Bayesian analysis for heavy-tailed nonlinear mixed effects models

Cibele M. Russo

in collaboration with Danilo A. Silva

Universidade de São Paulo, Brazil 오

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- Motivating example
- Nonlinear mixed effects models
- Heavy-tailed nonlinear mixed effects models
- Application
- Discussion and remarks
- Work in progress
- Bibliography

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- We assume that the random effects and errors jointly follow heavy tailed distributions for the random effects and errors.
- This assumption may produce more robust estimates against outlying or influential observations (see, for instance, Meza et al., 2011 and Russo et al., 2009).

Motivating example: Theophylline concentration

Kinetics of anti-asthmatic agent theophylline (Pinheiro & Bates 2000)

- Y: Theophylline Concentration (mg / L) (response variable)
- T: Time (h) (covariate)
- D: Applied dose (mg / kg)

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$$\mathsf{E}(Y) = D\exp(IK_e + IK_a - IC_l)\frac{[\exp(-e^{IK_e}T) - \exp(-e^{IK_a}T)]}{e^{IK_a} - e^{IK_e}}$$

- *IK_a*: logarithm of the absorption rate,
- *IK_e*: logarithm of the elimination rate
- IC_I : the logarithm of clearance.

Motivating example: Theophylline concentration

In an experiment, serum concentration (in mg/L) of the ophylline was measured in eleven times (in h) after the administration of D dose (in mg/kg) in each of twelve patients.

Motivating example: Theophylline data



Theophylline concentration versus time

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Bayesian analysis for nonlinear models

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For the *j*th measurement taken in the *i*th subject, Y_{ij} , in the time T_{ij} , a possible mixed effects nonlinear model would be

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For the *j*th measurement taken in the *i*th subject, Y_{ij} , in the time T_{ij} , a possible mixed effects nonlinear model would be

$$Y_{ij} = D \exp(\phi_{1i} + \phi_{2i} - \phi_{3i}) rac{[\exp(-e^{\phi_{1i}}T_{ij}) - \exp(-e^{\phi_{2i}}T_{ij})]}{e^{\phi_{2i}} - e^{\phi_{1i}}} + \epsilon_{ij}$$

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where

- $\phi_{1i} = IK_e + b_{1i}$
- $\phi_{2i} = IK_a + b_{2i}$
- $\phi_{3i} = IC_I + b_{3i}$
- IKe, IKa and ICI are fixed effects
- b_{1i} , b_{2i} and b_{3i} are random effects
- ϵ_{ij} are random errors.

It is usual to assume that the random effects and errors are independent and normally distributed.

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Here we assume the random effects and errors to jointly follow a multivariate scale mixture of normal distributions.

Interpretation of random effects in the nonlinear model



Interpretation of random effects in the nonlinear model

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Nonlinear mixed-effects models

Suppose that $\mathbf{y} = (\mathbf{y}_1^\top, \dots, \mathbf{y}_n^\top)^\top$ is a vector of observed continuous multivariate responses with \mathbf{y}_i a $(n_i \times 1)$ vector containing the observations for the experimental unit $i, i = 1, \dots, n$, such that

$$\mathbf{y}_i = \mathbf{g}(\phi_i, \mathbf{X}_i) + \epsilon_i, \ i = 1, \dots, n,$$

 $\phi_i = \boldsymbol{\beta} + \mathbf{b}_i,$ (1)

with

- X_i = (X_{i1},..., X_{ini})[⊤] a matrix of explanatory variables for the *i*-th unit,
- \mathbf{b}_i is a $(q \times 1)$ vector of random effects,
- ϵ_i is an $(n_i \times 1)$ vector of random errors values for i = 1, ..., n,
- β is a $(p \times 1)$ location vector.

Let a m-dimensional random vector **W** follows a scale mixture of normal distribution (SMN) in its stochastic form, that is

$$\mathbf{W} = \boldsymbol{\mu} + \kappa(U)^{1/2} \mathbf{Z},$$

where

- µ is the location vector,
- U is a positive random variable with cumulative distribution function (cdf) H(u, ν) and probability density function (pdf) h(u, ν),
- ν is a scalar or vector parameter indexing the distribution of U,
- $\kappa(U)$ is the weight function,
- Z \sim N(0, Σ)
- Z and U independent.

Characterization of some SMN distributions.							
Distribution	$\kappa(u)$	U					
$MN_m(\mu, \mathbf{\Sigma})$	1	U = 1					
$MSt_m(\mu, \mathbf{\Sigma}, \nu)$	$\frac{1}{u}$	$egin{aligned} & U \stackrel{ ext{ind.}}{\sim} Gamma\left(rac{ u}{2},rac{ u}{2} ight), \ & u > 0, u > 0 \end{aligned}$					
$MSI_m(oldsymbol{\mu}, oldsymbol{\Sigma}, u)$	$\frac{1}{u}$	$egin{array}{ll} U\sim {\it Beta}(u,1),\ 0< u<1, u>0 \end{array}$					
$MCN_m(\boldsymbol{\mu}, \boldsymbol{\Sigma}, \nu, \gamma)$	$\frac{1}{u}$	$egin{aligned} h(u, oldsymbol{ u}) &= u \mathbb{I}_{(u=\gamma)} + (1- u) \mathbb{I}_{(u=1)}, \ u &= \gamma, 1; 0 \leq u \leq 1, \ 0 \leq \gamma \leq 1, \end{aligned}$					
with $d = (\mathbf{y} - \boldsymbol{\mu})^{ op} \boldsymbol{\Sigma}^{-1} (\mathbf{y} - \boldsymbol{\mu})$							

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Scale mixture of normal distributions include symmetrical distributions with heavier tailed than the Gaussian distributions, including the normal itself.



Normal and Student-t distributions

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Bayesian analysis for nonlinear models

In the nonlinear model

$$\mathbf{y}_i = \mathbf{g}(\phi_i, \mathbf{X}_i) + \epsilon_i, \ i = 1, \dots, n,$$

 $\phi_i = \beta + \mathbf{b}_i,$

we assume that

$$\begin{pmatrix} \epsilon_i \\ \mathbf{b}_i \end{pmatrix} \stackrel{ind.}{\sim} \mathrm{SMN}_{n_i+q} \left(\begin{pmatrix} \mathbf{0} \\ \mathbf{0} \end{pmatrix}, \begin{pmatrix} \mathbf{\Sigma}_i & \mathbf{0} \\ \mathbf{0} & \mathbf{D} \end{pmatrix}; \mathrm{H} \right), \quad (2)$$

where **D** and Σ_i are positive-definite dispersion matrices. For simplicity, we assume that **D** = diag(τ) and $\Sigma_i = \sigma^2 \mathbf{I}_{n_i}$ for i = 1, ..., n and $\sigma > 0$ a scalar.

Application

In the nonlinear model for the theophylline data: $\boldsymbol{\theta} = (\boldsymbol{\beta}, \sigma^2, \boldsymbol{\tau})^\top = (\mathit{IKe}, \mathit{IKa}, \mathit{ICI}, \sigma^2, \tau_1, \tau_2, \tau_3)^\top$

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- $\beta_i \sim N(0, v)$,
- $\tau_i \sim \text{Gamma}(a_i, b_i)$,
- $\sigma^2 \sim \text{Gamma}(c, d)$.

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The Monte Carlo estimates using OpenBUGS were obtained generating chains of size 30000 spaced by 50 (burn-in of 5000). The posterior means and the corresponding standard deviations of the posterior distributions are presented in the following slides.

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Results: Theophylline data

	Normal		Student-t ₄		Slash ₄	
	mean	(sd)	mean	(sd)	mean	(sd)
IK _e	-2.465	(0.102)	-2.452	(0.099)	-2.450	(0.099)
IK _a	0.462	(0.142)	0.460	(0.146)	0.459	(0.147)
IC_{I}	-3.232	(0.095)	-3.221	(0.097)	-3.220	(0.094)
$(\sigma^2)^{-1}$	0.636	(0.104)	0.552	(0.145)	0.123	(0.035)
$(\tau_1)^{-1}$	5.580	(1.691)	5.053	(1.492)	5.052	(1.505)
$(\tau_2)^{-1}$	13.060	(3.458)	13.150	(3.451)	13.210	(3.451)
$(au_{3})^{-1}$	11.830	(3.087)	11.740	(3.082)	11.760	(3.085)
DIC	323.5		303.6		298.3	

Results: Theophylline data



Fitted curves

Fitted curves for theophylline problem under Slash₄ model.

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Convergence



Convergence of chains in Slash₄ case

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Implementation

• Easy implementation in OpenBUGS,

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Implementation

- Easy implementation in OpenBUGS,
- Computationally expensive,

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Implementation

- Easy implementation in OpenBUGS,
- Computationally expensive,
- For nonlinear mixed effects models, convergence may be difficult to reach depending on the initial values.

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- Considering a stochastic formulation in a bayesian approach, we propose the use of heavy-tailed distributions in a bayesian context to provide alternatives to the gaussian model.

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- Easy implementation in Bayesian approach.

- Nonlinear mixed effects models plays an important role in nonlinear problems with correlated data.
- Considering a stochastic formulation in a bayesian approach, we propose the use of heavy-tailed distributions in a bayesian context to provide alternatives to the gaussian model.
- Easy implementation in Bayesian approach.
- Estimation of individual profiles.

Work in progress

- Model diagnostics
- Robustness assessment
- Other nonlinear data sets.

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References

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Thank you!