GSERM - Oslo 2019 Introduction to Survival Data

January 9, 2019 (afternoon session)

Survival Analysis

- Models for time-to-event data.
- Roots in biostats/epidemiology, plus engineering, sociology, economics.
- Examples...
 - Political careers, confirmation durations, position-taking, bill cosponsorship, campaign contributions, policy innovation/adoption, etc.
 - Cabinet/government durations, length of civil wars, coalition durability, etc.
 - War duration, peace duration, alliance longevity, length of trade agreements, etc.
 - · Strike durations, work careers (including promotions, firings, etc.), criminal careers, marriage and child-bearing behavior, etc.

Characteristics of Time-To-Event Data

- *Discrete* events (i.e., not continuous),
- Take place over time,
- May not (or never) experience the event (i.e., possibility of censoring).

Survival Data Basics: Terminology

 Y_i = the duration until the event occurs,

 Z_i = the duration until the observation is "censored"

 $T_i = \min\{Y_i, Z_i\},$

 $C_i = 0$ if observation i is censored, 1 if it is not.

Survival Data Basics: The Density

$$f(t) = \Pr(T_i = t)$$

density function, probability T_i takes particular value t

Issues:

- $T_i = t$ iff $T_i > t 1$, t 2, etc.
- $C_i = 0$ (censoring)

up to the point T_i = if it does not equal t_before that (you need to be 21 years unmarried to get married at 22.

Survival Data Basics: Survivor Function

cdf

$$\Pr(T_i \leq t) \equiv F(t) = \int_0^t f(t) dt$$

$$Pr(T_i \ge t) \equiv S(t) = 1 - F(t)$$

$$= 1 - \int_0^t f(t) dt$$

Survival Data Basics: The Hazard

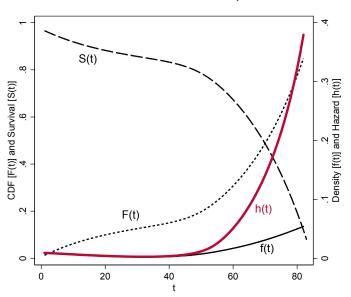
h is not a true probability, coz it can get bigger than 1 as f is bigger than S => correct term would be conditional risk.

$$Pr(T_i = t | T_i \ge t) \equiv h(t) = \frac{f(t)}{S(t)}$$

given that I have not been married until t, what is the probability that i get married at t (conditional)

$$= \frac{f(t)}{1 - \int_0^t f(t) dt}$$

Example: Human Mortality



solid right axis, dashed left axis.

Some Useful Equivalencies

$$f(t) = \frac{-\partial S(t)}{\partial t}$$

negative derivative

Implies

$$h(t) = \frac{\frac{-\partial S(t)}{\partial t}}{S(t)}$$
$$= \frac{-\partial \ln S(t)}{\partial t}$$

hazard is negative derivative of logged survivor function

More Useful Things: Integrated Hazard

Define

$$H(t) = \int_0^t h(t) dt.$$

cumulative risk.

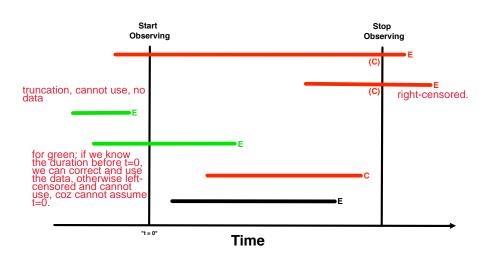
Implies

$$H(t) = \int_0^t \frac{-\partial \ln S(t)}{\partial t} dt$$
$$= -\ln[S(t)]$$

and

$$S(t) = \exp[-H(t)]$$

Censoring and Truncation





Censoring

• Defined by the researcher

(what is event, what is censoring "event", e.g. time to die, quit, get fired, etc. you need to define event of interest.)

- Conditionally independent of both T_i and X_i
- Doesn't mean that the observation provides no information

censoring is conditionally independent sounds unreasonable at first. but most covariates are typically conditionally dependent of X and you should control for it anyway. so, if you are confident in the specified model, there is also no conditional dependence of T_i.

Estimating S(t)

Kaplan-Meier estimator.

no ties = no 2 observations at the exact same time t.

absorbing events=event occurred, cant happen again.

Assume N observations, absorbing events, and no ties. Then define

 $n_t = \text{number of observations "at risk" for the event at } t$, and

 d_t = number of observations which experience the event

at time t.

Then

$$\widehat{S(t_k)} = \prod_{t \le t_k} \frac{n_t - d_t}{n_t}$$

descriptively we usually work with the survival function instead of hazard function.

proportion of at risk observations that had the event at timepoint t. for each period where no event happens, the curve stays the same, otherwise it decreases a little bit, i.e less surviving happening.

censoring changes the risk set, but not the d_t; so censoring does not change survival function. following no ties, d_t could be at most 1, but later we see d_t can also be bigger than 1.

Variance of $\widehat{S(t)}$

$$\mathsf{Var}[\widehat{S(t_k)}] = \left[\widehat{S(t_k)}\right]^2 \sum_{t \leq t_k} \frac{d_t}{n_t(n_t - d_t)}$$

Note:

- $Var[\widehat{S(t_k)}]$ is increasing in S(t),
- is also increasing in d_t , but
- is decreasing in n_t .

Estimating H(t)

"Nelson-Aalen":

$$\widehat{H(t_k)} = \sum_{t < t_k} \frac{d_t}{n_t}$$

it goes up by small amounts as d_t may be 1 and n_t large

...which gives an alternative estimator for the survival function equal to:

$$\widehat{S(t_k)} = \exp[-\widehat{H(t_k)}]$$

$$= \exp\left[-\sum_{t \le t_k} \frac{d_t}{n_t}\right]$$

kaplan-meier and nelson-aalen are asympotically equivalent, but as estimates they may be slightly different in selected samples

Bivariate Hypothesis Testing

(similar to chi-squared test)

	Treatment	Placebo	Total
Event	d_{1t}	d_{0t}	d_t
No Event	$n_{1t}-d_{1t}$	$n_{0t}-d_{0t}$	$n_t - d_t$
Total	n_{1t}	n _{0t}	n _t

Log-Rank Test:

$$Q = \frac{\left[\sum (d_{1t} - \frac{n_{1t}d_t}{n_t})\right]^2}{\left[\frac{n_{1t}n_{0t}d_t(n_t - d_t)}{n_t^2(n_t - 1)}\right]}$$
$$\sim \chi_1^2$$

A Diversion: Survival Models and Counting Processes

Assume

- Event is absorbing,
- Y_i is duration to the event
- Z_i is duration to censoring
- Observe $T_i = \min(Y_i, Z_i)$, and
- C_i:
 - $C_i = 0$ if $T_i = Z_i$,
 - $C_i = 1$ if $T_i = Y_i$.
- $T_i \neq T_j \ \forall \ i \neq j \ (\text{no "ties"})$

Three Key Variables

1. Counting Process Indicator:

$$N_i(t) = I(T_i \leq t, C_i = 1)$$

zero until event occured, then 1 for the rest.

2. Risk Indicator:

$$R_i(t) = I(T_i > t)$$

1 until event occuered, then 0 for the rest, overlaps with 1.

3. Intensity Process:

$$\lambda_i(t) dt = R_i(t)h(t)$$

hazard as long as you are at rest, if event occured, it goes to

Additional Things

With

$$\Lambda_i(t) = \int_0^t \lambda_i(t) dt$$

we can think of

$$N_i(t) = \Lambda_i(t) + M_i(t)$$

or

$$M_i(t) = N_i(t) - \Lambda_i(t).$$

kinda like "Residual = Observed - Expected"

it is a martingale process, so all the math can be used.

Martingales!

(memoryless process)

$$E(X_{t+s}|X_0, X_1, ...X_i, ...X_t) = X_t \ \forall \ s > 0$$

Data Structure and Organization: Non-Time-Varying

id	durat	censor	timein	timeout	Х
1	4	0	30	34	0.12
2	2	1	12	14	0.19
3	5	1	5	10	0.09
N	10	1	21	31	0.22

Time-Varying Data

id	durat	censor	${\tt timein}$	${\tt timeout}$	Х	Z
1	1	0	30	31	0.12	331
1	2	0	31	32	0.12	412
1	3	0	32	33	0.12	405
1	4	0	33	34	0.12	416
2	1	0	12	13	0.19	226
2	2	1	13	14	0.19	296
3	1	0	5	6	0.09	253
3	2	0	6	7	0.09	311
3	3	0	7	8	0.09	327
3	4	0	8	9	0.09	344
3	5	1	9	10	0.09	301

Analyzing Survival Data in R

```
survival object (non-time-varying):
library(survival)
NonTV<-read.csv(NonTVdata.csv)
NonTV.S<-Surv(NonTV$duration, NonTV$censor)

survival object (time-varying):
TV<-read.csv(TVdata.csv)
TV.S<-Surv(TV$starttime, TV$endtime, TV$censor)</pre>
```

An Example

OECD Cabinet survival [Strom (1985); King et al. (1990)],

N = 314 cabinets in 15 countries

Outcome: Duration of cabinet, in months

Covariates (all non-time varying):

- Fractionalization
- Polarization
- · Formation Attempts
- · Investiture
- · Numerical Status
- · Post-Election
- · Caretaker

Also: Indicator for whether the cabinet ended within 12 months of the end of the "constitutional inter-election period" $(\rightarrow$ censored)

(coz cabinet may call early elections shortly before mandated end)

KABL Data

> head(KABL)

	id	country	durat	ciep12	fract	polar	format	invest	numst2	${\tt eltime2}$	caretk2
1	1	1	0.5	1	656	11	3	1	0	1	0
2	2	1	3.0	1	656	11	2	1	1	0	0
3	3	1	7.0	1	656	11	5	1	1	0	0
4	4	1	20.0	1	656	11	2	1	1	0	0
5	5	1	6.0	1	656	11	3	1	1	0	0
6	6	1	7.0	1	634	6	4	1	1	1	0

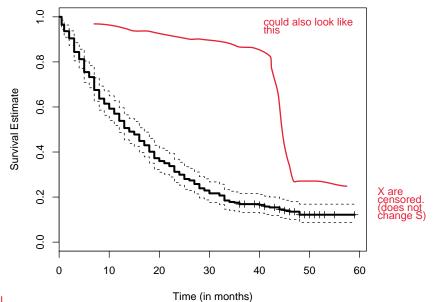
- > KABL.S<-Surv(KABL\$durat,KABL\$ciep12)
- > KABL.S[1:50,]

```
[1] 0.5 3.0 7.0 20.0 6.0 7.0 2.0 17.0 27.0 49.0+
        29.0 49.0+ 6.0
[11]
    4.0
                        23.0 41.0+ 10.0
                                       12.0
                                           2.0 33.0
[21]
    1.0 16.0 2.0
                    9.0
                        3.0 5.0 5.0 6.0 45.0+ 23.0
[31] 41.0
         7.0 49.0+ 46.0
                        9.0 51.0+ 10.0
                                       32.0
                                            28.0
                                                 3.0
[41] 53.0+ 17.0 59.0+ 9.0 52.0+ 3.0 23.0
                                       33.0
                                            1.0 30.0
```

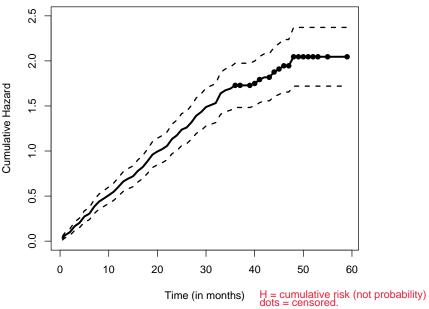
Example survfit Object

```
> KABL.fit<-survfit(KABL.S~1)
> str(KABL.fit)
List of 13
$ n : int 314
$ time : num [1:54] 0.5 1 2 3 4 5 6 7 8 9 ...
$ n.risk : num [1:54] 314 303 294 284 265 255 237 230 212 200 ...
$ n.event : num [1:54] 11 9 10 19 10 18 7 18 12 7 ...
$ n.censor : num [1:54] 0 0 0 0 0 0 0 0 0 ...
$ surv : num [1:54] 0.965 0.936 0.904 0.844 0.812 ...
$ type : chr "right"
$ std.err : num [1:54] 0.0108 0.0147 0.0183 0.0243 0.0271 ...
$ upper : num [1:54] 0.986 0.964 0.938 0.885 0.856 ...
$ lower : num [1:54] 0.945 0.91 0.873 0.805 0.77 ...
$ conf.type: chr "log"
$ conf.int : num 0.95
$ call : language survfit(formula = KABL.S ~ 1)
- attr(*, "class")= chr "survfit"
```

Plotting $\widehat{S(t)}$



Plotting $\widehat{H(t)}$



Comparing $\widehat{S(t)}$ s

```
Log-rank test:
```

> survdiff(KABL.S~invest,data=KABL,rho=0)

Call:

survdiff(formula = KABL.S ~ invest, data = KABL, rho = 0)

N Observed Expected (0-E)^2/E (0-E)^2/V invest=0 172 137 178.7 9.72 30.5 invest=1 142 134 92.3 18.81 30.5

Chisq= 30.5 on 1 degrees of freedom, p= 3.26e-08

formal test if these two survival functions are statistically different. you can also split the data and estimate two survival functions.

Comparing $\widehat{S(t)}$ s

