (a) Matching can eliminate the age, gender, location, and SES effects. Otherwise, we do not know the different is caused by WCV or other effects.

- (b) For the prior year, the odds ratio is

$$\hat{\theta} = \frac{509 \times 1593}{407 \times 1491} = 1.336.$$

For the post year

$$\hat{\theta} = \frac{672 \times 1625}{375 \times 1328} = 2.193.$$

- (c) The main effect of wcv reflects the odds ratio for prior year. When it is combined with the interaction effect, it reflects the odds ratio for the post year.
- (d) The interaction effect reflects the change of the odds ratio. We have

$$\log(2.192) - \log(1.336) = 0.4955$$

which is close to the interaction effect.

- (e) This model is a saturated model. It is equivalent to the method using original data.
6. (a) State is a fixed effect, Farm (State) and Cow (Farm\*State) are random effects.
- (b) There is no degree of freedom in the error terms because the estimation of Cow effect used all the observations. Actually, the model is a saturated model.
- (c) DF are 4, 120, 1750, respectively.
- (d) The F-statistic is

$$F^* = \frac{4289/4}{71463/1750} = 26.3$$

which is  $\geq \chi_{0.05,4} = 9.48$ . Thus, they are different.

- (e) The estimate of the variance is  $\hat{\sigma}^2 = 71463/1750 = 40.84$ . The variance of the mean is

$$\hat{Y}_{5..} = \frac{\hat{\sigma}^2}{375} = 0.1089$$

The 95% confidence interval is

$$34.70 \pm 1.96 \times \sqrt{0.1089} = [34.05, 35.35].$$