Evaluating the Impact of Baseline Hazard Function Misspecification on Treatment Effect Estimation

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Objectives

The goal of this study is to evaluate how misspecifying the baseline hazard function can influence the estimation of treatment effects in survival without censored observations (all subjects have an event throughout the study). This work focuses on conducting simulations to compare the exponential and Weibull proportional hazards models to the Cox proportional hazards model. We also discuss the impact of utilizing an overly complicated model (e.g., Cox) when a less complex model (e.g., exponential) is sufficient.

Statistical Methods

Survival analysis is used to analyze time-to-event data (e.g., time to symptom onset or time to mortality). Survival functions, S(t), measure the probability of an individual not experiencing an event past a certain time t. Similarly, hazard functions, h(t), measure the instantaneous risk of failure at a certain time t, given that the individual has not experienced an event until that time. The hazard function can be expressed as $\frac{f(t)}{S(t)}$, where f(t) is the distribution of survival times.

One purpose of proportional hazards modeling is to assess the effectiveness of a particular treatment (X) over survival time T, where the hazard ratio for patient i at time t is defined as $h_i(t) = h_0(t)e^{x_i\beta}$. Here, $h_0(t)$ denotes the pre-specified baseline hazard function, x_i indicates treatment allocation (0=control, 1=treatment), and β represents the log hazard ratio, or the hazard reduction among treated individuals compared to the control group. Thus, the proportional hazard can be expressed as $\frac{h(t|x_0)}{h(t|x_1)} = e^{\beta(x_0 - x_1)}$, which is independent of survival time t.

We consider three models: exponential, Weibull, and Cox. The exponential and Weibull models implement different baseline hazard functions, while the Cox model estimates β without this specification. Notably, even if the baseline hazard function is known, the Cox model is still expected to perform well due to its semi-parametric efficiency. It is important to note that the Exponential model is nested within the Weibull model, as an exponential distribution is a special case of the Weibull distribution where $\gamma = 1$. All models impose the restraint that the effect of the treatment be multiplicative on the hazard curve. The exponential and Weibull hazard curves can be seen below, in Table 1.

Simulation Design

All simulation data was generated using the simsurv function in the simsurv package. We defined a binomial treatment variable (trt) with p = 0.5 to ensure equal likelihood of random assignment to the treatment or control group. Parameters are set according to different distributions of hazards. The resulting dataset contains time of event(eventtime), status(status) and treatment(trt).

We simulated survival data from six different baseline hazard distributions. First, we simulated from an exponential distribution with $\lambda=1$, and from a Weibull with $\lambda=0.1$, and γ values of 0.5, 2, and 5. These values of γ were chosen to consider constant, monotone decreasing, linearly increasing, and monotone increasing hazard curves. From these simulations we aim to test the impact of utilizing an overly complex model, such as Cox, when an exponential or Weibull model would suffice.

We additionally simulated lognormal data with $\mu = 0$, $\sigma = 0.5$, which generates a non-monotone hazard curve, to understand the implications of mispecifying the underlying distribution when fitting a model. These first five hazard functions we simulated from can be seen in Table 1, below.

	Exponential	Weibull	Lognormal
$h_0(t)$	λ	$\lambda \gamma t^{\gamma-1}$	$\frac{\frac{1}{t\sigma}\phi(\frac{ln(t)}{\sigma})}{\Phi(\frac{-ln(t)}{\sigma})}$

Table 1. Baseline hazard functions, $h_0(t)$, for the continuous models considered in this study. The normal PDF and CDF are denoted by ϕ and Φ , respectively.

Lastly, we simulated from a discrete piecewise step function hazard curve, inspired by Brilleman et al. This hazard curve is visualized in Figure 1, below. It was included to further explore the implications of fitting an overly-simplistic model such as exponential or Weibull on data with a complex underlying distribution.

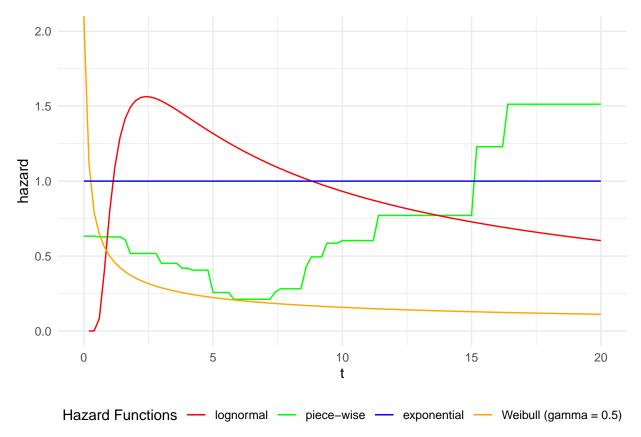


Figure 1: Baseline hazard functions used to generate data. Note Weibull distribution with $\gamma=2,5$ are excluded due to scale.

When simulating the data, we used $\beta = -0.5$ as the true effect size for all underlying distributions.

We set the sample size as 100 and for each underlying distribution sampled 1000 times to get the data frame of results. Each time we fitted exponential, Weibull and cox model to the data and extracted the β estimates provided by three models.

Results

To assess model performance, we plotted the survival probability versus time of all models with each simulation data. We also used mean-squared error (MSE) to check model performance.

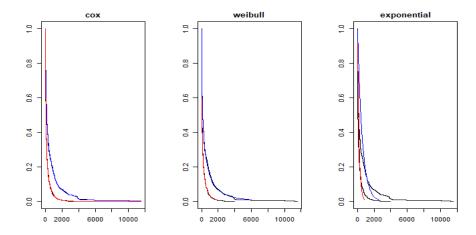


Figure 2. Survival probability versus time of each model, baseline hazard function follows weibull distribution with $\gamma = 0.5$.

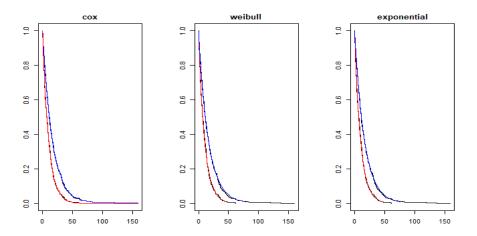


Figure 3. Survival probability versus time of each model, baseline hazard function follows weibull distribution with $\gamma = 1$.

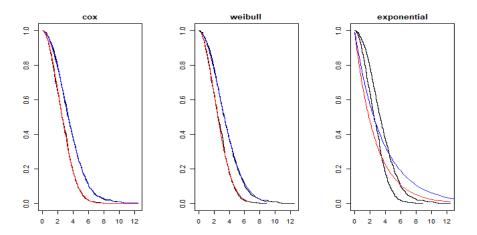


Figure 4. Survival probability versus time of each model, baseline hazard function follows weibull distribution with $\gamma=2$.

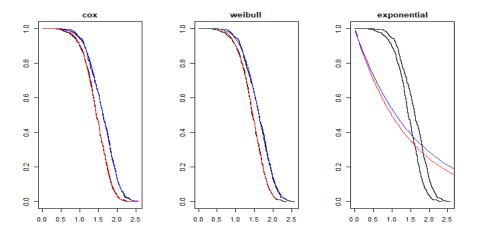
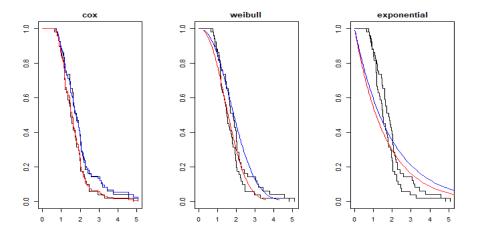


Figure 5. Survival probability versus time of each model, baseline hazard function follows weibull distribution with $\gamma = 5$.



 $Figure \ 6. \ Survival \ probability \ versus \ time \ of \ each \ model, \ baseline \ hazard \ function \ follows \ lognormal \ distribution.$

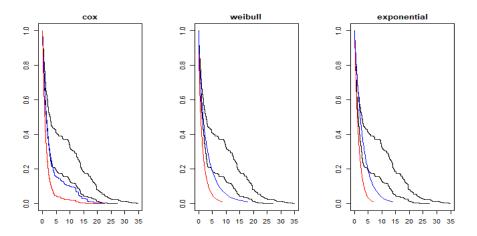


Figure 7. Survival probability versus time of each model, baseline hazard function follows piecewise distribution.

Model	$Gamma_0.5$	Gamma_1	${\rm Gamma_2}$	Gamma_5	Lognormal	Piecewise
Exponential Weibull Cox	0.3297817 1.7392631 0.3091898	0.1499803 0.1544043 0.1489322	0.0463993 0.0755829 0.0489004	$\begin{array}{c} 0.1615276 \\ 0.1617968 \\ 0.0461492 \end{array}$	0.0728173 0.0555354 0.0422765	$\begin{array}{c} 0.0558371 \\ 0.0563854 \\ 0.0422971 \end{array}$

Table 2. The MSE table of three models with each simulation data.

Conclusions

By comparing the Mean square error (MSE) and mean of β derived from three models in diffrent simulations, we found that misspecifying the baseline hazard function can lead to significant differences between actual value of β and the truth (which was set as -0.5). For example, we got high MSEs and biased means of β when applying exponential and Weibull model in data simulated by gamma distributions, which indicates that the treatment effect was not evaluated well. In this case, Cox model provide small MSEs, while means of β are quite close to the truth. Thus we conclude that Cox model can estimate the treatment effect with high accuracy and precision which avoids the impacts of misspecifying the baseline hazard function.

Case of the impact of fitting too complicated a model when an exponential is sufficient

Contributions

Charly Fowler worked on simulation functions, performed piecewise simulation, plotted baseline hazard curves, and created datasets of results. Hanfei Qi worked on formatting simulation functions and editing plotting functions. Robert Tumasian III worked on editing simulation functions, performing lognormal simulation, and creating datasets of results. Haoyang Yi worked on creating plotting functions and editing simulation functions. All members contributed equally to this project.

References

Brilleman, Samuel, Rory Wolfe, Margarita Moreno-Betancur, & Michael J. Crowther. "Simulating Survival Data Using the simsurv R Package." *Journal of Statistical Software* [Online], 97.3 (2021): 1 - 27. Web. 23 Feb. 2021