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An *R* implementation of bootstrap procedures for mixed models

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Outline

- Introduction and motivation
- Bootstrap methods for Mixed Models
- Implementation details
- Some examples
- Conclusions

(Generalized) Linear Mixed Models

- Repeated measures or Longitudinal data:

Response vector Y_i for i^{th} subject $Y_i = (Y_{i1}, \dots, Y_{in_i})'$

Observations on the same unit can be correlated

- Conditional / Hierarchical approach:

Between-subject variability explained by Random-effects b_i
usually with Normal distribution $b_i \sim N(0, D)$

$$E(Y_{ij} | b_i) = \mu_{ij}$$

$$g(\mu_{ij}) = \theta_{ij} = X_{ij}\beta + Z_{ij}b_i$$

Estimation in (G)LMM

- Random-effects are not directly observed
- Estimation of parameters based on Marginal Likelihood, after integration of Random-effects

$$L(\beta, D, \phi | Y) = \prod_{i=1}^n f_i(Y_i | \beta, D, \phi) = \\ \prod_{i=1}^n \int \prod_{j=1}^{n_i} f_{ij}(Y_{ij} | \beta, b_i, \phi) f(b_i | D) db_i$$

- MLE:
 - Analytic solution in the Normal case (Linear Mixed Models)
 - Approximations are needed in the general case.



- *lme4* package: common framework for L-GL-NL/MM
 - **Fast and efficient** estimation for ML and REML criteria via Laplace Approximation/Adaptive Gaussian Quadrature for GLMM.

Inference (G)LMM

- Wald-type and F-tests (summary)
 - Asymptotic standard errors for the fixed effects parameters
- Likelihood ratio test (anova)
 - Comparison of likelihood of two models
- Bayesian Inference (`mcmc samp`)
 - MCMC sampling procedure for posteriors on parameters
- Some drawbacks:
 - Asymptotic results
 - Degrees of freedom of the reference distribution in F-test
 - Likelihood Ratio test can be conservative under some conditions
 - Tests on Variance components close to the boundary of the parameter space.

Motivation

- Inference based on bootstrap for LMM and GLMM
- Inference on functions of parameters
 - i.e. confidence intervals and hypothesis test for ratio of variance components
- Robust approaches
 - i.e. in presence of influential data and outliers
- Effect of misspecification
 - i.e. non-gaussian random effects and/or residuals

Extension of the package `lmer`

- ***merBoot*** provides methods for Monte Carlo and bootstrap techniques in generalized and linear mixed-effects models
- The implementation is **object-oriented**
- It takes profit of specificities of the applied algorithms to enhance **efficiency**, using less time and memory.
- It has a **flexible** interface to design complex experiments.

Bootstrap in linear models

- For (Generalized) linear models (without random effects) there is only one random component → generation of the response variable according to the conditional mean.

$$\mu = X\beta \quad Y \sim N(\mu, \sigma)$$

$$\mu = g^{-1}(X\beta) \quad Y \sim F_\mu$$

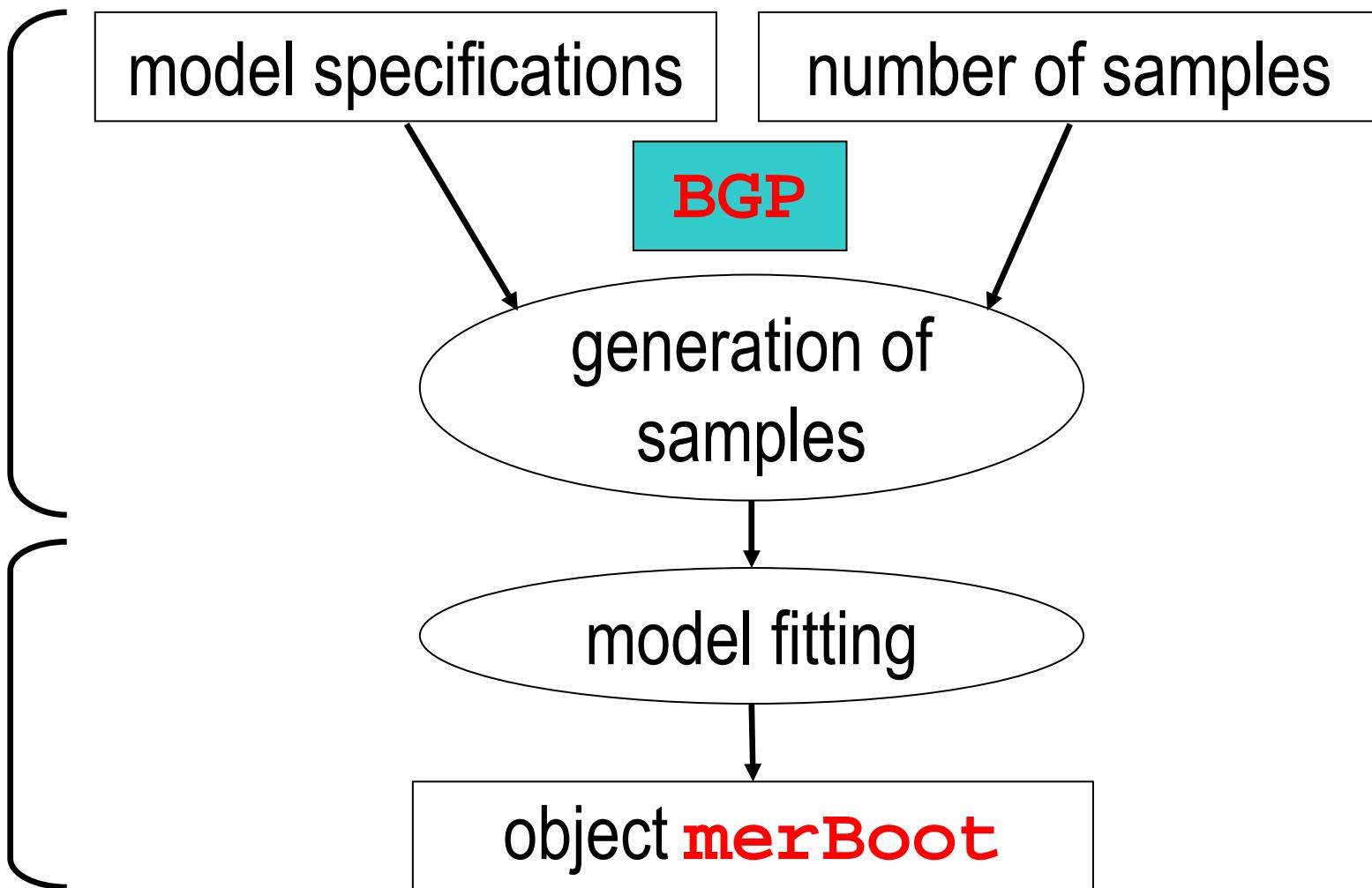
- Residual resampling:
 - Estimate parameters for the systematic part of the model
 - Resample random part of the model (parametric or empirical)
 - Some variants to deal with heterocedasticity (Wild bootstrap)

Bootstrap in Mixed Models

- In Mixed models, the systematic part has a random component → generation of the response variable in two steps:
 - Bootstrap of the conditional mean (function of the linear predictor)
 - Bootstrap of the response variable
$$\mu = X\beta + Zb \quad b \sim N(0, \theta) \quad Y \sim N(\mu, \sigma)$$
$$\mu = g^{-1}(X\beta + Zb) \quad b \sim N(0, \theta) \quad Y \sim F_\mu$$
- Two objects in the `merBoot` implementation:
 - **BGP**: Set-up for the Bootstrap Generation Process
 - **merBoot**: Coefficients for the resamples and methods for analysis

Implementation details

Step I Step II



Bootstrap Generation Process

Linear predictor level

- Fixed parameters β
- Design matrices $X_i Z_i$
- Random effects generator (b_i^*)
 - **Parametric**: generating b_i^* from a multivariate gaussian distribution
 - **Semiparametric/Empirical** (from a fitted object): sampling b_i^* from \hat{b}_i with replacement.
 - **User-defined**: any other distribution/criteria to generate b_i^*

$$\eta_{ij}^* = X_{ij}\beta + Z_{ij}b_i^*$$

Bootstrap Generation Process

Response level

- Family (distribution F + link function g)
- Response generator (Y_{ij}^*)
 - **Parametric**: $\mu_{ij}^* = g^{-1}(\eta_{ij}^*)$, sample $Y_{ij}^* \sim F(\cdot; \mu_{ij}^*)$
 - **Semiparametric/Empirical** (from a fitted object):
 - **Residual-based**: builds Y_{ij}^* like in linear heterocedastic models, depending on type of residuals
$$Y_{ij}^* = \mu_{ij}^* + \varepsilon_i^*$$
 - **Distribution-based**: resamples estimated quantiles

Residuals in GLM

- Raw residuals:

$$Y_{ij} - \hat{\mu}_{ij}$$

- Pearson residuals:

$$e_{ij} = \frac{Y_{ij} - \hat{\mu}_{ij}}{\sqrt{a_{ij}V(\hat{\mu}_{ij})}}$$

- Standardized Pearson residuals:

$$r_{ij} = \frac{e_{ij}}{\sqrt{1 - h(\hat{\mu}_{ij})}}$$

- Standardized residuals

on the linear predictor scale:

$$l_{ij} = \frac{g(Y_{ij}) - g(\hat{\mu}_{ij})}{\sqrt{a_{ij}g'(\hat{\mu}_{ij})^2V(\hat{\mu}_{ij})(1 - h_{ij})}}$$

- Deviance residuals:

$$r_{ij} = sign(Y_{ij} - \hat{\mu}_{ij})\sqrt{d_{ij}}$$

Empirical residual-based

- Standardized Pearson residuals:
 - Resample e_{ij}^* from centered e_{ij}
 - Calculate $Y_{ij}^* = \mu_{ij}^* + \left(\sqrt{\frac{\hat{\phi}}{a_i} V(\mu_{ij}^*)} \right) e_{ij}^*$
- Standardized Pearson residuals on the linear predictor scale:
 - Resample l_{ij}^* from l_{ij}
 - Calculate $Y_{ij}^* = g^{-1} \left(\eta_{ij}^* + \left(g'(\eta_{ij}^*) \sqrt{\frac{\hat{\phi}}{a_i} V(\mu_{ij}^*)} \right) l_{ij}^* \right)$

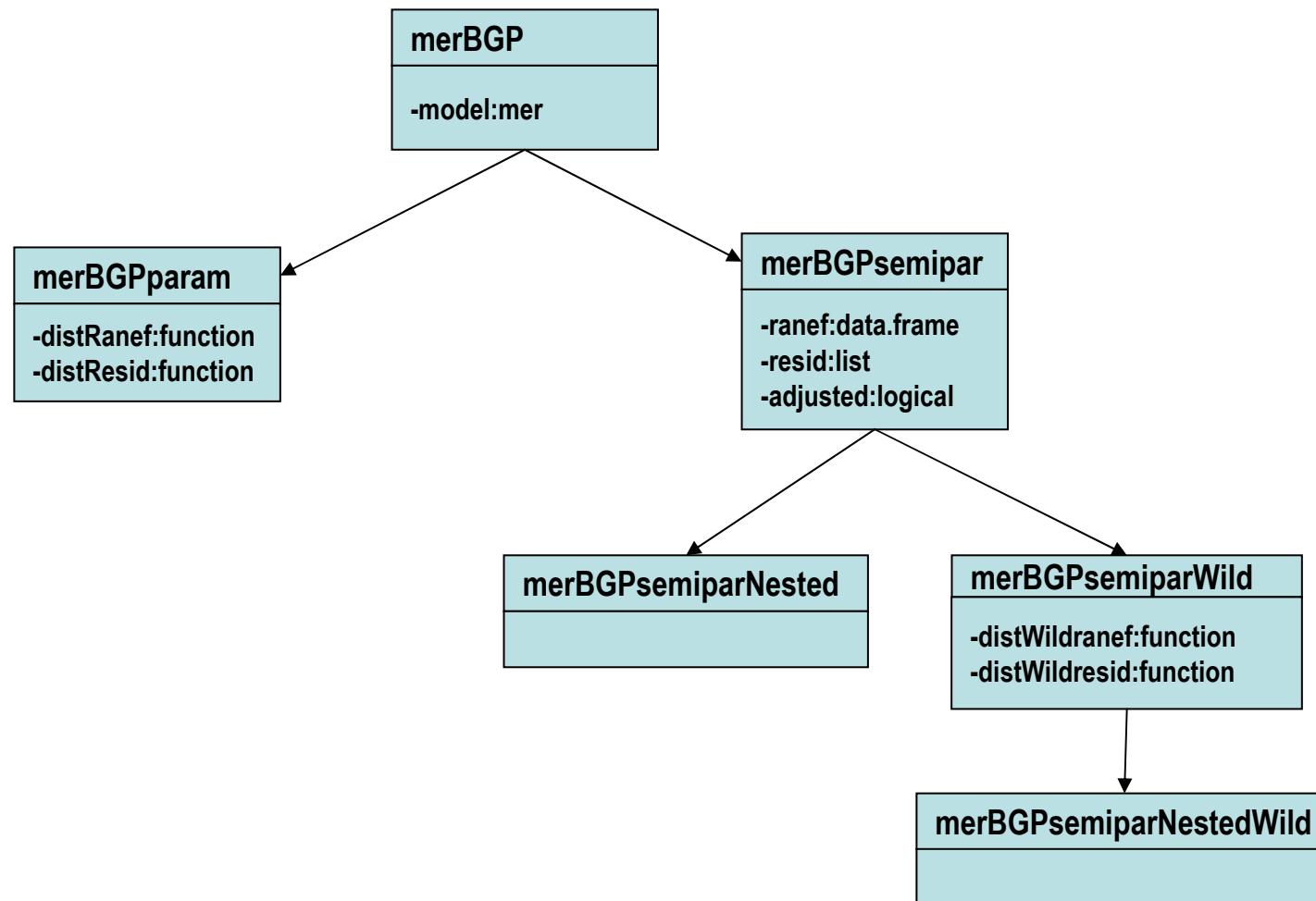
Empirical distribution-based

- Randomized Quantile residuals (Dunn & Smyth, 1996):
 - inverting the estimated distribution function for each observation to obtain exactly uniform (q_{ij}) or standard normal residuals ($\Phi^{-1}(q_{ij})$) for diagnostics. Randomization needed for discrete distributions.
- Resampling scheme with Quantiles Residuals for (G)LMM:
 - Calculate $\hat{q}_{ij} = F(Y_{ij}; \hat{\mu}_{ij})$
 - Sample q_{ij}^* with replacement from \hat{q}_{ij}
 - Generate $Y_{ij}^* = F^{-1}(q_{ij}^*; \hat{\mu}_{ij}^*)$

Response generation

- For Normal family and identity link function, all three strategies (pearson, linear predictor and quantile residuals) are the same.
- In all the schemes, response is rounded to the nearest valid value, according to the family considered.
- For discrete variables, **randomization** of the quantiles allows for continuous uniform residuals.
- Transformation of the random effects in order to have the first and second moments equal to the parameters (**adjusted bootstrap**).
- For all the schemes, if resample of residuals/quantiles is restricted to the subject obtained in the linear predictor level, a **nested bootstrap** is performed.

Bootstrap Generation Process



BGP Methods

- generateLinpred

- BGPparam: $b_i^* \sim F_{\hat{\theta}}$ $\eta_{ij} = X_{ij}\beta + Z_{ij}b_i^*$
- BGPsemipar: $b_i^* \sim \hat{F}_n(., b_i)$ $\eta_{ij} = X_{ij}\beta + Z_{ij}b_i^*$
- BGPsemiparWild: $b_i^* \sim \hat{F}_n(., b_i)$ $w_i^* \sim W$ $\eta_{ij} = X_{ij}\beta + Z_{ij}w_i^*b_i^*$

- generate (lmer) \rightarrow generateLinpred + Residual-based

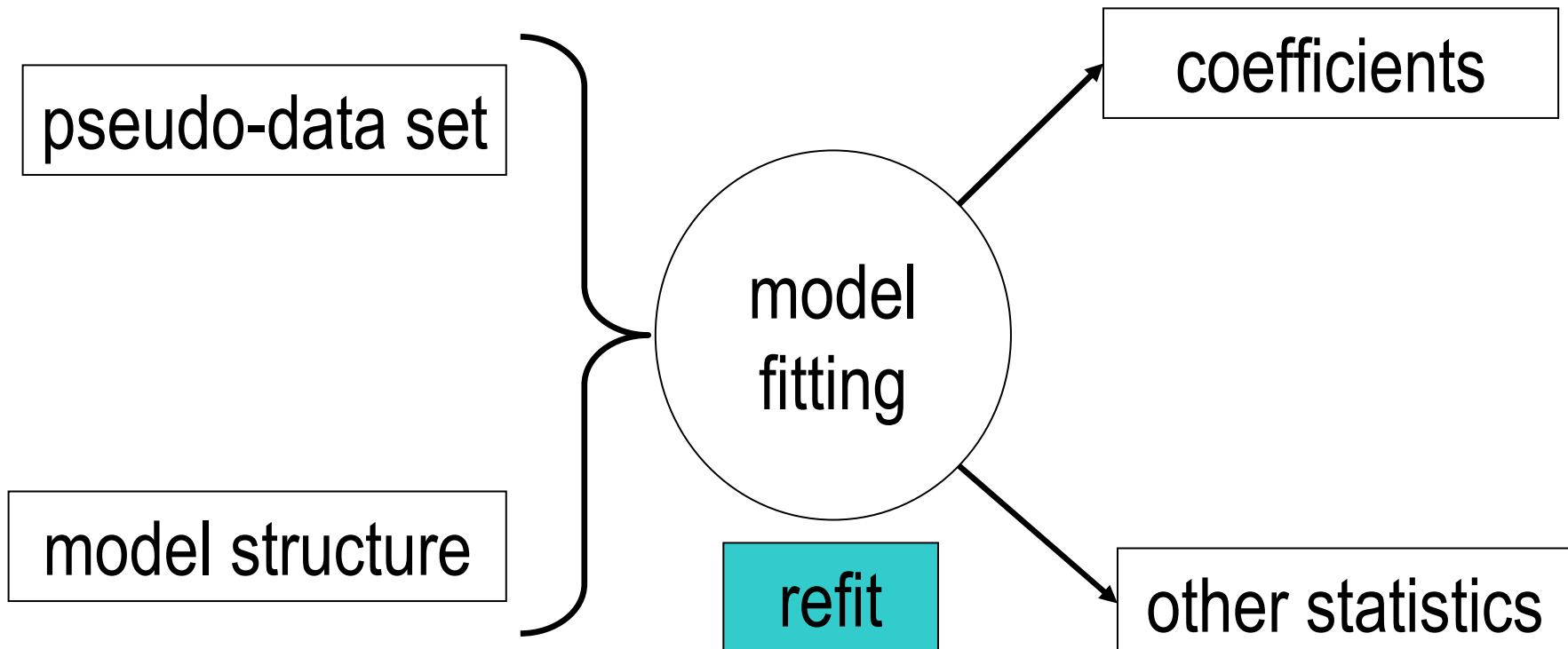
- BGPparam: $\varepsilon_{ij}^* \sim F_\theta$ $Y_{ij}^* = \mu_{ij}^* + \varepsilon_{ij}^*$
- BGPsemipar: $\varepsilon_{ij}^* \sim \hat{F}_n(., e_{ij})$ $Y_{ij}^* = \mu_{ij}^* + \varepsilon_{ij}^*$
- BGPsemiparWild: $\varepsilon_{ij}^* \sim \hat{F}_n(., e_{ij})$ $w_i^* \sim W$ $Y_{ij}^* = \mu_{ij}^* + w_i^* \varepsilon_{ij}^*$
- BGPsemiparNested $\varepsilon_{ij}^* \sim \hat{F}_n(., e_{i^*j})$ $Y_{ij}^* = \mu_{ij}^* + \varepsilon_{ij}^*$

- generate (glmer) \rightarrow generateLinpred + Distribution-based

- BGPparam:
- BGPsemipar: $q_{ij}^* \sim F_{\mu_{ij}}$ $Y_{ij}^* = F_{\mu_{ij}}^{-1}(q_{ij}^*)$
- BGPsemiparWild:
- BGPSemiparNested

Object `merBoot`

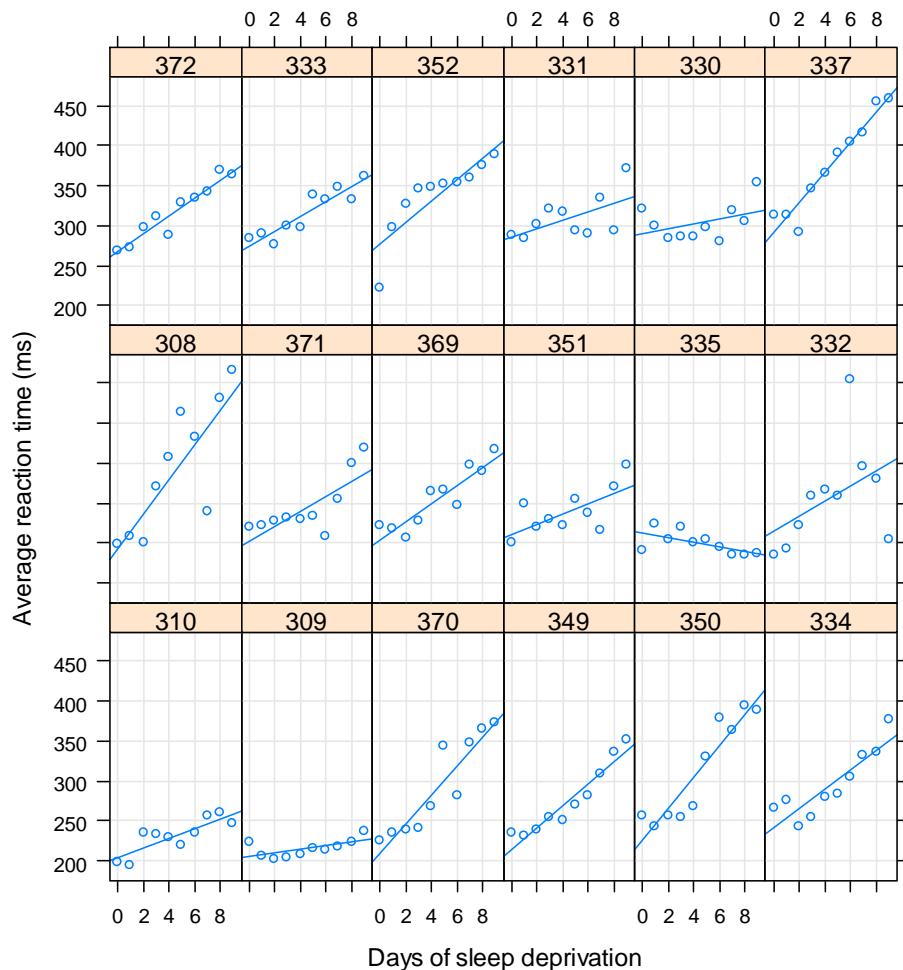
Step II: Extraction of coefficients



Bootstrap method for (g)lmer

object	<ul style="list-style-type: none">• Model specification to generate pseudo-data set (<i>glme</i> object or a list).• It contains the formula describing the structure, the parameters (β, D) and the design matrix (X, Z)
distref distres adj nest wild	<ul style="list-style-type: none">• Parameters to indicate how to generate variance components (random-effects) and response.• Strings (for pre-implemented options) or functions (user-defined methods)
B	<ul style="list-style-type: none">• Number of samples.
model2	<ul style="list-style-type: none">• Alternative model specification used to fit the pseudo-data. Default is same as object

Example: sleepstudy



```
model=lmer(Reaction~Days+(1|Subject)+  
(0+Days|Subject),sleepstudy)
```

Linear mixed model fit by REML

Formula: form

Data: sleepstudy

AIC BIC logLik deviance REMLdev
1754 1770 -871.8 1752 1744

Random effects:

Groups	Name	Variance	Std.Dev.
Subject	(Intercept)	627.568	25.0513
Subject	Days	35.858	5.9882
Residual		653.584	25.5653

Number of obs: 180, groups: Subject, 18

Fixed effects:

	Estimate	Std. Error	t value
(Intercept)	251.405	6.885	36.51
Days	10.467	1.559	6.71

Correlation of Fixed Effects:

(Intr)
Days -0.184

Methods `merBoot`: `print`

```
> sleep.boot=bootstrap(model,B=1000)
> print(sleep.boot)

lmer(formula = lmer(Reaction~Days+(1|Subject)+(0+Days|Subject),
data = sleepstudy)
```

Resampling Method: BGPparam

Bootstrap Statistics :

	original	bias	mean	std. error
(Intercept)	251.405105	0.04710710	251.452212	6.971430
Days	10.467286	-0.08913201	10.378154	1.627212
Subject.(Intercept)	25.051299	-0.56896799	24.482331	6.222008
Subject.Days	5.988173	-0.08323923	5.904934	1.231645
sigmaREML	25.565287	-0.04783798	25.517449	1.582567

Methods **merBoot**: intervals

```
> intervals(sleep.boot)
```

```
$norm
```

	lower	upper
(Intercept)	237.694245	265.021750
Days	7.367141	13.745695
Subject.(Intercept)	13.425355	37.815179
Subject.Days	3.657432	8.485393
sigmaREML	22.511352	28.714899

```
$basic
```

	lower	upper
(Intercept)	237.913080	265.836054
Days	7.351741	13.869182
Subject.(Intercept)	13.456205	37.647603
Subject.Days	3.600202	8.499955
sigmaREML	22.527143	28.691811

```
$perc
```

	lower	upper
(Intercept)	236.974156	264.897129
Days	7.065390	13.582831
Subject.(Intercept)	12.454995	36.646393
Subject.Days	3.476392	8.376144
sigmaREML	22.438764	28.603432

```
> HPDinterval(mcmcsamp(model,n=1000))
```

```
$fixef
```

	lower	upper
(Intercept)	241.741098	261.77774
Days	7.370217	13.88645
attr(,"Probability")		
[1] 0.95		

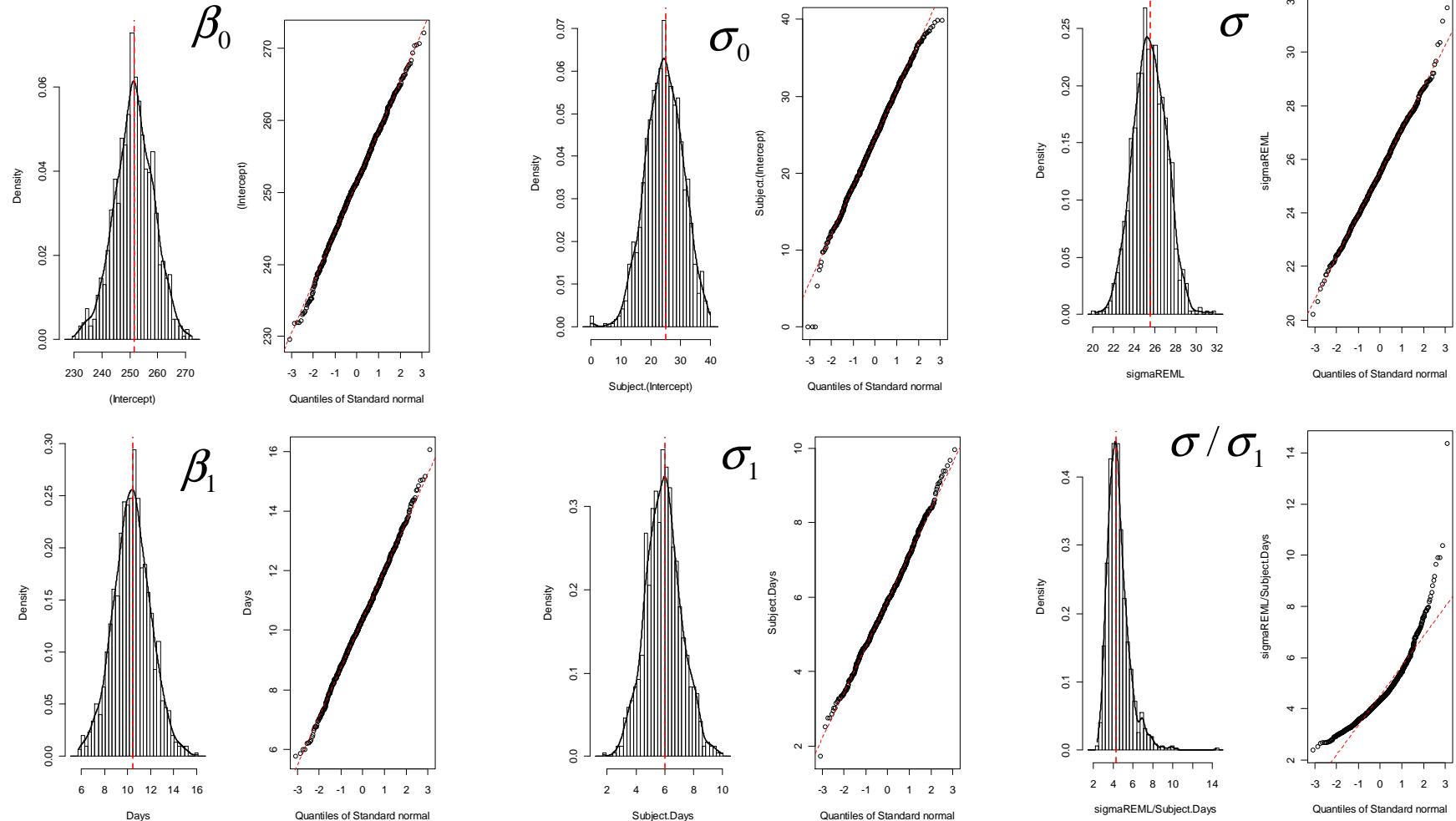
```
$ST
```

	lower	upper
[1,]	0.3117272	0.8478526
[2,]	0.1485967	0.3317120
attr(,"Probability")		
[1] 0.95		

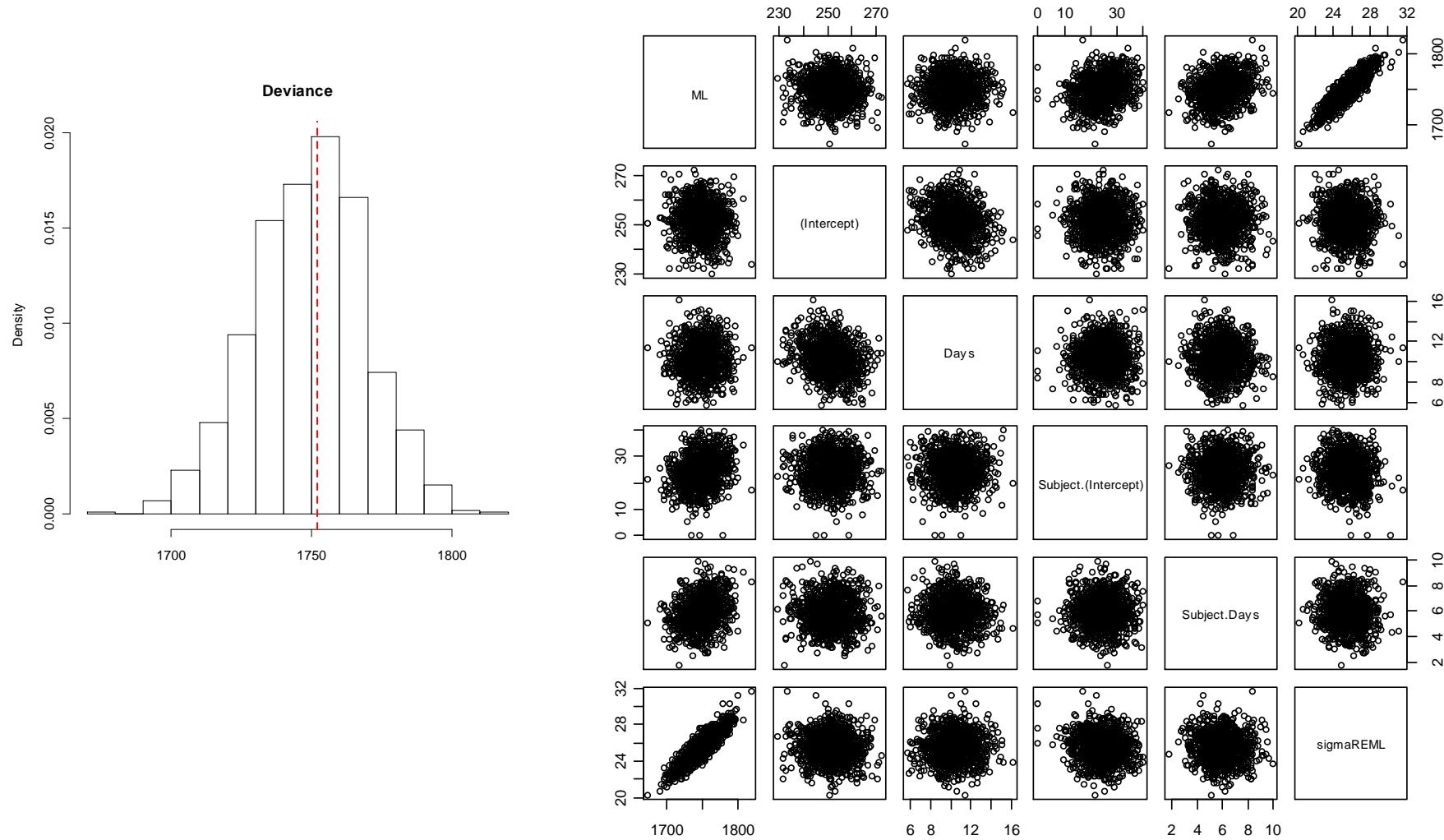
```
$sigma
```

	lower	upper
[1,]	23.88066	29.66232
attr(,"Probability")		
[1] 0.95		

Methods **merBoot**: plot



Methods **merBoot**: LR & pairs



Conclusions

- (G)LMM provide a framework to model longitudinal data for a wide range of situations (continuous, count and binary among others) but approximate estimation methods imply weaker inference.
- Monte-Carlo simulation and bootstrap can enhance inference providing empirical p-values and bootstrap estimators.
- The **BGP**/**merBoot** objects developed allow implementation of different bootstrap options and evaluation of the techniques via simulation.