A Bayesian Proportional-Hazards Model
In Survival Analysis

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1. Introduction. Suppose that a sample of n individuals has possible-censored survival times

$$Y_1 \leq Y_2 \leq \ldots \leq Y_n \tag{1.1}$$

Let $\delta_i = 1$ if the ith time Y_i is an observed death and $\delta_i = 0$ if it was a right-censored event: That is, the individual was alive at time Y_i, but was last seen at that time. If T_i ($1 \leq i \leq n$) are the true survival or failure times, then $Y_i = T_i$ if $\delta_i = 1$ and $Y_i < T_i$ if $\delta_i = 0$, in which case the true failure time T_i is unknown.

We also assume d-dimensional *covariate* vectors $X_1, X_2, \ldots, X_n$ for the n individuals in (1.1). The components of X_i might be age, income status, etc. The basic data for (1.1) is the set of triples (Y_i, δ_i, X_i) for $1 \leq i \leq n$. The most important statistical questions are connected with estimating the effect of the covariates X_i on the true survival times T_i.

Let

$$\widetilde{Y}_1 < \widetilde{Y}_2 < \ldots < \widetilde{Y}_m \tag{1.2}$$

be the *distinct* survival times in (1.1). At each time $Y = \widetilde{Y}_j$, let d_j be the number of observed deaths and a_j the number of censored events. Then $n = \sum_{j=1}^{m}(d_j + a_j)$ is the total sample size and $n_{\text{obs}} = \sum_{j=1}^{m} d_j = \sum_{i=1}^{n} \delta_i$ is the total number of observed deaths. The number of distinct observed death times is $r = \sum_{j=1}^{m} I_{[d_j > 0]} \leq m$.

The basic statistical model that we describe below is essentially due to Kalbfleisch (1978). See Clayton (1991) and Ibrahim et al. (2001) for additional discussion and details, and Lee and Wang (2003) for an introduction to survival analysis. The model described below is nonparametric in flavor, but still allows tied survival-time data to be handled in a natural way. The likelihood formula that we derive below for tied data appears to be new. Previous work on this model has mostly assumed survival times (1.1) that are either without ties or else with grouped survival times (Kalbfleisch 1978, Ibrahim et al. 2001).

2. A Survival Model. Let Y be the true lifetime of a random individual with covariates X. By definition, the survival function is

$$S_X(t) = P_X(Y > t) = \exp\big(-H_X(t)\big) = \exp\Big(-\int_0^t h_X(dy)\Big) \tag{2.1}$$

where $H_X(t)$ is a right-continuous increasing function with $H_X(0) = 0$ and $h_X(dy)$ is the related Lebesgue-Stieltjes measure. The function $H_X(t)$ is one form of the *cumulative hazard function* and $h_X(dy)$ the instantaneous hazard measure (or hazard rate). The *Proportional Hazards* assumption is

$$h_X(dy) = e^{\beta X} h(dy) \qquad \text{so that} \qquad H_X(t) = e^{\beta X} H(t) \tag{2.2}$$

for some d-dimensional vector of parameters β, where βX in (2.2) is the dot product. One of the purposes of the model is to estimate β from the data and to test each component of β to find out whether that component of X has a statistically significant effect on the survival times Y. A secondary goal is to estimate the baseline hazard density $h(dy)$, which would allow us to estimate the expected survival time distribution $S_X(t)$ for an individual whose covariates are X, even if X is not among the covariate vectors X_i in the data.

In principle, the likelihood of the data (Y_i, δ_i, X_i) in (1.1) is

$$
L = \left(\prod_{[\delta_i=0]} P_{X_i}(Y > Y_i) \right) \left(\prod_{[\delta_i=1]} P_{X_i}(Y = Y_i) \right) \tag{2.3}
$$

where $P_X(Y = Y_i)$ is with respect to some natural measure on the real line. Many inferential methods in statistics are based on finding the parameters that are the most likely for known data, in the sense of those parameters that have the largest value of L.

To derive an explicit formula for (2.3), choose numbers $\Delta_j > 0$ such that $\widetilde{Y_j} + \Delta_j < \widetilde{Y}_{j+1} - \Delta_{j+1}$ for all j and define the binned likelihood

$$
L_\Delta = \prod_{i=1}^{n} \begin{cases} P_{X_i}(Y > \widetilde{Y_j} + \Delta_j) & \text{if } \delta_i = 0 \\ P_{X_i}(\widetilde{Y_j} - \Delta_j < Y \leq \widetilde{Y_j} + \Delta_j) & \text{if } \delta_i = 1 \end{cases} \tag{2.4}
$$

By definition, the true lifetime $T_i > \widetilde{Y_j}$ for censored individuals with $Y_i = \widetilde{Y_j}$, so that (2.4) is the appropriate probability if the $\Delta_j > 0$ are sufficiently small. The likelihood (2.4) should be asymptotically proportional to (2.3) in the limit as $\Delta_j \to 0$.

We can write (2.4) in terms of the survival function $S_X(t)$ in (2.1) as

$$
\prod_{i=1}^{n} \begin{cases} S_{X_i}(\widetilde{Y_j} + \Delta_j) & \text{if } \delta_i = 0 \\ S_{X_i}(\widetilde{Y_j} - \Delta_j) - S_{X_i}(\widetilde{Y_j} + \Delta_j) & \text{if } \delta_i = 1 \end{cases}
$$

$$
= \prod_{i=1}^{n} \exp\left(-\int_0^{\widetilde{Y_j}-\Delta_j} h_{X_i}(dy)\right) \begin{cases} \exp\left(-\int_{\widetilde{Y_j}-\Delta_j}^{\widetilde{Y_j}+\Delta_j} h_{X_i}(dy)\right) & \text{if } \delta_i = 0 \\ 1 - \exp\left(-\int_{\widetilde{Y_j}-\Delta_j}^{\widetilde{Y_j}+\Delta_j} h_{X_i}(dy)\right) & \text{if } \delta_i = 1 \end{cases}
$$

Define

$$
Z_j = \int_{Y_{j-1}+\Delta_{j-1}}^{\widetilde{Y_j}-\Delta_j} h(dy) \qquad \text{and} \qquad Z_{j0} = \int_{\widetilde{Y_j}-\Delta_j}^{\widetilde{Y_j}+\Delta_j} h(dy) \tag{2.5}
$$

Then

$$
\int_0^{\widetilde{Y_j}-\Delta_j} h(dy) = Z_j + \sum_{k=1}^{j-1}(Z_k + Z_{k0}) = \sum_{k=1}^{j} Z_k + \sum_{k=1}^{j-1} Z_{k0}
$$

so that

$$
\sum_{i=1}^{n} \int_0^{\widetilde{Y_j}-\Delta_j} h_{X_i}(dy) = \sum_{i=1}^{n} e^{\beta X_i} \int_0^{\widetilde{Y_j}-\Delta_j} h(dy)
$$

$$
= \sum_{j=1}^{m} \left(\sum_{[Y_i=\widetilde{Y_j}]} e^{\beta X_i} \right) \left(\sum_{k=1}^{j} Z_k + \sum_{k=1}^{j-1} Z_{k0} \right)
$$

$$
= \sum_{k=1}^{m} Z_k \sum_{j=k}^{m} \left(\sum_{[Y_i=\widetilde{Y_j}]} e^{\beta X_i} \right) + \sum_{k=1}^{m} Z_{k0} \sum_{j=k+1}^{m} \left(\sum_{[Y_i=\widetilde{Y_j}]} e^{\beta X_i} \right)
$$

$$
= \sum_{j=1}^{m} \left(Z_j R_j(\beta) + Z_{j0} R_{j+1}(\beta) \right) \tag{2.6}
$$

In (2.6), $R_j(\beta)$ is the risk sum

$$R_j(\beta) \;=\; \sum_{k=j}^{m}\left(\sum_{[Y_i=\widetilde{Y}_k]} e^{\beta X_i}\right) \;=\; \sum_{[Y_i\geq\widetilde{Y}_j]} e^{\beta X_i} \tag{2.7}$$

corresponding to the individuals who are at risk immediately before time $\widetilde{Y}_j$. We can then write the binned likelihood (2.4) as

$$
\begin{aligned}
L_\Delta \;=\;& \exp\left(-\sum_{j=1}^{m}\Big(Z_j R_j(\beta) + Z_{j0} R_{j+1}(\beta)\Big)\right) \prod_{j=1}^{m}\prod_{\left[\substack{Y_i=\widetilde{Y}_j \\ \delta_i=0}\right]} \exp\big(-Z_{j0}e^{\beta X_i}\big) \\
&\times \prod_{j=1}^{m}\prod_{[Y_i=\widetilde{Y}_j,\delta_i=1]}\Big(1-\exp\big(-Z_{j0}e^{\beta X_i}\big)\Big) \\
\;=\;& \exp\left(-\sum_{j=1}^{m}\Big(Z_j R_j(\beta) + Z_{j0} R_j^0(\beta) + S_j(Z_{j0},\beta)\Big)\right)
\end{aligned}
\tag{2.8}
$$

where

$$S_j(Z_{j0},\beta) \;=\; \sum_{[Y_i=\widetilde{Y}_j,\delta_i=1]} \log\Big(1-\exp\big(-Z_{j0}e^{\beta X_i}\big)\Big) \tag{2.9}$$

is a sum over the observed deaths at times $Y_i = \widetilde{Y}_j$ and

$$R_j^0(\beta) \;=\; \sum_{[Y_i=\widetilde{Y}_j,\delta_i=0]} e^{\beta X_i} \;+\; \sum_{[Y_i>\widetilde{Y}_j]} e^{\beta X_i} \tag{2.10}$$

is the risk sum for individuals who are at risk exactly immediately after time $\widetilde{Y}_j$.

3. A Gamma-process Prior for $H(t) = \int_0^t h(dy)$. A useful way to estimate properties of the baseline hazard density $h(dy)$ is to assume a parameteric model for $H(t) = \int_0^t h(dy)$ and then estimate the parameters involved. A useful parametric probability distribution for the set of increasing functions $H(t)$ for $t \geq 0$ is the gamma process $Z(t)$. This is a stochastic process with independent increments whose increments have the gamma distribution

$$Z(t) - Z(s) \;\approx\; \mathcal{G}\Big(\theta\big(\alpha(t)-\alpha(s)\big), \lambda\Big) \tag{3.1}$$

where $\alpha(t)$ is some strictly-increasing function that is continuously differentiable for $t > 0$. In (3.1), $Z \approx \mathcal{G}(\theta,\lambda)$ means that Z is a random variable with the gamma probability density

$$\frac{\lambda^\theta}{\Gamma(\theta)}\, x^{\theta-1}e^{-\lambda x} \quad \text{for} \quad 0 \leq z < \infty$$

Examples of $\alpha(t)$ in (3.1) would be $\alpha(t) = t$ or $\alpha(t) = t^\sigma$ for some $\sigma > 0$. By (3.1),

$$E\big(Z(t) - Z(s)\big) \;=\; \Big(\theta\big(\alpha(t)-\alpha(s)\big)\Big)/\lambda \;=\; \mu\big(\alpha(t)-\alpha(s)\big) \quad \text{and}$$

$$\text{Var}\big(Z(t) - Z(s)\big) \;=\; \Big(\theta\big(\alpha(t)-\alpha(s)\big)\Big)/\lambda^2 \;=\; \mu\big(\alpha(t)-\alpha(s)\big)/\lambda \tag{3.2}$$

for $\mu = \theta/\lambda$.

If $\alpha(t) = t$, $E\big(Z(t)\big) = \mu t$ in (3.2), so that $\alpha(t) = t$ corresponds to "noisy exponential" baseline survival times. Similarly, if $\alpha(t) = t^\sigma$, then $E\big(Z(t)\big) = \mu t^\sigma$, corresponding to "noisy Weibull" survival distributions. The function $\alpha(t)$ is assumed fixed and θ and λ are parameters to be estimated. Given $\mu = \theta/\lambda$, $1/\lambda$ determines the variance of $H(t) = Z(t)$ about $E\big(H(t)\big) = \mu\alpha(t)$. Often θ or θ and λ are given preassigned values to improve estimation.

The sample paths of the gamma process $Z(t)$ are, with probability one, strictly-increasing purely-discontinuous functions of t, although the probability that any preassigned value of t is a jump is zero. This has the modeling advantage that tied survival-time values can occur with positive probability, even though the survival times themselves (*not* conditioned on the path $Z(t)$) have a continuous distribution, which means that any preassigned survival time has probability zero of being attained.

For any process $Z(t)$ with independent increments, the differences Z_j, Z_{j0} in (2.5) are independent random variables. By (3.1), the Z_j, Z_{j0} are independent random variables with gamma distributions

$$Z_j \approx \mathcal{G}(\theta W_j^\Delta, \lambda) \quad \text{where} \quad W_j^\Delta = \alpha(Y_j - \Delta_j) - \alpha(Y_{j-1} + \Delta_{j-1})$$

$$Z_{j0} \approx \mathcal{G}(\theta W_{j0}^\Delta, \lambda) \quad \text{where} \quad W_{j0}^\Delta = \alpha(Y_j + \Delta_j) - \alpha(Y_j - \Delta_j) \tag{3.3}$$

where we write $Y_j = \widetilde{Y_j}$ for $\widetilde{Y_j}$ in (1.2) for ease of notation.

If the Z_j, Z_{j0} are considered parameters or "hidden variables" in the data (Y_i, δ_i, X_i) with the probability distribution (3.3), then the parameters θ, λ in (3.1) are considered hyperparameters. In a Bayesian framework, the hyperparameters themselves are given probability (or prior) distributions. In this case, we assume gamma prior distributions $\theta, \lambda \approx \mathcal{G}(\epsilon, \epsilon)$ for some small $\epsilon > 0$ ($\epsilon = 0.001$ is the most common choice) and an uninformative normal prior for each component β_j of $\beta \in R^d$, specifically that the prior distributions of β_j are independent normal with means zero and standard deviation $1/\epsilon$ (Ibrahim et al. 2001). However, improper uniform priors for λ and the β_j would work just as well in this case.

4. The Full Likelihood L. Under these conditions, the full binned likelihood of the data, including the prior distributions for Z_j, Z_{j0} and θ, λ, β, corresponding to (2.8) is

$$L_\Delta = \frac{\epsilon^\epsilon}{\Gamma(\epsilon)} \theta^{\epsilon-1} e^{-\epsilon\theta} \frac{\epsilon^\epsilon}{\Gamma(\epsilon)} \lambda^{\epsilon-1} e^{-\epsilon\lambda} \prod_{a=1}^d \frac{1}{\sqrt{2\pi}} \exp\left(-\epsilon^2 \beta_a^2/2\right)$$

$$\times \prod_{j=1}^m \left(\frac{\lambda^{\theta W_j^\Delta}}{\Gamma(\theta W_j^\Delta)} Z_j^{\theta W_j^\Delta - 1} e^{-\lambda Z_j} e^{-Z_j R_j(\beta)} \right. \tag{4.1}$$

$$\times \left. \frac{\lambda^{\theta W_{j0}^\Delta}}{\Gamma(\theta W_{j0}^\Delta)} Z_{j0}^{\theta W_{j0}^\Delta - 1} e^{-\lambda Z_{j0}} e^{-Z_{j0} R_j^0(\beta)} \prod_{[Y_i = \widetilde{Y}_j, \delta_i = 1]} \left(1 - \exp(-Z_{j0} e^{\beta X_i})\right) \right)$$

As each $\Delta_j \to 0$,

$$W_j^\Delta \to W_j = \alpha(Y_j) - \alpha(Y_{j-1}) > 0 \quad \text{and} \quad W_{j0}^\Delta \to 0 \tag{4.2}$$

for $Y_j = \widetilde{Y_j}$ as before. The expressions in the first line of (4.1) vary continuously as $W_j^\Delta \to W_j > 0$. As $W_{j0}^\Delta \to 0$, the jth factor in the second line in (4.1) is asymptotic to $C(Z_{j0})\Gamma(\theta W_{j0}^\Delta)^{-1} \sim C(Z_{j0})\theta\alpha'(\widetilde{Y}_j)\Delta_j$ for $Z_{j0} > 0$ and $C(Z_{j0}) > 0$. There are two cases for the asymptotic behavior of the jth factor in the second line in (4.1):

If $d_j = 0$, the jth factor has a delta-function singularity at $Z_{j0} = 0$ as $\Delta_j \to 0$ and L_Δ does not need to be rescaled. In this case, the factors in (4.1) with Z_{j0} disappear in the limit as $\Delta_j \to 0$ (with $Z_{j0} = 0$).

If $d_j \geq 1$, the function $C(Z_{j0})$ is a bounded and continuous function of Z_{j0} for $Z_{j0} \geq 0$ and the jth factor in (4.1) is asymptotic to $C(Z_{j0})\theta\alpha'(Y_j)\Delta_j$ as $\Delta_j \to 0$.

Thus, ignoring constants that depend on Δ_j for $d_j > 0$, the limit of L_Δ in (4.1) as $\max_j \Delta_j \to 0$ is the limiting full likelihood

$$
\begin{aligned}
L \;=\; & C\,\lambda^{\epsilon-1}e^{-\epsilon\lambda}\left(\theta^{r+\epsilon-1}e^{-\epsilon\theta}\right)\exp\left(-\epsilon^2\sum_{a=1}^{d}\beta_a^2/2\right) \\
& \times \prod_{j=1}^{m}\left(\frac{\lambda^{\theta W_j}}{\Gamma(\theta W_j)}Z_j^{\theta W_j-1}\exp\left(-Z_j\left(\lambda+R_j(\beta)\right)\right)\right) \\
& \times \prod_{[d_j\geq 1]}^{m}\exp\left(-Z_{j0}\left(\lambda+R_j^0(\beta)\right)\right)\left(\frac{\prod_{[Y_i=\widetilde{Y}_j,\delta_i=1]}\left(1-\exp\left(-Z_{j0}e^{\beta X_i}\right)\right)}{Z_{j0}}\right)
\end{aligned}
\tag{4.3}
$$

In (4.3), C depends on ϵ and $\alpha'(Y_j)$ and r is the number of distinct times $Y_j = \widetilde{Y}_j$ with $d_j \geq 1$. As mentioned earlier, inferences about which parameter values are relatively more likely are based on finding relatively larger values of L in (4.3) for the data (Y_i, δ_i, X_i).

5. Estimating Parameters Using the Likelihood L.

We estimate the parameters and hidden variables $(\theta, \lambda, Z_j, Z_{j0}, \beta)$ in (4.3) by using Markov Chain Monte Carlo methods (Metropolis et al. 1953, Hastings 1970, Gilks et al. 1996).

Specifically, we define a Markov Chain Q_n that takes its values in the space of possible parameter vectors $(\theta, \lambda, Z_j, Z_{j0}, \beta)$ and which has a stationary or asymptotic distribution that is proportional to (4.3). This means that Q_n spends most of its time where the likelihood (4.3) is the largest. Mean or median values of components or functions of components of Q_n can be used to provide estimates of the parameters affecting the true survival times T_i.

The Markov chain Q_n proceeds by changing or updating each of the components of the vector $(\theta, \lambda, Z_j, Z_{j0}, \beta)$ in turn in a way that depends on the conditional probability distribution of that parameter value given the data and all the other parameters. We carry out these parameter changes or updates in the following way:

Updating θ : Ignoring multiplicative constants and also ignoring factors in (4.3) that do not depend on θ, the conditional density of θ given the data and the other parameters is

$$
\theta^{r+\epsilon-1}e^{-\epsilon\theta}\lambda^{\theta W}\prod_{j=1}^{m}\frac{Z_j^{\theta W_j}}{\Gamma(\theta W_j)}\qquad\text{where}\quad W\;=\;\sum_{j=1}^{m}W_j
\tag{5.1}
$$

for W_j in (4.2). The density (5.1) is asymptotic to $C\theta^{r+m+\epsilon-1}$ as $\theta \to 0$ and decays faster than exponentially at infinity, and can be updated efficiently by one step of a Metropolis random walk (Metropolis et al. 1953).

Alternatively, the density (5.1) is a log-concave function of θ, so that θ can be updated by a "Gibbs sampler" step that samples directly from the distribution (5.1) using one of the adaptive-rejection methods of Gilks and Wild (1992) or Gilks (1992). (See also Gilks et al. 1995.)

In general, a function $f(\theta)$ is called *log-concave* if $(d/d\theta)^2(\log f(\theta)) < 0$ for all θ, or, more generally, if $(d/d\theta)\log f(\theta)$ is decreasing in θ. The log-concavity of (5.1) follows from the identity

$$
\frac{d^2}{d\theta^2}\log\Gamma(\theta)\;=\;\text{Var}\left(\log\mathcal{G}(\theta,1)\right)\;>\;0
\tag{5.2}
$$

where $\mathcal{G}(\theta,1)$ represents a gamma-distributed random variable (as in (3.1)). (*Exercise*: Prove (5.2).)

Updating λ : Ignoring multiplicative constants and factors in (4.3) that do not depend on λ, the conditional density of λ given the data and the other parameters is

$$\lambda^{\theta W + \epsilon - 1} \exp\left(-\lambda\left(\epsilon + \sum_{j=1}^{m}\left(Z_j + Z_{j0}\right)\right)\right) \tag{5.3}$$

where $Z_{j0} = 0$ if $d_j = 0$. This can be updated by a Gibbs sampler step by sampling from the gamma distribution

$$\lambda \approx \mathcal{G}\left(\epsilon + \theta W, \; \epsilon + \sum_{j=1}^{m}\left(Z_j + Z_{j0}\right)\right)$$

See Fishman (1995) for algorithms for generating gamma-distributed random variates. Two other good references for statistical computation and for scientific computing in general are Devroye (1986) and Press et al. (1992).

Updating Z_j : Ignoring multiplicative constants and factors that do not depend on Z_j, the conditional density of Z_j given the other parameters is

$$Z_j^{\theta W_j - 1} e^{-Z_j\left(\lambda + R_j(\beta)\right)} \tag{5.4}$$

Thus Z_j can be updated by sampling from the gamma distribution

$$Z_j \approx \mathcal{G}\left(\theta W_j, \; \lambda + R_j(\beta)\right)$$

Updating Z_{j0} for $d_j \geq 1$: Ignoring multiplicative constants and factors that do not depend on Z_{j0}, the conditional density of Z_{j0} given the other parameters is

$$e^{-Z_{j0}\left(\lambda + R_j^0(\beta)\right)} \left(\frac{\prod_{[Y_i = \widetilde{Y}_j, \delta_i = 1]}\left(1 - \exp\left(-Z_{j0}e^{\beta X_i}\right)\right)}{Z_{j0}}\right) \tag{5.5}$$

The density (5.5) is normalizable in Z_{j0} and can be updated by a one step of a Metropolis random walk. Unfortunately, the density (5.5) is not log-concave in Z_{j0} due to the factor of Z_{j0} in the denominator.

Alternatively, a more general sampling technique can be used for (5.5) that does not require log concavity (Gilks et al. 1995). This method, called "Metropolis-within-Gibbs" sampling, is equivalent to independence Metropolis-Hastings sampling (Gilks et al. 1996) using, as the proposal distribution, an approximation of the density (5.5) based on the method of Gilks (1992). If the density that is being approximated is log concave, the method reduces to the adaptive-rejection method of Gilks (1992).

Technically speaking, the term "Metropolis-within-Gibbs" is not quite corrent, since independence sampling is not Metropolis sampling in the original sense. Metropolis et al. (1953) only described proposal distributions that are one step of a symmetric Markov chain. Independence sampling is contained in a generalization of Metropolis et al. (1953) due to Hastings (1970). The latter sampling scheme (or schemes) are usually called "Metropolis-Hastings" sampling.

Independence samplers can have extremely bad convergence properties if the proposal distribution is less singular or less heavy-tailed than the distribution being approximated (Gilks et al. 1996). In that case, a Metropolis random walk can be used instead.

Large values of βX_i can cause numerical underflows and overflows in the exponential and risk sums in (5.5), but do not cause arbitrarily large values in the likelihood (5.5). Depending on the compiler, computer programs may have to be adjusted to avoid program crashes in evaluating

$\exp\left(-Z_{j0}e^{\beta X_i}\right)$ in (5.5) if βX_i is large and positive. The term $\exp\left(-Z_{j0}e^{\beta X_i}\right)$ in (5.5) can be set explicitly to be equal to zero in programming code if $Z_{j0} > 0$ and $\beta X_i > 500$ and equal to one if $\beta X_i < -500$. Most modern computers replace exponential underflows (that is, smaller positive values than the program can handle) by zero without a program warning or crash. If numerical underflows in exponentials can also cause program crashes, program adjustments may also have to be made if $Z_{j0}e^{\beta X_i}$ by itself is large.

Updating β : Ignoring multiplicative constants and factors that do not depend on β, the conditional density of β_a in (4.3) given the data and other parameters is

$$\exp\left(-\sum_{j=1}^{m}\left(Z_j R_j(\beta) + Z_{j0}R_j^0(\beta) - S_j(Z_{j0}, \beta)\right) - \frac{1}{2}\epsilon^2\beta_a^2\right) \tag{5.6}$$

where

$$S_j(Z_{j0}, \beta) \;=\; \sum_{[Y_i = \widetilde{Y}_j, \delta_i = 1]} \log\left(1 - \exp\left(-Z_{j0}e^{\beta X_i}\right)\right)$$

is a sum over the observed deaths at times $Y_i = \widetilde{Y}_j$ as in (2.9). If there are no observed deaths at time $Y_i = \widetilde{Y}_j$, then $S_j(Z_{j0}, \beta) = 0$ and $Z_{j0} = 0$, and the second two terms in the sums in (5.6) do not appear.

   Baring linear dependencies among the sample covariates, the conditional likelihood in (5.6) is normalizable in each component β_j, so that each β_j can be updated efficiently by one step of a Metropolis random walk.

   Alternatively, the density (5.6) is log-concave in β_a, so that Gibbs sampler updates can be made using the adaptive rejection methods of Gilks and Wild (1992) or Gilks (1992). Gilks has programming code in C on a Web site for carrying out Metropolis-within-Gibbs sampling that reduces to Gilks (1992) if a parameter is set. This C code can be used for non-Metropolis updates of θ, Z_{j0}, and β.

6. A Likelihood For (θ, λ, β) . The advantage of the Markov Chain Monte Carlo (MCMC) procedure of the previous section is that it also gives us information about the conditional distribution of the baseline cumulative hazards

$$Z_j \;\approx\; H(Y_j-) - H(Y_{j-1}) \quad \text{and} \quad Z_{j0} \;\approx\; dH(Y_j) = h(dY_j)$$

given the observed data. If we are primarily interested the parameters (θ, λ, β) and not in the baseline hazard density $h(dy)$, the Z_j, Z_{j0} can be integrated out of the likelihood (4.3) to obtain a marginal likelihood that depends only on (θ, λ, β).

   Evaluating the integrals $\int L(Z_j)\,dZ_j$ in (4.3) in succession yields

$$L \;=\; C\,\lambda^{\epsilon-1}e^{-\epsilon\lambda}\,\theta^r \prod_{j=1}^{m}\left(\frac{\lambda}{\lambda + R_j(\beta)}\right)^{\theta W_j} \times \exp\left(-\epsilon^2\sum_{a=1}^{d}\beta_a^2/2\right) \tag{6.1}$$

$$\times \prod_{[d_j \geq 1]}^{m} \exp\left(-Z_{j0}\left(\lambda + R_j^0(\beta)\right)\right)\left(\frac{\prod_{[Y_i = Y_j, \delta_i = 1]}\left(1 - \exp\left(-Z_{j0}e^{\beta X_i}\right)\right)}{Z_{j0}}\right)$$

While λ no longer has a simple gamma update, the parameter θ now has a gamma update, specifically

$$\theta \;\approx\; \mathcal{G}\left(r + 1, \sum_{j=1}^{m} W_j \log\left((\lambda + R_j(\beta))/\lambda\right)\right) \tag{6.2}$$

The parameters Z_{j0} can be integrated by using the identity

$$\int_0^\infty \frac{e^{-at} - e^{-bt}}{t} \, dt = \int_0^\infty \int_a^b e^{-\theta t} \, d\theta dt = \int_a^b \frac{d\theta}{\theta} = \log \frac{b}{a} \tag{6.3}$$

for $b > a > 0$. Thus if $d_j = 1$, the jth factor in the second line of (6.1) integrates to

$$\log \left(\frac{\lambda + R_j^0(\beta) + e^{\beta X_i}}{\lambda + R_j^0(\beta)} \right) = \log \left(\frac{\lambda + R_j(\beta)}{\lambda + R_j^0(\beta)} \right)$$

In particular, if $d_j \leq 1$ for all j, so that there are no ties among observed death times, then evaluating the integrals $\int L(Z_{j0}) \, dZ_{j0}$ for $d_j = 1$ in (6.1) leads to the more compact form

$$L = L(\theta, \lambda, \beta) = C \, \lambda^{\epsilon-1} e^{-\epsilon\lambda} \, \theta^r \prod_{j=1}^m \left(\left(\frac{\lambda}{\lambda + R_j(\beta)} \right)^{\theta W_j} \log \left(\frac{\lambda + R_j(\beta)}{\lambda + R_j^0(\beta)} \right) \right) \tag{6.4}$$

ignoring the prior terms in β. If $d_j = 0$, then $R_j^0(\beta) = R_j(\beta)$ and the logarithmic factor does not appear. Analogous expressions can be found for $d_j \geq 2$ by expanding the last product in (6.1) into a linear combination of differences of exponentials and applying (6.3).

The likelihood (6.4) no longer has information about the baseline hazards Z_j, Z_{j0}, although the conditional density of Z_j, Z_{j0} is given by (5.4) and (5.5) if β is known precisely. See Kalbfleisch (1978) for a different derivation if $d_j \leq 1$ for all j.

7. The Posterior Distribution of the Hazard Function $H(t)$.
(In Bayesian terminology, "posterior" means "conditional on the observed data for a given prior".)

For any j and any partition $(Y_{j-1}, Y_j) = \bigcup_{a=1}^{A_j} (Y_{j,a-1}, Y_{ja})$ of (Y_{j-1}, Y_j), define $Z_j = H(Y_j) - H(Y_{j-1}) = \sum_{a=1}^{A_j} Z_{ja}$ for $Z_{ja} = H(Y_{ja}-) - H(Y_{j,a-1})$. The same argument as in (2.5) to (4.3) shows that the posterior distribution (4.3) is still valid with Z_{ja} in place of Z_j, with of course $d_{ja} = 0$ unless there is an actual observed death at Y_{ja}. This implies that the posterior distribution of the random variables Z_{ja} is that they are independent gamma-distributed random variables with distributions

$$Z_{ja} \approx \mathcal{G} \Big(\theta \big(\alpha(Y_{ja}) - \alpha(Y_{j,a-1}) \big), \lambda + R_j(\beta) \Big) \tag{7.1}$$

This in turn implies that, for each j, the posterior distribution of the process $Z(t) - Z(Y_{j-1}) = H(t) - H(Y_{j-1})$ for $Y_{j-1} < t < Y_j$ is that of a gamma process in (Y_{j-1}, Y_j) with scale parameter $\lambda_j = \lambda + R_j(\beta)$, with jumps $Z(\widetilde{Y_j}+) - Z(\widetilde{Y_j}-) \approx Z_{j0}$ in the posterior distribution of $Z(t)$ at observed death times $\widetilde{Y_j}$. As before, $Z_{j0} = 0$ if $d_j = 0$. If $d_j > 0$, Z_{j0} has the density (5.5). (See also Kalbfleisch 1978 and Clayton 1991.)

8. Simulating Data for the Model in Sections 1–3.
We can simulate survival data (Y_i, δ_i, X_i) for the model (2.1)–(2.2)–(3.1) as follows:

First, choose a sample size n, the number of covariates d, and, for $i \leq i \leq n$, covariates $X_i \in R^d$. As in most regression models, these are assumed to be deterministic and are arbitrary. Choose arbitrary parameters values $\theta, \lambda > 0$ and risk parameters $\beta \in R^d$. Also, choose a strictly-increasing continuously-differentiable function $\alpha(t)$ with $\alpha(0) = 0$, for example $\alpha(t) = t$.

The first goal is to define failure times Y_i satisfying (2.1)–(2.2)–(3.1), that is

$$P(Y_i > t) = \exp\big(-H_{X_i}(t)\big) = \exp\big(-e^{\beta X_i} H(t)\big), \quad t \geq 0 \tag{8.1}$$

where $H(t) = Z(t)$ is a realization of the gamma process

$$Z(t) \approx \mathcal{G}\big(\theta\alpha(t), \lambda\big) \approx (1/\lambda)\mathcal{G}\big(\theta\alpha(t), 1\big) \tag{8.2}$$

The final step will be to modify the construction so that some of the observations Y_i can be censored.

The sample paths of $Z(t)$ are right-continuous with jumps in every time interval (t_1, t_2) with $0 \le t_1 < t_2$. This implies

$$P(Y_i > t) = P\big(Z(Y_i) > Z(t)\big) = \exp\big(-e^{\beta X_i} Z(t)\big)$$

so that

$$P(Z(Y_i) > s) = \exp\big(-e^{\beta X_i} s\big) \tag{8.3}$$

whenever $s = Z(t)$. This suggests that $Z(Y_i)$ might have an exponential distribution with mean $e^{-\beta X_i}$, but this is not correct. In fact, given $Z(t)$, the values of $Z(Y_i)$ are restricted to the range of $Z(t)$, which is the complement of an open dense set of real numbers since $Z(t)$ is increasing with jumps in every open interval. This means that if the random variable $Z(Y_i)$ has a probability distribution with a density $g(s)$, then $g(s) = 0$ on an open dense set of real numbers s. Thus $Z(Y_i)$ cannot have a probability distribution with a continuous density.

If the variables $Z(Y_i)$ were exponentially distributed, then we could simulate $Y_i \approx Z^{-1}(Z_i)$ where $Z_i \approx Z(Y_i)$ had a known distribution. However, we can do essentially the same even though the $Z(Y_i)$ are not exponentially distributed.

Let Z_i be independent exponentially distribution random variables with mean $e^{-\beta X_i}$, as incorrectly suggested for $Z(Y_i)$ by (8.3). The Z_i can be simulated as

$$Z_i \approx e^{-\beta X_i}\big(-\log(U_i)\big)$$

where U_i are independent uniforms for $0 \le U_i \le 1$. Define

$$Y_i = \min\{\, t : Z(t) \ge Z_i \,\} \tag{8.4}$$

Then $Y_i \le t_2$ if and only if $Z(t_2) \ge Z_i$, so that

$$P(Y_i > t) = P(Z(t) < Z_i) = \exp\big(-e^{\beta X_i} Z(t)\big) \tag{8.5}$$

which is exactly (8.1). If follows from (8.3) and (8.5) that $P\big(Z(Y_i) \le s\big) = P(Z_i \le s)$ whenever s is a value attained by $Z(t)$, but $Z(Y_i)$ and Z_i have different probability distributions.

To simulate Y_i from (8.4), we need an approximate sample path of $Z(t)$. Define independent gamma-distributed random variables

$$Q_j \approx \mathcal{G}\big(\theta\Delta(j,m), 1\big) \quad \text{for} \quad 1 \le j \le mT \tag{8.6}$$

where $\Delta(j,m) = \alpha(j/m) - \alpha((j-1)/m)$ and m and T are large. In particular, $\Delta(j,m) = 1/m$ if $\alpha(t) = t$. In general, by (8.2) and (8.6),

$$Z(k/m) \approx \mathcal{G}\big(\theta\alpha(k/m), \lambda\big) \approx (1/\lambda)\mathcal{G}\big(\theta\alpha(k/m), 1\big) \approx (1/\lambda)\sum_{j=1}^{k} Q_j$$

Thus we can simulate Y_i in (8.4) by

$$Y_i = \min\{\, k/m : (1/\lambda)\sum_{j=1}^{k} Q_k \ge Z_i \,\} = \frac{1}{m}\min\{\, k : \sum_{j=1}^{k} Q_k \ge \lambda Z_i \,\}$$

or, equivalently, by

$$Y_i \;=\; \frac{1}{m}\,\min\{\,k : \sum_{j=1}^{k} Q_k \geq \widetilde{Z}_i\,\} \quad \text{where} \tag{8.7}$$

$$\widetilde{Z}_i \;\approx\; \lambda \exp(-\beta X_i)\big(-\log(U_i)\big) \;\approx\; \lambda Z_i$$

To include censoring, we define *censoring times*

$$Y_i^c \;=\; \frac{1}{m}\min\{\,k : \sum_{j=1}^{k} Q_j \geq \widetilde{Z}_i^c\,\} \quad \text{for} \quad \widetilde{Z}_i^c \;\approx\; \mu e^{-\beta X_i}\big(-\log(U_i)\big)$$

in the same way for some constant $\mu > 0$. Define $\delta_i = 1$ (that is, the true failure time $Y_i = T_i$ is observed) if $Y_i < Y_i^c$ and $\delta_i = 0$ (that is, $Y_i < T_i$ and Y_i is censored) if $Y_i^c < Y_i$. The last observed times (observed failure or censoring times) are

$$Y_i^o = \min\{\,Y_i, Y_i^c\,\} = \frac{1}{m}\min\{\,k : \sum_{j=1}^{k} Q_j \geq \widetilde{Z}_i^o\,\}, \quad \widetilde{Z}_i^o = \min\{\,\widetilde{Z}_i, \widetilde{Z}_i^c\,\} \tag{8.8}$$

In general, if X_1 and X_2 are independent exponentials with $E(X_1) = \mu_1$ and $E(X_2) = \mu_2$, then $X_3 = \min\{X_1, X_2\}$ is exponential with $E(X_3) = \mu_1\mu_2/(\mu_1 + \mu_2)$ and $P(X_1 < X_2) = \mu_2/(\mu_1 + \mu_2)$. Morever, X_3 and the event $\{X_1 < X_2\}$ are independent. (*Exercise*: Prove these three statements.)

This implies that the triple (Y_i^o, δ_i, X_i) for Y_i^o in (8.8) satisfies the conditions of the model (2.1)–(2.2)–(3.1) with λ replaced by $\lambda\mu/(\lambda + \mu)$. Moreover, the variables $\delta_i = I_{\{\widetilde{Z}_i < \widetilde{Z}_i^c\}}$ are independent with $P(\delta_i = 0) = P(\widetilde{Z}_i^c < \widetilde{Z}_i) = \lambda/(\lambda + \mu)$, and the δ_i are independent of $\widetilde{Z}_i^o$.

This means that if we choose θ, λ and $0 < q < 1$ and define

$$Y_i \;=\; \frac{1}{m}\min\{\,k : \sum_{j=1}^{k} Q_j > \widetilde{Z}_i\,\} \quad \text{for} \quad \widetilde{Z}_i \;\approx\; \lambda(1 - q)e^{-\beta X_i}\big(-\log(U_i)\big)$$

and then, for each i, independently of the value of Y_i, call Y_i *censored* ($\delta_i = 0$) with probability q and *observed* ($\delta_i = 1$) with probability $1 - q$, then (Y_i, δ_i, X_i) satisfy the conditions of Sections 1–3 with the original value of λ. (*Exercise*: Prove this. Note that if $\mu = ((1 - q)/q)\lambda$, then $\lambda\mu/(\lambda + \mu) = (1 - q)\lambda$ and $\lambda/(\lambda + \mu) = q$.)

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