

# Empirical Likelihood Ratio Confidence Interval Estimation of Best Linear Combinations of Biomarkers

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## Abstract

A novel smoothed empirical likelihood (EL) approach that incorporates kernel estimation of the area under the receiver operating characteristic curve (AUC) to construct nonparametric confidence intervals of AUC based on the best linear combination (BLC) of biomarkers is proposed. The method has several advantages including the feasibility to use gradient-based techniques for fast computation of BLC coefficients and to employ powerful likelihood methods without specification of underlying data distributions. Simulation results show that the new method performs well even when the distribution of biomarkers is skewed, a situation commonly met in practice. A data set from a clinical experiment related to atherosclerotic coronary heart disease is used to illustrate the efficiency of the proposed method.

*Keywords:* Area under the ROC curve; Best linear combination; Empirical likelihood; Kernel; Receiver operating characteristic curve (ROC).

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## 1. Introduction

2 The receiver operating characteristic (ROC) curve methodology is often ap-  
3 plied to evaluate the performance of diagnostic markers that classify subjects  
4 into two populations: diseased and nondiseased. Suppose that a biomarker,

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5 measured on a continuous scale, is performed on  $n$  diseased subjects, yielding  
6 independent, identically distributed (i.i.d.) measurements  $\mathbf{X}_i$ ,  $i = 1, \dots, n$ , with  
7 distribution function  $F$ , and on  $m$  nondiseased subjects, yielding i.i.d. measure-  
8 ments  $\mathbf{Y}_j$ ,  $j = 1, \dots, m$ , with distribution function  $G$ . The ROC curve is a plot  
9 of  $R(t) = 1 - F\{G^{-1}(1 - t)\}$  against  $t$ ,  $0 \leq t \leq 1$  (Pepe, 1997; Faraggi and  
10 Reiser, 2002). The area under the ROC curve (AUC), as a common measure  
11 of the diagnostic performance of a biomarker, is equal to  $P(X > Y)$  (Bamber,  
12 1975).

13 In practice, different markers are usually related to the disease in various  
14 magnitudes and different directions. For example, low levels of high density  
15 lipoprotein (HDL)-cholesterol and high levels of thiobarbuturic acid reacting  
16 substances (TBARS) are biomarkers of oxidative stress and antioxidant sta-  
17 tus, as indicators of coronary heart disease (Schisterman et al., 2002). When  
18 multiple biomarkers are available, we are interested in seeking a simple best  
19 linear combination (BLC) of biomarkers such that the combined score achieves  
20 the maximum AUC over all possible linear combinations. The maximum AUC  
21 measures the ability to discriminate between the control and the disease groups  
22 (Pepe and Thompson, 2000; McIntosh and Pepe, 2002).

23 A variety of publications have focused on the BLC of multiple biomarkers  
24 in both parametric and nonparametric cases. Su and Liu (1993) investigated  
25 the BLC and the corresponding maximum AUC under multivariate normal as-  
26 sumptions. Based on the point estimator given by Su and Liu (1993), Reiser  
27 and Faraggi (1997) derived confidence intervals for the BLC-based AUC under  
28 normal assumptions, which is useful when the sample size in the two groups is  
29 small and equal or moderate and unequal, respectively. In practice, however,  
30 the distributions of biomarker measurements commonly deviate from the normal  
31 assumptions (Limpert et al., 2001). For example, epidemiological studies have  
32 demonstrated that TBARS has a distribution with heavy tails (Schisterman  
33 et al., 2001). In this case, loss of efficiency of the BLC coefficients constructed  
34 under normality assumptions can be expected and alternative approaches are  
35 needed to accommodate different data distributions.

36 Pepe et al. (2006) considered the empirical AUC and noticed that the empir-  
37 ical BLC estimator is consistent under the generalized linear model assumption.  
38 They noted that sophisticated computational algorithms are required because  
39 the empirical AUC is not a continuous function. Ma and Huang (2005, 2007)  
40 also noticed the computational complexity in maximizing the empirical AUC  
41 and proposed a corresponding sigmoid approximation. Under the generalized  
42 linear model assumption, it is proved that the sigmoid AUC converges to the  
43 maximized theoretical AUC and the corresponding BLC coefficients are asymp-  
44 totically normally distributed (Ma and Huang, 2007). Instead of the specific  
45 sigmoid AUC, Vexler et al. (2006) demonstrated consistency, uniqueness, and  
46 ease of computation of the kernel-smoothed AUC estimator. Furthermore, they  
47 provided an upper confidence bound for the maximum AUC. The distribution-  
48 free confidence interval estimation of the maximum AUC remains unsolved.

49 As a nonparametric alternative to the optimal parametric likelihood method  
50 (Markatou et al., 1998), the EL methodology (Owen, 1990, 2001) is an efficient  
51 way to construct confidence intervals. In the context of a single biomarker,  
52 Qin and Zhou (2006) proposed an EL-based confidence interval for the AUC  
53 and demonstrated that it outperforms the existing normal approximation-based  
54 intervals and bootstrap intervals, particularly when the AUC is close to one.

55 In this paper, we extend and modify the method of Qin and Zhou (2006) to  
56 adjust for multiple biomarkers. Incorporating a kernel distribution estimation  
57 (Azzalini, 1981; Nadaraya, 1964) into the EL-based confidence interval estima-  
58 tion, we propose smoothed EL-based confidence intervals for BLC-based AUCs.  
59 The advantages of the proposed method include: (1) the smooth estimate of the  
60 AUC is differentiable, making gradient-based methods feasible and the compu-  
61 tation of BLC coefficients simple, fast and unique; (2) it is robust to underlying  
62 distributions of biomarkers' measurements; (3) it employs the EL methodology,  
63 that is known to be an approximate nonparametric most powerful tool based  
64 on likelihood ratios (Lazar and Mykland, 1998); (4) the EL-based confidence  
65 interval is range preserving (Hall and La Scala, 1990) and therefore always lies  
66 between 0.5 and 1. The proposed method is illustrated with a study of the ac-

67 curacy of biomarkers related to the atherosclerotic coronary heart disease. An  
68 EL interval for AUC is derived based on a linear combination of measurements  
69 of four important biomarkers related to the atherosclerotic coronary heart dis-  
70 ease including lutein, TBARS, HDL cholesterol and uric acid, obtained from a  
71 12-hour fasting blood specimen for biochemical analysis at baseline.

72 The paper is organized as follows. Section 2 develops the smoothed EL-based  
73 confidence interval for the BLC-based AUC. Section 3.2 presents simulation  
74 studies to compare the relative performance of the proposed smoothed EL-based  
75 interval with the existing intervals for the BLC-based AUC. In Section 4, we  
76 apply our method to a real example related to atherosclerotic coronary heart  
77 disease. Section 5 presents a broader discussion on deriving linear combinations  
78 of biomarkers to improve the diagnostic accuracy. Conditions and proofs are  
79 deferred to the Appendix.

## 80 2. Methods

81 Consider a study with  $d$  continuous-scale biomarkers yielding measurements  
82  $\mathbf{X}_i = (X_{1i}, \dots, X_{di})^T$ ,  $i = 1, \dots, n$ , on  $n$  diseased subjects and measurements  
83  $\mathbf{Y}_j = (Y_{1j}, \dots, Y_{dj})^T$ ,  $j = 1, \dots, m$ , on  $m$  nondiseased patients, respectively.  
84 We are interested in constructing effective one-dimensional combined scores of  
85 biomarkers measurements, say,  $X(\mathbf{a}) = \mathbf{a}^T \mathbf{X}$  and  $Y(\mathbf{a}) = \mathbf{a}^T \mathbf{Y}$ , such that the  
86 AUC based on these scores is maximized over all possible linear combinations  
87 of biomarkers. Define  $A(\mathbf{a}) = P(X(\mathbf{a}) > Y(\mathbf{a}))$ ; the statistical problem is to  
88 estimate the maximum AUC defined as  $A = A(\mathbf{a}_0)$ , where  $\mathbf{a}_0$  are the BLC co-  
89 efficients satisfying  $\mathbf{a}_0 = \arg \max_{\mathbf{a}} A(\mathbf{a})$ . For simplicity, we assume that the first  
90 component of the vector  $\mathbf{a}$  equals to 1 throughout the paper; see Pepe et al.  
91 (2006) and Ma and Huang (2007).

### 92 2.1. Confidence interval for a single biomarker-based AUC

93 Qin and Zhou (2006) developed an EL-approach for constructing confidence  
94 intervals for the AUC using a single biomarker (in our notation  $d = 1$  and

95  $\mathbf{a} = \mathbf{a}_0 = 1)$ . The confidence interval estimation is executed via construction  
 96 of the empirical likelihood ratio (ELR) test statistic for testing the hypothesis  
 97  $H_0 : A = A_0$  versus  $H_a : A \neq A_0$ .

Qin and Zhou (2006) based the ELR test statistic on the concept of placement value of a diseased subject (Pepe and Cai, 2004). The placement value is defined as  $U = 1 - G(X)$ , where  $U$  can be interpreted as the proportion of non-diseased subjects with their biomarker measurements greater than  $X$ . Noticing that  $E(1 - U) = A$ , where  $A$  denotes the AUC based on a single marker, EL inference for the AUC for the case of a single marker is developed. Specifically, let  $\mathbf{p} = (p_1, p_2, \dots, p_n)^T$  be a probability weight vector,  $\sum_{i=1}^n p_i = 1$  and  $p_i \geq 0$  for all  $i = 1, \dots, n$ . The profile EL for the AUC, evaluated at the true value  $A_0$  of AUC, can be defined as

$$\tilde{L}(A_0) = \sup\left\{\prod_{i=1}^n p_i : \sum_{i=1}^n p_i = 1, \sum_{i=1}^n p_i(1 - U_i) = A_0\right\}, \quad (1)$$

where  $U_i = 1 - G(X_i)$ ,  $i = 1, \dots, n$ . When the distribution function  $G$  of the nondiseased population is unknown, the empirical distribution  $\hat{G}$  is used. Accordingly, replacing  $U_i$  by its estimator  $\hat{U}_i = 1 - \hat{G}(X_i)$  in equation (1), and using Lagrange multipliers, one can obtain that

$$p_i = \frac{1}{n} \frac{1}{1 + \lambda(1 - \hat{U}_i - A_0)}, i = 1, \dots, n.$$

Furthermore, the Lagrange multiplier  $\lambda$  is the root of

$$\frac{1}{n} \sum_{i=1}^n \frac{1 - \hat{U}_i - A_0}{1 + \lambda(1 - \hat{U}_i - A_0)} = 0.$$

The corresponding empirical log-likelihood ratio for the AUC is

$$l(A_0) = 2 \sum_{i=1}^n \log(1 + \lambda(1 - \hat{U}_i - A_0)). \quad (2)$$

98 Thus, an EL-based confidence interval for the AUC of a single biomarker is also  
 99 constructed based on the empirical log-likelihood ratio, which has an asymptotic  
 100 scaled Chi-square distribution (Qin and Zhou, 2006).

101 However, this approach cannot be directly applied when the estimation of  
 102 best linear combination of multiple biomarkers is involved. In the next section,

103 we will use kernels to construct the EL-based confidence interval estimation  
 104 for the BLC-based maximum AUC. Incidentally, Chen and Hall (1993) showed  
 105 that the kernel approach is very efficient when EL is used to estimate confidence  
 106 intervals for quantiles.

107 *2.2. Confidence interval for the BLC-based AUC*

108 When multiple biomarkers are available, e.g.,  $d \geq 2$ , we generalize the  
 109 method of Qin and Zhou (2006) considering a BLC of biomarkers. First the  
 110 placement values are defined as  $U_i(\mathbf{a}_0) = 1 - G_{\mathbf{a}_0}(\mathbf{a}_0^T \mathbf{X}_i)$ ,  $i = 1, \dots, n$ , where  
 111  $G_{\mathbf{a}_0}$  is the cumulative distribution function of the combined score  $Y(\mathbf{a}_0)$  from  
 112 the nondiseased group.

113 In the case that  $\mathbf{a}_0$  is known, we can simply replace  $U_i$  in equation (1) by  
 114  $U_i(\mathbf{a}_0)$  or its corresponding empirical distribution estimator. However, the de-  
 115 pendence of placement values on not only the unknown cumulative distribution  
 116 function  $G_{\mathbf{a}_0}$  but also the unknown  $\mathbf{a}_0$  further complicates construction of the  
 117 ELR test statistic.

118 When the best linear combination coefficient  $\mathbf{a}_0$  is unknown, the estima-  
 119 tion of  $\mathbf{a}_0$  will be associated with a maximization of the AUC and we will  
 120 need a simple and unique solution with respect to the  $A_0$  estimation. The use  
 121 of the empirical distribution function is problematic because of its discontinu-  
 122 ity, so we consider kernel estimation of the distribution function. Let  $k$  be a  
 123 symmetric kernel function and define  $K_h(x) = \int_{-\infty}^{x/h} k(u)du$ ; see details related  
 124 to the definitions in Appendix A. Then, the kernel estimator of  $1 - U_i(\mathbf{a}_0)$  is  
 125  $v_i(\mathbf{a}_0) = \sum_{j=1}^n K_h(\mathbf{a}_0^T \mathbf{X}_i - \mathbf{a}_0^T \mathbf{Y}_j)/m$ ,  $i = 1, \dots, n$ .

The EL for the BLC-based AUC is given as

$$L(A_0) = \sup\left\{\prod_{i=1}^n p_i : \sum_{i=1}^n p_i = 1, \sum_{i=1}^n p_i v_i(\hat{\mathbf{a}}_0) = A_0\right\}, \quad (3)$$

where  $\hat{\mathbf{a}}_0$  satisfies  $\sum_{i=1}^n p_i \partial v_i(\mathbf{a})/\partial \mathbf{a} |_{\mathbf{a}=\hat{\mathbf{a}}_0} = \mathbf{0}$ . One can show (using Lagrange multipliers) that

$$p_i = \frac{1}{n} \frac{1}{1 + \lambda (v_i(\hat{\mathbf{a}}_0) - A_0)}, i = 1, \dots, n, \quad (4)$$

where the Lagrange multiplier  $\lambda$  is the root of

$$\frac{1}{n} \sum_{i=1}^n \frac{v_i(\hat{\mathbf{a}}_0)}{1 + \lambda (v_i(\hat{\mathbf{a}}_0) - A_0)} = A_0. \quad (5)$$

Under the alternative hypothesis  $H_a$ , we have just the constraint  $\sum_{i=1}^n p_i = 1$ , and hence  $L(A_0) = (1/n)^n$  at  $p_i = 1/n$ . Therefore, the empirical log-likelihood ratio test statistic is

$$l(A_0) = -2 \log ELR(A_0) = 2 \sum_{i=1}^n \log(1 + \lambda(v_i(\hat{\mathbf{a}}_0) - A_0)). \quad (6)$$

126 The objective of confidence interval construction based on the empirical likeli-  
 127 hood ratio  $l(A_0)$  is to include all values of  $A$  in the interval  $R_\alpha(A) = \{A : l(A) \leq c_\alpha\}$   
 128 for pre-specified  $\alpha$  that satisfy  $P(A_0 \in R_\alpha(A)) = 1 - \alpha$ . The value of the con-  
 129 stant  $c_\alpha > 0$  depends on the coverage probability of the interval. For future use,  
 130 we define  $A_{m,n}^K(\mathbf{a}) = \sum_{i=1}^n v_i(\mathbf{a})/n$  and  $\hat{\mathbf{a}}_K = \arg \max_{\mathbf{a}} A_{m,n}^K(\mathbf{a})$ . The following  
 131 proposition establishes the asymptotic distribution of the statistic  $l(A_0)$  that is  
 132 needed to construct the confidence interval for the maximum AUC.

**Proposition 1.** *Let  $\mathbf{X}_i$ ,  $i = 1, \dots, n$ , and  $\mathbf{Y}_j$ ,  $j = 1, \dots, m$ , be i.i.d. measurements for diseased subjects and nondiseased subjects, respectively. Assume that regularity conditions (C1)-(C7) presented in Appendix A are satisfied. Then, under  $H_0 : A = A_0$ , the asymptotic distribution of  $l(A_0)$  is a scaled Chi-square distribution with one degree of freedom, i.e.,*

$$\gamma(A_0)l(A_0) \xrightarrow{d} \chi_1^2, \text{ as } n, m \rightarrow \infty, \quad (7)$$

where

$$\gamma(A_0) = \frac{m\hat{\sigma}^2}{(m+n)s^2}, \hat{\sigma}^2 = \sum_{i=1}^n (v_i(\hat{\mathbf{a}}_K) - \sum_{j=1}^n v_j(\hat{\mathbf{a}}_K)/n)^2/n, s^2 = \frac{m\hat{\sigma}_{10}^2 + n\hat{\sigma}_{01}^2}{m+n},$$

$$\sigma_{10}^2 = \text{Cov}(K_h(\mathbf{a}_0^T \mathbf{X}_1 - \mathbf{a}_0^T \mathbf{Y}_1), K_h(\mathbf{a}_0^T \mathbf{X}_1 - \mathbf{a}_0^T \mathbf{Y}_2)),$$

$$\sigma_{01}^2 = \text{Cov}(K_h(\mathbf{a}_0^T \mathbf{X}_1 - \mathbf{a}_0^T \mathbf{Y}_1), K_h(\mathbf{a}_0^T \mathbf{X}_2 - \mathbf{a}_0^T \mathbf{Y}_1)),$$

133 and  $\hat{\sigma}_{10}^2$ ,  $\hat{\sigma}_{01}^2$  are the corresponding estimates.

134 *Proof.* It is outlined in the Appendix B. □

Thus, the  $100(1 - \alpha)\%$  EL confidence interval for the maximum AUC based on the BLC of multiple markers can be constructed as

$$R_\alpha(A_0) = \{A_0 : \gamma(A_0)l(A_0) \leq \chi_1^2(1 - \alpha)\},$$

135 where  $\chi_1^2(1 - \alpha)$  is the  $(1 - \alpha)$ th quantile of the Chi-square distribution with 1  
136 degree of freedom.

137 **Remark 1.** *The conditions that the kernel function needs to satisfy for the*  
138 *results of Proposition 1 to be true are reasonable and generally used in density*  
139 *estimation. An example of a second order kernel that satisfies these conditions*  
140 *is the Gaussian kernel. Here, the random vectors  $\mathbf{X}$  and  $\mathbf{Y}$  should satisfy  $E|\mathbf{X} -$*   
141  *$\mathbf{Y}|^t < \infty$  and  $\delta \in (1/2t, 1/4)$  for  $t > 2$ .*

142 We further discuss this point in Appendix C.

143 **Remark 2.** *Following Vealer et al. (2006), the estimators of the best linear*  
144 *coefficients  $\hat{\mathbf{a}}_K$  and the corresponding AUC estimator  $A_{m,n}^K(\hat{\mathbf{a}}_K)$  exist uniquely*  
145 *if (1) the distributions of biomarkers in the case and control group are both*  
146 *absolutely continuous; (2) all biomarkers have discriminatory abilities and there*  
147 *is one biomarker has a better discriminant ability than other markers and (3)*  
148 *the second derivative of  $A_{m,n}^K(\mathbf{a})$  exists and is nonzero.*

149 Assumption (2) can be satisfied via variable selection procedures such as  
150 stepwise selection, the best subset selection and LASSO (Tibshirani, 1996).  
151 Condition (C7) presented in Appendix A guarantees that assumption (3) holds.

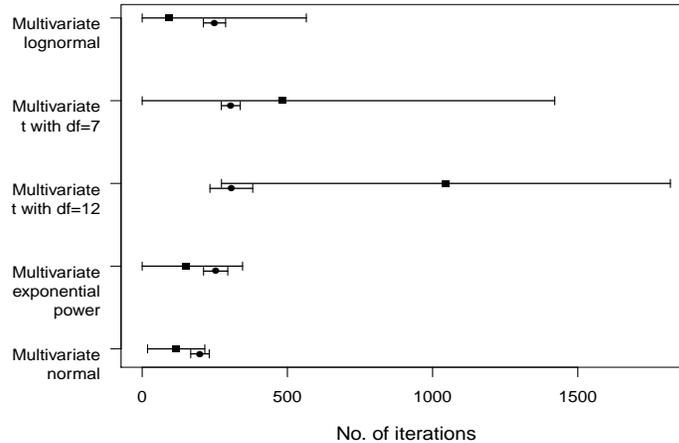
### 152 3. Monte Carlo Simulations

#### 153 3.1. Computational strategy

154 Computations are conducted using R (R Development Core Team, 2013).  
155 The EL log-likelihood ratio test statistic can be obtained using the function  
156 *el.test* in the *emplik* package. The maximization of the smooth AUC objective

157 function is carried out based on gradient-based methods such as the quasi-  
 158 Newton algorithm (Shanno, 1970) using the basic procedure *optim* in R. The  
 159 gradient of the kernel-based AUC objective function can be calculated in a  
 160 simple and fast manner. The related computer codes used to carry out the  
 161 numerical results presented in this paper are provided in the Supplementary  
 162 material.

Figure 1: Number of iterations (x-axis) of the maximization procedure with  $m = n = 100$  and  $d = 10$ . The y-axis presents the distributions in ascending order of their empirical skewness. The filled squares represent the number of iterations of the kernel-based AUC method and the filled circles represent the number of iterations used by the empirical AUC.



163 On the contrary, the optimization of the empirical AUC requires a brutal  
 164 search or the development of special algorithms. For example, the Nelder-Mead  
 165 method (Nelder and Mead, 1965) can be used, and it works reasonably well  
 166 for non-differentiable functions. For comparison purpose, the number of iterations  
 167 for the maximization of the empirical AUC and the kernel-based AUC are  
 168 calculated for different data distributions.

169 Figure 1 plots the median of the number of iterations (x-axis) each maxi-  
 170 mization requires versus ascending ordering of distributions according to their  
 171 average skewness coefficient (y-axis) over 500 simulations. The simulated data

172 sets for both the control group and the case group consist of  $d = 10$  biomark-  
 173 ers with  $m = n = 100$  and with underlying distributions being multivariate  
 174 normal distribution, multivariate exponential power distribution, multivariate t  
 175 distribution and multivariate lognormal distribution (Johnson, 2013). As Fig-  
 176 ure 1 illustrates, the kernel-based AUC requires a smaller number of iterations  
 177 for most of the selected distributions, indicating the computational advantage  
 178 of the proposed method. For details about the parameter settings and related  
 179 concepts, please see section 1 of the supplementary material.

### 180 3.2. Simulation studies

181 In this section, we report the simulation studies for evaluating coverage accu-  
 182 racy and interval length of the newly proposed EL interval for the optimal AUC  
 183 based on multiple biomarkers. The proposed smoothed EL-based confidence  
 184 interval is compared with the normality-based confidence intervals derived by  
 185 Reiser and Faraggi (1997).

186 To study the performance of our method, we computed the coverage prob-  
 187 ability and associated lengths of simulated confidence intervals, setting the  
 188 theoretical coverage probability equal to 95% at a nominal significance level  
 189  $\alpha = 0.05$ . We simulated the cases of two and four biomarkers with different un-  
 190 derlying distributions, including multivariate normal distribution, multivariate  
 191 lognormal distribution and multivariate power exponential distribution (John-  
 192 son, 2013). For future use, denote  $\Sigma = \{\sigma_{ij}\}$  where  $\sigma_{ii} = 1$  and  $\sigma_{ij} = \rho$  for  
 193  $i \neq j, i, j = 1, \dots, d$ .

194 In each study, we replicated the simulation experiment  $MC = 10,000$  times  
 195 so the simulation error defined as  $z_{1-\alpha/2}(\alpha(1-\alpha)/MC)^{1/2}$  is 0.0043, where  
 196  $z_{1-\alpha/2}$  is the  $100(1-\alpha/2)$ th quantile of a standard normal distribution. Con-  
 197 fidence intervals are calculated for values of AUC equal to 0.75, 0.85, 0.90 and  
 198 0.95 in balanced and unbalanced designs with both small samples and large  
 199 samples. We considered the commonly used Gaussian kernel and the optimal  
 200 Epanechnikov kernel which yields the minimum asymptotic mean-squared-error  
 201 for the estimation of the density function.

Table 1: Coverage probabilities and lengths (in parentheses) of the normality-based and EL-based confidence intervals (CI) with a 95% nominal level for various AUC values in the case of four biomarkers with a multivariate normal distribution and  $\rho = 0$ .

AUC ( $m, n$ )	Normality based CI	EL-based CI with Gaussian kernels			EL-based CI with Epanechnikov kernels		
		$\delta = 1/3$	$\delta = 1/5$	$\delta = 1/7$	$\delta = 1/3$	$\delta = 1/5$	$\delta = 1/7$
0.75							
(25,25)	0.951(0.214)	0.873(0.220)	0.892(0.217)	0.904(0.214)	0.891(0.216)	0.907(0.212)	0.916(0.208)
(75,25)	0.948(0.178)	0.924(0.193)	0.933(0.192)	0.937(0.189)	0.923(0.191)	0.939(0.187)	0.931(0.184)
(25,75)	0.950(0.178)	0.919(0.195)	0.928(0.194)	0.936(0.191)	0.930(0.194)	0.935(0.190)	0.948(0.186)
(75,75)	0.952(0.127)	0.932(0.144)	0.936(0.144)	0.941(0.142)	0.940(0.143)	0.947(0.141)	0.946(0.138)
(200,50)	0.950(0.124)	0.937(0.141)	0.940(0.141)	0.942(0.139)	0.940(0.141)	0.948(0.139)	0.942(0.136)
(200,100)	0.951(0.100)	0.942(0.111)	0.946(0.111)	0.948(0.110)	0.946(0.111)	0.948(0.109)	0.939(0.107)
(200,200)	0.950(0.086)	0.942(0.092)	0.944(0.092)	0.948(0.091)	0.947(0.092)	0.953(0.090)	0.938(0.089)
0.85							
(25,25)	0.951(0.187)	0.853(0.166)	0.874(0.167)	0.887(0.166)	0.874(0.166)	0.891(0.166)	0.905(0.165)
(75,25)	0.953(0.150)	0.911(0.148)	0.922(0.148)	0.929(0.148)	0.916(0.148)	0.913(0.146)	0.918(0.145)
(25,75)	0.950(0.151)	0.921(0.153)	0.927(0.153)	0.932(0.152)	0.933(0.153)	0.933(0.151)	0.931(0.150)
(75,75)	0.950(0.110)	0.931(0.113)	0.933(0.113)	0.935(0.112)	0.935(0.113)	0.940(0.112)	0.925(0.111)
(200,50)	0.948(0.107)	0.941(0.111)	0.943(0.111)	0.940(0.110)	0.937(0.111)	0.941(0.110)	0.940(0.109)
(200,100)	0.951(0.086)	0.942(0.087)	0.944(0.087)	0.941(0.087)	0.951(0.087)	0.950(0.087)	0.923(0.086)
(200,200)	0.950(0.072)	0.942(0.072)	0.945(0.072)	0.944(0.072)	0.948(0.072)	0.946(0.071)	0.917(0.071)
0.90							
(25,25)	0.954(0.157)	0.820(0.128)	0.848(0.130)	0.868(0.131)	0.854(0.131)	0.873(0.132)	0.883(0.132)
(75,25)	0.952(0.127)	0.890(0.116)	0.904(0.117)	0.912(0.117)	0.899(0.116)	0.912(0.117)	0.905(0.117)
(25,75)	0.948(0.127)	0.910(0.121)	0.919(0.122)	0.927(0.122)	0.922(0.122)	0.927(0.123)	0.914(0.122)
(75,75)	0.946(0.091)	0.924(0.090)	0.929(0.090)	0.930(0.009)	0.936(0.091)	0.931(0.091)	0.919(0.090)
(200,50)	0.947(0.088)	0.940(0.088)	0.943(0.088)	0.941(0.088)	0.940(0.088)	0.937(0.088)	0.927(0.088)
(200,100)	0.951(0.070)	0.939(0.070)	0.941(0.070)	0.945(0.070)	0.948(0.070)	0.942(0.070)	0.913(0.070)
(200,200)	0.950(0.058)	0.942(0.058)	0.944(0.058)	0.944(0.058)	0.952(0.058)	0.940(0.058)	0.890(0.058)
0.95							
(25,25)	0.951(0.112)	0.739(0.077)	0.788(0.082)	0.822(0.085)	0.786(0.082)	0.836(0.086)	0.840(0.091)
(75,25)	0.953(0.089)	0.844(0.071)	0.865(0.073)	0.878(0.075)	0.854(0.072)	0.886(0.076)	0.881(0.077)
(25,75)	0.953(0.088)	0.868(0.078)	0.890(0.080)	0.904(0.081)	0.888(0.080)	0.895(0.081)	0.891(0.083)
(75,75)	0.952(0.061)	0.906(0.058)	0.915(0.059)	0.915(0.060)	0.916(0.059)	0.919(0.060)	0.885(0.061)
(200,50)	0.951(0.060)	0.926(0.057)	0.932(0.058)	0.934(0.058)	0.930(0.058)	0.924(0.059)	0.902(0.060)
(200,100)	0.948(0.046)	0.933(0.046)	0.936(0.046)	0.934(0.047)	0.942(0.047)	0.924(0.047)	0.887(0.048)
(200,200)	0.949(0.038)	0.942(0.038)	0.944(0.038)	0.940(0.039)	0.943(0.038)	0.928(0.039)	0.869(0.039)

202 Choices of bandwidths have already been well studied in the density or  
 203 distribution function estimation literature (Lloyd, 1998; Lloyd and Yong, 1999;  
 204 Zou et al., 1997). In the simulations, we considered bandwidths of order  $h =$   
 205  $(n + m)^{-\delta}$  with  $\delta = 1/3, 1/5$  and  $1/7$ . For a second order kernel such as the  
 206 Gaussian kernel, it is easy to check that conditions (C1)-(C7) stated in Appendix  
 207 A are satisfied when  $\delta = 1/5$  and  $1/7$ . The case that  $\delta = 1/3$  shows the results  
 208 when the conditions are slightly violated.

209 Table 1 shows the simulation results for four biomarkers ( $d = 4$ ) follow-  
 210 ing multivariate normal distributions with  $\boldsymbol{\mu}_{\mathbf{X}} = (\mu, 0.5, 0, -0.2)^T$ ,  $\boldsymbol{\mu}_{\mathbf{Y}} =$   
 211  $(0.2, 0, 0.5, 0)^T$  and  $\boldsymbol{\Sigma}_{\mathbf{X}} = \boldsymbol{\Sigma}_{\mathbf{Y}} = \boldsymbol{\Sigma}$ , where  $\rho = 0$ . Here the covariance ma-  
 212 trices are assumed to be equal for simplicity but without loss of generality. The  
 213 value of  $\mu \in \mathbf{R}$  varies to reach pre-specified values of AUC based on the point  
 214 estimator of Su and Liu (1993). Table 1 suggest that the Gaussian kernel is  
 215 not very sensitive to the choice of bandwidths, yielding equivalent performance  
 216 in terms of achieved coverage probability and associated confidence interval  
 217 lengths. Furthermore, while one would have expected that, for normally dis-  
 218 tributed data, the Gaussian kernel would perform considerably better in terms  
 219 of coverage probability, our simulations indicate that the Epanechnikov kernel  
 220 shows equivalent performance across all values of bandwidths and sample sizes.  
 221 Same results can be observed for the case of two biomarkers. For details, please  
 222 see the supplementary materials. As shown in Tables 2 and 3, these results  
 223 are also confirmed for the case where  $\log \mathbf{X}$  and  $\log \mathbf{Y}$  are normally distributed  
 224 according to the aforementioned normal distributions, making the distribution  
 225 of the markers a lognormal distribution. However, as expected, normality based  
 226 confidence intervals have very low coverage probabilities.

227 In addition, the scenarios where the off-diagonal elements of  $\boldsymbol{\Sigma}$  are  $\rho =$   
 228  $0.4, 0.7$  for the aforementioned multivariate normal distribution and the multi-  
 229 variate lognormal distribution are also considered. Tables 1 through 8 of the  
 230 supplementary material present the results. Here we note that higher corre-  
 231 lations yield slightly smaller coverage probabilities but the proposed method  
 232 performs well even in skew data for moderate to large sample sizes.

Table 2: Coverage probabilities and lengths (in parentheses) of the normality-based and EL-based confidence intervals (CI) with a 95% nominal level for various AUC values in the case of two biomarkers with a multivariate lognormal distribution and  $\rho = 0$ .

AUC ( $m, n$ )	Normality based CI	EL-based CI with Gaussian kernels			EL-based CI with Epanechnikov kernels		
		$\delta = 1/3$	$\delta = 1/5$	$\delta = 1/7$	$\delta = 1/3$	$\delta = 1/5$	$\delta = 1/7$
0.75							
(25,25)	0.947(0.238)	0.913(0.232)	0.916(0.229)	0.917(0.227)	0.912(0.232)	0.917(0.229)	0.917(0.227)
(75,25)	0.916(0.216)	0.928(0.198)	0.931(0.196)	0.932(0.194)	0.928(0.198)	0.930(0.196)	0.932(0.194)
(25,75)	0.898(0.196)	0.936(0.199)	0.938(0.197)	0.939(0.195)	0.936(0.199)	0.938(0.197)	0.938(0.195)
(75,75)	0.817(0.160)	0.940(0.147)	0.940(0.145)	0.941(0.144)	0.941(0.147)	0.940(0.145)	0.940(0.144)
(200,50)	0.808(0.162)	0.945(0.143)	0.946(0.142)	0.945(0.141)	0.945(0.143)	0.946(0.142)	0.945(0.141)
(200,100)	0.657(0.127)	0.947(0.113)	0.948(0.112)	0.946(0.111)	0.947(0.113)	0.948(0.112)	0.946(0.111)
(200,200)	0.448(0.101)	0.948(0.093)	0.950(0.092)	0.948(0.091)	0.948(0.093)	0.950(0.092)	0.947(0.091)
0.85							
(25,25)	0.807(0.261)	0.893(0.177)	0.900(0.176)	0.904(0.176)	0.894(0.177)	0.901(0.176)	0.905(0.176)
(75,25)	0.758(0.245)	0.913(0.153)	0.916(0.152)	0.916(0.151)	0.913(0.153)	0.916(0.152)	0.916(0.151)
(25,75)	0.431(0.176)	0.930(0.157)	0.932(0.157)	0.934(0.157)	0.930(0.157)	0.933(0.157)	0.935(0.157)
(75,75)	0.274(0.154)	0.942(0.116)	0.945(0.115)	0.944(0.115)	0.942(0.116)	0.945(0.115)	0.944(0.115)
(200,50)	0.380(0.175)	0.947(0.113)	0.946(0.112)	0.942(0.111)	0.947(0.113)	0.946(0.112)	0.943(0.111)
(200,100)	0.085(0.129)	0.950(0.089)	0.947(0.089)	0.944(0.088)	0.950(0.089)	0.947(0.089)	0.944(0.088)
(200,200)	0.005(0.095)	0.946(0.073)	0.947(0.073)	0.948(0.073)	0.946(0.073)	0.947(0.073)	0.948(0.073)
0.90							
(25,25)	0.494(0.258)	0.870(0.137)	0.880(0.137)	0.884(0.138)	0.869(0.137)	0.880(0.138)	0.884(0.138)
(75,25)	0.431(0.249)	0.902(0.119)	0.905(0.119)	0.908(0.119)	0.903(0.119)	0.905(0.119)	0.908(0.119)
(25,75)	0.072(0.161)	0.919(0.125)	0.924(0.126)	0.924(0.126)	0.919(0.125)	0.924(0.126)	0.924(0.126)
(75,75)	0.024(0.150)	0.937(0.092)	0.937(0.092)	0.936(0.092)	0.937(0.092)	0.937(0.092)	0.937(0.092)
(200,50)	0.076(0.177)	0.942(0.089)	0.943(0.089)	0.942(0.089)	0.942(0.089)	0.943(0.089)	0.941(0.089)
(200,100)	0.001(0.128)	0.945(0.071)	0.943(0.071)	0.940(0.071)	0.945(0.071)	0.943(0.071)	0.940(0.071)
(200,200)	0.000(0.093)	0.947(0.059)	0.945(0.059)	0.942(0.059)	0.947(0.059)	0.945(0.059)	0.943(0.059)
0.95							
(25,25)	0.136(0.250)	0.814(0.084)	0.824(0.086)	0.833(0.087)	0.814(0.084)	0.824(0.086)	0.832(0.087)
(75,25)	0.100(0.248)	0.860(0.074)	0.866(0.075)	0.873(0.075)	0.862(0.074)	0.865(0.075)	0.873(0.075)
(25,75)	0.001(0.150)	0.883(0.081)	0.890(0.082)	0.896(0.083)	0.882(0.081)	0.890(0.082)	0.897(0.083)
(75,75)	0.000(0.147)	0.917(0.060)	0.920(0.060)	0.922(0.060)	0.917(0.060)	0.920(0.060)	0.923(0.060)
(200,50)	0.002(0.177)	0.928(0.058)	0.929(0.058)	0.930(0.058)	0.929(0.058)	0.929(0.058)	0.930(0.058)
(200,100)	0.000(0.127)	0.941(0.047)	0.942(0.047)	0.940(0.047)	0.941(0.047)	0.941(0.047)	0.939(0.047)
(200,200)	0.000(0.085)	0.942(0.039)	0.944(0.039)	0.941(0.039)	0.942(0.039)	0.944(0.039)	0.941(0.039)

Table 3: Coverage probabilities and lengths (in parentheses) of the normality-based and EL-based confidence intervals (CI) with a 95% nominal level for various AUC values in the case of four biomarkers with a multivariate lognormal distribution and  $\rho = 0$ .

AUC ( $m, n$ )	Normality based CI	EL-based CI with Gaussian kernels			EL-based CI with Epanechnikov kernels		
		$\delta = 1/3$	$\delta = 1/5$	$\delta = 1/7$	$\delta = 1/3$	$\delta = 1/5$	$\delta = 1/7$
0.75							
(25,25)	0.957(0.251)	0.87(0.219)	0.875(0.216)	0.881(0.214)	0.870(0.219)	0.877(0.216)	0.882(0.214)
(75,25)	0.920(0.200)	0.916(0.194)	0.92(0.192)	0.924(0.190)	0.916(0.194)	0.920(0.192)	0.925(0.190)
(25,75)	0.937(0.231)	0.912(0.192)	0.917(0.191)	0.923(0.19)	0.912(0.192)	0.919(0.191)	0.922(0.190)
(75,75)	0.855(0.161)	0.93(0.144)	0.934(0.143)	0.935(0.142)	0.931(0.144)	0.934(0.143)	0.934(0.142)
(200,50)	0.799(0.141)	0.939(0.143)	0.943(0.142)	0.943(0.14)	0.940(0.143)	0.942(0.142)	0.942(0.140)
(200,100)	0.686(0.117)	0.941(0.112)	0.945(0.111)	0.947(0.11)	0.941(0.112)	0.945(0.111)	0.948(0.110)
(200,200)	0.567(0.101)	0.939(0.092)	0.941(0.092)	0.946(0.091)	0.939(0.092)	0.941(0.092)	0.945(0.091)
0.85							
(25,25)	0.792(0.282)	0.850(0.162)	0.859(0.162)	0.866(0.162)	0.848(0.163)	0.862(0.163)	0.868(0.163)
(75,25)	0.474(0.183)	0.903(0.147)	0.908(0.146)	0.912(0.146)	0.902(0.147)	0.908(0.146)	0.913(0.146)
(25,75)	0.723(0.262)	0.914(0.150)	0.922(0.150)	0.925(0.150)	0.915(0.150)	0.922(0.150)	0.926(0.150)
(75,75)	0.278(0.156)	0.932(0.112)	0.933(0.112)	0.934(0.112)	0.932(0.112)	0.934(0.112)	0.934(0.112)
(200,50)	0.070(0.117)	0.943(0.111)	0.945(0.110)	0.945(0.110)	0.943(0.111)	0.944(0.110)	0.944(0.110)
(200,100)	0.019(0.103)	0.948(0.088)	0.948(0.087)	0.948(0.087)	0.948(0.088)	0.947(0.087)	0.948(0.087)
(200,200)	0.004(0.095)	0.945(0.072)	0.944(0.072)	0.941(0.072)	0.946(0.072)	0.943(0.072)	0.940(0.072)
0.90							
(25,25)	0.544(0.280)	0.808(0.124)	0.821(0.125)	0.833(0.126)	0.809(0.124)	0.823(0.126)	0.834(0.126)
(75,25)	0.114(0.166)	0.874(0.114)	0.882(0.114)	0.885(0.114)	0.875(0.114)	0.882(0.114)	0.884(0.114)
(25,75)	0.448(0.268)	0.893(0.118)	0.904(0.120)	0.910(0.120)	0.894(0.119)	0.906(0.120)	0.912(0.121)
(75,75)	0.040(0.152)	0.920(0.089)	0.924(0.089)	0.926(0.090)	0.919(0.089)	0.925(0.090)	0.926(0.090)
(200,50)	0.000(0.105)	0.932(0.088)	0.934(0.088)	0.936(0.088)	0.932(0.088)	0.934(0.088)	0.936(0.088)
(200,100)	0.000(0.097)	0.941(0.070)	0.941(0.070)	0.942(0.070)	0.941(0.070)	0.941(0.070)	0.942(0.070)
(200,200)	0.000(0.093)	0.942(0.058)	0.944(0.058)	0.944(0.058)	0.942(0.058)	0.944(0.058)	0.943(0.058)
0.95							
(25,25)	0.161(0.272)	0.744(0.076)	0.764(0.078)	0.779(0.08)	0.748(0.077)	0.765(0.079)	0.781(0.081)
(75,25)	0.002(0.153)	0.846(0.071)	0.851(0.072)	0.859(0.072)	0.846(0.071)	0.852(0.072)	0.858(0.072)
(25,75)	0.112(0.270)	0.861(0.077)	0.874(0.078)	0.883(0.079)	0.861(0.077)	0.875(0.078)	0.884(0.079)
(75,75)	0.000(0.149)	0.905(0.058)	0.910(0.059)	0.912(0.059)	0.905(0.058)	0.910(0.059)	0.912(0.059)
(200,50)	0.000(0.092)	0.925(0.057)	0.926(0.058)	0.926(0.058)	0.926(0.058)	0.926(0.058)	0.927(0.058)
(200,100)	0.000(0.089)	0.935(0.046)	0.935(0.047)	0.933(0.047)	0.935(0.046)	0.935(0.047)	0.934(0.047)
(200,200)	0.000(0.087)	0.942(0.039)	0.941(0.039)	0.939(0.039)	0.943(0.039)	0.940(0.039)	0.939(0.039)

Table 4: Coverage probabilities and lengths (in parentheses) of the normality-based and EL-based confidence intervals (CI) with a 95% nominal level for various AUC values in the case of two biomarkers with marginal Chi-square distributions coupled together via a normal copula with correlation  $\rho = 0.7$ .

AUC ( $m, n$ )	Normality based CI	EL-based CI with Gaussian kernels			EL-based CI with Epanechnikov kernels		
		$\delta = 1/3$	$\delta = 1/5$	$\delta = 1/7$	$\delta = 1/3$	$\delta = 1/5$	$\delta = 1/7$
0.75							
(25,25)	0.840(0.163)	0.899(0.226)	0.900(0.226)	0.899(0.226)	0.899(0.226)	0.900(0.226)	0.900(0.226)
(75,25)	0.832(0.155)	0.941(0.201)	0.941(0.201)	0.942(0.201)	0.940(0.201)	0.941(0.201)	0.942(0.201)
(25,75)	0.642(0.105)	0.918(0.191)	0.918(0.191)	0.918(0.191)	0.917(0.191)	0.918(0.191)	0.918(0.191)
(75,75)	0.532(0.088)	0.947(0.146)	0.947(0.146)	0.947(0.146)	0.947(0.146)	0.947(0.146)	0.947(0.146)
(200,50)	0.636(0.107)	0.941(0.148)	0.941(0.148)	0.941(0.148)	0.941(0.148)	0.941(0.148)	0.941(0.148)
(200,100)	0.361(0.051)	0.942(0.115)	0.942(0.115)	0.942(0.115)	0.942(0.115)	0.942(0.115)	0.942(0.115)
(200,200)	0.052(0.001)	0.945(0.092)	0.945(0.092)	0.946(0.092)	0.945(0.092)	0.945(0.092)	0.946(0.092)
0.85							
(25,25)	0.840(0.163)	0.853(0.166)	0.854(0.166)	0.855(0.166)	0.853(0.166)	0.855(0.166)	0.855(0.166)
(75,25)	0.832(0.155)	0.931(0.158)	0.931(0.158)	0.931(0.158)	0.931(0.158)	0.931(0.158)	0.932(0.158)
(25,75)	0.642(0.105)	0.860(0.140)	0.861(0.140)	0.860(0.140)	0.860(0.140)	0.861(0.140)	0.861(0.140)
(75,75)	0.532(0.088)	0.936(0.115)	0.935(0.115)	0.935(0.115)	0.935(0.115)	0.935(0.115)	0.936(0.115)
(200,50)	0.636(0.107)	0.949(0.123)	0.950(0.123)	0.950(0.123)	0.950(0.123)	0.950(0.123)	0.950(0.123)
(200,100)	0.361(0.051)	0.951(0.096)	0.951(0.096)	0.951(0.096)	0.951(0.096)	0.951(0.096)	0.951(0.096)
(200,200)	0.052(0.001)	0.951(0.074)	0.951(0.074)	0.951(0.074)	0.951(0.074)	0.951(0.074)	0.951(0.074)
0.90							
(25,25)	0.840(0.163)	0.845(0.133)	0.844(0.132)	0.845(0.132)	0.846(0.133)	0.844(0.133)	0.845(0.132)
(75,25)	0.832(0.155)	0.920(0.123)	0.919(0.123)	0.919(0.123)	0.920(0.123)	0.920(0.123)	0.920(0.123)
(25,75)	0.642(0.105)	0.884(0.118)	0.885(0.118)	0.884(0.118)	0.885(0.118)	0.885(0.118)	0.884(0.118)
(75,75)	0.532(0.088)	0.932(0.092)	0.932(0.092)	0.933(0.092)	0.932(0.092)	0.933(0.092)	0.933(0.092)
(200,50)	0.636(0.107)	0.947(0.094)	0.948(0.094)	0.948(0.094)	0.947(0.094)	0.947(0.094)	0.948(0.094)
(200,100)	0.361(0.051)	0.949(0.073)	0.949(0.073)	0.949(0.073)	0.949(0.073)	0.949(0.073)	0.949(0.073)
(200,200)	0.052(0.001)	0.950(0.059)	0.951(0.059)	0.950(0.059)	0.950(0.059)	0.950(0.059)	0.950(0.059)
0.95							
(25,25)	0.840(0.163)	0.782(0.079)	0.782(0.080)	0.781(0.080)	0.784(0.080)	0.783(0.080)	0.783(0.080)
(75,25)	0.832(0.155)	0.905(0.078)	0.905(0.078)	0.905(0.078)	0.905(0.078)	0.905(0.078)	0.905(0.078)
(25,75)	0.642(0.105)	0.818(0.072)	0.819(0.072)	0.819(0.072)	0.818(0.072)	0.819(0.072)	0.818(0.072)
(75,75)	0.532(0.088)	0.909(0.059)	0.909(0.059)	0.909(0.059)	0.909(0.059)	0.910(0.059)	0.909(0.059)
(200,50)	0.636(0.107)	0.940(0.063)	0.940(0.063)	0.940(0.063)	0.940(0.063)	0.940(0.063)	0.940(0.063)
(200,100)	0.361(0.051)	0.940(0.050)	0.940(0.050)	0.940(0.050)	0.940(0.050)	0.940(0.050)	0.940(0.050)
(200,200)	0.052(0.001)	0.943(0.039)	0.943(0.039)	0.943(0.039)	0.943(0.039)	0.943(0.039)	0.943(0.039)

233 Furthermore, we constructed a bivariate distribution whose marginals are  
234 Chi-square distributions, coupled together via a normal copula with correlation  
235  $\rho$ . The marginal distributions of the control group are Chi-square distributions  
236 with degrees of freedom 2 and 1, respectively, and they are coupled together via  
237 a normal copula with  $\rho = 0.4$ . For the case group, the degrees of freedom of the  
238 marginal Chi-square distribution vary to reach pre-specified values of the AUC,  
239 and the scenarios  $\rho = 0, 0.4, 0.7$  are considered. Table 4 shows the simulation  
240 results for the scenario where  $\rho = 0.7$ . Tables 10 and 11 of the supplementary  
241 material present the results for the scenario where  $\rho = 0$  and 0.4, respectively.  
242 Similar to the multivariate lognormal scenario, the proposed method works well,  
243 whereas the method based on the normal assumptions fails.

244 The family of multivariate power exponential distribution is also investi-  
245 gated. We considered the case  $\kappa = 1$  and  $\Sigma_{\mathbf{X}} = \Sigma_{\mathbf{Y}} = \Sigma$  where  $\rho = 0, 0.4, 0.7$ .  
246 Note that the multivariate normal distribution is also included in the family of  
247 multivariate power exponential distribution, corresponding to  $\kappa = 2$ . Detailed  
248 simulation results are shown in the Supplementary materials (see tables 12 to  
249 17). The extensive Monte Carlo studies confirm the excellent performance and  
250 need for the proposed method over normality based confidence intervals when  
251 the data distribution is skewed.

#### 252 4. Data Example

253 We illustrate the application of the proposed EL interval for AUC based on  
254 a linear combination of biomarkers in the study of the accuracy of biomarkers  
255 related to atherosclerotic coronary heart disease. In the pathogenesis of vari-  
256 ous diseases including atherosclerosis, oxygen radicals play an important role.  
257 Different biomarkers of individual oxidative stress and antioxidant status quan-  
258 tify different phases of the oxidative stress and antioxidant status process of an  
259 individual.

260 The study sample consists of 105 individuals with myocardial infarction and  
261 437 controls. The sampling frame for adults between the ages of 35 and 65

262 was the New York State Department of Motor Vehicles drivers license rolls.  
 263 Also a randomly selected elderly sample (age 65 – 79) from the Health Care  
 264 Financing Administration database was taken (Schisterman et al., 2001). A  
 265 number of biomarkers were examined from a 12-hour fasting blood specimen for  
 266 biochemical analysis at baseline.

267 *4.1. Uric acid, TBARS and HDL cholesterol*

To improve diagnostic accuracy, we combine three important biomarkers related to the atherosclerotic coronary heart disease in an efficient manner. The measurements are uric acid, TBARS and HDL cholesterol. The p-value of the Shapiro-Wilk test for multivariate normality for the case group is  $2.1 \times 10^{-15}$  and for the control group is  $7.6 \times 10^{-27}$ , indicating the non-normal character of the data. For these three markers, the estimated means and variances for the diseased and the nondiseased are

$$\hat{\boldsymbol{\mu}}_{\mathbf{X}} = (6.106 \ 1.385 \ 45.655)^T, \hat{\boldsymbol{\mu}}_{\mathbf{Y}} = (5.516 \ 1.372 \ 52.327)^T,$$

$$\hat{\boldsymbol{\Sigma}}_{\mathbf{X}} = \begin{pmatrix} 2.878 & 0.021 & -8.592 \\ 0.021 & 0.251 & -0.316 \\ -8.592 & -0.316 & 162.957 \end{pmatrix}, \text{ and } \hat{\boldsymbol{\Sigma}}_{\mathbf{Y}} = \begin{pmatrix} 1.917 & 0.007 & -6.715 \\ 0.007 & 0.156 & -1.322 \\ -6.715 & -1.322 & 218.930 \end{pmatrix},$$

268 respectively.

269 The best linear combination estimates derived by the normality-based method  
 270 of Su and Liu (1993)  $\hat{\mathbf{a}}_N$  is uric acid  $-0.416 \times$  TBARS  $-0.189 \times$  HDL cholesterol,  
 271 with a maximum AUC estimate of 0.6467. The normality based 95% confidence  
 272 interval of Reiser and Faraggi (1997) for the maximum AUC is [0.583, 0.695]  
 273 with a length of 0.112. Recall that the the Shapiro-Wilk test does not justify the  
 274 normality assumption. Our method yields the best linear combination estimates  
 275 of uric acid  $+0.542 \times$  TBARS  $-0.199 \times$  HDL cholesterol. It gives a maximum  
 276 AUC estimate of 0.6520, and the smooth EL-based 95% confidence interval for  
 277 the maximum AUC is [0.594, 0.705] with a length of 0.111. Compared to the  
 278 normality-based method, the proposed method yields similar numerical results  
 279 for all practical purposes. However, judging from the BLC estimates, the pro-

280 posed method yields a more realistic result than the normality based method.  
 281 It is well-known that low levels of HDL-cholesterol and high levels of TBARS  
 282 as well as uric acid indicate higher risk of coronary heart disease (Schisterman  
 283 et al., 2002). Different from the normality based result, the signs of the point  
 284 BLC estimates by the proposed method confirm the negative correlation between  
 285 HDL-cholesterol and the risk of the disease as well as the positive correlation  
 286 between TBARS and uric acid and the risk of the disease.

To explain the numerical results obtained above and to evaluate the obtained confidence interval estimators, we fixed the intervals  $[0.583, 0.695]$  and  $[0.594, 0.705]$ , and bootstrap 1,000 times from the original case and control data sets with the same sample sizes as the original case and control data, calculating the empirical estimator

$$\tilde{\theta}_{n,m} = \underset{a_1, a_2}{\operatorname{argmax}} \frac{1}{nm} \sum_{i=1}^n \sum_{j=1}^m I \{ \text{uric acid}_i + a_1 \text{TBARS}_i + a_2 \text{HDL cholesterol}_i > \text{uric acid}_j + a_1 \text{TBARS}_j + a_2 \text{HDL cholesterol}_j \},$$

287 where  $n = 105$  and  $m = 437$ . The proportion of the event that the proposed CI  
 288 contains  $\tilde{\theta}_{105,437}$  was observed to be 0.943, whereas the normality-based Reiser  
 289 and Faraggi (1997)'s method provided a frequency of 0.890. This proportion is  
 290 expected to be 95%.

291 Additionally, in order to evaluate the proposed confidence interval estimation  
 292 versus Reiser and Faraggi (1997)'s approach, we conduct the following cross-  
 293 validation type study. In each stage of this study, we randomly splitted the  
 294 original data sets into two parts. One with fixed samples sizes, consists of  
 295 40 observations for the case group and 177 observations for the control group,  
 296 and the proposed confidence interval estimator and Reiser and Faraggi (1997)'s  
 297 confidence interval estimator are obtained. The second part of data sets consists  
 298 of the remaining 65 observations for the case group and 260 observations for  
 299 the control group. The empirical estimator  $\tilde{\theta}_{65,260}$  is calculated based on the  
 300 second part of data sets. Then we checked the event that  $\tilde{\theta}_{65,260}$  belongs to the  
 301 corresponding confidence intervals. This stage was repeated in a cross-validation

302 manner 1,000 times. The frequency of  $\tilde{\theta}_{65,260}$  belonging to the proposed CI  
 303 was observed to be 0.945, whereas for Reiser and Faraggi (1997)'s method,  
 304 we observed the frequency as 0.890. This proportion is expected to be 95%.  
 305 Thus, despite almost numerical equivalence in the estimation of the AUC and  
 306 confidence intervals, the problem of the Reiser and Faraggi (1997)'s application  
 307 may be described by the invalidation of the normal assumptions.

308 *4.2. Blood glucose, vitamin E and body mass index*

For one additional simple example, we randomly selected the data with 25  
 observations in the case group and 100 observations in the control group, and  
 evaluate markers including blood glucose, vitamin E (a fat-soluble antioxidant  
 vitamin) and body mass index (BMI). For these three markers, the estimated  
 means and variances for the diseased and the nondiseased are

$$\hat{\boldsymbol{\mu}}_{\mathbf{X}} = (111.400 \ 13.806 \ 27.954)^T, \hat{\boldsymbol{\mu}}_{\mathbf{Y}} = (109.260 \ 13.695 \ 27.947)^T,$$

$$\hat{\boldsymbol{\Sigma}}_{\mathbf{X}} = \begin{pmatrix} 680.583 & -7.781 & 4.815 \\ -7.781 & 28.975 & 3.652 \\ 4.815 & 3.652 & 14.771 \end{pmatrix}, \text{ and } \hat{\boldsymbol{\Sigma}}_{\mathbf{Y}} = \begin{pmatrix} 2184.881 & 4.284 & 50.747 \\ 4.284 & 30.417 & 3.469 \\ 50.747 & 3.469 & 24.169 \end{pmatrix},$$

309 respectively.

310 The Shapiro-Wilk test for multivariate normality applied to data from the  
 311 case group and control group rejects normality with p-values of  $1.1 \times 10^{-5}$   
 312 and  $6.7 \times 10^{-28}$ , respectively. Based on the normality assumption that is not  
 313 justified by the Shapiro-Wilk multivariate normality test, the maximum AUC  
 314 estimate is 0.517, whereas the maximum AUC estimate by the proposed method  
 315 is 0.659. Though the proposed method is not focused on the point estimator  
 316 of the maximum AUC, it yields a much higher maximum AUC than Su and  
 317 Liu (1993)'s estimate. The 95% confidence interval based on the normality as-  
 318 sumption (Reiser and Faraggi, 1997) is [0.500, 0.539], and the proposed smooth  
 319 EL-based 95% confidence interval for the maximum AUC is [0.533, 0.762]. In  
 320 addition, the best linear combination estimates derived by Su and Liu (1993)  
 321 is glucose + 2.675  $\times$  vitamin E - 1.681  $\times$  BMI, whereas the BLC derived by the

322 proposed method is  $\text{glucose} + 0.098 \times \text{vitamin E} + 0.351 \times \text{BMI}$ . The sign of BMI  
323 in the BLC estimates between the proposed method and Su and Liu (1993)'s  
324 method is reversed. It is shown that higher BMI measures are associated with a  
325 higher risk of a heart disease (Lamon-Fava et al., 1996), indicating that a neg-  
326 ative coefficient associated with BMI goes against scientific findings. Judging  
327 from the BLC estimates, the proposed method yields a realistic result that is  
328 not provided by the use of the normality-based methods.

329 **Remark 3.** *A natural question pertains to the use of all markers to compute the*  
330 *AUC and associated confidence interval. This aspect of the problem is related*  
331 *to biomarker selection and, given the mathematical challenges associated with it*  
332 *constitutes a topic of our future work.*

## 333 5. Discussion

334 In this paper, we have proposed an EL-based confidence interval for the max-  
335 imum area under the ROC curve based on a linear combination of biomarkers  
336 measured on a continuous scale. We show that the proposed method has good  
337 theoretical properties regardless of the data distribution. Simulation results  
338 indicate the proposed EL-based interval has a similar performance with nor-  
339 mality based confidence intervals when normality is present, and outperforms  
340 normality based confidence intervals in the case of skewed data.

341 Future work may focus on approaches to extend the proposed method to cat-  
342 egorical biomarkers. Since empirical likelihood regions are Bartlett correctable,  
343 additional future work may consider the Bartlett correction to reduce the cov-  
344 erage error of EL confidence intervals.

## 345 Supplementary Materials

346 The supplementary material includes some details considered in the compu-  
347 tational strategy, simulation results reporting the coverage probability and the  
348 length of the confidence intervals based on the proposed method for a variety of

349 distributions. Furthermore, the R code for obtaining the EL-based confidence  
 350 interval for the maximum AUC based on multiple biomarkers is provided.

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 354 significant improvements to the paper.

## 355 Appendix

### 356 Appendix A. Conditions

357 Throughout the paper, we assume the following:

(C1) Let  $h = h_{n,m} > 0$  be a bandwidth sequence converging to zero as  $n$   
 and  $m$  converge to infinity and  $k(\cdot)$  is a compactly supported  $r$ th order kernel  
 ( $r \geq 2$ ). That is, for some integer  $r \geq 2$  and a finite constant  $\xi \neq 0$ ,

$$\int u^j k(u) du = \begin{cases} 1, & j = 0, \\ 0, & j \in [1, r-1], \\ \xi, & j = r. \end{cases}$$

358 (C2) The kernel function  $k(u)$  is symmetric about  $u = 0$  with the first  
 359 derivative being bounded and continuous.

360 (C3) For the density function  $g_{\mathbf{a}}(x) := \partial G_{\mathbf{a}} / \partial x$ ,  $g_{\mathbf{a}}^{(r-1)}(x)$  exists and is  
 361 continuous.

362 (C4) As  $m$  and  $n$  converge to infinity,  $m/(m+n) \rightarrow q$ , where  $0 < q < 1$ .

363 (C5) (i) The true parameter value  $\mathbf{a}_0$  is an interior point of  $\Lambda$ , which is a  
 364 compactor subset of  $\mathbf{R}^d$ ; (ii) The partial derivative of the marginal density of  
 365  $\mathbf{a}_0^T(\mathbf{X}^T, \mathbf{Y}^T)^T$  exists and is bounded.

366 (C6)  $E|\mathbf{X} - \mathbf{Y}|^2 < \infty$  and  $(m+n)^{1/2}h^2 \rightarrow \infty$  as  $m, n \rightarrow \infty$ .

367 (C7) There exist  $t > 2$  such that  $(m+n)^{1/2}h^t \rightarrow 0$ ,  $E|\mathbf{X} - \mathbf{Y}|^t < \infty$ , the  $t$ th  
 368 derivative  $K^{(t)}$  exists and is continuous with the  $l$ th derivative  $|K^{(l)}(u)| \leq c$  for  
 369 all  $l \leq t$ .

370 The kernel function given in (C1) is commonly used in nonparametric density  
 371 estimation. Condition (C2) allows the corresponding scaled symmetric distri-  
 372 bution function as an approximation to the indicator function. Condition (C3)  
 373 requires that the density of the combined scores in the nondiseased group be  
 374 sufficiently smooth. Condition (C4) states the growth rate of two sample sizes  
 375 should be of equal order. Condition (C5) is needed for establishing the consis-  
 376 tency of  $\hat{\mathbf{a}}_K$  and the uniform convergence of  $A_{m,n}^K(\mathbf{a})$  to its population version.  
 377 Condition (C6) ensures that  $A_{m,n}^K(\hat{\mathbf{a}}_K) - A_{m,n}^K(\mathbf{a}_0)$  converges to 0 at the rate of  
 378 at least  $(m+n)^{-1/2}$ . Condition (C7) guarantees that  $EA_{m,n}^K(\mathbf{a}_0) - A_0$  converge  
 379 to 0 at the rate of at least  $(m+n)^{-1/2}$ .

## 380 Appendix B. Proofs

381 The proof of Proposition 1 will be based on the following steps: (1) Lemma 1  
 382 gives the derivation of  $l(A_0)$  with respect to  $A_0$  up to order 3. The first and  
 383 second derivative show the equivalence between  $A_{m,n}^K(\hat{\mathbf{a}}_K)$  and  $A_0$ , as well as  
 384  $\hat{\mathbf{a}}_0 = \hat{\mathbf{a}}_K$ ; (2) we will show  $\hat{\mathbf{a}}_K$  converges to  $\mathbf{a}_0$  at the rate of at least  $(m+n)^{-1/2}$   
 385 in Lemma 2 and  $A_{m,n}^K(\hat{\mathbf{a}}_K) - A_{m,n}^K(\mathbf{a}_0)$  converges to 0 at the rate of at least  
 386  $(m+n)^{-1/2}$  with a proper choice of bandwidths in Lemma 3; (3) the asymptotic  
 387 distribution of  $A_{m,n}^K(\mathbf{a}_0) - EA_{m,n}^K(\mathbf{a}_0)$  is shown in Lemma 4 based on the theory  
 388 of U-statistics.

389 **Lemma 1.** *The maximum smoothed empirical likelihood estimator (MELE) of*  
 390 *the maximum AUC is  $A_{m,n}^K(\hat{\mathbf{a}}_K)$ . At the smoothed MELE  $A_{m,n}^K(\hat{\mathbf{a}}_K)$ , we have*  
 391  $l(A_0)|_{A_0=A_{m,n}^K(\hat{\mathbf{a}}_K)} = l'(A_0)|_{A_0=A_{m,n}^K(\hat{\mathbf{a}}_K)} = 0$ ,  $l''(A_0)|_{A_0=A_{m,n}^K(\hat{\mathbf{a}}_K)} = 2n\hat{\sigma}^{-2}$ ,  
 392  $l'''(A_0)|_{A_0=\tilde{A}} = O(n)$ , *where  $\hat{\sigma}^2$  is stated in the Proposition 1, and  $\tilde{A}$  is some*  
 393 *real number between  $A_{m,n}^K(\hat{\mathbf{a}}_K)$  and  $A_0$ .*

*Proof.* By the constraints defined in equation (3), the definition of  $\hat{\mathbf{a}}_0$ , and  
 equation (6), we have

$$l'(A_0) = 2n \frac{\partial \lambda}{\partial A_0} \sum_{i=1}^n p_i (v_i(\hat{\mathbf{a}}_0) - A_0) + 2n\lambda \sum_{i=1}^n p_i \frac{\partial v_i(\hat{\mathbf{a}}_0)}{\partial \hat{\mathbf{a}}_0^T} \frac{\partial \hat{\mathbf{a}}_0}{\partial A_0} - 2n\lambda = -2n\lambda.$$

Since the MELE minimizes  $l(A_0)$ , we have  $\lambda = 0$  and  $p_i = n^{-1}$  by equation (4). Therefore, by equation (5), the MELE of the maximum AUC is  $A_{m,n}^K(\hat{\mathbf{a}}_K)$  and the corresponding  $\hat{\mathbf{a}}_0$  is equivalent to  $\hat{\mathbf{a}}_K$ . Therefore, we have

$$l(A_0)|_{A_0=A_{m,n}^K(\hat{\mathbf{a}}_K)} = l'(A_0)|_{A_0=A_{m,n}^K(\hat{\mathbf{a}}_K)} = 0.$$

To derive the second derivative of  $l(A_0)$  with respect to  $A_0$  at the MELE  $A_{m,n}^K(\hat{\mathbf{a}}_K)$ , we take the first derivative of both sides of the equation (5) with respect to  $A_0$ . It leads to

$$-n\partial\lambda/\partial A_0 \sum_{i=1}^n p_i^2 (v_i(\hat{\mathbf{a}}_0) - A_0)^2 - n\lambda \sum_{i=1}^n p_i^2 \partial v_i(\hat{\mathbf{a}}_0)/\partial A_0 (v_i(\hat{\mathbf{a}}_0) - A_0) = 1$$

At the MELE  $A_{m,n}^K(\hat{\mathbf{a}}_K)$ , we have

$$\partial\lambda/\partial A_0|_{A_0=A_{m,n}^K(\hat{\mathbf{a}}_K)} = -n / \sum_{i=1}^n [v_i(\hat{\mathbf{a}}_K) - A_0]^2 = -\hat{\sigma}^{-2}.$$

394 The second derivative  $l''(A_0)|_{A_0=A_{m,n}^K(\hat{\mathbf{a}}_K)} = 2n\hat{\sigma}^{-2} > 0$  confirms that the  
 395 MELE minimizes  $l(A_0)$ . Based on the fact  $\max_{1 \leq i \leq n} |v_i(\hat{\mathbf{a}}_0)| = O(1)$ , we have  
 396  $l'''(A_0)|_{A_0=\tilde{A}} = -2n\partial^2\lambda/\partial A^2|_{A_0=\tilde{A}} = O(n)$  for some real number  $\tilde{A}$  between  
 397  $A_{m,n}^K(\hat{\mathbf{a}}_K)$  and  $A_0$ . □

398 **Lemma 2.** *Suppose (C1)-(C3) and (C5) hold. Then  $\hat{\mathbf{a}}_K - \mathbf{a}_0 = O_p((m+n)^{1/2}\mathbf{1})$ ,*  
 399 *as  $m, n \rightarrow \infty$ .*

400 *Proof.* This lemma follows from Theorem 2 of Ma and Huang (Biometrics,  
 401 2007). □

**Lemma 3.** *Suppose (C1)-(C3) and (C6) hold. Then*

$$\sqrt{m+n} (A_{m,n}^K(\hat{\mathbf{a}}_K) - A_{m,n}^K(\mathbf{a}_0)) \rightarrow 0, \text{ as } m, n \rightarrow \infty.$$

*Proof.* The first derivative of  $A_{m,n}^K(\mathbf{a})$  with respect to  $\mathbf{a}$  at  $\mathbf{a} = \hat{\mathbf{a}}_K$  is 0 and  $K(\cdot)$  has continuous second partial derivatives. Apply the Taylor expansion of  $A_{m,n}^K(\mathbf{a}_0)$  at  $\hat{\mathbf{a}}_K$  to obtain

$$A_{m,n}^K(\mathbf{a}_0) = A_{m,n}^K(\hat{\mathbf{a}}_K) + (\mathbf{a}_0 - \hat{\mathbf{a}}_K)^T \partial A_{m,n}^K(\mathbf{a})/\partial \mathbf{a}|_{\mathbf{a}=\hat{\mathbf{a}}_K} + R_{m,n} = A_{m,n}^K(\hat{\mathbf{a}}_K) + R_{m,n},$$

where  $R_{m,n} = \frac{1}{2}(\hat{\mathbf{a}}_K - \mathbf{a}_0)^T \frac{\partial^2 A_{m,n}^K(\mathbf{a})}{\partial \mathbf{a} \partial \mathbf{a}^T} \Big|_{\mathbf{a}=\tilde{\mathbf{a}}} (\hat{\mathbf{a}}_K - \mathbf{a}_0)$  with  $\tilde{\mathbf{a}} \in (\mathbf{a}_0, \hat{\mathbf{a}}_K)$ . By (C1)-(C3) and (C6), we have

$$\begin{aligned} \left| \frac{\partial^2 A_{m,n}^K(\mathbf{a})}{\partial \mathbf{a} \partial \mathbf{a}^T} \Big|_{\mathbf{a}=\tilde{\mathbf{a}}} \right| &\leq \frac{1}{nm} \sum_{i=1}^n \sum_{j=1}^m \left| k' \left( \frac{\tilde{\mathbf{a}}^T (\mathbf{X}_i - \mathbf{Y}_j)}{h} \right) \right| \left| \frac{\mathbf{X}_i - \mathbf{Y}_j}{h} \right|^2 \\ &\leq \frac{c}{nm} \sum_{i=1}^n \sum_{j=1}^m \left| \frac{\mathbf{X}_i - \mathbf{Y}_j}{h} \right|^2. \end{aligned}$$

402 In addition, Lemma 1 states that  $\hat{\mathbf{a}}_K - \mathbf{a}_0 = O_p((m+n)^{-1/2} \mathbf{1})$  and thus  
 403  $\sqrt{m+n} R_{m,n} = O_p((m+n)^{-1/2} h^{-2})$ . Lemma 3 follows when  $(m+n)^{1/2} h^2 \rightarrow$   
 404  $\infty$  as  $m, n \rightarrow \infty$ .  $\square$

**Lemma 4.** *Suppose that (C4) holds, then*

$$\sqrt{\frac{mn}{m+n}} \frac{A_{m,n}^K(\mathbf{a}_0) - \mathbb{E} A_{m,n}^K(\mathbf{a}_0)}{s} \xrightarrow{d} N(0, 1),$$

405 where  $s$  is defined in Proposition 1.

*Proof.* Let  $\psi(x, y) = K_h(x - y)$ . Based on the theory of U statistics (Serfling, 2009), we have

$$\frac{\sqrt{m+n}(A_{m,n}^K(\mathbf{a}_0) - \mathbb{E} A_{m,n}^K(\mathbf{a}_0))}{(\sigma_{10}^2/(1-q) + \sigma_{01}^2/q)^{1/2}} \xrightarrow{d} N(0, 1),$$

406 and therefore Lemma 4 holds.  $\square$

**Proof of Proposition 1.** Apply Taylor expansion of  $l(A_0)$  at  $A_0 = A_{m,n}^K(\hat{\mathbf{a}}_K)$ . Based on the first and the second derivative of  $l(A_0)$  as shown in Lemma 1, we have

$$\begin{aligned} l(A_0) &= l(A_{m,n}^K(\hat{\mathbf{a}}_K)) + (A_0 - A_{m,n}^K(\hat{\mathbf{a}}_K)) l'(A_{m,n}^K(\hat{\mathbf{a}}_K)) \\ &\quad + (A_0 - A_{m,n}^K(\hat{\mathbf{a}}_K))^2 l''(A_{m,n}^K(\hat{\mathbf{a}}_K))/2 + r_{m,n} \\ &= (A_{m,n}^K(\hat{\mathbf{a}}_K) - A_0)^2 n \hat{\sigma}^{-2} + r_{m,n}, \end{aligned}$$

407 where  $\hat{\sigma}^2$  is defined in Proposition 1, and  $r_{m,n} = O_p((A_0 - A_{m,n}^K(\hat{\mathbf{a}}_K))^3 l'''(A_0)|_{A_0=\tilde{A}})$

408 for some real number  $\tilde{A}$  between  $A_{m,n}^K(\hat{\mathbf{a}}_K)$  and  $A_0$ .

To derive the asymptotic distribution of  $A_{m,n}^K(\hat{\mathbf{a}}_K) - A_0$ , we use the following decomposition,

$$A_{m,n}^K(\hat{\mathbf{a}}_K) - A_0 = (A_{m,n}^K(\hat{\mathbf{a}}_K) - A_{m,n}^K(\mathbf{a}_0)) + (A_{m,n}^K(\mathbf{a}_0) - \mathbb{E} A_{m,n}^K(\mathbf{a}_0)) + (\mathbb{E} A_{m,n}^K(\mathbf{a}_0) - A_0).$$

Apply Taylor theorem of  $K_h(u - Y(\mathbf{a}_0))$  around  $h = 0$  to the  $t$ th order. Under assumption (C7), we have

$$\begin{aligned} |K_h(u - Y(\mathbf{a}_0)) - 1| I \{u - Y(\mathbf{a}_0) > 0\} &= h^t \left| K_{\theta_1 h}^{(t)}(u - Y(\mathbf{a}_0)) \right| I \{u - Y(\mathbf{a}_0) > 0\} \leq h^t c, \\ |K_h(u - Y(\mathbf{a}_0))| I \{u - Y(\mathbf{a}_0) \leq 0\} &= h^t \left| K_{\theta_2 h}^{(t)}(u - Y(\mathbf{a}_0)) \right| I \{u - Y(\mathbf{a}_0) \leq 0\} \leq h^t c, \end{aligned}$$

and

$$\begin{aligned} &|K_h(u - Y(\mathbf{a}_0)) - I \{Y(\mathbf{a}_0) < u\}| \\ &= |K_h(u - Y(\mathbf{a}_0)) - 1| I \{Y(\mathbf{a}_0) < u\} + |K_h(u - Y(\mathbf{a}_0))| I \{Y(\mathbf{a}_0) \geq u\} \leq 2h^t c, \end{aligned}$$

where  $0 < \theta_1, \theta_2 < 1$ , and thus

$$\begin{aligned} |A_0 - A_{m,n}^K(\mathbf{a}_0)| &= \left| \int_{-\infty}^{\infty} G_{\mathbf{a}_0} dF_{\mathbf{a}_0} - \int_{-\infty}^{\infty} G_{\mathbf{a}_0}^K dF_{\mathbf{a}_0}^n \right| \\ &= \left| \int_{-\infty}^{\infty} G_{\mathbf{a}_0} d(F_{\mathbf{a}_0} - F_{\mathbf{a}_0}^n) + \int_{-\infty}^{\infty} (G_{\mathbf{a}_0} - G_{\mathbf{a}_0}^m) dF_{\mathbf{a}_0}^n + \int_{-\infty}^{\infty} (G_{\mathbf{a}_0}^m - G_{\mathbf{a}_0}^K) dF_{\mathbf{a}_0}^n \right| \\ &\leq \sup_{-\infty < u < \infty} |F_{\mathbf{a}_0}(u) - F_{\mathbf{a}_0}^n(u)| + \sup_{-\infty < u < \infty} |G_{\mathbf{a}_0}(u) - G_{\mathbf{a}_0}^m(u)| + 2h^t c, \end{aligned}$$

where  $F_{\mathbf{a}_0}$  and  $G_{\mathbf{a}_0}$  is the distribution function of  $X(\mathbf{a}_0)$  and  $Y(\mathbf{a}_0)$ ,  $F_{\mathbf{a}_0}^n$  and  $G_{\mathbf{a}_0}^m$  is the empirical distribution function of  $X(\mathbf{a}_0)$  and  $Y(\mathbf{a}_0)$ , and  $G_{\mathbf{a}_0}^K(u) = \int_{-\infty}^{\infty} \int_{-\infty}^{(u-y)/h} k(w) dw dG_{\mathbf{a}_0}^m(y)$ . Since

$$\sqrt{m+n} |A_0 - EA_{m,n}^K(\mathbf{a}_0)| \leq \sqrt{m+n} E |A_0 - A_{m,n}^K(\mathbf{a}_0)|,$$

409 it follows that when  $\sqrt{m+n}h^t \rightarrow 0$ , we have  $\sqrt{m+n} |EA_{m,n}^K(\mathbf{a}_0) - A_0| \rightarrow 0$ .

In addition, by Lemma 1-Lemma 3, it follows that  $\sqrt{m+n} (A_{m,n}^K(\hat{\mathbf{a}}_K) - A_0)$  and  $\sqrt{m+n} (A_{m,n}^K(\mathbf{a}_0) - EA_{m,n}^K(\mathbf{a}_0))$  shares the same distribution. Thus by Lemma 4, we have

$$\sqrt{\frac{mn}{m+n}} \frac{A_{m,n}^K(\mathbf{a}_0) - EA_{m,n}^K(\mathbf{a}_0)}{s} \xrightarrow{d} N(0, 1).$$

Since  $|A_0 - A_{m,n}^K(\hat{\mathbf{a}}_K)| = O_p(n^{-1/2})$  and  $l'''(A_0)|_{A_0=\bar{A}} = O(n)$  shown in Lemma 1, we have  $r_{m,n} = O_p(n^{-1/2})$  and hence,

$$\gamma(A_0)l(A_0) \doteq \left( \sqrt{\frac{mn}{m+n}} \frac{A_{m,n}^K(\hat{\mathbf{a}}_K) - A_0}{s} \right)^2.$$

410 Therefore,  $\gamma(A_0)l(A_0)$  has the  $\chi_1^2$  distribution asymptotically. The proof of

411 Proposition 1 is thus completed.  $\square$

412 **Appendix C. Discussion on the Remark**

For the Gaussian kernel, the first derivative is  $k'(u) = -\frac{u}{\sqrt{2\pi}} \exp(-u^2/2)$ . To show the boundedness for  $k'(u)$ , we calculate the first and the second derivative of  $k'(u)$ , obtaining

$$k''(u) = (u^2 - 1) \exp(-u^2/2) / \sqrt{2\pi},$$

and

$$k'''(u) = (3u - u^3) \exp(-u^2/2) / \sqrt{2\pi}.$$

413 Set  $k''(u)$  to be 0. At  $u = -1$ ,  $k'''(u) < 0$ , therefore  $k'$  reaches to the  
 414 maximum of  $\exp(-1/2) / \sqrt{2\pi}$ . At  $u = 1$ ,  $k'''(u) > 0$ ,  $k'$  reaches the minimum  
 415 of  $-\exp(-1/2) / \sqrt{2\pi}$ . Similarly, we can show the  $l$ th derivative  $K^{(l)}(u)$  is  
 416 bounded for all  $l \geq 2$ . If there exist  $t > 2$  such that  $t = \sup_l \{l : E|\mathbf{X} - \mathbf{Y}|^l <$   
 417  $\infty\}$ , then Proposition 1 is satisfied with proper selections of bandwidths where  
 418  $\delta \in (1/2t, 1/4)$ .

419 **References**

- 420 Azzalini, A., 1981. A note on the estimation of a distribution function and  
 421 quantiles by a kernel method. *Biometrika* 68, 326–328.
- 422 Bamber, D., 1975. The area above the ordinal dominance graph and the area  
 423 below the receiver operating characteristic graph. *Journal of Mathematical*  
 424 *Psychology* 12, 387–415.
- 425 Chen, S.X., Hall, P., 1993. Smoothed empirical likelihood confidence intervals  
 426 for quantiles. *The Annals of Statistics* 21, 1166–1181.
- 427 Faraggi, D., Reiser, B., 2002. Estimation of the area under the ROC curve.  
 428 *Statistics in Medicine* 21, 3093–3106.
- 429 Hall, P., La Scala, B., 1990. Methodology and algorithms of empirical likelihood.  
 430 *International Statistical Review/Revue Internationale de Statistique* 58, 109–  
 431 127.

- 432 Johnson, M.E., 2013. Multivariate Statistical Simulation: A Guide to Selecting  
433 and Generating Continuous Multivariate Distributions. John Wiley & Sons.
- 434 Lamon-Fava, S., Wilson, P.W., Schaefer, E.J., 1996. Impact of body mass index  
435 on coronary heart disease risk factors in men and women the Framingham  
436 offspring study. *Arteriosclerosis, Thrombosis, and Vascular Biology* 16, 1509–  
437 1515.
- 438 Lazar, N., Mykland, P.A., 1998. An evaluation of the power and conditionality  
439 properties of empirical likelihood. *Biometrika* 85, 523–534.
- 440 Limpert, E., Stahel, W.A., Abbt, M., 2001. Log-normal distributions across  
441 the sciences: Keys and clues on the charms of statistics, and how mechanical  
442 models resembling gambling machines offer a link to a handy way to character-  
443 ize log-normal distributions, which can provide deeper insight into variability  
444 and probability—normal or log-normal: That is the question. *BioScience* 51,  
445 341–352.
- 446 Lloyd, C.J., 1998. Using smoothed receiver operating characteristic curves to  
447 summarize and compare diagnostic systems. *Journal of the American Statis-  
448 tical Association* 93, 1356–1364.
- 449 Lloyd, C.J., Yong, Z., 1999. Kernel estimators of the ROC curve are better than  
450 empirical. *Statistics and Probability Letters* 44, 221–228.
- 451 Ma, S., Huang, J., 2005. Regularized ROC method for disease classification and  
452 biomarker selection with microarray data. *Bioinformatics* 21, 4356–4362.
- 453 Ma, S., Huang, J., 2007. Combining multiple markers for classification using  
454 ROC. *Biometrics* 63, 751–757.
- 455 Markatou, M., Basu, A., Lindsay, B.G., 1998. Weighted likelihood equations  
456 with bootstrap root search. *Journal of the American Statistical Association*  
457 93, 740–750.

- 458 McIntosh, M.W., Pepe, M.S., 2002. Combining several screening tests: opti-  
459 mality of the risk score. *Biometrics* 58, 657–664.
- 460 Nadaraya, E.A., 1964. Some new estimates for distribution functions. *Theory*  
461 *of Probability & Its Applications* 9, 497–500.
- 462 Nelder, J.A., Mead, R., 1965. A simplex method for function minimization. *The*  
463 *Computer Journal* 7, 308–313.
- 464 Owen, A.B., 1990. Empirical likelihood ratio confidence regions. *The Annals of*  
465 *Statistics* 18, 90–120.
- 466 Owen, A.B., 2001. *Empirical Likelihood*. CRC press.
- 467 Pepe, M.S., 1997. A regression modelling framework for receiver operating  
468 characteristic curves in medical diagnostic testing. *Biometrika* 84, 595–608.
- 469 Pepe, M.S., Cai, T., 2004. The analysis of placement values for evaluating  
470 discriminatory measures. *Biometrics* 60, 528–535.
- 471 Pepe, M.S., Cai, T., Longton, G., 2006. Combining predictors for classification  
472 using the area under the receiver operating characteristic curve. *Biometrics*  
473 62, 221–229.
- 474 Pepe, M.S., Thompson, M.L., 2000. Combining diagnostic test results to in-  
475 crease accuracy. *Biostatistics* 1, 123–140.
- 476 Qin, G., Zhou, X.H., 2006. Empirical likelihood inference for the area under the  
477 ROC curve. *Biometrics* 62, 613–622.
- 478 R Development Core Team, 2013. *R: A Language and Environment for Statis-*  
479 *tical Computing*. R Foundation for Statistical Computing. Vienna, Austria.  
480 URL: <http://www.R-project.org/>. ISBN 3-900051-07-0.
- 481 Reiser, B., Faraggi, D., 1997. Confidence intervals for the generalized ROC  
482 criterion. *Biometrics* 53, 644–652.

- 483 Schisterman, E.F., Faraggi, D., Browne, R., Freudenheim, J., Dorn, J., Muti,  
484 P., Armstrong, D., Reiser, B., Trevisan, M., 2001. Tbars and cardiovascular  
485 disease in a population-based sample. *Journal of Cardiovascular Risk* 8, 219–  
486 225.
- 487 Schisterman, E.F., Faraggi, D., Browne, R., Freudenheim, J., Dorn, J., Muti,  
488 P., Armstrong, D., Reiser, B., Trevisan, M., 2002. Minimal and best linear  
489 combination of oxidative stress and antioxidant biomarkers to discriminate  
490 cardiovascular disease. *Nutrition, Metabolism, and Cardiovascular Diseases:*  
491 *NMCD* 12, 259–266.
- 492 Serfling, R.J., 2009. *Approximation Theorems of Mathematical Statistics*. Vol-  
493 ume 162. John Wiley & Sons.
- 494 Shanno, D.F., 1970. Conditioning of quasi-newton methods for function mini-  
495 mization. *Mathematics of Computation* 24, 647–656.
- 496 Su, J.Q., Liu, J.S., 1993. Linear combinations of multiple diagnostic markers.  
497 *Journal of the American Statistical Association* 88, 1350–1355.
- 498 Tibshirani, R., 1996. Regression shrinkage and selection via the lasso. *Journal*  
499 *of the Royal Statistical Society, Series B* 58, 267–288.
- 500 Vexler, A., Liu, A., Schisterman, E.F., Wu, C., 2006. Note on distribution-free  
501 estimation of maximum linear separation of two multivariate distributions.  
502 *Nonparametric Statistics* 18, 145–158.
- 503 Zou, K.H., Hall, W., Shapiro, D.E., 1997. Smooth non-parametric receiver op-  
504 erating characteristic (ROC) curves for continuous diagnostic tests. *Statistics*  
505 *in Medicine* 16, 2143–2156.