

Statistical Practice in Epidemiology 2018

Survival analysis with competing risks

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Points to be covered

1. Survival or time to event data & censoring.
2. Competing risks: event-specific cumulative incidences & hazards.
3. Kaplan–Meier and Aalen–Johansen estimators.
4. Regression modelling of hazards: Cox model.
5. Packages `survival`, `mstate`, `cmprisk`.
6. Functions `Surv()`, `survfit()`, `plot.survfit()`, `coxph()`, `Cuminc()`.

Survival time – time to event

Time spent (lex.dur) in a given **state** (lex.Cst) from its beginning till a certain *endpoint* or *outcome event* (lex.Xst) or *transition* occurs, changing the state to another.

Examples of such times and outcome events:

- ▶ lifetime: birth \rightarrow death,
- ▶ duration of marriage: wedding \rightarrow divorce,
- ▶ healthy exposure time:
start of exposure \rightarrow onset of disease,
- ▶ clinical survival time:
diagnosis of a disease \rightarrow death.

Ex. Survival of 338 oral cancer patients

Important variables:

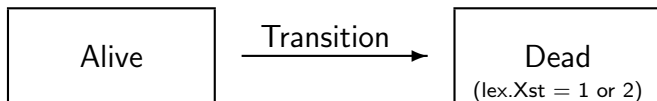
- ▶ `time` = duration of patientship from diagnosis (**entry**) till death (`death`) or censoring (`Alive`), (`lex.Cst` is (`Alive`))
- ▶ `event` = indicator for the outcome and its observation at the end of follow-up (**exit**):
0 = censoring,
1 = death from oral cancer

Special features:

- ▶ Two possible endpoints
- ▶ Censoring – incomplete observation of the survival time.

Set-up of classical survival analysis

- ▶ **Two-state model:** only one type of event changes the initial state.
- ▶ Major applications: analysis of lifetimes since birth and of survival times since diagnosis of a disease until death from any cause.



- ▶ **Censoring:** Death and final lifetime not observed for some subjects due to emigration or closing the follow-up while they are still alive

Distribution concepts: hazard function

The **hazard rate** or **intensity** function $\lambda(t)$

$$\lambda(t) = P(t < T \leq t + \Delta | T > t) / \Delta, \text{ for small } \Delta$$

\approx the conditional probability that the event occurs in a short interval $(t, t + \Delta]$, given that it does not occur before t , divided by interval length.

In other words, during a short interval

$$\text{risk of event} \approx \text{hazard} \times \text{interval length}$$

Distribution concepts: survival and cumulative hazard functions

Survival function

$$S(t) = P(T > t),$$

= probability of avoiding the event at least up to t
(the event occurs only after t).

The **cumulative hazard** (or integrated intensity):

$$\Lambda(t) = \int_0^t \lambda(u) du$$

Connections between the functions:

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

Observed data on survival times

For individuals $i = 1, \dots, n$ let

T_i = time to outcome event,

U_i = time to censoring.

Censoring is assumed **noninformative**, *i.e.*
independent from occurrence of events.

We observe

$y_i = \min\{T_i, U_i\}$, *i.e.* the exit time, and

$\delta_i = 1_{\{T_i < U_i\}}$, indicator (1/0) for the outcome event
occurring first, before censoring.

Censoring must properly be taken into account in the
statistical analysis.

Approaches for analysing survival time

- ▶ **Parametric model** (like Weibull, gamma, etc.) on hazard rate $\lambda(t) \rightarrow$ Likelihood:

$$L = \prod_{i=1}^n \lambda(y_i)^{\delta_i} S(y_i)$$

- ▶ **Piecewise constant rate** model on $\lambda(t)$
 - see Bendix's lecture on time-splitting (Poisson likelihood).
- ▶ **Non-parametric** methods, like Kaplan–Meier (KM) estimator of survival curve $S(t)$ and Cox proportional hazards model on $\lambda(t)$.

R package survival

Tools for analysis with one outcome event.

- ▶ `Surv(time, event) -> sobj`
creates a **survival object** `sobj` assuming that all start at 0, containing pairs (y_i, δ_i) ,
- ▶ `Surv(entry, exit, event) -> sobj2`
creates a survival object from entry and exit times,
- ▶ `survfit(sobj ~ x) -> sfo`
creates a **survfit** object `sfo` containing KM or other non-parametric estimates (also from a fitted Cox model),
- ▶ `plot(sfo)`
plot method for survival curves and related graphs,
- ▶ `coxph(sobj ~ x1 + x2)`
fits a Cox model with covariates `x1` and `x2`.
- ▶ `survreg()` – parametric survival models.

Ex. Oral cancer data (cont'd)

```
> orca$suob <- Surv(orca$time, 1*(orca$event > 0) )

> orca$suob[1:7]      # + indicates censored observation
[1] 5.081+ 0.419  7.915  2.480  2.500  0.167  5.925+

> km1 <- survfit( suob ~ 1, data = orca)
> km1                # brief summary
records    n.max n.start  events  median 0.95LCL 0.95UCL
 338.00    338.00  338.00   229.00   5.42    4.33    6.92

> summary(km1)        # detailed KM-estimate
```

time	n.risk	n.event	survival	std.err	lower 95% CI	upper 95% CI
0.085	338	2	0.9941	0.00417	0.9859	1.000
0.162	336	2	0.9882	0.00588	0.9767	1.000
0.167	334	4	0.9763	0.00827	0.9603	0.993
0.170	330	2	0.9704	0.00922	0.9525	0.989
0.246	328	1	0.9675	0.00965	0.9487	0.987
...						

Ex. Oral cancer KM estimates

