

Dealing with competing risks in survival analysis

Per Kragh Andersen
Section of Biostatistics
University of Copenhagen, Denmark

On behalf of TG8 - survival analysis

STRATOS meeting, Banff, 2016

Summary

1. Repetition of a few well-known facts from survival analysis
2. Competing risks
3. Applied papers on competing risks
4. Competing risks and other topic groups
5. Pseudo-observations
6. Concluding remarks

TG8 members: Michal Abrahamowicz, Per Kragh Andersen, Richard Cook, Coraline Danieli, Pierre Joly, Torben Martinussen, Maja Pohar Perme, Jeremy Taylor, Terry Therneau

Target population and censoring

When doing statistical inference, data are considered a *sample* from some *target population* to which parameters refer.

In *survival analysis*, parameters (like the survival function $S(t)$ or the hazard function $h(t)$) refer to a potentially completely observed population, i.e. one *without censoring*.

The object of survival analysis is then (the ambitious one) of drawing inference for such parameters based on *incomplete data*.

This requires an assumption of *independent censoring*.

Independent censoring

Independent censoring means that individuals censored at any given time t should not be a biased sample of those who are *at risk* at time t .

Stated in other words: the hazard

$$h(t) \approx P(T \leq t + dt \mid T > t)/dt$$

gives the *event rate* at time t , i.e. the failure rate given that the subject is still alive ($T > t$).

Independent censoring then means that the extra information that the subject is not only alive, but also uncensored at time t does not change the hazard.

Typically, independent censoring cannot be tested from the available data - it is a matter of discussion. However, it is crucial that the complete population (without censoring) is *well-defined*.

Inference for independently censored data

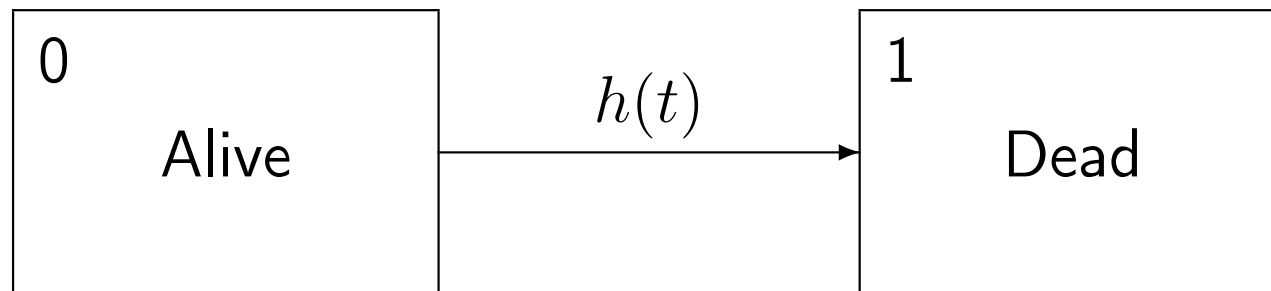
For independent observations $(X_i, \Delta_i, i = 1, \dots, n)$, with $X_i = \min(T_i, U_i)$, $\Delta_i = I(T_i \leq U_i)$, the likelihood is conveniently expressed via the hazard (and cumulative hazard) functions:

$$L(\theta) = \prod_i h_\theta(X_i)^{\Delta_i} \exp(-H_\theta(X_i)).$$

This leads (with suitable definition of NPMLE) to the *Nelson-Aalen estimator* for $H(\cdot)$ and to the *Cox partial likelihood*.

Using the relation between $S(t)$ and $H(t)$ leads to the *Kaplan-Meier estimator* as a plug-in estimator (because the product-integral relation $S(t) = \exp(-H(t))$ for absolutely continuous distributions becomes a finite product for a discrete distribution).

Target population



In the target population with complete observation, every one ends up in state 1 and the probability of being in state 1 at time t (the 'failure risk') is given uniquely from the hazard:

$$F(t) = 1 - S(t) = 1 - \exp(-H(t)).$$

Competing risks

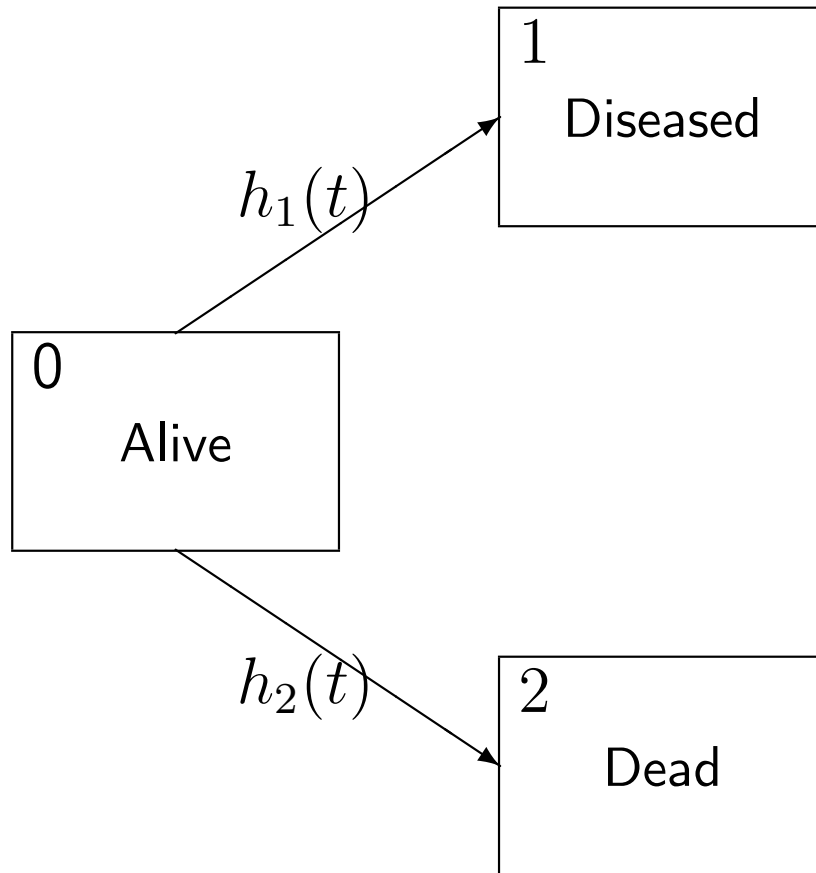
Suppose now that the event of interest is not overall mortality (or a composite end-point that includes death) but rather something like the occurrence of a given disease.

Then, in the target population even with complete follow-up, every one will *not* experience the event: some subjects will die without ever getting the disease.

Deaths without the disease should then *not* be thought of as an 'independent censoring' because a target population 'without censoring' (i.e., one where subjects are not allowed to die without the disease) is completely hypothetical.

We have *competing risks*.

Target population



Relationship between rates and risks

The cause-specific hazards $j = 1, 2$ are the transition intensities:

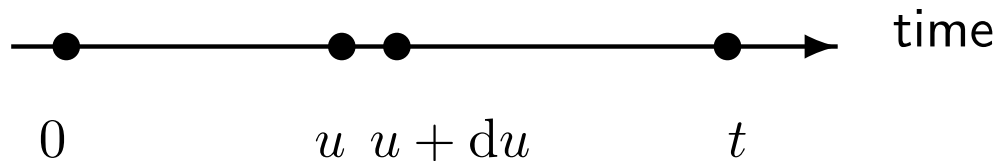
$$h_j(t) \approx P(\text{state } j \text{ time } t + dt \mid \text{state } 0 \text{ time } t) / dt.$$

The state occupation probabilities include the *overall survival function*:

$$S(t) = P(\text{alive time } t) = \exp(-(H_1(t) + H_2(t)))$$

and the *cumulative incidences* ('risks') for causes $j = 1, 2$:

$$F_j(t) = P(\text{state } j \text{ time } t) = \int_0^t S(u) h_j(u) du.$$



Both rates are needed to compute one risk!

Inference for independently censored data

For independent observations $(X_i, \Delta_i \cdot D_i, i = 1, \dots, n)$, with $X_i = \min(T_i, U_i)$, $\Delta_i = I(T_i \leq U_i)$, D_i = final state, the likelihood may again be expressed via the cause-specific hazard (and cumulative hazard) functions:

$$L(\theta) = \prod_i h_{1,\theta}(X_i)^{\Delta_i I(D_i=1)} h_{2,\theta}(X_i)^{\Delta_i I(D_i=2)} \exp(-H_{1\theta}(X_i) - H_{2\theta}(X_i))$$

which factorizes:

$$L(\theta) = \prod_i h_{1,\theta}(X_i)^{\Delta_i I(D_i=1)} \exp(-H_{1\theta}(X_i)) \\ \times \prod_i h_{2,\theta}(X_i)^{\Delta_i I(D_i=2)} \exp(-H_{2\theta}(X_i)).$$

Consequences of likelihood factorization

- Each factor has the form it would have if only cause j events were events and events of the other type were treated as (independent) censorings
- In particular, the *Nelson-Aalen estimator* for $H_j(\cdot)$, $j = 1, 2$ and the *Cox partial likelihood* for each cause separately work perfectly well
- Cumulative incidences may be estimated by plug-in (the 'Aalen-Johansen estimator'), while the '1-Kaplan-Meier estimator' for cause j estimates

$$\int_0^t \exp(-H_j(u)) h_j(u) du$$

and is (upwards) biased for $F_j(t)$

- First of all, it leads to *confusion* and *misunderstanding*

Confusion

Frequent question:

“why can we use the Cox model with competing risks but not the Kaplan-Meier estimator?”

Questions like that have lead to *numerous* (more or less) *pedagogical* papers in the applied (mostly medical) literature trying to explain these difficulties. These include:

Satagopan et al. (2004). *Br. J. Cancer*, 91, 1229-1235.

Southern et al. (2006). *J. Clin. Epidemiol.*, 59, 1110-1114.

Kim (2007). *Clin. Cancer Res.*, 13, 559-565.

Biau et al. (2007). *Clin. Orthopaed. and Rel. Res.*, 462, 229-233.

Scrucca et al. (2007). *Bone Marrow Transpl.*, 40, 381-387.

Zhang et al. (2008). *Expert Rev. Clin. Pharmacol.* 1, 391 .

Dignam et al. (2008). *J. Clin. oncol.*, 26, 4027-4034.

Shiels et al. (2009). *J. Clin. Epidemiol.*, 63, 459-467.

Wolbers et al. (2009). *Epidemiol.*, 20, 555-561.

Lau et al. (2009). *Amer. J. Epidemiol.*, 170, 244-256.

Allignol et al. (2011). *BMC Med. Res. Methodol.*, 11, 86.

Andersen et al. (2012). *Int. J. Epidemiol.*, 41, 861-870.

Austin et al. (2016). *Circulation*, 133, 601-619.

Much of the confusion seems to arise from not clearly distinguishing between censoring and competing risks (defining the target population) and between 'rates' and 'risks'.

Rates vs. risks?

Competing risks 'analogy':



Gladiators

Suppose that a gladiator may lose via two quite different mechanisms: lions or fellow gladiators.

When training a gladiator, he should both be prepared to face a lion or a fellow gladiator, and special skills may be needed to face a lion (even in the presence of the competing risk) and, similarly, special skills may be needed to beat a fellow gladiator. The cause-specific hazards describe how these mechanisms depend on properties and equipment of the gladiator.

For Caesar to predict the number of remaining gladiators still around at time t , both risks must be considered. He needs the cumulative incidences given the distribution of properties and equipment of the population of gladiators.

Conclusion

- Cause-specific hazards describe the mechanisms by which subjects may fail.
- Cumulative incidences describe the fraction of the population that fails from given causes.
- Both are useful (and both are needed?) for a complete description of the competing risks situation.

Latouche A., Allignol A., Beyersmann J., Labopin M., Fine J.P.: A competing risks analysis should report results on all cause-specific hazards and cumulative incidence functions. *J. Clin. Epidemiol.* (2013) **66**, 648-653.

Competing risks and TG8

Competing risks is, therefore, crucial for STRATOS TG8 and we have to find our niche in this vast literature.

The above-mentioned papers correspond to 'STRATOS-level 1'.

Can/should we contribute more to that?

Or should we focus on level 2?

(I am quite sceptical towards STRATOS-level 3.)

Competing risks and other topic groups

TG1 (missing data) There is a steadily increasing literature on how to do multiple imputation in competing risks analysis with missing data (e.g., Bartlett & Taylor, *Biostatistics*, 2016; Delord & Genin, *J. Statist. Comp. Sim.*, 2016).

Methods depend on whether covariates or cause of failure is missing. Important to distinguish between censoring and failure from a competing cause.

TG2 (variable selection and functional forms of a dose-response relationship) Special regression models are typically used for survival analysis (possibly with competing risks) though working with models with a linear predictor (LP) is quite similar for different types of outcome variable.

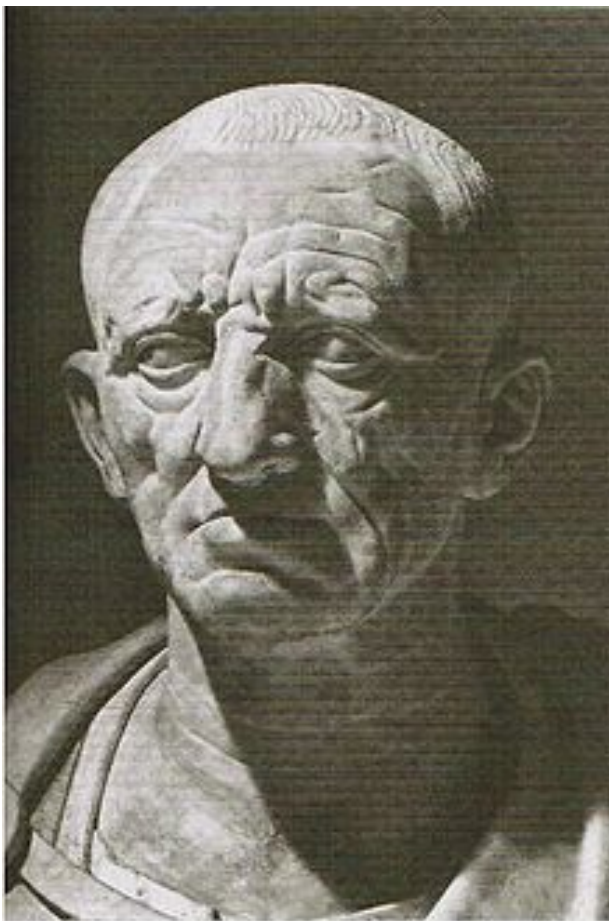
More below.

TG6 (evaluating diagnostic tests and prediction models) When assessing predictive accuracy, special care is needed for right-censored outcomes, including situations with competing risks.

More below.

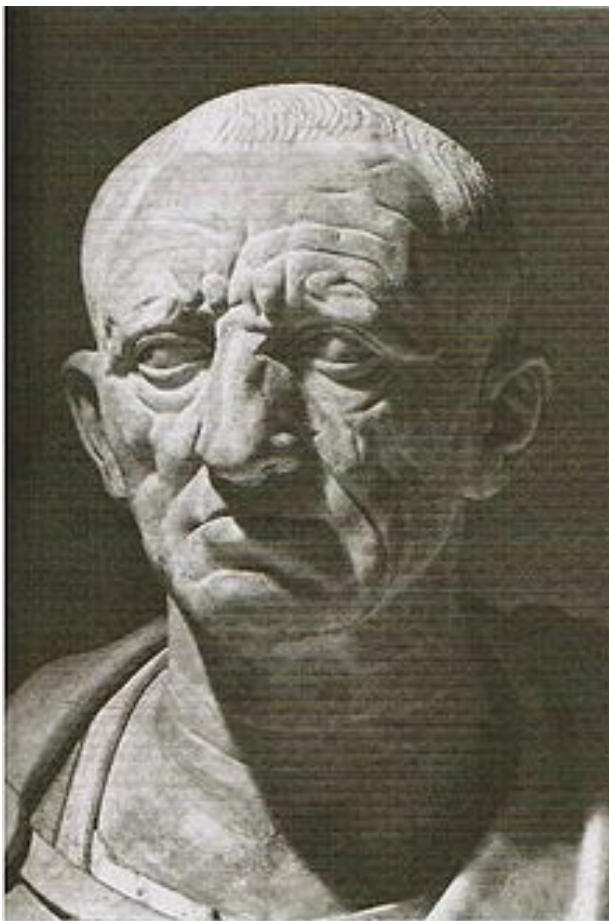
TG7 (causal inference) Both when using IPTW and when using the g-formula, special techniques are needed in the presence of competing risks.

More below.



Marcus Porcius Cato (“Cato the elder”, \approx 234 BC - 149 BC): Roman soldier, consul, senator etc.

“Ceterum censeo Carthaginem esse delendam”



Marcus Porcius Cato (“Cato the elder”, \approx 234 BC - 149 BC): Roman soldier, consul, senator etc.

“Moreover, I advise that Carthage must be destroyed”

Pseudo-observations

“Moreover, I think that pseudo-observations are useful in survival analysis”

Consider censored data and a mean value parameter $\theta = \mathbb{E}(f(T))$.

Possible mean value parameters, θ of interest are:

- survival probability $S(t) = \mathbb{E}(I(T_i > t))$,
- t -restricted mean $\mu_t = \mathbb{E}(T_i \wedge t)$,
- cumulative incidence in competing risks

$$F_j(t) = \mathbb{E}(I(T_i \leq t, D_i = j)).$$

Regression analysis for $E(f(T) | Z)$ may be performed using *pseudo-observations* as targets in a GEE:

$$\theta_i = n\hat{\theta} - (n - 1)\hat{\theta}^{-i}.$$

Here, $\hat{\theta}$ is a consistent estimator for the marginal mean (e.g., the Aalen-Johansen estimator for the cumulative incidence) and $\hat{\theta}^{-i}$ is the same estimator applied to the sample of size $n - 1$ obtained by eliminating subject i .

With no censoring, θ_i in the examples is simply, respectively:

- $I(T_i > t)$,
- $T_i \wedge t$,
- $I(T_i \leq t, D_i = j)$.

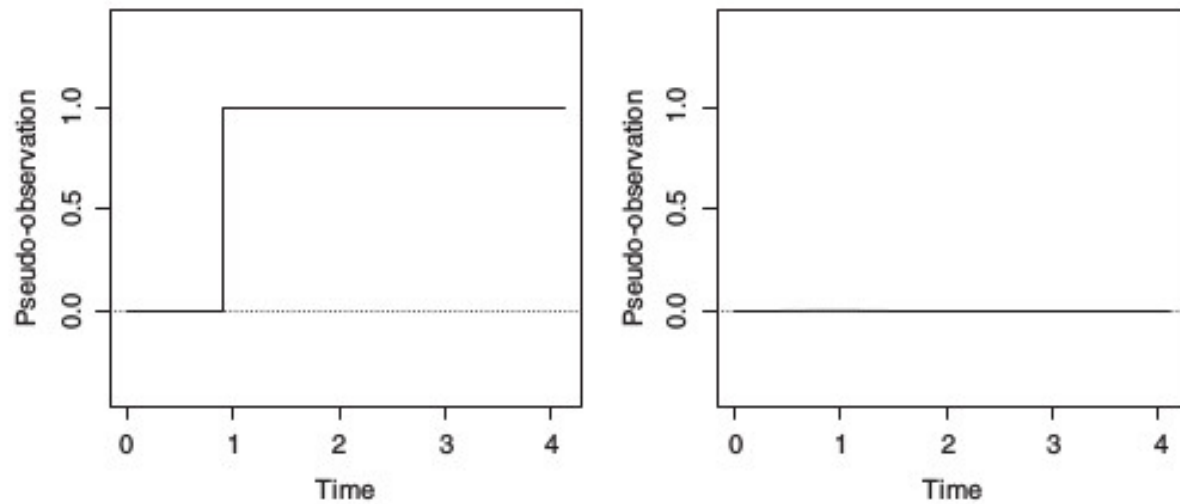


Figure 6 The pseudo-observations for the cumulative incidence function (for cause 1) in time for the case of no censoring. (a) Individual dying from cause 1; (b) Individual dying from cause 2.

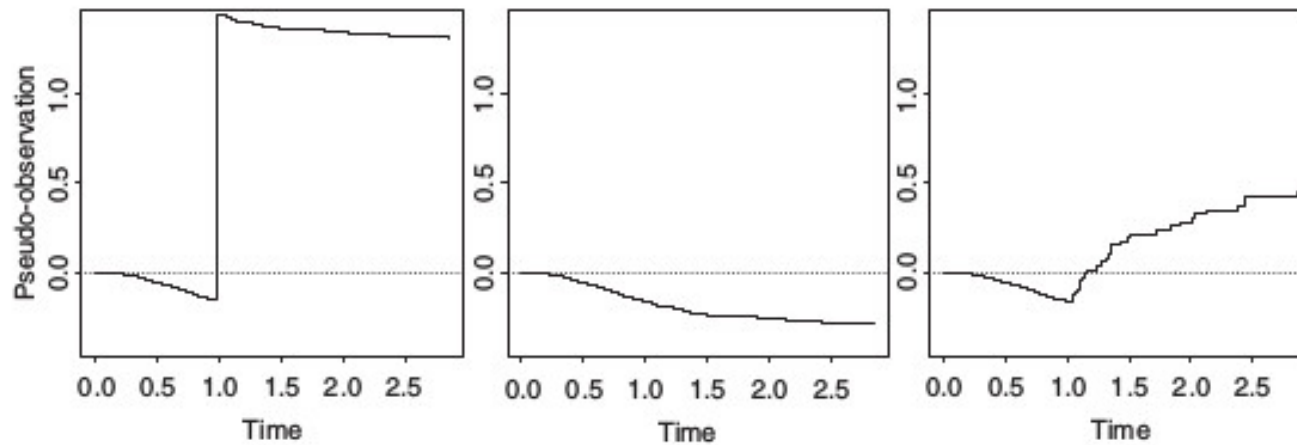


Figure 7 The pseudo-observations for the cumulative incidence function (for cause 1) in time for a censored data set. (a) Individual failing from cause 1; (b) Individual failing from cause 2; (c) Censored individual.

Properties of pseudo-observations

Graw, Gerds and Schumacher (*LIDA*, 2009) showed (for the competing risks cumulative incidence function) that if censoring is independent of covariates then, uniformly in $i = 1, \dots, n$ as $n \rightarrow \infty$:

$$\mathbb{E}(F_{ji}(t) \mid Z_i) = \mathbb{E}(I(T_i \leq t, D_i = j) \mid Z_i) + o_P(1).$$

(For covariate-dependent censoring - see Binder, Gerds and Andersen, *LIDA*, 2014)

Later asymptotic results by Jacobsen and Martinussen, (*Scand. J. Statist.*, 2016) and Overgaard, Parner and Pedersen (submitted), show that the estimated SD using the sandwich estimator when pseudo-values are used as targets in GEE may be (slightly) conservative.

Variable selection and functional forms

Competing risks regression for the cumulative incidence (at one or a few time points) may be performed using pseudo-observations (Klein & Andersen, *Biometrics*, 2005; Gerds et al., *Stat. in Med.*, 2012):

$$\sum_i \left(\frac{\partial}{\partial \beta} g^{-1}(LP_i) \right)^T V_i^{-1} (F_{ji} - g^{-1}(LP_i)) = 0.$$

Pros:

- Easy to apply different link functions, g not only the cloglog link
- Scatterplots and residual plots are available (Pohar Parme & Andersen, *Stat. in Med.*, 2008)

Cons:

- Not fully efficient
- Not (yet) possible to do for all time points simultaneously

Evaluation of prediction models

Pseudo-observations may be used when estimating the Brier score for a competing risks regression model for the cumulative incidence (Cortese et al., *Stat.in Med.*, 2013). The estimated value for complete data:

$$\hat{B}_j(s, t) = \frac{1}{m_s} \sum_{i \in R(s)} (I(X_i(t) = j) - r_j(t | Z_s))^2$$

(where $r_j(t | Z_s)$ is the predictor used at time s) is then replaced by

$$\tilde{B}_j(s, t) = \frac{1}{m_s} \sum_{i \in R(s)} (F_{ji}(t | s)(1 - 2r_j(t | Z_s)) + r_j(t | Z_s)^2)$$

Evaluation of prediction models

Pseudo-observations may also be used for making a calibration plot for a prediction model $r_j(t | z)$ (Gerds et al., *Stat. in Med.*, 2014).

The calibration curve is the graph of $p \rightarrow C(p, t, r)$:

$$C_j(p, t, r) = \mathbb{E}_{X,Z}(I(X(t) = j) | Z \in G_r(t; p))$$

where $G_r(t; p)$: set of covariates Z for which r predicts an event probability of p at t :

$$G_r(t; p) = \{z \in \mathbf{R}^d : r_j(t | z) = p\}.$$

The calibration curve may be estimated by:

$$\hat{C}_{j, a_n}(p, t, r) = \frac{1}{n_{a_n}} \sum_{i=1}^n F_{ji}(t) K_{a_n}(p, r_j(t | Z_i)).$$

Causal inference

The 'g-formula' for a completely observed outcome, Y is:

$$\frac{1}{n} \sum_{i=1}^n (\widehat{E}(Y_i | A_i := 1, Z_i) - \widehat{E}(Y_i | A_i := 0, Z_i)).$$

The propensity score weighted outcome to be used in a marginal structural model is:

$$\tilde{Y}_i = \frac{A_i Y_i}{\widehat{e}(Z_i)} + \frac{(1 - A_i) Y_i}{(1 - \widehat{e}(Z_i))}$$

(where $e(Z) = P(A = 1 | Z)$ is the propensity score). For a time-to-event outcome, e.g. the competing risks cumulative cause j incidence at time point t , Andersen et al. (submitted) showed that the outcome Y_i in these expressions may be replaced by the pseudo-value $F_{ji}(t)$.

Concluding remarks

- Competing risks is a central topic for TG8
- (However, there are many other topics of interest for TG8)
- Many examples of overlap between TG8 and other TG's
- Here, exemplified via competing risks
- Pseudo-values are useful (and fun!)