

Survival Models: Week 6

Example: Partial Likelihood

The times of death, or censoring (*), following a particular operation are given below. Information on each person's weight and gender is also provided.

	time	Gender	Weight	$\beta^T x$
	4	M	60	$\beta_1 + 60\beta_2$
censored ←	5*	F	80	$80\beta_2$
tie death {	6	F	50	$50\beta_2$
	6	M	70	$\beta_1 + 70\beta_2$
	7*	M	100	$\beta_1 + 100\beta_2$

Note: Gender will be coded as 1 for Male and 0 for female.

Example: Partial Likelihood

Based on the information in the table, the first two terms in the PL will be:

$$\frac{\exp(\beta_1 + 60\beta_2)}{\exp(\beta_1 + 60\beta_2) + \exp(80\beta_2) + \exp(50\beta_2) + \exp(\beta_1 + 70\beta_2) + \exp(\beta_1 + 100\beta_2)}$$

and,

$$\frac{\exp((1 + 0)\beta_1 + (50 + 70)\beta_2)}{[\exp(50\beta_2) + \exp(\beta_1 + 70\beta_2) + \exp(\beta_1 + 100\beta_2)]^2}.$$

R Example - Recidivism

Call:

```
coxph(formula = Surv(week, arrest) ~ fin + age + race + wexp +  
      mar + prio, data = Rossi)
```

n= 432, number of events= 114

	coef	exp(coef)	se(coef)	z	Pr(> z)	
finyes	-0.37352	0.68831	0.19082	-1.957	0.050295	.
age	-0.05640	0.94516	0.02184	-2.583	0.009796	**
raceother	-0.30983	0.73357	0.30780	-1.007	0.314133	
wexpyes	-0.15331	0.85786	0.21218	-0.723	0.469957	
marnot married	0.44339	1.55799	0.38136	1.163	0.244958	
prio	0.09336	1.09785	0.02832	3.296	0.000981	***

$$\lambda(t, \beta) = \lambda_0(t) \cdot e^{\beta^T x}$$

\downarrow \downarrow
 $1 \times p$ $p \times 1$

e.g. $\lambda_1(t; \beta) = \lambda_0(t) \cdot e^{\beta^T x_1}$

$\lambda_2(t; \beta) = \lambda_0(t) \cdot e^{\beta^T x_2}$

Holding either variable constant let the first variable $x_{2,1} = x_{1,1} + 1$

\downarrow first variable
 2nd individual

\downarrow first variable
 1st individual

$$\frac{\lambda_1(t; \beta)}{\lambda_2(t; \beta)} = \frac{e^{\beta_1 x_{2,1}}}{e^{\beta_1 x_{1,1}}} = \boxed{e^{\beta_1}}$$

1 unit \uparrow in x_1 , will increase $\lambda(t; \beta)$ to e^{β_1} times

\downarrow
first variable

For example

$e^{\beta_1} \approx 69\%$ for "fin"

compared with fin=0, fin=1

$\lambda(t, \beta)$ will be decreased to

$e^{\beta_1} = 69\%$

$\lambda_2(t; \beta) \leftarrow \text{fin}=1$
 $\lambda_1(t; \beta) \leftarrow \text{fin}=0$
 $= 69\%$

alternatively, $\lambda(t, \beta)$ will be decreased by $1 - e^{\beta_1} = 31\%$
 b/c $\lambda_2 - \lambda_1 = -31\% \lambda_1$

R Example - Recidivism

The *relative risk* for a not-married individual relative to a married individual is $\exp(0.44) = 1.55$.

⇒ similar effect to maximize likelihood

The parameter estimates obtained from maximizing the PL have an approximately Normal sampling distribution with mean β and covariance matrix that can be found by inverting the information matrix. Tests of whether particular covariates are important can be conducted by comparing the statistic:

null: $\beta = 0$

alt: $\beta \neq 0$

$$\Rightarrow Z = \frac{\hat{\beta} - \beta}{SE(\hat{\beta})}$$

$$\frac{\hat{\beta}}{SE(\hat{\beta})} \sim Z\text{-statistic}$$

to the $N(0, 1)$ distribution. For example, the test-statistic for *married* is 1.16 with a p-value of 0.24. This suggests that married is not an important explanatory variable.

Overall Significance

Testing the overall significance of the Cox model equates to testing the following:

$$\text{null: } \beta_1 = \beta_2 = \dots \beta_p = 0$$

$$\text{alt: at least one of the } \beta_i \neq 0 \ i = 1, \dots, p.$$

One way to conduct this test is too look at the likelihoods correspond to the models specified under the null and alternative. Assuming the null hypothesis is correct the quantity $-2(LL(null) - LL(alt))$ has an approximate chi-square distribution with degrees of freedom equal to the difference in the number of parameters in the models specified under the null and alternative.

R Example - Recidivism

```
library(RcmdrPlugin.survival)
data(Rossi) #see J Fox notes for more details
cox.null<-coxph(Surv(week,arrest)~1,data=Rossi)
cox.alt<-coxph(Surv(week,arrest)~fin+age+race+wexp+mar+prio,data=Rossi)
cox.null$loglik
[1] -675.3806
cox.alt$loglik
[1] -675.3806 -658.8411
TS<--2*(cox.null$loglik[1]-cox.alt$loglik[2])
TS
[1] 33.07898
1-pchisq(TS,df=6)
[1] 1.012523e-05
```


R Example - Recidivism

```
> summary(cox.alt)
Call:
coxph(formula = Surv(week, arrest) ~ fin + age + race + wexp +
      mar + prio, data = Rossi)

n= 432, number of events= 114
```

```
#I have deleted some output here##
```

	exp(coef)	exp(-coef)	lower .95	upper .95
finyes	0.6883	1.4528	0.4735	1.0005
age	0.9452	1.0580	0.9056	0.9865
raceother	0.7336	1.3632	0.4013	1.3410
wexpyes	0.8579	1.1657	0.5660	1.3003
marnot married	1.5580	0.6419	0.7378	3.2898
prio	1.0979	0.9109	1.0386	1.1605

```
Concordance= 0.642 (se = 0.027 )
Rsquare= 0.074 (max possible= 0.956 )
Likelihood ratio test= 33.08 on 6 df, p=1.013e-05
Wald test = 32.01 on 6 df, p=1.625e-05
Score (logrank) test = 33.43 on 6 df, p=8.68e-06
```

R Example - Recidivism

How about testing whether *race* and *married* are needed in a model that contains the other four covariates? This corresponds to the following test:

$$\text{null: } \beta_{\text{race}} = \beta_{\text{married}} = 0$$

$$\text{alt: at least one of } \beta_{\text{race}} \text{ or } \beta_{\text{married}} \neq 0.$$

This test is easily conducted using a likelihood ratio test. The following slide shows how the test is conducted. Note: in this R code we need to make sure that *race* and *married* are included as the last two variables in the `cox.alt` model.

R Example - Recidivism

```
cox.null<-coxph(Surv(week,arrest)~fin+age+wexp+prio,data=Rossi)
cox.alt<-coxph(Surv(week,arrest)~fin+age+wexp+prio+race+mar,data=Rossi)
cox.null$loglik
[1] -675.3806 -660.2845
cox.alt$loglik
[1] -675.3806 -658.8411
TS<--2*(cox.null$loglik[2]-cox.alt$loglik[2])
TS
[1] 2.886664
1-pchisq(TS,df=2)
[1] 0.2361397
anova(cox.alt,cox.null)
Analysis of Deviance Table

Cox model: response is  Surv(week, arrest)
Model 1: ~ fin + age + wexp + prio + race + mar
Model 2: ~ fin + age + wexp + prio
   loglik  Chisq Df P(>|Chi|)
1 -658.84
2 -660.28 2.8867  2    0.2361
```

Survival Curves

Estimated survival curves can also be produced based on a fitted Cox regression model. These estimates are derived using the following relationship:

$$S_0(t)^{\exp(\beta^T x)},$$

and plugging in the coefficient estimates obtained from the Cox regression model. The baseline survival function, $S_0(t)$, is also estimated based on the Cox regression (See O'Neill notes page 27 for details). Often estimated survival curves are plotted based on the covariates being set at their mean or median levels.

#R code for plot on next slide.

```
cox.full<-coxph(Surv(week,arrest)~fin+age+race+wexp+mar+prio,data=Rossi)  
plot(survfit(cox.full))
```

R Example - Recidivism

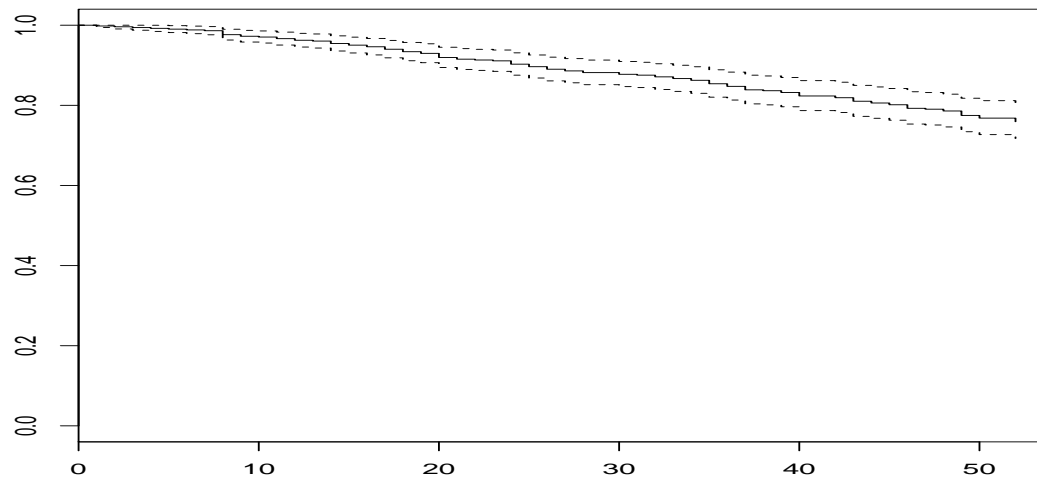


Figure 1: Estimated survival curves for Recidivism data. All covariates at their mean levels.

R Example - Recidivism

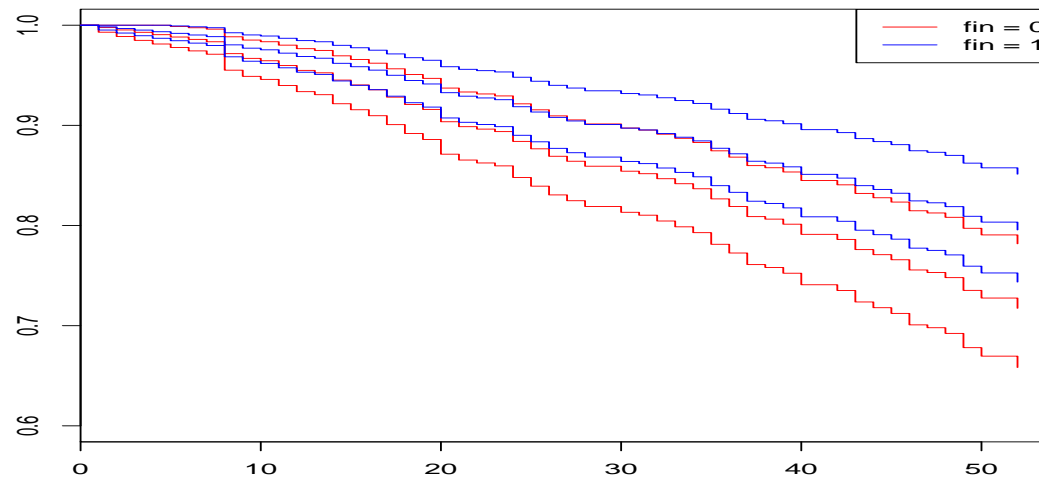


Figure 2: Estimated survival curves based on financial aid for Recidivism data. All other covariates at their mean levels.

R Example - Recidivism

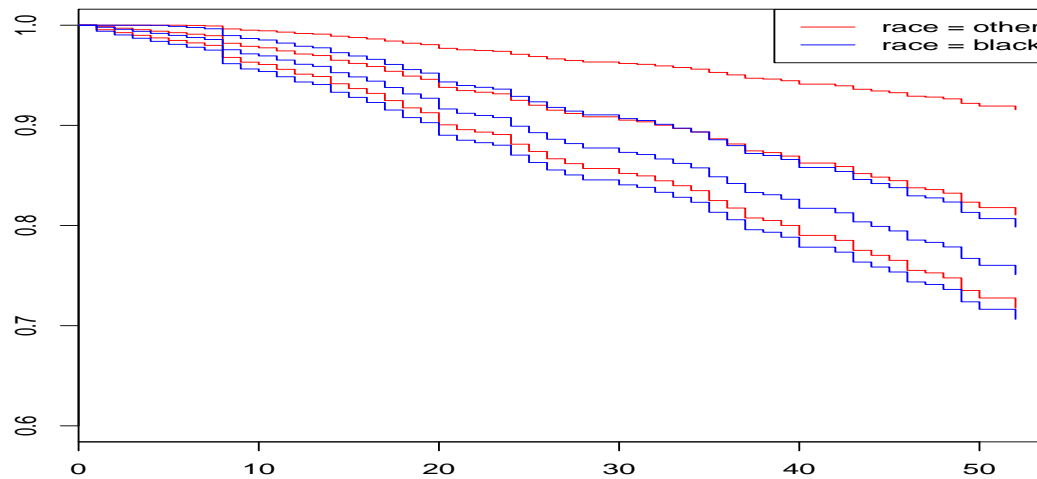


Figure 3: Estimated survival curves based on race for Recidivism data. All other covariates at their mean levels.

R Example - Addict

In week 5 we produced the following survival curves using KM.

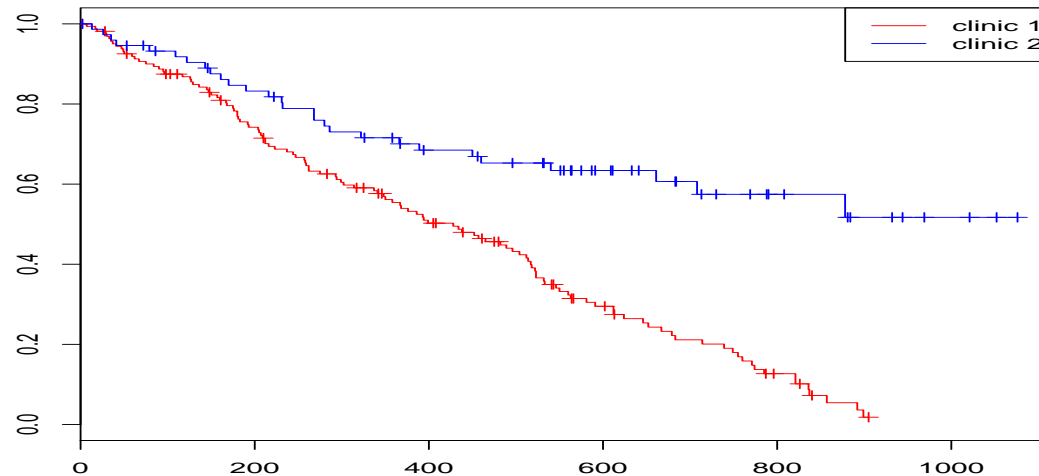


Figure 4: Estimated KM survival curves for Addict data

R Example - Addict

Fitting a Cox regression model to this data gives the following output:

```
attach(addict)
cox.mod<-coxph(Surv(time,status)~as.factor(clinic)+prison+dose,data=addict)
summary(cox.mod)
Call:
coxph(formula = Surv(time, status) ~ as.factor(clinic) + prison +
      dose, data = addict)
```

n= 238, number of events= 150

	coef	exp(coef)	se(coef)	z	Pr(> z)	
as.factor(clinic)2	-1.009896	0.364257	0.214889	-4.700	2.61e-06	***
prison	0.326555	1.386184	0.167225	1.953	0.0508	.
dose	-0.035369	0.965249	0.006379	-5.545	2.94e-08	***

Signif. codes: 0 *** 0.001 ** 0.01 * 0.05 . 0.1 1

	exp(coef)	exp(-coef)	lower .95	upper .95
as.factor(clinic)2	0.3643	2.7453	0.2391	0.5550
prison	1.3862	0.7214	0.9988	1.9238
dose	0.9652	1.0360	0.9533	0.9774

Concordance= 0.665 (se = 0.026)
Rsquare= 0.238 (max possible= 0.997)
Likelihood ratio test= 64.56 on 3 df, p=6.228e-14
Wald test = 54.12 on 3 df, p=1.056e-11
Score (logrank) test = 56.32 on 3 df, p=3.598e-12

R Example - Addict

Is there an interaction between prison and dose?

```
cox.mod<-coxph(Surv(time,status)~as.factor(clinic)+prison+dose+I(prison*dose),data=addict)
summary(cox.mod)
```

Call:

```
coxph(formula = Surv(time, status) ~ as.factor(clinic) + prison +
      dose + I(prison * dose), data = addict)
```

n= 238, number of events= 150

	coef	exp(coef)	se(coef)	z	Pr(> z)	
as.factor(clinic)2	-0.995839	0.369413	0.215784	-4.615	3.93e-06	***
prison	-0.206669	0.813289	0.766871	-0.269	0.788	
dose	-0.038905	0.961842	0.008086	-4.811	1.50e-06	***
I(prison * dose)	0.009084	1.009126	0.012727	0.714	0.475	

Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

R Example - Addict

Estimated effect of dose is -0.038905 if prison=0 and $-0.038905 + 0.009084$ if prison=1. However, the interaction term is not significant (p-value=0.475).

We could also include a squared term to model the effect of dose. Perhaps the impact of dose increases up to a certain point and then “flattens out”.

```
cox.mod<-coxph(Surv(time,status)~as.factor(clinic)+prison+dose+I(dose^2),data=addict)
summary(cox.mod)
```

R Example - Addict

Compare KM estimate Cox estimate

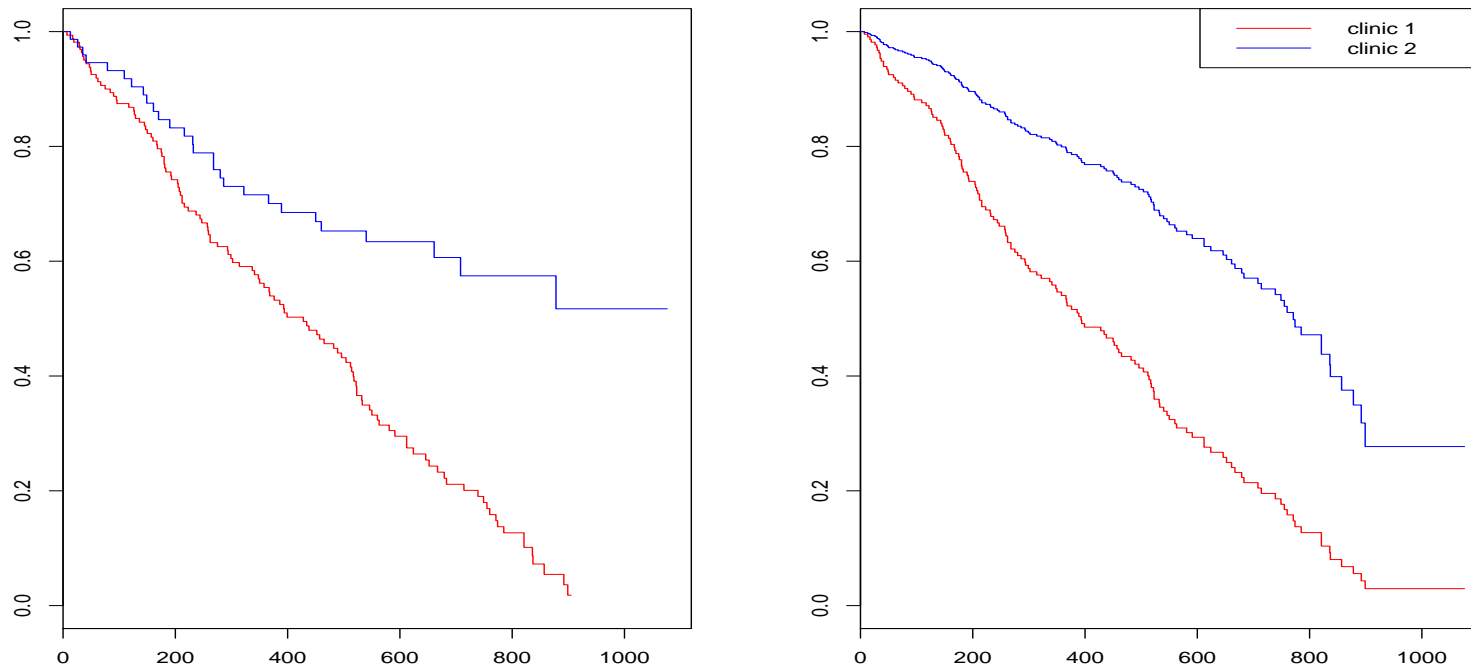


Figure 5: Estimated KM (left) and Cox (right) survival curves for Addict data

Diagnostic Checks

It is possible to conduct diagnostic checks to see whether the following two assumptions hold:

- Proportional Hazards assumption
- Linearity.
- Influential observations