

Discount Survival Models for No Covariate Case

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Abstract

For the survival data analysis of no covariate the discount survival model is proposed to estimate the time-varying hazard rate and the survival function recursively. In comparison with the covariate case it provide the distributionally explicit evolution of hazard rate between time intervals under the assumption of a conjugate gamma distribution. Also forecasting of the hazard rate in the next time interval is suggested, which leads to the forecasted survival function.

1. Introduction

West, Harrison and Migon(1985) developed the dynamic generalized linear model which allows the use of one dimensional exponential family distribution. The prior distribution is chosen to be a conjugate family member having same first two moments as the linearized function of a parameter vector. Gamerman(1991) proposed the dynamic Bayesian model for the analysis of survival data with the hazard rate which is a function of a time-varying parameter vector modeling effects of covariates. It is assumed that the time-variation of a parameter vector is determined through the evolution of a parameter vector between time intervals in the linear pattern, where its distribution is partially specified in terms of the mean vector and the variance-covariance matrix. Ameen and Harrison(1985) developed the normal discount Bayesian model to overcome some practical disadvantages of dynamic linear model, where the discount factor d_i ($0 < d_i < 1$) is introduced for an increase of $100(1-d_i)/d_i$ percent in the variance of the time interval I_i . Bolstad(1995) proposed the multiprocess dynamic Poisson model for the no covariate case by applying the multiprocess dynamic approach to the time-varying Poisson parameter.

In this article, under the assumption of a conjugate gamma distribution of the hazard rate in each time interval, the time-variation of the hazard rate between time intervals is determined through the multiplication of a gamma parameters by discount factors.

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The recursive estimation of the hazard rate and the survival function based on the information gathered until the end of each time interval is provided in Section 2. Forecasting of the hazard rate and the survival function in the next time interval is provided in Section 3. The performance of the discount survival model for no covariate case is illustrated via the gastric carcinoma data in Section 4.

2. Model descriptions

Here we assume that the survival time has a piecewise exponential distribution which has the constant hazard rate in each time interval, $\lambda(t) = \lambda_i$ for $t \in I_i = (\tau_{i-1}, \tau_i]$, where τ_0 is usually set to 0 and $I_s = (\tau_{s-1}, \infty)$. The survival function and the probability function for the current time interval I_i given survival up to the end of previous time interval can be easily calculated due to the lack of memory property of exponential distributions. Thus we obtain the likelihood for λ_i under the assumption that the random censoring time has no relation with the survival time,

$$l(\lambda_i | t) = \lambda_i^\delta \exp(-\lambda_i(t - \tau_{i-1})) \quad \text{for } t \in I_i$$

where δ is the indicator function of a death of the individual in the time interval I_i . We denote the hazard rate for the time interval I_i by λ_i and the number of individuals alive at the beginning of the time interval I_i by n_i . Let T_{ij} be the survival time of the j -th individual alive at the beginning of the time interval I_i . Also let t_{ij} be an observed failure time corresponding to T_{ij} , which can be interpreted as the minimum of the survival time of the j -th individual and the corresponding censoring time. Let D_i be a set of the information from all observations of each time interval, I_1, \dots, I_i , and let $D_{i-1(j)}$ be a set containing D_{i-1} and information from first j observations of the time interval I_i so that $D_{i-1(0)} = D_{i-1}$ and $D_{i-1(n_i)} = D_i$.

Then the discount survival model for the no covariate case in the time interval I_i can be defined as follows.

i) observation equation ;

$$T_{ij} \sim \exp(\lambda_i) \quad \text{for } i=1, \dots, s.$$

ii) evolution equation;

$$(\lambda_i | D_{i-1}) \sim \text{Ga}(d_i b_{i-1}, d_i r_{i-1}),$$

where b_{i-1} and r_{i-1} are parameters of the posterior distribution of λ_{i-1} given D_{i-1} , d_i ($0 < d_i < 1$) is the discount factor of the time interval I_i .

3. Inference Procedures

The process is started with the initial distribution of λ_0 , $Ga(b_0, r_0)$, where b_0 and r_0 are given prior to the time interval I_1 . At the end of each time interval I_{i-1} the posterior distribution of λ_{i-1} is obtained as

$$(\lambda_{i-1} | D_{i-1}) \sim Ga(b_{i-1}, r_{i-1}),$$

which leads to the prior distributions of λ_i at the beginning of each time interval I_i ,

$$(\lambda_i | D_{i-1}) \sim Ga(b_{i0}, r_{i0}),$$

where

$$b_{i0} = d_i b_{i-1}, \quad r_{i0} = d_i r_{i-1}$$

with the discount factor of the time interval I_i , d_i ($0 < d_i < 1$).

With the information from first j observations, the joint prior distribution of the updated distribution of λ_i is obtained as

$$(\lambda_i | D_{i-1(j)}) \sim Ga(b_{i\bar{j}}, r_{i\bar{j}}),$$

where

$$b_{i\bar{j}} = b_{i,j-1} + \delta_{i\bar{j}}, \quad r_{i\bar{j}} = r_{i,j-1} + t_{i\bar{j}} - \tau_{i-1}$$

and $\delta_{i\bar{j}}$ is the indicator function of a death of the j -th individual alive at the beginning of the time interval I_i .

When all individuals in the time interval I_i are observed, the posterior distribution is obtained as

$$(\lambda_i | D_i) \sim Ga(b_i, r_i),$$

where

$$D_i = D_{i-1(n_i)}, \quad b_i = b_{i,n_i} \quad \text{and} \quad r_i = r_{i,n_i}.$$

At the end of each time interval the posterior distribution of λ_i given D_i is used to obtain the survival function under the assumption of a conjugate gamma distribution of the hazard rate λ_i .

Using that

$$\begin{aligned} P(T > t | T > \tau_{i-1}, D_i) &= \int P(T > t - \tau_{i-1}, \lambda_i | D_i) d\lambda_i \\ &= \int P(T > t - \tau_{i-1} | \lambda_i, D_i) p(\lambda_i | D_i) d\lambda_i \\ &= \left(1 + \frac{t - \tau_{i-1}}{r_i}\right)^{-b_i} \quad \text{for } t \in I_i, i = 1, \dots, s, \end{aligned}$$

at the end of the time interval I_i the survival function is obtained as

$$P(T > t | D_i) = \left(1 + \frac{t - \tau_{i-1}}{r_i}\right)^{-b_i} \prod_{k=1}^{i-1} \left(1 + \frac{\tau_k - \tau_{k-1}}{r_k}\right)^{-b_k} \quad \text{for } t \in I_i.$$

At the beginning of the time interval I_{i+1} the prior distribution of λ_{i+1} given D_i is

obtained, which is used to obtain the forecasted survival function of the time interval I_i given D_i ,

$$(\lambda_{i+1}|D_i) \sim Ga(d_{i+1}b_i, d_{i+1}r_i).$$

Thus the forecasted survival function of the time interval I_{i+1} based on the information gathered until the end of the time interval I_i is obtained as

$$P(T > t | D_i) = \left(1 + \frac{t - \tau_i}{d_{i+1}r_i}\right)^{d_{i+1}b_i} \prod_{k=1}^i \left(1 + \frac{\tau_k - \tau_{k-1}}{r_k}\right)^{-b_k} \quad \text{for } t \in I_{i+1}.$$

4. Illustrations

In this section, the performance of the estimation proposed in previous sections is illustrated via the data set in Table 1, which consists of survival times of 45 gastric carcinoma patients of treatment of the chemotherapy.

Table 1. Survival Times in Days

1	63	105	129	182	216	250	262	301	301	342	354	356	358	380	381c	383	383	388	394	408	
460	489	499	524	529c	535	562	675	676	748	748	778	786	797	945c	955	968	1180c	1245			
1271	1277c	1397c	1512c	1519c																	

c: censored, source: Stablein et al.(1981)

In the discount survival model, the survival time of the individuals alive at the beginning of time interval I_i is assumed to have the exponential distribution with the hazard rate λ_i . The end point of each time interval is taken as the multiple of 30 days. We start the analysis with the initial distribution of $(\lambda_0 | D_0) \sim Ga(1, 1362)$, where values of parameters are obtained by equating the PL-estimate(Kaplan and Meier,1958) of the time interval I_1 to the survival function of the time interval I_1 , where the PL-estimate is introduced not to be compared with but to be used as a criterion of the estimates under the discount survival model. The value of the discount factor is tentatively obtained by comparing estimated survival functions from different values of the discount factor - 0.85, 0.9, 0.95. The value $d_i = 0.9$ for $i=1, \dots, 51$, which gives the closest estimated survival function to the PL-estimate, is chosen. Figure 1 shows the cumulative hazard rate obtained under the discount survival model and the cumulative hazard rate estimated from the PL-estimate - the cumulative hazard rate at time i is the negative value of the logarithm of the PL-estimate. In the figure the estimates show the similar pattern in the increase of the cumulative hazard rate. Figure 2 shows the 1-step

ahead forecasted survival function and the PL-estimate. In the figure the 1-step ahead forecasted survival function under the discount survival model looks close to the PL-estimate.

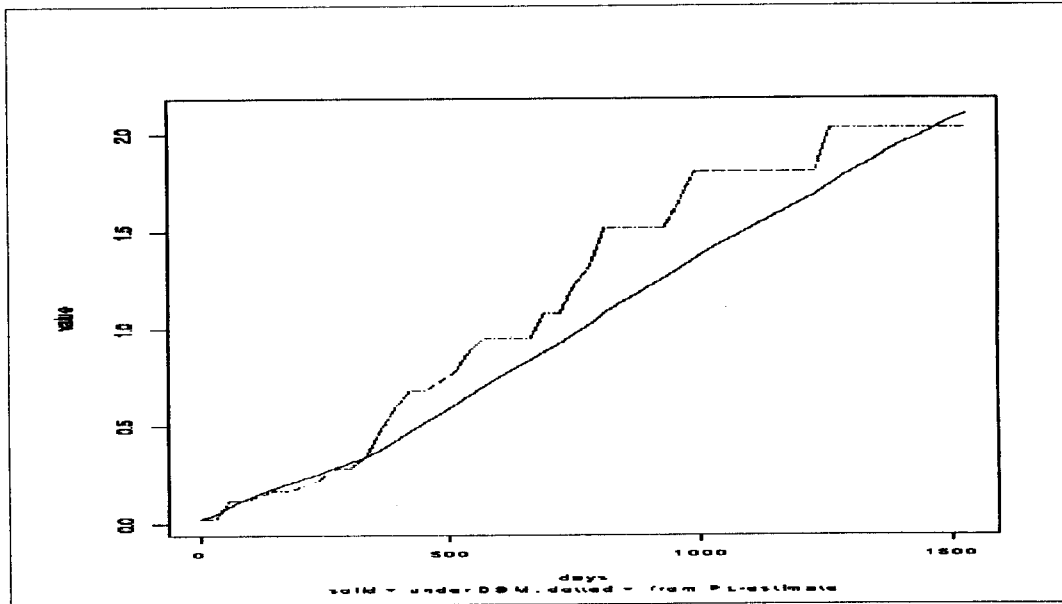


Figure 1. Estimated Cumulative Hazard Rate

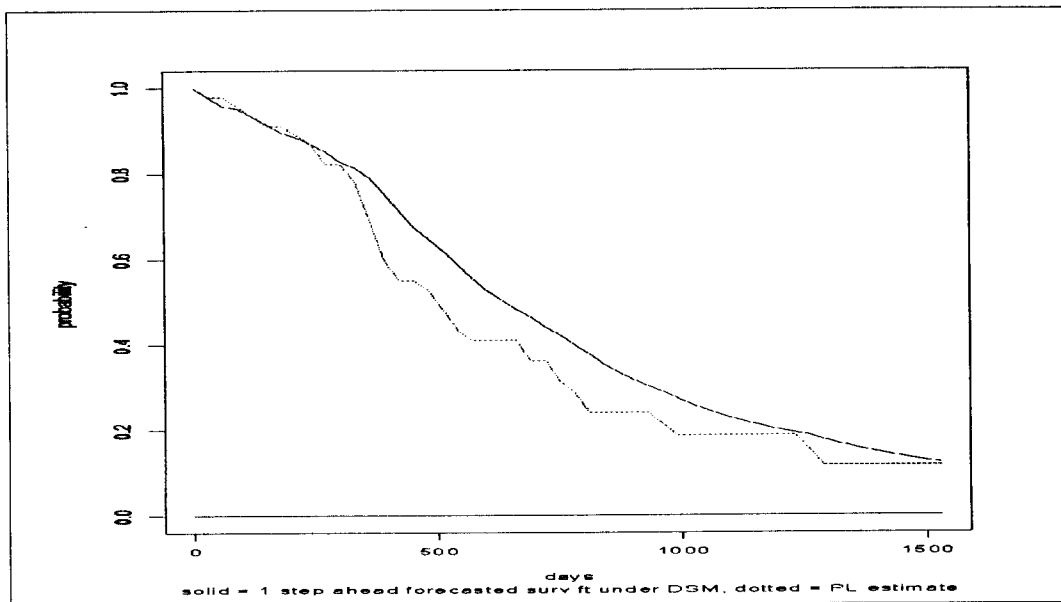


Figure 2. Forecasted Survival Function under DSM and PL-estimate

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