#### Aldo Gardini<sup>1</sup>, Carlo Trivisano<sup>1</sup> and Enrico Fabrizi<sup>2</sup> aldo.gardini2@unibo.it

<sup>1</sup>Dipartimento di Scienze Statistiche 'P. Fortunati', Università di Bologna <sup>2</sup>DISES, Università Cattolica del S. Cuore

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### Outline of the presentation

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2 Methodological aspects in SAE context

- The random intercept model
- Prior specification

3 Simulation studies

- Model based
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Research motivation

### Log-normality and Bayesian inference

Let us consider the simple **unconditional mean** estimation of an i.i.d. sample  $X_1, ..., X_n$  under log-normality assumption:

$$X_i \sim \log \mathcal{N}\left(\xi, \sigma^2
ight), \; i = 1, ..., n.$$

The functional to estimate is  $\theta = \exp{\{\xi + 0.5\sigma^2\}}$ . Is it possible to compute its posterior mean?

- With usual non-informative priors on σ<sup>2</sup>, p(θ|X) belongs to the log-t family. Its moments coincide with evaluations of the t distribution moment generating function: they are not finite.
- To overcome this issue, Fabrizi and Trivisano (2012) proposed the generalized inverse Gaussian (GIG) prior for σ<sup>2</sup>:

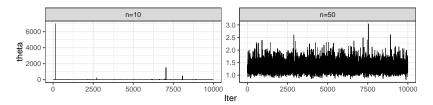
$$f_X(x) = \left(\frac{\gamma}{\delta}\right)^{\lambda} \frac{1}{2K_{\lambda}(\delta\gamma)} x^{\lambda-1} \exp\left\{-\frac{1}{2}\left(\frac{\delta^2}{x} + \gamma^2 x\right)\right\}, \quad x > 0.$$

Research motivation

### What does moment infiniteness implies?

**Checking the moments existence is often ignored in practice:** usually MCMC methods are applied without checking for the existence conditions. **Do they send warnings about this issue?** 

- *Small datasets (n* < 15): running reasonably long chains, anomalies can be observed.
- Larger datasets: the chains seem to converge to the posterior distribution without issues. But, which is the meaning of a numerical estimate of an integral that is analytically infinite?



Research motivation

### Bayesian log-normal linear mixed model

Returning to the log-normal context, the same problems are registered in estimating data scale quantities from the general log-normal mixed model.

This model arises when a normal mixed model is assumed on the logarithm of the response variable  $\mathbf{w} = \log \mathbf{y}$ . This procedure is largely used in practice.

The model could be expressed in the following Bayesian hierarchical formulation:

$$\begin{split} \mathbf{w} | \mathbf{u}, \boldsymbol{\beta}, \sigma^2 &\sim \mathcal{N}_n \left( \mathbf{X} \boldsymbol{\beta} + \mathbf{Z} \mathbf{u}, \mathbf{I}_n \sigma^2 \right); \\ \mathbf{u} | \tau_1^2, ..., \tau_q^2 &\sim \mathcal{N}_m \left( \mathbf{0}, \mathbf{D} \right), \ \mathbf{D} = \oplus_{s=1}^q \mathbf{I}_{m_s} \tau_s^2; \quad (\mu, \sigma^2) \sim p(\mu, \sigma^2); \\ \boldsymbol{\tau}^2 &\sim p(\tau_1^2, ..., \tau_q^2). \end{split}$$

#### The moments of the following posterior distributions require care:

- Conditioned expectations with respect to covariates and/or random effects,
- Posterior predictive distribution  $p(\tilde{\mathbf{y}}|\mathbf{y})$ .

L The random intercept model

### Small area estimation: notation

We considered the classical small area estimation framework:

- **Population** U of N units. It is partitioned into D sub-populations  $U_1, ..., U_D$  having dimensions  $N_1, ..., N_D$ ,
- A random sample s with size n is drawn from U. D sub-samples  $s_1, ..., s_D$  with size  $n_1, ..., n_D$  are obtained.
- The unsampled unites  $r_1, ..., r_D$  are  $N_d n_d$  in each area.
- The subscript s denotes quantities related to the sampled units, whereas r refers to the unsampled units characteristics.
- **y**<sub>s</sub>  $\in \mathbb{R}^n$  denotes the observed values for the response variable, whereas **w**<sub>s</sub> is the log-transformed vector.
- The values of p covariates are stored in  $X_s$ .

Methodological aspects in SAE context

L The random intercept model

### Random intercept model definition

The classical BHF unit-level model is specified for  $\mathbf{w}_s$  in the Bayesian framework:

$$\begin{split} \mathbf{w}_{s} | \mathbf{u}, \boldsymbol{\beta}, \sigma^{2} &\sim \mathcal{N}_{n} \left( \mathbf{X}_{s} \boldsymbol{\beta} + \mathbf{Z}_{s} \mathbf{u}, \sigma^{2} \mathbf{I}_{n} \right); \\ \mathbf{u} | \tau^{2} &\sim \mathcal{N}_{D} \left( 0, \tau^{2} \mathbf{I}_{D} \right), \ (\boldsymbol{\beta}, \sigma^{2}) &\sim p(\boldsymbol{\beta}, \sigma^{2}); \\ \tau^{2} &\sim p(\tau^{2}); \end{split}$$

where:

- **Z**<sub>s</sub>  $\in \mathbb{R}^{n \times D}$  is the random effects design matrix that defines the random intercept model,
- $\beta \in \mathbb{R}^p$  contains the fixed effects coefficients and  $u \in \mathbb{R}^D$  the area-specific random effects.

The target inferential quantities are the area means:

$$\hat{ar{y}}_d(oldsymbol{ heta}) = N_d^{-1} \left[ \sum_{i \in s_d} y_{di} + \sum_{i \in r_d} \hat{y}_{di} 
ight].$$

The random intercept model

### Target quantity: area mean

In the hierarchical Bayes context, it is natural to estimate the out-of-sample elements exploiting the posterior predictive distribution:

$$\hat{\tilde{Y}}_{d}^{HB} = \frac{1}{N_{d}} \left( \sum_{i \in s_{d}} y_{di} + \sum_{i \in r_{d}} \mathbb{E} \left[ y_{di} | \mathbf{y}_{s} \right] \right).$$

Theorem (Area mean posterior moments existence)

The posterior moments of  $\hat{\tilde{Y}}_{d}^{HB}$  are defined up to order r if the prior of  $\sigma^2$  includes an exponential term  $\exp\{-k\sigma^2\}$  and the hyperparameters are fixed in order to satisfy the inequality:

$$k > r^2 + r^2 h_d^{max}, \quad h_d^{max} = \max_{i \in r_d} \mathbf{x}_{di}^T (\mathbf{X}_s^T \mathbf{X}_s) \mathbf{x}_{di}.$$

L The random intercept model

### Target quantity: poverty measures

The hierarchical Bayes approach is appealing in case of non-linear data transformation target functionals. An example is represented by the family FGT (Foster et al. 1984) of poverty measures, that is defined for subject i of area d as:

$$F_{di,\alpha} = \left(\frac{c - y_{di}}{c}\right)^{\alpha} \mathbf{1}_{\{y_{di} < c\}}(Y_{di});$$

where c is the poverty line and  $\alpha = 0, 1, 2$ .

The area mean poverty measure HB estimate is:

$$\hat{F}_{d,\alpha}^{HB} = \frac{1}{N_d} \left[ \sum_{i \in s_d} F_{di,\alpha} + \sum_{i \in \bar{s}_d} \mathbb{E} \left[ F_{di,\alpha} | \mathbf{y}_s \right] \right]$$

**Existence condition:** same of previous theorem but with  $r' = r\alpha$ . The prior used in Molina et al. (2014) does not preserve the posterior moments existence.

Prior specification

## Prior specification: general ideas

- **Location parameter:** improper flat prior  $p(\beta) \propto 1$ .
- Variance components: how to specify a weakly informative prior preserving the moments existence conditions?

#### Variance components

**Distribution choice:** the flexibility of the three-parameters GIG distribution has been privileged. The existence condition affects the tail parameter  $\gamma$  that is fixed equal to  $\gamma_0^2 = (r+1)^2 + (r+1)^2 h^{max}$ .

General strategy:

- Preserving the posterior moments existence;
- Saving the balance among the variance components evaluating the marginal prior on the intraclass correlation coefficient  $\rho$ . Following the idea of *uniform shrinkage prior* (Chaloner, 1987), a uniform prior on  $\rho$  is the target to reach.

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Methodological aspects in SAE context

Prior specification

### Prior specification: some details

Briefly, the main steps covered to formulate our proposal are:

**1** Two independent GIG priors having equal hyperparameters are fixed for the variance components:

$$p(\sigma^2) \sim GIG(\lambda, \delta, \gamma_0); \quad p(\tau_s^2) \sim GIG(\lambda, \delta, \gamma_0).$$

**2** The marginal prior on  $\rho = \tau^2 (\sigma^2 + \tau^2)^{-1}$  has the following form:

$$p(\rho) = \frac{K_{2\lambda} \left( \gamma_0^2 \delta^2 \left[ \frac{1}{\rho} + \frac{1}{1-\rho} \right] \right)}{2 \left[ K_\lambda \left( \gamma_0 \delta \right) \right]} \left[ \rho(1-\rho) \right]^{-1}, \quad \rho \in (0,1).$$

- 3 To remove the tail parameter effect, the limiting case  $\delta \rightarrow 0$  might be considered. If  $\lambda > 0$ , then the gamma distribution is obtained.
- **4** Fixing  $\lambda = 1$  a uniform prior in the interval [0,1] is assumed for  $\rho$ .

$$p(\sigma^2) \sim \mathcal{G}(1, \gamma_0/2); \quad p(\tau_s^2) \sim \mathcal{G}(1, \gamma_0/2).$$

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Simulation studies

Model - based

### Model based simulation setting

Evaluation of the frequentist properties of the area estimates: compared EB and HB (both under GIG priors and uniform shrinkage priors) methods. Design of the simulation study inspired by Berg et al. (2016). At each iteration (B = 5000), a population is randomly generated from the model:

$$w_{di} = \beta_0 + \beta_1 x_{di} + u_d + e_{di};$$
  
 $u_d \sim \mathcal{N}(0, \tau^2), \ e_{di} \sim \mathcal{N}(0, \sigma^2); \ d = 1, ..., 10; \ i = 1, ..., N_D,$ 

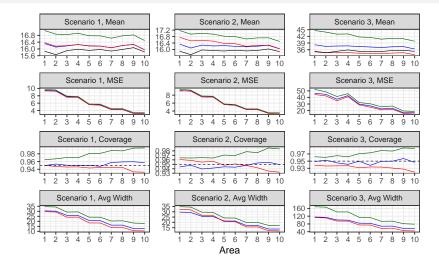
and a stratified random sample is drawn. Two equivalent area for each dimension are considered, with  $N_d = (41, 81, 161, 323, 645)$  and  $n_d = (3, 5, 10, 20, 40)$ .

Three scenarios for the variance components  $(\sigma^2, \tau^2)$ : (0.6, 0.3), (0.78, 0.12) and (2, 0.5).

#### -Simulation studies

Model - based

### Model based simulation results



Method - EB - HB GIG - HB naive US - True

Simulation studies

Design - based

### Design based simulation: AAGIS data

It is useful to investigate the design properties of the considered solutions.

The synthetic finite population of size N = 81982 based on the Australian Agricultural Grazing Industries Survey (AAGIS) has been already used to investigate log-normal SAE methods.

At each iteration (B = 1000), a stratified simple random sample (n = 1686) is drawn from the 29 areas. The variable of interest is the *annual firm cost* and the available auxiliary information is the logarithm of *firm size*. The same BHF model as before was fitted.

#### Simulation results:

	EB	HB <sub>GIG</sub>
Average RRMSE	0.145	0.145
Average bias	0.011	0.011

Conclusion and future work

### Conclusion and future work

- Bayesian analysis of mixed models on the log scale requires care:
  - Existence of posterior moments should not be taken for granted.
  - We proposed a prior distribution that allows the moments existence for usual target quantity of the SAE framework.

#### Future work:

- Implement scalable MCMC algorithms in order to be able to apply Bayesian methods to large input datasets.
- Work on approximations of the posterior predictive distribution to allow the analysis of whole population as out-of-samples.

Conclusion and future work

### References

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