

# Multilevel Models for Big Data

## Approaches for handling very large datasets

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# Table of Contents

## ① Introduction

- Multilevel Models
- Issues with Large Datasets

## ② The split-sample approach

- Pseudo Likelihood
- Graphical representation
- Independent subsamples
- Dependent subsamples
- Overlapping subsamples

## ③ R Packages

- lme4 and mgcv
- Why use bam()?
- When to use bam()?

## ④ Summary

## ⑤ References

## What are multilevel models?

- Hierarchical Structure
- Residual Components
- Variance Partitioning

## Why use multilevel models?

- Correct Inferences
- Group Effects Estimation
- Simultaneous Estimation
- Generalization Beyond Sample <sup>1</sup>

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<sup>1</sup>Centre for Multilevel Modelling 2025.

# Introduction — Issues with Large Datasets

As hierarchical data scales ( $N$  groups  $\times$   $n$  individuals), massive datasets create the following issues:

- High number of groups
- Large group sizes<sup>2</sup>
- Design matrix construction<sup>3</sup>

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<sup>2</sup>Clark 2019; Speelman, Heylen, and Geeraerts 2018.

<sup>3</sup>S. Wood, Goude, and Shaw 2015.

# Table of Contents

## ① Introduction

- Multilevel Models
- Issues with Large Datasets

## ② The split-sample approach

- Pseudo Likelihood
- Graphical representation
- Independent subsamples
- Dependent subsamples
- Overlapping subsamples

## ③ R Packages

- lme4 and mgcv
- Why use bam()?
- When to use bam()?

## ④ Summary

## ⑤ References

# The split-sample approach – Pseudo Likelihood

- Consider the log-likelihood function  $\ell(\boldsymbol{\theta}) = \sum_i \ell(\mathbf{y}_i | \boldsymbol{\theta})$  where  $\mathbf{y}_i$  is the vector of all observations in group  $i$
- Replaces the log-likelihood contribution  $\ell(\mathbf{y}_i | \boldsymbol{\theta})$  by a weighted sum of log-likelihood contributions for sub-vectors  $\mathbf{Y}_i^{(s)}$
- More specifically, the pseudo-log-likelihood function:

$$p\ell(\boldsymbol{\psi}) = \sum_i \sum_s \delta_s \ell(\mathbf{y}_i^{(s)} | \boldsymbol{\psi})$$

is maximized instead with respect to  $\boldsymbol{\psi}$ , which is not necessarily identical to  $\boldsymbol{\theta}$

- Although  $\hat{\boldsymbol{\psi}}$  is not the MLE estimate, it still has similar properties such as consistency and asymptotic normality<sup>4</sup>

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<sup>4</sup>Clark 2019.

# The split-sample approach – Graphical representation

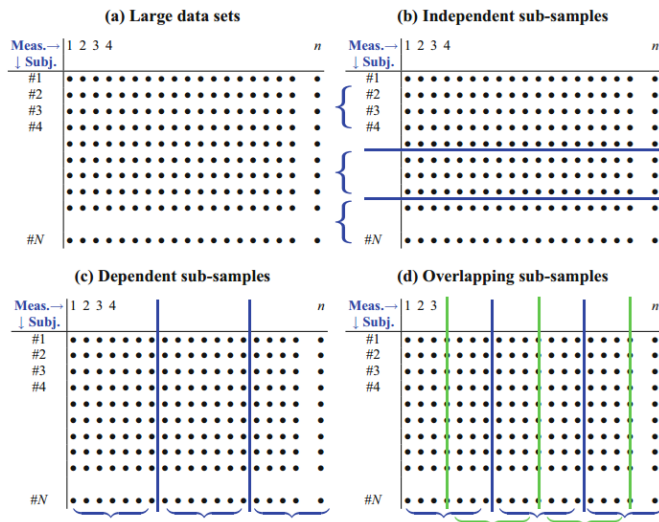


Figure 1: Graphical representation of different ways to split large samples

# The split-sample approach — Independent subsamples

- Shown in panel (b) of Figure 1, dataset with large  $N$  is partitioned into  $M$  independent sets  $S_m$  of groups, where  $m = 1, \dots, M$
- In each subsample, the model is fitted, yielding an estimate  $\hat{\theta}_m$  of  $\theta$ , equivalent to maximizing

$$p\ell(\psi) = \sum_m \sum_{i \in S_m} \ell(\mathbf{y}_i | \theta_m)$$

with respect to  $\psi = \{\theta_1, \dots, \theta_M\}$

- All  $\theta_m$  are equal to  $\theta$ , therefore the estimates  $\hat{\theta}_m$  can be averaged to obtain an overall estimate  $\hat{\theta}$



# The split-sample approach – Dependent subsamples

- Shown in panel (c) of Figure 1, dataset with large  $n$  is partitioned into  $M$  (not independent) sets  $S_m$  of groups, where  $m = 1, \dots, M$
- Fitting the model on each subsample, equivalent to maximizing

$$p\ell(\psi) = \sum_m \sum_i \ell(\mathbf{Y}_i^{(m)} | \theta_m)$$

with respect to  $\psi = \{\theta_1, \dots, \theta_M\}$ , where  $\mathbf{Y}_i^{(m)}$  is the observations in  $\mathbf{Y}_i$  belonging to subsample  $S_m$ .

- All  $\theta_m$  are not necessarily equal to  $\theta$ , therefore the combination of all  $\hat{\theta}_m$  into a single estimator  $\hat{\theta}$  depends on the precise model and data structure.

# The split-sample approach — Overlapping subsamples

- Shown in panel (d) of Figure 1, dataset with large  $n$  is partitioned similarly to dependent subsamples, but association between observations is accounted for by letting the subsamples overlap
- Denoting the parameters in pair  $\{\mathbf{Y}_i^{(p)}, \mathbf{Y}_i^{(q)}\}$  by  $\theta_{p,q}$ , fitting the models on all pairs is equivalent to maximizing

$$p\ell(\psi) = \sum_{p < q} \sum_i \ell(\mathbf{Y}_i^{(p)}, \mathbf{Y}_i^{(q)} | \theta_{p,q})$$

with respect to  $\psi = \{\theta_{1,2}, \theta_{1,3}, \dots, \theta_{Q-1,Q}\}$ , where  $\mathbf{Y}_i^{(p)}$  and  $\mathbf{Y}_i^{(q)}$  are the observations in  $\mathbf{Y}_i$  belonging to subsamples  $S_p$  and  $S_q$ , respectively.

- Similarly, the combination of all  $\hat{\theta}_{p,q}$  into a single estimator  $\hat{\theta}$  depends on the precise model and data structure.

# Table of Contents

## ① Introduction

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## ② The split-sample approach

- Pseudo Likelihood
- Graphical representation
- Independent subsamples
- Dependent subsamples
- Overlapping subsamples

## ③ R Packages

- lme4 and mgcv
- Why use bam()?
- When to use bam()?

## ④ Summary

## ⑤ References

## lme4

- an R package for fitting linear and generalized linear mixed-effects (multilevel) models<sup>5</sup>
- efficient, able to handle large sample sizes for simple model, and process hundreds of thousands observations on a typical laptop
- Modeling functions: `lmer()` and `glmer()`

## mgcv

- an R package for fitting generalized additive model and generalized additive mixed models<sup>6</sup>
- Modeling functions: `gam()` and `bam()`

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<sup>5</sup>Bates et al. 2015.

<sup>6</sup>S. N. Wood 2011.

# R Packages – Why use bam()?

- Same underlying model between `gam()` and `lme4`, with differences in parameter estimation
- How `bam()` works:
  - QR decomposition<sup>7</sup>
  - (i) Efficient fitting algorithm, (ii) Parallel computation, and (iii) Covariate discretization<sup>8</sup>
  - Efficient crossproduct matrix  $X^T W X$  computation<sup>9</sup>
- Discretization on large datasets leads to tradeoff between accuracy and speed

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<sup>7</sup>S. Wood, Goude, and Shaw 2015.

<sup>8</sup>S. Wood, Li, et al. 2017.

<sup>9</sup>Li and S. Wood 2020.

# R Packages – When to use `bam()`?

- In general, `lme4` is preferred due to easy syntax and robust estimation
- `bam()` is particularly useful for:
  - Complex models that exceed `lme4`'s capabilities
  - Incorporating smooth (nonlinear) terms
  - Large datasets with memory issues
  - Leveraging parallel computing resources

# Table of Contents

## ① Introduction

- Multilevel Models
- Issues with Large Datasets

## ② The split-sample approach

- Pseudo Likelihood
- Graphical representation
- Independent subsamples
- Dependent subsamples
- Overlapping subsamples

## ③ R Packages

- lme4 and mgcv
- Why use bam()?
- When to use bam()?

## ④ Summary

## ⑤ References

- Large multilevel datasets pose significant computational challenges
- The split-sample approach offers a practical solution
- R packages like `lme4` and `mgcv` provide robust tools for fitting multilevel models
- Approach and tools selection depends on dataset and research questions



# Table of Contents

## ① Introduction

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- Issues with Large Datasets

## ② The split-sample approach

- Pseudo Likelihood
- Graphical representation
- Independent subsamples
- Dependent subsamples
- Overlapping subsamples







## ③ R Packages

- lme4 and mgcv
- Why use bam()?
- When to use bam()?

## ④ Summary

## ⑤ References

# References I

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