

# Joint Analysis of Longitudinal and Survival AIDS Data with a Spatial Fraction of Long-term Survivors: A Bayesian Approach

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## Overview

The objective is to model longitudinal and survival data jointly taking into account the dependence between the two responses in a real HIV/AIDS dataset. We employ here a Bayesian hierarchical approach to jointly model spatial-clustered survival data with a fraction of long-term survivors along with longitudinal measurements of CD4<sup>+</sup>T lymphocyte counts for a random sample of 500 HIV/AIDS individuals collected in all the 27 states of Brazil during the period 2002–2006.

In order to accommodate a more flexible choice for the longitudinal model we propose to use a specification via penalized B-Splines [2]. To deal with the possible fact that only a subpopulation undergoes the event we propose a spatial cure model, which helps in explaining the behavior of chronic or potentially terminal diseases in different regions. Classical survival models (e.g. Cox model) are not-well suited to take into account long-term survivors.

Results show that using a cure fraction allows us to improve the results comparatively to a survival analysis without it. The inclusion of spatial frailties eases in mapping the heterogeneity in the risk among the Brazilian states and helps in the explanation of the hazard.

## Dataset

- **Data origin:** Brazilian database on HIV;
- **Period (years):** 2002–2006;
- **Sample size:**  $n = 500$  individuals
- **Response variables:**  $y = \sqrt{\text{CD4}^+\text{T}}$  lymphocyte counts and survival time (years since the patient's entry in the study until death)
- **Explanatory variables:** age ( $<50=0$ ,  $\geq 50=1$ ); gender (Female=0, Male=1); PrevOI (previous opportunistic infection at study entry=1, no previous infection=0); region of residence (one of the 27 Brazilian States)
- 34 deaths. 88% of the patients were between 15 and 49 years old; 60% were males. The CD4 counts initial median was 245 cells/mm<sup>3</sup> (men - 226 cells/mm<sup>3</sup>; women - 263 cells/mm<sup>3</sup>).

## References

- [1] Cooner, F., Banerjee, S., and McBean, A.M. (2006) Modelling geographically referenced survival data with a cure fraction. *Stat Meth Med Res*, **15**(4):307–324;
- [2] Lang, S., Brezger, A. (2004) Bayesian p-splines. *J Comp Graph Stat*, **13**(1):183–212;
- [3] Martins, R., Silva, G.L., and Andreozzi, V. (2016) Bayesian joint modeling of longitudinal and spatial survival AIDS data. *Stat Med*, doi: 10.1002/sim.6937.
- [4] Rizopoulos, D., Ghosh, P. (2011) A Bayesian semi-parametric multivariate joint model for multiple longitudinal outcomes and a time-to-event. *Stat Med*, **30**(12):1366–1380;

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## Model

Consider the CD4 repeated measurements,  $\mathbf{y}_{ik} = (y_{ik1}, \dots, y_{in_{ik}})$ , and the observed (possibly right censored) time-to-death,  $T_{ik}$ , for the  $i$ th individual living in the  $k$ th region,  $k = 1, \dots, K$ ,  $i = 1, \dots, n_k$ . **Longitudinal data** is described by a **nonlinear mixed effects model**,

$$y_{ikj} | \mathbf{b}_{ik}, \sigma_e^2 \sim \mathcal{N}(y_{ik}^*(t_{ikj}), \sigma_e^2), \quad j = 1, \dots, n_{ik}, \quad (1)$$

$$y_{ik}^*(t_{ikj}) = (\beta_{10} + b_{ik0}) + \mathbf{z}_{ik}^\top \boldsymbol{\beta}_1 + m_p(t_{ikj}; \boldsymbol{\beta}_2) + m_{ik}(t_{ikj}; \mathbf{b}_{ik}), \quad (2)$$

where  $m_p(\cdot)$  and  $m_{ik}(\cdot)$  are the overall and individual nonlinear effects, respectively, considered here as **Penalized cubic B-Splines**.  $\beta_{10}$  and  $b_{ik0}$  are the common and subject-specific intercept, respectively.  $\boldsymbol{\beta}_1 = (\beta_{11}, \dots, \beta_{1r})^\top$ , whose elements are assumed to be mutually independent, represents the fixed effects of the subject-specific vector of covariates,  $\mathbf{z}_{ik}$ ;  $\boldsymbol{\beta}_2 = (\beta_{21}, \dots, \beta_{2Q})^\top$  and  $\mathbf{b}_{ik} = (b_{ik1}, \dots, b_{ikQ})^\top$  are, respectively, population and individual-specific **regression parameters for each basis function, penalized through a second order random-walk**.

Assuming the time-to-death for the non-cured group is Weibull distributed,  $\mathcal{W}(\rho, e^{\eta(t)})$ , the **spatial cure model** [1] is described as,

$$S_p(t) = (1 - \theta_k) + \theta_k S(t) = (1 - \theta_k) + \theta_k \exp \left\{ -t^\rho e^{\eta(t)} \right\}, \quad (3)$$

being  $S_p$  and  $S$  the survival functions for the entire population and for the non-cured group, respectively.  $(1 - \theta_k)$  is a region-specific cure fraction. The baseline covariates and the longitudinal information will be introduced through the scale parameter,  $e^{\eta(t)}$ , allowing it to **vary across individuals and regions**:

$$\eta_{ik}(t) = \mathbf{x}_{ik}^\top \boldsymbol{\beta}_3 + \gamma y_{ik}^*(t) + W_k, \quad (4)$$

where  $\gamma$  is a parameter quantifying the effect of the CD4 values to the survival;  $\mathbf{x}_{ik}$  is a vector of baseline covariates (can coincide with  $\mathbf{z}_{ik}$ ),  $\boldsymbol{\beta}_3$  is the respective vector of coefficients and  $W_k$  is a region-specific frailty,  $W_k | \sigma_W^2 \sim \text{ICAR}(\sigma_W^2)$ .

## Application

Several scenarios for the Penalized splines were tried. Namely the internal knots were fixed every: 1 month, 2 months, 3 months, 4 months, 6 months and 12 months providing, respectively, 59, 29, 19, 14, 9 and 4 internal knots. The best fit was always achieved with 19 internal knots, corresponding roughly to a knot every 3 months during the 5 years, resulting in  $Q = 23$  cubic basis functions.

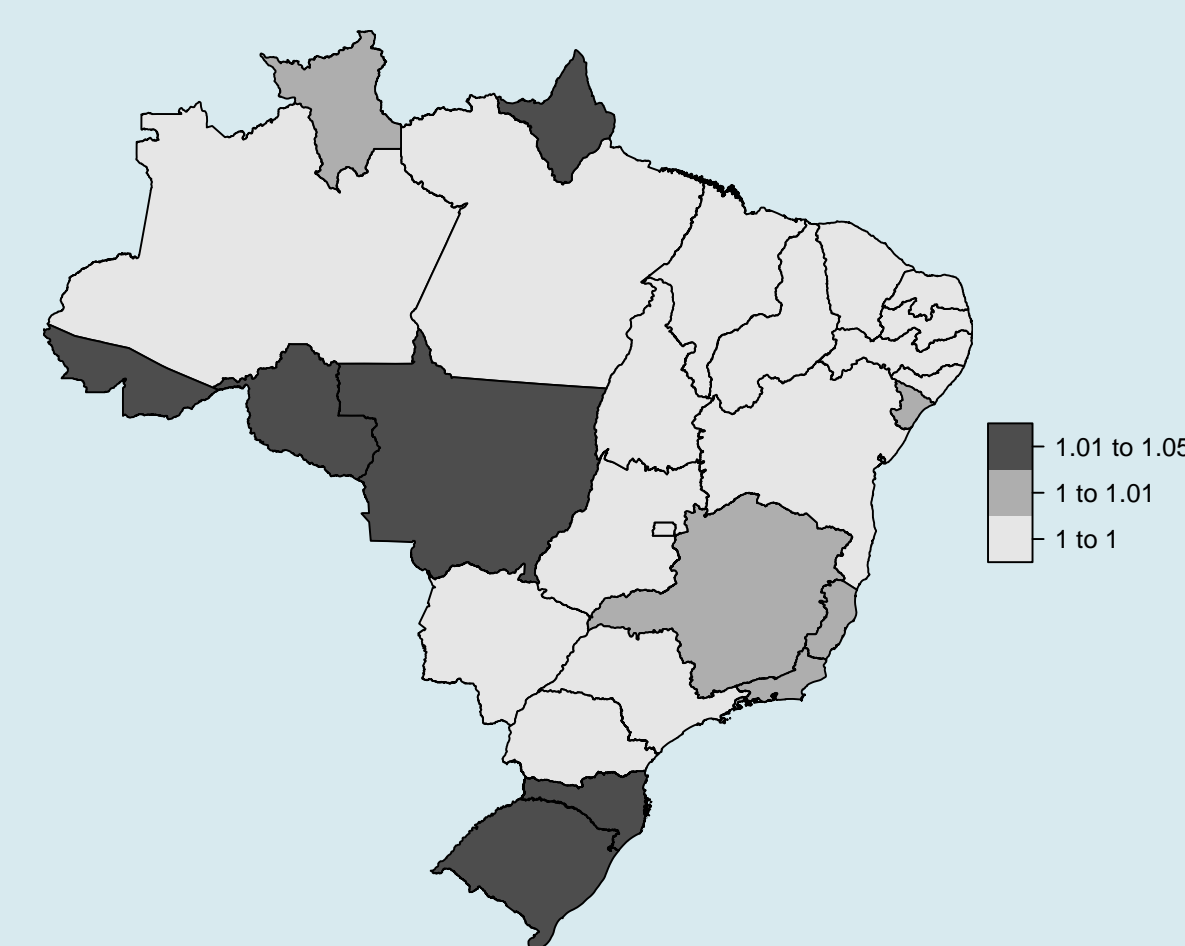
$$y_{ikj}^* = (\beta_{10} + b_{ik0}) + \beta_{11} \text{sex}_{ik} + \beta_{12} \text{age}_{ik} + \beta_{13} \text{PrevOI}_{ik} + \sum_{q=1}^{Q=23} \beta_{2q} B_q(t_{ikj}) + \sum_{q=1}^{Q=23} b_{ikq} B_q(t_{ikj}) \quad (5)$$

$$\eta_{ik}(t) = \beta_{31} + \beta_{32} \text{sex}_{ik} + \beta_{33} \text{age}_{ik} + \beta_{34} \text{PrevOI}_{ik} + \gamma y_{ik}^*(t) + W_k \quad (6)$$

## Results

$\pi(\theta)$	$W_k$	Measure
<b>Common Cure</b> $\theta_k = \theta, \forall k$ $\theta \sim \mathcal{U}(0, 1)$	0	14867 -7473
	$W_k$	<b>14540(SM)</b> -7453(SM)
<b>Regional Cure</b> $\theta_k \neq \theta_l, \forall k \neq l$ $\theta_k \sim \mathcal{U}(0, 1)$	0	14578 -7482
	$W_k$	14596 -7487
<b>No Cure</b> $1 - \theta_k = 0, \forall k$	0	14575 -7461
	$W_k$	14577 -7496

**Model choice:** Watanabe-Akaike Information Criterion (WAIC); Logarithm of the Pseudo Marginal Likelihood (LPML).



**Spatial frailties** - Brazil's map presenting the posterior median of the region-specific relative risks,  $\exp\{W_k\}$ .

	Mean	95% CI
$(\beta_{10})$	14.65	(8.37, 18.78)
$(\beta_{11})$ sex	-0.15	(-2.66, 2.08)
$(\beta_{12})$ age	-1.30	(-4.25, 1.30)
$(\beta_{13})$ PrevOI	-1.84	(-3.67, -0.01)
$\sigma_e^2$	4.99	(4.54, 5.26)
$(\beta_{31})$	-3.14	(-4.42, -1.87)
$(\beta_{32})$ sex	0.47	(-0.35, 1.36)
$(\beta_{33})$ age	0.58	(-0.45, 1.61)
$(\beta_{34})$ PrevOI	0.67	(-0.15, 1.49)
$\gamma$	-0.17	(-0.25, -0.11)
$\sigma_W^2$	0.015	(0.0002, 0.13)
$\rho$	2.34	(1.67, 3.11)
$1 - \theta$	0.17	(0.01, 0.47)

**Parameters:** posterior mean and its 95% Credibility Interval (CI).

The model choice measures, WAIC and LPML, agree in the the selected model (SM)  $\Rightarrow$  Considering a common cure fraction for all patients and a spatial frailty outperforms the traditional joint model approach for the HIV/AIDS data, which does not considers a fraction of long-term survivors. The estimated cure fraction,  $(1 - \theta)$ , indicates that nearly 17% of the HIV/AIDS patients in this study may be considered long-term survivors.