

Semiparametric Transformation Models for Survival Data with a Cure Fraction

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We propose a class of transformation models for survival data with a cure fraction. The class of transformation models is motivated by biological considerations, and it includes both the proportional hazards and the proportional odds cure models as two special cases. An efficient recursive algorithm is proposed to calculate the maximum likelihood estimators. Furthermore, the maximum likelihood estimators for the regression coefficients are shown to be consistent and asymptotically normal, and their asymptotic variances attain the semiparametric efficiency bound. Simulation studies are conducted to examine the finite sample properties of the proposed estimators. The method is illustrated on data from a clinical trial involving the treatment of melanoma.

KEYWORDS: Cure model; Linear transformation models; Proportional hazards model; Proportional odds model; Semiparametric efficiency.

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1 Introduction

In time-to-event data arising from cancer and AIDS clinical trials, it is often observed that a proportion of subjects will never fail. For analyzing such data, cure rate models have been proposed and studied extensively. One type of commonly used cure rate model is the so-called *two-component mixture cure model* (Berkson and Gage, 1952), which treats the whole population as a mixture of cured subjects and non-cured subjects. This mixture model has been studied by many authors, including Gray and Tsiatis (1989), Sposto, Sather, and Baker (1992), Laska and Meisner (1992), Kuk and Chen (1992), Taylor (1995), Sy and Taylor (2000), Lu and Ying (2004) among others. The book by Maller and Zhou (1996) gives an detailed discussion of frequentist methods of inference for the two-component mixture cure model.

Although the mixture cure model is intuitively attractive, it does have several drawbacks, both from a Bayesian and frequentist perspective, as pointed out by Chen, Ibrahim and Sinha (1999) and Ibrahim, Chen and Sinha (2001). An alternative cure rate model with desirable properties, called the *promotion time cure model*, has been proposed and studied by Yakovlev and Tsodikov (1996), Tsodikov (1998), and Chen, Ibrahim and Sinha (1999). In this model, the cured subjects are assumed to have survival time equal to infinity and the survival distribution for either cured subjects or non-cured subjects can be integrated into one single formulation: for the i th individual with covariate \mathbf{X}_i in the population, the survival function of subject i is given by

$$S(t|\mathbf{X}_i) = \exp\{-\theta(\mathbf{X}_i)F(t)\}, \quad (1)$$

where $\theta(\cdot)$ is a known link function and $F(t)$ is a distribution function. Under the promotion time cure model (1), the cure rate is $S(\infty|\mathbf{X}_i) = \exp\{-\theta(\mathbf{X}_i)\}$ and the hazard rate at time t for subject i is equal to $\theta(\mathbf{X}_i)f(t)$, where $f(t) = dF(t)/dt$. Thus, we see that model (1) has the proportional hazards structure when the covariates are modeled through $\theta(\cdot)$. Moreover, when $\theta(\mathbf{X}_i) = \exp(\boldsymbol{\beta}^T \mathbf{X}_i)$ and $\boldsymbol{\beta}$ contains an intercept term β_0 , model (1) becomes the usual Cox (1972) proportional hazards model subject to the restriction of a *bounded* cumulative baseline hazard function, given by $\Lambda(t) = F(t) \exp(\beta_0)$. Thus, any cure rate model has a bounded cumulative hazard, leading to an improper survival function (i.e., $S(\infty) > 0$), whereas non-cure models, such as the Cox model (Cox, 1972), have an unbounded cumulative hazard, thus leading to a proper survival function (i.e., $S(\infty) = 0$).

Yakovlev and Tsodikov (1996) and Chen, Ibrahim and Sinha (1999) provide a biological derivation for model (1). The motivation comes from studying the time to relapse of cancer for patients with or without tumor cells. Specially, the promotion time cure model is derived as follows. For the i th

subject, let N_i denote the number of tumor cells that have the potential of metastasizing, i.e., the number of metastasis-competent tumor cells. The N_i 's are unobservable latent variables. We assume that N_i has a Poisson distribution with Poisson rate (mean) $\theta(\mathbf{X}_i)$. We denote the promotion time for the k th tumor cell by \tilde{T}_k ($k = 1, \dots, N_i$) which is the time for the k th metastasis-competent tumor cell to produce a detectable tumor mass. The \tilde{T}_k 's are also unobservable quantities. Conditional on N_i , the \tilde{T}_k 's are independent and identically distributed (i.i.d.) as F , where F is sometimes referred to as the promotion time cumulative distribution function. Then the time to relapse of cancer, defined as $T = \min(\tilde{T}_1, \dots, \tilde{T}_{N_i})$, which is the observed event time, has the survival function

$$\begin{aligned} S(t|\mathbf{X}_i) &= P(N_i = 0) + \sum_{k \geq 1} P(\tilde{T}_1 > t, \dots, \tilde{T}_k > t | N_i = k) P(N_i = k) \\ &= \exp\{-\theta(\mathbf{X}_i)\} + \sum_{k=1}^{\infty} \{1 - F(t)\}^k \frac{\theta(\mathbf{X}_i)^k \exp\{-\theta(\mathbf{X}_i)\}}{k!} \\ &= \exp\{-\theta(\mathbf{X}_i)F(t)\}. \end{aligned}$$

In the derivation of (1), one critical assumption is that conditional on the number of tumor cells $N_i = k$, $(\tilde{T}_1, \dots, \tilde{T}_k)$ are mutually independent. This assumption may be unrealistic since $(\tilde{T}_1, \dots, \tilde{T}_k)$ are unobserved random variables taken on the same subject. One possible relaxation and remedy of this assumption is to introduce a subject-specific frailty ξ_i such that conditional on both $N_i = k$ and ξ_i , $(\tilde{T}_1, \dots, \tilde{T}_k)$ are mutually independent with distribution function $F(t)$. Moreover, we assume that conditional on \mathbf{X}_i and ξ_i , N_i has a Poisson distribution with rate $\xi_i \theta(\mathbf{X}_i)$; thus, ξ_i represents the heterogeneity of the Poisson rates in the N_i 's. Following the same derivation as before, we then obtain that the survival function for the time to relapse, T , is

$$S(t|\mathbf{X}_i) = E_{\xi_i} \left[e^{-\theta(\mathbf{X}_i)F(t)\xi_i} \right],$$

where E_{ξ_i} denotes the expectation with respect to ξ_i . For example, when ξ_i has a gamma distribution with mean one, that is, ξ_i has the density $\{\gamma^{1/\gamma} \Gamma(1/\gamma)\}^{-1} \xi_i^{1/\gamma-1} \exp(-\xi_i/\gamma)$, then after simple algebra, we obtain

$$S(t|\mathbf{X}_i) = \{1 + \gamma \theta(\mathbf{X}_i) F(t)\}^{-1/\gamma}.$$

Equivalently, we can write

$$S(t|\mathbf{X}_i) = G_\gamma\{\theta(\mathbf{X}_i)F(t)\}, \tag{2}$$

where $G_\gamma(\cdot)$ is the transformation

$$G_\gamma(x) = \begin{cases} (1 + \gamma x)^{-1/\gamma} & \gamma > 0 \\ e^{-x} & \gamma = 0. \end{cases} \tag{3}$$

Through (2) and (3), we obtain a very general class of transformation cure models, and note that the proportional hazards cure rate model in (1) is a special case of this class which corresponds to $\gamma = 0$. There are also other interesting special cases arising from (2) and (3). When $\gamma = 1$, we obtain a proportional odds type of cure model, similar in flavor to the proportional odds models with proper survival functions considered by Pettitt (1982) and Bennett (1983). Moreover, the general form of the class in (2) not only has a strong biological motivation, but also it can reduce to the usual linear transformation models studied by Cheng, Wei and Ying (1995) under a special choice of $\theta(\cdot)$. For instance, if we choose $\theta(\mathbf{X}_i) = \exp(\beta_0 + \beta_1^T \mathbf{Z}_i)$ with $\mathbf{X}_i = (1, \mathbf{Z}_i^T)^T$, $\beta = (\beta_0, \beta_1^T)^T$ and β_0 being the intercept term in the regression, then model (2) is equivalent to $S(t|\mathbf{Z}_i) = G_\gamma\{\exp(\beta_1^T \mathbf{Z}_i)\Lambda(t)\}$ where $\Lambda(t) = F(t)\exp(\beta_0)$ is the cumulative baseline hazard. However, when $\theta(\mathbf{X}_i)$ has a form other than $\theta(\mathbf{X}_i) = \exp(\beta^T \mathbf{X}_i)$, for example, if $\theta(\mathbf{X}_i) = \exp(\beta^T \mathbf{X}_i)/\{1 + \exp(\beta^T \mathbf{X}_i)\}$, then model (2) is quite different from the linear transformation model.

When γ , which specifies transformations in (3), is treated as an unknown parameter, the model parameters may not be identifiable. For example, suppose that $\theta(\mathbf{X}) = \exp(\beta_0)$. Then for any $\gamma \neq \tilde{\gamma}$, we can find a $\tilde{\beta}_0$, different from β_0 , such that

$$\left\{1 + \gamma e^{\beta_0}\right\}^{-1/\gamma} = \left\{1 + \tilde{\gamma} e^{\tilde{\beta}_0}\right\}^{-1/\tilde{\gamma}}.$$

Thus, for any distribution function $F(t)$, we define $\tilde{F}(t)$ so that

$$\left\{1 + \gamma e^{\beta_0} F(t)\right\}^{-1/\gamma} = \left\{1 + \tilde{\gamma} e^{\tilde{\beta}_0} \tilde{F}(t)\right\}^{-1/\tilde{\gamma}}.$$

Clearly, $\tilde{F}(t)$ is also a distribution function. Consequently, the two sets of parameters (γ, β_0, F) and $(\tilde{\gamma}, \tilde{\beta}_0, \tilde{F})$ give the same survival function so they are not distinguishable from the observed data. More identifiability results are given in Section 4. Additionally, in most practical applications, there is little information in the data to estimate γ with a reasonable degree of precision for small to even moderately large sample sizes. In these situations, the likelihood function of γ is flat. Our experience shows that γ can be well estimated when the sample size is very large, such as $n = 1500$ or larger. Due to these limitations, we will focus on the γ fixed case throughout the development of our model and asymptotic theory. However, in Section 4, we will discuss estimation of γ when it is identifiable and also suggest a model selection strategy for choosing γ in the γ fixed case.

The transformation in (2) may not necessarily be from the family (3); different transformations are possible when ξ takes other distributions. For example, we may consider the following Box-Cox type transformations:

$$G_\gamma(x) = \begin{cases} \exp\left\{-\frac{(1+x)^\gamma - 1}{\gamma}\right\} & \gamma > 0 \\ 1/(1+x) & \gamma = 0. \end{cases} \quad (4)$$

In this family, $\gamma = 1$ yields the proportional hazards model while $\gamma = 0$ yields the proportional odds model. In this paper, we study general classes of transformations $G(\cdot)$ and link functions $\theta(\cdot)$, and examine inference based on maximum likelihood estimation. However, for ease and clarity of exposition, we will focus on the class in (3) or (4), and $\theta(\mathbf{X}_i) = \exp(\boldsymbol{\beta}^T \mathbf{X}_i)$ in the examples of Section 5. In addition, the promotion time cumulative distribution functions, $F(t)$, will be completely unspecified, and thus estimated nonparametrically throughout.

The rest of the paper is organized as follows. In Section 2, we introduce notation and propose an efficient computational algorithm for the maximum likelihood estimation procedure. In Section 3, we derive the asymptotic properties of the parameter estimates, including consistency and asymptotic normality. In Section 4, we discuss important issues of model selection, including estimation of γ when it is identifiable as well as the selection of γ when it is treated as fixed. In Section 5, we conduct simulation studies to evaluate the finite sample properties of the estimators and also illustrate the proposed model with a real dataset. Some concluding remarks are given in Section 6. Technical details for the proofs of the theorems are given in the Appendix.

2 Maximum Likelihood Estimation

Suppose that there are n i.i.d. right-censored observations, $\{Y_i = T_i \wedge C_i, \mathbf{X}_i, \Delta_i = I(T_i \leq C_i); i = 1, \dots, n\}$, where $T_i \wedge C_i = \min(T_i, C_i)$ and $I(\cdot)$ is the indicator function. We assume that the follow-up time is infinite and a proportion of subjects never experience failure or right-censoring, that is $Y_i = \infty$ (so $C_i = \infty$) with probability one for some subjects. The right-censoring time C_i is assumed to be conditionally independent of T_i given \mathbf{X}_i and has a finite hazard rate almost everywhere. We assume that model (2) is used to link T_i with the covariate vector \mathbf{X}_i , where $\theta(\mathbf{X}_i) = \eta(\boldsymbol{\beta}^T \mathbf{X}_i)$, $\eta(\cdot)$ is a known and strictly positive link function, and $\boldsymbol{\beta}$ includes an intercept term.

Thus, the observed-data likelihood function of the parameters $(\boldsymbol{\beta}, F)$ is given by

$$\prod_{i=1}^n \left\{ \left[\{-G'(\eta(\boldsymbol{\beta}^T \mathbf{X}_i)F(Y_i))\eta(\boldsymbol{\beta}^T \mathbf{X}_i)f(Y_i)\}^{\Delta_i} \{G(\eta(\boldsymbol{\beta}^T \mathbf{X}_i)F(Y_i))\}^{(1-\Delta_i)} \right]^{I(Y_i < \infty)} \times [G(\eta(\boldsymbol{\beta}^T \mathbf{X}_i))]^{I(Y_i = \infty)} \right\}, \quad (5)$$

where $G'(x)$ denotes the derivative of G with respect to x , and $f(\cdot)$ is the density function corresponding to the distribution function $F(\cdot)$ with respect to Lebesgue measure. We wish to maximize the above likelihood function to obtain the maximum likelihood estimates for $\boldsymbol{\beta}$ and F . However, this maximum does not exist since one can choose $f(Y_i) = \infty$ for some Y_i with $\Delta_i = 1$. Thus, we

apply a nonparametric maximum likelihood estimation (NPMLE) approach, where F is allowed to be a right-continuous function. Instead of maximizing (5), we maximize the following modified function,

$$\prod_{i=1}^n \left\{ \left[\{-G'(\eta(\beta^T \mathbf{X}_i)F(Y_i))\eta(\beta^T \mathbf{X}_i)F\{Y_i\}\}^{\Delta_i} \{G(\eta(\beta^T \mathbf{X}_i)F(Y_i))\}^{(1-\Delta_i)} \right]^{I(Y_i < \infty)} \times [G(\eta(\beta^T \mathbf{X}_i))]^{I(Y_i = \infty)} \right\}, \quad (6)$$

where $F\{Y_i\}$ is the jump size of F at Y_i . The maximum likelihood estimate for F is termed the NPMLE for F and it is easy to show that the estimate for F must be a distribution function only with point masses at the observed Y_i with $\Delta_i = 1$. In order to estimate $F(t)$ nonparametrically, we must decide upon a follow-up time such that all censored observations beyond that follow-up time, called the *cure threshold*, are treated as “ $Y_i = \infty$ ” (i.e., observed to be cured) and all observations lower than this threshold are treated as $Y_i < \infty$ (i.e., observed to be either failures or right-censored). This assumption is needed so that the model is identifiable in (β, F) , as shown in Section 3. Note that if a parametric form is assumed for F , as in Ibrahim, Chen and Sinha (2001), then the condition that some of the Y_i ’s are observed to be infinity is not needed.

To compute the maximum likelihood estimates, we first derive the F which maximizes (6) for fixed β . Equivalently, we maximize the logarithm of (6) which is equal to

$$\sum_{i=1}^n I(Y_i < \infty) [\Delta_i \log p_i + \Delta_i \log \{-G'(\eta(\beta^T \mathbf{X}_i)F_i)\} + (1 - \Delta_i) \log G(\eta(\beta^T \mathbf{X}_i)F_i)],$$

subject to the constraint $\sum_j \Delta_j I(Y_j < \infty) p_j = 1$, where $p_i = F\{Y_i\}$ denotes the jump size of F at Y_i and $F_i = \sum_{Y_j \leq Y_i, \Delta_j = 1} p_j$. If we order the observed failure times from the smallest to the largest and use the indices $(1), \dots, (m)$ for the ordered times, $Y_{(1)} < \dots < Y_{(m)}$, where $m = \sum_i \Delta_i I(Y_i < \infty)$, then after introducing the Lagrange-multiplier λ , we obtain $p_{(i)}$ by solving the equation

$$\frac{1}{p_{(i)}} + \sum_{j=1}^n \left\{ \Delta_j \frac{G''(\eta(\beta^T \mathbf{X}_j)F_j)\eta(\beta^T \mathbf{X}_j)I(Y_{(i)} \leq Y_j < \infty)}{G'(\eta(\beta^T \mathbf{X}_j)F_j)} + (1 - \Delta_j) \frac{G'(\eta(\beta^T \mathbf{X}_j)F_j)\eta(\beta^T \mathbf{X}_j)I(Y_{(i)} \leq Y_j < \infty)}{G(\eta(\beta^T \mathbf{X}_j)F_j)} \right\} - \lambda = 0,$$

where $G''(x)$ denotes the second derivative of G with respect to x . Thus, it follows that

$$\frac{1}{p_{(i+1)}} = \frac{1}{p_{(i)}} + \sum_{Y_{(i)} \leq Y_j < Y_{(i+1)}} \left\{ \Delta_j \frac{G''(\eta(\beta^T \mathbf{X}_j)F_j)\eta(\beta^T \mathbf{X}_j)}{G'(\eta(\beta^T \mathbf{X}_j)F_j)} + (1 - \Delta_j) \frac{G'(\eta(\beta^T \mathbf{X}_j)F_j)\eta(\beta^T \mathbf{X}_j)}{G(\eta(\beta^T \mathbf{X}_j)F_j)} \right\}.$$

Equivalently,

$$\frac{1}{p_{(i+1)}} = \frac{1}{p_{(i)}} + \frac{G''(\eta(\beta^T \mathbf{X}_{(i)})F_{(i)})\eta(\beta^T \mathbf{X}_{(i)})}{G'(\eta(\beta^T \mathbf{X}_{(i)})F_{(i)})} + \sum_{Y_{(i)} < Y_j < Y_{(i+1)}} \frac{G'(\eta(\beta^T \mathbf{X}_j)F_{(i)})\eta(\beta^T \mathbf{X}_j)}{G(\eta(\beta^T \mathbf{X}_j)F_{(i)})}, \quad (7)$$

where $F_{(i)} = p_{(1)} + \dots + p_{(i)}$. Using the fact that $\sum_{i=1}^m p_{(i)} = 1$, we can also write (7) as

$$\begin{aligned} \frac{1}{p_{(i)}} &= \frac{1}{p_{(i+1)}} - \frac{G''(\eta(\beta^T \mathbf{X}_{(i)})(1 - S_{(i+1)}))\eta(\beta^T \mathbf{X}_{(i)})}{G'(\eta(\beta^T \mathbf{X}_{(i)})(1 - S_{(i+1)}))} \\ &\quad - \sum_{Y_{(i)} < Y_j < Y_{(i+1)}} \frac{G'(\eta(\beta^T \mathbf{X}_j)(1 - S_{(i+1)}))\eta(\beta^T \mathbf{X}_j)}{G(\eta(\beta^T \mathbf{X}_j)(1 - S_{(i+1)}))}, \end{aligned} \quad (8)$$

where $S_{(i+1)} = p_{(i+1)} + p_{(i+2)} + \dots + p_{(m)}$. From (7), we obtain a recursive formula of calculating $p_{(i+1)}$ from $p_{(i)}$ and $F_{(i)}$; while from (8), we obtain another recursive formula of calculating $p_{(i)}$ from $p_{(i+1)}$ and $S_{(i+1)}$. When $G'' > 0$ and $G' < 0$, we prefer to using (8) since it ensures that $0 < p_{(i)} < p_{(i+1)}$ once $p_{(i+1)} > 0$ and $S_{(i+1)} < 1$.

Hence, from (8), we can treat β , $\alpha \equiv p_{(m)} > 0$ and λ as independent parameters, and $p_{(1)}, \dots, p_{(m-1)}$ are functions of β and α . Then, the constrained maximum likelihood equations for β and $p_{(1)}, \dots, p_{(m)}$ can be reduced to solving the following score equations for β , α and λ ,

$$\begin{aligned} 0 &= \sum_{i=1}^m \frac{1}{p_{(i)}} \frac{\partial}{\partial \beta} p_{(i)} + \sum_{i=1}^m \frac{G''(\eta(\beta^T \mathbf{X}_{(i)})F_{(i)})}{G'(\eta(\beta^T \mathbf{X}_{(i)})F_{(i)})} \left\{ \eta'(\beta^T \mathbf{X}_{(i)})\mathbf{X}_{(i)}F_{(i)} + \eta(\beta^T \mathbf{X}_{(i)}) \frac{\partial}{\partial \beta} F_{(i)} \right\} \\ &\quad + \sum_{i=1}^m \sum_{Y_{(i)} < Y_j < Y_{(i+1)}} \frac{G'(\eta(\beta^T \mathbf{X}_j)F_{(i)})}{G(\eta(\beta^T \mathbf{X}_j)F_{(i)})} \left\{ \eta'(\beta^T \mathbf{X}_j)\mathbf{X}_jF_{(i)} + \eta(\beta^T \mathbf{X}_j) \frac{\partial}{\partial \beta} F_{(i)} \right\} \\ &\quad + \sum_{j=1}^n \Delta_j \mathbf{X}_j + \sum_{j=1}^n I(Y_j = \infty) \frac{G'(\eta(\beta^T \mathbf{X}_j))}{G(\eta(\beta^T \mathbf{X}_j))} \eta'(\beta^T \mathbf{X}_j)\mathbf{X}_j - \lambda \sum_{i=1}^m \frac{\partial}{\partial \beta} p_{(i)}, \\ 0 &= \sum_{i=1}^m \frac{1}{p_{(i)}} \frac{\partial}{\partial \alpha} p_{(i)} + \sum_{i=1}^m \frac{G''(\eta(\beta^T \mathbf{X}_{(i)})F_{(i)})}{G'(\eta(\beta^T \mathbf{X}_{(i)})F_{(i)})} \eta(\beta^T \mathbf{X}_{(i)}) \frac{\partial}{\partial \alpha} F_{(i)} \\ &\quad + \sum_{i=1}^m \sum_{Y_{(i)} < Y_j < Y_{(i+1)}} \frac{G'(\eta(\beta^T \mathbf{X}_j)F_{(i)})}{G(\eta(\beta^T \mathbf{X}_j)F_{(i)})} \eta(\beta^T \mathbf{X}_j) \frac{\partial}{\partial \alpha} F_{(i)} - \lambda \sum_{i=1}^m \frac{\partial}{\partial \alpha} p_{(i)}, \\ 0 &= \sum_{i=1}^m p_{(i)} - 1. \end{aligned} \quad (9)$$

After eliminating λ from the first two equations, the Newton-Raphson algorithm can be used to solve the system of equations in (9). The first and second derivatives of $p_{(i)}$ with respect to β and α can be computed using the recursive formula (8).

We denote the maximum likelihood estimators for β and α by $\hat{\beta}_n$ and $\hat{\alpha}_n$, respectively. We can estimate the asymptotic variance of $(\hat{\beta}_n, \hat{\alpha}_n)$ based on the profile log-likelihood function for (β, α) , which is defined as the maximum value of the logarithm of (6) for any fixed (β, α) and is denoted by $pl_n(\beta, \alpha)$. The asymptotic variance of $(\hat{\beta}_n, \hat{\alpha})$ can be estimated using the negative inverse of the

curvature of $pl_n(\boldsymbol{\beta}, \alpha)$ at $(\hat{\boldsymbol{\beta}}_n, \hat{\alpha}_n)$, i.e.,

$$- \left(\begin{array}{cc} \frac{\partial^2}{\partial \boldsymbol{\beta}^2} pl_n(\boldsymbol{\beta}, \alpha) & \frac{\partial^2}{\partial \boldsymbol{\beta} \partial \alpha} pl_n(\boldsymbol{\beta}, \alpha) \\ \frac{\partial^2}{\partial \alpha \partial \boldsymbol{\beta}} pl_n(\boldsymbol{\beta}, \alpha) & \frac{\partial^2}{\partial \alpha^2} pl_n(\boldsymbol{\beta}, \alpha) \end{array} \right)^{-1} \Big|_{\boldsymbol{\beta}=\hat{\boldsymbol{\beta}}_n, \alpha=\hat{\alpha}_n}.$$

Specifically, the second derivative of $pl_n(\boldsymbol{\beta}, \alpha)$ with respect to $\boldsymbol{\beta}$ and α can be calculated based on the following chain rule and the recursive formula (8):

$$\begin{aligned} \frac{\partial}{\partial \boldsymbol{\beta}} pl_n(\boldsymbol{\beta}, \alpha) &= \frac{\partial}{\partial \boldsymbol{\beta}} l_n(\boldsymbol{\beta}, F) + \sum_{i=1}^{m-1} \frac{\partial l_n(\boldsymbol{\beta}, F)}{\partial p_{(i)}} \frac{\partial p_{(i)}}{\partial \boldsymbol{\beta}}, \\ \frac{\partial}{\partial \alpha} pl_n(\boldsymbol{\beta}, \alpha) &= \frac{\partial}{\partial \alpha} l_n(\boldsymbol{\beta}, F) + \sum_{i=1}^{m-1} \frac{\partial l_n(\boldsymbol{\beta}, F)}{\partial p_{(i)}} \frac{\partial p_{(i)}}{\partial \alpha}, \end{aligned}$$

where $l_n(\boldsymbol{\beta}, F)$ is the logarithm value of (6). The justification of the above variance estimation method is based on the profile likelihood theory in Murphy and van der Vaart (2000), and is discussed in the Appendix.

3 Asymptotic Properties

In this section, we establish theorems characterizing the asymptotic properties of $(\hat{\boldsymbol{\beta}}_n, \hat{\alpha}_n)$. In order to achieve consistency and asymptotic normality, we first need the following assumptions:

- (C1) The covariate \mathbf{X} is bounded with probability one, and if there exists a vector $\tilde{\boldsymbol{\beta}}$ such that $\tilde{\boldsymbol{\beta}}^T \mathbf{X} = 0$ with probability one, then $\tilde{\boldsymbol{\beta}} = 0$.
- (C2) Conditional on \mathbf{X} , the right-censoring time C is independent of T , and $P(C = \infty | \mathbf{X}) > 0$.
- (C3) The true value of $\boldsymbol{\beta}$, denoted as $\boldsymbol{\beta}_0$, belongs to the interior of a known compact set \mathcal{B}_0 , and the true promotion time cumulative distribution function F_0 is differentiable with $F'_0(x) > 0$ for all $x \in R^+$.
- (C4) The link function $\eta(\cdot)$ is strictly increasing and twice-continuously differentiable with $\eta(\cdot) > 0$. Furthermore, the transformation G satisfies

$$G(0) = 1, G(x) > 0, G'(x) < 0, G^{(3)}(x) \text{ exists and is continuous,}$$

where $G^{(3)}(x)$ is the third derivative of $G(x)$.

Condition (C1) is the usual condition for a design matrix in regression settings. The condition $P(C = \infty | \mathbf{X})$ in (C2) ensures that at least some cured subjects are not right-censored; otherwise, if

all subjects either fail or are right-censored, then intuitively, one would be unable to identify the cure rate. In (C3), β is assumed to be bounded. Such an assumption is often imposed in semiparametric inference, as practical calculation is always performed within a reasonable bounded set. Many link functions $\eta(\cdot)$ and $G(\cdot)$ satisfy the conditions in (C4). Some examples of $\eta(\cdot)$ include $\eta(x) = e^x$, $\eta(x) = e^x/(1+e^x)$, $\eta(x) = \Phi(x)$ where Φ is the cumulative distribution function of the standard normal distribution. Examples of transformations satisfying (C4) include the transformations $(1 + \gamma x)^{-1/\gamma}$ for $\gamma > 0$ and $\exp(-x)$ for $\gamma = 0$, as well as some others, such as $G(x) = \{1 + \log(1 + x)\}^{-\gamma}$ for $\gamma > 0$ and $G(x) = \exp\{-(1 + x)^\gamma - 1\}/\gamma$ for $\gamma > 0$.

Before stating the main results, we first show that under conditions (C1) - (C4), the parameters β and F are identifiable. Suppose that two sets of parameters (β, F) and $(\tilde{\beta}, \tilde{F})$ give the same likelihood function for the observed data. We claim that $\beta = \tilde{\beta}$ and $F = \tilde{F}$. Since

$$\begin{aligned} & \left[\left\{ -G'(\eta(\beta^T \mathbf{X})F(Y))\eta(\beta^T \mathbf{X})f(Y) \right\}^\Delta \left\{ G(\eta(\beta^T \mathbf{X})F(Y)) \right\}^{(1-\Delta)} \right]^{I(Y < \infty)} \\ & \quad \times \left[G(\eta(\beta^T \mathbf{X})) \right]^{I(Y = \infty)} \\ = & \left[\left\{ -G'(\eta(\tilde{\beta}^T \mathbf{X})\tilde{F}(Y))\eta(\tilde{\beta}^T \mathbf{X})\tilde{f}(Y) \right\}^\Delta \left\{ G(\eta(\tilde{\beta}^T \mathbf{X})\tilde{F}(Y)) \right\}^{(1-\Delta)} \right]^{I(Y < \infty)} \\ & \quad \times \left[G(\eta(\tilde{\beta}^T \mathbf{X})) \right]^{I(Y = \infty)}, \end{aligned} \quad (10)$$

we choose $Y = \infty$. Then from the monotonicity of both G and η , it follows that $\beta^T \mathbf{X} = \tilde{\beta}^T \mathbf{X}$. Thus, condition (C1) gives $\beta = \tilde{\beta}$. Furthermore, by letting $\Delta = 1$ and $Y = y$ and integrating both sides of (10) from 0 to y , we have $G(\eta(\beta^T \mathbf{X})F(y)) = G(\eta(\tilde{\beta}^T \mathbf{X})\tilde{F}(y))$. Therefore, $F(y) = \tilde{F}(y)$.

The following theorem establishes the consistency of the maximum likelihood estimator.

Theorem 1. Under conditions (C1) - (C4), with probability one,

$$|\hat{\beta}_n - \beta_0| \rightarrow 0 \quad \text{and} \quad \sup_{t \in R^+} |\hat{F}_n(t) - F_0(t)| \rightarrow 0,$$

i.e., both $\hat{\beta}_n$ and \hat{F}_n are strongly consistent.

The basic idea in proving Theorem 1 is as follows: Suppose that $\hat{\beta}_n$ and \hat{F}_n converge to β^* and F^* , respectively. We first construct an empirical distribution function \tilde{F}_n converging to F_0 . Then since $\left\{ l_n(\hat{\beta}_n, \hat{F}_n) - l_n(\beta_0, \tilde{F}_n) \right\} / n \geq 0$, where $l_n(\beta, F)$ denotes the observed log-likelihood function at (β, F) , and this difference converges to the negative Kullback-Leibler divergence between (β^*, F^*) and (β_0, F_0) , the identifiability result gives $\beta^* = \beta_0$ and $F^* = F_0$. This establishes the consistency result in Theorem 1. Constructing the empirical function \tilde{F}_n and using the Kullback-Leibler divergence to prove consistency has been used by many others in semiparametric theory, including Murphy (1994), Murphy, Rossini and van der Vaart (1997), Parner (1998), Slud and Vonta (2004), and Kosorok, Lee

and Fine (2004) among others. However, observing the fact that \hat{F}_n is a distribution function, proving the convergence of the Kullback-Leibler divergence is not trivial in our case, as shown in the Appendix.

Our second result concerns the joint asymptotic distribution of $\hat{\beta}_n$ and \hat{F}_n . In order to obtain the joint asymptotic distribution for $(\hat{\beta}_n, \hat{F}_n)$, we first introduce the set

$$\mathcal{H} = \left\{ (\mathbf{h}_1, h_2) : \mathbf{h}_1 \in R^d, \|\mathbf{h}_1\| < 1, \right.$$

$h_2 \text{ is a function in } [0, \infty) \text{ with its total variation bounded by } 1 \}.$

Here, the total variation of a function h_2 is defined as the supremum of $\sum_{i=1}^m |h_2(t_{i+1}) - h_2(t_i)|$ over all finite partitions $0 = t_1 < t_2 < \dots < t_{m+1} = \infty$. We use $\|h_2\|_V$ to denote the total variation of h_2 . Then $\sqrt{n}(\hat{\beta}_n - \beta_0, \hat{F}_n - F_0)$ can be treated as a linear functional in $l^\infty(\mathcal{H})$, the space of all bounded linear functionals on \mathcal{H} , which is defined as

$$\sqrt{n}(\hat{\beta}_n - \beta_0, \hat{F}_n - F_0)[\mathbf{h}_1, h_2] = \sqrt{n}(\hat{\beta}_n - \beta_0)^T \mathbf{h}_1 + \sqrt{n} \int h_2(t) d(\hat{F}_n - F_0).$$

The next theorem establishes the asymptotic distribution of $\sqrt{n}(\hat{\beta}_n - \beta_0, \hat{F}_n - F_0)$ in the metric space $l^\infty(\mathcal{H})$.

Theorem 2. Under conditions (C1)-(C4), $\sqrt{n}(\hat{\beta}_n - \beta_0, \hat{F}_n - F_0)$ converges weakly to a zero-mean Gaussian process in $l^\infty(\mathcal{H})$. Furthermore, $\hat{\beta}_n$ is efficient; equivalently, its asymptotic variance attains the semiparametric efficiency bound for β_0 .

The covariance matrix of the asymptotic Gaussian process is given in the Appendix. The definition of the semiparametric efficiency bound can be found in Chapter 3 of Bickel, Klaassen, Ritov and Wellner (1993). Thus, Theorem 2 establishes that the maximum likelihood estimators are asymptotically normal and efficient. The proof of Theorem 2 is standard in most of the current semiparametric literature, including Murphy (1995), Parner (1998) and Kosorok et al. (2004). The proof relies on the linearization of the likelihood equations for $\hat{\beta}_n$ and \hat{F}_n and uses Theorem 3.3.1 of van der Vaart and Wellner (1996). In the proof, the verification of some Donsker classes and proving the invertibility of the information operator are the key steps. Both issues are discussed in detail in the Appendix for the proposed model.

Theorem 2 has many useful applications. By letting $h_2(\cdot) = I(\cdot \leq t)$ for any $t \geq 0$, we obtain that $\sqrt{n}(\hat{\beta}_n - \beta_0, \hat{F}(t) - F_0(t))$ converges weakly to a zero-mean Gaussian process in $l^\infty(R^d \times [0, \infty))$. As a result, for fixed t_0 , $\sqrt{n}(\hat{F}_n(t_0) - F_0(t_0))$ has an asymptotic normal distribution with mean zero. If its asymptotic variance can be estimated, one can easily construct a confidence interval for $F_0(t_0)$. Special choices of t_0 can be the quantiles of F_0 . Furthermore, when interest is to test whether the

true promotion distribution function is equal to a given distribution function F_0 , we can construct a test statistic $\sqrt{n} \sup_{t \geq 0} |\hat{F}_n(t) - F_0(t)|$, similar to the Kolmogorov-Smirnov statistic. Then Theorem 2 implies that such a statistic has an asymptotic distribution which is the same as the supremum of a Gaussian process. We remark that in the above cases, the asymptotic covariance function of the Gaussian process in Theorem 2 needs to be estimated. One practical way of estimating this function is through a bootstrapping approach. The justification of the bootstrapping procedure can be shown using the same techniques as in Kosorok et al. (2004). We will not pursue this issue further here but rather focus only on inference for regression coefficients in the subsequent development.

4 Estimation of the Transformation $G(\cdot)$

In the forgoing sections, the transformation $G(\cdot)$ was assumed to be known. One important practical issue is how to estimate $G(\cdot)$ using the observed data. We discuss two possible methods to estimate this transformation.

The first approach is to consider $G(\cdot)$ from a parametric transformation family $\{G_\gamma : \gamma \in \Gamma\}$, where Γ is a compact set in Euclidean space. For example, G_γ arises from the family given in (3) or (4). Using the observed data, we then estimate γ along with β and F . However, as noted in the introduction, one serious problem with this approach is the possible nonidentifiability of γ . However, for some special families of transformations, the parameters (γ, β, F) are identifiable, as given in the following proposition.

Proposition 1. Let $\mathbf{X} = (1, \mathbf{W}^T)^T$ and β_0 as $(\beta_{01}, \beta_{0w}^T)^T$, respectively. Assume that \mathbf{W} has support containing a nonempty open interior and $\beta_{0w}^T \mathbf{W} \neq 0$. Then for transformations from the family (3) and $\eta(x) = \exp(x)$, β_0, F_0 and γ_0 are identifiable.

Proof. Suppose that $(\tilde{\beta}, \tilde{F}, \tilde{\gamma})$ gives the same observed likelihood function as (β_0, F_0, γ_0) . That is,

$$\begin{aligned} & \left[\left\{ -G'_{\gamma_0}(\eta(\beta_0^T \mathbf{X}) F_0(Y)) \eta(\beta_0^T \mathbf{X}) f_0(Y) \right\}^\Delta \left\{ G_{\gamma_0}(\eta(\beta_0^T \mathbf{X}) F_0(Y)) \right\}^{(1-\Delta)} \right]^{I(Y < \infty)} \\ & \quad \times \left[G_{\gamma_0}(\eta(\beta_0^T \mathbf{X})) \right]^{I(Y = \infty)} \\ &= \left[\left\{ -G'_{\tilde{\gamma}}(\eta(\tilde{\beta}^T \mathbf{X}) \tilde{F}(Y)) \eta(\tilde{\beta}^T \mathbf{X}) \tilde{f}(Y) \right\}^\Delta \left\{ G_{\tilde{\gamma}}(\eta(\tilde{\beta}^T \mathbf{X}) \tilde{F}(Y)) \right\}^{(1-\Delta)} \right]^{I(Y < \infty)} \\ & \quad \times \left[G_{\tilde{\gamma}}(\eta(\tilde{\beta}^T \mathbf{X})) \right]^{I(Y = \infty)}, \end{aligned} \tag{11}$$

where $G_\gamma(x) = (1 + \gamma x)^{-1/\gamma}$. We choose $Y = \infty$ in (11) and obtain

$$\left\{ 1 + \tilde{\gamma} \exp(\tilde{\beta}^T \mathbf{X}) \right\}^{1/\tilde{\gamma}} = \left\{ 1 + \gamma_0 \exp(\beta_0^T \mathbf{X}) \right\}^{1/\gamma_0}.$$

Since both sides are analytic in \mathbf{W} , this equality holds for any \mathbf{W} in real space. If $\gamma_0 < \tilde{\gamma}$, then from the monotonicity of $(1 + \gamma x)^{1/\gamma}$, we have $\beta_0^T \mathbf{X} > \tilde{\beta}^T \mathbf{X}$ for any \mathbf{X} . Immediately, we conclude that $\tilde{\beta}_{0w} = \beta_{0w}$ and $\beta_{01} < \tilde{\beta}_{01}$. As a result, we have

$$\left\{1 + \tilde{\gamma} \exp(\tilde{\beta}_{01} + \beta_{0w}^T \mathbf{W})\right\}^{1/\tilde{\gamma}} = \left\{1 + \gamma_0 \exp(\beta_{01} + \beta_{0w}^T \mathbf{W})\right\}^{1/\gamma_0},$$

and this holds for any real \mathbf{W} . Letting $\beta_{0w}^T \mathbf{W} \rightarrow \infty$, we then obtain $\gamma_0 = \tilde{\gamma}$ and $\beta_{01} = \tilde{\beta}_{01}$. Furthermore, choosing $\Delta = 1$ and $Y = y$ and integrating from 0 to y in (11), we obtain $\tilde{F}(y) = F_0(y)$.

Proposition 1 states that if a continuous covariate has a non-zero effect, then γ can be identified. When model parameters are identifiable, with some additional regularity conditions beyond (C.1)-(C.4), the nonparametric maximum likelihood estimators for β, F and γ are strongly consistent and asymptotically normal. The details are given in the remarks of the Appendix. This approach utilizes the observed data to estimate the transformation parameter and our proposed algorithm can be easily adapted to incorporate this extra parameter estimation. However, this approach may not be useful for practical applications for the following reasons: First, with no prior knowledge about the true covariate effects, there is always the concern regarding identifying all of the parameters in the model since nonidentifiability can cause numerical instability in the computations. Second, even if the parameters are identifiable, our experience indicates that for small samples, the likelihood function is typically quite flat as a function of γ . Thus, to obtain an accurate estimate of γ , a very large sample size is required and this may not be practical in many biomedical studies. Third, when the choices of transformations are from multiple families of transformations which are parameterized differently, this approach is no longer feasible.

Hence, we suggest the following approach for estimating the transformation G in practice. When many transformations are under consideration, we can calculate the NPMLEs under each transformation then choose the transformation which maximizes the Akaike information criterion (AIC). The AIC is defined as the twice log-likelihood function minus twice the number of parameters. In some applications, to obtain algebraically simple transformations, one may also penalize the complexity of the transformation. Some possible choices of a penalty can be the maximal difference between $G(x)$ and $\exp(-x)$, so that we can choose a model close to the proportional hazards model; or, the choice can be the maximal difference between $G(x)$ and $1/(1+x)$, so that we can choose a model close to the proportional odds model. However, the determination of the transformation complexity remains an unsolved issue so we defer further discussion to future work. Besides the AIC criterion, other criteria can also be used as well, including the Bayesian information criterion (BIC, Schwarz, 1978), the L measure (Ibrahim and Laud, 1994), or likelihood-based cross-validation. As an additional note, in

most practice, the inference is solely based on the selected model; therefore, the variance estimate does not reflect the variation due to the model selection procedure. The correction of the variance estimate, sometimes called post-model-selection inference, is still an open problem in semiparametric inference.

In the subsequent simulation study, we will examine the performance of the NPMLEs for a fixed transformation; while, in the data application, the AIC will be used to select the best transformation to fit the data.

5 Numerical Studies

5.1 Simulation

We conducted simulation studies to examine the small-sample performance of our proposed methodology. In the first simulation study, the transformation cure model had survival function of the form

$$S(t|X_1, X_2) = \{1 + \gamma \exp(\beta_0 + \beta_1 X_1 + \beta_2 X_2) F(t)\}^{-1/\gamma},$$

where X_1 was a uniformly distributed random variable in $[0, 1]$, X_2 was a Bernoulli random variable, $\beta_0 = 0.5, \beta_1 = 1, \beta_2 = -0.5$, and $F(t) = 1 - \exp(-t)$. We chose γ to vary from 0 to 1. Moreover, each subject had a 40% chance of being right-censored and the censoring time was generated from an exponential distribution with mean 1. The censoring proportions varied from 17% to 22% as γ changed from 0 to 1; while the cure rate could be as low as 8% when $\gamma = 0$ and became 20% when $\gamma = 1$. For each simulated dataset, the proposed method of Section 2 was implemented to calculate the maximum likelihood estimates of β and its corresponding variance estimate. In solving the score equations using the Newton-Raphson iterations, the initial values for β were set to zero and the initial value for α was set to $1/n$, with n being the sample size. Other initial values were also tested in the simulation study and results were very robust to those choices. The convergence of each simulation was fast and often obtained within ten iterations.

Table 1 summarizes the results from 1000 replications for each combination of γ and n : the column labeled “Estimate” denotes the average values of the estimates; “SE” is the sample standard error of the estimates; “ESE” is the average of the estimated standard errors; and “CP” is the coverage proportion of 95% confidence intervals constructed based on the asymptotic normal approximation. The results in Table 1 indicate that the proposed estimation method performs well with sample sizes of 100 and 200: the biases are small, the estimated standard errors agree well with the sample standard errors, and the coverage probabilities are accurate.

In the second simulation study, we generated the failure time from the transformation cure model with survival function

$$S(t|X_1, X_2) = \exp[-\{(1 + \gamma \exp(\beta_0 + \beta_1 X_1 + \beta_2 X_2)F(t))^\gamma - 1\} / \gamma],$$

where $F(t) = 1 - \exp(-t)$ and the covariates and censoring time were generated using the same distributions as in the first simulation. In this setting, we also varied γ from 0 to 1, where $\gamma = 0$ corresponds to the proportional odds cure model and $\gamma = 1$ corresponds to the proportional hazards cure model. The censoring proportion and the cure rate were 22% and 20% when $\gamma = 0$ and became 17% and 8% when $\gamma = 1$. The results based on 1000 repetitions for sample sizes 100 and 200 are summarized in Table 2. From Table 2, we obtain the same conclusions as in the first simulation study. Thus, we conclude that the maximum likelihood estimation procedure proposed here not only provides an asymptotically efficient estimator, but also yields good inferential properties for small sample sizes.

Since the proportional hazards cure model and the proportional odds cure model are commonly used in practice, we also conducted a simulation study to examine the performance of the estimates based on these two models when data were generated from a different model. Specifically, we utilized the same setting for generating the covariates and censoring time as in the other two simulations described above, while we generated the survival time either from the model with a transformation $(1 + x/2)^{-2}$ or $\exp\{-2((1 + x)^{1/2} - 1)\}$; equivalently, $\gamma = 1/2$ in both classes of (3) and (4). Both choices corresponded to a model between the proportional hazards cure model and the proportional odds cure model. The results based on 1000 replications are reported in Table 3. From Table 3, we observe that both the proportional hazards cure model and proportional odds cure models produce notable bias. Interestingly, both models estimate the direction of the coefficients correctly and the proportional hazards cure model tends to bias towards zero while the opposite is observed for the proportional odds cure model. The bias for the intercept term in both models is large but the bias for other covariate effects are relatively small. We also observe that even with sizable bias, standard error estimates of the regression coefficients corresponding to the covariates appear to be correct.

Finally, we considered the estimation of γ . We generated failure times using the cure model for the transformation class $G(x) = (1 + \gamma x)^{-1/\gamma}$. The simulation study, which is not shown here, indicates that the performance of the NPMLEs is poor and the convergence in calculating the NPMLEs is often problematic with a sample size of $n = 400$. This is due to the fact that the likelihood function tends to be flat when γ varies around the true value.

5.2 Application to Melanoma Data

As an illustration, we applied the transformation cure model in (2) to a phase III melanoma clinical trial conducted by the Eastern Cooperative Oncology Group (ECOG), labeled E1690 (Kirkwood et al., 2000). This trial consisted of two treatment arms with a total of $n = 427$ patients on the combined treatment arms, of which 241 patients experienced the event (cancer relapse). The response variable was relapse-free survival (RFS) time (in years). The covariates included in this analysis were treatment (high-dose interferon=1, observation=0), age (a continuous variable which ranged from 19.13 to 78.05 with a mean of 47.93 years), sex (female=1, male=0) and nodal category (taking a value of 0 if there were zero positive nodes, or 1 if there were one or more positive nodes). The median follow-up time for this study was 4.33 years, which is considered as a sufficient length of follow-up for this disease. The solid and dotted curves in Figure 1 show the Kaplan-Meier survival curves for the two treatment arms. We see that a reasonable plateau has been reached at the tails of the survival curves, and it appears that based on this period of follow-up, a cure rate model would be a suitable approach for the data. Cure rate models for the E1690 data were also considered in Chen, Harrington and Ibrahim (2002), and were shown to fit better than proper survival models. Based on Figure 1, we considered the subjects as “cured” if they were censored at 5.5 years or beyond. In the dataset, 30 subjects had censored RFS times greater than or equal to 5.5 years ($Y_i = \infty$). Patients with observed times less than 5.5 years were either failures or right-censored, and some of those right-censored subjects might indeed have been “cured” patients, but we cannot determine that due to the right-censoring.

We fit the proposed model in (2), where $G(x)$ comes from the family (3) as well as the family (4). We considered values of γ in $[0, 2]$. The maximum likelihood estimates for the regression coefficients of the proposed class of semiparametric transformation cure models were computed using the proposed method. Furthermore, we selected the best transformation among these two classes as the one that maximized the AIC criterion, which is equivalent to the observed log-likelihood function in this case since the number of parameters is constant. Figure 2 plots the observed log-likelihood functions obtained using the two classes of transformations. Interestingly, both classes select out the same best transformation, which corresponds to the proportional hazards cure model.

Consequently, we report the results from the proportional hazards cure model in the second panel of Table 4. The results show that both interferon treatment and sex did not significantly affect RFS, while age and nodal category did. Younger patients or those with zero positive nodes had significantly better RFS and thus were more likely to be “cured”, that is, not to have recurrence of melanoma. The results can also be used to estimate the cure rate for each group. For example, the estimated cure

rates for a 50 year old female patient with positive nodes under the interferon treatment is 41.0%. Furthermore, we plot in Figure 1 the fitted survival function within each treatment group, where the survival function is calculated as the empirical average of the predicted survival functions within each group. The dashed and dot-dashed lines in Figure 1 present the predicted survival functions and they agree with the Kaplan-Meier curves quite well.

As noted earlier, we treated censored subjects with RFS times 5.5 years or greater as “cured” to estimate the parameters. The choice of such a threshold value can be artificial unless it has some biological meaning. Thus, we also studied the sensitivity of the estimates to the choice of this threshold value. To do this, we varied the threshold value larger than the last failure (5 years), using values of 5.1, 5.5, 6, 6.5 and 7 years. The estimates of the coefficients only differ in the third decimal point, as shown in Table 4.

6 Discussion

We have proposed a class of semiparametric transformation cure models which are motivated by a specific biological process. This class is quite broad and it includes the well-known proportional hazards and proportional odds structures as two special cases. We have provided an efficient algorithm for calculating the maximum likelihood estimates. The maximum likelihood estimation procedure yields efficient estimators of the regression parameters. As one by-product, since model (2) reduces to a linear transformation model with a special choice of the link function $\theta(\cdot)$, the algorithm in Section 2 provides a simple way of calculating the maximum likelihood estimates for linear transformation models in general. Specifically, for a linear transformation model with $S(t|\mathbf{Z}_i) = G\{\exp(\beta_1^T \mathbf{Z}_i)\Lambda(t)\}$, we can reparameterize to make it a cure rate model by defining $F(t) = \Lambda(t)/\Lambda(\tau)$ and adding an intercept term $\log \Lambda(\tau)$ into the regression. Here, τ refers to the termination time of the study. Thus, treating any subjects censored at time τ as “cured”, we then implement our proposed algorithm to calculate the maximum likelihood estimates of the parameters.

The cure threshold for the E1690 melanoma data was taken to be 5.5 years. The choice of this cutoff value heavily depends on the dataset at hand and other practical elements, including the type of disease, the severity or stage, the corresponding treatment, and other patient prognostic factors that require expert opinion from the physician. A simple guideline is that there should not be any failures after the cure threshold. In fact, the estimates from the proposed method are very robust with respect to the choice of this threshold, as shown in Table 4.

The transformation $G(x)$ can be misspecified in practice due to limited knowledge or complex

relationships between the covariates and the time-to-event variable. Kosorok et al. (2004) gives some examples in univariate survival data showing that the regression parameters can be estimated up to the correct direction even if $G(x)$ is misspecified. The same ideas can be extended to our proposed model. However, the computation of such estimable quantities in the presence of nonidentifiable parameters is a very challenging problem.

In deriving (2), we assumed that the promotion time survival function, $S^*(t) = 1 - F(t)$, is the same for all tumor cells. One possible generalization to this is to incorporate covariates into $S^*(t)$, for example, to allow them to be different across treatments. In this case, the survival function of the tumor cell for the i th subject would be $\exp\{-\Lambda(t)e^{\boldsymbol{\zeta}^T \mathbf{Z}_i}\}$, where \mathbf{Z}_i is a covariate vector for treatment and other risk factors, and \mathbf{Z}_i may share the same components as \mathbf{X}_i . Thus, the population survival function of interest for subject i is

$$S(t|\mathbf{X}_i, \mathbf{Z}_i) = G\{(1 - e^{-\Lambda(t)e^{\boldsymbol{\zeta}^T \mathbf{Z}_i}})\theta(\mathbf{X}_i)\}.$$

Issues regarding model identifiability and maximum likelihood estimation in these general models are currently being investigated.

Appendix

A.1 Proof of Theorem 1

We introduce the following notation. Let \mathbf{P}_n and \mathbf{P} denote the empirical measure of n i.i.d observations and the expectation, respectively; i.e., for any measurable function $g(\Delta, Y, \mathbf{X})$ in $L_2(P)$,

$$\mathbf{P}_n[g(\Delta, Y, \mathbf{X})] = \frac{1}{n} \sum_{i=1}^n g(\Delta_i, Y_i, \mathbf{X}_i), \quad \mathbf{P}[g(\Delta, Y, \mathbf{X})] = E[g(\Delta, Y, \mathbf{X})].$$

From the Lagrange-multiplier calculation, \hat{F}_n satisfies the equation that for $Y_i < \infty$,

$$\begin{aligned} \frac{\Delta_i}{F\{Y_i\}} + \sum_{\infty > Y_j \geq Y_i} \left\{ \Delta_j \frac{G''(\eta(\hat{\boldsymbol{\beta}}_n^T \mathbf{X}_j)F(Y_j))\eta(\hat{\boldsymbol{\beta}}_n^T \mathbf{X}_j)}{G'(\eta(\hat{\boldsymbol{\beta}}_n^T \mathbf{X}_j)F(Y_j))} \right. \\ \left. + (1 - \Delta_j) \frac{G'(\eta(\hat{\boldsymbol{\beta}}_n^T \mathbf{X}_j)F(Y_j))\eta(\hat{\boldsymbol{\beta}}_n^T \mathbf{X}_j)}{G(\eta(\hat{\boldsymbol{\beta}}_n^T \mathbf{X}_j)F(Y_j))} \right\} = n\hat{\lambda}_n. \end{aligned}$$

We multiply both sides by $\hat{F}_n\{Y_i\}$ and sum over Y_i such that $Y_i < \infty$. We get

$$\hat{\lambda}_n = \frac{1}{n} \sum_{i=1}^n \Delta_i I(Y_i < \infty) + \int_0^\infty H_n(y, \hat{\boldsymbol{\beta}}_n, \hat{F}_n) d\hat{F}_n(y), \quad (\text{A.1.1})$$

where

$$H_n(y, \hat{\beta}_n, \hat{F}_n) = \frac{1}{n} \left[\sum_{Y_j < \infty} \left\{ \Delta_j \frac{G''(\eta(\hat{\beta}_n^T \mathbf{X}_j) \hat{F}_n(Y_j)) \eta(\hat{\beta}_n^T \mathbf{X}_j) I(Y_j \geq y)}{G'(\eta(\hat{\beta}_n^T \mathbf{X}_j) \hat{F}_n(Y_j))} \right. \right. \\ \left. \left. + (1 - \Delta_j) \frac{G'(\eta(\hat{\beta}_n^T \mathbf{X}_j) \hat{F}_n(Y_j)) \eta(\hat{\beta}_n^T \mathbf{X}_j) I(Y_j \geq y)}{G(\eta(\hat{\beta}_n^T \mathbf{X}_j) \hat{F}_n(Y_j))} \right\} \right].$$

Hence, $\hat{F}_n\{Y_i\} = \Delta_i/n(\hat{\lambda}_n - H_n(Y_i, \hat{\beta}_n, \hat{F}_n))$. Obviously, from (A.1.1), $\hat{\lambda}_n$ should be bounded by a constant with probability one. Thus, by choosing a subsequence, still indexed by $\{n\}$, we assume $\hat{\lambda}_n \rightarrow \lambda^*$. By choosing a further subsequence, we assume $\hat{\beta}_n \rightarrow \beta^*$ and $\hat{F}_n \rightarrow F^*$ pointwise.

We consider the following class

$$\mathcal{A}_1 = \left\{ \Delta \frac{G''(\eta(\beta^T \mathbf{X}) F(Y)) \eta(\beta^T \mathbf{X}) I(\infty > Y \geq y)}{G'(\eta(\beta^T \mathbf{X}) F(Y))} + (1 - \Delta) \frac{G'(\eta(\beta^T \mathbf{X}) F(Y)) \eta(\beta^T \mathbf{X}) I(\infty > Y \geq y)}{G(\eta(\beta^T \mathbf{X}) F(Y))} : \right. \\ \left. F \text{ is a distribution function, } \beta \in \mathcal{B}_0, y \in [0, \infty) \right\}.$$

First, $\{\beta^T \mathbf{X} : \beta \in \mathcal{B}_0\}$ and $\{F(Y) : F \text{ is a distribution function}\}$ are both Donsker classes, where the latter follows from Theorem 2.7.5 of van der Vaart and Wellner (1996). Since G , G' , G'' and η are continuously differentiable functions, the preservation of the Donsker property based on Theorem 2.10.6 of van der Vaart and Wellner (1996) implies that the classes

$$\left\{ G^{(k)}(\eta(\beta^T \mathbf{X}) F(Y)) : \beta \in \mathcal{B}_0, F \text{ is a distribution function} \right\}, \quad k = 0, 1, 2,$$

and $\{\eta(\beta^T \mathbf{X}) : \beta \in \mathcal{B}_0\}$ are Donsker classes. Furthermore, we note $G'(x)$ and $G(x)$ are both bounded away from zero when x is in a compact set. Thus, the preservation of the Donsker property under the summation, product and quotient, as given in Examples 2.10.7-2.10.9 of van der Vaart and Wellner (1996) gives that the class \mathcal{A}_1 is a Donsker class so is also a Glivenko-Cantelli class. As a result of the Glivenko-Cantelli theorem and the bounded convergence theorem, we conclude that uniformly in y , $H_n(y, \hat{\beta}_n, \hat{F}_n) \rightarrow H^*(y)$, where

$$H^*(y) = E \left[\Delta \frac{G''(\eta(\beta^{*T} \mathbf{X}) F^*(Y)) \eta(\beta^{*T} \mathbf{X}) I(\infty > Y \geq y)}{G'(\eta(\beta^{*T} \mathbf{X}) F^*(Y))} \right. \\ \left. + (1 - \Delta) \frac{G'(\eta(\beta^{*T} \mathbf{X}) F^*(Y)) \eta(\beta^{*T} \mathbf{X}) I(\infty > Y \geq y)}{G(\eta(\beta^{*T} \mathbf{X}) F^*(Y))} \right].$$

Moreover, the right-hand side of (A.1.1) converges to

$$\lambda^* = E\{\Delta I(Y < \infty)\} + E\{I(Y < \infty) \int_0^Y H^*(y) dF^*(y)\}.$$

Now we wish to show that $|\lambda^* - H^*(y)| > \delta^*$ for some positive constant δ^* . To see that, we first note that from $\sum_{i=1}^n \hat{F}_n\{Y_i\} = 1$, then

$$\begin{aligned} 1 &= \sum_{i=1}^n I(Y_i < \infty) \frac{\Delta_i}{n(\hat{\lambda}_n - H_n(Y_i, \hat{\beta}_n, \hat{F}_n))} = \sum_{i=1}^n I(Y_i < \infty) \frac{\Delta_i}{n|\hat{\lambda}_n - H_n(Y_i, \hat{\beta}_n, \hat{F}_n)|} \\ &\geq \frac{1}{n} \sum_{i=1}^n I(Y_i < \infty) \frac{\Delta_i}{|\hat{\lambda}_n - H_n(Y_i, \hat{\beta}_n, \hat{F}_n)| + \epsilon}, \end{aligned} \quad (\text{A.1.2})$$

for any positive constant ϵ . Since $H_n(y, \hat{\beta}_n, \hat{F}_n)$ converges uniformly to $H^*(y)$,

$$\frac{1}{n} \sum_{i=1}^n I(Y_i < \infty) \frac{\Delta_i}{|\hat{\lambda}_n - H_n(Y_i, \hat{\beta}_n, \hat{F}_n)| + \epsilon} - \frac{1}{n} \sum_{i=1}^n I(Y_i < \infty) \frac{\Delta_i}{|\lambda^* - H^*(Y_i)| + \epsilon} \rightarrow 0.$$

Then after taking limits on both sides, we obtain $1 \geq E\{\Delta I(Y < \infty)/(|\lambda^* - H^*(Y)| + \epsilon)\}$. Letting $\epsilon \rightarrow 0$, by the monotone convergence theorem, we have

$$1 \geq \int_0^\infty \frac{c_0 dy}{|\lambda^* - H^*(y)|}, \quad (\text{A.1.3})$$

where c_0 is a positive constant. Thus, if $\inf_y |\lambda^* - H^*(y)| = 0$, we claim that there exists a finite y_0 such that $H^*(y_0) = \lambda^*$. Otherwise, $H^*(\infty) = \lambda^* = 0$. Then, for large y , $|\lambda^* - H^*(y)| < 1$ which makes (A.1.3) impossible. Now suppose that there exists a finite y_0 such that $\lambda^* = H^*(y_0)$. Then (A.1.3) becomes $1 \geq c_0 \int_0^\infty dy/|H^*(y_0) - H^*(y)|$. This is impossible since $H^*(y)$ is continuously differentiable in a neighborhood of y_0 . Therefore, there exists a positive constant δ^* such that $|\lambda^* - H^*(y)| > \delta^*$. This implies that when n is large, $|\hat{\lambda}_n - H_n(y, \hat{\beta}_n, \hat{F}_n)| > \delta^*$. Note that $\hat{F}_n(y) = n^{-1} \sum_{i=1}^n \Delta_i I(Y_i \leq y)/|\hat{\lambda}_n - H_n(Y_i, \hat{\beta}_n, \hat{F}_n)|$ so $\hat{F}_n(y)$ converges uniformly to $F^*(y) = E\{\Delta I(Y \leq y)/|\lambda^* - H^*(Y)|\}$.

We now show that $\beta^* = \beta_0, F^* = F_0$. To do so, we construct another function \tilde{F} which has jumps only at Y_i such that $\Delta_i = 1$ and $Y_i < \infty$. Moreover,

$$\tilde{F}_n\{Y_i\} = \frac{1}{nc_n} \frac{\Delta_i}{\tilde{\lambda}_n - H_n(Y_i, \beta_0, F_0)},$$

where $\tilde{\lambda}_n$ satisfies a similar equation to (A.1.1) and it is given by

$$\tilde{\lambda}_n = \frac{1}{n} \sum_{i=1}^n \Delta_i I(Y_i < \infty) + \int_0^\infty H_n(y, \beta_0, F_0) dF_0(y)$$

and c_n is a constant such that $\sum_{i=1}^n \tilde{F}_n\{Y_i\} = 1$. Furthermore, using the argument of the Glivenko-Cantelli property as before, we can easily show that uniformly in y , $H_n(y, \beta_0, F_0)$ converges to

$$\tilde{H}(y) = E \left\{ \Delta \frac{G''(\eta(\beta_0^T \mathbf{X}) F_0(Y)) \eta(\beta_0^T \mathbf{X}) I(\infty > Y \geq y)}{G'(\eta(\beta_0^T \mathbf{X}) F_0(Y))} \right\}$$

$$+(1-\Delta)\frac{G'(\eta(\beta_0^T \mathbf{X})F_0(Y))\eta(\beta_0^T \mathbf{X})I(\infty > Y \geq y)}{G(\eta(\beta_0^T \mathbf{X})F_0(Y))}\Big\},$$

which, after integration by part, is equal to $E[\eta(\beta_0^T \mathbf{X})G'(\eta(\beta_0^T \mathbf{X})F_0(y))S_c(y|\mathbf{X})]$, where S_c is the conditional survival function of the censoring time. Consequently, direct calculation gives that $\tilde{\lambda}_n$ converges to 0. Furthermore, from equation

$$c_n \tilde{F}_n(y) = \frac{1}{n} \sum_{i=1}^n \frac{\Delta_i I(Y_i \leq y)}{\tilde{\lambda}_n - H_n(Y_i, \beta_0, F_0)},$$

we obtain that uniformly in y , $c_n \tilde{F}_n(y)$ converges to

$$E \left[\frac{\Delta I(Y \leq y)}{-E \{S_c(\tilde{y}|\mathbf{X})\eta(\beta_0^T \mathbf{X})G'(\eta(\beta_0^T \mathbf{X})F_0(\tilde{y}))\} |_{\tilde{y}=Y}} \right] = F_0(y).$$

Hence, $c_n \rightarrow 1$ and $\tilde{F}_n(y)$ converges to $F_0(y)$ uniformly.

Note that \hat{F}_n is absolutely continuous with respect to $\tilde{F}_n(y)$ with

$$\hat{F}_n(y) = \int_0^y \frac{|\tilde{\lambda}_n - \tilde{H}_n(t, \beta_0, F_0)|}{|\hat{\lambda}_n - H_n(t, \hat{\beta}_n, \hat{F}_n)|} d\tilde{F}_n(t). \quad (\text{A.1.4})$$

From the forgoing arguments, the integrand in (A.1.4) is bounded and uniformly converges to $|\tilde{H}(t)|/|\lambda^* - H^*(t)|$. We conclude that $F^*(y) = \int_0^y |\tilde{H}(t)|dF_0(t)/|\lambda^* - H^*(t)|$. This implies that F^* is absolutely continuous with respect to F_0 . Therefore, F^* is also differentiable and we denote its density function by f^* .

On the other hand, since the observed log-likelihood function at $(\hat{\beta}_n, \hat{F}_n)$ is larger than or equal to the observed log-likelihood function at (β_0, \tilde{F}_n) , we have

$$\begin{aligned} & \frac{1}{n} \sum_{i=1}^n I(Y_i < \infty) \Delta_i \log \frac{\hat{F}_n\{Y_i\}}{\tilde{F}_n\{Y_i\}} + \frac{1}{n} \sum_{i=1}^n \left\{ I(Y_i = \infty) \log \frac{G(\eta(\hat{\beta}_n^T \mathbf{X}_i))}{G(\eta(\beta_0^T \mathbf{X}_i))} \right\} \\ & + \frac{1}{n} \sum_{i=1}^n I(Y_i < \infty) \left\{ \Delta_i \log \frac{G'(\eta(\hat{\beta}_n^T \mathbf{X}_i)\hat{F}_n(Y_i))\eta(\hat{\beta}_n^T \mathbf{X}_i)}{G'(\eta(\beta_0^T \mathbf{X}_i)\tilde{F}_n(Y_i))\eta(\beta_0^T \mathbf{X}_i)} + (1 - \Delta_i) \log \frac{G(\eta(\hat{\beta}_n^T \mathbf{X}_i)\hat{F}_n(Y_i))}{G(\eta(\beta_0^T \mathbf{X}_i)\tilde{F}_n(Y_i))} \right\} \geq 0. \end{aligned}$$

We take limits on both sides and note that

$$\frac{1}{n} \sum_{i=1}^n \Delta_i I(Y_i < \infty) \log \frac{\hat{F}_n\{Y_i\}}{\tilde{F}_n\{Y_i\}} \rightarrow E \left\{ \Delta I(Y < \infty) \log \frac{f^*(Y)}{f_0(Y)} \right\}.$$

We obtain $-K((\beta^*, F^*), (\beta_0, F_0)) \geq 0$, where $K(\cdot, \cdot)$ denotes the Kullback-Leibler information of (β^*, F^*) with respect to the true parameters. Immediately, we obtain

$$\begin{aligned} & \left\{ -G'(\eta(\beta^{*T} \mathbf{X})F^*(Y))\eta(\beta^{*T} \mathbf{X})f^*(Y) \right\}^{\Delta I(Y < \infty)} \left\{ G(\eta(\beta^{*T} \mathbf{X})F^*(Y)) \right\}^{(1-\Delta)I(Y < \infty) + I(Y = \infty)} \\ & = \left\{ -G'(\eta(\beta_0^T \mathbf{X})F_0(Y))\eta(\beta_0^T \mathbf{X})f_0(Y) \right\}^{\Delta I(Y < \infty)} \left\{ G(\eta(\beta_0^T \mathbf{X})F_0(Y)) \right\}^{(1-\Delta)I(Y < \infty) + I(Y = \infty)} \quad (\text{A.1.5}) \end{aligned}$$

for almost every (Δ, X, Y) in its support. According to the second paragraph in Section 3, we obtain $\beta^* = \beta_0$ and $F^* = F_0$.

We have shown that for almost every sample in the probability space, we can always choose a subsequence of $(\hat{\beta}_n, \hat{F}_n)$ so that it converges to (β_0, F_0) . Hence, with probability one, $\hat{\beta}_n \rightarrow \beta_0$ and $\hat{F}_n(y) \rightarrow F_0(y)$ for every $y \in [0, \infty)$. Particularly, we obtain $\sup_y |\hat{F}_n(y) - F_0(y)| \rightarrow 0$ due to the continuity of F_0 .

Remark A.1 When transformation G depends on some unknown parameter γ , where γ belongs to a compact set Γ , the proof of the consistency applies when assumptions (C1) and (C3) are replaced by (C1'). Parameters (β_0, γ_0, F_0) are identifiable.

(C3'). $G_\gamma(x)$ is three times differentiable with respect to γ and x and all the derivatives are uniformly bounded with $G'_\gamma(x) > 0$.

Especially, (C3') ensures the classes of random functions in the above proof to be the Glivenko-Cantelli classes while (C1') ensures the limit of $(\hat{\beta}_n, \hat{\gamma}_n, \hat{F}_n)$ must be the true parameters.

A.2 Proof of Theorem 2

To prove the asymptotic properties of $(\hat{\beta}_n, \hat{F}_n)$, we recall the definition of \mathcal{H} in Section 3. Furthermore, we abbreviate $l(\beta, F)$ as the log-likelihood function of (5), given by

$$\begin{aligned} l(\beta, F) &= I(Y < \infty) [\Delta \log f + \Delta \log \{-G'(\eta(\beta^T \mathbf{X})F(Y))\eta(\beta^T \mathbf{X})\} \\ &\quad + (1 - \Delta) \log G(\eta(\beta^T \mathbf{X})F(Y))] + I(Y = \infty) \log G(\eta(\beta^T \mathbf{X})). \end{aligned}$$

Denote $l_\beta(\beta, F)$ as the derivative of $l(\beta, F)$ with respect to β and denote $l_F(\beta, F)[\int (h_2 - Q_F[h_2])dF]$ as the derivative of $l(\beta, F)$ along the path $(\beta, F_\epsilon = F + \epsilon \int Q_F(h_2)dF), \epsilon \in (-\epsilon_0, \epsilon_0)$ for a small constant ϵ_0 , where $Q_F[h_2] = h_2(t) - \int_0^\infty h_2(t)dF(t)$. Additionally, we can define the derivative of $l_\beta(\beta, F)$ with respect to β , denoted by $l_{\beta\beta}(\beta, F)$, the derivative of $l_\beta(\beta, F)$ with respect to F along the path $F + \epsilon(\hat{F}_n - F)$, denoted by $l_{\beta F}[\hat{F}_n - F]$, and the derivative of $l_F(\beta, F)[\int Q_F(h_2)dF]$ with respect to β , denoted by $l_{F\beta}(\beta, F)[\int Q_F(h_2)dF]$, the derivative $l_F(\beta, F)[\int Q_F(h_2)dF]$ with respect to F along the path $F + \epsilon(\hat{F}_n - F)$, denoted by $l_{FF}(\beta, F)[\int Q_F(h_2)dF, \hat{F}_n - F]$. Furthermore, we define

$$\begin{aligned} \Psi_1(\Delta, Y, \mathbf{X}) &= I(Y < \infty) \Delta \left\{ \frac{G^{(3)}(\eta(\beta^T \mathbf{X})F(Y))}{G'(\eta(\beta^T \mathbf{X})F(Y))} - \frac{G''(\eta(\beta^T \mathbf{X})F(Y))^2}{G'(\eta(\beta^T \mathbf{X})F(Y))^2} \right\} \\ &\quad + \left\{ (1 - \Delta)I(Y < \infty) + I(Y = \infty) \right\} \frac{G''(\eta(\beta^T \mathbf{X})F(Y))}{G(\eta(\beta^T \mathbf{X})F(Y))} \\ &\quad - \left\{ (1 - \Delta)I(Y < \infty) + I(Y = \infty) \right\} \frac{G'(\eta(\beta^T \mathbf{X})F(Y))^2}{G(\eta(\beta^T \mathbf{X})F(Y))^2}, \end{aligned}$$

$$\begin{aligned}\Psi_2(\Delta, Y, \mathbf{X}) &= I(Y < \infty) \Delta \frac{G''(\eta(\beta^T \mathbf{X})F(Y))}{G'(\eta(\beta^T \mathbf{X})F(Y))} \\ &\quad + \left\{ (1 - \Delta)I(Y < \infty) + I(Y = \infty) \right\} \frac{G'(\eta(\beta^T \mathbf{X})F(Y))}{G(\eta(\beta^T \mathbf{X})F(Y))}.\end{aligned}$$

Since $(\hat{\beta}_n, \hat{F}_n)$ maximizes $\mathbf{P}_n l(\beta, F)$, for any $(\mathbf{h}_1, h_2) \in \mathcal{H}$, it follows that

$$\mathbf{P}_n \left\{ l_{\beta}(\hat{\beta}_n, \hat{F}_n)^T \mathbf{h}_1 + l_F(\hat{\beta}_n, \hat{F}_n) \left[\int Q_{\hat{F}_n}(h_2) d\hat{F}_n \right] \right\} = 0.$$

Note that $\mathbf{P} \left\{ l_{\beta}(\beta_0, F_0)^T \mathbf{h}_1 + l_F(\beta_0, F_0) \left[\int Q_{F_0}(h_2) dF_0 \right] \right\} = 0$. Thus, we obtain

$$\begin{aligned}& \sqrt{n}(\mathbf{P}_n - \mathbf{P}) \left\{ l_{\beta}(\hat{\beta}_n, \hat{F}_n)^T \mathbf{h}_1 + l_F(\hat{\beta}_n, \hat{F}_n) \left[\int Q_{\hat{F}_n}(h_2) d\hat{F}_n \right] \right\} \\ &= -\sqrt{n}\mathbf{P} \left\{ l_{\beta}(\hat{\beta}_n, \hat{F}_n)^T \mathbf{h}_1 + l_F(\hat{\beta}_n, \hat{F}_n) \left[\int Q_{\hat{F}_n}(h_2) d\hat{F}_n \right] \right\} \\ &\quad + \sqrt{n}\mathbf{P} \left\{ l_{\beta}(\beta_0, F_0)^T \mathbf{h}_1 + l_F(\beta_0, F_0) \left[\int Q_{F_0}(h_2) dF_0 \right] \right\}.\end{aligned}\tag{A.2.1}$$

First, by the same arguments in the consistency proof, the classes of

$$\begin{aligned}\mathcal{A}_2 &= \left\{ \frac{G'(x)}{G(x)}, \frac{G''(x)}{G'(x)} \Big|_{x=\eta(\beta^T \mathbf{X})F(Y)} : \|\beta - \beta_0\| < \delta_0, \sup_y |F(y) - F_0(y)| < \delta_0 \right\}, \\ \mathcal{A}_3 &= \left\{ \eta'(\beta^T \mathbf{X})F(Y), \eta(\beta^T \mathbf{X})F(Y) : \|\beta - \beta_0\| < \delta_0, \sup_y |F(y) - F_0(y)| < \delta_0 \right\}\end{aligned}$$

are P-Donsker. Additionally, it is clear to see both classes $\{Q_F(h_2) : \|h_2\|_V \leq 1, \sup_y |F(y) - F_0(y)| < \delta_0\}$ and $\left\{ \int_0^Y Q_F(h_2) dF : \|h_2\|_V \leq 1, \sup_y |F(y) - F_0(y)| < \delta_0 \right\}$ contain the functions of Y with bounded variations so they are also P-Donsker. Therefore, from the explicit expression of l_{β} and l_F , the preservation of the Donsker classes under algebraic operations implies that the class

$$\mathcal{A}_4 = \left\{ l_{\beta}(\beta, F)^T \mathbf{h}_1 + l_F(\beta, F) \left[\int Q_F(h_2) dF \right] : \|\mathbf{h}_1\| \leq 1, \|h_2\|_V \leq 1, \|\beta - \beta_0\| + \sup_y |F(y) - F_0(y)| < \delta_0 \right\}$$

is P-Donsker. On the other hand, it is straightforward to show that

$$l_{\beta}(\hat{\beta}_n, \hat{F}_n)^T \mathbf{h}_1 + l_F(\hat{\beta}_n, \hat{F}_n) \left[\int Q_{\hat{F}_n}(h_2) d\hat{F}_n \right] \rightarrow l_{\beta}(\beta_0, F_0)^T \mathbf{h}_1 + l_F(\beta_0, F_0) \left[\int Q_{F_0}(h_2) dF_0 \right]$$

uniformly in $(\mathbf{h}_1, h_2) \in \mathcal{H}$. Thus, the left-hand side of (A.2.1) is equal to

$$\sqrt{n}(\mathbf{P}_n - \mathbf{P}) \left\{ l_{\beta}(\beta_0, F_0)^T \mathbf{h}_1 + l_F(\beta_0, F_0) \left[\int Q_{F_0}(h_2) dF_0 \right] \right\} + o_p(1),$$

where $o_p(1)$ is a random variable that converges to zero in probability in the metric space $l^\infty(\mathcal{H})$. As a result, the left-hand side of (A.2.1) converges weakly to a zero-mean Gaussian process in $l^\infty(\mathcal{H})$.

Second, simple algebra shows that uniformly in $(\mathbf{h}_1, h_2) \in \mathcal{H}$,

$$\begin{aligned}
& \left| l_{\boldsymbol{\beta}}(\hat{\boldsymbol{\beta}}_n, \hat{F}_n)^T \mathbf{h}_1 + l_F(\hat{\boldsymbol{\beta}}_n, \hat{F}_n) \left[\int Q_{\hat{F}_n}(h_2) d\hat{F}_n \right] - l_{\boldsymbol{\beta}}(\boldsymbol{\beta}_0, F_0)^T \mathbf{h}_1 - l_F(\boldsymbol{\beta}_0, F_0) \left[\int Q_{F_0}(h_2) dF_0 \right] \right. \\
& \quad - \left\{ (\hat{\boldsymbol{\beta}}_n - \boldsymbol{\beta}_0)^T l_{\boldsymbol{\beta}\boldsymbol{\beta}}(\boldsymbol{\beta}_0, F_0) \mathbf{h}_1 + (\hat{\boldsymbol{\beta}}_n - \boldsymbol{\beta}_0)^T l_{F\boldsymbol{\beta}}(\boldsymbol{\beta}_0, F_0) \left[\int Q_{F_0}(h_2) dF_0 \right] \right. \\
& \quad \left. \left. + \mathbf{h}_1^T l_{\boldsymbol{\beta}F}[\hat{F}_n - F_0] + l_{FF}(\boldsymbol{\beta}_0, F_0) \left[\int Q_{F_0}(h_2) dF_0, \hat{F}_n - F_0 \right] \right\} \right| \\
& \leq o_p \left\{ \|\hat{\boldsymbol{\beta}}_n - \boldsymbol{\beta}_0\| + \|\hat{F}_n - F_0\|_{l^\infty} \right\}.
\end{aligned}$$

Thus, combining with the expressions of $l_{\boldsymbol{\beta}\boldsymbol{\beta}}, l_{\boldsymbol{\beta}F}, l_{F\boldsymbol{\beta}}$ and l_{FF} , we obtain that the right-hand side of (A.2.1) equals

$$\begin{aligned}
& -\sqrt{n} \left\{ (\hat{\boldsymbol{\beta}}_n - \boldsymbol{\beta}_0)^T \Omega_{\boldsymbol{\beta}}(\mathbf{h}_1, Q_{F_0}(h_2)) + \int_0^\infty \Omega_F(\mathbf{h}_1, Q_{F_0}(h_2)) d(\hat{F}_n - F_0)(y) \right\} \\
& \quad + o \left\{ \sqrt{n} (\|\hat{\boldsymbol{\beta}}_n - \boldsymbol{\beta}_0\| + \|\hat{F}_n - F_0\|_{l^\infty}) \right\},
\end{aligned}$$

where

$$\begin{aligned}
\Omega_{\boldsymbol{\beta}}(\mathbf{h}_1, Q_{F_0}(h_2)) &= E \left[I(Y < \infty) \Delta \frac{\eta''(\boldsymbol{\beta}^T \mathbf{X}) \eta(\boldsymbol{\beta}^T \mathbf{X}) - \eta'(\boldsymbol{\beta}^T \mathbf{X})^2}{\eta(\boldsymbol{\beta}^T \mathbf{X})^2} \mathbf{X} \mathbf{X}^T \mathbf{h}_1 \right] \\
& \quad + E \left[\{ \Psi_1^0(\Delta, Y, \mathbf{X}) \eta'(\boldsymbol{\beta}_0^T \mathbf{X})^2 F_0(Y)^2 + \Psi_2^0(\Delta, Y, \mathbf{X}) \eta''(\boldsymbol{\beta}_0^T \mathbf{X}) F_0(Y) \} \mathbf{X} \mathbf{X}^T \mathbf{h}_1 \right] \\
& \quad + E \left[\{ \Psi_1^0(\Delta, Y, \mathbf{X}) \eta(\boldsymbol{\beta}_0^T \mathbf{X}) \eta'(\boldsymbol{\beta}_0^T \mathbf{X}) F_0(Y) + \Psi_2^0(\Delta, Y, \mathbf{X}) \eta'(\boldsymbol{\beta}_0^T \mathbf{X}) F_0(Y) \} \mathbf{X} \right. \\
& \quad \left. \times \int_0^Y Q_{F_0}(h_2) dF_0 \right], \\
\Omega_F(\mathbf{h}_1, Q_{F_0}(h_2)) &= -E \left[I(Y < \infty) \Delta + \Psi_2^0(\Delta, Y, \mathbf{X}) \eta(\boldsymbol{\beta}_0^T \mathbf{X}) \{ F_0(Y) - I(Y \geq y) \} \right] Q_{F_0}[h_2] \\
& \quad + E \left[\{ \Psi_1^0(\Delta, Y, \mathbf{X}) \eta(\boldsymbol{\beta}_0^T \mathbf{X}) \eta'(\boldsymbol{\beta}_0^T \mathbf{X}) F_0(Y) + \Psi_2^0(\Delta, Y, \mathbf{X}) \eta'(\boldsymbol{\beta}_0^T \mathbf{X}) F_0(Y) \} \right. \\
& \quad \left. \times \mathbf{X}^T \mathbf{h}_1 I(Y \geq y) \right] \\
& \quad + E \left[I(Y \geq y) \Psi_1^0(\Delta, Y, \mathbf{X}) \eta(\boldsymbol{\beta}_0^T \mathbf{X})^2 \int_0^Y Q_{F_0}(h_2) dF_0 \right],
\end{aligned}$$

where Ψ_1^0 and Ψ_2^0 have the same expressions as Ψ_1 and Ψ_2 respectively but with $\boldsymbol{\beta}$ and F replaced by $\boldsymbol{\beta}_0$ and F_0 .

Third, the linear operator $(\Omega_{\boldsymbol{\beta}}, \Omega_F)$ is a bounded linear operator from the linear space

$$\mathcal{S} = R^d \times \left\{ \tilde{h}_2 : \|\tilde{h}_2\|_V < \infty, \int_0^\infty \tilde{h}_2(y) dF_0(y) = 0 \right\}$$

to itself. We wish to show that $(\Omega_{\boldsymbol{\beta}}, \Omega_F)$ is invertible. From the direct calculation, we have

$$-E \left[I(Y < \infty) \Delta + \Psi_2^0(\Delta, Y, \mathbf{X}) \eta(\boldsymbol{\beta}_0^T \mathbf{X}) \{ F_0(Y) - I(Y \geq y) \} \right] = E \left[G'(\eta(\boldsymbol{\beta}_0^T \mathbf{X}) F_0(y)) \eta(\boldsymbol{\beta}_0^T \mathbf{X}) S_c(y|\mathbf{X}) \right],$$

which is negative. Thus, $(\Omega_{\beta}, \Omega_F)$ can be written as the summation of an invertible operator and a compact operator. By Rudin (1973), to prove the invertibility of $(\Omega_{\beta}, \Omega_F)$, it is sufficient to show that $(\Omega_{\beta}, \Omega_F)$ is one-to-one. That is, if there exists some $(\mathbf{h}_1, \tilde{h}_2) \in \mathcal{S}$ such that $\Omega_{\beta}(\mathbf{h}_1, \tilde{h}_2) = 0$ and $\Omega_F(\mathbf{h}_1, \tilde{h}_2) = 0$, we need to show $\mathbf{h}_1 = 0$ and $\tilde{h}_2 = 0$. However, we notice that according to the derivation of Ω 's, it holds that

$$\mathbf{h}_1^T \Omega_{\beta}(\mathbf{h}_1, \tilde{h}_2) + \int_0^{\infty} \Omega_{\beta}(\mathbf{h}_1, \tilde{h}_2) \tilde{h}_2 dF_0 = -E \left\{ l_{\beta}(\beta_0, F_0)^T \mathbf{h}_1 + l_F(\beta_0, F_0)[\tilde{h}_2] \right\}^2.$$

We thus obtain that with probability one,

$$l_{\beta}(\beta_0, F_0)^T \mathbf{h}_1 + l_F(\beta_0, F_0)[\tilde{h}_2] = 0.$$

Particularly, we choose $Y = \infty$ and obtain $\mathbf{h}_1 = 0$; then we let $Y < \infty$ and $\Delta = 1$ and obtain a homogeneous integral equation for \tilde{h}_2 . Such an equation has one trivial solution $\tilde{h}_2 = 0$.

Finally, using the inverse of $(\Omega_{\beta}, \Omega_F)$, denoted by $(\tilde{\Omega}_{\beta}, \tilde{\Omega}_F)$, the equation (A.2.1) can be written as

$$\begin{aligned} & \sqrt{n} \left\{ (\hat{\beta}_n - \beta_0)^T \mathbf{h}_1 + \int_0^{\infty} \tilde{h}_2 d(\hat{F}_n - F_0) \right\} \\ = & -\sqrt{n}(\mathbf{P}_n - \mathbf{P}) \left\{ l_{\beta}(\beta_0, F_0)^T \tilde{\Omega}_{\beta}(\mathbf{h}_1, \tilde{h}_2) + l_F(\beta_0, F_0)^T \tilde{\Omega}_F(\mathbf{h}_1, \tilde{h}_2) \right\} \\ & + o_p(1) \left\{ \sqrt{n}(\|\hat{\beta}_n - \beta_0\| + \|\hat{F}_n - F_0\|_{l^{\infty}}) \right\}, \end{aligned}$$

where $o_p(1)$ converges to zero in probability uniformly in $(\mathbf{h}_1, \tilde{h}_2) \in \mathcal{S}_0$, where \mathcal{S}_0 contains all $(\mathbf{h}_1, \tilde{h}_2) \in \mathcal{S}$ such that $\|\mathbf{h}_1\| \leq 1$ and $\|\tilde{h}_2\|_V \leq 1$. This immediately implies that

$$\sqrt{n}(\|\hat{\beta}_n - \beta_0\| + \|\hat{F}_n - F_0\|_{l^{\infty}}) = O_p(1).$$

Hence,

$$\begin{aligned} & \sqrt{n} \left\{ (\hat{\beta}_n - \beta_0)^T \mathbf{h}_1 + \int_0^{\infty} \tilde{h}_2 d(\hat{F}_n - F_0) \right\} \\ = & -\sqrt{n}(\mathbf{P}_n - \mathbf{P}) \left\{ l_{\beta}(\beta_0, F_0)^T \tilde{\Omega}_{\beta}(\mathbf{h}_1, \tilde{h}_2) + l_F(\beta_0, F_0)^T \tilde{\Omega}_F(\mathbf{h}_1, \tilde{h}_2) \right\} + o_p(1). \end{aligned} \quad (\text{A.2.2})$$

Then $\sqrt{n} \left\{ (\hat{\beta}_n - \beta_0)^T \mathbf{h}_1 + \int_0^{\infty} \tilde{h}_2 d(\hat{F}_n - F_0) \right\}$ converges weakly to a Gaussian process, denoted by $GP(\mathbf{h}_1, \tilde{h}_2)$. The covariance between $GP(\mathbf{h}_1, \tilde{h}_2)$ and $GP(\mathbf{h}_1^*, \tilde{h}_2^*)$ is given by

$$\begin{aligned} & E \left[\left\{ l_{\beta}(\beta_0, F_0)^T \tilde{\Omega}_{\beta}(\mathbf{h}_1, \tilde{h}_2) + l_F(\beta_0, F_0)[\tilde{\Omega}_F(\mathbf{h}_1, \tilde{h}_2)] \right\} \right. \\ & \quad \left. \times \left\{ l_{\beta}(\beta_0, F_0)^T \tilde{\Omega}_{\beta}(\mathbf{h}_1^*, \tilde{h}_2^*) + l_F(\beta_0, F_0)[\tilde{\Omega}_F(\mathbf{h}_1^*, \tilde{h}_2^*)] \right\} \right]. \end{aligned}$$

Since for any h_2 , $\int h_2 d(\hat{F}_n - F_0) = \int Q_{F_0}(h_2) d(\hat{F}_n - F_0)$, the above convergence result also implies the weak convergence result in Theorem 2.

Specifically, if we choose in equation (A.2.2) that $\tilde{h}_2 = 0$, then we conclude that $\hat{\beta}_n^T \mathbf{h}_1$ is an asymptotic linear estimator for $\beta_0^T \mathbf{h}_1$ with its influence function given by

$$l_{\beta}(\beta_0, F_0)^T \tilde{\Omega}_{\beta}(\mathbf{h}_1, 0) + l_F(\beta_0, F_0)[\tilde{\Omega}_F(\mathbf{h}_1, 0)].$$

This implies that $\hat{\beta}_n$ is semiparametrically efficient since the influence function is on the linear space spanned by the score functions for β_0 and F_0 .

Remark A.2.1. When the transformation depends on some parameter γ , the above proof can be easily adapted to this case by introducing one more parameter γ . The results hold if γ_0 is assumed to belong to the interior of Γ , (C1) and (C3) are replaced by assumptions (C1') and (C3'), and the following assumption also holds:

(C5') If with probability one,

$$G'_{\gamma}(\eta(\beta_0^T \mathbf{X}))\eta'(\beta_0^T \mathbf{X})\mathbf{X}^T \mathbf{h}_1 + \dot{G}_{\gamma}(\eta(\beta_0^T \mathbf{X}))h_3 = 0,$$

where \mathbf{h}_1 and h_3 are constant vectors and \dot{G}_{γ} denotes the derivative with respect to γ , then $\mathbf{h}_1 = \mathbf{0}$ and $h_3 = 0$.

Note that (C5') is particularly used for proving the invertibility of Ω 's.

Remark A.2.2. The profile likelihood function can be used to give a consistent estimate for the asymptotic variance of $\hat{\beta}_n$. Its justification follows from verifying all the conditions of Theorem 1 in Murphy and van der Vaart (2000). Especially, from the invertibility of Ω 's, we conclude that the information operator for (β_0, F_0) is invertible. Therefore, there exists a vector of functions \mathbf{h} with bounded variation such that $l_F^* l_F[\int Q_{F_0}(\mathbf{h})dF_0] = l_F^* l_{\beta}$, where l_F^* is the dual operator of l_F . The function $\int Q_{F_0}(\mathbf{h})dF_0$ is called the least favorable direction in Murphy and van der Vaart (2000). We then consider the submodel (ϵ, F_{ϵ}) where (ϵ, F_{ϵ}) , where $F_{\epsilon} = F + (\epsilon - \beta) \int Q_F(\mathbf{h})dF$ and $\epsilon \in R^d$. It is clear that such submodel satisfies conditions (8) and (9) in Murphy and van der Vaart (2000). Furthermore, for any $\tilde{\beta}_n$, we let \tilde{F}_n be the distribution function maximizing (6) in which $\beta = \tilde{\beta}_n$. From the proof of Theorem 1, the same arguments imply that \tilde{F}_n uniformly converges to F_0 with probability one. We thus verify condition (10) in Murphy and van der Vaart (2000). As in the proof of Theorem 2, we linearize the likelihood function for \tilde{F}_n , which is equal to

$$0 = \mathbf{P}_n \left\{ l_F(\tilde{\beta}_n, \tilde{F}_n) \left[\int Q_{\tilde{F}_n}(h_2)d\tilde{F}_n \right] \right\}.$$

Following the same expansion and using the P-Donsker property as in proving Theorem 2, we obtain

$$\sqrt{n} \int \Omega_F(0, Q_{F_0}(h_2))d(\tilde{F}_n - F_0) = \sqrt{n}(\mathbf{P}_n - \mathbf{P}) \left\{ l_F(\beta_0, F_0) \left[\int Q_{F_0}[h_2]dF_0 \right] \right\}$$

$$-\sqrt{n}\mathbf{P}\left[l_F(\tilde{\beta}_n, F_0)\left[\int Q_{F_0}[h_2]dF_0\right] - l_F(\beta_0, F_0)\left[\int Q_{F_0}[h_2]dF_0\right]\right] + o_p(1).$$

From the invertibility of $\Omega_F(0, \cdot)$ and noting that

$$\left|\mathbf{P}\left[l_F(\tilde{\beta}_n, F_0)\left[\int Q_{F_0}[h_2]dF_0\right] - l_F(\beta_0, F_0)\left[\int Q_{F_0}[h_2]dF_0\right]\right]\right| \leq O_p(\|\tilde{\beta}_n - \beta_0\|).$$

We obtain $\sqrt{n}\|\tilde{F}_n - F_0\|_{l^\infty} = O_p(\sqrt{n} + \sqrt{n}\|\tilde{\beta}_n - \beta_0\|)$. This immediately implies condition (11), i.e., the no-bias condition, in Murphy and van der Vaart (2000). Furthermore, by the same arguments as in proving Theorem 1, it is straightforward to check that the class

$$\left\{\frac{\partial}{\partial \epsilon}l(\epsilon, F\epsilon) : \|\epsilon - \beta_0\| + \|\beta - \beta_0\| + \|F - F_0\| < \delta_0\right\}$$

is P-Donsker and the class

$$\left\{\frac{\partial^2}{\partial \epsilon^2}l(\epsilon, F\epsilon) : \|\epsilon - \beta_0\| + \|\beta - \beta_0\| + \|F - F_0\| < \delta_0\right\}$$

is P-Glivenko-Cantelli. Thus, all the conditions in Theorem 1 of Murphy and van der Vaart (2000) hold so the results of Theorem 1 in Murphy and van der Vaart (2000) are true. One conclusion of this theorem shows the consistency of the variance estimator based on the profile likelihood function.

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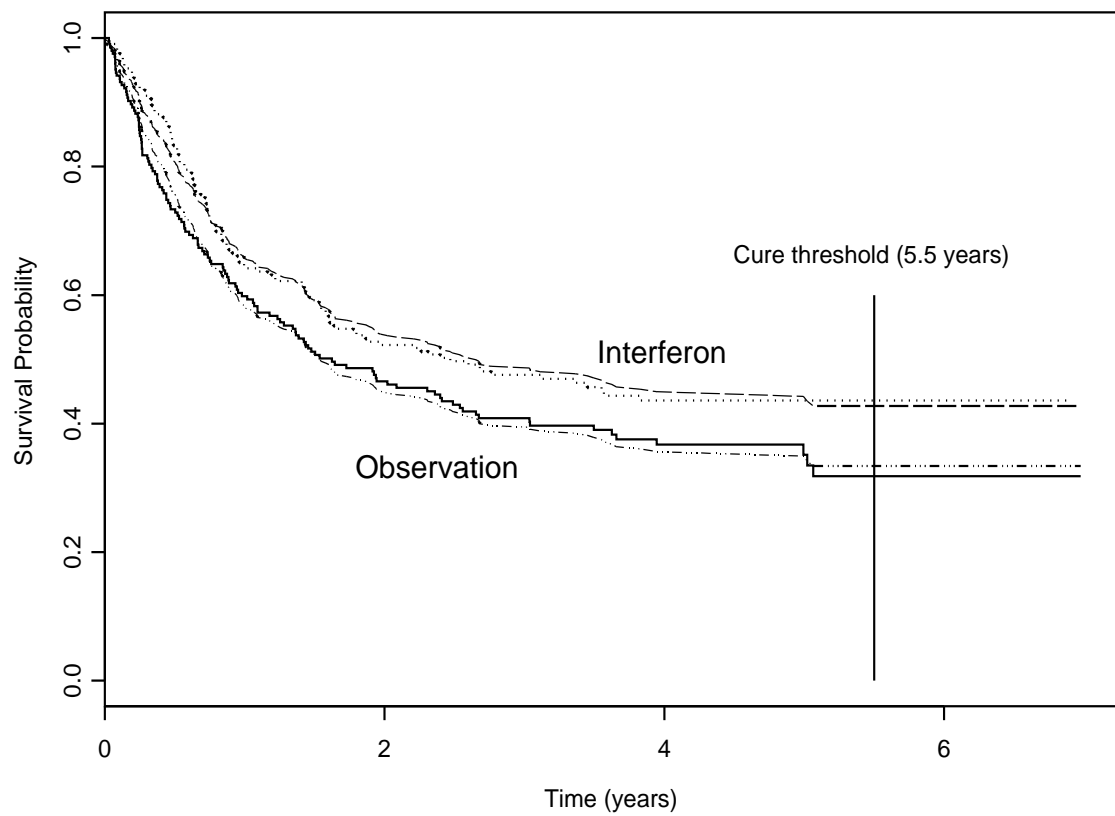


Figure 1: Kaplan-Meier curves and predicted survival curves of the interferon and observation groups in the E1690 data: the solid line and the dotted line are the Kaplan-Meier curves; the dashed line and the dot-dashed line are the predicted survival curves, respectively.

Table 1: Simulation results from 1000 replications under the transformation $G(x) = (1 + \gamma x)^{-1/\gamma}$

Model	n	Parameter	True Value	Estimate	SE	ESE	CP (%)
$\gamma = 0$	100	β_0	0.5	0.490	0.289	0.312	97.7
		β_1	1	1.033	0.433	0.427	94.9
		β_2	-0.5	-0.519	0.242	0.242	95.7
	200	β_0	0.5	0.502	0.200	0.218	96.1
		β_1	1	1.019	0.300	0.296	94.6
		β_2	-0.5	-0.509	0.167	0.168	95.9
$\gamma = 0.25$	100	β_0	0.5	0.476	0.341	0.350	95.6
		β_1	1	1.036	0.512	0.493	94.0
		β_2	-0.5	-0.490	0.280	0.281	96.0
	200	β_0	0.5	0.499	0.236	0.245	95.6
		β_1	1	1.006	0.356	0.344	95.1
		β_2	-0.5	-0.507	0.194	0.197	95.5
$\gamma = 0.5$	100	β_0	0.5	0.477	0.380	0.388	96.3
		β_1	1	1.022	0.550	0.554	95.4
		β_2	-0.5	-0.518	0.320	0.318	95.1
	200	β_0	0.5	0.488	0.271	0.273	95.5
		β_1	1	1.015	0.400	0.388	94.9
		β_2	-0.5	-0.505	0.225	0.222	95.1
$\gamma = 0.75$	100	β_0	0.5	0.487	0.410	0.423	95.7
		β_1	1	0.995	0.601	0.607	95.1
		β_2	-0.5	-0.491	0.359	0.348	94.2
	200	β_0	0.5	0.486	0.284	0.298	96.5
		β_1	1	1.022	0.426	0.425	94.7
		β_2	-0.5	-0.494	0.241	0.244	95.4
$\gamma = 1$	100	β_0	0.5	0.455	0.426	0.458	96.7
		β_1	1	1.043	0.637	0.658	96.1
		β_2	-0.5	-0.498	0.375	0.378	95.4
	200	β_0	0.5	0.482	0.310	0.321	95.4
		β_1	1	1.015	0.458	0.460	94.8
		β_2	-0.5	-0.502	0.258	0.264	95.8

Table 2: Simulation results from 1000 replications under the transformation $G(x) = \exp[-\{(1+x)^\gamma - 1\}/\gamma]$

Model	n	Parameter	True Value	Estimate	SE	ESE	CP (%)
$\gamma = 0$	100	β_0	0.5	0.465	0.442	0.458	96.6
		β_1	1	1.026	0.632	0.658	96.4
		β_2	-0.5	-0.510	0.387	0.378	94.8
	200	β_0	0.5	0.498	0.318	0.321	95.4
		β_1	1	0.995	0.474	0.461	93.9
		β_2	-0.5	-0.504	0.263	0.264	95.0
$\gamma = 0.25$	100	β_0	0.5	0.500	0.391	0.406	95.2
		β_1	1	0.994	0.568	0.585	96.3
		β_2	-0.5	-0.501	0.328	0.335	95.7
	200	β_0	0.5	0.489	0.283	0.285	94.8
		β_1	1	1.010	0.397	0.409	95.9
		β_2	-0.5	-0.502	0.237	0.235	94.7
$\gamma = 0.5$	100	β_0	0.5	0.459	0.356	0.364	95.8
		β_1	1	1.081	0.545	0.523	94.7
		β_2	-0.5	-0.500	0.297	0.299	95.8
	200	β_0	0.5	0.502	0.247	0.256	96.3
		β_1	1	1.005	0.360	0.365	95.4
		β_2	-0.5	-0.502	0.214	0.209	93.6
$\gamma = 0.75$	100	β_0	0.5	0.471	0.318	0.332	96.8
		β_1	1	1.069	0.479	0.469	93.9
		β_2	-0.5	-0.505	0.264	0.267	95.3
	200	β_0	0.5	0.506	0.228	0.233	95.8
		β_1	1	1.000	0.327	0.326	94.8
		β_2	-0.5	-0.500	0.192	0.187	94.2
$\gamma = 1$	100	β_0	0.5	0.509	0.289	0.314	97.8
		β_1	1	1.008	0.419	0.423	95.5
		β_2	-0.5	-0.516	0.245	0.242	94.2
	200	β_0	0.5	0.508	0.205	0.219	97.1
		β_1	1	1.010	0.296	0.296	95.2
		β_2	-0.5	-0.508	0.172	0.168	94.1

Table 3: Simulation results from 1000 replications under misspecified transformation ($n = 100$)

Model	Parameter	True Value	Estimate	SE	ESE	CP (%)
True Transformation: $G(x) = (1 + x/2)^{-2}$						
Proportional Hazards Model	β_0	0.5	0.165	0.290	0.311	82.6
	β_1	1	0.799	0.450	0.446	92.5
	β_2	-0.5	-0.404	0.248	0.255	94.2
Proportional Odds Model	β_0	0.5	0.818	0.456	0.466	90.8
	β_1	1	1.240	0.672	0.654	93.0
	β_2	-0.5	-0.578	0.375	0.373	95.1
True Transformation: $G(x) = \exp[-2\{(1 + x)^{1/2} - 1\}]$						
Proportional Hazards Model	β_0	0.5	0.189	0.304	0.311	84.1
	β_1	1	0.868	0.464	0.442	91.9
	β_2	-0.5	-0.411	0.254	0.252	92.7
Proportional Odds Model	β_0	0.5	0.960	0.463	0.472	84.4
	β_1	1	1.205	0.650	0.652	94.8
	β_2	-0.5	-0.606	0.363	0.373	95.0

Table 4: Estimates of regression coefficients in the proportional hazards cure model for the E1690 data

Cure Threshold	Covariate	Estimate	Std. Err.	<i>p</i> -value
5.1 years	Intercept	-0.7977	0.3147	0.0113
	Treatment	-0.2200	0.1298	0.0901
	Age	0.0115	0.0050	0.0220
	Sex	-0.2209	0.1371	0.1072
	Nodal category	0.5519	0.1599	0.0006
5.5 years	Intercept	-0.8027	0.3156	0.0110
	Treatment	-0.2197	0.1300	0.0911
	Age	0.0115	0.0050	0.0225
	Sex	-0.2208	0.1374	0.1081
	Nodal category	0.5520	0.1603	0.0006
6 years	Intercept	-0.7988	0.3151	0.0112
	Treatment	-0.2199	0.1298	0.0902
	Age	0.0115	0.0050	0.0220
	Sex	-0.2209	0.1372	0.1074
	Nodal category	0.5519	0.1600	0.0006
6.5 years	Intercept	-0.7969	0.3147	0.0113
	Treatment	-0.2200	0.1297	0.0898
	Age	0.0115	0.0050	0.0219
	Sex	-0.2210	0.1371	0.1070
	Nodal category	0.5518	0.1599	0.0006
7 years	Intercept	-0.7972	0.3148	0.0113
	Treatment	-0.2200	0.1297	0.0898
	Age	0.0115	0.0050	0.0219
	Sex	-0.2209	0.1371	0.1071
	Nodal category	0.5518	0.1599	0.0006

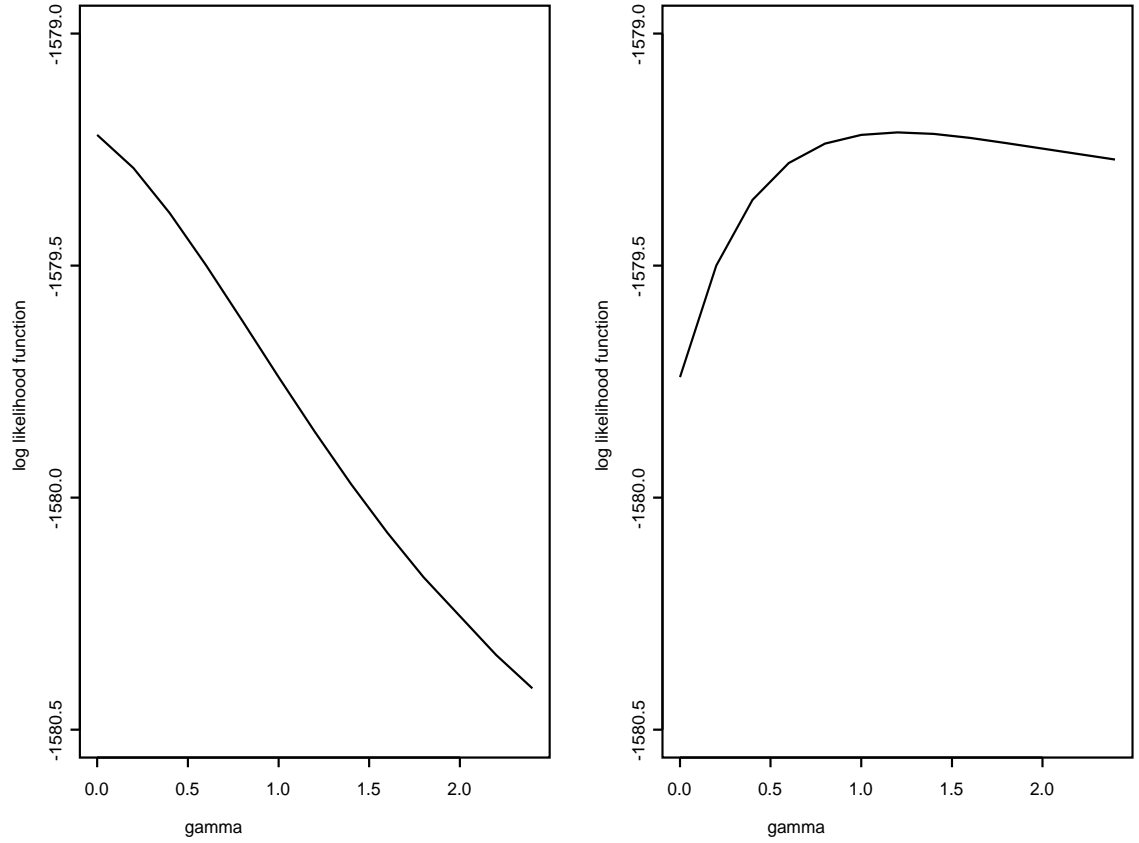


Figure 2: The observed log-likelihood functions from different transformations in the E1690 data: the left plot is the log-likelihood functions from transformations $G(x) = (1 + \gamma x)^{-1/\gamma}$; the right plot is the log-likelihood functions from transformations $G(x) = \exp\{-((1 + x)^\gamma - 1)/\gamma\}$.