Introduction to survival analysis and Cox proportional hazards models

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Introduction to Cox proportional hazards

(These slides are primarily based on chapter 3 of Kleinbaum and Klein (2012))



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- Uses covariates which do not vary with time
- An extension for time-dependent variables exists (but is often inadvisable to use) (See Prentice (1982))

Hazard function

The Cox proportional hazards model is typically presented via the hazard function:

$$h(t,X) = h_0(t) \cdot e^{\sum_{i=1}^{p} \beta_i x_i}$$
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- ► The baseline hazard is dependent only on t whilst the second term is dependent only on X and a set of regression coefficients.
- h₀(t) is unspecified but is robust: closely approximates the "correct" parametric model

Maximum (partial) likelihood estimates 1

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- Actually partial likelihood due to only considering probabilities for subjects who met the end-point (non-censored).
- Can be written as product of several likelihoods (one for each event time). If we rank event times, then each contribution to the likelihood is the probability of individual j failing at the jth event time given there is one event from the group of individuals still at risk at this time point (the risk set).

$$L = L_1 \times L_2 \times \dots L_k = \prod_{j=1}^k L_j$$
 (2)

Let $X_j = (X_{j1}, ... X_{jp})$ be the values of the covariates for subject j. Then L_j can be written as:

$$L_{j} = \frac{h(Y_{j}|X_{j})}{\sum_{g:Y_{g} \geq y_{j}} h(Y_{j}|X_{g})} = \frac{h_{0}(Y_{j})\theta_{j}}{\sum_{g:Y_{g} \geq y_{j}} h_{0}(Y_{j})\theta_{g}} = \frac{\theta_{j}}{\sum_{g:Y_{g} \geq y_{j}} \theta_{g}}$$

where $\theta_j = \exp(X_j\beta)$.

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We maximise by solving

$$\frac{\partial \ln L}{\partial \beta_i} = 0 \tag{3}$$

for i = 1, ..., p (p parameters)



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$$HR = \frac{h(t, X = treatment)}{h(t, X = placebo)}$$

The group with the largest hazard is typically the numerator as hazard ratios ≥ 1 are easier to interpret.

$$HR = \frac{h_0(t) \cdot \exp\left(\sum_{i=1}^{p} \beta_i x_i^*\right)}{h_0(t) \cdot \exp\left(\sum_{i=1}^{p} \beta_i x_i\right)}$$
$$= \exp\left(\sum_{i=1}^{p} \beta_i \left(x_i^* - x_i\right)\right)$$

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If x_i is a (0,1) exposure variable, then $HR = e^{\beta_i}$ is the marginal effect size (in terms of HR) of an exposure - provided all other covariates are constant and there are no interaction terms.

Hazard ratio confidence intervals

A 95% confidence interval for the hazard ratio corresponding to the i^{th} covariate can be predicted using

$$\exp\left(\hat{eta}_i \pm 1.96\sqrt{\hat{Var}\hat{eta}_i}
ight)$$

where
$$s_{\hat{eta}_i} = \sqrt{\hat{Var}\hat{eta}_i}$$

This is simple to calculate when there are no interaction effects in the model, but the computational formula for the estimated standard error is more complex when there are interaction terms. Fortunately, this is already implemented in most statistical software!

Cox survival curves

From a Cox model, survival curves can be obtained which adjust for explanatory variables. A Cox hazard function can be converted to the below survival function by using

$$S(t) = \exp\left(-\int_0^t h(t)dx\right)$$

which gives us

$$S(t,X) = [s_0(t)]^{\exp\left(\sum_{i=1}^p \beta_i x_i\right)}$$

The proportional hazards assumption

HR is assumed to be constant over time. Recall that a hazard ratio does not depend on t.

$$HR = \frac{h(t, x^*)}{h(t, x)} = \exp\left(\sum_{i=1}^{p} \beta_i (x_i^* - x_i)\right)$$

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$$HR = \frac{h(t, x^*)}{h(t, x)} = \exp\left(\sum_{i=1}^{p} \beta_i (x_i^* - x_i)\right)$$

We would therefore expect

$$\frac{h(t,x^*)}{h(t,x)}$$

to not depend on time. If this property is not satisfied then we should consider alternatives such as stratified Kaplan-Meier or accelerated failure time models.

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- A popular method is via Schoenfeld residuals.
- For each predictor in a model, Schoenfeld residuals exist for each individual who experiences an event.

Suppose subject f has an event at time Y_f , then the Schoenfeld residual for a parameter, say x_1 , is $x_{f1} - W_{1f}$ where W_{1f} is a weighted average of x_1 for subjects still at risk at time Y_f . The weights used are each subject's hazard.

If the proportional hazards assumption is satisfied for a covariate then the Schoenfeld residuals for that covariate will not be related to survival time.

We implement the test as follows,

- 1. Obtain Schoenfeld residuals for each predictor
- Rank event times
- Test the correlation between Schoenfeld residuals and respective ranks

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$$h_0(t) = \frac{-d}{dt} \log e^{-\lambda t} = \lambda$$

and the hazard becomes:

$$h(t,X) = \lambda \cdot e^{\sum_{i=1}^{p} \beta_i x_i}$$



A popular choice of distribution for survival analysis is the Weibull. If survival times follow a Weibull distribution then

$$S(t) = \exp\left(-\left(\lambda t\right)^{p}\right)$$

for $p, \lambda > 0$. Noting that:

$$h(t) = \frac{-d}{dt} \log S(t)$$

we have:

$$h_0(t) = \frac{-d}{dt} - (\lambda t)^p$$
$$= p\lambda(\lambda t)^{p-1}$$

Multiplying the baseline hazard by $\theta = \exp(X\beta)$ gives:

$$h(t,X) = h_0(t)\theta = p\lambda(\lambda t)^{p-1} \cdot \theta$$
$$= p\lambda(\lambda t)^{p-1} \cdot \theta^{\left(\frac{1}{p} + 1 - \frac{1}{p}\right)}$$
$$= p\left(\lambda \theta^{\frac{1}{p}}\right) \left(\lambda \theta^{\frac{1}{p}}t\right)^{p-1}$$

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If we set $k=\lambda\theta^{\frac{1}{p}}$, we can clearly see that this is still a Weibull model which demonstrates the Weibull family is closed under proportional hazards

$$h(t,X) = pk(kt)^{p-1}$$



References I

- Cox, D. R. (1972). Regression models and life-tables. Journal of the Royal Statistical Society. Series B (Methodological), 34(2):187–220.
- Kleinbaum, D. G. and Klein, M. (2012). *Survival Analysis*. Springer New York.
- Prentice, R. L. (1982). Covariate measurement errors and parameter estimation in a failure time regression model. *Biometrika*, 69(2):331–342.