

Bayesian high-dimensional variable selection in non-linear mixed-effects models using the SAEM algorithm

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Journée AppliBUGS 10 Juin 2022







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Framework: repeated measurement data

* Mixed-effects models: analyse observations collected repeatedly on several individuals.

(m) 200 90 150 100 50 400 800 1200 1600 Age (days)

Circumference of five orange trees

- Same overall behaviour but with individual variations.
- Non-linear growth.
- Are these variations due to known characteristics?
 - ► E.g.: growing conditions, genetic markers, ...





1) Description of intra-individual variability: For all $i \in \{1, ..., n\}$, $j \in \{1, ..., J\}$,

$$y_{ij} = g(\varphi_i, \psi, t_{ij}) + \varepsilon_{ij}, \ \varepsilon_{ij} \stackrel{\text{i.i.d.}}{\sim} \mathcal{N}(0, \sigma^2)$$

- $y_{ij} \in \mathbb{R}$: response of individual *i* at time t_{ij} (observation).
- $\varphi_i \in \mathbb{R}$: individual parameter, **not observed**.
- $\psi \in \mathbb{R}^q$: fixed effects, unknown.
- g: non-linear function with respect to φ_i (known).

2) Description of inter-individual variability:

$$\varphi_i = \mu + {}^{\mathrm{t}}\beta V_i + \xi_i, \ \xi_i \overset{\mathrm{i.i.d.}}{\sim} \mathcal{N}(0, \Gamma^2)$$

- $\mu \in \mathbb{R}$: intercept, unknown.
- $V_i \in \mathbb{R}^p$: covariates for individual *i* (known).
- $\beta = {}^{t}(\beta_{1}, \dots, \beta_{p}) \in \mathbb{R}^{p}$ covariate fixed effects vector, unknown.

Population parameters: $\theta = (\mu, \beta, \psi, \sigma^2, \Gamma^2)$



Marion Naveau Bayesian high-dimensional variable selection in NLMEM

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Variable selection

Aim: identify the most relevant covariates to characterise inter-individual variability.

✤ Active/Non-active covariates: covariates that are actually influential/non-influential for the characteristic under consideration.

Description of inter-individual variability:

$$\varphi_i = \mu + {}^{\mathrm{t}}_{\beta} V_i + \xi_i, \ \xi_i \stackrel{\mathrm{i.i.d.}}{\sim} \mathcal{N}(0, \Gamma^2)$$

- $\beta_{\ell} = 0 \iff$ covariate ℓ has no effect on parameter φ_i
- $\beta_{\ell} \neq 0 \iff$ covariate ℓ gives some information on parameter φ_i

• Model selection: variable selection \iff model selection among all the possible supports of β :

$$\mathcal{S}_{eta} = igg\{\ell \in \{1,\ldots,p\} igg| eta_\ell
eq 0igg\}.$$



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High-dimensional covariate selection in NLMEM

- Goal: identify the non-zero components of β .
- Specificity of the problem: p >> n
- Main difficulties:
 - High-dimensional variable selection:
 - \blacktriangleright parsimonious estimation of β
 - Non-explicit likelihood
 - The φ_i 's are not observed (latent variables model)
 - ▶ g is non-linear

$$p(y;\theta) = \int p(y|\varphi;\theta)p(\varphi;\theta)d\varphi = \prod_{i=1}^{n} \int p(y_{i}|\varphi_{i};\theta)p(\varphi_{i};\theta)d\varphi_{i}$$
$$= C_{\sigma^{2},\Gamma^{2}}\prod_{i=1}^{n} \int \exp\left(-\sum_{j=1}^{J} \frac{(y_{ij} - g(\varphi_{i},\psi,t_{ij}))^{2}}{2\sigma^{2}} - \frac{(\varphi_{i} - \mu - {}^{t}\beta V_{i})^{2}}{2\Gamma^{2}}\right)d\varphi_{i}$$



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State of	the art for <mark>l</mark>	nigh-dimensi	onal varia	ble selecti	on in

Frequentist framework:

mixed-effects models

- LMEM: both theoretical results and algorithmic developments for regularised methods (Schelldorfer et al., 2011; Fan and Li, 2012).
- NLMEM: algorithmic contribution (Ollier, 2021).
- Bayesian framework:
 - Linear regression (without random effects): $y_i = \alpha + {}^{t}\beta X_i + \epsilon_i$ theoretical and algorithmic developments using various sparsity-inducing priors (cf book Tadesse and Vannucci (2021)).
 - NLMEM: (Lee, 2022) advocated the Bayesian approach for this model but this is only a review, without implementation, does not focus on the high-dimension.

Proposed approach

Association of a Bayesian *spike-and-slab* prior for variable selection with a stochastic version of the EM algorithm, called MCMC-SAEM, for inference



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Spike-and-slab prior for the coefficients of β

♣ Introduction of latent variables δ_{ℓ} , $1 \leq \ell \leq p$:

 $\delta_\ell = \left\{ \begin{array}{ll} \mathbf{1} & \text{if covariate } \ell \text{ is to be included in the model}, \\ \mathbf{0} & \text{otherwise}. \end{array} \right.$

Spike-and-slab prior on β (George and McCulloch, 1997):

 $\pi(\beta|\delta) = \mathcal{N}_{\rho}(0, \mathsf{diag}((1 - \delta_{\ell})\nu_{0} + \delta_{\ell}\nu_{1})), \ 0 \leq \nu_{0} < \nu_{1} \ \mathsf{fixed},$

i.e. β_{ℓ} are independent and:

• $\beta_{\ell}|(\delta_{\ell}=0) \sim \mathcal{N}(0,\nu_0)$: "spike" distribution, ν_0 small

• $\beta_{\ell}|(\delta_{\ell}=1) \sim \mathcal{N}(0, \nu_1)$: "slab" distribution, ν_1 large





Figure: Spike-and-slab prior. Source: Deshpande et al. (2019) Marion Naveau Bayesian high-dimensional variable selection in NLMEM







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Proposed	method				

Idea: explore different levels of sparsity in β by varying the value of ν_0 in a grid Δ .

- 1. Creation of a model collection: for each $\nu_0 \in \Delta$,
 - Compute $\widehat{\Theta}$ by a MCMC-SAEM algorithm (Kuhn and Lavielle, 2004):

 $\widehat{\Theta}_{\nu_0}^{MAP} = \operatorname*{argmax}_{\Theta \in \Lambda} \pi(\Theta|y)$

► Estimate
$$\hat{\delta}$$
 (Ročková and George, 2014):
 $\hat{\delta} = \underset{\delta}{\operatorname{argmax}} P(\delta|\hat{\Theta}_{\nu_0}^{MAP}) \text{ such as } \hat{\delta}_{\ell} = 1 \iff \mathbb{P}(\delta_{\ell} = 1|\hat{\Theta}_{\nu_0}^{MAP}) \ge 0.5$
 $\iff \text{Define } \widehat{S}_{\nu_0} = \left\{ \ell \in \{1, \dots, p\} \mid |(\widehat{\beta}_{\nu_0}^{MAP})_{\ell}| \ge s_{\beta}(\nu_0, \nu_1, \widehat{\alpha}_{\nu_0}^{MAP}) \right\}$

2. Select the "best" model among $(\widehat{S}_{\nu_0})_{\nu_0 \in \Delta}$ by a fast criterion, eBIC (Chen and Chen, 2008):

$$\hat{\nu}_{0} = \operatorname*{argmin}_{\nu_{0} \in \Delta} \left\{ -2\log\left(p(y; \hat{\theta}_{\nu_{0}}^{MLE})\right) + B_{\nu_{0}} \times \log(n) + 2\log\left(\binom{p}{B_{\nu_{0}}}\right) \right\}$$

with $B_{
u_0}$: number of free parameters in the model $\widehat{S}_{
u_0}$.

3. Return $\widehat{S}_{\hat{\nu}_0}$.

A



Spike-and-slab regularisation plot



Figure: n = 200, J = 10, p = 500, $\Gamma^2 = 200$, $\sigma^2 = 30$, $\nu_1 = 12000$, $\mu = 1200$, $\beta = {}^{t}(100, 50, 20, 0, \dots, 0)$





- Let's go back to the first step of the proposed method:
 - \blacktriangleright Compute the MAP estimator of Θ
 - ▶ Goal: maximise $\pi(\Theta|y) = \int_{\mathcal{Z}} \pi(\Theta, Z|y) dZ$ with

$$\pi(\Theta, Z|y) = \frac{p(y|\Theta, Z)p(\Theta, Z)}{\int_{\mathcal{Z}} \int_{\Lambda} p(y|\Theta, Z)p(\Theta, Z)d\Theta dZ}$$

Non-explicit integral





Reference: Dempster et al. (1977)

- 1. Initialisation: choose $\Theta^{(0)}$.
- 2. Iteration $k \ge 0$:
 - E-step (Expectation): compute

$$Q(\Theta|\Theta^{(k)}) = \mathbb{E}_{Z|(y,\Theta^{(k)})} \left[\log(\pi(\Theta, Z|y)) \middle| y, \Theta^{(k)} \right]$$

• M-step (Maximisation): compute

$$\Theta^{(k+1)} = \underset{\Theta \in \Lambda}{\operatorname{argmax}} \ Q(\Theta | \Theta^{(k)}).$$

3. $\hat{\Theta} = \Theta^{(K)}$, for K large enough.



SAEM and MCMC-SAEM algorithms

Methodology

References: Delyon et al. (1999), Kuhn and Lavielle (2004)

- 1. Initialisation: choose $\Theta^{(0)}$ and $Q_0(\Theta) = 0$,
- 2. Iteration $k \ge 0$:

Δ0,

- S-step (Simulation): simulate Z^(k) using the result of one iteration of an MCMC procedure with π(Z|y, Θ^(k)) for target distribution,
- SA-step (Stochastic Approximation): compute an approximation of Q(Θ|Θ^(k)) according to:

 $Q_{k+1}(\Theta) = Q_k(\Theta) + \frac{\gamma_k}{\log \pi(\Theta, Z^{(k)}|y)} - Q_k(\Theta)),$

• M-step (Maximisation): compute

$$\Theta^{(k+1)} = \underset{\Theta \in \Lambda}{\operatorname{argmax}} Q_{k+1}(\Theta),$$

3. $\hat{\Theta} = \Theta^{(\kappa)}$, for K large enough,

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where $(\gamma_k)_k$ a step sizes sequence decreasing towards 0 such that $\forall k$, $\gamma_k \in [0, 1]$, $\sum_k \gamma_k = \infty$ and $\sum_k \gamma_k^2 < \infty$.

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Specifics in Spike-and-Slab-NLMEM

\bullet Decomposition of Q:

$$Q(\Theta|\Theta^{(k)}) = \mathbb{E}_{(\varphi,\delta)|(y,\Theta^{(k)})}[\log(\pi(\Theta,\varphi,\delta|y))|y,\Theta^{(k)}]$$

= $C + \underbrace{\mathbb{E}_{\varphi|y,\Theta^{(k)}}\left[\widetilde{Q}_{1}(y,\varphi,\theta,\Theta^{(k)})\Big|y,\Theta^{(k)}\right]}_{\text{non-explicit}} + \underbrace{\widetilde{Q}_{2}(\alpha,\Theta^{(k)})}_{\text{explicit}}$

M-step:

- \blacktriangleright θ and α estimated separately.
- ▶ $\hat{\alpha}$ updated as in an EM algorithm with $\tilde{Q}_2(\alpha, \Theta^{(k)})$.
- $\blacktriangleright \ \widehat{\theta}$ updated via stochastic approximation of:

$$\mathbb{E}_{\varphi|y,\Theta^{(k)}}\left[\left. \stackrel{\sim}{Q}_{1}(y,arphi, heta,\Theta^{(k)}) \right| y,\Theta^{(k)}
ight].$$



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MCMC-SAEM algorithm in SSNLMEM

- 1. Initialisation: choose $\Theta^{(0)}$ and $Q_{1,0}(\theta) = 0$,
- 2. Iteration $k \ge 0$:
 - S-step (Simulation): simulate φ^(k) using the result of one iteration of an MCMC procedure with π(φ|y, Θ^(k)) for target distribution,
 - SA-step (Stochastic Approximation): compute

$$Q_{1,k+1}(\theta) = Q_{1,k}(\theta) + \gamma_k (\tilde{Q}_1(y,\varphi^{(k)},\theta,\Theta^{(k)}) - Q_{1,k}(\theta)),$$

and $\widetilde{Q}_2(\alpha, \Theta^{(k)})$, • M-step (Maximisation):

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$$\theta^{(k+1)} = \operatorname*{argmax}_{\theta \in \Lambda_{ heta}} Q_{1,k+1}(\theta) \text{ and } \alpha^{(k+1)} = \operatorname*{argmax}_{\alpha \in [0,1]} \tilde{Q}_{2}(\alpha, \Theta^{(k)}),$$

3. $\hat{\Theta} = \Theta^{(K)}$, for K large enough, where $(\gamma_k)_k$ a step sizes sequence decreasing towards 0 such that $\forall k$, $\gamma_k \in [0, 1]$, $\sum_k \gamma_k = \infty$ and $\sum_k \gamma_k^2 < \infty$.

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Convergence graphs





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Logistic §	growth mod	el			



Figure: Simulated data

• Size of plant $i \in \{1, ..., n\}$ at time t_{ij} , $j \in \{1, ..., 10\}$: $y_{ij} = g(\varphi_i, \psi, t_{ij}) + \varepsilon_{ij}, \ \varepsilon_{ij} \stackrel{\text{iid}}{\sim} \mathcal{N}(0, \sigma^2)$ where:

$$g(arphi_i,\psi,t_{ij}) = rac{\psi_1}{1+\exp\left(-rac{t_{ij}-arphi_i}{\psi_2}
ight)}$$

 $\psi = (\psi_1, \psi_2)$ fixed effects.

• φ_i : characteristic time $\varphi_i = \mu + {}^{t}\beta V_i + \xi_i, \ \xi_i \stackrel{\text{iid}}{\sim} \mathcal{N}(0, \Gamma^2)$

$$\boldsymbol{\theta} = (\boldsymbol{\mu}, \boldsymbol{\beta}, \boldsymbol{\psi}, \sigma^2, \boldsymbol{\Gamma}^2)$$



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Simulatio	n design				

Parameters:

- $n \in \{100, 200\}$ individuals,
- $p \in \{500, 2000, 5000\}$ simulated covariates according to $V_i \sim \mathcal{N}(0, \Sigma)$:
 - ► Scenario i.i.d.: $\Sigma = Id$ ► Correlated scenarios: $\Sigma \neq Id$
- $\beta = {}^{t}(100, 50, 20, 0, \dots, 0)$ covariate fixed effects vector,
- $\Gamma^2 \in \{200, 1000, 2000\}$ inter-individual variance,
- $\mu = 1200$, $\sigma^2 = 30$, $\psi = (\psi_1, \psi_2) = (200, 300)$.

* Spike-and-slab hyperparameters:

• $\nu_1 = 12000$ slab variance,

•
$$\log_{10}(\Delta) = \left\{ -2 + k imes rac{4}{19}, k \in \{0, \dots, 19\}
ight\}$$
 grid of u_0 values.

► For each combination of (n, p, Γ^2) , the method is applied on 100 different simulated datasets.





Results for independent covariates



Figure: Empirical probability of correct model selection.



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Summary	of the resu	llts			

• Uncorrelated covariates $V_i \sim \mathcal{N}(0, I_p)$:

- Results improve as *n* increases.
- Degradation of results when p or Γ^2 increases.
- When the procedure fails, it is most often because it under-selects:
 - ▶ "Cautious" approach, few false positives!
- Correlated covariates $V_i \sim \mathcal{N}(0, \Sigma)$:
 - Fairly similar good performance.
 - More false positives and/or false negatives in some correlation scenarios:

 \blacktriangleright + false positives: correlations between active and non-active covariates.

▶ + false negatives: correlated active covariates.







- Both methods have an execution time that grows **polynomially** with *p*.
- The proposed inference method can browse grid of about 20 values of ν_0 while adaptive MCMC explores a single value.



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Presentat	ion of the	dataset		

Wheat leaf senescence data.

✤ Panel: n = 216 soft wheat varieties subjected to nitrogen stress, observed J = 18 times.

Varieties respond differently to stress: for example, some of them tolerate stress better and senescence is delayed.

* Aim: select molecular markers, from among p = 34838 markers, which could be associated with this tolerance.



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Data repr	esentation:	percentage c	of desiccate	ed leaves	







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SAINT_EX









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Modelling	<u>y</u>		

$$\begin{cases} y_{ij} = g(\phi_i, t_{ij}) + \varepsilon_{ij} &, \varepsilon_{ij} \stackrel{\text{i.i.d.}}{\sim} \mathcal{N}(0, \sigma^2), \text{with } \phi_i = (\varphi_i, \psi_i) \in \mathbb{R}^2\\ \varphi_i = \mu + {}^{t} \lambda v_i + {}^{t} \beta V_i + \xi_i &, \xi_i \stackrel{\text{i.i.d.}}{\sim} \mathcal{N}(0, \Gamma^2)\\ \psi_i = \eta + \omega_i &, \omega_i \stackrel{\text{i.i.d.}}{\sim} \mathcal{N}(0, \Omega^2) \end{cases}$$

where:

•
$$g(\phi_i, t_{ij}) = rac{100}{1 + \exp\left(-rac{t_{ij} - \varphi_i}{\psi_i}
ight)},$$

- v_i: covariates not subject to selection, allows the inclusion of sub-populations in the model,
- V_i: molecular markers, subject to selection, which contains QTLs identified by biologists and markers associated with heading date which is highly correlated with φ_i.

$$\theta = (\mu, \lambda, \beta, \eta, \sigma^2, \Gamma^2, \Omega^2)$$

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Data pro	cessing		

* p >> n: ultra-high dimensional problem.

• Molecular markers \implies strong correlations/collinearity between covariates.

Covariates have few modalities:

> table(nb_mod_cov)
nb_mod_cov
 1 2 3

1 2 3 4 45 9237 19712 5844

- ✤ With "too many" 0's or "too many" 1's for some covariates, we remove:
 - markers filled in the same way for all individuals,
 - markers entered as the exact opposite of another marker (marker1=1-marker2).
 - markers whose minimum and maximum modalities are not represented at least 10 times.
 - markers that have a correlation > 0.7.

p = 6164



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Results



- Selected support size: 20
- Number of covariates selected at least once along the grid: 90
- "Peak" structure could be explained by correlations between the covariates.



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Summary:

- Development of an original method that combines SAEM and Bayesian variable selection.
- Very encouraging numerical results on simulated data.
- Faster method than a full MCMC implementation.

 \Rightarrow **Preprint:** Naveau and al. (2022). Bayesian high-dimensional covariate selection in non-linear mixed-effects models using the SAEM algorithm. <u>arXiv:2206.01012</u>.

Perspectives:

- Provide theoretical guarantees: selection consistency.
- Apply our method to a real dataset (in progress).
- Consider a multidimensional individual parameter.



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Thanks for your attention!



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• Decomposition of Q:

$$Q(\Theta|\Theta^{(k)}) = \mathbb{E}_{(\varphi,\delta)|(y,\Theta^{(k)})}[\log(\pi(\Theta,\varphi,\delta|y))|y,\Theta^{(k)}]$$

= $\mathbb{E}_{\varphi|(y,\Theta^{(k)})}\left[\mathbb{E}_{\delta|(\varphi,y,\Theta^{(k)})}\left[\log(\pi(\Theta,\varphi,\delta|y))|\varphi,y,\Theta^{(k)}\right]|y,\Theta^{(k)}\right]$
= $\mathbb{E}_{\varphi|(y,\Theta^{(k)})}\left[\widetilde{Q}(y,\varphi,\Theta,\Theta^{(k)})|y,\Theta^{(k)}\right]$
= $C + \underbrace{\mathbb{E}_{\varphi|y,\Theta^{(k)}}\left[\widetilde{Q}_{1}(y,\varphi,\theta,\Theta^{(k)})|y,\Theta^{(k)}\right]}_{\text{non-explicit}} + \underbrace{\widetilde{Q}_{2}(\alpha,\Theta^{(k)})}_{\text{explicit}}$



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Results for uncorrelated covariates

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Correlated	l covariates	$V_i \sim \mathcal{N}(0,$	Σ)		

Scenario	Σ
iid	I_p
1	$\left(\begin{array}{c c c} I_3 & 0_{3,p-3} \\ \hline 0_{p-3,3} & (\rho_{\Sigma}^{ i-j })_{i,j \in \{4,,p\}} \end{array} \right)$
2	$\left(\begin{array}{c c} I_3 & A \\ \hline & I_{A} & I_{p-3} \end{array}\right), \text{ with } A = \left(\begin{array}{cccc} 0 & 0 & \dots & 0 \\ 0 & 0 & \dots & 0 \\ & & (\rho_{\Sigma}^{ 3-j })_{j \in \{4,\dots,p\}} \end{array}\right)$
3	$\left(\begin{array}{c c} (\rho_{\Sigma}^{ i-j })_{i,j\in\{1,,3\}} & 0_{3,p-3} \\ \hline 0_{p-3,3} & I_{p-3} \end{array}\right)$
4	$(ho_{\Sigma}^{ i-j })_{i,j\in\{1,,p\}}$

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Results for $\rho_{\Sigma} = 0.3$



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