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



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# Giving context: defining where we are and what we did



## **Object**

 Handle clustered competing risks data (a kind of failure time data) through the cumulative incidence function (CIF).

#### Goal

 Perform maximum likelihood estimation in terms of a full likelihood formulation based on Cederkvist et al. (2019)'s CIF specification (Scheike's).

#### Contribution

- The full likelihood formulation is in terms of a generalized linear mixed model (GLMM) a conditional approach (with fixed and random/latent effects);
- The optimization and inference are tacked down via an efficient model implementation with the use of *state-of-art* computational libraries (Kristensen et al. (2016)'s TMB).

## **Outline**



- 1 Data;
- 2 Model;
- 3 TMB: Template Model Builder;
- 4 Simulation study;
- **5** Conclusion;
- 6 References.

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# Clustered competing risk data



#### Key ideas:

- Clustered: groups with a dependence structure (e.g. families);
- 2 Causes competing by something;
- 3 Occurrence time of this something.

### Something?

- Failure of an industrial or electronic component;
- Occurrence or cure of a disease or some biological process;

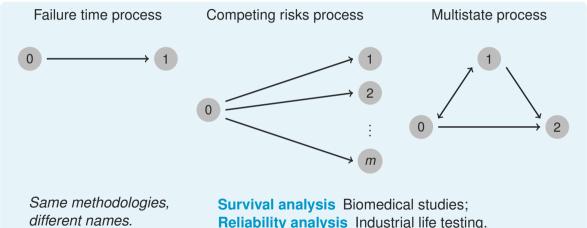
 Progress of a patient clinic state.

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/time-to-event outcomes





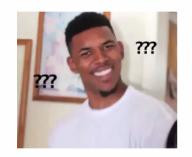


A comprehensive reference is Kalbfleisch and Prentice (2002)'s book.

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## Modeling clustered competing risks data









What? Why? How?

## Modeling failure time data



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.

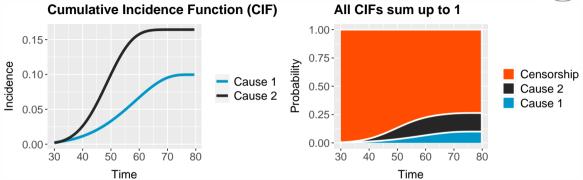
Frailty-based models for (multiple) survival experiences turn out in challengeable likelihood functions with inference routines mostly done via

 Elaborated and slow expectation—maximization (EM) algorithms;  Inefficient Markov chain Monte Carlo (MCMC) schemes.

2 Not usually, the probability scale.

## $\textbf{Probability scale} \rightarrow \textbf{Cause-specific CIF}$





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

Common applications: family studies.

↓ Keywords: within-family/cluster dependence; age at disease onset; populations.

## 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;
- 3  $g(t) = \operatorname{arctanh}(2t/\delta 1), \quad t \in (0, \delta), \quad g(t) \in (-\infty, \infty).$
- In Cederkvist et al. (2019), this CIF specification is modeled under a 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 j, in time t, we have

$$\begin{aligned} y_{ijt} \mid \underbrace{\{u_{1j}, u_{2j}, \eta_{1j}, \eta_{2j}\}}_{\text{latent effects}} &\sim \text{Multinomial}(p_{1ijt}, p_{2ijt}, p_{3ijt}) \\ \begin{bmatrix} u_1 \\ u_2 \\ \eta_1 \\ \eta_2 \end{bmatrix} &\sim \text{Multivariate} \begin{pmatrix} \begin{bmatrix} 0 \\ 0 \\ 0 \\ 0 \end{bmatrix}, \begin{bmatrix} \sigma_{u_1}^2 & \text{cov}(u_1, u_2) & \text{cov}(u_1, \eta_1) & \text{cov}(u_1, \eta_2) \\ \sigma_{u_2}^2 & \text{cov}(u_2, \eta_1) & \text{cov}(u_2, \eta_2) \\ \sigma_{\eta_1}^2 & \text{cov}(\eta_1, \eta_2) \end{bmatrix} \\ p_{kijt} &= \frac{\partial}{\partial t} F_k(t \mid \boldsymbol{x}, \boldsymbol{u}, \eta_k) \\ &= \frac{\exp\{\boldsymbol{x}_{kij}\beta_k + u_{kj}\}}{1 + \sum_{m=1}^{K-1} \exp\{\boldsymbol{x}_{mij}\beta_m + u_{mi}\}} \end{aligned}$$

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

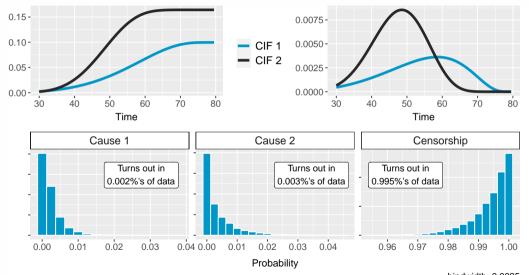
(3)

## Simulating from the model



dCIF 1

dCIF 2



# Marginal likelihood function for two competing causes



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

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

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.,

$$\Sigma = \begin{bmatrix} \sigma_{u_1}^2 & \cos_{u_1, u_2} & 0 & 0 \\ & \sigma_{u_2}^2 & 0 & 0 \\ & & \sigma_{\eta_1}^2 & \cos_{\eta_1, \eta_2} \\ & & & \sigma_{\eta_2}^2 \end{bmatrix}$$

#### 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  $\approx 0.15$ ;

**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 500 samples, totalizing  $72 \times 500 = 36000$  model fittings.



#### First of all, the **time**.

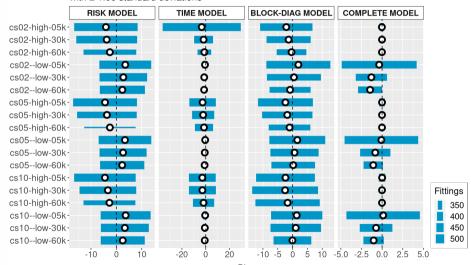
- The non-complete models (2D Laplace aprox.) are kind of fast, taking always less than 5 min.
- In the most expensive scenarios (30K 4D Laplaces),
   the complete model takes 30 min.
   In a full R implementation with 10K 4D Laplaces, it took 30hrs. TMB is fast.
- We also did a Bayesian analysis via Stan/NUTS-HMC (Stan Development Team 2020).
  - 1 week of parallelized processing for a 2500 size 2 clusters scenario with tuned NUTS.
     This just reinforces the MCMC impracticability for some complex models.

#### Parameters estimation.

The non-complete models fail to learn the data.
 They appear to be not structured enough to capture the data characteristics.

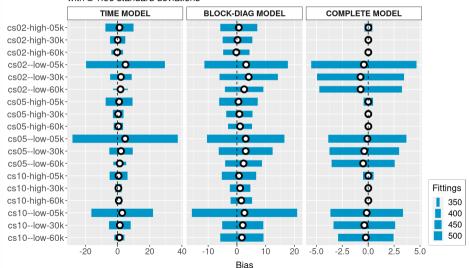
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Parameter: β<sub>1</sub>



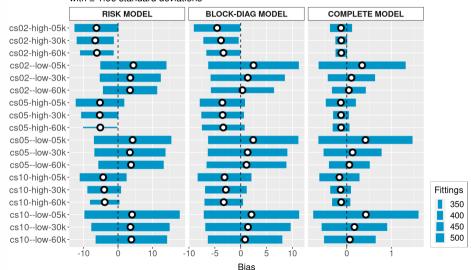
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Parameter:  $log(\sigma_4^2)$ 



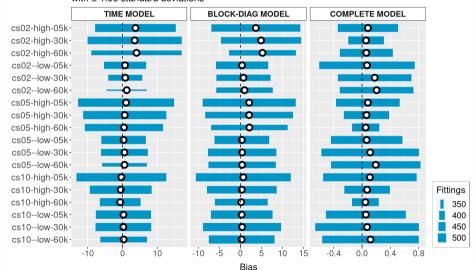
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Parameter:  $z(\rho_{12})$ 

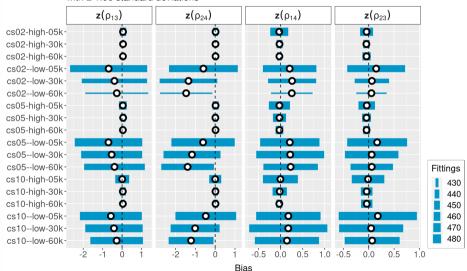


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Parameter:  $z(\rho_{34})$ 



#### Complete model's cross-correlations



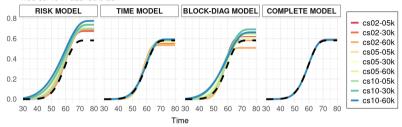


## Simulation study results: High CIF scenario



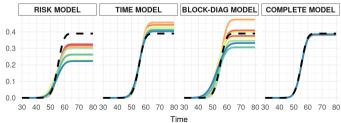


True curve in dashed black



#### CIF of failure cause 2

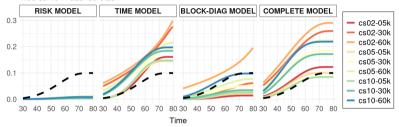
True curve in dashed black



## Simulation study results: Low CIF scenario

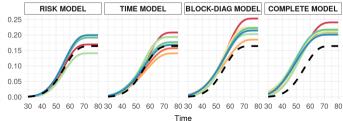
#### CIF of failure cause 1

True curve in dashed black



#### CIF of failure cause 2

True curve in dashed black



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## Take-home message



The complete model works. It's not magnificent, but it works.

- 1 It works better in the high CIF scenarios;
- 2 As expected, as the sample size increases the results get better;
- 3 We do not see any considerable performance difference between cluster/family sizes;
- 4 Satisfactory full likelihood analysis under the maximum likelihood estimation framework (the estimates bias-variance could be smaller).

#### What else can we do?

- 1 Instead of a conditional approach (latent effects model), we can try a marginal approach e.g., an McGLM (Bonat and Jørgensen 2016);
- We can also try a copula (Embrechts 2009), on maybe two fronts:1) for a full specification; 2) to accommodate the within-cluster dependence.



For more read Laureano (2021) master thesis.

## Thanks for watching and have a great day



Special thanks to



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Joint work with

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