



Problems of INLA in generalized linear mixed models for binary responses

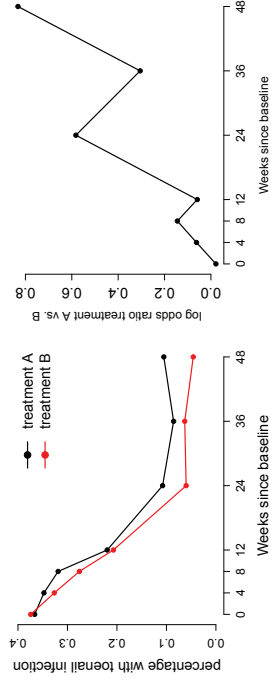
11. September 2013

Rafael Sauter and Leonhard Held



Toenail infection data (De Backer et al., 1998)

- Randomized, double-blinded study comparing two oral treatments for toenail infection.
- Binary outcome: absent, mild (0) or moderate, severe (1).
- $N = 289$ patients with 1903 observations.





GLMM for toenail infection data

- Patients $i = 1, \dots, N$, sessions $j = 1, \dots, n_i$.
 $y_{ij} \mid \mathbf{x}_{ij}, \mathbf{z}_{ij}, \mathbf{b}_i \sim \text{Bernoulli}(\mu_{ij})$
 $\text{logit}(\mu_{ij}) = \mathbf{x}_{ij}^T \boldsymbol{\beta} + \mathbf{z}_{ij}^T \mathbf{b}_i$
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– time_{ij} is measured in months since baseline.
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- Random intercept, random slope: $\mathbf{z}_{ij} = (1, t_{ij})^\top$
covariance matrix $\mathbf{D} = \begin{pmatrix} \sigma_{z_1}^2 & \rho\sigma_{z_1}\sigma_{z_2} \\ \rho\sigma_{z_2}\sigma_{z_1} & \sigma_{z_2}^2 \end{pmatrix}$.



ML-estimation with Laplace Approximation

Marginal likelihood contribution for patient i :

$$L_i(\mathbf{y}_i \mid \boldsymbol{\beta}, \mathbf{D}) = \int \pi(\mathbf{y}_i \mid \boldsymbol{\beta}, \mathbf{b}_i) \pi(\mathbf{b}_i \mid \mathbf{D}) d\mathbf{b}_i$$



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Iterate:

1. Find the mode $\hat{\mathbf{b}}_i = \arg \max_{\mathbf{b}_i} \pi(\mathbf{y}_i | \hat{\boldsymbol{\beta}}, \mathbf{b}_i) \pi(\mathbf{b}_i | \hat{\mathbf{D}})$ with penalized iterative reweighted least squares.
2. Maximize Laplace-approximation of $L_i(\mathbf{y}_i | \boldsymbol{\beta}, \mathbf{D})$ around $\hat{\mathbf{b}}_i$ and obtain $\hat{\boldsymbol{\beta}}, \hat{\mathbf{D}}$.



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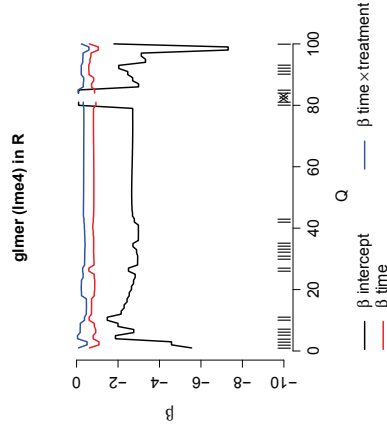
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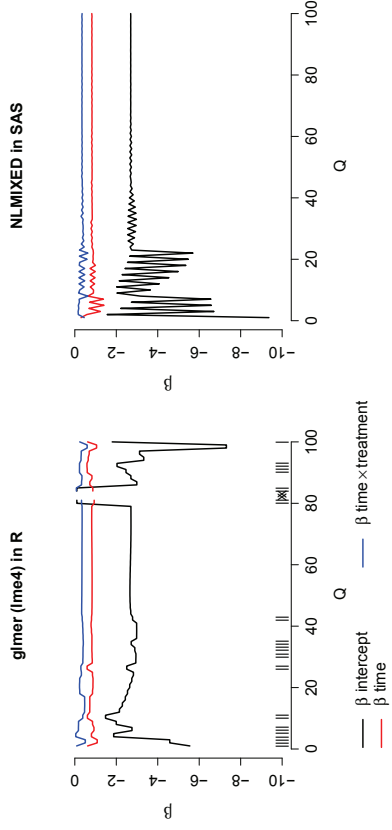
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- Zhang et al. (2011) simulation study comparing statistical packages for GLMM with binary response.

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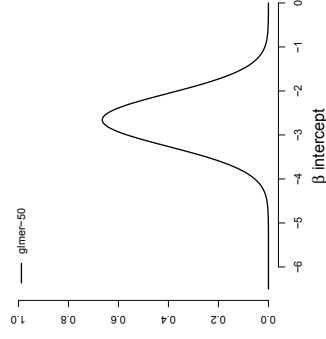
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2. $\tilde{\pi}(\gamma_k | \boldsymbol{\theta}_u, \mathbf{y})$ Gaussian, simplified or full Laplace approximation for $\pi(\gamma_k | \boldsymbol{\theta}_u, \mathbf{y})$.

Intercept for GLMER, INLA, MCMC fixed θ

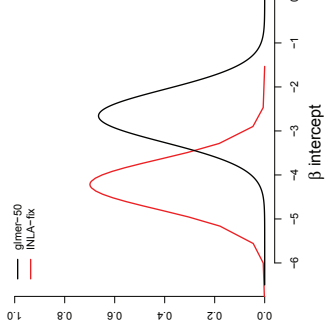
- **glmer-50**: ML-estimation with GH approximation using `g_lmer` in package `lme4` by Bates et al. (2011) with $Q=50$.





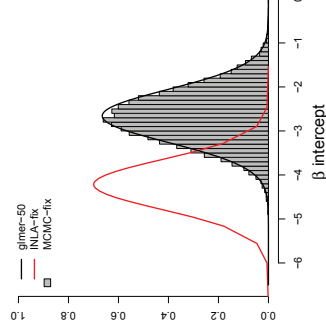
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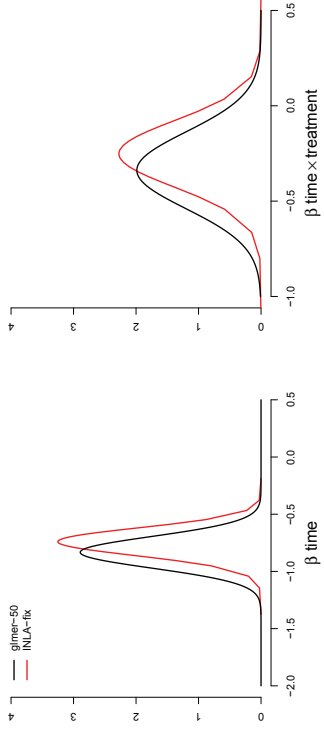
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- **MCMC**: MCMC sampling with fixed θ at ML-estimates.

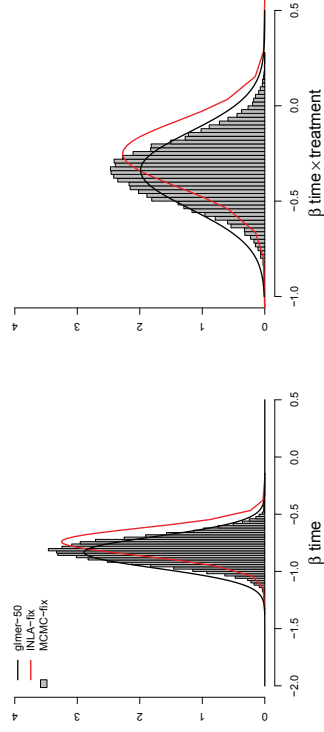




Time, time:treatment for GLMER, INLA, MCMC fixed θ

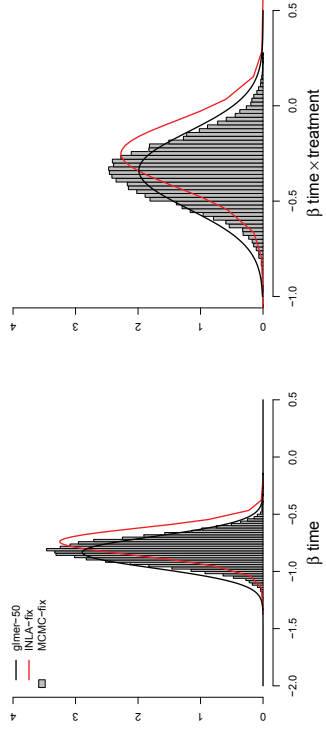


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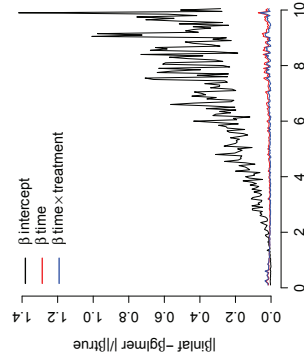
Time, time:treatment for GLMER, INLA, MCMC fixed θ



- 95%-CI GLMER: [-0.73, 0.06]
- 95%-CI INLA: [-0.60, 0.09]
- 95%-CI MCMC: [-0.66, -0.03]



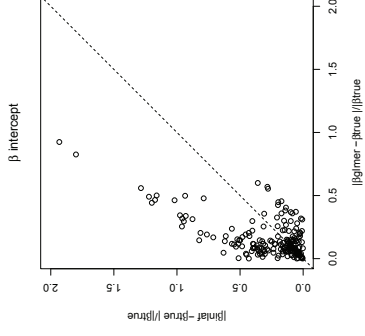
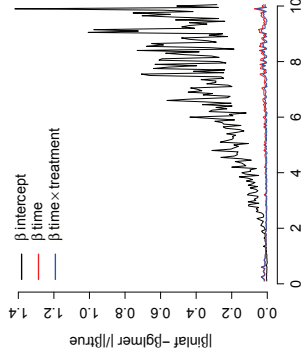
Simulation for a RI model



- Fixed effects: $\beta_0 = -2.5$, $\beta_1 = -0.8$, $\beta_2 = 0.8$.
- Random effect standard deviation σ_{ϵ_1} : 0.1 to 10 by 0.05.
- 290 patients for 7 visits at times 0, 1, 2, 3, 6, 9, 12.
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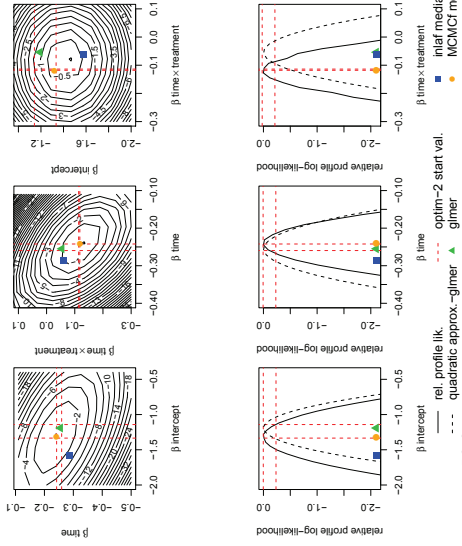
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- Is there a better way to approximate the likelihood?
- Use INLA with fixed hyperparameters and compare...
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- Approximation of the Likelihood with INLA resulted in shifted marginal posterior distributions.



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Profiled relative log-likelihood surface



Computational time for GLMER and INLA

