## Harvard University Harvard University Biostatistics Working Paper Series

*Year* 2006 *Paper* 43

# Using Profile Likelihood for Semiparametric Model Selection with Application to Proportional Hazards Mixed Models

Ronghui Xu<sup>∗</sup> Anthony Gamst† Michael Donohue‡ Florin Vaida∗∗ David P. Harrington††

<sup>∗</sup>University of California, San Diego, rxu@math.ucsd.edu

†University of California, San Diego, agamst@ucsd.edu

‡University of California, San Diego, mdonohue@ucsd.edu

∗∗University of California, San Diego, vaida@ucsd.edu

††Dana-Farber Cancer Institute and Harvard School of Public Health, dph@hsph.harvard.edu

This working paper is hosted by The Berkeley Electronic Press (bepress) and may not be commercially reproduced without the permission of the copyright holder.

http://biostats.bepress.com/harvardbiostat/paper43

Copyright  $\odot$  2006 by the authors.

# Using Profile Likelihood for Semiparametric Model Selection with Application to Proportional Hazards Mixed Models

Ronghui Au<sup>-,-,</sup> , Anthony Gamst-, Michael Donohue-, Florin Vaidaand David P. Harrington<sup>3</sup>

1Division of Biostatistics and Bioinformatics

Department of Family and Preventive Medicine and 2Mathematics

University of California, San Diego

3Department of Biostatistics and Computational Biology

Dana-Farber Cancer Institute and Harvard School of Public Health

Correspondence: rxu@ucsd.edu

May 28, 2006

### A bstract

Proportional hazards mixed effects model (PHMM) was recently proposed, which incorporates general random effects of arbitrary covariates and includes the univariate frailty model as a special case. In this paper we establish the asymptotic properties of the nonparametric maximum likelihood estimator under PHMM. The asymptotic **Collection of Biostatistics** 1 **Research Archive** 

results allow us to use the prole likelihood for selection of both nested and non-nested PHMMs. We define both a profile likelihood ratio test and a profile Akaike information for general models with nuisance parameters. Asymptotic quadratic expansion of the log profile likelihood allows derivation of the asymptotic null distribution of the likelihood ratio statistic including the boundary cases, as well as unbiased estimation of the Akaike information by an Akaike information criterion. For computation of the likelihood under PHMM we apply three algorithms: Laplace approximation, reciprocal importance sampling and bridge sampling. We compare the three algorithms under different data structures, and apply the methods to a multi-center lung cancer clinical trial.

Key words: Akaike information, asymptotic efficiency, consistency, profile likelihood, likelihood ratio test, testing on the boundary, Laplace approximation, reciprocal importance sampling, bridge sampling.



#### **Motivation** 1

In recent years random effects models for failure time data have been applied in various areas, for unobserved heterogeneity, for dependence induced by clustering in, for instance, familial studies, and in settings where some effects, such as center effects in a multi-center trial, are best thought of as sampled from a wider population. The work in this paper, although developed under the more general semiparametric models, has been motivated by the random effects models for failure time data. Like linear and generalized linear models, these random effects models have provided a natural way to model many within-cluster correlations. For example, Vaida and Xu (2000) showed how such models can be used to understand institutional variation in outcomes of a multi-center lung cancer trial conducted by the Eastern Cooperative Oncology Group. The use of random effects survival models in clinical trials was also advocated in Glidden and Vittinghoff (2004), Murray et al. (2004) and Sylvester et al. (2002). Liu et al. (2004ab), on the other hand, used variance components to identify the genetic contribution to the age of onset of alcohol dependence and alcohol abuse. The full power and flexibility of the random effects models, however, has not yet been extended to regression methods for right-censored data.

Vaida and Xu (2000) studied the proportional hazards model with mixed effects (PHMM). It includes the more classical 'frailty' models with random effects on the baseline hazard, but also allows random covariate effects. In this way it is able to model covariate by cluster interactions, such as varying treatment effects in a multicenter clinical trial. The model is of the form

$$
\lambda_{ij}(t) = \lambda_0(t) \exp(\beta' \mathbf{Z}_{ij} + \mathbf{b}'_i \mathbf{W}_{ij}), \qquad (1)
$$

where  $\lambda_{ij}(t)$  is the hazard function of the j-th observation from the i-th cluster,  $\mathbf{b}_i$  is a vector of random effects for the *i*-th cluster, and  $\mathbf{Z}_{ij}$ ,  $\mathbf{W}_{ij}$  are the covariate vectors for the fixed and random effects. This model contains a multivariate random effect with **Collection of Biostatistics** 3

**Research Archive** 

arbitrary design matrix in the log relative risk, in a way similar to the linear, generalized linear and nonlinear mixed models. Vaida and Xu developed the nonparametric maximum likelihood estimator (NPMLE) of the parameters in this model, computed using the EM algorithm and Markov Chain Monte Carlo (MCMC) methods. However, the asymptotic properties of the NPMLE remain unproven under the PHMM.

As in any regression setting, model selection is an important aspect of data analysis. In particular, in the application of model (1), it often needs to be decided whether a random effect term should be incorporated into the model. From the testing point of view, the null hypothesis is that the corresponding variance component is zero. Although the standard errors of the estimated variance components are obtained in Vaida and Xu (2000), they cannot be used directly for testing zero variance components, because the null hypothesis lies on the boundary of the parameter space. Gray (1995) and Commenges and Andersen (1995) proposed a score test of homogeneity for this purpose. The score test, however, is restricted to the null hypothesis of no random effects. In addition, no tests are readily available for testing more than one parameter at a time, such as for testing the signicance of a categorical covariate with more than two categories. In this paper we develop a likelihood ratio test in the general semiparametric setting that, under PHMM, allows arbitrary testing on the mixed model, so a data analyst could test for the significance of a specified subset of the random and/or fixed effects.

Another approach to model selection is via information criteria (Linhart and Zucchini, 1986), which easily handles the comparison of non-nested models, and avoids the boundary problem in the case of selection of random effects. The Akaike information criterion (AIC; Akaika, 1973; deLeuw, 1992; Burnham and Anderson, 2002) is among the most commonly used in practice. It has a simple interpretation as penalized loglikelihood, as well as an information-theoretic foundation. Under the Cox model with no random effects, an AIC has been used in association with the partial likelihood **Collection of Biostatistics** 

**Research Archive** 

(Verweij and van Houwelingen, 1995). However, partial likelihoods do not universally exist for semiparametric models; in particular, strictly speaking it does not apply to PHMM (1). Here we aim to give a meaningful derivation of the AIC for general models with nuisance parameters, and in particular to semiparametric models where only the finite dimensional parameters are of interest.

In the next section we prove the consistency and asymptotic normality of the NPMLE under PHMM. In Section 3 we study the profile likelihood for general semiparametric models, and use it to derive the profile likelihood ratio test including the boundary case; we also develop an  $AIC$  using the profile likelihood. In Section 4 we apply the profile likelihood ratio test and the profile  $AIC$  to  $PHMM$ , and consider three algorithms to compute the maximized likelihood under PHMM. Simulation studies are carried out in Section 5 and an example is given in Section 6 to illustrate the methods. Section 7 contains some further discussion. But first, we review the proportional hazards mixed model in some detail below.

## 1.1 Proportional hazards mixed model

Assume that the data consist of possibly right-censored event time observations from n clusters, with  $n_i$  observations in each cluster,  $i = 1...n$ . Within a cluster the observations are dependent, but conditional on the cluster-specic d 1 vector of random effects  $\mathbf{b}_i$ , the survival times  $T_{ij}$  are independent and their hazard functions follow PHMM (1). In (1)  $\mathbf{W}_{ij}$  is often a subset of  $\mathbf{Z}_{ij}$ , apart from possibly a '1' which represents the cluster effect on the baseline hazard. To insure identifiability, we assume that  $E(\mathbf{b}_i) = \mathbf{0}$ . For distribution of the random effects we also assume that

$$
\mathbf{b}_i \stackrel{iid}{\sim} N(\mathbf{0}, \mathbf{\Sigma}) \tag{2}
$$

as in Vaida and Xu (2000). Note that the other commonly used frailty distribution, the gamma distribution, is not suitable under the general random effects model  $(1)$ . **Collection of Biostatistics** 5 **Research Archive** 

This is because it is not scale-invariant so that the inference is not invariant under a change of measuring unit for covariates of the random effects.

The data from subject j in cluster i can be written  $y_{ij} = (X_{ij}, \delta_{ij}, \mathbf{Z}_{ij}, \mathbf{W}_{ij})$ , where  $X_{ij}$  is the possibly right-censored failure time and  $\delta_{ij}$  is the failure-event indicator. Let  $y_i = (y_{i1},..., y_{in_i})$  be the data for cluster *i*. For cluster *i*, conditional on the random effect  $\mathbf{b}_i$ , the log-likelihood is

$$
l_i = l_i(\boldsymbol{\beta}, \lambda_0; \mathbf{y}_i | \mathbf{b}_i) = \sum_{j=1}^{n_i} \{ \delta_{ij} \log \lambda_0(X_{ij}) + \delta_{ij} (\boldsymbol{\beta}' \mathbf{Z}_{ij} + \mathbf{b}_i' \mathbf{W}_{ij}) - \Lambda_0(X_{ij}) e^{\boldsymbol{\beta}' \mathbf{Z}_{ij} + \mathbf{b}_i' \mathbf{W}_{ij}} \},
$$
(3)

where  $\mathbf{u} \cdot \mathbf{v} = \mathbf{u} \cdot \mathbf{u}$  $\int_0^t \lambda_0(s) ds$ . We rewrite the parameter for the baseline hazard in the following as  $\lambda$ , to be consistent with the general semiparametric model framework that we will use. The likelihood of the observed data is then

$$
L(\theta) = \prod_{i=1}^{n} \int \exp(l_i) p(\mathbf{b}_i; \Sigma) d\mathbf{b}_i,
$$
\n(4)

where  $\theta = (\beta, \Sigma, \lambda)$  and  $p(\cdot)$  is the multivariate normal distribution. Usually no closedform expression is available for  $L(\theta)$  and its calculation involves d-dimensional integration.

#### $\overline{2}$ Asymptotic theory under PHMM

We assume the following conditions on the data.

- C1. Conditional on the covariates  $\mathbf{z}_{ij}$  and  $\mathbf{w}_{ij}$ , the fatent censoring time  $C_{ij}$  is independent of the failure time  $T_{ij}$  and random effects  $\mathbf{b}_i$ .
- C<sub>2</sub>. There exists some positive constant  $\epsilon$  such that  $\Gamma$ ( $C_{ij} \geq \tau |\mathbf{Z}_{ij}, \mathbf{W}_{ij} \rangle \geq \epsilon$  almost surely.
- C3.  $\mathbf{Z}_{ij}$  and  $\mathbf{W}_{ij}$  are bounded. In addition, if there exists a constant vector **c** and a symmetric matrix  $\Sigma$  such that

A BEPRESS REPOSITION of Blostatistics

\nResecarloA Archive

\n1. 
$$
Z'_{ij}I' + W'_{ij} \Sigma W_{ij} = 0, \quad j = 1, \ldots, n_i
$$

and

$$
\mathbf{W}'_{ij}\mathbf{\Sigma}\mathbf{W}_{ij'}=0, \quad j\neq j'; j,j'=1,\ldots,n_i
$$

almost surely, then  $c = 0$  and  $\Sigma = 0$ .

- C4. The true cumulative hazard  $\Lambda_0(t)$  is strictly increasing and continuously differentiable in [0,  $\tau$ ]. Also,  $\Lambda_0(\tau) < \infty$ .
- C5. The true values of  $\beta$  and  $\Sigma$ ,  $\beta_0$  and  $\Sigma_0$ , belong to the interior of a known compact set,

 $\mathcal{S}$  is functionally in the some constant  $\mathcal{S}$  for some constant  $\mathcal{S}$  , and if  $\mathcal{S}$  $\Sigma$  is positive definite and its eigenvalues are bounded and 1g  $\sim$  1g  $\$ 

 $C$ 6. The cluster sizes  $\{e_i\}$  are in a random variables and  $\{e_i\}$  and  $\{e_i\}$   $\equiv$   $\{e_i\}$   $\equiv$   $\{e_i\}$  and  $\{e_i\}$ 

**Theorem 1** Under conditions  $C_1 = C_0$ ,  $||p_n - p_0|| \to 0$ ,  $||\angle p_n - \angle p_0|| \to 0$  and  $\sup_{t \in [0, \tau]} ||\Delta n(t) |\Lambda_0(t)| \to 0$  almost surely where  $\|\cdot\|$  is the Euclidean norm.

Theorem 2 Under conditions  $C1-C6$ 

$$
\sqrt{n}(\hat{\boldsymbol{\beta}}_n'-\boldsymbol{\beta}_0',\hat{\boldsymbol{\Sigma}}_n'-\boldsymbol{\Sigma}_0',\hat{\Lambda}_n(\cdot)-\Lambda_0(\cdot))'
$$

converges to a zero mean Gaussian process in  $\mathbf{R}^2 \times \mathbf{R}^2 \times \mathbb{R}^2 \times \mathbb{R}^2$  [0,7] where  $\mathbf{Z}_n$ and  $\Sigma_0$  are treated as extended column vectors consisting of the upper triangle elements and  $l^{\infty}[0, \tau]$  is the space of all bounded functions on  $[0, \tau]$  with the sup norm on  $[0, \tau]$ . Furthermore,  $\mu_n$  and  $\omega_n$  are asymptotically efficient.

**Theorem 3** Let  $V(\mathbf{h}_1, \mathbf{h}_2, h_3)$  be the asymptotic variance of

$$
\sqrt{n}\{\mathbf{h}'_1(\hat{\boldsymbol{\beta}}_n - \boldsymbol{\beta}_0) + \mathbf{h}'_2(\hat{\boldsymbol{\Sigma}}_n - \boldsymbol{\Sigma}_0) + \int_0^{\tau} h_3(t) d(\hat{\boldsymbol{\Lambda}}_n(t) - \boldsymbol{\Lambda}_0(t))\};
$$
  
Collection of Blostatistics  
Research Archive

 $\mathbf{h}_n$  be the vector  $\mathbf{h}_1$ ,  $\mathbf{h}_2$ , and  $h_3(X_{ij})$  for which  $\delta_{ij} = 1$ ; and  $\mathbf{J}_n$  be the negative Hessian matrix of  $\log L_n(\sigma)$  with respect to  $(\rho, \Delta)$  and the jump sizes of ix at  $X_{ij}$  for which  $\sigma_{ij} =$ 1. Then under C1-C6, the variance estimator  $n\mathbf{h}'_n J_n^{-1} \mathbf{h}_n \to V(\mathbf{h}_1, \mathbf{h}_2, h_3)$  uniformly in probability.

The proofs of the above theorems are given in the Appendix.

## 3 Profile likelihood for model selection

In this section we discuss the profile likelihood in the general context of semiparametric models, using the quadratic expansion of Murphy and van der Vaart (2000). Assume that the data consists of a random sample of n observations,  $y_1, \ldots, y_n$ , from a distribution depending on parameters  $\varphi$  and  $\lambda$ . We assume that  $\varphi \in \Psi$ , a subset of  $\mathbf{R}^r$ , and  $\lambda$  is a nuisance parameter, possibly of infinite dimension. The log-likelihood of the data is  $l(\boldsymbol{\phi}, \lambda) = \sum_{i=1}^n l_i(\boldsymbol{\phi}, \lambda)$ , and  $l_i$  is the log-likelihood for  $\mathbf{y}_i$ . The log profile likelihood function for  $\phi$ , with the nuisance parameter  $\lambda$  'profiled out', is

$$
pl(\boldsymbol{\phi}) = \sup_{\lambda} l(\boldsymbol{\phi}, \lambda).
$$
 (5)

Following Murphy and van der Vaart (2000), under suitable conditions the log profile likelihood behaves as a quadratic function asymptotically; i.e. for any random sequence  $\phi_n$  such that  $\|\phi_n - \phi_0\| = O_p(1/\sqrt{n})$  where  $\phi_0$  is the true parameter value,

$$
\frac{1}{n}\left\{\text{pl}(\boldsymbol{\phi}_n)-\text{pl}(\boldsymbol{\phi}_0)\right\}=(\boldsymbol{\phi}_n-\boldsymbol{\phi}_0)'\mathbf{A}-\frac{1}{2}(\boldsymbol{\phi}_n-\boldsymbol{\phi}_0)'\mathbf{I}(\boldsymbol{\phi}_n-\boldsymbol{\phi}_0)+o_p\left(\frac{1}{n}\right),\qquad(6)
$$

where  $\mathbf{A} = \sum_{1}^{n} \mathbf{s}(\mathbf{y}_i)/n$ , s is the efficient score for  $\boldsymbol{\phi}$ , i.e. the ordinary observed score function minus its orthogonal projection onto the closed linear span of the score functions for the nuisance parameter  $\lambda$ , and **I**, its covariance matrix, is the efficient Fisher information matrix (Murphy and van der Vaart, 2000; Severini and Wong, 1992). We will derive the results of this section for semiparametric models that satisfy  $(6)$ .

**Collection of Biostatistics Research Archive** 

8

## 3.1 Prole likelihood ratio test

The likelihood ratio statistic for two nested parametric models, when the parameter space of the smaller model lies entirely in the interior of that of the larger model, has a chi-squared null distribution with the number of degrees of freedom equal to the difference of those of the two models. For a semiparametric model such as  $(1)$ , the number of degrees of freedom of the model itself is not well defined, since there is at least one infinite dimensional parameter. However, if the infinite dimensional parameter is a nuisance parameter, then under certain conditions the likelihood ratio statistic can be defined via the profile likelihoods, with the number of degrees of freedom calculated using the finite dimensional parameters.

For two nested models let - be the parameter space under the larger model, and the parameter space under the smaller model, or equivalently, under the null hypothesis  $H_0$ . We assume that  $H_0$  places no additional restrictions on the nuisance parameter  $\lambda$ . Denote L the likelihood, and let

$$
LR = \frac{\sup_{\Theta_0} L(\phi, \lambda)}{\sup_{\Theta} L(\phi, \lambda)}.
$$
 (7)

Then LR is the ratio of the maximized likelihoods under the two models. The above can also be viewed as the ratio of the maximized profile likelihoods, with the nuisance parameter  $\lambda$  'profiled out'. So

$$
-2\log LR = -2\{\sup_{\Phi_0} \text{pl}(\boldsymbol{\phi}) - \sup_{\Phi} \text{pl}(\boldsymbol{\phi})\},\tag{8}
$$

where  $\Phi_0$  and  $\Phi$  are the corresponding parameter spaces for  $\phi$  under the two models. Murphy and van der Vaart (2000) showed that as result of the quadratic expansion (6), when  $\phi_0$  lies in the interior of the parameter space, the profile likelihood ratio test for  $H_0$ :  $\phi = \phi_0$  has asymptotically chi-squared null distribution with the number of degrees of freedom equal to the dimension of  $\phi$ .

Testing on the boundary

As mentioned in Section 1, the challenging problem in hypothesis testing under model (1) is when the null hypothesis lies on the boundary of the parameter space, such as testing against zero variances of the random effects. We show in the following that the asymptotic expansion (6) enables us to obtain results on the null distribution of the prole likelihood ratio statistic similar to those in Self and Liang (1987). First we obtain a result similar to that of Theorem 1 in Self and Liang (1987), on the  $\sqrt{n}$ consistency of the maximum (profile) likelihood estimator when  $\phi_0$  is on the boundary of  $\Phi$ , given the  $\sqrt{n}$ -consistency when  $\phi_0$  lies in the interior of  $\Phi$ .

**Theorem 4** Given the quadratic expansion (6), with probability tending to 1 as  $n \to \infty$ ! 1 there exists a sequence of points in  $\mathbf{x}, \varphi_n$ , at which local maxima of  $\mu_n(\varphi)$  occur, that converges to  $\boldsymbol{\phi}_0$  in probability. Moreover,  $\sqrt{n}(\boldsymbol{\phi}_n - \boldsymbol{\phi}_0) = O_p(1)$ .

See Appendix for proof.

Notice that (6) is equal to

$$
\frac{1}{2}\mathbf{A}'\mathbf{I}^{-1}\mathbf{A}-\frac{1}{2}\{\mathbf{z}_n-(\boldsymbol{\phi}_n-\boldsymbol{\phi}_0)\}^{\prime}\mathbf{I}\{\mathbf{z}_n-(\boldsymbol{\phi}_n-\boldsymbol{\phi}_0)\}+o_p\left(\frac{1}{n}\right),
$$
\n(9)

where  $z_n = I^{-1}A$ . Therefore the same representation of the asymptotic distribution of  $-2 \log LR$  as that from Chernoff (1954) and Self and Liang (1987) is obtained, which can then be used to calculate the null distribution of the likelihood ratio statistics. Specifically, assume that  $\Phi$  and  $\Phi_0$  are regular enough to be approximated by cones with vertices at  $\phi_0$  (for definition see Self and Liang (1987) or Chernoff (1954)), we have

Theorem 5 Let Z be a random variable with a multivariate Gaussian distribution of mean  $\varphi$  and covariance matrix  $\mathbf{1} = (\varphi_0)$ , and let  $\mathfrak{C}_\Phi$  and  $\mathfrak{C}_{\Phi_0}$  be non-empty cones approximating  $\Phi$  and  $\Phi_0$  at  $\phi_0$ , respectively. Then the asymptotic distribution of the likelihood ratio statistic,  $-2 \log LR$ , is the same as the distribution of the likelihood ratio test of  $\mathcal{I}$  ,  $\mathcal{I}$  constructed on a single realization of  $\mathcal{I}$  when  $\mathcal{I}$ **Collection of Biostatistics** 

**Research Archive** 

## 3.2 Profile Akaike information

In this subsection we construct the Akaike information and its associated criterion, AIC, for models with nuisance parameters. Since the relevant quantity is the profile likelihood, we term the criterion profile AIC.

Considered a family of models  $\bullet$  ,  $\bullet$  parameterized by  $\circ$  ,  $\circ$ the parameter of interest, and  $p$  is the nuisance parameter, possibly of interest, possibly of interest,  $p$ dimension. The view we take here, similar to Claeskens and Hjort (2003), is that we are interested in selecting the ' $\phi$  part' of the modelling, while leaving the parameter space the same across all competing models. In this way, for model selection purposes  $\mu$  is purposed as  $\mu$ really indexed by  $\phi$  alone. Assume that the data vector y, consisting of n independent observations  $y_1, ..., y_n$ , is generated by a distribution with density f. The classical 'distance' from the true distribution f to a member  $g_{\theta} = g(\cdot | \phi, \lambda)$  of M is given by the Kullback-Leibler information  $(KL)$ ,  $I(f, g_{\theta}) = E_f\{\log f(\mathbf{y}) - \log g_{\theta}(\mathbf{y})\}\$ . When the focus is on  $\phi$  alone, the relevant distance is that between f and the subfamily of models for  $\partial \psi_i \wedge \cdots \equiv -1$ . The map  $\partial \psi_i \wedge \cdots \wedge \partial \psi_i$ some  $\Delta = \Delta(\psi)$  for each  $\psi$ . Following Severini and Wong (1992),  $\Delta(\psi)$  is in fact a least favorable curve under smoothness conditions (see also Fan and Wong, 2000). We denote  $g_{\phi} = g(\phi, \wedge(\psi))$ . Ignoring the constant term  $E_1$ log  $f(y)$  in  $I(f, \phi)$ , we have that

$$
E\{\log g_{\boldsymbol{\phi}}(\mathbf{y})\} = \max_{\lambda} E\{\log g_{\boldsymbol{\phi},\lambda}(\mathbf{y})\};
$$

the expectations here and in the rest of this section are with respect to the true distribution f. Therefore  $g_{\phi}$  is the theoretical equivalent of the profile likelihood.

Minimum KL is attained at  $\phi_0$  such that  $I(f, g_{\phi_0}) = \min_{\phi} I(f, g_{\phi})$ , or, equivalently,

$$
E\{\log g_{\boldsymbol{\phi}_0}(\mathbf{y})\} = \max_{\boldsymbol{\phi}} E\{\log g_{\boldsymbol{\phi}}(\mathbf{y})\}.
$$

 $J(\psi)$  is the function term approximation to f within the family of models  $J(\psi)$ model is corrected in the correct  $j$  and  $j$  and  $j$   $\theta_{\theta}$ . In practice  $f_{0}$  is estimated in a set of Blocketicisc 11 **Research Archive** 

 $\mathcal{D}(\mathbf{y})$  which maximizes the profile intentional.

$$
pl(\mathbf{y}|\hat{\boldsymbol{\phi}}) = \max_{\boldsymbol{\phi}} pl(\mathbf{y}|\boldsymbol{\phi}) = \max_{\boldsymbol{\phi}, \lambda} \log g(\mathbf{y}|\boldsymbol{\phi}, \lambda).
$$

Note that  $(\varphi, \lambda)$  is the MLE for  $(\varphi, \lambda)$ . The predictive value of pl( $\psi$ ) is given by the expected  $\bm{\Lambda}$ L for predicting new data  $\bm{y}$  , independent of but from the same distribution as y. Ignoring the constant term, we define the profile Akaike Information

$$
\text{pAI} = -2E_{f(\mathbf{y})}E_{f(\mathbf{y}^*)}\text{pl}(\mathbf{y}^*|\hat{\boldsymbol{\phi}}(\mathbf{y})).\tag{10}
$$

It is important to note that  $\mathsf{p}(\mathbf{y} \mid \boldsymbol{\varphi}(\mathbf{y}))$  is different from the log-fikelihood function computed at the  $MLE$  ( $\phi$ ,  $\lambda$ ), since it allows maximizing the inclinioud over  $\lambda$  based on the new data  ${\bf y}$  . The following result shows that  ${\rm p}$ AI can be estimated by a corresponding profile  $AIC$ , where the number in the correction term is  $p$ , the dimension of  $\phi$ .

 $\theta_0$  in the interior of the parameter space. Further, assume that  $y, y^*$  consist of n i.i.d. vectors, and  $\varphi$  is consistent for  $\varphi_0$ . Then the projue AIC

$$
pAIC = -2pl(\mathbf{y}|\hat{\boldsymbol{\phi}}(\mathbf{y})) + 2p \tag{11}
$$

is an approximately unbiased estimator of pAI, in the sense that

$$
pAI = E(pAIC) + E(r),
$$

where  $r = o_p(1)$  as  $n \to \infty$ . If in addition r is uniformly integrable, then  $E(r) = o(1)$ , and pAIC is asymptotically unbiased for pAI.

See Appendix for proof.

Note that in proving the above we assume that the family of models under consideration contains the operating model  $f$ , so that the parameters lie in the interior **Collection of Biostatistics** 12 **Research Archive** 

of the parameter space. This is generally the case in the theory of AIC. Incidentally, for model selection this avoids the boundary problem encountered in likelihood ratio testing for nested models, since the AIC is computed assuming that the model in each case holds. We also hoted earlier that with hew data **y** the pronie likelihood function at  $\varphi(y)$  is not the same as the intermood function at the MLE based on data  $y$ . However, when computing the pAIC, the observed profile likelihood in  $(11)$  is the same as the maximized intermode at  $v$ . The correction term,  $2p$ , depends on the definition of the parameter of interest. In particular, if  $\lambda$  has finite dimension q, the classic AIC for  $\nu = (\psi, \lambda)$  is  $\psi = 2\iota(\nu) + 2(\nu + q)$ , while the profile AIC for  $\psi$  is  $\psi = 2\iota(\nu) + 2\mu$ .

#### 4 Application to PHMM 4

Under PHMM our parameter of interest is  $\phi = (\beta, \Sigma)$ , whereas the baseline hazard  $\lambda$  is seen as a nuisance parameter. Asymptotic normality of the MLE established in Section 2 implies that the likelihood surface is asymptotically quadratic near the true parameter values, which in turn implies that the same holds for the profile likelihood (Murphy and van der Vaart, 2000; Li, 2000) . The asymptotic properties of the MLE have also been established for the gamma frailty models (Murphy, 1994, 1995; Parner, 1998), and Maple  $et \ al.$  (2002) verified empirically that the contours of the profile likelihood under PHMM are elliptic.

## 4.1 Profile likelihood ratio test under PHMM

The representation given in Theorem 5 only involves the finite dimensional parameter  $\phi$  under the PHMM, so for the cases of null distributions considered by Self and Liang, or by Stram and Lee  $(1994, 1995)$  for linear mixed effects model, the results are exactly the same.

In the following we list the cases which are the most likely to be encountered in practice, and correct an error in the existing literature. Denote in the following  $d$  as the dimension of b.

Case 1:  $d = q + 1$  and

$$
\mathbf{\Sigma} = \left(\begin{array}{cc} \mathbf{\Sigma}_{11} & \sigma_{12} \\ & \\ \sigma_{12} & \sigma_{22} \end{array}\right),
$$

where  $11$  is  $1$  is the asymptotic null distribution of  $2$  log LR for testing  $2$  $H_0: \sigma_{22} = 0$  (and therefore  $\sigma_{12} = 0$ ) against  $\omega$  positive semidefinite is  $(\chi_q + \chi_{q+1})/2$ . when  $q = 0$ , the above distribution is a 50:50 mixture of a point mass at 0 and  $\chi_1^2$ ; note that in this case the maximum likelihood estimator of the variance components has a positive probability of being zero. Our *Case 1* corresponds to cases 1-3 of Stram and Lee (1994).

Case 2: Same as in Case 1, but the test also includes a r-dimensional subvector of fixed effects,  $\beta_2$ , i.e.,  $H_0: \sigma_{22} = 0, \sigma_{12} = 0, \beta_2 = 0$  against  $\Sigma$  positive semidefinite and  $\Omega$  and as in the asymptotic distribution of 2 log Left in (A,4 ii +  $\Lambda$ ,4 ii +  $\mu$  )  $\pm$   $\mu$ 

<u>Case 3</u>:  $d=q+k$  and

$$
\mathbf{\Sigma} = \left(\begin{array}{cc} \mathbf{\Sigma}_{11} & \mathbf{\Sigma}_{12} \\ \\ \mathbf{\Sigma}_{12}' & \mathbf{\Sigma}_{22} \end{array}\right),
$$

where  $1$  is q  $\alpha$  is  $2$  is the asymptotic null distribution of  $\alpha$  is the asymptotic number of  $\alpha$  is  $\alpha$ for testing  $H_0$ :  $\Sigma_{22} = 0$  (and therefore  $\Sigma_{12} = 0$ ) against  $\Sigma$  positive semidefinite is a mixture of  $\chi^2$  distributions with degrees of freedom  $s, s + 1, \ldots, s + k$ , where  $s = 1$ 

This corresponds to Case 4 of Stram and Lee (1994). Note, however, that the degrees of freedom for the mixture indicated in their paper was in error. In Stram and Lee (1995) they corrected the maximum degrees of freedom to  $s + k$ , but not the minimum degrees of freedom. To see why the correct mixture is the one we stated above, reparameterize  $\Sigma = \text{diag}(\sigma) \text{R} \text{diag}(\sigma)$ , where  $\sigma$  is the vector of standard deviations,

i.e. the square roots of the diagonal values of  $\Sigma$ , and  $\mathbf{R} = (\rho_{ij})$  is the correlation matrix. Testing  $\Sigma_{22} = 0$  and  $\Sigma_{12} = 0$  is equivalent to testing  $\sigma_{q+1} = \ldots = \sigma_{q+k} = 0$ , and  $\rho_{ij} = 0, i > j > k$ ; that is, k variance parameters tested on the boundary and s unconstrained correlation parameters. The result then follows along the same lines as in Case 7 of Self and Liang (1987). The mixing probabilities, however, are not directly available in general, and simulation methods may be used to estimate the mixing probabilities, or to estimate the null distribution itself. See Self and Liang (1987) and Stram and Lee (1994) for further discussion.

If, in addition, the condition  $\beta_1 = 0$  is part of the null hypothesis, then the asymptotic distribution of  $-2 \log LR$  is a  $\chi^2$  mixture with degrees of freedom  $s+r,\ldots,s+r+k$ . Case 4: Another situation of interest is when in the full model  $\Sigma_{12} = 0$  and  $\Sigma_{22}$  is diagonal. Similarly to Case 3, the asymptotic null distribution for testing  $\Sigma_{22} = 0$  is a  $\chi^2$  mixture with degrees of freedom 0 through k.

Remark The above asymptotic results are obtained under the assumption that the number of clusters,  $n$ , goes to infinite. For small  $n$ , the approximation by the mixture distributions given above may not be accurate. Crainiceanu and Ruppert (2004) showed that, for balanced linear one-way ANOVA with a single variance component, the mass at zero is larger than 0.5 when  $n$  is finite. We further discuss this issue in the simulation section.

#### 4.2 Profile AIC under PHMM

The PHMM was our original motivation for developing the profile AIC. When the focus is on the fixed effects  $\beta$  and the variance components  $\Sigma$ , the pAIC is given by (11), where p is the number of parameters in  $\beta$  and  $\Sigma$ . Computation of the likelihood term in (11) is addressed in the next subsection.

As a special case, when there are only fixed effects in the proportional hazards

model, the profile  $AIC$  is also given by (11), where p is the dimension of the regression parameter  $\beta$ . The profile likelihood in this case is the partial likelihood (Cox, 1975; Murphy and van der Vaart, 2000). This AIC has been previously used, for example, by Verweij and van Houwelingen (1995), although no formal justication has been given as an unbiased estimate of a dened Akaike information. Murphy and van der Vaart  $(2000)$  verified the conditions for the quadratic expansion  $(6)$  in this case. The validity of this AIC as an unbiased estimate of an Akaika information can also be shown directly, using the facts that asymptotically the partial likelihood score has zero expectation, and the second derivative of the log partial likelihood gives the observed information for  $\rho$  (Andersen and Gill, 1982).

## 4.3 Computing the likelihood under PHMM

For the FHMM we computed  $\sigma$  using an EM-type algorithm, see valua and Xu (2000). To compute the likelihood ratio statistic and the pAIC, only the maximum of the full likelihood function given in (4) is needed, since  $p_1(\psi) = \log L(\psi)$ . The likelihood function (4) is, in general, an intractable integral of dimension d. Here we consider three  $\min$  intertributing  $\iota(\nu) = \log L(\nu)$ . Laplace approximation, reciprocal importance sampling (RIS, Gelfand and Day, 1994), and bridge sampling (BS, Meng and Wong, 1996). Laplace approximation is computationally simple, but it is less accurate when  $n_i$ , the number of observations per cluster, is small. RIS and BS provide a numerically unbiased estimator for  $\iota(\nu)$  regardless of  $n_i$ , at an additional computational expense. We will compare the performance of the three methods in simulations and data analysis.

In the following we denote  $\mathbf{b} = (\mathbf{b}_1, ..., \mathbf{b}_n)$  and  $\mathbf{y} = (\mathbf{y}_1, ..., \mathbf{y}_n)$ .

Laplace approximation. This general method of computing integrals (see, e.g., Tierney and Kadane, 1986) is based on a normal approximation to the posterior distribution of the non-normalized integrand in (4),  $p(\mathbf{y}_i)p(\mathbf{b}_i|\mathbf{y}_i)$ , and is justified asymp-

**Collection of Biostatistics** 

**Research Archive** 

totically, as  $n_i \to \infty$ . The approxmation for cluster i is given by the formula:

$$
l_L^{(i)} = (d/2) \log(2\pi) + (1/2) \log |\hat{V}_i| + \log p(\mathbf{y}_i|\hat{\mathbf{b}}_i, \hat{\theta}) + \log p(\hat{\mathbf{b}}_i|\hat{\Sigma}),
$$
 (12)

where  $\mathbf{v}_i = \mathbf{E}(\mathbf{v}_i | \mathbf{y}_i, v)$ ,  $v_i = \text{var}(\mathbf{v}_i | \mathbf{y}_i, v)$  are the posterior mean and variance or the random eneros (DiCiccio et al., 1991). We compute  $\mathbf{v}_i$  and  $v_i$  using MCMC sample averages after convergence of the EM algorithm. Alternatively,  $\mathbf{v}_i, v_i$  can be taken as the posterior mode and inverse negative curvature or  $p(\mathbf{b}_i | \mathbf{y}_i, v)$ , respectively. We compute the Laplace approximation separately for each cluster, and let

$$
l_L = \sum_{i=1}^{n} l_L^{(i)} = (nd/2) \log(2\pi) + (1/2) \log |\hat{V}| + \log p(\mathbf{y}|\hat{\mathbf{b}}, \hat{\theta}) + \log p(\hat{\mathbf{b}}|\hat{\Sigma}), \qquad (13)
$$

where  $\mathbf{v} = \mathbf{E}(\mathbf{v}|\mathbf{y}, v)$  and  $\mathbf{v} = \mathbf{v}$ ar( $\mathbf{v}$ jy;  $v$ ). Note that Ripatti and Palmgren (2000) and Therneau and Grambsch (2000) used Laplace approximation for estimation of  $\theta$ in PHMM.

**Reciprocal importance sampling.** Let  $p_0(\mathbf{b})$  be the density of a fully specified approximating distribution to  $p(\mathbf{b}|\mathbf{y}, \theta)$ , for example, the normal density  $p_0(\mathbf{b})$  from  $N(\mathbf{D}, V)$ . If  $\mathbf{D}^{(1)}$ ,...,  $\mathbf{D}^{(2)}$  is a MCMC sample from  $p(\mathbf{D}|\mathbf{y}, \sigma)$ , then the reciprocal  $\min$  portance sampling estimator of  $\ell(\nu)$  is

$$
l_R = l_L - \log A,\tag{14}
$$

where

$$
A = \frac{1}{M} \sum_{k=1}^{M} \exp\{v(\mathbf{b}^{(k)})\}
$$
(15)

and

$$
v(\mathbf{b}) = l_L + \log p_0(\mathbf{b}) - \log p(\mathbf{y}, \mathbf{b}|\hat{\theta}).
$$
\n(16)

For numerical accuracy, the computations are done on the logarithmic scale as in (16). The orientation is defined in the case of  $\mathcal{L}$  and  $\mathcal{L}$  are case log A.However, using  $\mathcal{L}$ the Laplace approximation  $l<sub>L</sub>$  as a "point of reference" in (16) greatly improves the

numerical accuracy of  $l_R$ . A simple probabilistic argument shows that indeed A in (15) is a monte Carlo unbiased estimator of  $\exp\{tL - \epsilon(v)\}\,$ , see Geliand and Day (1994) for details.

The sampling and computation for  $l_R$  are straightforward to implement. The following result shows that in practice it is more efficient to compute  $l_R$  separately for each cluster.

**Proposition 1** Assume that  $l_R$  is computed as in  $(14)$  over the whole dataset, and  $l_R$  is the same except computed cluster-by-cluster. More precisely,  $l_R = \sum_{i=1}^n l_R^{(i)}$ , where  $l_R^{(i)}=l_L^{(i)}-\log A_i$ ,  $l_L^{(i)}$  is given by (12), and  $A_i=\sum_k \exp\{v(\mathbf{b}_i^{(k)})\}/M$ . Put  $A~=~\prod_{i=1}^n A_i,~so~that~l_R~=~l_L~-~\mathrm{log}\,A. ~~~Then~ both~l_R~and~l_R~converge~to~l(\theta)~with~$ probability one, and the sampling variance of A is at least as large as the sampling variance of  $A$ .

See Appendix for proof.

**Bridge sampling.** Assume that the Monte Carlo samples  $D^{(1)}, \ldots, D^{(m)}$  from  $p(\mathbf{b}|\mathbf{y},\sigma)$  and  $\mathbf{u}^{\vee},\dots,\mathbf{u}^{\vee}$  from  $p_0(\mathbf{b})$  are both available, where  $p_0(\mathbf{b})$  is a fully  $\mathbf s$  pecified approximation to  $p(\mathbf b|\mathbf y, v)$ , as described for RIS above. The bridge sampling  $\mu$  and wong, 1990) estimator for  $\mu$   $\mu$  is given by

$$
l_B = \log(B) - \log(C) + l_L, \tag{17}
$$

where

$$
B = \frac{1}{M_0} \sum_{k=1}^{M_0} [1 + \exp\{v(\mathbf{u}^{(k)})\}]^{-1}
$$
(18)

$$
C = \frac{1}{M} \sum_{k=1}^{M} [1 + \exp\{-v(\mathbf{b}^{(k)})\}]^{-1}.
$$
 (19)

It is again more efficient to compute  $l_B$  separately for each cluster and then combine the results, as in Proposition 1.

The three methods will be compared using simulation experiments in the next section. **RESS REPOSITORY** 

## 5 Simulation experiments

In this section we carry out simulations to compare the accuracy of the three methods described above for calculating the likelihood values, and to study the finite sample distribution of the likelihood ratio statistic.

We simulate data under model (1) with a single binary covariate Z,  $\beta = 1.5$ ,  $\lambda_0(t) = 1$ , and no random effects. The censoring distribution is Uniform  $(0, \tau)$ , where  $\tau$  is chosen to achieve about 15% censoring. We then fit model (1) with a random intercept, i.e.  $\lambda_{ij}(t) = \lambda_0(t) \exp(\beta Z_{ij} + b_i)$ . Different combinations of numbers of clusters and clusters sizes (n  $\ldots$ ) are used. In Figure 1 the used interesting successive computed using the three methods described in the last section. We see that reciprocal importance sampling (RIS) and bridge sampling (BS) have extremely close agreement in computing the likelihood (ratio) for all cases. For the number of observations per cluster  $n_i = 20$  Laplace approximation also gives similar results to RIS and BS. For  $n_i = 2$ , however, there are discrepancies between Laplace approximation and RIS or BS. The discrepancies increase with the number of clusters n since the log likelihood is the sum of that from each cluster, and the overall discrepancies are the sums of the discrepancies from each cluster.

In Figure 1 the ordered likelihood ratio statisitcs from 100 simulations are plotted against the theoretical mixture distribution quantiles. The asymptotic results for the null distribution of the likelihood ratio statistic requires that the number of clusters n is the 1000 for 1000 (lower panels) we compare the distribution of the distribution of the likelihood of the ratio statistic with its asymptotic distribution given in Case 1 of Section 4.1, i.e. a 50:50 mixture of point mass at zero and  $\chi_1^2$ . In Figure 1 'p0' denotes the probability of point mass at zero. For  $n = 10$  (upper panels) the asymptotic distribution does not appear to provide good approximation, and we use the result of Crainiceanu and Ruppert (2004) on linear mixed models (balanced one-way ANOVA) as a guideline,

i.e. a 65:35 mixture of point mass at zero and  $\chi_1^2$ . Note that their result requires the cluster size  $n_i \to \infty$  while keeping the number of clusters *n* fixed.

There is a clear effect of the number of observations per cluster on the null distribution of the likelihood ratio. For  $n_i = 20$  the empirical distributions of the computed likelihood ratio statistics agree reasonably well with their theoretical distributions according to the plots, for both  $n = 100$  and  $n = 10$ . But for  $n_i = 2$  even the distributions of the likelihood ratio values computed using RIS and BS have a clear departure from the theoretical mixtures. As mentioned before, for  $n = 10$  Crainiceanu and Ruppert's result requires that  $n_i$  be reasonably large. It is interesting to note that the departure also exists for  $n_i = 2$  and  $n = 100$ . The asymptotic mixture of 50:50 is theorectically asymptotic distribution does seem to provide a reasonable approximation for  $n = 100$ and  $n_i = 20$ . For  $n_i = 2$  we noticed (data not shown here) that the distribution of the likelihood ratio statisitcs (computed using RIS and BS) is much better approximated by the  $50:50$  mixture when n is as large as 250.

#### An example 6

In this section we consider the multi-center non-small cell lung cancer trial that was used as an example in Vaida and Xu (2000). The trial enrolled 579 patients from 31 institutions. The primary endpoint was patient death. There were two randomized treatment arms in the trial, a standard chemotherapy (CAV) arm and an alternating regimen (CAV-HEM) arm. Other important covariates that affected patient survival were: presence or absence of bone metastases, presence or absence of liver metastases, performance status at study entry and whether there was weight loss prior to entry. Gray (1995) used a score test for the existence of random treatment effect, and found it to be signicant.

In the following we mainly consider the three nested models of Vaida and Xu (2000); they are named Models 1-3 in Table 1. They all include the fixed effects of the five covariates. Model 1 includes no random effect; Model 2 includes a random treatment effect; and Model 3 includes random treatment and random bone metastases effects. The estimate of the other variance components corresponding to potential random effects for the rest three of the covariates, as well as random center effect on the baseline hazard function (see also Gray, 1995), converged to zero during the EM algorithm (Vaida and Xu, 2000). The parameter estimates under the three models were given in Table 1 of Vaida and Xu (2000). Table 1 here gives minus twice the log likelihood values for the models, computed using Laplace approximation, reciprocal importance sampling and bridge sampling for models 2 and 3. Note that the likelihood can be computed directly when there are no random effects, and such is the case for Models 1 and 0 (see below). The likelihood values for Models 2 and 3 are computed after 50 EM steps where the maximum likelihood estimate has converged; the sample sizes for Gibbs sampler during MCEM are 100 initially and increased to 1000 for the last 10 EM steps. The Monte Carlo sample sizes for RIS and BS are 1000, respectively. >From the table we see that the values of the log likelihoods agree well among the three computational methods.

As seen in the table, if we are to test Model 2 versus Model 1 using the likelihood ratio statistic, its sampling distribution under Model 1 is asymptotically  $(\chi_0 + \chi_1)/2,$ according to Case 1 of Section 4.1, with critical value of 2.71 at .05 significance level. Model 1 is then rejected in favor of Model 2. Similarly, to test Model 3 versus Model 2, the likelihood ratio statistic is again asymptotically  $(\chi_0^- + \chi_1^-)/2$  under Model 2. This is a special case of Case 4, and the mixing probabilities can be derived directly as in Case 1. Therefore Model 2 is rejected in favor of Model 3. Note that the finite sample distribution we considered in Section 5 puts more point mass at zero, leading to even smaller critical values for the likelihood ratio statistic.

**Collection of Biostatistics Research Archive** 

21

We can also compare Models 1 and 3 directly. Under Model 1 the asymptotic distribution of the likelihood ratio statistic is a mixture of  $\chi_{0}^{-},\,\chi_{1}^{-}$  and  $\chi_{2}^{-}.$  This is againn Case  $4$  in Section 4.1. The mixing probabilities are not straightforward to compute; however, given that the 0.95 quantile of  $\chi_2^-$  is 5.99, and that the same quantile for the mixture is smaller, Model 1 is therefore rejected in favor of Model 3.

Finally, Model 0 is the Cox model with only fixed effects for the 4 covariates other than treatment. The comparison of Model 0 versus Model 2 provides an illustration for Case 2 of Section 4.1, i.e. neither the fixed nor the random treatment effect is significant. Here  $q = 0$  and  $r = 1$ , so the null asymptotic distribution of the likelihood ratio statistic is  $(\chi_1^* + \chi_2^*)/2$ . It is again easy to see that Model 0 is rejected in favor of Model 2 at 0.05 signcance level.

Alternatively, we can use the profile  $AIC$  to compare the nested models. From the table it is also clear that the larger models are chosen by the criterion.

#### **Discussion**  $\overline{7}$

In this paper we established the asymptotic properties of the nonparametric maximum likelihood estimator under the proportional hazards mixed effects model. Motivated by model selection problems under PHMM, we developed the profile likelihood ratio test and a prole Akaike information criterion that are generally applicable to models with nuisance parameters. The development was based on the asymptotic quadratic expansion of the log profile likelihood function. The profile likelihood ratio test for the null hypothesis that lies in the interior of the parameter space was given in Murphy and van der Vaart (2000); here we further developed it for testing on the boundary. The prole AIC has not been previously proposed in the literature, to our best knowledge. It applies to both parametric and semiparamtric models, and for the latter type of models the focus is on the finite dimenstional parameter. The AIC approach does

not encounter the boundary problem as in hypothesis testing. The profile  $AIC$  also provides a theoretical justification for the use of the partial likelihood in the AIC under the classic Cox model.

Model selection has been an area of growing interest in the recent years. In this paper we restricted our attention to the classic derivation of the Akaike information criterion. However we acknowledge, as Longford  $(2005)$  pointed out, that, whatever the selection criterion, single-model based inference can be inherently biased. Alternatives may include the use of a mixture of plausible models, and the focused information criteria of Claeskens and Hjort (2003). The associated new challenges of such improvements in practice are model interpretability and variability of inferences following the model averaging or selection.

For computation of the maximized likelihood, the Laplace approximation is the most straightforward but is only accurate when the cluster sizes are reasonably large. In view of the  $MCEM$  algorithm that is used to fit the PHMM, the additional computation of RIS or BS is often comparable to one step of the MCEM. Therefore we include RIS and BS as default in our computational program.

Finally, under linear mixed models when the interest lies in the inference of the random effects themselves, Vaida and Blanchard  $(2005)$  propose a conditional AIC using the notion of effective degrees of freedom. The usefulness of conditional inference carries over to PHMM, and it is our future work to develop a conditional AIC under the PHMM. Additionally, the finite sample distribution of the likelihood ratio statistic for testing zero variance components is another area that requires further work.

PROOF OF THEOREM 1. To prove consistency we follow methods used by  $m$ urphy (1994) and Zeng et al. (2000). First prove  $\Lambda_n(\cdot)$  is bounded on  $[0, t]$ . We **Collection of Biostatistics Research Archive** 

then invoke the compactness of the parameter space and Helly's selection theorem to conclude the existence of convergent subsequence of  $\{\theta_n\}$ . Finally we show the limit of this subsequence must be  $\theta_0$ .

 $\mathcal{L}$  be show  $\Lambda_n(\cdot)$  has an upper bound int  $[0, t]$ . First let

$$
\bar{\Lambda}_n(t) = \sum_{ij} \frac{\delta_{ij} (1 - Y_{ij}(t))}{\sum_{kl} Y_{kl}(X_{ij}) e^{\beta'_0 \mathbf{Z}_{kl}} \mathbf{E}_{\theta}(e^{\mathbf{b}'_k \mathbf{W}_{kl}} | \mathbf{y}_k)},
$$
\n
$$
a_i(t) = \sum_{j=1}^{n_i} \int_{u=0}^t \{ dN_{ij}(u) - Y_{ij}(u) e^{\beta'_0 \mathbf{Z}_{ij}} \mathbf{E}_{\theta}(e^{\mathbf{b}'_i \mathbf{W}_{ij}} | \mathbf{y}_i) d\Lambda_0(u) \},
$$
\n
$$
f_n(u) = n^{-1} \sum_{i=1}^n \sum_{j=1}^{n_i} Y_{ij}(u) e^{\beta'_0 \mathbf{Z}_{ij}} \mathbf{E}_{\theta}(e^{\mathbf{b}'_i \mathbf{W}_{ij}} | \mathbf{y}_i).
$$

We show  $\sup_{t\in[0,\tau]} |h_n(t) - h_0(t)| \to 0$  almost surely.

Note that  $\{a_i(t): i = 1, 2, \ldots\}$  is a mean zero independent sequence for fixed t. Also, by the boundedness assumptions on  $\mathbf{W}_{ij}$  and  $n_i$ :

$$
\mathrm{E}_\theta(e^{\mathbf{b}_i'\mathbf{W}_{ij}}|\mathbf{y}_i)
$$

for some constant  $B_{\Sigma_0}$ . Similarly  $e^{\rho_0 L_{ij}} < B_{\beta_0} < \infty$ , and since  $a_i(t)$  is bounded for any  $t \in [0, \tau]$  we have  $\text{Var}(a_i(t))$  is bounded and by the SLLN  $n^{-1} \sum_i a_i(t) \to 0$  almost surely.

Similarly,  $f_n(u) - \mathrm{E}(f_n(u)) \to 0$  almost surely. Since  $\mathrm{E}[Y_{ij}(u)e^{D_0 L_{ij}} \mathrm{E}_\theta(e^{\mathbf{D}_i \cdot \mathbf{W}_{ij}} | \mathbf{y}_i)]$ is bounded and  $\gamma$  from zero, there exists some c1  $\ell$  0 such that eventually fn(u)  $\pm$  1 almost surely. Likewise, since  $\alpha$  is the since  $\alpha$  is the case of  $\alpha$  is a such that for  $\alpha$   $\alpha$ 

 $c_2$ .

Now consider

$$
\sum_{ij} \int_{u=0}^t \left\{ dN_{ij}(u) - Y_{ij}(u) e^{\beta'_0 \mathbf{Z}_{ij}} \mathbf{E}_{\theta}(e^{\mathbf{b}'_i \mathbf{W}_{ij}} | \mathbf{y}_i) d\bar{\Lambda}_n(u) \right\} = 0, \tag{20}
$$

since by switching the order of summation,

$$
LHS = \sum_{ij} \left\{ \delta_{ij} (1 - Y_{ij}(t)) - \sum_{kl} \frac{Y_{ij}(X_{kl}) e^{\beta'_0 \mathbf{Z}_{ij}} E_{\theta}(e^{\mathbf{b}'_i \mathbf{W}_{ij}} | \mathbf{y}_i) \delta_{kl} (1 - Y_{kl}(t))}{\sum_{rs} Y_{rs}(X_{kl}) e^{\beta'_0 \mathbf{Z}_{rs}} E_{\theta}(e^{\mathbf{f}' \mathbf{W}_{rs}} | \mathbf{y}_r)} \right\}
$$
  
= 
$$
\sum_{ij} \delta_{ij} (1 - Y_{ij}(t)) - \sum_{kl} \left\{ \frac{\sum_{ij} Y_{ij}(X_{kl}) e^{\beta'_0 \mathbf{Z}_{ij}} E_{\theta}(e^{\mathbf{b}'_i \mathbf{W}_{ij}} | \mathbf{y}_i) \delta_{kl} (1 - Y_{kl}(t))}{\sum_{rs} Y_{rs}(X_{kl}) e^{\beta'_0 \mathbf{Z}_{rs}} E_{\theta}(e^{\mathbf{f}' \mathbf{W}_{rs}} | \mathbf{y}_r)} \right\}
$$
  
= 0.

Now by adding and subtracting  $dN_{ij}(u)$  in (20) we have for fixed t

$$
\int_{u=0}^{t} f_n(u) d(\Lambda_0 - \bar{\Lambda}_n)(u) = n^{-1} \sum_{ij} \int_{u=0}^{t} Y_{ij}(u) e^{\beta'_0 \mathbf{Z}_{ij}} \mathbf{E}_{\theta}(e^{\mathbf{b}'_i \mathbf{W}_{ij}} | \mathbf{y}_i) d(\Lambda_0 - \bar{\Lambda}_n)(u)
$$
  
\n
$$
= n^{-1} \sum_{ij} \int_{u=0}^{t} \{dN_{ij}(u) - Y_{ij}(u) e^{\beta'_0 \mathbf{Z}_{ij}} \mathbf{E}_{\theta}(e^{\mathbf{b}'_i \mathbf{W}_{ij}} | \mathbf{y}_i) d\Lambda_0(u) \}
$$
  
\n
$$
= n^{-1} \sum_{i=1}^{n} a_i(t)
$$
  
\n
$$
\rightarrow 0 \text{ a.s.,}
$$

by SLLN. Futhermore

$$
c_1 \int_{u=0}^t d(\Lambda_0 - \bar{\Lambda}_n)(u) \le \int_{u=0}^t f_n(u) d(\Lambda_0 - \bar{\Lambda}_n)(u) \to 0 \text{ a.s.}
$$

and

$$
c_2 \int_{u=0}^t d(\Lambda_0 - \bar{\Lambda}_n)(u) \ge \int_{u=0}^t f_n(u) d(\Lambda_0 - \bar{\Lambda}_n)(u) \to 0 \text{ a.s.}.
$$

Since  $c_1$  and  $c_2$  are both positive, we must have

$$
\int_{u=0}^t d(\Lambda_0 - \bar{\Lambda}_n)(u) \to 0 \text{ a.s.},
$$

which implies  $\Lambda_n(t) \to \Lambda_0(t)$  a.s. for all  $t \in [0, t]$ . Followise convergence of nondecreasing functions to a continuous limit implies local (on  $[0, \tau]$  in particular) uniform continuity.



Since  $\mu_n$ ,  $\mathbf{z}_n$ ,  $\mathbf{z}_{kl}$ , and Wkl are in compact sets, there exists some ninte, possibly negative C such that

$$
\hat{\boldsymbol{\beta}}_n^{\prime} \mathbf{Z}_{kl} + \log \mathrm{E}_{\hat{\theta}_n}[e^{\mathbf{b}_k^{\prime} \mathbf{W}_{kl}}|\mathbf{y}_k] \geq \boldsymbol{\beta}_0^{\prime} \mathbf{Z}_{kl} + \log \mathrm{E}_{\theta_0}[e^{\mathbf{b}_k^{\prime} \mathbf{W}_{kl}}|\mathbf{y}_k] + C.
$$

Therefore

$$
\hat{\Lambda}_n(\tau) = \sum_{ij} \frac{\delta_{ij}(1 - Y_{ij}(\tau))}{\sum_{kl} Y_{kl}(X_{ij}) \exp{\{\hat{\beta}'_n \mathbf{Z}_{kl} + \log E_{\hat{\theta}_n}[e^{\mathbf{b}'_k \mathbf{W}_{kl}}|\mathbf{y}_k]\}}}
$$
\n
$$
\leq \sum_{ij} \frac{\delta_{ij}(1 - Y_{ij}(\tau))}{\sum_{kl} Y_{kl}(X_{ij}) \exp{\{\beta'_0 \mathbf{Z}_{kl} + \log E_{\theta_0}[e^{\mathbf{b}'_k \mathbf{W}_{kl}}|\mathbf{y}_k] + C\}}}
$$
\n
$$
= e^{-C} \bar{\Lambda}_n(\tau) \to e^{-C} \Lambda_0(\tau).
$$

 $\mathcal{S}$ ic $\mu$   $\alpha$ . Since  $\Lambda$  has an upper bound almost surely, and  $\mu_n$  and  $\Delta_n$  are in compact sets, we can use Helly's selection theorem to establish a convergent subsequence which we now denote by  $\sigma_n = (\Lambda_n, \rho_n, \Sigma_n)$  with limit  $\sigma$ .

Taking limits of both sides of

$$
\hat{\Lambda}_n(t) = \int_0^t \frac{\sum_{kl} Y_{kl}(u) \exp\{\beta'_0 \mathbf{Z}_{kl} + \log E_{\theta_0}[e^{\mathbf{b}'_k} \mathbf{W}_{kl} | \mathbf{y}_k]\}}{\sum_{kl} Y_{kl}(u) \exp\{\hat{\beta}'_n \mathbf{Z}_{kl} + \log E_{\hat{\theta}_n}[e^{\mathbf{b}'_k} \mathbf{W}_{kl} | \mathbf{y}_k]\}} d\bar{\Lambda}_n(u)
$$
(21)

we see that  $\Lambda$  is absolutely continuous with respect to  $\Lambda_0$ . Furthermore,  $\Lambda$  (t) is differentiable with respect to t and  $a\Lambda_n(t)/a\Lambda_n(t)$  converges to  $a\Lambda_-(t)/a\Lambda_0(t)$ . Note that the finite sample likelihood as expressed via (3) has no finite maximum, since  $\lambda$ is free to go to infinity at any  $X_{ij}$ . We restrict  $\Lambda$  to be right continuous with jumps at  $X_{ij}$ ; and for cluster *i*, conditional on the random effect  $\mathbf{b}_i$ , we let the log-likelihood be

$$
l_i = l_i(\boldsymbol{\beta}, \lambda; \mathbf{y}_i | \mathbf{b}_i) = \sum_{j=1}^{n_i} \{ \delta_{ij} \log \Lambda \{ X_{ij} \} + \delta_{ij} (\boldsymbol{\beta}' \mathbf{Z}_{ij} + \mathbf{b}'_i \mathbf{W}_{ij}) - \Lambda (X_{ij}) e^{\boldsymbol{\beta}' \mathbf{Z}_{ij} + \mathbf{b}'_i \mathbf{W}_{ij}} \}, (22)
$$

where  $\Lambda\{t\}$  is the size of the jump in  $\Lambda$  at t. The likelihood of the observed data,  $L_n(\theta)$ , is still as defined in (3) and we let  $l_n(\theta) = \log L_n(\theta)$ . In place of  $\Lambda_0$ , which is

continuous at  $X_{ij}$ , we use  $X_{ij}$ . In particular we have.

$$
0 \leq n^{-1} \{ l_n(\hat{\boldsymbol{\beta}}_n, \hat{\boldsymbol{\Sigma}}_n, \hat{\boldsymbol{\Lambda}}_n) - l_n(\boldsymbol{\beta}_0, \boldsymbol{\Sigma}_0, \bar{\boldsymbol{\Lambda}}_n) \}
$$
  
\n
$$
= n^{-1} \sum_{i=1}^n \log \left\{ \int_{\mathbf{b}} R_i(\hat{\boldsymbol{\beta}}_n, \hat{\boldsymbol{\Lambda}}_n, \mathbf{b}) \mathbf{p}(\mathbf{b}, \hat{\boldsymbol{\Sigma}}_n) d\mathbf{b} \right\}
$$
  
\n
$$
- n^{-1} \sum_{i=1}^n \log \left\{ \int_{\mathbf{b}} R_i(\boldsymbol{\beta}_0, \bar{\boldsymbol{\Lambda}}_n, \mathbf{b}) \mathbf{p}(\mathbf{b}, \boldsymbol{\Sigma}_0) d\mathbf{b} \right\}
$$
  
\n
$$
+ n^{-1} \sum_{i=1}^n \sum_{j=1}^{n_i} \delta_{ij} \log(\hat{\boldsymbol{\Lambda}}_n \{X_{ij}\} / \bar{\boldsymbol{\Lambda}}_n \{X_{ij}\})
$$

where

$$
R_i(\boldsymbol{\beta}, \Lambda, \mathbf{b}) = \prod_{j=1}^{n_i} \exp[\delta_{ij}(\boldsymbol{\beta}' \mathbf{Z}_{ij} + \mathbf{b}' \mathbf{W}_{ij}) - \Lambda(X_{ij}) \exp(\boldsymbol{\beta}' \mathbf{Z}_{ij} + \mathbf{b}' \mathbf{W}_{ij})].
$$

Letting  $n \to \infty$  we have

$$
0 \leq E \log \left\{ \int_{\mathbf{b}} R_i(\boldsymbol{\beta}^*, \Lambda^*, \mathbf{b}) \mathrm{p}(\mathbf{b}, \boldsymbol{\Sigma}^*) d\mathbf{b} \prod_{j=1}^{n_i} \lambda^*(X_{ij})^{\delta_{ij}} \times \left( \int_{\mathbf{b}} R_i(\boldsymbol{\beta}_0, \Lambda_0, \mathbf{b}) \mathrm{p}(\mathbf{b}, \boldsymbol{\Sigma}_0) d\mathbf{b} \prod_{j=1}^{n_i} \lambda_0 (X_{ij})^{\delta_{ij}} \right)^{-1} \right\}.
$$

Because the right side is negative the Kullback-Leibler information we have

$$
\int_{\mathbf{b}} R_i(\boldsymbol{\beta}^*, \Lambda^*, \mathbf{b}) \mathrm{p}(\mathbf{b}, \boldsymbol{\Sigma}^*) d\mathbf{b} \prod_{j=1}^{n_i} \lambda^* (X_{ij})^{\delta_{ij}} = \int_{\mathbf{b}} R_i(\boldsymbol{\beta}_0, \Lambda_0, \mathbf{b}) \mathrm{p}(\mathbf{b}, \boldsymbol{\Sigma}_0) d\mathbf{b} \prod_{j=1}^{n_i} \lambda_0 (X_{ij})^{\delta_{ij}}
$$

or

$$
\int_{\mathbf{b}} \prod_{j=1}^{n_i} \lambda^* (X_{ij})^{\delta_{ij}} \exp[\delta_{ij} (\boldsymbol{\beta}^{*'} \mathbf{Z}_{ij} + \mathbf{b}' \mathbf{W}_{ij}) - \Lambda^* (X_{ij}) \exp(\boldsymbol{\beta}^{*'} \mathbf{Z}_{ij} + \mathbf{b}' \mathbf{W}_{ij})] p(\mathbf{b}, \mathbf{\Sigma}^*) d\mathbf{b}
$$
\n
$$
= \int_{\mathbf{b}} \prod_{j=1}^{n_i} \lambda_0 (X_{ij})^{\delta_{ij}} \exp[\delta_{ij} (\boldsymbol{\beta}_0' \mathbf{Z}_{ij} + \mathbf{b}' \mathbf{W}_{ij}) - \Lambda_0 (X_{ij}) \exp(\boldsymbol{\beta}_0' \mathbf{Z}_{ij} + \mathbf{b}' \mathbf{W}_{ij})] p(\mathbf{b}, \mathbf{\Sigma}_0) d\mathbf{b}
$$
\n(23)

Now we use techniques adapted from Zeng et al. (2005) to conclude  $\sigma = \sigma_0$ . Fix some k in  $1, \ldots, n_i$ . For  $j = 1, \ldots, k$ , let  $\delta_{ij} = 1, X_{ij} = 0$  in (23) and note that we **Collection of Biostatistics** 27 Research Archive

assume  $\Lambda$  (0)  $\Xi$   $\Lambda$ <sub>0</sub>(0)  $\Xi$  0. If  $j = \kappa + 1, \ldots, n_i$  and  $\sigma_{ij} = 0$ , we replace  $\Lambda_{ij}$  with  $\tau$ . Otherwise, if  $j = k + 1, \ldots, n_i$  and  $\delta_{ij} = 1$ , we integrate  $X_{ij}$  from 0 to  $\tau$ . We get:

$$
\int_{\mathbf{b}} \prod_{j=1}^{k} \lambda^{*}(0) \exp[\boldsymbol{\beta}^{*'} \mathbf{Z}_{ij} + \mathbf{b}' \mathbf{W}_{ij}]
$$
\n
$$
\times \prod_{j=k+1}^{n_i} \{ \exp[-\Lambda^{*}(\tau) \exp(\boldsymbol{\beta}^{*'} \mathbf{Z}_{ij} + \mathbf{b}' \mathbf{W}_{ij})] \}^{1-\delta_{ij}}
$$
\n
$$
\times \prod_{j=k+1}^{n_i} \left\{ \int_{y=0}^{\tau} \lambda^{*}(y) \exp[\boldsymbol{\beta}^{*'} \mathbf{Z}_{ij} + \mathbf{b}' \mathbf{W}_{ij} - \Lambda^{*}(y) \exp(\boldsymbol{\beta}^{*'} \mathbf{Z}_{ij} + \mathbf{b}' \mathbf{W}_{ij})] dy \right\}^{\delta_{ij}} p(\mathbf{b}, \mathbf{\Sigma}^{*}) d\mathbf{b}
$$
\n
$$
= \int_{\mathbf{b}} \prod_{j=1}^{k} \lambda_{0}(0) \exp[\boldsymbol{\beta}_{0}' \mathbf{Z}_{ij} + \mathbf{b}' \mathbf{W}_{ij}]
$$
\n
$$
\times \prod_{j=k+1}^{n_i} \left\{ \exp[-\Lambda_{0}(\tau) \exp(\boldsymbol{\beta}_{0}' \mathbf{Z}_{ij} + \mathbf{b}' \mathbf{W}_{ij})] \right\}^{1-\delta_{ij}}
$$
\n
$$
\times \prod_{j=k+1}^{n_i} \left\{ \int_{y=0}^{\tau} \lambda_{0}(y) \exp[\boldsymbol{\beta}_{0}' \mathbf{Z}_{ij} + \mathbf{b}' \mathbf{W}_{ij} - \Lambda_{0}(y) \exp(\boldsymbol{\beta}_{0}' \mathbf{Z}_{ij} + \mathbf{b}' \mathbf{W}_{ij})] dy \right\}^{\delta_{ij}} p(\mathbf{b}, \mathbf{\Sigma}_{0}) d\mathbf{b}
$$



Z <sup>b</sup> <sup>Y</sup> (0) exp[0Zij <sup>+</sup> b0Wij ] j=1 Yni ( ) exp(0Zij <sup>+</sup> <sup>b</sup>0Wij )] 1ij exp[ j=k+1 Yni ( ) exp(0Zij <sup>+</sup> <sup>b</sup>0Wij )] ij p(b; exp[ 1 )db j=k+1 Z <sup>b</sup> <sup>Y</sup> 0(0) exp[00Zij <sup>+</sup> <sup>b</sup>0Wij ] = j=1 Yni 0Zij <sup>+</sup> b0Wij )] 1ij exp[0( ) exp(0 j=k+1 Yni 0Zij <sup>+</sup> b0Wij )] ij p(b; 0)db: (24) 1 exp[0( ) exp(0 j=k+1 

Because  $\delta_{ij}$  are arbitrary, we sum the two sides of (24) over all possible  $\delta_{ij}$  to yield:

$$
\int_{\mathbf{b}} \prod_{j=1}^k \lambda^*(0) \exp[\boldsymbol{\beta}^{*'} \mathbf{Z}_{ij} + \mathbf{b}' \mathbf{W}_{ij}] p(\mathbf{b}, \boldsymbol{\Sigma}^*) d\mathbf{b} = \int_{\mathbf{b}} \prod_{j=1}^k \lambda_0(0) \exp[\boldsymbol{\beta}_0' \mathbf{Z}_{ij} + \mathbf{b}' \mathbf{W}_{ij}] p(\mathbf{b}, \boldsymbol{\Sigma}_0) d\mathbf{b}
$$

and

$$
\exp\left\{\sum_{j=1}^k \beta^{*'} \mathbf{Z}_{ij} + \frac{(\sum_{j=1}^k \mathbf{W}_{ij})' \mathbf{\Sigma}^*(\sum_{j=1}^k \mathbf{W}_{ij})}{2}\right\} \lambda^*(0)^k
$$
  
= 
$$
\exp\left\{\sum_{j=1}^k \beta_0' \mathbf{Z}_{ij} + \frac{(\sum_{j=1}^k \mathbf{W}_{ij})' \mathbf{\Sigma}_0 (\sum_{j=1}^k \mathbf{W}_{ij})}{2}\right\} \lambda_0(0)^k
$$

We assume  $\lambda$  (0)  $>0$ . Since the index set can be replaced by any subset of  $1,\ldots,n_i$ we have

$$
\mathbf{W}_{ij}' \mathbf{\Sigma}^* \mathbf{W}_{ij'} = \mathbf{W}_{ij}' \mathbf{\Sigma}_0 \mathbf{W}_{ij'}, \ j \neq j' : j, j' = 1, \ldots, n_i,
$$

and

$$
\beta^*'\mathbf{Z}_{ij} + \frac{\mathbf{W}'_{ij}\mathbf{\Sigma}^*\mathbf{W}_{ij}}{2} + \log \lambda^*(0)
$$
  
=  $\beta_0'\mathbf{Z}_{ij} + \frac{\mathbf{W}'_{ij}\mathbf{\Sigma}_0\mathbf{W}_{ij}}{2} + \log \lambda_0(0), \ j = 1, \dots, n_i$ 

Therefore, under C<sub>2</sub>,  $\boldsymbol{\Sigma} = \boldsymbol{\Sigma}_0$ ,  $\boldsymbol{\beta} = \boldsymbol{\beta}_0$ , and  $\boldsymbol{\lambda}(0) = \lambda_0(0)$ .

To show  $\Lambda^* = \Lambda_0$ , we manipulate the terms of (23) again. Let  $\delta_{i1} = 1$  and integrate  $X_{i1}$  from 0 to t. Also for  $j = 2, \ldots, n_i$ , if  $\delta_{ij} = 0$ , replace  $X_{ij}$  with  $\tau$  and if  $\delta_{ij} = 1$ integrate  $X_{ij}$  from 0 to  $\tau$ . Summing the result over all possible  $\{\delta_{ij} : j = 2,\ldots,n_i\},\$ this time we get

$$
\int_{\mathbf{b}} 1 - \exp[-\Lambda^*(t) \exp(\beta_0' Z_{i1} + \mathbf{b}' \mathbf{W}_{i1})] p(\mathbf{b}, \Sigma_0) d\mathbf{b}
$$
  
= 
$$
\int_{\mathbf{b}} 1 - \exp[-\Lambda_0(t) \exp(\beta_0' Z_{i1} + \mathbf{b}' \mathbf{W}_{i1})] p(\mathbf{b}, \Sigma_0) d\mathbf{b}.
$$
 (25)

Because both sides of (25) are strictly monotone in  $\Lambda$  (t) and  $\Lambda_0(t)$ , we have  $\Lambda_1(t) =$  $\Lambda_0(t)$ . Since  $\Lambda_0$  is non-decreasing and continuous, the pointwise convergence can be extended to uniform convergence on  $[0, \tau]$ .

29

**PROOF OF THEOREM 2.** To prove asymptotic normality and efficiency we invoke methods of Murphy (1995) and Zeng et al. (2005). Consider the set

$$
\mathcal{H} = \{ (\mathbf{h}_1, \mathbf{h}_2, h_3) : \mathbf{h}_1 \in \mathbf{R}^{d_1}, \mathbf{h}_2 \in \mathbf{R}^{d_2(d_2+1)/2},
$$
  

$$
h_3(\cdot) \text{ is a function on } [0, \tau]; \|\mathbf{h}_1\|, \|\mathbf{h}_2\|, \|\mathbf{h}_3\|_V \le 1 \}
$$
 (26)

where  $||h_3||_V$  denotes the total variation of  $h_3(\cdot)$  in  $[0, \tau]$ . We define a sequence of maps  $S \cap \{0\}$  and  $S \cap \{0\}$  and  $S \cap \{0\}$  of  $\{0,1\}$  and  $\{0,1\}$  $(\boldsymbol{\beta}, \boldsymbol{\Sigma}, \Lambda)$  into  $l^{\infty}(\mathcal{H})$  as:

$$
S_n(\boldsymbol{\beta}, \boldsymbol{\Sigma}, \Lambda)[\mathbf{h}_1, \mathbf{h}_2, h_3]
$$
  
\n
$$
\equiv n^{-1} \frac{d}{d\epsilon} l_n \left( \boldsymbol{\beta} + \epsilon \mathbf{h}_1, \boldsymbol{\Sigma} + \epsilon \mathbf{h}_2, \Lambda(t) + \epsilon \int_0^t h_3(s) d\Lambda(s) \right) \Big|_{\epsilon=0}
$$
  
\n
$$
\equiv A_{n1}[\mathbf{h}_1] + A_{n2}[\mathbf{h}_2] + A_{n3}[h_3]
$$

where  $\Sigma$  is treated as extended column vector consisting of the upper triangle elements; and  $A_{np}$ ,  $p = 1, 2, 3$ , are linear functionals on  $\mathbf{R}^{n_1}, \mathbf{R}^{n_2 \backslash n_2 + 2 \gamma - 2}$  and  $DV[0, \tau]$  (the space of functions with finite total variation in  $[0, \tau]$ ). If we let  $l_{\beta}$ ,  $l_{\Sigma}$  and  $l_{\Lambda}$  be the score functions for  $\beta$ ,  $\Sigma$ , and  $\Lambda$  (along  $\int_0^t 1 + \epsilon h_3(s) d\Lambda(s)$ ) for a single cluster, then

$$
A_{n1}[\mathbf{h}_1] = \mathcal{P}_n[\mathbf{h}'_1 l_\beta], A_{n2}[\mathbf{h}_2] = \mathcal{P}_n[\mathbf{h}'_2 l_\mathbf{\Sigma}], \text{ and } A_{n3}[h_3] = \mathcal{P}_n[l_\Lambda[h_3]]
$$

where  $P$  is empirical measure based on n independent clusters. We now indepe seek explicit expression for  $A_{np}$ . Recall the log likelihood

$$
n^{-1}l_n(\theta) = n^{-1} \sum_{i=1}^n \log \left\{ \int_{\mathbf{b}} R_i(\boldsymbol{\beta}, \Lambda, \mathbf{b}) \mathrm{p}(\mathbf{b}, \boldsymbol{\Sigma}) d\mathbf{b} \right\} + n^{-1} \sum_{i=1}^n \sum_{j=1}^{n_i} \delta_{ij} \log \Lambda \{ X_{ij} \}
$$

$$
R_i(\boldsymbol{\beta}, \Lambda, \mathbf{b}) = \exp \left\{ \sum_{j=1}^{n_i} \delta_{ij} (\boldsymbol{\beta}' \mathbf{Z}_{ij} + \mathbf{b}' \mathbf{W}_{ij}) - \Lambda(X_{ij}) \exp(\boldsymbol{\beta}' \mathbf{Z}_{ij} + \mathbf{b}' \mathbf{W}_{ij}) \right\}.
$$

Note that  $\hfill\blacksquare$ 

$$
\frac{\partial}{\partial \epsilon} R_i(\boldsymbol{\beta} + \epsilon \mathbf{h}_1, \Lambda, \mathbf{b})\big|_{\epsilon=0} = R_i(\boldsymbol{\beta}, \Lambda, \mathbf{b}) \sum_{j=1}^{n_i} \mathbf{h}'_1 \mathbf{Z}_{ij} (\delta_{ij} - \Lambda(X_{ij}) \exp(\boldsymbol{\beta}' \mathbf{Z}_{ij} + \mathbf{b}' \mathbf{W}_{ij}))
$$
  
Collection of Blostatistics  
Research Archive 30

Furthermore let  $\Lambda_{\epsilon}(t) = \int_0^t 1 + \epsilon h_3 d\Lambda$ , then  $\frac{\partial}{\partial \epsilon} \Lambda_{\epsilon}(t) = \int_0^t h_3(s) d\Lambda(s)$  and

$$
\frac{\partial}{\partial \epsilon} R_i (\boldsymbol{\beta}, \Lambda_{\epsilon}, \mathbf{b}) \big|_{\epsilon=0} = -R_i (\boldsymbol{\beta}, \Lambda, \mathbf{b}) \sum_{j=1}^{n_i} \int_0^{X_{ij}} h_3(s) d\Lambda(s) \exp(\boldsymbol{\beta}' \mathbf{Z}_{ij} + \mathbf{b}' \mathbf{W}_{ij}).
$$

Also  $\Lambda_{\epsilon}\lbrace t \rbrace = (1 + \epsilon h_3(t)) \Lambda \lbrace t \rbrace$ , so

$$
\frac{d}{d\epsilon}\log\Lambda_{\epsilon}\{t\}\big|_{\epsilon=0} = \frac{h_3(t)\Lambda\{t\}}{\Lambda_{\epsilon}\{t\}}\big|_{\epsilon=0} = h_3(t).
$$

If we fix  $\sim$  (H2) denote the matrix corresponding to the extended vector h2 and denote the  $\cdots$  operation on two matrices  $\bf{m}_1$  and  $\bf{m}_2$  to be trace( $\bf{m}_1\bf{m}_1$ ), then

$$
\frac{\partial}{\partial \epsilon} \mathbf{p}(\mathbf{b}; \mathbf{\Sigma} + \epsilon \mathbf{h}_2)|_{\epsilon=0} = \frac{\partial}{\partial \epsilon} |\mathbf{\Sigma} + \epsilon \mathbf{h}_2|^{-1/2} e^{-\mathbf{b}'(\mathbf{\Sigma} + \epsilon \mathbf{h}_2)^{-1} \mathbf{b}/2} |_{\epsilon=0}
$$

$$
= \left\{ \mathbf{b}' \mathbf{\Sigma}^{-1} \mathcal{D}(\mathbf{h}_2) \mathbf{\Sigma}^{-1} \mathbf{b}/2 - \mathbf{\Sigma}^{-1} \cdot \mathcal{D}(\mathbf{h}_2)/2 \right\} e^{-\mathbf{b}' \mathbf{\Sigma}^{-1} \mathbf{b}/2}.
$$

Finally, we can explicitly write  $\mathcal{A}_{np}$  as

$$
A_{n1}[\mathbf{h}_1] = n^{-1} \sum_{i=1}^n \left( \int_{\mathbf{b}} \sum_{j=1}^{n_i} \mathbf{h}_1^j \mathbf{Z}_{ij} \left( \delta_{ij} - \Lambda(X_{ij}) e^{\beta' \mathbf{Z}_{ij} + \mathbf{b}' \mathbf{W}_{ij}} \right) \times R_i(\boldsymbol{\beta}, \Lambda, \mathbf{b}) e^{-\mathbf{b}' \mathbf{\Sigma}^{-1} \mathbf{b}/2} d\mathbf{b} \right)
$$
  
\n
$$
\times \left( \int_{\mathbf{b}} R_i(\boldsymbol{\beta}, \Lambda, \mathbf{b}) e^{-\mathbf{b}' \mathbf{\Sigma}^{-1} \mathbf{b}/2} d\mathbf{b} \right)^{-1}
$$
  
\n
$$
A_{n2}[\mathbf{h}_2] = n^{-1} \sum_{i=1}^n \left( \int_{\mathbf{b}} {\mathbf{b}' \mathbf{\Sigma}^{-1} \mathcal{D}(\mathbf{h}_2) \mathbf{\Sigma}^{-1} \mathbf{b}/2 - \mathbf{\Sigma}^{-1} \cdot \mathcal{D}(\mathbf{h}_2)/2} \right)
$$
  
\n
$$
\times R_i(\boldsymbol{\beta}, \Lambda, \mathbf{b}) e^{-\mathbf{b}' \mathbf{\Sigma}^{-1} \mathbf{b}/2} d\mathbf{b}
$$
  
\n
$$
\times \left( \int_{\mathbf{b}} R_i(\boldsymbol{\beta}, \Lambda, \mathbf{b}) e^{-\mathbf{b}' \mathbf{\Sigma}^{-1} \mathbf{b}/2} d\mathbf{b} \right)^{-1}
$$
  
\n
$$
A_{n3}[h_3] = n^{-1} \sum_{i=1}^n \sum_{j=1}^{n_i} \delta_{ij} h_3(X_{ij}) - \int_0^{X_{ij}} h_3(s) d\Lambda(s)
$$
  
\n
$$
\times \int_{\mathbf{b}} e^{\boldsymbol{\beta}' \mathbf{Z}_{ij} + \mathbf{b}' \mathbf{W}_{ij}} R_i(\boldsymbol{\beta}, \Lambda, \mathbf{b}) e^{-\mathbf{b}' \mathbf{\Sigma}^{-1} \mathbf{b}/2} d\mathbf{b}
$$
  
\n
$$
\times \left( \int_{\mathbf{b}} R_i(\boldsymbol{\beta}, \Lambda, \mathbf{b}) e^{-\mathbf{b}' \mathbf{\Sigma}^{-1} \mathbf{b}/2} d\math
$$

$$
A_{n1}[\mathbf{h}_1] = n^{-1} \sum_{i=1}^n \int_{\mathbf{b}} \sum_{j=1}^{n_i} \mathbf{h}'_1 \mathbf{Z}_{ij} \left( \delta_{ij} - \Lambda(X_{ij}) e^{\beta' \mathbf{Z}_{ij} + \mathbf{b}' \mathbf{W}_{ij}} \right) d\mu_i(\mathbf{b})
$$
  
\n
$$
A_{n2}[\mathbf{h}_2] = n^{-1} \sum_{i=1}^n \int_{\mathbf{b}} \left\{ \mathbf{b}' \mathbf{\Sigma}^{-1} \mathcal{D}(\mathbf{h}_2) \mathbf{\Sigma}^{-1} \mathbf{b}/2 - \mathbf{\Sigma}^{-1} \cdot \mathcal{D}(\mathbf{h}_2)/2 \right\} d\mu_i(\mathbf{b})
$$
  
\n
$$
A_{n3}[h_3] = n^{-1} \sum_{i=1}^n \sum_{j=1}^{n_i} \delta_{ij} h_3(X_{ij}) - \int_0^{X_{ij}} h_3(s) d\Lambda(s) \int_{\mathbf{b}} e^{\beta' \mathbf{Z}_{ij} + \mathbf{b}' \mathbf{W}_{ij}} d\mu_i(\mathbf{b})
$$

where

$$
d\mu_i(\mathbf{b}) = \frac{R_i(\boldsymbol\beta, \Lambda, \mathbf{b}) e^{-\mathbf{b}'\mathbf{\Sigma}^{-1}\mathbf{b}/2} d\mathbf{b}}{\int_{\mathbf{b}} R_i(\boldsymbol\beta, \Lambda, \mathbf{b}) e^{-\mathbf{b}'\mathbf{\Sigma}^{-1}\mathbf{b}/2} d\mathbf{b}}
$$

we define the minimal  $\beta$ .  $(\beta, 2, \Lambda)$ [**ii**], **ii**<sub>2</sub>,  $n_3$ ]  $\rightarrow$  (*ii*) as

$$
S(\boldsymbol{\beta},\boldsymbol{\Sigma},\Lambda)[\mathbf{h}_1,\mathbf{h}_2,h_3]=A_1[\mathbf{h}_1]+A_2[\mathbf{h}_2]+A_3[h_3]
$$

where the linear functionals  $A_p$  are obtained by replacing the empirical sum in  $A_{np}$  by the expectation. By construction,  $\mathcal{S}_n(\mathcal{P}_n, \mathcal{Z}_n, \Lambda_n) = 0$  and  $\mathcal{S}(\mathcal{P}_0, \mathcal{Z}_0, \Lambda_0) = 0$ .

Asymptotic normality will follow as desired by verifying the four conditions of Theorem 2 in Murphy (1995). First,  $\sqrt{n}(S_n(\boldsymbol{\beta}_0,\boldsymbol{\Sigma}_0,\Lambda_0)-S(\boldsymbol{\beta}_0,\boldsymbol{\Sigma}_0,\Lambda_0))$  weakly converges to a tight Gaussian process on  $l^{\infty}(\mathcal{H})$ , because H is a Donsker class and the functionals Anp are bounded Lipschitz functionals with respect to H. The approximation condition condit that

$$
\sup_{(\mathbf{h}_1, \mathbf{h}_2, h_3) \in \mathcal{H}} |(S_n - S)(\hat{\beta}_n, \hat{\Sigma}_n, \hat{\Lambda}_n) - (S_n - S)(\beta_0, \Sigma_0, \Lambda_0)|
$$
  
= 
$$
o_p \left( n^{-1/2} \vee \left\{ \|\hat{\beta}_n - \beta_0\| + \|\hat{\Sigma}_n - \Sigma_0\| + \sup_{t \in [0, \tau]} |\hat{\Lambda}_n(t) - \Lambda_0(t)| \right\} \right)
$$

can be proved in a manner similar to Lemma 1 in the appendix of Murphy (1995). By the smoothness of  $S(\beta, \Sigma, \Lambda)$ , the Fréchet differentiability condition holds and the derivative of  $S(\boldsymbol{\beta}, \boldsymbol{\Sigma}, \Lambda)$  at  $(\boldsymbol{\beta}_0, \boldsymbol{\Sigma}_0, \Lambda_0)$  by  $\dot{S}(\boldsymbol{\beta}_0, \boldsymbol{\Sigma}_0, \Lambda_0)$ . We consider  $\dot{S}(\boldsymbol{\beta}_0, \boldsymbol{\Sigma}_0, \Lambda_0)$  to be a map,  $T$ , from the space

 $\{(\boldsymbol{\beta} - \boldsymbol{\beta}_0, \boldsymbol{\Sigma} - \boldsymbol{\Sigma}_0, \Lambda - \Lambda_0): (\boldsymbol{\beta}, \boldsymbol{\Sigma}, \Lambda) \text{ is in the neighborhood } \mathcal{U} \text{ of } (\boldsymbol{\beta}_0, \boldsymbol{\Sigma}_0, \Lambda_0)\}$ **Collection of Biostatistics** 32 **Research Archive** 

or

to  $l^{\infty}(\mathcal{H})$ . Lastly, we need to show the linear map, T, is continuously invertible on its range.

Now we can write

$$
T(\boldsymbol{\beta} - \boldsymbol{\beta}_0, \boldsymbol{\Sigma} - \boldsymbol{\Sigma}_0, \boldsymbol{\Lambda} - \boldsymbol{\Lambda}_0) = (\boldsymbol{\beta} - \boldsymbol{\beta}_0)' \mathcal{Q}_1(\mathbf{h}_1, \mathbf{h}_2, h_3) + (\boldsymbol{\Sigma} - \boldsymbol{\Sigma}_0)' \mathcal{Q}_2(\mathbf{h}_1, \mathbf{h}_2, h_3) + \int_0^{\tau} \mathcal{Q}_3(\mathbf{h}_1, \mathbf{h}_2, h_3) d(\boldsymbol{\Lambda} - \boldsymbol{\Lambda}_0)
$$

where the  $\mathcal{Q}_i$  are the respective particles of the respective of S with respect to  $p$  , , , , and . . . . The Qi are of the form

$$
Q_1(\mathbf{h}_1, \mathbf{h}_2, h_3) = B_1 \begin{pmatrix} \mathbf{h}_1 \\ \mathbf{h}_2 \end{pmatrix} + \int_0^{\tau} h_3(t) D_1(t) dt,
$$
  

$$
Q_2(\mathbf{h}_1, \mathbf{h}_2, h_3) = B_2 \begin{pmatrix} \mathbf{h}_1 \\ \mathbf{h}_2 \end{pmatrix} + \int_0^{\tau} h_3(t) D_2(t) dt,
$$

and

$$
Q_3(\mathbf{h}_1, \mathbf{h}_2, h_3) = B_3 \binom{\mathbf{h}_1}{\mathbf{h}_2} + b_4 h_3(t) + \int_0^{\tau} h_3(t) D_3(t) dt;
$$

where  $B_1$ ,  $B_2$ , and  $B_3$  are constant matrices;  $D_1(t)$ ,  $D_2(t)$ ,  $D_3(t)$  are continuously differentiable functions; and  $b_4 > 0$ ; each of which depends on  $\theta_0$ . Therefore the operator  $\mathcal{Q} = (\mathcal{Q}_1, \mathcal{Q}_2, \mathcal{Q}_3)$  can be considered the sum of a continuously invertible operator and a compact operator from H to itself.

To prove T is invertible, we need only show the invertibility of the linear operator  $\mathcal{Q}(\mathbf{h}_1, \mathbf{h}_2, h_3)$ ; or equivalently that  $\mathcal Q$  is one-to-one (Zeng et al. 2005; Rudin 1973, pp. 99-103). Suppose  $\mathcal{Q}(\mathbf{h}_1, \mathbf{h}_2, h_3) = \mathbf{0}$ , then  $T(\boldsymbol{\beta} - \boldsymbol{\beta}_0, \boldsymbol{\Sigma} - \boldsymbol{\Sigma}_0, \Lambda - \Lambda_0)[\mathbf{h}_1, \mathbf{h}_2, h_3] = \mathbf{0}$  for any (i.e. ) in the neighborhood U. In particular,  $\alpha$  is some small constant  $\alpha$  small constant  $\alpha$ 

$$
\beta = \beta_0 + \epsilon \mathbf{h}_1, \quad \Sigma = \Sigma_0 + \epsilon \mathbf{h}_2,
$$
  

$$
\Lambda(t) = \Lambda_0(t) + \epsilon \int_0^t h_3(t) d\Lambda_0(t).
$$

By definition of  $T$ , we have

 $\mathbb{P} \left( \begin{array}{ccc} 0 & -\sqrt{2} & 0 \ 0 & 0 & 0 \end{array} \right) = \mathbb{P} \left( \begin{array}{ccc} 0 & -\sqrt{2} & =\epsilon E\left\{\left(l\beta_0\left[\mathbf{n}_1\right]+l\mathbf{n}_0\left[\mathbf{n}_2\right]+l\Lambda_0\left[n_3\right]\right)\right\},\$ **Collection of Biostatistics** 33 **Research Archive** 

 $\rho_{01}$  is expression to the contract surface surface  $\rho_{-1}$ 

$$
0 = \sum_{j=1}^{n_i} \int_{\mathbf{b}} \mathbf{h}'_1 \mathbf{Z}_{ij} \left( \delta_{ij} - \Lambda_0(X_{ij}) e^{\beta'_0 \mathbf{Z}_{ij} + \mathbf{b}' \mathbf{W}_{ij}} \right) R_{2i}(\boldsymbol{\beta}_0, \Lambda_0, \mathbf{b}) d_{\mathbf{b}} N(\mathbf{0}, \boldsymbol{\Sigma}_0)
$$
  
+ 
$$
\int_{\mathbf{b}} \left\{ \mathbf{b}' \mathbf{\Sigma}_0^{-1} \mathcal{D}(\mathbf{h}_2) \mathbf{\Sigma}_0^{-1} \mathbf{b}/2 - \mathbf{\Sigma}_0^{-1} \cdot \mathcal{D}(\mathbf{h}_2)/2 \right\} R_{2i}(\boldsymbol{\beta}_0, \Lambda_0, \mathbf{b}) d_{\mathbf{b}} N(\mathbf{0}, \boldsymbol{\Sigma}_0)
$$
  
+ 
$$
\sum_{j=1}^{n_i} \int_{\mathbf{b}} \left( \delta_{ij} h_3(X_{ij}) - \int_0^{X_{ij}} h_3(s) d \Lambda_0(s) e^{\beta'_0 \mathbf{Z}_{ij} + \mathbf{b}' \mathbf{W}_{ij}} \right) R_{2i}(\boldsymbol{\beta}_0, \Lambda_0, \mathbf{b}) d_{\mathbf{b}} N(\mathbf{0}, \boldsymbol{\Sigma}_0)
$$
(27)

where

$$
R_{2i}(\boldsymbol{\beta}_0, \Lambda_0, \mathbf{b}) = R_i(\boldsymbol{\beta}_0, \Lambda_0, \mathbf{b}) \prod_{j=1}^{n_i} \{ \lambda_0(X_{ij}) \}^{\delta_{ij}}
$$
  
= 
$$
\prod_{j=1}^{n_i} \exp[\delta_{ij}(\boldsymbol{\beta}_0' \mathbf{Z}_{ij} + \mathbf{b}' \mathbf{W}_{ij}) - \Lambda_0(X_{ij}) \exp(\boldsymbol{\beta}_0' \mathbf{Z}_{ij} + \mathbf{b}' \mathbf{W}_{ij})] \{ \lambda_0(X_{ij}) \}^{\delta_{ij}}.
$$

Using techniques from Zeng *et al.* (2005) similar to the identifiability step of the consistency proof, we show that (27) implies  $\mathbf{h}_1 = \mathbf{0}$ ,  $\mathbf{h}_2 = \mathbf{0}$ , and  $h_3 = 0$ . Let  $\mathbf{Z}_{ij}$  and  $\mathbf{W}_{ij}$  be fixed. Then for fixed integer k in  $1, \ldots, n_i$ , we define measures  $\mu_1, \ldots, \mu_{n_i}$  on  $t - \frac{1}{2}$  1g  $t - \frac{1}{2}$ 

$$
\mu_m(\{0\} \times A) = 0, \quad \mu_m(\{1\} \times A) = I(0 \in A), \quad m \le k,
$$

and

$$
\mu_m(\{0\} \times A) = I(\tau \in A), \quad \mu_m(\{1\} \times A) = \int I_A dx, \quad m > k,
$$

where A is any Borel set in  $[0, \tau]$ . We integrate both sides of (27) with respect to  $\{(\delta_{i1}, X_{i1}), \ldots, (\delta_{in_i}, X_{in_i})\}$  and the product measure  $d\mu_1, \ldots, d\mu_{n_i}$ . That is, we let im and  $\alpha$  implies the state of  $\alpha$  for all m  $\alpha$  in  $\alpha$  in  $\alpha$  is the  $\alpha$  in  $\alpha$  is the  $\alpha$  implies  $\alpha$  is the  $\alpha$ integrate  $X_{im}$  from 0 to  $\tau$  if  $\delta_{im} = 1$ , then sum over  $\delta_{ij} \in \{0, 1\}$ . Then we sum all of the equalities of (27) for all possible combinations of  $\{o_{i1},\ldots,o_{in_i}\}\in\{0,1\}$ .

We compute the integral of each term on the right side of (27) with respect to the

 $\lambda$   $\sim$   $\lambda$ 

**Collection of Biostatistics Research Archive** 

product measure,  $\prod_{m=1}^{n_i} \mu_m$ , the sum of which must be 0. First note, for any **b**,

$$
\int R_{2i}(\beta_0, \Lambda_0, \mathbf{b}) d\left(\prod_{m=1}^{n_i} \mu_m\right)
$$
\n
$$
= \prod_{m \leq k} {\{\lambda_0(0) e^{\beta'_0 \mathbf{Z}_{im} + \mathbf{b}' \mathbf{W}_{im}}\}}
$$
\n
$$
\times \sum_{\delta_{im} \in \{0,1\} \ m > k} {\prod_{m > k} (\exp[-\Lambda_0(\tau) \exp(\beta'_0 \mathbf{Z}_{im} + \mathbf{b}' \mathbf{W}_{im})])^{1-\delta_{im}}}
$$
\n
$$
\times \left\{ \int_{y=0}^{\tau} \exp[\beta'_0 \mathbf{Z}_{im} + \mathbf{b}' \mathbf{W}_{im} - \Lambda_0(y) \exp(\beta'_0 \mathbf{Z}_{im} + \mathbf{b}' \mathbf{W}_{im})] \lambda_0(y) dy \right\}^{\delta_{im}}
$$
\n
$$
= \prod_{m \leq k} {\{\lambda_0(0) e^{\beta'_0 \mathbf{Z}_{im} + \mathbf{b}' \mathbf{W}_{im}}\}}
$$
\n
$$
\times \sum_{\delta_{im} \in \{0,1\} \ m > k} {\prod_{m > k} (\exp[-\Lambda_0(\tau) \exp(\beta'_0 \mathbf{Z}_{im} + \mathbf{b}' \mathbf{W}_{im})])^{1-\delta_{im}}}
$$
\n
$$
\times (1 - \exp[-\Lambda_0(\tau) \exp(\beta'_0 \mathbf{Z}_{im} + \mathbf{b}' \mathbf{W}_{im})])^{\delta_{im}}
$$
\n
$$
= \prod_{m \leq k} {\{\lambda_0(0) e^{\beta'_0 \mathbf{Z}_{im} + \mathbf{b}' \mathbf{W}_{im}}\}}
$$

For the rst term of (27), if j k, then for any b:

 0(Xij )e00Zij+b0Wij R2i(0; 0; b) <sup>d</sup> Yni m! Z h01Zij ij m=1 h01ZijR2i(0; 0; b) d Yni m! Z = m=1 =h01Zij <sup>Y</sup> f0(0)e00Zim+b0Wimg 

If  $j>k$ , then

$$
\int \mathbf{h}'_{1} \mathbf{Z}_{ij} \left( \delta_{ij} - \Lambda_{0} (X_{ij}) e^{\beta_{0}' \mathbf{Z}_{ij} + \mathbf{b}' \mathbf{W}_{ij}} \right) R_{2i}(\beta_{0}, \Lambda_{0}, \mathbf{b}) d \left( \prod_{m=1}^{n_{i}} \mu_{m} \right)
$$
\n
$$
= \mathbf{h}'_{1} \mathbf{Z}_{ij} \prod_{m \leq k} \left\{ \lambda_{0}(0) e^{\beta_{0}' \mathbf{Z}_{im} + \mathbf{b}' \mathbf{W}_{im}} \right\}
$$
\n
$$
\times \sum_{\substack{\delta_{im} \in \{0,1\} \\ m > k, m \neq j}} \prod_{m \leq k} \left( \exp[-\Lambda_{0}(\tau) \exp(\beta_{0}' \mathbf{Z}_{im} + \mathbf{b}' \mathbf{W}_{im}) ] \right)^{1 - \delta_{im}}
$$
\n
$$
\times \left( 1 - \exp[-\Lambda_{0}(\tau) \exp(\beta_{0}' \mathbf{Z}_{im} + \mathbf{b}' \mathbf{W}_{im}) ] \right)^{\delta_{im}}
$$
\n
$$
\times \sum_{\delta_{ij} \in \{0,1\}} (1 - \delta_{ij}) \left( -\Lambda_{0}(\tau) e^{\beta_{0}' \mathbf{Z}_{ij} + \mathbf{b}' \mathbf{W}_{ij}} \right) \exp[-\Lambda_{0}(\tau) \exp(\beta_{0}' \mathbf{Z}_{ij} + \mathbf{b}' \mathbf{W}_{ij}) ]
$$
\n
$$
+ \delta_{ij} \int_{y=0}^{\tau} \left( 1 - \Lambda_{0}(y) e^{\beta_{0}' \mathbf{Z}_{ij} + \mathbf{b}' \mathbf{W}_{ij}} \right) \exp[\beta_{0}' \mathbf{Z}_{ij} + \mathbf{b}' \mathbf{W}_{ij} - \Lambda_{0}(y) \exp(\beta_{0}' \mathbf{Z}_{ij} + \mathbf{b}' \mathbf{W}_{ij}) ] \lambda_{0}(y) dy
$$
\n
$$
= \mathbf{h}'_{1} \mathbf{Z}_{ij} \prod_{m \leq k} \left\{ \lambda_{0}(0) e^{\beta_{0}' \mathbf{Z}_{im} + \mathbf{b}' \mathbf{W}_{im}} \right\}
$$
\n
$$
\times \sum_{\delta_{ij} \in \{0,1\}} (1 - \delta_{ij}) \left
$$

Therefore

$$
\int \sum_{j=1}^{n_i} \int_{\mathbf{b}} \mathbf{h}'_1 \mathbf{Z}_{ij} \left( \delta_{ij} - \Lambda_0(X_{ij}) e^{\beta'_0 \mathbf{Z}_{ij} + \mathbf{b}' \mathbf{W}_{ij}} \right) R_{2i}(\boldsymbol{\beta}_0, \Lambda_0, \mathbf{b}) d_{\mathbf{b}} N(\mathbf{0}, \boldsymbol{\Sigma}_0) d \left( \prod_{m=1}^{n_i} \mu_m \right)
$$
  
= 
$$
\sum_{j \le k} \mathbf{h}'_1 \mathbf{Z}_{ij} \int_{\mathbf{b}} \prod_{m \le k} \{ \lambda_0(0) e^{\beta'_0 \mathbf{Z}_{im} + \mathbf{b}' \mathbf{W}_{im}} \} d_{\mathbf{b}} N(\mathbf{0}, \boldsymbol{\Sigma}_0).
$$
 (28)

Likewise, from the second term of (27):

$$
\int \int_{\mathbf{b}} \left\{ \mathbf{b}^{\prime} \mathbf{\Sigma}_{0}^{-1} \mathcal{D}(\mathbf{h}_{2}) \mathbf{\Sigma}_{0}^{-1} \mathbf{b}/2 - \mathbf{\Sigma}_{0}^{-1} \cdot \mathcal{D}(\mathbf{h}_{2})/2 \right\} R_{2i}(\beta_{0}, \Lambda_{0}, \mathbf{b}) d_{\mathbf{b}} N(\mathbf{0}, \Sigma_{0}) d \left( \prod_{m=1}^{n_{i}} \mu_{m} \right)
$$
  
\n
$$
= \int_{\mathbf{b}} \left\{ \mathbf{b}^{\prime} \mathbf{\Sigma}_{0}^{-1} \mathcal{D}(\mathbf{h}_{2}) \mathbf{\Sigma}_{0}^{-1} \mathbf{b}/2 - \mathbf{\Sigma}_{0}^{-1} \cdot \mathcal{D}(\mathbf{h}_{2})/2 \right\} \prod_{m \leq k} \left\{ \lambda_{0}(0) e^{\beta_{0}' \mathbf{Z}_{im} + \mathbf{b}^{\prime} \mathbf{W}_{im}} \right\} d_{\mathbf{b}} N(\mathbf{0}, \Sigma_{0}).
$$
  
\nA BERRES REPOSITION  
\nCollection of Blostatistics  
\nResearch Archive  
\n36  
\n(29)

Furthermore, from the third term of (27), if j <sup>k</sup> then

$$
\int \left( \delta_{ij} h_3(X_{ij}) - \int_0^{X_{ij}} h_3(s) d\Lambda_0(s) e^{\beta'_0 \mathbf{Z}_{ij} + \mathbf{b}' \mathbf{W}_{ij}} \right) R_{2i}(\boldsymbol{\beta}_0, \Lambda_0, \mathbf{b}) d \left( \prod_{m=1}^{n_i} \mu_m \right)
$$
  
=  $h_3(0) \prod_{m \le k} {\lambda_0(0) e^{\beta'_0 \mathbf{Z}_{im} + \mathbf{b}' \mathbf{W}_{im}}};$  (30)

if  $j > k$ , then

$$
\int \left( \delta_{ij} h_3(X_{ij}) - \int_0^{X_{ij}} h_3(s) d\Lambda_0(s) e^{\beta_0' \mathbf{Z}_{ij} + \mathbf{b}' \mathbf{W}_{ij}} \right) R_{2i}(\beta_0, \Lambda_0, \mathbf{b}) d \left( \prod_{m=1}^{n_i} \mu_m \right)
$$
\n
$$
= \prod_{m \leq k} \left\{ \lambda_0(0) e^{\beta_0' \mathbf{Z}_{im} + \mathbf{b}' \mathbf{W}_{im}} \right\}
$$
\n
$$
\times \sum_{\delta_{ij} \in \{0,1\}} \left\{ -(1 - \delta_{ij}) \int_0^{\tau} h_3(s) d\Lambda_0(s)
$$
\n
$$
\times \exp[\beta_0' \mathbf{Z}_{ij} + \mathbf{b}' \mathbf{W}_{ij} - \Lambda_0(t) \exp(\beta_0' \mathbf{Z}_{ij} + \mathbf{b}' \mathbf{W}_{ij})]
$$
\n
$$
+ \delta_{ij} \int_{y=0}^{\tau} \left( h_3(y) - \int_{s=0}^{y} h_3(s) d\Lambda_0(s) e^{\beta_0' \mathbf{Z}_{ij} + \mathbf{b}' \mathbf{W}_{ij}} \right)
$$
\n
$$
\times \exp[\beta_0' \mathbf{Z}_{ij} + \mathbf{b}' \mathbf{W}_{ij} - \Lambda_0(t) \exp(\beta_0' \mathbf{Z}_{ij} + \mathbf{b}' \mathbf{W}_{ij})] \lambda_0(y) dy \right\}
$$
\n
$$
= \prod_{m \leq k} \left\{ \lambda_0(0) e^{\beta_0' \mathbf{Z}_{im} + \mathbf{b}' \mathbf{W}_{im}} \right\}
$$
\n
$$
\times \sum_{\delta_{ij} \in \{0,1\}} \left\{ -(1 - \delta_{ij}) \int_0^{\tau} h_3(s) d\Lambda_0(s)
$$
\n
$$
\times \exp[\beta_0' \mathbf{Z}_{ij} + \mathbf{b}' \mathbf{W}_{ij} - \Lambda_0(t) \exp(\beta_0' \mathbf{Z}_{ij} + \mathbf{b}' \mathbf{W}_{ij})]
$$
\n
$$
+ \delta_{ij} \int_{s=0}^{\tau} h_3(s) d\Lambda_0(s) \exp[\beta_0' \mathbf{Z}_{ij} + \mathbf{b}' \mathbf{W
$$

Thus, the contract of the contract of  $\mathbb{R}^n$ 

$$
\int \sum_{j=1}^{n_i} \int_{\mathbf{b}} \left( \delta_{ij} h_3(X_{ij}) - \int_0^{X_{ij}} h_3(s) d\Lambda_0(s) e^{\beta'_0 \mathbf{Z}_{ij} + \mathbf{b}' \mathbf{W}_{ij}} \right) R_{2i}(\beta_0, \Lambda_0, \mathbf{b}) d_{\mathbf{b}} N(\mathbf{0}, \Sigma_0) d \left( \prod_{m=1}^{n_i} \mu_m \right)
$$
  
= 
$$
\sum_{j \le k} h_3(0) \int_{\mathbf{b}} \prod_{m \le k} {\{\lambda_0(0) e^{\beta'_0 \mathbf{Z}_{im} + \mathbf{b}' \mathbf{W}_{im}} d_{\mathbf{b}} N(\mathbf{0}, \Sigma_0)}
$$
(31)

Combining (28), (29), and (31) and integrating over b, we obtain

$$
\sum_{\mathbf{A} \in \mathbb{R} \setminus \mathbb{R}} \mathbf{h}'_1 \mathbf{Z}_{ij} + \frac{1}{2} \left( \sum_{j=1}^k \mathbf{W}_{ij} \right)' \mathcal{D}(\mathbf{h}_2) \left( \sum_{j=1}^k \mathbf{W}_{ij} \right) + kh_3(0) = 0.
$$

Since the index set  $j = 1, \ldots, k$  is arbitrary, we conclude

$$
\sum_{j=k_1+1}^{k_2} \mathbf{h}'_1 \mathbf{Z}_{ij} + \frac{1}{2} \left( \sum_{j=k_1+1}^{k_2} \mathbf{W}_{ij} \right)' \mathcal{D}(\mathbf{h}_2) \left( \sum_{j=k_1+1}^{k_2} \mathbf{W}_{ij} \right) + (k_2 - k_1) h_3(0) = 0.
$$

for any  $1 \leq \kappa_1 \leq \kappa_2 \leq n_i$ . Therefore  ${\bf w}_{ij} \nu({\bf n}_2)$   ${\bf w}_{ij'} = 0$  for  $j \neq j$  and  ${\bf z}_{ij} {\bf n}_1 +$  $\mathbf{W}'_{ij}\mathcal{D}(\mathbf{h}_2)\mathbf{W}_{ij}/2 + h_3(0) = 0$ . Condition C3 yields  $\mathcal{D}(\mathbf{h}_2) = \mathbf{0}$ , and it follows that  $\mathbf{h}_1 = \mathbf{0}, \, \mathbf{h}_2 = \mathbf{0}, \text{ and } h_3(0) = 0.$ 

 $\mathbb{R}^n$   $\mathbb{R}^n$   $\mathbb{R}^n$   $\mathbb{R}^n$   $\mathbb{R}^n$   $\mathbb{R}^n$   $\mathbb{R}^n$   $\mathbb{R}^n$   $\mathbb{R}^n$  to  $\mathbb{R}^n$  to  $\mathbb{R}^n$   $\mathbb{R}^n$  to  $\mathbb{R}^n$  to  $\mathbb{R}^n$  to  $\mathbb{R}^n$ 

$$
h_3(X_{i1})=\frac{\int_0^{X_{i1}} h_3(s) d \Lambda_0(s) \int_{\mathbf{b}} e^{\boldsymbol{\beta}_0^t Z_{i1}+\mathbf{b}' \mathbf{W}_{i1}} R_{2i}(\boldsymbol{\beta}_0,\Lambda_0,\mathbf{b}) d_{\mathbf{b}} N(\mathbf{0},\boldsymbol{\Sigma}_0)}{\int_{\mathbf{b}} R_{2i}(\boldsymbol{\beta}_0,\Lambda_0,\mathbf{b}) d_{\mathbf{b}} N(\mathbf{0},\boldsymbol{\Sigma}_0)}.
$$

 $\mathbb{R}$  so the expression group  $\mathbb{R}$  of  $\mathbb{R}$  , we have the expression group  $\mathbb{R}$  $\int_0^y h_3(t) d\Lambda_0(t)$  satisfies the homogeneous equation

$$
\frac{g'(y)}{\lambda_0(y)}-g(y)\frac{\int_{\mathbf{b}}e^{\boldsymbol{\beta}_0^tZ_{i1}+\mathbf{b}'\mathbf{W}_{i1}}R_{2i}(\boldsymbol{\beta}_0,\Lambda_0,\mathbf{b})d_{\mathbf{b}}N(\mathbf{0},\boldsymbol{\Sigma}_0)}{\int_{\mathbf{b}}R_{2i}(\boldsymbol{\beta}_0,\Lambda_0,\mathbf{b})d_{\mathbf{b}}N(\mathbf{0},\boldsymbol{\Sigma}_0)}=0
$$

with a condition given given given given good  $\mathcal{Y}(y)$  for  $\mathcal{Y}(y)$  is one-to-one-to-one-to-one-to-one-to-one,  $\mathcal{Y}(y)$  is one-to-one-to-one-to-one,  $\mathcal{Y}(y)$  is one-to-one-to-one-to-one-to-one-to-one-to-one-to-one and  $S(\boldsymbol{\beta}_0, \boldsymbol{\Sigma}_0, \Lambda_0)$  is invertible.

Asymptotic normality follows from Theorem 2 of Murphy (1995) and the proof of asymptotic emerging for  $\mu_n$  and  $\omega_n$  is identical to Zeng et al. (2005).

**PROOF OF THEOREM 3.** The proof is analogous to the proof of Theorem 3 in Zeng et al. (2005).

PROOF OF THEOREM 4. The proof is similar to the proof of Theorem 1 in Self and Liang (1987), except that the Taylor series expansion cited in Lehmann (1983, pp.429-432) is now replaced by  $(6)$ .

**PROOF OF THEOREM 5**. From Theorem 1 we have that  $\sqrt{n}(\phi - \phi_0) = O_p(1)$ . Applying (0) for the sequence  $\varphi_n = \varphi$ , we get

$$
\text{pl}(\mathbf{y}^*|\hat{\boldsymbol{\phi}}(\mathbf{y})) = \text{pl}(\mathbf{y}^*|\boldsymbol{\phi}_0) + s(\mathbf{y}^*|\boldsymbol{\phi}_0)'(\hat{\boldsymbol{\phi}} - \boldsymbol{\phi}_0) - n(\hat{\boldsymbol{\phi}} - \boldsymbol{\phi}_0)' \mathcal{I}_0(\hat{\boldsymbol{\phi}} - \boldsymbol{\phi}_0)/2 + r_1, \quad (32)
$$

where  $r_1 = o_p(1)$ . The main result (5) from Murphy and van der Vaart (2000) implies that  $Es(\mathbf{y}^*|\boldsymbol{\phi}_0) = 0$  (divide by  $\sqrt{n}$  and take limits on both sides of (5), and then apply **Collection of Biostatistics** Research Archive

the strong law of large numbers). Therefore, taking expectations on both sides of the equality in  $(32)$ , the first-order term vanishes and we get

$$
E_{f(\mathbf{y}^*)} \text{pl}(\mathbf{y}^* | \hat{\phi}(\mathbf{y})) = E \text{pl}(\phi_0) - n(\hat{\phi} - \phi_0)' \mathcal{I}_0(\hat{\phi} - \phi_0) / 2 + E r_1.
$$
 (33)

Taking expectation one more time, with respect to  $\bf{y}$  on both sides of (33), we have

$$
\begin{array}{lll} \mathrm{pAI} & = & -2E\mathrm{pl}(\mathbf{y}|\boldsymbol{\phi}_0) + E\{n(\hat{\boldsymbol{\phi}} - \boldsymbol{\phi}_0)' \mathcal{I}_0(\hat{\boldsymbol{\phi}} - \boldsymbol{\phi}_0)\} + Er_1 \\ \\ & = & -2E\mathrm{pl}(\mathbf{y}|\hat{\boldsymbol{\phi}}(\mathbf{y})) + 2E\{\mathrm{pl}(\mathbf{y}|\hat{\boldsymbol{\phi}}(\mathbf{y})) - \mathrm{pl}(\mathbf{y}|\boldsymbol{\phi}_0)\} + E\{n(\hat{\boldsymbol{\phi}} - \boldsymbol{\phi}_0)' \mathcal{I}_0(\hat{\boldsymbol{\phi}} - \boldsymbol{\phi}_0)\} + Er_1. \end{array}
$$

>From Corollary 2 and 1 of Murphy and van der Vaart (2000), the middle term and the last term under expectation signs in the last equation above have a  $\chi_p$  distribution, except for remainder terms of  $o_p(1)$ . Collecting all the remainder terms in  $r_2 = o_p(1)$ , we get that

$$
\text{pAI} = -2E\text{pl}(\mathbf{y}|\hat{\boldsymbol{\phi}}(\mathbf{y})) + 2p + Er
$$

which proves the theorem. If r is uniformly integrable, then  $E(r) = o(1)$ , and pAIC is asymptotically unbiased for pAI.

**PROOF OF PROPOSITION 1.** The consistency part is immediate by applying the strong law of large numbers to  $A$  and  $A_i$ .

To show the variance inequality, note that  $A = \sum_k \exp\{\sum_i v(\mathbf{b}_i^{(k)})\}/M$  . Assume for simplicity that  $n=2$  (the general case follows by induction). Put  $\exp\{v(\mathbf{b}_i)^{(n)}\}=\xi_i^{(n)}$ , for  $i = 1, 2$ . Then  $A = \zeta_1 \zeta_2$ , and  $A = \zeta_1 \zeta_2$ , where the bar denotes sample average of M observations. Let  $\mu_i, \sigma_i$  denote respectively the mean and variance of  $\zeta_i$ ,  $i=1,2$ . Then

$$
\begin{array}{rcl}\n\text{Var}(\overline{\xi_1 \xi_2}) & = & \text{Var}(\xi_1 \xi_2) / M \\
& = & \sigma_1^2 \sigma_2^2 / M + \mu_1^2 \sigma_2^2 / M + \mu_2^2 \sigma_1^2 / M \\
\text{Var}(\overline{\xi_1 \xi_2}) & = & (\sigma_1^2 / M)(\sigma_2^2 / M) + \mu_1^2 \sigma_2^2 / M + \mu_2^2 \sigma_1^2 / M\n\end{array}
$$

The mst term in  $var(\xi|\xi_2)$  is no smaller than the corresponding term in  $var(\xi|\xi_2)$ , while the other two terms are identical, so the result follows.

## References

- Akaike, H. (1973). Information theory and an extension of the maximum likelihood principle. In *Breakthroughs in statistics (1992)*, vol. 1, 610–624. Springer-Verlag.
- Andersen, P. and Gill, R. (1982). Cox's regression model for counting processes: a large sample study. Annals of Statistics  $10$ , 1100-20.
- Burnham, K. P. and Anderson, D. R. (2002). *Model Selection and Multimodel* Inference: A Practical Information - Theoretic Approach. Springer, 2nd edn.
- Chernoff, H. (1954). On the distribution of the likelihood ratio. Annals of Mathematical Statistics  $25, 573-578$ .
- Claeskens, G. and Hjort, N. L. (2003). Focused information criterion (with discussion). Journal of the American Statistical Association  $98, 900-945$ .
- Commenges, D. and Andersen, P. (1995). Score test of homogeneity for survival data. Lifetime Data Analysis 1, 145-156.
- Cox, D. (1975). Partial likelihood. *Biometrika*  $62$ , 269–276.
- Crainiceanu, C. and Ruppert, D. (2004). Likelihood ratio tests in linear mixed models with one variance component. Journal of the Royal Statistical Society, Series B  $66, 165-185.$
- deLeeuw, J. (1992). Introduction to Akaike (1973) 'Information theory and an extension of the maximum likelihood principle'. In Breakthroughs in Statistics, vol. 1, 599–609. New York: Springer.
- DiCiccio, T. J., Kass, R. E., Raftery, A., and Wasserman, L. (1997). Computing Bayes factors by combining simulation and asymptotic approximations. Journal of the American Statistical Association  $92, 903{-}915.$
- Fan, J. H. and Wong, W. H. (2000). Discussion of 'On profile likelihood', by Murphy, S. A. and van der Vaart, A. W. Journal of the American Statistical Association 95,  $468 - 471$ .

- Gelfand, A. and Day, D. (1994). Bayesian model choice: asymptotics and exact calculations. Journal of the Royal Statistical Society, Series  $B$  56, 501-514.
- Glidden, D. and Vittinghoff, E. (2004). Modelling clustered survival data from multicenter clinical trials. Statistics in Medicine 23, 369-388.
- Gray, R. (1995). Tests for variation over groups in survival data. *Journal of the* American Statistical Association  $90, 198-203$ .
- Lehmann, E. L. (1983). Theory of Point Estimation. John Wiley, New York.
- Li, B. (2000). Comment on 'on profile likelihood'. Journal of the American Statistical Association 95, 472-474.
- Linhart, H. and Zucchini, W. (1986). *Model Selection*. Wiley, New York.
- Liu, I., Blacker, D., Xu, R., Fitzmaurice, G., Lyons, M., and Tsuang, M. (2004a). Genetic and environmental contributions to the development of alcohol dependence in male twins. Archives of General Psychiatry  $61, 897-903$ .
- Liu, I., Blacker, D., Xu, R., Fitzmaurice, G., Tsuang, M., and Lyons, M. J. (2004b). Genetic and environmental contributions to age of onset of alcohol dependence symptoms in male twins.  $Addiction$  99, 1403-1409.
- Longford, N. T. (2005). Model selection and efficiency  $-$  is 'Which model...?' the right question? Journal of the Royal Statistical Association, Series A 168, 469{472.
- Maple, J., Murphy, S., and Axinn, W. (2002). Two-level proportional hazards models. *Biometrics* 58, 754-763.
- Meng, X.-L. and Wong, W. (1996). Simulating ratios of normalizing constants via a simple identity. *Statistica Sinica* 6, 831-860.
- Murphy, S. (1994). Consistency in a proportional hazards model incorporating a random effect. Annals of Statistics  $22$ ,  $2$ ,  $712-731$ .
- Murphy, S. (1995). Asymptotic theory for the frailty model. Annals of Statistics  $23, 1, 182 - 198.$

- Murphy, S. and van der Vaart, A. (2000). On profile likelihood. Journal of the American Statistical Association 95, 449-485.
- Murray, D., Varnell, S., and Blitstein, J. (2004). Design and analysis of grouprandomized trials: a review of recent methodological developments. American Journal of Public Health  $94, 423-432$ .
- Parner, E. (1998). Asymptotic theory for the correlated Gamma-frailty model. Annals of Statistics  $26$ , 1, 183-214.
- Ripatti, S. and Palmgren, J. (2000). Estimation of multivariate frailty models using penalized partial likelihood. *Biometrics* 56, 1016-1022.
- Rudin, W. (1973). Functional Analysis. New York: McGraw-Hill.
- Self, S. and Liang, K.-Y. (1987). Asymptotic properties of maximum likelihood estimators and likelihood ratio tests under nonstandard conditions. Journal of the American Statistical Association 82, 398, 605-610.
- Severini, T. A. and Wong, W. H. (1992). Profile likelihood and conditionally parametric models. Annals of Statistics 20, 1768-1802.
- Stram, D. and Lee, J. (1994). Variance components testing in the longitudinal mixed effects model. Biometrics 50, 1171-1177.
- Stram, D. and Lee, J. (1995). Correction to \Variance component testing in the longitudinal mixed effects model". *Biometrics* 51, 1196.
- Sylvester, R., van Glabbeke, M., Collette, L., Suciu, S., Baron, B., Legrand, C., Gorlia, T., Collins, G., Coens, C., Declerck, L., and Therasse, P. (2002). Statistical methodology of phase iii cancer clinical trials: advances and future perspectives. European Journal of Cancer 38, S162-S168.
- Therneau, T. and Grambsch, P. (2000). *Modelling Survival Data: Extending the* Cox Model. Springer Verlag, New York, USA.
- Tierney, L. and Kadane, J. B. (1986). Accurate approximations for posterior moments and marginal densities. Journal of the American Statistical Association  $81,\,82\hbox{--}86.$ REPOSITORY

- Vaida, F. and Blanchard, S. (2005). Conditional akaike information for mixedeffects models. Biometrika  $92, 351-370$ .
- Vaida, F. and Xu, R. (2000). Proportional hazards model with random effects. Statistics in Medicine  $19, 3309 - 3324$ .
- Verweij, P. and van Houwelingen, H. (1995). Time-dependent effects of fixed covariates in Cox regression. Biometrics  $51$ , 1550-56.
- Zeng, D., Lin, D. Y., and Yin, G. (2005). Maximum likelihood estimation for proportional odds model with random effects. Journal of the American Statistical Association 100, 470-483.



	Model Laplace	RIS	BS.	$pAIC^+$
$0^*$	7241.76	7241.76	7241.76	7249.76
$1^*$	7232.80 (8.96) 7232.80		7232.80	7242.80
$2^{\circ}$		7228.98 (3.82) 7228.80 (4.00) 7228.78 (4.02)		7240.80
		7222.72 (6.26) 7222.55 (6.25) 7222.60 (6.18) 7236.55		

Table 1: 2- Log likelihood values from the lung cancer data

RIS - reciprocal importance sampling, BS - bridge sampling.

<sup>+</sup> computed using RIS.

likelihood computed directly when there are no random eects.

In () are the likelihood ratio statistics between the model and its immediate submodel (3 vs. 2, <sup>2</sup> vs. 1, etc.).





Figure 1: Q-Q plots of likelihood ratio statistics from simulated data

