Introduction to survival analysis

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Plan for today

Gentle introduction to survival analysis.

Main source: Harrell, F. E., Jr. (2015). *Regression Modeling Strategies*, 2nd edition. Springer

Chapters: 17, 18, and 20.

Survival analysis (SA)

Data: For which the *time until the event* is of interest.

► This goes beyond *logistic regression*, which focuses on the *occurrence* of the event.

Outcome variable:

- ▶ T = Time until the event.
- ▶ Often referred to as *survival time, failure time,* or *event time*.

Examples

Survival time: Time until...

▶ death, desease, relapse.

Failure time: Time until...

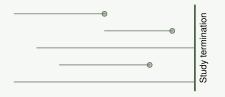
▶ product malfunction.

Event time: Time until...

▶ graduation, marriage, divorce.

Advantages of SA over typical regression models

 SA allows modeling units that did not fail up to data collection (*censored on the right* data).



- Regression could be considered to model the expected survival time. *But*:
- ► Survival time is often not normally distributed.
- P(survival > t) is often more interesting than $\mathbb{E}($ survival time).

Censoring

Some subjects:

- ▶ Did not experiment the event up to the end of data collection;
- ► Withdrew from study;
- ► Were lost to follow-up.

These data are right-censored.

Define random variables for the *i*th subject:

•
$$T_i$$
 = time to event

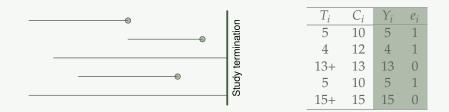
• C_i = censoring time

• e_i = event indicator = $\begin{cases}
1 & \text{if event is observed } (T_i \leq C_i) \\
0 & \text{if event is not observed } (T_i > C_i)
\end{cases}$

• $Y_i = \min(T_i, C_i)$ = what occurred first (failure or censoring)

Variables $\{Y_i, e_i\}$ include all the necessary information.

Typical data set



Observe the flexibility of SA data:

- ▶ Subjects may join the study at different moments.
- Censoring times may differ among subjects.

 $\{Y_i, e_i\}$ does include all the necessary information.

But, *assumption*: Censoring is **non-informative**, i.e., it is independent of the risk of the event.

Three main functions

T =time until event.

► Survival function:

$$S(t) = P(T > t) = 1 - F(t),$$

where $F = P(T \le t)$ is the distribution function of *T*.

Cumulative hazard function:

$$\Lambda(t) = -\log(S(t))$$

► Hazard function:

$$\lambda(t) = \Lambda'(t)$$

Survival function

$$S(t) = P(T > t) = 1 - F(t)$$

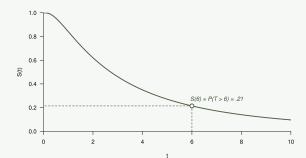
Example:

If event = death, then S(t) = prob. death occurs after time t.

Properties:

• $S(0) = 1, S(\infty) = 0.$

▶ Non-increasing function of *t*.



Cumulative hazard function

$$\Lambda(t) = -\log(S(t))$$

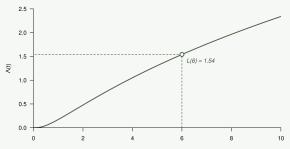
Idea:

Accumulated risk up until time *t*.

Properties:

 $\blacktriangleright \Lambda(0) = 0.$

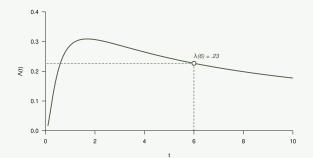
▶ Non-decreasing function of *t*.



Hazard function

$$\lambda(t) = \Lambda'(t)$$

Idea: Instantaneous event rate at time *t*.



Relation between the three functions

All functions are related:

Any two functions can be derived from the third function.

► The three functions are equivalent ways of describing the same random variable (*T* = time until event).

More generally, all the following functions give mathematically equivalent specifications of the distribution of *T*:

- F(t): Distribution function
- f(t): Density function
- \blacktriangleright *S*(*t*): Survival function
- ► $\lambda(t)$: Hazard function
- $\Lambda(t)$: Cumulative hazard function.

Examples

Next are two primary examples of parametric survival distributions:

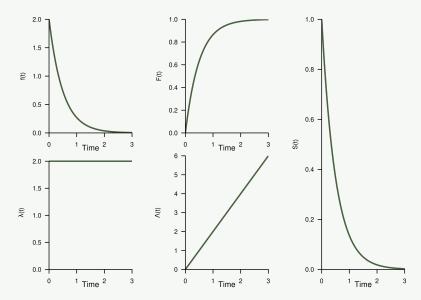
- ▶ the exponential distribution;
- ► the Weibull distribution.

These models (still) include no covariates, thus:

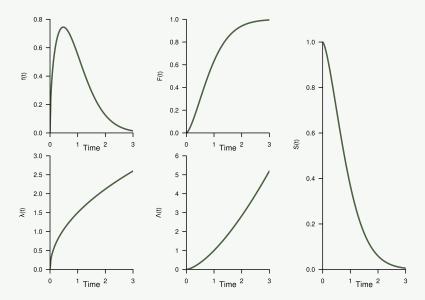
► Each subject in the sample is assumed to have the same distribution of *T*.

No formulas for now. Instead: Let's plot.

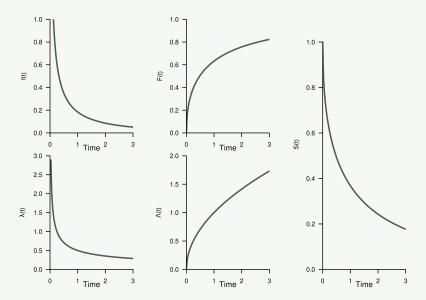
Exponential survival distribution



Weibull survival distribution (I)

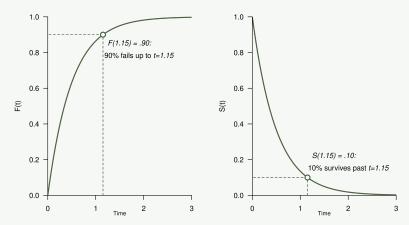


Weibull survival distribution (II)



Quantiles

Q: What is the time by which (100q)% of the population will fail? *A*: Value t_q such that $F(t_q) = q$, or, equiv., $S(t_q) = 1 - q$.



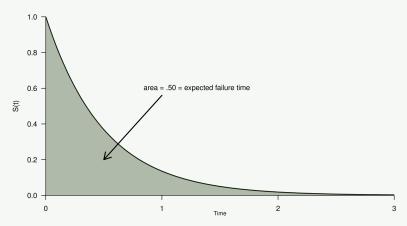
In particular, median survival time = $t_{.50}$.

Expected failure time

(Note: *T* is skewed, so the mean is not the best summary. Better use median.)

Q: What is the expected failure time?

A: It is the area under the survival function.



Various estimation approaches

There are several options available to estimate the survival function (and friends).

Here we will briefly go through only a few:

- ▶ Not parametric and homogeneous (i.e., without predictors):
 - ✓ Kaplan-Meier estimator
 - $\checkmark~$ Altschuler-Nelson estimator.
- ► Parametric:
 - ✓ Homogeneous (i.e., no predictors): Exponential, Weibull, normal, logistic, log-normal, log-logistic,...
 - ✓ Proportional hazards models
 - ✓ Semi-parametric:
 - Cox proportional hazards regression model.

After a brief intro to each, I will use them all on an empirical dataset.

Kaplan-Meier estimator

- ► Also known as the *product-limit* estimator.
- ▶ Non parametric, and super simple to do even manually.
- ▶ Key ingredient: *Conditional probabilites*.

Assume t = 0, 1, 2, ...We have that S(0) = P(T > 0) = 1. For $t \ge 1$ we then have that

$$P(T > t | T > t - 1) = \frac{P(T > t, T > t - 1)}{P(T > t - 1)} = \frac{P(T > t)}{P(T > t - 1)}$$

and so

$$P(T > t) = P(T > t - 1) \times P(T > t | T > t - 1),$$

or in terms of the survival function,

$$S(t) = S(t-1) \times P(T > t | T > t-1)$$

$$\overline{S(t) = S(t-1) \times (1 - P(T \le t | T > t-1))}$$

Kaplan-Meier estimator – Example

Data: Seven subjects; failure times T = 1, 3, 3, 3+, 6+, 9, 10+.

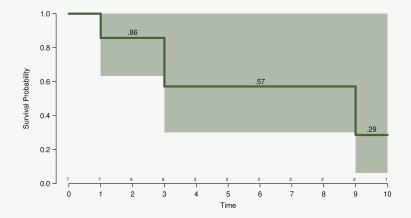
Day	No. subjects	Deaths	Censored	$S(t) = S(t-1) \times$
	at risk			$\times (1 - P(T \le t T > t - 1))$
1	7	1	0	$1 \times (1 - 1/7) = 6/7$
3	7 - (1 + 0) = 6	2	1	$6/7 \times (1 - 2/6) = 4/7$
6	6 - (2 + 1) = 3	0	1	$4/7 \times (1 - 0/3) = 4/7$
9	3 - (0 + 1) = 2	1	0	$4/7 \times (1 - 1/2) = 2/7$
10	2 - (1 + 0) = 1	0	1	$2/7 \times (1 - 0/1) = 2/7$

Hence:

$$S(t) = \begin{cases} 1, & 0 \le t < 1\\ 6/7 = .86, & 1 \le t < 3\\ 4/7 = .57, & 3 \le t < 9\\ 2/7 = .29, & 9 \le t < 10\\ \text{undefined}^*, & t \ge 10 \end{cases}$$

*Not everyone failed by t = 10, so we cannot tell what happened after that.

Kaplan-Meier estimator – Example



Altschuler-Nelson estimator

- ▶ Non parametric, also simple.
- Similar to Kaplan-Meier, but based on $\Lambda(t)$.

Recall that $\Lambda(t) =$ accumulated risk up until time *t*. Hence it makes sense to estimate $\Lambda(t)$ by

$$\widehat{\Lambda}(t) = \sum_{i:t_i \le t} \frac{\text{\# failures at } t_i}{\text{\# subjects at risk at } t_i}$$

Then,

$$\widehat{S}(t) = \exp(-\widehat{\Lambda}(t)).$$

Interesting property: $\sum_{i} \widehat{\Lambda}(Y_i) = \text{total number of events.}$

Altschuler-Nelson estimator – Example

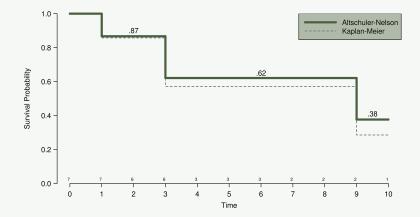
Data: Seven subjects; failure times T = 1, 3, 3, 3+, 6+, 9, 10+.

Day	No. subjects	Deaths	Censored	$\Lambda(t)$
	at risk			
1	7	1	0	1/7
3	7 - (1 + 0) = 6	2	1	1/7 + 2/6 = 10/21
6	6 - (2 + 1) = 3	0	1	10/21 + 0/3 = 10/21
9	3 - (0 + 1) = 2	1	0	10/21 + 1/2 = 41/42
10	2 - (1 + 0) = 1	0	1	41/42 + 0/1 = 41/42
		$\sum_i = 4$		$\sum_i = 4$

Hence:

$$S(t) = \exp(-\Lambda(t)) = \begin{cases} \exp(0) = 1, & 0 \le t < 1\\ \exp(-1/7) = .87, & 1 \le t < 3\\ \exp(-10/21) = .62, & 3 \le t < 9\\ \exp(-41/42) = .38, & 9 \le t < 10\\ & \text{undefined}, & t \ge 10 \end{cases}$$

Altschuler-Nelson estimator – Example



Homogeneous parametric models

Q: How about *continuous*, parametric, counterparts to KM and AN? Still incorporating no predictors?

A: There are really *a lot* of possibilities.

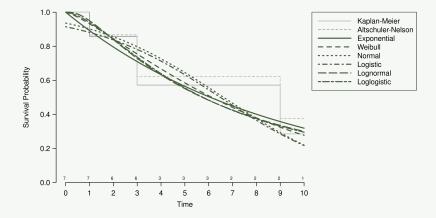
Most common examples:

- ► Exponential
- ▶ Weibull
- Normal
- ► Logistic
- ► Log-normal
- ► Log-logistic
- ▶ ...

My advice: Just fit several of these and compare. There is no 'best' model, it depends on the data.

Homogeneous parametric models

Data: T = 1, 3, 3, 3+, 6+, 9, 10+.



Assessing model fit

I like Harrell's take on this:

- ▶ To assess model fit, use graphical methods.
- ▶ No significance tests at this point, great!
- ▶ But there are some test options, see e.g. Chapter 20.

We show an example: Assess the fit of the exponential model.

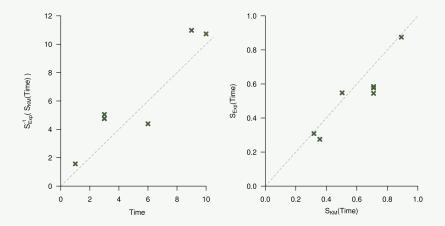
Two plotting options, akin to QQ-plots:

▶ Plot
$$S_{\text{Exp}}^{-1}(S_{\text{KM}}(T))$$
 versus *T*;

▶ Plot $S_{\text{Exp}}(T)$ versus $S_{\text{KM}}(T)$.

Assessing model fit

Data: T = 1, 3, 3, 3+, 6+, 9, 10+.



First model until now that allows incorporating predictor variables $X = \{X_1, X_2, \dots, X_k\}.$

• X_i can be continuous, dichotomous, polytomous, etc.

The proportional hazards (PH) model generalizes the hazard function $\lambda(t)$:

$$\lambda(t|X) = \lambda(t) \overbrace{\exp(\underbrace{\beta_0 + \beta_1 X_1 + \beta_2 X_2 + \dots + \beta_x X_k}_{X\beta})}^{\text{relative hazard function}} = \lambda(t) \exp(X\beta)$$

- ► $\lambda(t|X)$ = hazard function for *T* given the predictors *X*.
- ► $\lambda(t)$ = 'underlying' hazard function (for a subject with $X\beta = 0$).
- $\exp(X\beta)$ describes the *relative* effects of the predictors.

Note: The intercept β_0 may be omitted (kind of 'absorbed' into $\lambda(t)$).

$$\lambda(t|X) = \lambda(t) \exp(X\beta)$$

Here are the 'friends':

$$\Lambda(t|X) = \Lambda(t) \exp(X\beta)$$
$$S(t|X) = S(t)^{\exp(X\beta)}$$

- ∧(t) = 'underlying' cumulative hazard function (for a subject with Xβ = 0).
- S(t) = 'underlying' survival function (for a subject with Xβ = 0).

It is easier to consider the log-model versions:

$$\log \lambda(t|X) = \log \lambda(t) + X\beta$$
$$\log \Lambda(t|X) = \log \Lambda(t) + X\beta$$
$$\log S(t|X) = \underbrace{\log S(t)}_{\text{time}} \times \underbrace{\exp(X\beta)}_{\text{predictors}}$$

 Observe that we separated the time and the predictors components.

Important consequence due to the separability of *t* and *X*:

- ▶ The effect of *X* is assumed to be the same at all values of *t*.
- I.e.: We assume no $t \times X$ interaction effect.

How to interpret regression coefficient β_j (j = 1, ..., k)?

$$\log \lambda(t|X) = \log \lambda(t) + (\beta_0 + \beta_1 X_1 + \beta_2 X_2 + \dots + \beta_x X_k)$$
$$\log \Lambda(t|X) = \underbrace{\log \Lambda(t)}_{\text{time}} + \underbrace{(\beta_0 + \beta_1 X_1 + \beta_2 X_2 + \dots + \beta_x X_k)}_{\text{predictors}}$$

Additive interpretation:

log λ(t|X) increases by β_j units when X_j increases by 1 unit at any time point t, holding all the other predictors constant:

$$\log \lambda(t|\ldots, X_j+1,\ldots) = \log \lambda(t|\ldots, X_j,\ldots) + \beta_j.$$

Same for $\log \Lambda(t|X)$.

How to interpret regression coefficient β_j (j = 1, ..., k)?

$$\begin{split} \lambda(t|X) &= \lambda(t) & \times \exp(\beta_0 + \beta_1 X_1 + \beta_2 X_2 + \dots + \beta_x X_k) \\ \Lambda(t|X) &= \Lambda(t) & \times \exp(\beta_0 + \beta_1 X_1 + \beta_2 X_2 + \dots + \beta_x X_k) \\ \log S(t|X) &= \underbrace{\log S(t)}_{\text{time}} \times \underbrace{\exp(\beta_0 + \beta_1 X_1 + \beta_2 X_2 + \dots + \beta_x X_k)}_{\text{predictors}} \end{split}$$

Multiplicative interpretation:

 λ(t|X) is multiplied by exp(β_j) units when X_j increases by 1 unit at any time point t, holding all the other predictors constant:

$$\underbrace{\frac{\lambda(t|\ldots,X_j+1,\ldots)}{\lambda(t|\ldots,X_j,\ldots)}}_{(j,j,\ldots)} = \exp(\beta_j).$$

hazard ratio

Same for Λ(t|X).
Same for log S(t|X).

Hazard ratio

$$\mathrm{HR} = \frac{\lambda(t|...,X_j+1,...)}{\lambda(t|...,X_j,...)}$$

▶ HR = 1: No effect, i.e., X_i is unrelated to P(event).

- ▶ HR < 1: Hazard reduction, i.e., X_j is negatively associated with P(event). Larger survival time.
- ► *HR* > 1: Hazard increase, i.e., *X_j* is positively associated with *P*(event). Smaller survival time.

Proportional hazards assumption

The hazards ratio for any two subjects is independent of time:

$$\frac{\lambda(t|X_{Sub1})}{\lambda(t|X_{Sub2})} = \underbrace{\exp\left[\beta(X_{Sub1} - X_{Sub2})\right]}_{\text{no }t \text{ here!}}$$

In particular:

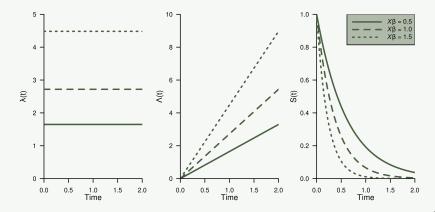
► The hazard curves for different groups (e.g., sex groups) should be proportional and thus cannot cross.

Example: Exponential PH survival model $X\beta = \beta_0 + \beta_1 X_1 + \beta_2 X_2 + \dots + \beta_x X_k.$

$$\lambda(t|X) = \exp(X\beta)$$

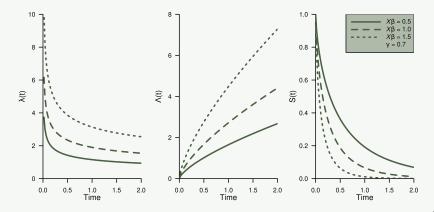
$$\Lambda(t|X) = t \exp(X\beta)$$

$$S(t|X) = \exp(-t)^{\exp(X\beta)}$$



Example: Weibull PH survival model $X\beta = \beta_0 + \beta_1 X_1 + \beta_2 X_2 + \dots + \beta_x X_k.$

$$\lambda(t|X) = \gamma t^{\gamma-1} \exp(X\beta)$$
$$\Lambda(t|X) = t^{\gamma} \exp(X\beta)$$
$$S(t|X) = \exp(-t^{\gamma})^{\exp(X\beta)}$$



Cox proportional hazards model

Seemingly the most popular survival model used.

The Cox PH model:

$$\lambda(t|X) = \lambda(t) \exp(X\beta)$$

- ► Looks the same as the general PH model!
- ▶ But, it is semiparametric:
 - ✓ It makes a parametric assumption in $X\beta = \beta_1 X_1 + \cdots + \beta_x X_k$. (NB: No intercept is typical for the Cox PH model.)
 - ✓ But, it assumes no parametric model for the hazard function $\lambda(t)$. Actually, it won't even be estimated!

Rationale:

- The true hazard function $\lambda(t)$ may be too complex.
- The effect of the predictors is more relevant than the shape of $\lambda(t)$.

The Cox PH model allows by passing $\lambda(t)$.

Cox proportional hazards model

But how does this magic work?

▶ Use the rank ordering of *T*.

Advantages:

- ▶ Better protection against outliers.
- ► The Cox PH model is more efficient than parametric PH models when parametric assumptions are strongly violated.
- Surprisingly, the Cox PH model is as efficient as parametric PH models even when parametric assumptions hold.

Final worked out example

I will use the *lung* dataset from the *survival* package in R.

- ▶ The data concern survival in patients with advanced lung cancer.
- ▶ These data have been analyzed *ad nauseam*, e.g.:
- ► Tutorial 1
- Tutorial 2
- Tutorial 3
- Using Bayesian statistics and Stan!

I will just run some basics.

Want something else to play afterwards?

- Check other datasets in the *survival* R package, it has plenty (e.g., *ovarian, veteran*).
- ▶ Bayesian analysis on mastectomy data (HSAUR R package)
- Recidivism data (carData R package)
- ▶ ...

Lung data

Time and censoring

Predictors

$Y_i = time$	e_i =status	age	sex	ph.ecog	ph.karno	wt.loss
306	2	74	1	1	90	NA
455	2	68	1	0	90	15
1010	1	56	1	0	90	15
210	2	57	1	1	90	11
883	2	60	1	0	100	0
1022	1	74	1	1	50	0
:	:	:	:	:	:	:
			•			

- ► *time*: Survival time in days
- status: Censoring
 (1=censored, 2=dead)

- ► *age*: Age in years
- ▶ *sex*: Male=1, Female=2
- *ph.ecog*: ECOG performance score (0=good, ..., 5=dead)
- ▶ *ph.karno*: 0-100 performance score (physician)
- ▶ *wt.loss*: Weight loss in last 6 months

In R

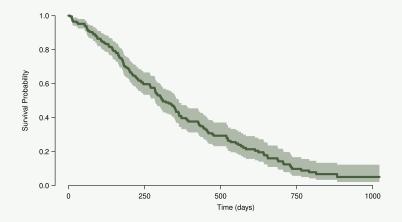
There are loads of packages and options to go about:



- ▶ ...

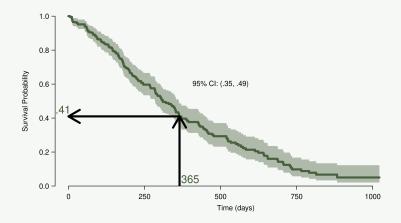
I'll use functions from various packages depending on functionality and eye-candyness.

Lung data: Kaplan-Meier



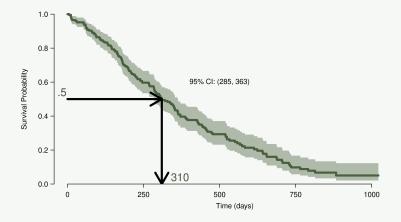
Lung data: Kaplan-Meier

What is P(t > 365 days)?



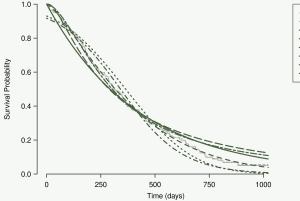
Lung data: Kaplan-Meier

What is the average (median) survival time?



Lung data: Homogeneous parametric models

With no predictors.





First we regress T on sex.

term	estimate	std.error	statistic	p.value	conf.low	conf.high
sex	-0.53	0.17	-3.18	0	-0.86	-0.2

Hence

$$\lambda(t|sex) = \lambda(t) \exp(-.53sex)$$

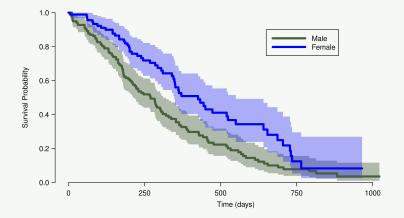
and

hazard ratio =
$$\frac{\lambda(t|\text{sex} = 2)}{\lambda(t|\text{sex} = 1)} = \frac{\lambda(t|\text{Female})}{\lambda(t|\text{Male})} = \exp(-.53) \simeq .59.$$

(Multiplicative) Interpretation:

▶ .59 times as many females are dying as males, at any time *t*. *or, equivalently,*

▶ Being female reduces the hazard by (1 - .59)100% = 41%.



Adding more predictors.

term	estimate	std.error	statistic	p.value	conf.low	conf.high
sex	-0.63	0.18	-3.56	0.00	-0.98	-0.28
age	0.02	0.01	1.55	0.12	0.00	0.03
ph.ecog	0.74	0.19	3.87	0.00	0.37	1.12
ph.karno	0.02	0.01	1.56	0.12	0.00	0.03
wt.loss	-0.01	0.01	-1.39	0.17	-0.02	0.00

Holding the other predictors constant, ...

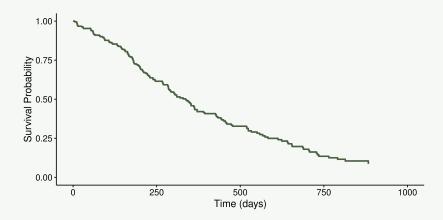
•
$$HR_{sex} = \exp(-.63) = .53$$
:

... being female *reduces* the hazard rate by a factor of .53 (47%), at any time t.

$$\blacktriangleright HR_{age} = \exp(.02) = 1.02:$$

... each extra year *increases* the hazard rate by a factor of 1.02 (2%), at any time t.

^{► ...}



Conclusion

Survival analysis offers a plethora of statistical models suitable to analyze 'time to event' data.

Much more is to be said on this topic. For instance:

- More on assumption checking (a lot of plotting options are available, also a few tests...).
- ► More plotting possibilities.
- Model comparison.
- ► Bayesian survival analysis!
- ► Accelerated failure time models.
- ► More complex models.
- ► ...