

# Problem Set 4

## Applied Stats II

Due: April 16, 2023

### Question 1

We're interested in modeling the historical causes of infant mortality. We have data from 5641 first-born in seven Swedish parishes 1820-1895. Using the "infants" dataset in the `eha` library, fit a Cox Proportional Hazard model using mother's age and infant's gender as covariates. Present and interpret the output.

```
1
2 # Load data
3 data(child)
4 head(child)
5
6 # Build a survival object out of the 'child' data.frame
7 child_surv <- with(child, Surv(enter, exit, event))
8
9 # Run a Cox Proportional Hazard regression
10 cox <- coxph(child_surv ~ m.age + sex, data = child)
11 summary(cox)
12 drop1(cox, test = "Chisq")
13 stargazer(cox, type = "text")
14
```

We can plot this survival function using a Kaplan Meyer plot:

```
1 # Plot
2 km <- survfit(child_surv ~ 1, data = child)
3 summary(km, times = seq(0, 15, 1))
4 plot(km, main = "Kaplan-Meier Plot", xlab = "Years", ylim = c(0.7, 1))

1 # Run a Cox Proportional Hazard regression
2 cox <- coxph(child_surv ~ m.age + sex, data = child)
3 summary(cox)
4 drop1(cox, test = "Chisq")
5 stargazer(cox, type = "text")
```

Please view output on next page:

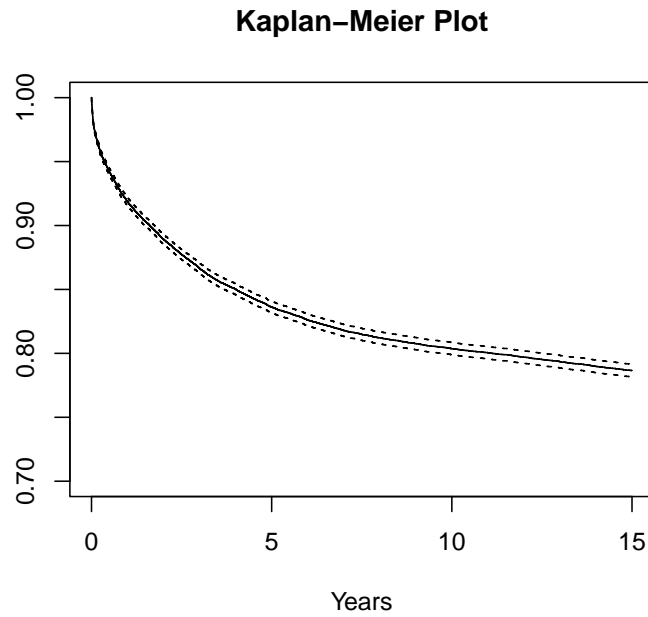


Figure 1: Child Survival Kaplan Meier plot

Table 1:

<i>Dependent variable:</i>	
	child_surv
m.age	0.008*** (0.002)
sexfemale	−0.082*** (0.027)
Observations	26,574
R <sup>2</sup>	0.001
Max. Possible R <sup>2</sup>	0.986
Log Likelihood	−56,503.480
Wald Test	22.520*** (df = 2)
LR Test	22.518*** (df = 2)
Score (Logrank) Test	22.530*** (df = 2)
<i>Note:</i>	*p<0.1; **p<0.05; ***p<0.01

**Analysis:** The results show that both mother's age and infant's gender are statistically significant predictors of child mortality, with p-values less than 0.01.

For mother's age, the coefficient estimate of 0.008 indicates that for each one-year increase in mother's age, the hazard of child mortality increases by a factor of 1.008 ( $\exp(0.008)$ ).

For infant's gender, the coefficient estimate of -0.082 indicates that the hazard of mortality is lower for female infants than for male infants by a factor of 0.921 ( $\exp(-0.082)$ ).

The `drop1()` function allows us to assess the model quality by removing each predictor one at a time from the model and comparing the fit of the reduced model to the original model using an LRT.

```
> drop1(cox, test = "Chisq")
Single term deletions

Model:
child_surv ~ m.age + sex
Df    AIC      LRT Pr(>Chi)
<none>    113011
m.age    1 113022 12.7946 0.0003476 ***
sex      1 113018  9.4646 0.0020947 **
---
Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

Here we can see that removing either mother's age or infant's gender from the model leads to a significant deterioration in the model fit based on the likelihood ratio test. This indicates that both predictors are important in predicting child survival.