

Evaluating the log-likelihood in nonlinear mixed models

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Outline

1 Computation for LMMs

Introduction

- Nonlinear mixed-effects models (NLMMs) are widely used in the analysis of pharmacokinetic/pharmacodynamic data.
- Over the years many algorithms for determining the parameter estimates in these models have been proposed and several implementations of algorithms have been developed.
- Many of these algorithms concentrate on the details of optimization of something like the log-likelihood. It is not always clear that such approaches should be expected to yield estimates close to the maximum likelihood estimates.
- Recent developments in computational methods for linear mixed-effects models (LMMs) provide effective ways of evaluating good approximations to the log-likelihood of an NLMM.

Outline

1 Computation for LMMs

Evaluating the deviance function

- The *profiled deviance* function for a LMM can be expressed as a function the variance-component parameters only. We can profile out the residual variance and the fixed-effects parameters.
- We describe the probability model in terms of the n -dimensional response random variation, \mathbf{Y} , whose value, \mathbf{y} , is observed, and the q -dimensional, unobserved random effects variable, \mathbf{B} , with distributions

$$(\mathbf{Y}|\mathbf{B} = \mathbf{b}) \sim \mathcal{N}(\mathbf{Z}\mathbf{b} + \mathbf{X}\boldsymbol{\beta}, \sigma^2 \mathbf{I}_n), \quad \mathbf{B} \sim \mathcal{N}(\mathbf{0}, \boldsymbol{\Sigma}_\theta),$$

- We never really form $\boldsymbol{\Sigma}_\theta$; we always work with the *relative covariance factor*, $\boldsymbol{\Lambda}_\theta$, defined so that

$$\boldsymbol{\Sigma}_\theta = \sigma^2 \boldsymbol{\Lambda}_\theta \boldsymbol{\Lambda}_\theta^\top.$$

Note that we must allow for $\boldsymbol{\Lambda}_\theta$ to be less than full rank.

Orthogonal or “unit” random effects

- We will define a q -dimensional “spherical” or “unit” random-effects vector, \mathbf{u} , such that

$$\mathbf{u} \sim \mathcal{N}(\mathbf{0}, \sigma^2 \mathbf{I}_q), \mathbf{B} = \mathbf{\Lambda}_\theta \mathbf{u} \Rightarrow \text{Var}(\mathbf{B}) = \sigma^2 \mathbf{\Lambda}_\theta \mathbf{\Lambda}_\theta^\top = \mathbf{\Sigma}_\theta.$$

- The linear predictor expression becomes

$$\mathbf{Z}\mathbf{b} + \mathbf{X}\boldsymbol{\beta} = \mathbf{Z}\mathbf{\Lambda}_\theta \mathbf{u} + \mathbf{X}\boldsymbol{\beta} = \mathbf{U}_\theta \mathbf{u} + \mathbf{X}\boldsymbol{\beta}$$

where $\mathbf{U}_\theta = \mathbf{Z}\mathbf{\Lambda}_\theta$.

- The key to evaluating the log-likelihood is the Cholesky factorization

$$\mathbf{L}_\theta \mathbf{L}_\theta^\top = \mathbf{P} (\mathbf{U}_\theta^\top \mathbf{U}_\theta + \mathbf{I}_q) \mathbf{P}^\top$$

(\mathbf{P} is a fixed permutation that has practical importance but can be ignored in theoretical derivations). The sparse, lower-triangular \mathbf{L}_θ can be evaluated and can be updated when $\boldsymbol{\theta}$ is changed, even when q is in the millions and the model involves random effects for several factors.

The profiled deviance

- The Cholesky factor, \mathbf{L}_θ , allows evaluation of the conditional mode $\tilde{\mathbf{u}}_{\theta,\beta}$ (also the conditional mean for linear mixed models) from

$$(\mathbf{U}_\theta^\top \mathbf{U}_\theta + \mathbf{I}_q) \tilde{\mathbf{u}}_{\theta,\beta} = \mathbf{P}^\top \mathbf{L}_\theta \mathbf{L}_\theta^\top \mathbf{P} \tilde{\mathbf{u}}_{\theta,\beta} = \mathbf{U}_\theta^\top (\mathbf{y} - \mathbf{X}\boldsymbol{\beta})$$

Let $r^2(\boldsymbol{\theta}, \boldsymbol{\beta}) = \|\mathbf{y} - \mathbf{X}\boldsymbol{\beta} - \mathbf{U}_\theta \tilde{\mathbf{u}}_{\theta,\beta}\|^2 + \|\tilde{\mathbf{u}}_{\theta,\beta}\|^2$.

- $\ell(\boldsymbol{\theta}, \boldsymbol{\beta}, \sigma | \mathbf{y}) = \log L(\boldsymbol{\theta}, \boldsymbol{\beta}, \sigma | \mathbf{y})$ can be written

$$-2\ell(\boldsymbol{\theta}, \boldsymbol{\beta}, \sigma | \mathbf{y}) = n \log(2\pi\sigma^2) + \frac{r^2(\boldsymbol{\theta}, \boldsymbol{\beta})}{\sigma^2} + \log(|\mathbf{L}_\theta|^2)$$

- The conditional estimate of σ^2 is

$$\widehat{\sigma^2}(\boldsymbol{\theta}, \boldsymbol{\beta}) = \frac{r^2(\boldsymbol{\theta}, \boldsymbol{\beta})}{n}$$

producing the *profiled deviance*

$$-2\tilde{\ell}(\boldsymbol{\theta}, \boldsymbol{\beta} | \mathbf{y}) = \log(|\mathbf{L}_\theta|^2) + n \left[1 + \log \left(\frac{2\pi r^2(\boldsymbol{\theta}, \boldsymbol{\beta})}{n} \right) \right]$$

Profiling the deviance with respect to β

- Because the deviance depends on β only through $r^2(\theta, \beta)$ we can obtain the conditional estimate, $\hat{\beta}_\theta$, by extending the PLS problem to

$$r^2(\theta) = \min_{\mathbf{u}, \beta} \left[\|\mathbf{y} - \mathbf{X}\beta - \mathbf{U}_\theta \mathbf{u}\|^2 + \|\mathbf{u}\|^2 \right]$$

with the solution satisfying the equations

$$\begin{bmatrix} \mathbf{U}_\theta^\top \mathbf{U}_\theta + \mathbf{I}_q & \mathbf{U}_\theta^\top \mathbf{X} \\ \mathbf{X}^\top \mathbf{U}_\theta & \mathbf{X}^\top \mathbf{X} \end{bmatrix} \begin{bmatrix} \tilde{\mathbf{u}}_\theta \\ \hat{\beta}_\theta \end{bmatrix} = \begin{bmatrix} \mathbf{U}_\theta^\top \mathbf{y} \\ \mathbf{X}^\top \mathbf{y} \end{bmatrix}$$

- The profiled deviance, which is a function of θ only, is

$$-2\tilde{\ell}(\theta) = \log(|\mathbf{L}_\theta|^2) + n \left[1 + \log \left(\frac{2\pi r^2(\theta)}{n} \right) \right]$$