## BASIC PHD WRITTEN EXAMINATION IN BIOSTATISTICS

### THEORY, SECTION 2

# (9:00 AM- 1:00 PM Thursday, August 12, 2010)

#### INSTRUCTIONS:

- a) This is a CLOSED-BOOK examination.
- b) The time limit for this Examination is four hours.
- c) Answer any TWO (2) (BUT ONLY TWO) of the THREE (3) questions that follow.
- d) Put the answers to different questions on separate sets of paper.
- e) Put your code letter, **NOT YOUR NAME**, on each page. The same code will be used for Section 1 and Section 2 of the PhD Theory Exam. Please keep the code confidential and do not share this information with any students or faculty.
- f) Return the examination with a signed statement of the UNC honor pledge, separately from your answers. The pledge statement is given on the last page of the exam handout.
- g) In the questions to follow, you are required to answer only what is asked, and not to tell all you know about the topics involved.

- 1. Consider independent observations  $(X_1, Y_1), \dots, (X_n, Y_n)$ , where  $Y_i$  takes values 0 and 1. Suppose that  $X_i|Y_i=m \sim N(\mu_m, \sigma^2)$  and  $P(Y_i=m)=\pi_m$  for m=0,1, where  $\pi_0+\pi_1=1$  and  $\pi_0 \in (0,1)$ .
  - (a) Show that  $P(Y_i = m|X_i), m = 0, 1$  satisfies a logistic model, that is

$$logit(P(Y_i = 1|X_i, \alpha)) = \alpha_0 + \alpha_1 X_i$$

where logit(u) = log(u/(1-u)),  $\alpha = (\alpha_0, \alpha_1)$ , and  $\alpha_0$  and  $\alpha_1$  are unknown parameters. Derive the explicit form of  $\alpha = g(\theta)$  as a function of  $\theta = (\pi_1, \mu_0, \mu_1, \sigma^2)$ .

- (b) Based on the logistic model in (a), please give the explicit form of the Newton-Raphson algorithm for calculating the maximum likelihood estimate of  $\alpha$ , denoted by  $\hat{\alpha} = (\hat{\alpha}_0, \hat{\alpha}_1)$ , and derive the asymptotic covariance matrix of  $\hat{\alpha}$ .
- (c) Please write down the joint distribution of  $\{(X_i, Y_i) : i = 1, \dots, n\}$  and calculate the maximum likelihood estimate of  $\theta$ , denoted by  $\hat{\theta}_F$ , and its asymptotic covariance matrix.
- (d) Calculate the asymptotic covariance matrix of  $g(\hat{\theta}_F)$ .
- (e) In this part, suppose that  $\mu_0 = \mu_1$ . Show that  $\operatorname{Cov}(\hat{\alpha})^{-1}\operatorname{Cov}(g(\hat{\theta}_F))$  converges to a matrix, which does not depend on  $\theta$ . Please interpret the results.
- (f) Now, suppose that  $\pi_1$  is known. Will the results in (b)-(e) be changed? Please explain. If so, then please derive the corresponding results and compare with those obtained above.

#### 2. Consider the following model:

$$Y_i = X_{i1}\beta_1 + X_{i2}\beta_2 + \ldots + X_{in}\beta_n + U + \epsilon_i, \tag{0.1}$$

 $i=1\ldots n$ , where  $\beta_1,\ldots,\beta_p$  are unknown parameters,  $Y=(Y_1,\ldots,Y_n)$  is the vector of responses, and  $X_{ij}, i=1\ldots n, j=1\ldots p$ , are fixed covariates. Assume that  $\epsilon_i \sim N(0,\sigma^2)$ ,  $U \sim N(\alpha,k\sigma^2)$ , where  $\alpha$  and  $\sigma^2 > 0$  are unknown, k>0 is known, and  $\epsilon_i$  are independent of each other and of U. Assume further that the  $(n \times p)$  matrix with entries  $X_{ij} - \overline{X}_{.j}$  has rank p, where  $\overline{X}_{.j} = \frac{1}{n} \sum_{i=1}^{n} X_{ij}$ .

- (a) Find the distribution of Y, and show that the variance-covariance matrix for Y is positive-definite. (Hint: For a constant c the inverse of a matrix I + cJ is in the form of I + dJ for certain constant d, where I is the  $(n \times n)$  identity matrix and J is the  $(n \times n)$  matrix with all entries equal to 1.)
- (b) Show that  $\beta_1, \ldots, \beta_p$  and  $\alpha$  are estimable.
- (c) Let  $\theta = (\alpha, \beta_1, \dots, \beta_p)^T$ . Derive the maximum likelihood estimator for  $\theta$ , denoted by  $\hat{\theta} = (\hat{\alpha}, \hat{\beta}_1, \dots, \hat{\beta}_p)^T$ . What is the distribution of  $\hat{\theta}$ ?
- (d) Let  $\tilde{\theta} = (\tilde{\alpha}, \tilde{\beta}_1, \dots, \tilde{\beta}_p)$  be the value of the vector  $\theta$  minimizing the sum of squares

$$\sum_{i=1}^{n} [Y_i - (\alpha + X_{i1}\beta_1 + \ldots + X_{ip}\beta_p)]^2.$$

Is it true that  $Var(\hat{\alpha}) < Var(\tilde{\alpha})$ ? Carefully justify your answer.

3. To evaluate the diagnostic performance using two continuous biomarkers, we randomly select n diseased subjects and m non-diseased subjects. Let  $X_1 = (X_{11}, X_{12})', ..., X_n = (X_{n1}, X_{n2})'$  be these two measured biomarkers for the diseased subjects and  $Y_1 = (Y_{11}, Y_{12})', ..., Y_m = (Y_{m1}, Y_{m2})'$  be the same two measured biomarkers for the non-diseased subjects. We aim to find an optimal linear combination of these two biomarkers to maximize some measure of the diagnostic performance. In particular, we need to find  $\beta = (\beta_1, \beta_2)'$  such that the area under the receiver operating characteristics curve, defined by  $AUC(\beta) \equiv P(\beta'\mathbf{X}_1 \geq \beta'\mathbf{Y}_1)$ , is maximized.

Assume  $X_1, ..., X_n$  are i.i.d from  $MN(\mu_1, \Sigma)$  and  $Y_1, ..., Y_m$  are i.i.d from  $MN(\mu_2, \Sigma)$ , where  $\mu_1 = (\mu_{11}, \mu_{21})', \mu_2 = (\mu_{12}, \mu_{22})', \Sigma = \begin{pmatrix} \sigma_{11} & \sigma_{12} \\ \sigma_{12} & \sigma_{22} \end{pmatrix}$  are unknown parameters and  $\Sigma$  is a positive definite matrix. Moreover, assume  $m = \tau n$  for a fixed constant  $\tau > 0$ .

- (a) Show  $AUC(\beta) = \Phi\left(\beta'(\mu_1 \mu_2)/\sqrt{2\beta'\Sigma\beta}\right)$ , where  $\Phi(x)$  is the cumulative distribution function of N(0,1).
- (b) Show that the maximum of  $AUC(\beta)$ , denote as  $A^{optimal}$ , is

$$\Phi\left(\left[(\mu_1-\mu_2)'\Sigma^{-1}(\mu_1-\mu_2)/2\right]^{1/2}\right).$$

Hint: the  $\beta$  maximizing  $AUC(\beta)$  is unique up to some multiplicative scale.

- (c) Calculate the maximum likelihood estimator for  $A^{optimal}$  and denote it by  $\hat{A}$ .
- (d) Describe how you will obtain the asymptotic distribution of  $\sqrt{n}(\hat{A} A^{optimal})$ . You do not need to give the explicit expression of the asymptotic variance.
- (e) To test whether the combination of the two biomarkers is useful for diagnosis, we formulate the hypothesis  $H_0: A^{optimal} = 1/2$  vs  $H_1: A^{optimal} > 1/2$  and reject  $H_0$  when  $\hat{A} > c_n$  for some threshold value  $c_n$  (depending on n).
  - i. Determine  $c_n$  such that the type I error converges to a given level  $\alpha$ , where  $c_n$  is a constant depending only on n and  $\alpha$ ; that is,  $\lim_{n\to\infty} P(\hat{A} > c_n|H_0) = \alpha$ .
  - ii. Calculate the asymptotic power of this test at a local alternative  $H_1: A^{optimal} = 1/2 + \delta/\sqrt{n}$  where  $\delta$  is a fixed positive constant.

# 2010 PhD Theory Exam, Section 2

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Statement of the UNC honor pledge: