

Problem Set 4

Applied Stats II

Due: April 4, 2022

Instructions

- Please show your work! You may lose points by simply writing in the answer. If the problem requires you to execute commands in R, please include the code you used to get your answers. Please also include the .R file that contains your code. If you are not sure if work needs to be shown for a particular problem, please ask.
- Your homework should be submitted electronically on GitHub in .pdf form.
- This problem set is due before class on Monday April 4, 2022. No late assignments will be accepted.
- Total available points for this homework is 80.

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.

```
Installing packages
install.packages("eha")
install.packages("stargazer")
library(stargazer)
library(eha)
library(survival)
library(ggplot2)
library(tidyverse)
data("infants")
```

```
summary(infants)
```

```
fitting a cox model with infants gender and mothers age as covariates
add-mor j- coxph(Surv(enter, exit, event) ~ sex + age, data = infants)
summary(add-mor)
```

There is a 0.49 decrease in the expected log
of the hazard of male infants compared to females,
holding mother's age constant.

There is a 0.04 decrease in the expected log
of the hazard for infants of difference
in mother's age, holding the infants sex constant.

Gender hazard ratio

```
exp(-0.48518)
```

```
0.6155864
```

The hazard ratio of male infants

is 0.62 that of female infants,

ie. male infants are less likely to die

(62 male infants die for every 100 female infants;

male deaths are 38 percent lower)

where mother's age is equal across infants.

At a given instant in time,

male infants are 0.616 times as likely to die

as females within a 95 percent confidence interval

(0.2587, 1.465), adjusting for mother's age.

From likelihood ratio test,

we can see that the model

is not statistically significant

since pvalue = 0.4.

pvalue needs to be less than 0.05 to be significant

Neither sex nor age are significant to the model, since pvalues \geq 0.05.

```
drop1(add-mor, test = "Chisq")
```

Summary of finding above, no new information. Tidies the output.

```
stargazer(add-mor, type = "text")
```

Adding an interaction

```
cox.int <- coxph(Surv(enter, exit, event) ~ sex * age, data = infants)
```

```
summary(cox.int)
```

Read for 3 'coef' values in output

There is a 1.89 increase in the expected

log of the hazard of male infants compared to females,

holding mother's age constant.

There is a -0.001519 decrease in the expected log

of the hazard for infants

of difference in mother's age,

holding the infants sex constant.

there is a -0.09128 decrease in the expected log

of the hazard of male infants

when both the gender and the mothers

age are taken into account

Model not sig, since $p \geq 0.05$.

```
drop1(cox.int, test = "Chisq")
```

Interaction term not sig since p greater than 0.05.

```
stargazer(cox.int, type = "text")
```

```

plot of the cumulative hazard function
plot-CoxPH j- coxreg(Surv(enter, exit, event) sex + age, data = infants)
plot(plot-CoxPH)

```

```

Plotting model
cox-fit j- survfit(add-mor)
autoplot(cox-fit)

```

```

newdat j- with(infants, data.frame( sex = c("boy", "girl"), age ="26" ) )
newdat

```

All of the code above allows us to produce figure 1 plot

```

plot(survfit(add-mor, newdata = newdat), xscale = 12,
conf.int = T,
ylim = c(0.6, 1),
col = c("red", "blue"),
xlab = "Time",
ylab = "Survival proportion",
main = "")
legend("bottomleft",
legend=c("boy", "girl"),
lty = 1,
col = c("red", "blue")
, text.col = c("red", "blue"))

```

running this code you get the following plots:
if you include the following code instead
of running the original packages it allows

autoplot to work and you get the autoplot
figure 2 that is shown below

```

rm(list=ls())
detach all libraries
detachAllPackages j- function()
basic.packages j- c("package:stats"
"package:graphics", "package:grDevices",
"package:utils", "package:datasets",
"package:methods", "package:base")
package.list j-
search()[ifelse(unlist(gregexpr("package:",
search()))==1, TRUE, FALSE)]
package.list j- setdiff(package.list, basic.packages)
if (length(package.list)>0) for (package in package.list) detach(package, character.only=TRUE)

detachAllPackages()

load libraries

```

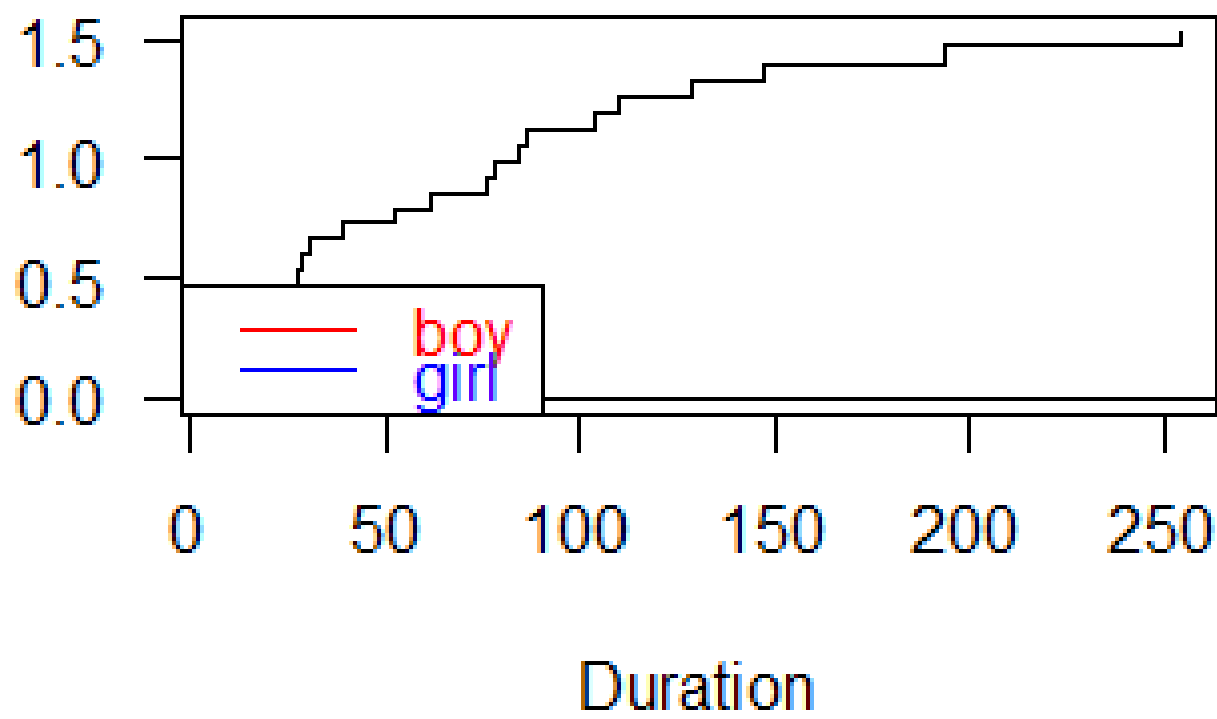


Figure 1: Caption

```
pkgTest <- function(pkg)
new.pkg <- pkg[!(pkg %in% installed.packages()[, "Package"])]
if (length(new.pkg))
install.packages(new.pkg, dependencies = TRUE)
sapply(pkg, require, character.only = TRUE)
```

here is where you load any necessary packages

ex: stringr

```
lapply(c("stringr"), pkgTest)
```

```
lapply(c("survival", "eha", "tidyverse", "ggfortify", "stargazer"), pkgTest)
```

1 Introduction

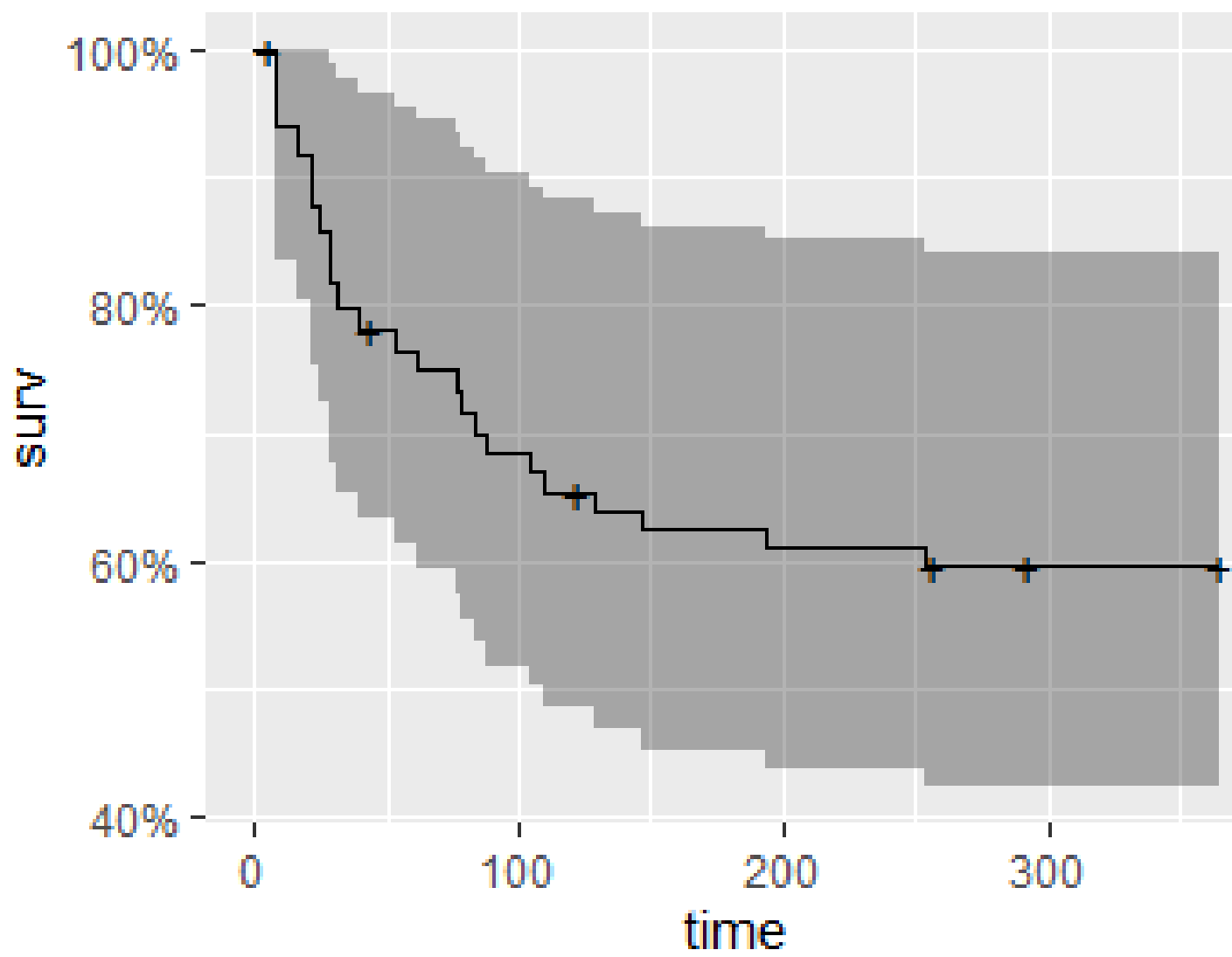


Figure 2: Caption