Modeling the cumulative incidence function of clustered competing risks data: a multinomial GLMM approach

master thesis defense



Henrique Laureano (.github.io)

LEG @ UFPR

April 11, 2021

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Clustered competing risk data



Key terms:

- Olustered: groups with a dependence structure (e.g. families);
- 2 Causes competing by something.

Something?

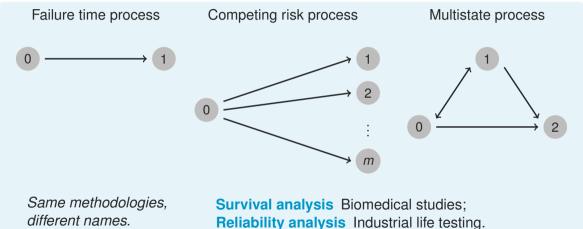
- Failure of an industrial or electronic component;
- Occurence or cure of a disease or some biological process;
- Progress of a patient clinic state.

Independent of the application, always the same framework

Cluster	ID	Cause 1	Cause 2	Censorship	Time	Feature
1	1	Yes	No	No	10	Α
1	2	No	No	Yes	8	Α
2	1	No	No	Yes	7	В
2	2	No	Yes	No	5	Α

Big picture: Failure time data





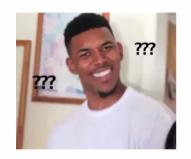


A comprehensive reference is Kalbfleisch and Prentice (2002)'s book.

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Modeling clustered competing risks data









What? Why? How?

Failure time data → Survival models



First of all, we have to choose which scale we model the survival experience.

1 Usually, is in the

hazard (failure rate) scale :
$$\lambda(t \mid \text{features}) = \lambda_0(t) \times c(\text{features})$$
. (1)

We have a Equation 1 for each competing cause.

The cluster dependence is something actually not measured...

Not measured dependence \rightarrow random/latent effects \rightarrow Frailty models.

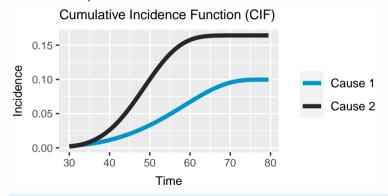
Full likelihood analysis with frailty models for competing risks data is generally complicated, when not impracticable.

2 Not usually, the probability scale.

Probability scale \rightarrow Cause-specific CIF



Besides the within-cluster dependence, there is an often interest in describing the time at event onset, directly described by the cause-specific



i.e., $CIF = \mathbb{P}[$ failure time $\leq t$, a given cause | features & latent effects].

Formally,



for a cause-specific of failure k, the cumulative incidence function (CIF) is defined as

$$F_k(t \mid \mathbf{x}) = \mathbb{P}[T \leqslant t, \ K = k \mid \mathbf{x}]$$

$$= \int_0^t f_k(z \mid \mathbf{x}) \, \mathrm{d}z \quad (f_k(t \mid \mathbf{x}) \text{ is the (sub)density for the time to a type } k \text{ failure})$$

$$= \int_0^t \underbrace{\lambda_k(z \mid \mathbf{x})}_{\text{cause-specific hazard function}} \underbrace{S(z \mid \mathbf{x})}_{\text{overall survival function}} z_{\text{unvival function}} t > 0, \quad k = 1, \dots, K.$$



Again, a comprehensive reference is Kalbfleisch and Prentice (2002)'s book.



Cederkvist et al. (2019)'s CIF specification



For two competing causes of failure, the cause-specific CIFs are specified in the following manner

$$F_k(t \mid \mathbf{x}, u_1, u_2, \eta_k) = \underbrace{\pi_k(\mathbf{x}, u_1, u_2)}_{\text{cluster-specific risk level}} \times \underbrace{\Phi[w_k g(t) - \mathbf{x} \gamma_k - \eta_k]}_{\text{cluster-specific failure time trajectory}}, \quad t > 0, \quad k = 1, 2, \quad (2)$$

with

$$\mathbf{1} \pi_k(\mathbf{x}, \mathbf{u}) = \exp\{\mathbf{x}\beta_k + u_k\} / \left(1 + \sum_{m=1}^{K-1} \exp\{\mathbf{x}\beta_m + u_m\}\right), \quad k = 1, 2, \quad K = 3;$$

 \bullet $\Phi(\cdot)$ is the cumulative distribution function of a standard Gaussian distribution;

In Cederkvist et al. (2019), this CIF specification is modeled under a *complicated* pairwise composite likelihood approach (Lindsay 1988; Varin, Reid, and Firth 2011).

Our contribution: a full likelihood analysis



For two competing causes of failure, a subject i, in the cluster i, in time t, we have

$$y_{ijt} \mid \underbrace{\{u_{1j}, u_{2j}, \eta_{1j}, \eta_{2j}\}}_{\text{latent effects}} \sim \text{Multinomial}(p_{1ijt}, p_{2ijt}, p_{3ijt})$$

$$\begin{bmatrix} u_1 \\ u_2 \\ \eta_1 \\ \eta_2 \end{bmatrix} \sim \begin{array}{l} \text{Multivariate} \\ \text{Normal} \end{array} \begin{pmatrix} \begin{bmatrix} 0 \\ 0 \\ 0 \\ 0 \end{bmatrix}, \begin{bmatrix} \sigma_{u_1}^2 & \text{cov}(u_1, u_2) & \text{cov}(u_1, \eta_1) & \text{cov}(u_1, \eta_2) \\ \sigma_{u_2}^2 & \text{cov}(u_2, \eta_1) & \text{cov}(u_2, \eta_2) \\ \sigma_{\eta_1}^2 & \sigma_{\eta_2}^2 \end{bmatrix} \end{pmatrix}$$

$$p_{kijt} = \frac{\partial}{\partial t} F_k(t \mid \boldsymbol{x}, \boldsymbol{u}, \eta_k)$$

$$= \frac{\exp\{\boldsymbol{x}_{kij}\beta_k + u_{kj}\}}{1 + \sum_{m=1}^{K-1} \exp\{\boldsymbol{x}_{mij}\beta_m + u_{mi}\}}$$

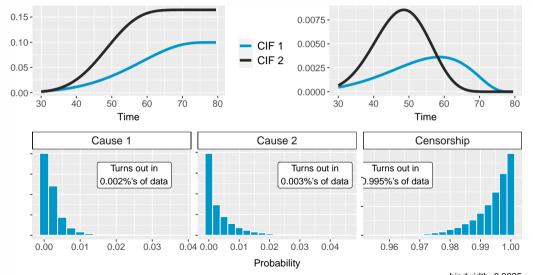
$$\times w_k \frac{\delta}{2\delta t - 2t^2} \phi \left(w_k \operatorname{arctanh} \left(\frac{t - \delta/2}{\delta/2} \right) - \boldsymbol{x}_{kij} \gamma_k - \eta_{kj} \right), \quad k = 1, 2.$$

Simulating from the model



dCIF 1

dCIF 2



Marginal likelihood function for two competing causes



$$L(\boldsymbol{\theta}; \boldsymbol{y}) = \prod_{j=1}^{J} \int_{\mathfrak{R}^{4}} \pi(\boldsymbol{y}_{j} \mid \boldsymbol{r}_{j}) \times \pi(\boldsymbol{r}_{j}) \, d\boldsymbol{r}_{j}$$

$$= \prod_{j=1}^{J} \int_{\mathfrak{R}^{4}} \left\{ \prod_{i=1}^{n_{j}} \prod_{t=1}^{n_{ij}} \left(\frac{(\sum_{k=1}^{K} y_{kijt})!}{y_{1ijt}! y_{2ijt}! y_{3ijt}!} \prod_{k=1}^{K} p_{kijt}^{y_{kijt}} \right) \right\} \times$$
fixed effect component
$$(2\pi)^{-2} |\Sigma|^{-1/2} \exp\left\{ -\frac{1}{2} \boldsymbol{r}_{j}^{\top} \Sigma^{-1} \boldsymbol{r}_{j} \right\} d\boldsymbol{r}_{j}$$
latent effect component

$$= \prod_{j=1}^{J} \int_{\mathfrak{R}^4} \left\{ \prod_{j=1}^{n_j} \prod_{t=1}^{n_{ij}} \prod_{k=1}^{K} p_{kijt}^{y_{kijt}} \right\} \underbrace{(2\pi)^{-2} |\Sigma|^{-1/2} \exp\left\{-\frac{1}{2} \boldsymbol{r}_j^\top \Sigma^{-1} \boldsymbol{r}_j\right\}}_{\boldsymbol{\mathfrak{R}}^{-1}} \mathrm{d}\boldsymbol{r}_j,$$

latent effect component

with p_{kijt} from Equation 3 and where $\theta = [\beta \ \gamma \ \mathbf{w} \ \sigma^2 \ \rho]^{\top}$ is the parameters vector.

fixed effect

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TMB: Automatic Differentiation and Laplace Approximation





Kristensen et al. (2016).

An R (R Core Team 2021) package for the quickly implementation of complex random effect models through simple C++ templates.

Workflow

- Write your objective function in a .cpp through a #include <TMB.hpp>;
- 2 Compile and load it in R via TMB::compile() and base::dyn.load(TMB::dynlib());
- 3 Compute your objective function derivatives with obj <- TMB::MakeADFun();</p>
- 4 Perform the model fitting, opt <- base::nlminb(obj\$par, obj\$fn, obj\$gr);</pre>
- **5** Compute the parameters standard deviations, TMB::sdreport(obj).



For details about TMB, AD, and Laplace approximation: Laureano (2021).

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Simulation study model designs



Risk model

Latent effects only on the risk level i.e.,

$$\Sigma = \begin{bmatrix} \sigma_{u_1}^2 & \mathsf{COV}_{u_1, u_2} \\ & \sigma_{u_2}^2 \end{bmatrix}.$$

Block-diag model

Latent effects on the risk and time levels without cross-correlations i.e.,

$$\Sigma = egin{bmatrix} \sigma_{u_1}^2 & \operatorname{cov}_{u_1,u_2} & 0 & 0 \ & \sigma_{u_2}^2 & 0 & 0 \ & & \sigma_{\eta_1}^2 & \operatorname{cov}_{\eta_1,\eta_2} \ & & & \sigma_{\eta_2}^2 \end{bmatrix}$$

Time model

Latent effects only on the failure time trajectory level i.e.,

$$\Sigma = egin{bmatrix} \sigma_{\eta_1}^2 & \mathsf{cov}_{\eta_1,\eta_2} \ & \sigma_{\eta_2}^2 \end{bmatrix}.$$

Complete model

A complete latent effects structure i.e..

$$\Sigma = \begin{bmatrix} \sigma_{u_1}^2 & \text{cov}_{u_1, u_2} & 0 & 0 \\ & \sigma_{u_2}^2 & 0 & 0 \\ & & \sigma_{\eta_1}^2 & \text{cov}_{\eta_1, \eta_2} \\ & & & \sigma_{\eta_2}^2 \end{bmatrix}. \qquad \Sigma = \begin{bmatrix} \sigma_{u_1}^2 & \text{cov}_{u_1, u_2} & \text{cov}_{u_1, \eta_1} & \text{cov}_{u_1, \eta_2} \\ & \sigma_{u_2}^2 & \text{cov}_{u_2, \eta_1} & \text{cov}_{u_2, \eta_2} \\ & & & \sigma_{\eta_1}^2 & \text{cov}_{\eta_1, \eta_2} \\ & & & & \sigma_{\eta_2}^2 \end{bmatrix}.$$

Simulation study setup

Four latent effects structures:



Risk model;

2 Time model;

3 Block-diag model;

4 Complete model.

Two CIF configurations:

Low max incidence ≈ 0.15 ;

High max incidence \approx 0.60.

For each of those $4 \times 2 = 8$ scenarios, we vary the sample and cluster sizes:

5000 data points

- 2500 clusters of size 2;
- 1000 clusters of **size 5**;
- 500 clusters of size 10.

30000 data points

- 15000 clusters of size 2;
- 6000 clusters of size 5;
- 3000 clusters of size 10.

60000 data points

- 30000 clusters of size 2;
- 12000 clusters of **size 5**;
- 6000 clusters of **size 10**.

Totalizing, $\mathbf{8} \times \mathbf{3} \times \mathbf{3} = \mathbf{72}$ scenarios.

For each scenario, we simulate 400 samples, totalizing $72 \times 400 = 28800$ model fittings.

Simulation study results



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Thanks for watching and have a great day



Special thanks to



PPGMNE

Programa de Pós-Graduação em Métodos Numéricos em Engenharia





Joint work with

Wagner H. Bonat http://leg.ufpr.br/~wagner

ner

Paulo Justiniano Ribeiro Jr. http://leg.ufpr.br/~paulojus



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