Confidence interval estimation of small area parameters shrinking both means and variances

by Sarat C. Dass, Tapabrata Maiti, Hao Ren and Samiran Sinha

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- r revised
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Confidence interval estimation of small area parameters shrinking both means and variances

Sarat C. Dass, Tapabrata Maiti, Hao Ren and Samiran Sinha

Abstract

We propose a new approach to small area estimation based on joint modelling of means and variances. The proposed model and methodology not only improve small area estimators but also yield “smoothed” estimators of the true sampling variances. Maximum likelihood estimation of model parameters is carried out using EM algorithm due to the non-standard form of the likelihood function. Confidence intervals of small area parameters are derived using a more general decision theory approach, unlike the traditional way based on minimizing the squared error loss. Numerical properties of the proposed method are investigated via simulation studies and compared with other competitive methods in the literature. Theoretical justification for the effective performance of the resulting estimators and confidence intervals is also provided.

Key Words: EM algorithm; Empirical Bayes; Hierarchical models; Rejection sampling; Sampling variance; Small area estimation.

1. Introduction

Small area estimation and related statistical techniques have become a topic of growing importance in recent years. The need for reliable small area estimates is felt by many agencies, both public and private, for making useful policy decisions. An example where small area techniques are used in practice is in the monitoring of socio-economic and health conditions of different age-sex-race groups where the patterns are observed over small geographical areas.

It is now widely recognized that direct survey estimates for small areas are usually unreliable due to their typically large standard errors and coefficients of variation. Hence, it becomes necessary to obtain improved estimates with higher precision. Model-based approaches, either explicit or implicit, are elicited to connect the small areas and improved precision is achieved by “borrowing strength” from similar areas. The estimation technique is also known as shrinkage estimation since the direct survey estimates are shrunk towards the overall mean. The survey based direct estimates and sample variances are the main ingredients for building aggregate level small area models. The typical modeling strategy assumes that the sampling variances are known while a suitable linear regression model is assumed for the means. For details of these developments, we refer to reader to Ghosh and Rao (1994), Pfeffermann (2002) and Rao (2003). The typical area level models are subject to two main criticisms. First, in practice, the sampling variances are estimated quantities, and hence, are subject to substantial errors. This is because they are often based on equivalent sample sizes from which the direct estimates are calculated. Second, the assumption of known and fixed sampling variances of typical small area models does not take into account the uncertainty in the variance estimation into the overall inference strategy.

Previous attempts have been made to model only the sampling variances; see, for example, Maples, Bell and Huang (2009), Gershunskaya and Lahiri (2005), Huff, Eltinge and Gershunskaya (2002), Cho, Eltinge, Gershunskaya and Huff (2002), Valliant (1987) and Otto and Bell (1995). The articles Wang and Fuller (2003) and Rivest and Vandal (2003) extended the asymptotic mean square error (MSE) estimation of small area estimators when the sampling variances are estimated as opposed to the standard assumption of known variances. Additionally, You and Chapman (2006) considered the modelling of the sampling variances with inference using full Bayesian estimation techniques.

The necessity of variance modelling has been felt by many practitioners. The latest developments in this area are nicely summarized in a recent article by William Bell of the United States Census Bureau 2008. He carefully examined the consequences of these issues in the context of MSE estimation of model based small area estimators. He also provided numerical evidence of MSE estimation for Fay-Herriot models (given in Equation 1) when sampling variances are assumed to be known. The developments in the small area literature so far can be “loosely” viewed as (i) smoothing the direct sampling error variances to obtain more stable variance estimates with low bias and (ii) (partial) accounting of the uncertainty in sampling variances by extending the Fay-Herriot model.

As evident, lesser or no attention has been given to account for the sampling variances effectively while modeling the mean compared to the volume of research that has been done for modeling and inferring the means. There is a lack of systematic development in the small area literature that

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includes “shrinking” both means and variances. In other words, we like to exploit the technique of “borrowing strength” from other small areas to “improve” variance estimates as we do to “improve” the small area mean estimates. We propose a hierarchical model which uses both the direct survey and sampling variance estimates to infer all model parameters that determine the stochastic system. Our methodological goal is to develop the dual “shrinkage” estimation for both the small area means and variances, exploiting the structure of the mean-variance joint modelling so that the final estimators are more precise. Numerical evidence shows the effectiveness of dual shrinkage on small area estimates of the mean in terms of the MSE criteria.

Another major contribution of this article is to obtain confidence intervals of small area means. The small area literature is dominated by point estimates and their associated standard errors; it is well known that the standard practice of [point estimate ± q × standard error], where q is the Z (standard normal) or t cut-off point, does not produce accurate coverage probabilities of the intervals; see Hall and Maiti (2006) and Chatterjee, Lahiri and Li (2008) for more details. Previous work is based on the bootstrap procedure and has limited use due to the repeated estimation of model parameters. We produce confidence intervals for the means from a decision theory perspective. The construction of confidence intervals is easy to implement in practice.

The rest of the article is organized as follows. The proposed hierarchical model for the sample means and variances is developed in Section 2. The estimation of model parameters via the EM algorithm is developed in Section 3. Theoretical justification for the proposed confidence interval and coverage properties are presented in Section 4. Sections 5 and 6 present a simulation study and a real data example, respectively. Some discussion and concluding remarks are presented in Section 7. An alternative model formulation for small area as well as mathematical details are provided in the Appendix.

2. Proposed model

Suppose n small areas are in consideration. For the i th small area, let \( (X_i, S_i^2) \) be the pair of direct survey estimate and sampling variance, for \( i = 1, 2, \ldots, n \). Let \( Z_i = (Z_{1i}, \ldots, Z_{pi})^T \) be the vector of p covariates available at the estimation stage for the i th small area. We propose the following hierarchical model:

\[
\begin{align*}
X_i \mid \theta_i, \sigma_i^2 & \sim \text{Normal}(\theta_i, \sigma_i^2) \\
\theta_i & \sim \text{Normal}(Z_i^T \beta, \tau^2)
\end{align*}
\]

(1)

independently for \( i = 1, 2, \ldots, n \). In the model elicitation, \( n_i \) is the sample size for a simple random sample (SRS) from the i th area, \( \beta = (\beta_1, \ldots, \beta_p)^T \) is the \( p \times 1 \) vector of regression coefficients, and \( B = (a, b, \beta, \tau^2)^T \) is the collection of all unknown parameters in the model. Also, \( \Gamma(a, b) \) is the Gamma density function with positive shape and scale parameters \( a \) and \( b \), respectively, defined as \( f(x) = (b^a x^{a-1} e^{-bx})/\Gamma(a) \) for \( x > 0 \), and 0 otherwise. The unknown \( \sigma_i^2 \) is the true variance of \( X_i \) and is usually estimated by the sample variance \( S_i^2 \). Although \( S_i^2 \)'s are assumed to follow a chi-square distribution with \( (n_i - 1) \) degrees of freedom (as a result of normality and SRS), we note that for complex survey designs, the degree of freedom needs to be determined carefully [e.g., Maples et al. 2009]. More importantly, the role of the sample sizes in shrinkage estimation of \( \sigma_i^2 \) is as follows: For low values of \( n_i \), the estimate of \( \sigma_i^2 \) is shrunk more towards the overall mean (ab) compared to higher \( n_i \) values. Thus, for variances, sample sizes play the same role as precision in shrinkage estimation of the small area mean estimates. We note that You and Chapman (2006) also considered the second level of the sampling variance modelling. However, the hyperparameters related to prior of \( \sigma_i^2 \) are not data driven, they are rather chosen in such a way that the prior will be vague. Thus, their model can be viewed as the Bayesian version of the models considered in Rivest and Vandal (2003) and Wang and Fuller (2003). The second level modelling of \( \sigma_i^2 \) in (2) can be further extended to \( \sigma_i^2 \sim \text{Gamma}(a, b) \) so that \( E(\sigma_i^2) = \exp(Z_i^T \beta) \) for another set of p regression coefficients \( \beta \), to accommodate covariate information in the variance modeling.

Although our model is motivated by Hwang, Qiu and Zhao (2009), we like to mention that Hwang et al. (2009) considered shrinking means and variances in the context of microarray data where they prescribed an important solution by plugging in a shrinkage estimator of variance into the mean estimator. The shrinkage estimator of the variance in Hwang et al. (2009) is a function of \( S_i^2 \) only, and not of both \( X_i \) and \( S_i^2 \); see Remarks 2 and 3 in Section 2. Thus, inference of the mean does not take into account the full uncertainty in the variance estimation. Further, their model does not include any covariate information. The simulation study described subsequently indicate that our method of estimation performed better than Hwang et al. (2009).

In the above model formulation, inference for the small area mean parameter \( \theta_i \) can be made based on the conditional distribution of \( \theta_i \) given all of the data \{\( (X_i, S_i^2, Z_i) \), \( i = 1, \ldots, n \}\}. Under our model set up, the conditional

\[
\begin{align*}
\frac{(n_i - 1)S_i^2}{\sigma_i^2} & \sim \text{Chi}^2_{n_i - 1} \\
\sigma_i^{-1} & \sim \text{Gamma}(a, b)
\end{align*}
\]

(2)
distribution of \( \theta_i \) is a non-standard distribution and does not have a closed form, thus requiring numerical methods, such as Monte Carlo and the EM algorithm, for inference, and the details are provided in the next section.

3. Inference methodology

3.1 Estimation of unknown parameters via EM algorithm

In practice, \( \mathbf{B} = (a, b, \mathbf{\beta}, \tau^2)^T \) is unknown and has to be estimated from the data \( \{(X_i, S_i^2, Z_i), i = 1, 2, ..., n\} \). Our proposal is to estimate \( \mathbf{B} \) by the marginal maximum likelihood method: Estimate \( \mathbf{B} \) by \( \hat{\mathbf{B}} \) where \( \hat{\mathbf{B}} \) maximizes the marginal likelihood \( L_M(\mathbf{B}) = \prod_{i=1}^{n} L_{M,i}(\mathbf{B}) \), where

\[
L_{M,i} = \frac{\Gamma(n_i/2+a)}{\Gamma(a)b^a} \int \exp\left[-\frac{(\theta_i - Z_i^T \mathbf{\beta})^2}{2\tau^2}\right] \psi_i^{(n_i/2+a)} d\theta_i,
\]

and

\[
\psi_i = \left\{0.5(X_i - \theta_i)^2 + 0.5(n_i - 1)S_i^2 + \frac{1}{b}\right\}.
\]

The marginal likelihood \( L_M \) involves integrals that cannot be evaluated in closed-form, and hence, one has to resort to numerical methods for its maximization. One such algorithm is the EM (Expectation-Maximization) iterative procedure which is used when such integrals are present. The EM algorithm involves augmenting the observed likelihood \( L_M(\mathbf{B}) \) with missing data; in our case, the variables of the form \( \theta_i \), \( i = 1, 2, ..., n \), constitute this missing information. Given \( \mathbf{\theta} = \{\theta_1, \theta_2, ..., \theta_n\} \), the complete data log likelihood \( \ell_c(\mathbf{B}, \mathbf{\theta}) \) can be written as

\[
\ell_c(\mathbf{B}, \mathbf{\theta}) = \sum_{i=1}^{n} \log \left[ \frac{\Gamma(n_i/2+a)}{\Gamma(a)b^a} \int \exp\left[-\frac{(\theta_i - Z_i^T \mathbf{\beta})^2}{2\tau^2}\right] \psi_i^{(n_i/2+a)} d\theta_i\right] - a\log(b) - 0.5\log(\tau^2) - \frac{(\theta_i - Z_i^T \mathbf{\beta})^2}{2\tau^2} - (n_i/2+a)\log(\psi_i),
\]

where the expression of \( \psi_i \) is given in Equation (4). Starting from an initial value of \( \mathbf{B}_0 \), the EM algorithm iteratively performs a maximization with respect to \( \mathbf{B} \). At the \( t \)th step the objective function maximized is

\[
Q(\mathbf{B} | \mathbf{B}^{(t-1)}) = E(\ell_c(\mathbf{B}, \mathbf{\theta}))
\]

\[
= \sum_{i=1}^{n} \log \left[ \frac{\Gamma(n_i/2+a)}{\Gamma(a)b^a} \int \exp\left[-\frac{(\theta_i - Z_i^T \mathbf{\beta})^2}{2\tau^2}\right] \psi_i^{(n_i/2+a)} d\theta_i\right] - a\log(b) - 0.5\log(\tau^2) - \frac{E(\theta_i - Z_i^T \mathbf{\beta})^2}{2\tau^2} - (n_i/2+a)E[\log(\psi_i)],
\]

The expectation in \( Q(\mathbf{B} | \mathbf{B}^{(t-1)}) \) is taken with respect to the conditional distribution of each \( \theta_i \) given the data, \( \pi(\theta_i | X_i, S_i^2, Z_i, \mathbf{B}^{(t-1)}) \), which is

\[
\pi(\theta_i | X_i, S_i^2, Z_i, \mathbf{B}) \propto \exp[-0.5(\theta_i - Z_i^T \mathbf{\beta})^2 / \tau^2] \psi_i^{(n_i/2+a)}.
\]

One challenge here is that the expectations are not available in closed form. Thus, we resort to a Monte Carlo method for evaluating the expressions. Suppose that \( R \) iid samples of \( \theta_i \) are available, say \( \theta_{i,1}, \theta_{i,2}, ..., \theta_{i,R} \). Then, each expectation of the form \( E[h(\theta_i)] \) can be approximated by the Monte Carlo mean

\[
E[h(\theta_i)] \approx \frac{1}{R} \sum_{r=1}^{R} h(\theta_{i,r}).
\]

However, drawing random numbers from the conditional distribution \( \pi(\theta_i | X_i, S_i^2, Z_i, \mathbf{B}^{(t-1)}) \) is also not straightforward since this is not a standard density. Samples are drawn using the accept-reject procedure (Robert and Casella 2004): For a sample from the target density \( f \), sample \( x \) from the proposal density \( g \), and accept the sample as a sample from \( f \) with probability \( f(x)/Mg(x) \) where \( M = \sup_x f(x)/g(x) \). One advantage of the accept-reject method is that the target density \( f \) only needs to be known up to a constant of proportionality which is the case for \( \pi(\theta_i | X_i, S_i^2, Z_i, \mathbf{B}^{(t-1)}) \) in (5); due to the non-standard form of the density, the normalizing constant cannot be found in a closed form. For the accept-reject algorithm, we used the normal density \( g(\theta_i) \propto \exp[-0.5(\theta_i - Z_i^T \mathbf{\beta})^2 / \tau^2] \) as the proposal density. The acceptance probability is calculated to be \( [(1/b + 0.5(n_i - 1)S_i^2) / (1/b + 0.5(n_i - 1)S_i^2 + 0.5(\theta_i - X_i))^2)]^{n_i/2+a} \). One can choose a better proposal distribution to increase acceptance probability or different algorithm (such as the adaptive rejection sampling or envelope accept-reject algorithms) but our chosen proposal worked satisfactorily in the studies we conducted.

The maximizer of \( Q(\mathbf{B} | \mathbf{B}^{(t-1)}) \) at the \( t \)th step can be described explicitly. The solutions for \( \mathbf{\beta} \) and \( \tau^2 \) are available in closed form as

\[
\mathbf{\beta}^{(t)} = \left( \sum_{i=1}^{n} Z_i Z_i^T \right)^{-1} \left( \sum_{i=1}^{n} Z_i E(\theta_i) \right)
\]

and

\[
(\tau^2)^{(t)} = \frac{1}{n} \sum_{i=1}^{n} E(\theta_i - Z_i^T \mathbf{\beta})^2,
\]

respectively. Also, \( a^{(t)} \) and \( b^{(t)} \) are obtained by solving \( S_a = \partial Q(\mathbf{B} | \mathbf{B}^{(t-1)}) / \partial a = 0 \) and \( S_b = \partial Q(\mathbf{B} | \mathbf{B}^{(t-1)}) / \partial b = 0 \) using the Newton-Raphson method where
\[ S_a = \sum_{i=1}^{n} \frac{a}{\hat{\sigma}} \log(\Gamma(n_i/2 + a)) \]
\[ - n \left( \frac{a}{\hat{\sigma}} \log(\Gamma(a)) \right) - n \log(b) - \sum_{i=1}^{n} E[\log(\psi_i)] \]
and
\[ S_b = - \frac{na}{b} + \sum_{i=1}^{n} \left( \frac{n_i}{2} + a \right) \frac{E[\psi_i]}{b^2} \]

We set \( B^{(i)} = (a^{(i)}, b^{(i)}, \hat{\theta}^{(i)}, (\tau^{(i)})^2) \) and proceed to the \((t+1)\)-st step. This maximization procedure is repeated until the estimate \( B^{(i)} \) converges. The MLE of \( B, \hat{B} = B^{(\infty)} \), once convergence is established.

### 3.2 Point estimate and confidence interval for \( \theta_i \)

Following the standard technique, the small area estimator of \( \theta_i \) is taken to be
\[ \hat{\theta}_i = E(\theta_i | X_i, S^2_i, Z_i, B) \bigg|_{b=b^*} \]  
(7)

the expectation of \( \theta_i \) with respect to the conditional density \( \pi(\theta_i | X_i, S^2_i, Z_i, B) \) with the maximum likelihood estimate \( \hat{B} \) plugged in for \( B \). The estimate \( \hat{\theta}_i \) is calculated numerically using the Monte Carlo procedure (6) described in the previous section. Subsequently, all quantities involving the unknown \( B \) will be plugged in by \( \hat{B} \) although we still keep using the notation \( B \) for simplicity.

Further, we develop a confidence interval for \( \theta_i \) based on a decision theory approach. Following Joshi (1969), Casella and Hwang (1991), Hwang et al. (2009), consider the loss function associated with the confidence interval \( C \) given by \( k/\sigma L(C) - I_c(0) \) where \( k \) is a tuning parameter independent of the model parameters, \( L(C) \) is the length of \( C \) and \( I_c(0) \) is the indicator function taking values 1 or 0 depending on whether \( \theta \in C \) or not. Note that this loss function takes into account both the coverage probability as well as the length of the interval; the positive quantity \( (k/\sigma) \) serves as the relative weight of the length compared to the coverage probability of the confidence interval. If \( k = 0 \), the length of the interval is not under consideration, which leads to the optimal \( C \) to be \((\infty, \infty)\) with coverage probability 1. On the other hand, if \( k = \infty \), then the coverage probability is 0, leading to optimal \( C \) to be a point set. The Bayes confidence interval for \( \theta_i \) is obtained by minimizing the risk function (the expected loss) \( E[(k/\sigma) L(C) - I_c(0)] | X_i, S^2_i, Z_i, B \). The optimal choice of \( C \) is given by
\[ C_i(B) = \{ \theta_i : k E(\sigma^{-1}_i | X_i, S^2_i, Z_i, B) < \pi(\theta_i | X_i, S^2_i, Z_i, B) \} \]  
(8)

Since \( C_i(B) \) is obtained by minimizing the posterior risk, one may like to interpret this as a Bayesian credible set. However, following Casella and Berger (1990, page 470), we will continue naming \( C_i(B) \) as a confidence interval. From an empirical Bayes perspective also, this terminology is more appropriate. How the tuning parameter \( k \) determines the confidence level of \( C_i(B) \) will be shown explicitly in Section 3.3.

Assuming \( k \) is known for the moment, we follow the steps below to calculate \( C_i(B) \). The conditional densities of \( \sigma^2_i \) and \( \theta_i \) are given by
\[ \pi(\sigma^2_i | X_i, S^2_i, Z_i, B) \propto \exp \left[ - \frac{0.5(X_i - Z_i^T \beta)^2}{(\sigma^2_i + \tau^2)} - \frac{0.5(n_i - 1)S_i^2 + 1}{b} \frac{1}{\sigma^2_i} \right] \]  
(9)

and (5), respectively, which as mentioned before, are not available in closed form. Thus, similar to the case of \( \theta_i \), \( E(\sigma^{-1}_i | X_i, S^2_i, Z_i, B) \) is computed numerically using the Monte Carlo method by approximating the expected value with the mean \( 1/N \sum_{i=1}^{N} \sigma^{-1}_{i,r} \) where \( \sigma^2_{i,r} \), \( r = 1, 2, ..., R \) are \( R \) samples from the conditional density \( \pi(\sigma^2_i | X_i, S^2_i, Z_i, B) \). The accept reject procedure is used to draw random numbers from \( \pi(\sigma^2_i | X_i, S^2_i, Z_i, B) \) with a proposal density given by the inverse Gamma
\[ \exp \left[ - \left( \frac{0.5(n_i - 1)S_i^2 + 1}{b} \frac{1}{\sigma^2_i} \right) \right] \]  
\[ \left( \sigma^2_i \right)^{(n_i-1)/2} \]  
and the acceptance probability
\[ \exp \left[ - \frac{0.5(X_i - Z_i^T \beta)^2}{(\sigma^2_i + \tau^2)} \right] \times \exp(0.5) \times |X_i - Z_i^T \beta|. \]

The next step is to determine the boundary values of \( C_i(B) \) by finding two \( \theta_i \) values that satisfy the equation \( k E(\sigma^{-1}_i | X_i, S^2_i, Z_i, B) - \pi(\theta_i | X_i, S^2_i, Z_i, B) = 0 \). This requires the normalizing constant in (5)
\[ D_i = \int_{-\infty}^{\infty} \exp(-0.5(\theta_i - Z_i^T \beta)^2 / \tau^2) \psi_{\sigma^{-1}} \psi_{\sigma^{-1}} d\theta_i \]

to be evaluated numerically. This is obtained using the Gauss-Hermite integration with 20 nodes.

### 3.3 Choice of \( k \)

The choice of the tuning parameter \( k \) in (8) is taken to be

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\[
k = k(B) = u_{1,0} \phi \left( t_{\alpha/2, \frac{n_i + 2\alpha + 2}{n_i - 1}} \right)
\]
where \( \phi \) is the standard normal distribution, \( t_{\alpha/2} \) is the \((1 - \alpha / 2)\)th percentile of \( t \) distribution with \((n_i - 1)\) degrees of freedom, and \( u_{1,0} = \sqrt{1 + \sigma_i^2 / \tau^2} \). Since \( u_{1,0} \) involves \( \sigma_i^2 \) which is unknown, an estimated version \( \hat{u}_{1,0} \) is obtained by plugging in the maximum a posteriori estimate
\[
\hat{\sigma}_i^2 = \hat{\sigma}_i^2(\hat{B}) = \arg \max_{\sigma_i^2} \pi(\sigma_i^2 | X_i, S_i, Z_i, B) |_{B = \hat{B}}
\]
in place of \( \sigma_i^2 \). Also, \( B \) is replaced by \( \hat{B} \) in (11). We demonstrate that the coverage probability of \( C_i(B) \) with this choice of \( k \) is close to \( 1 - \alpha \). Theoretical justifications are provided in Section 4.

### 3.4 Other related methods for comparison

Our method will be denoted as Method I. Three other methods to be compared are briefly described below.

**Method II:** Wang and Fuller (2003) considered the Fay-Herriot small area estimation model given by (1). Their primary contribution is the construction of the mean squared error estimation formulae for small area estimators with estimated sampling variances. In the process, they had constructed two formulae denoted by MSE_1 and MSE_2. We use MSE_1 for our comparisons, which was derived following the bias correction approach of Prasad and Rao (1990). The basic difference with our approach is that they did not smooth the sampling variances, only taking uncertainty into account while making inference on the small area parameters. The method of parameter estimation, which is moment based for all the model parameters, is also different from ours.

**Method III:** Hwang et al. (2009) considered the log-normal and inverse Gamma models for \( \sigma^2 \) in (2) for microarray data analysis. Their simulation study showed improved performance of confidence intervals for small area estimators under the log-normal model compared to the inverse gamma. We thus modified their log-normal model to add covariates and for unequal sample sizes \( n_j \) as follows:
\[
X_i | \theta_i, \sigma_i^2 \sim \text{Normal}(\theta_i, \sigma_i^2) \\
\theta_i \sim \text{Normal}(Z_i^T \beta, \tau^2)
\]
\[
\log S_i^2 = \log(\sigma_i^2) + \delta_i; \delta_i \sim N(m_i, \sigma^2_{\delta,i})
\]
\[
\log(\sigma_i^2) \sim N(\mu_i, \tau^2_i)
\]
independently for \( i = 1, 2, ..., n \). Note that the model for the means in (12) is identical to (1). The quantities \( \tau^2, m_i \) and \( \sigma^2_{\delta,i} \) are assumed to be known and are given by \( m_i = E[\log(\chi^2_{n_i-1}/(n_i-1))] \) and \( \sigma^2_{\delta,i} = \text{Var}[\log(\chi^2_{n_i-1}/(n_i-1))] \).

Thus, the sample size \( n_i \)'s determine the shape of the \( \chi^2 \) distribution via its degrees of freedom parameter. More importantly, as mentioned earlier, the different sample sizes account for different degrees of shrinkage for the corresponding true variance parameter. Similar to their estimation approach, the unknown model parameters \( \theta_i \) and \( \tau^2_i \) are estimated using a moment based approach in an empirical Bayes framework giving \( \hat{\mu}_i \) and \( \hat{\tau}^2_i \), respectively.

Note that in Hwang et al. (2009), these estimates are obtained based on the hierarchical model for \( \sigma^2_i \) of (13) only without regard to the modelling (1) of the mean. We refer to the Section 5 of their paper for details of the estimation of the hyper-parameters. We follow the same procedure using only (13) to estimate \( \mu_i \) and \( \tau^2_i \), in the case of unequal sample sizes.

The Bayes estimate of \( \sigma^2_i \) is derived to be
\[
\hat{\sigma}^2_{i,B} = \exp[\text{E}[\log(\sigma^2_i) | \text{Var}(S_i^2)]]
\]
\[
= \left\{ \frac{\hat{S}_i^2}{\text{exp}(m_i)} \right\}^{\hat{M}_{\tau,i}} \exp(\mu_i(1 - M_{\tau,i}))
\]
where \( M_{\tau,i} = \tau^2 / (\tau^2 + \sigma^2_{\hat{\mu}_i}) \) and with estimates plugged in for the unknown quantities. The conditional distribution of \( \theta_i \) given \( (X_i, S_i^2) \), is
\[
\pi(\theta_i | X_i, S_i^2) = \int_0^\infty \pi(\theta_i | X_i, S_i^2, \sigma_i^2) \pi(\sigma_i^2 | X_i, S_i^2) d\sigma_i^2,
\]
is approximated as \( \pi(\theta_i | X_i, S_i^2) \approx \int_0^\infty \pi(\theta_i | X_i, S_i^2, \hat{\sigma}^2_{i,B}) \pi(\hat{\sigma}^2_{i,B} | X_i, S_i^2) d\hat{\sigma}^2_{i,B} \). This suggests the approximate Bayes estimator of the small area parameters given by
\[
\hat{\theta}_i = E(\theta_i | X_i, \hat{\sigma}^2_{i,B}) = M_i X_i + (1 - M_i) Z_i^T \hat{\beta},
\]
where \( M_i = \tau^2 / (\tau^2 + \hat{\sigma}^2_{i,B}) \). The confidence interval for \( \theta_i \) is obtained as
\[
C_i^{\text{IH}} = \left\{ \theta_i : \frac{\theta_i - \hat{\theta}_i}{M_i \hat{\sigma}^2_{i,B}} < -2\ln\{k / 2\pi\} - \ln(M_i) \right\}.
\]

In Section 3 of Hwang et al. (2009) pages 269-271, the interval \( C_i^{\text{IH}} \) is matched with the 100(1 - \( \alpha \))% \( t \)-interval \([|\theta_i - X_i| / t S_i] \) to obtain the expression of \( k \) as \( k = k_i = \text{exp}(-t^2 / 2) \text{exp}(m_i / (\sqrt{2} \pi)) \).

**Method IV:** This method comprises of a special case of the Fay-Herriot model in (1) but with the estimation of model parameters adopted from Qiu and Hwang (2007). Qiu and Hwang (2007) considered the model
\[
X_i | \theta_i, \sigma^2 \sim \text{Normal}(\theta_i, \sigma^2)
\]
\[
\theta_i \sim \text{Normal}(0, \tau^2),
\]
Due to equation (9), the shrinkage $S_i$ depends on $2(\theta_i - \mu_i + \gamma_i)$ as defined in (11). In a similar way, $\hat{\gamma}_i = (\Sigma_{i=1}^n X_i^2)^{1/2}$. The confidence interval for $\theta_i$ is $\hat{\theta}_i \pm \tau_i (M_i)$, where $v_i (M) = \sigma^2 M \gamma_i$ with $\gamma_i$ denoting the standard normal cut-off point corresponding to desired level of confidence coefficient and $v_i(0) = 0$. Here For the purpose of comparisons with our method, the first level of the hierarchical model in (16) is modified as follows:

$$X_i = Z_i^T \beta + v_i + e_i$$

where $v_i \sim \text{Normal}(0, \tau^2)$ and $e_i \sim \text{Normal}(0, S_i^2)$ independently for $i = 1, 2, ..., n$, and $S_i^2$ is treated as known. Following Qiu and Hwang (2007), $\tau^2$ is estimated by

$$\xi^2 = \frac{1}{n-p} \left[ \sum_i \hat{u}_{i}^2 - \sum_i S_i^2 \left\{ 1 - Z_i^T \left( \sum_i Z_i Z_i^T \right)^{-1} Z_i^T \right\} \right]$$

and $\hat{\tau}^2 = \max(\xi^2, 1/n)$ where $u_i = X_i - Z_i^T \hat{\beta}$ and $\hat{\beta} = (\Sigma_{i=1}^n Z_i Z_i^T)^{-1}(\Sigma_{i=1}^n Z_i X_i)$. Next, define $\hat{M}_{0i} = \tau^2 / (\hat{\tau}^2 + S_i^2)$ and $\hat{M}_i = \max(M_{0i}, M_i)$ where in the latter expression, $\hat{M}_{0i}$ is truncated by $M_{ii} = 1 - Q_\alpha / (n_i - 2)$, and $Q_\alpha$ is the $\alpha$th quantile of a chi-squared distribution with $n_i$ degrees of freedom. This $\hat{M}_i$ is used in the formula of the confidence interval for $\theta_i$ given earlier. When applying this method in our simulation study and real data analysis, we modified the model to accommodate such unequal sample sizes and covariate information mentioned earlier.

**Remark 1.** Hwang et al. (2009) choose $k$ by equating (15) to the $t$ interval based on only $X_i$ for the small area parameters $\theta_i$. Note that $X_i$ is the direct survey estimator. Consequently, this choice of $k$ does not have any direct control over the coverage probability of the interval constructed under shrinkage estimation. On the other hand, our proposed choice of $k$ has been derived to maintain nominal coverage under, specifically, shrinkage estimation.

**Remark 2.** Note that without any hierarchical modelling assumption, $S_i$ and $X_i$ are independent as $S_i^2$ and $X_i$ are, respectively, ancillary and the complete sufficient statistics for $\theta_i$. However, under models (1) and (2) the conditional distribution of $\sigma^2_i$ and $\theta_i$ involve both $X_i$ and $S_i^2$ which is seen from (5) and (9).

**Remark 3.** In Hwang et al. (2009), the shrinkage estimator for $\sigma^2_i$ is based only on the information on $S_i^2$ and not of both $X_i$ and $S_i^2$. The Bayes estimator of $\sigma^2_i$ is plugged into the expression for the Bayes estimator of small area parameters. Thus, Hwang et al.’s small area estimator is written as $E(\theta_i | X_i, \hat{\sigma}^2_{i,b})$ in (14) where $\hat{\sigma}^2_{i,b}$ is the Bayes estimator of $\sigma^2_i$. Due to equation (9), the shrinkage estimator of $\sigma^2_i$ depends on $(X_i - Z_i^T \beta)^2$ in addition to $S_i^2$ in contrast to Hwang et al. (2009). We believe this could be the reason for improved performance of our method compared to Hwang et al. (2009).

**Remark 4.** As mentioned previously, the degree of freedom associated with the $\chi^2$ distribution for the sampling variance need not to be simply $n_i - 1$, $n_i$ being the sample size for $i$th area. There is no sound theoretical result for determining the degree of freedom when the survey design is complex. The article Wang and Fuller (2003) approximated the $\chi^2$ with a normal based on the Wilson-Hilferty approximation. If one knows the exact sampling design then the simulation based guideline of Maples et al. (2009) could be useful. For county level estimation using the American Community Survey, Maples et al. (2009) suggested the estimated degrees of freedom of 0.36 $\times \sqrt{n_i}$.

### 4. Theoretical justification

Theoretical justification for the choice of $k$ according to equation (10) is presented in this section. As in Hwang et al. (2009), the conditional distribution of $\theta_i$ given $X_i$ and $S_i^2$ can be approximated as $\pi(\theta_i | X_i, S_i^2, B) \approx \pi(\theta_i | X_i, S_i^2, \hat{B})$ where $\hat{B}$ is as defined in (11). In a similar way, approximate $E(\sigma^2_i | X_i, S_i^2, B)$ by $E(\sigma^2_i | X_i, S_i^2, \hat{B}) \approx \hat{\sigma}^2_i$. Based on these approximations, we have $C_i(B) \approx \hat{C}_i(B)$ where $\hat{C}_i(B)$ is the confidence interval for $\theta_i$ given by $\hat{C}_i(B) = \{ \theta_i: \pi(\theta_i | X_i, S_i^2, B, \hat{\sigma}^2_i) \geq k \hat{\sigma}^2_i \}$. From (1), it follows that the conditional density $\pi(\theta_i | X_i, S_i^2, B, \sigma^2_i)$ is a normal with mean $\mu_i$ and variance $v_i$, where $\mu_i$ and $v_i$ are given by the expressions

$$\mu_i = w_i X_i + (1 - w_i) Z_i^T \hat{\beta},$$

$$v_i = \left( \frac{1}{\sigma^2_i} + \frac{1}{\hat{\tau}^2} \right)^{-1} = \sigma^2_i \left( 1 + \frac{\sigma^2_i}{\hat{\tau}^2} \right)^{-1},$$

and

$$w_i = \frac{1}{1 + \sigma^2_i + \frac{1}{\hat{\tau}^2}}.$$
where \( \hat{\mu}_i \) is the expression for \( \mu_i \) in (17) with \( \sigma_i^2 \) replaced by \( \hat{\sigma}_i^2 \). Now consider the behavior of \( \hat{\sigma}_i^2 \equiv \hat{\sigma}_i^2(B) \) as \( \tau^2 \) ranges between 0 and \( \infty \). When \( \tau^2 \to \infty \), \( \hat{\sigma}_i^2 \) converges to

\[
\hat{\sigma}_i^2(\infty) = \frac{(n_i - 1)S_i^2 + \frac{1}{2}b}{(n_i - 1)S_i^2 + \frac{2}{b}}.
\]

Similarly, when \( \tau^2 \to 0 \), \( \hat{\sigma}_i^2 \) converges to

\[
\hat{\sigma}_i^2(0) = \frac{(X_i - Z_i^T \beta)^2 + (n_i - 1)S_i^2 + \frac{2}{b}}{n_i + 2a + 2}.
\]

For all intermediate values of \( \tau^2 \), we have \( \min(\hat{\sigma}_i^2(0), \hat{\sigma}_i^2(\infty)) \leq \hat{\sigma}_i^2 \leq \max(\hat{\sigma}_i^2(0), \hat{\sigma}_i^2(\infty)) \). Therefore, it is sufficient to consider the following two cases: (i) \( \hat{\sigma}_i^2 \geq \hat{\sigma}_i^2(\infty) \), where it follows that \( (n_i + 2a + 2)\hat{\sigma}_i^2 = (n_i + 2a + 1)\hat{\sigma}_i^2 + \hat{\sigma}_i^2 \geq (n_i - 1)S_i^2 + 2/b + \hat{\sigma}_i^2 \geq (n_i - 1)S_i^2 \), and (ii) \( \hat{\sigma}_i^2 \geq \hat{\sigma}_i^2(0) \), where it follows that \( (n_i + 2a + 2)\hat{\sigma}_i^2 = (X_i - Z_i^T \beta)^2 + (n_i - 1)S_i^2 + 2/b \geq (n_i - 1)S_i^2 \). So, in both cases (i) and (ii),

\[
(n_i + 2a + 2)\hat{\sigma}_i^2 \geq (n_i - 1)S_i^2.
\]

Since \( \theta_i - \mu_i \sim N(0, \sigma_i^2\tau^2/(\sigma_i^2 + \tau^2)) \) and \( (n_i - 1)S_i^2/\sigma^2 \sim \chi^2_{(n_i - 1)} \), the confidence interval

\[
D_i = \left\{ \theta_i : u_0 \frac{|\theta_i - \mu_i|}{S_i} \leq t_{a/2} \right\}
\]

has coverage probability \( 1 - \alpha \). Thus, if \( u_0 \) and \( \hat{\mu}_i \) are replaced by \( u_0 \) and \( \hat{\mu}_i \), it is expected that the resulting confidence interval \( \hat{D}_i \), say, will have coverage probability of approximately \( 1 - \alpha \). From (19), we have

\[
P(\hat{C}_i(B)) \geq P(\hat{D}_i) \approx 1 - \alpha,
\]

establishing an approximate lower bound of \( 1 - \alpha \) for the confidence level of \( \hat{C}_i(B) \).

In (21), \( B \) was assumed to be fixed and known. When \( B \) is unknown, we replace \( B \) by its marginal maximum likelihood estimate \( \hat{B} \). Since (21) holds regardless of the true value of \( B \), substituting \( \hat{B} \) for \( B \) in (21) will involve an order \( O(1/\sqrt{N}) \) of error where \( N = \sum_{i=1}^{n_u} n_i \). Compared to each single \( n_i \), this pooling of \( n_i \)'s is expected to reduce the error significantly so that \( \hat{C}_i(B) \) is sufficiently close to \( \hat{C}_i(B) \) to satisfy the lower bound of \( 1 - \alpha \) in (21).

5. A simulation study

5.1 Simulation setup

We considered a simulation setting using a subset of parameter configurations from Wang and Fuller (2003). Each sample in the simulation study was generated from the following steps: First, generate observations using the model

\[ X_{ij} = \beta + u_i + e_{ij}, \]

where \( u_i \sim N(0, \tau^2) \) and \( e_{ij} \sim N(0, \sigma_i^2) \), independently for \( j = 1, ..., n_i \) and \( i = 1, ..., n \). Then, the random effects model for the small area mean, \( X_i \), is

\[ X_i = \beta + u_i + e_i, \]

where \( X_i = \bar{X}_i = \frac{1}{n_i} \sum_{j=1}^{n_i} X_{ij} \) and \( e_i = \bar{e}_i = \frac{1}{n_i} \sum_{j=1}^{n_i} e_{ij} \). Therefore, \( X_i \sim N(\theta_i, \sigma_i^2) \) where \( \theta_i = \beta + u_i, \theta_i \sim N(\beta, \tau^2) \) and \( e_i \sim N(0, \sigma_i^2) \). We estimated \( \sigma_i^2 \) with the unbiased estimator

\[ S_i^2 = (n_i - 1)^{-1} \sum_{j=1}^{n_i} (X_{ij} - \bar{X}_i)^2, \]

and it follows that \( (n_i - 1)S_i^2/\sigma_i^2 \sim \chi^2_{(n_i - 1)} \), independently for \( i = 1, 2, ..., n \). Note that the simulation layout has ignored the second level modeling of sampling variances in (2). Thus, our result will indicate robustness with respect to the variance model misspecification.

The above steps produced the data \( (X_i, S_i^2), i = 1, ..., n \). To simplify the simulation, we do not choose any covariate information \( Z_i \). Similar to Wang and Fuller (2003), we set all \( n_i \)'s equal to \( m \) to ease programming efforts. However, the true sampling variances are still chosen to be unequal: One-third of the \( \sigma_i^2 \) are set to 1, another one-third are set to 4, and the remaining one-third are set to 16. We take \( \beta = 10 \) and three different choices of \( \tau^2 = 0.25, 1 \) and \( 4 \). These parameter values are chosen from Qiu and Hwang (2007). For each of \( \tau^2 \), we generated 200 samples for the two combinations \( (m, n) = (9, 36) \) and \( (18, 180) \).

In the simulation study, we compare the proposed method with the methods of Wang and Fuller (2003), Hwang et al. (2009) and Qiu and Hwang (2007) which are referred to as Methods I, II, III, and IV, respectively, based on bias, mean squared error (MSE), coverage probability (CP) of the confidence intervals and the length of the confidence intervals (ALCI). Table 1 contains the parameter estimates for \( a, b, \beta \) and \( \tau^2 \). The numerical results indicate good performance of the maximum likelihood estimates for the model parameters; the estimated values of \( \beta \) and \( \tau^2 \) are close to the true values indicating good robustness properties with respect to distributional misspecification in the second level of (2). Statistically significant estimates for both \( a \) and \( b \) indicate that “shrunk” sampling variances are incorporated in the proposed method. Tables 2, 3 and 4 provide numerical results averaged over areas within each group having the same true sampling variances. The results in the Tables are based on 200 replications.
Table 1
Simulation results for the model parameters, $a$ (top left panel), $b$ (top right panel), $\beta$ (bottom left panel) and $\tau^2$ (bottom right panel). Here SD represents the standard deviation over 200 replicates. We took $\beta = 10$ and $\tau^2 = 0.25$, $1$ and $4$

<table>
<thead>
<tr>
<th>$\tau^2$</th>
<th>$n = 36$, $m = 9$</th>
<th>$n = 180$, $m = 18$</th>
<th>$\beta$</th>
<th>$\tau^2$</th>
<th>$n = 36$, $m = 9$</th>
<th>$n = 180$, $m = 18$</th>
</tr>
</thead>
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<td>SD</td>
<td>Mean</td>
<td>SD</td>
<td>Mean</td>
<td>SD</td>
</tr>
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<td>1.0328</td>
<td>0.0442</td>
<td>0.25</td>
<td>0.3992</td>
</tr>
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</tr>
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<td>9.9970</td>
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</tr>
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Table 2
Simulation results for prediction when $\tau^2 = 0.25$. Here MSE, ALCI, CP represent the mean squared error, average confidence interval width, and coverage probability, respectively

<table>
<thead>
<tr>
<th>$\sigma^2_i$</th>
<th>Method</th>
<th>$n = 36$, $m = 9$</th>
<th>Method</th>
<th>$n = 180$, $m = 18$</th>
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</thead>
<tbody>
<tr>
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<tr>
<td>bias</td>
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<td>-0.0158</td>
<td>-0.0109</td>
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<tr>
<td>MSE</td>
<td>1</td>
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<td>0.0096</td>
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</tr>
<tr>
<td>ALCI</td>
<td>1</td>
<td>0.3066</td>
<td>0.3805</td>
<td>0.2258</td>
</tr>
<tr>
<td>CP</td>
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</tr>
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<td>16</td>
<td>0.9365</td>
<td>0.9933</td>
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</table>

Table 3
Simulation results for prediction when $\tau^2 = 1$. Here MSE, ALCI, CP represent the mean squared error, average confidence interval width and coverage probability, respectively

<table>
<thead>
<tr>
<th>$\sigma^2_i$</th>
<th>Method</th>
<th>$n = 36$, $m = 9$</th>
<th>Method</th>
<th>$n = 180$, $m = 18$</th>
</tr>
</thead>
<tbody>
<tr>
<td>Relative</td>
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<tr>
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<td></td>
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Bias Comparisons: In most cases, the bias of the four methods are comparable. There is no clear evidence of significant differences between them in terms of the bias. High sampling variance gives more weight to the population mean by construction that makes the estimator closer to the mean at the second level. On the other hand, Methods I - III use shrinkage estimators of the sampling variances which would be less than the maximum of all sampling variances. Thus, Methods I - III tend to have little more bias. However, due to shrinkage in sampling variances, one may expect a gain in the variance of the estimators which, in turn, makes the MSE smaller. Among Methods I - III, Method I performed better compared to Methods II and III, which were quite similar to each other. The maximum gain using Method I compared to Method II is 99%.

MSE Comparisons: In terms of the MSE, Method I performed consistently better than the other three in all cases except when the ratio of $\sigma_i^2$ to $\tau^2$ is the lowest: $(\sigma_i^2 = 1) / (\tau^2 = 4) = 0.25$. In this case, the variance between small areas (model variance) is much higher than the variance within the areas (sampling variance). When using our method to estimate $\theta_i$, the information “borrowed” from other areas may misdirect the estimation: The estimated mean of the Gamma distribution for $\sigma_i^2$ from the second level in (2) is $\hat{a} \hat{b}$ which equals 0.44 approximately for both the $(m, n)$ combinations of (9, 36) and (18, 180) (the true value is $ab = 0.4$). Thus, $E(\sigma_i^2 | X_i, S_i^2, \hat{B})$ is significantly smaller than 1 due to shrinkage towards the mean for the group which has the true value of $\sigma_i^2 = 1$. Also, since $\sigma_i^2$ is smaller than $\tau^2$, the weight of $X_i$ should be much more compared to $\hat{B}$, the overall mean. However, due to underestimation of $\sigma_i^2$ in this case, the resulting estimator puts less weight on $X_i$ which leads to higher MSE. However, this underestimation will decrease for large sample sizes due to the consistency of Bayes estimators. This fact is actually observed when the sample size increases from $n = 36$ to $n = 180$ for the case $\sigma_i^2 = 1$ and $\tau^2 = 4$. Compared to Method II, Method I shows gains in most of the simulation cases: the maximum gain is 30% while the only loss is 9% for the combination $\sigma_i^2 = 1$ and $\tau^2 = 4$ for $n = 36$ and $m = 9$. Similarly, for Method III, the maximum gain of Method I is 77% and the only loss of 11% is for the same parameter and sample size specifications.

ACP Comparisons: We obtained confidence intervals with confidence level 95%. Methods I and III do not indicate any under-coverage. This is expected from their optimal confidence interval construction. Method I meets the nominal coverage rate more frequently than any other methods. Method II has some under-coverage and can go as low as 82%.

ALCI Comparisons: Method I produced considerably shorter confidence intervals in general. Method IV produced comparable lengths as the other methods in all cases except when $\sigma_i^2$ was high, in which case, the lengths were considerably higher. The confidence interval proposed in Qiu and Hwang (2007) does not have good finite sample properties, particularly for small $\tau^2$. To avoid low coverage, they proposed to truncate $M_\theta = \tau^2/(\tau^2 + \sigma_i^2)$ with a positive number $M_\theta = 1 - Q_{\alpha/n}(v - 2)$ for known $\sigma_i^2$, where $Q_{\alpha/n}$ is the $\alpha/n$-quantile of a chi-squared distribution with $v$ degrees of freedom. When the ratio of sampling
variance to model variance, \( \sigma_i^2 / \tau^2 \), is high, \( M_i \) tends to be higher than \( M_{\nu} \). This results in a nominal coverage but with larger interval lengths. For example, in case of \((\sigma_i^2, \tau^2) = (16, 0.25)\), the ALCI is 11.13 for Method IV whereas ALCI is only 2.78 and 4.56 for Methods I and II.

5.2 Robustness study

In order to study the robustness of the proposed method with respect to departures from the normality assumption in the errors, we conducted the following simulation study. Data was generated as before but with \( e_{ij} \)'s drawn from a double-exponential (Laplace) and an uniform distribution. The estimators from Methods II and III had little effect. This is perhaps due to the fact that these methods used moment based estimation for model parameter estimation. Method IV resulted in larger relative bias, MSE and ALCI, and lower coverage probability. The MSE from Method I is always lower than that from Method II. For \( \tau^2 = 0.25 \) and 1, ALCI is smaller for Method I compared to Method II for \((n = 36, m = 9)\) but the results are opposite when \((n = 180, m = 18)\). In terms of CP, Method II has some under coverage (lowest is 80%). However, Method I did not have any under-coverage. In order to save space we only provide the results for parameters \( a \), \( b \), \( \beta \) and \( \tau^2 \) under the Laplace errors (see Table 5).

6. Real data analysis

We illustrate our methodology based on a widely studied example. The data set is from the U.S. Department of Agriculture and was first analyzed by Battese (1988). The data set is on corn and soybeans productions in 12 Iowa counties. The sample sizes for these areas are small, ranging from 1 to 5. We shall consider corn only to save space. For the proposed model, the sample sizes \( n_i \geq 1 \) necessarily. Therefore, modified data from You and Chapman (2006) with \( n_i \geq 2 \) are used. The mean reported crop hectares for corn \((X_i)\) are the direct survey estimates and are given in Table 6. Table 6 also gives the sample variances which are calculated based on the original data assuming simple random sampling. The sample standard deviation varies widely, ranging from 5.704 to 53.999 (the coefficient of variation varies from 0.036 to 0.423). Two covariates are considered in Table 6: \( Z_{1i} \), the mean of pixels of corn, and \( Z_{2i} \), the mean of pixels of soybean, from the LANDSAT satellite data.

### Table 5
Simulation results for the model parameters, \( a \) (top left panel), \( b \) (top right panel), \( \beta \) (bottom left panel) and \( \tau^2 \) (bottom right panel) when the errors follow a laplace distribution. Here SD represents the standard deviation over 200 replicates. We took \( \beta = 10 \) and \( \tau^2 = 0.25, 1 \) and 4

<table>
<thead>
<tr>
<th>( \tau^2 )</th>
<th>( n = 36, m = 9 )</th>
<th>( n = 180, m = 18 )</th>
<th>( \tau^2 )</th>
<th>( n = 36, m = 9 )</th>
<th>( n = 180, m = 18 )</th>
</tr>
</thead>
<tbody>
<tr>
<td></td>
<td>Mean</td>
<td>SD</td>
<td>Mean</td>
<td>SD</td>
<td>Mean</td>
</tr>
<tr>
<td>( a )</td>
<td>( 0.25 )</td>
<td>( 0.9624 )</td>
<td>( 0.1632 )</td>
<td>( 0.5793 )</td>
<td>( 0.1733 )</td>
</tr>
<tr>
<td></td>
<td>( 1 )</td>
<td>( 0.9628 )</td>
<td>( 0.1657 )</td>
<td>( 0.5816 )</td>
<td>( 0.1777 )</td>
</tr>
<tr>
<td></td>
<td>( 4 )</td>
<td>( 0.9689 )</td>
<td>( 0.1694 )</td>
<td>( 0.5758 )</td>
<td>( 0.1796 )</td>
</tr>
<tr>
<td>( b )</td>
<td>( 0.25 )</td>
<td>( 9.9736 )</td>
<td>( 0.3775 )</td>
<td>( 0.2696 )</td>
<td>( 0.0882 )</td>
</tr>
<tr>
<td></td>
<td>( 1 )</td>
<td>( 9.9753 )</td>
<td>( 0.3709 )</td>
<td>( 1.0508 )</td>
<td>( 0.2501 )</td>
</tr>
<tr>
<td></td>
<td>( 4 )</td>
<td>( 9.9736 )</td>
<td>( 0.4835 )</td>
<td>( 3.9624 )</td>
<td>( 1.1719 )</td>
</tr>
</tbody>
</table>

### Table 6
Corn data from You and Chapman (2006)

<table>
<thead>
<tr>
<th>County</th>
<th>( n_i )</th>
<th>( X_i )</th>
<th>( Z_{1i} )</th>
<th>( Z_{2i} )</th>
<th>( \sqrt{S_i^2} )</th>
</tr>
</thead>
<tbody>
<tr>
<td>Franklin</td>
<td>3</td>
<td>158.623</td>
<td>318.21</td>
<td>188.06</td>
<td>5.704</td>
</tr>
<tr>
<td>Pocahontas</td>
<td>3</td>
<td>102.523</td>
<td>257.17</td>
<td>247.13</td>
<td>43.406</td>
</tr>
<tr>
<td>Winnebago</td>
<td>3</td>
<td>112.773</td>
<td>291.77</td>
<td>185.37</td>
<td>30.547</td>
</tr>
<tr>
<td>Wright</td>
<td>3</td>
<td>144.297</td>
<td>301.26</td>
<td>221.36</td>
<td>53.999</td>
</tr>
<tr>
<td>Webster</td>
<td>4</td>
<td>117.595</td>
<td>262.17</td>
<td>247.09</td>
<td>21.298</td>
</tr>
<tr>
<td>Hancock</td>
<td>5</td>
<td>109.382</td>
<td>314.28</td>
<td>198.66</td>
<td>15.661</td>
</tr>
<tr>
<td>Kossuth</td>
<td>5</td>
<td>110.252</td>
<td>298.65</td>
<td>204.61</td>
<td>12.112</td>
</tr>
<tr>
<td>Hardin</td>
<td>5</td>
<td>120.054</td>
<td>325.99</td>
<td>177.05</td>
<td>36.807</td>
</tr>
</tbody>
</table>

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The estimates of $B$ are as follows: $a = 1.707$, $b = 0.00135$, $\tau^2 = 90.58$ and $\beta = (-186.0, 0.7505, 0.4100)$. The estimated prior mean of $1/\sigma_i^2$ which is the mean of the Gamma distribution with parameters $a$ and $b$ is $ab = 0.002295$ with a square root of 0.048 (note that $1/0.048 = 20.85$ consistent with the range of the sample standard deviations between 5.704 and 53.999). The small area estimates and their confidence intervals are summarized in Table 7 and Figure 1. Point estimates of all 4 methods are comparable: the summary measures comprising of the mean, median, and range of the small area parameter estimates for Methods I, II, III, and IV are (121.9, 124.1, 122.2, 122.6), (125.2, 120.4, 115.0, 114.5) and (23.1, 53.0, 58.4, 56.6), respectively. The distribution of $\hat{\theta}_i$ (plotted based on considering all the $i$'s) are summarized in Figure 2 which shows that there is a significant difference in their variability. Method I has the lowest variability and is superior in this sense. Further, smoothing sampling variiances has strong implication in measuring uncertainty and hence in the interval estimation. The proposed method has the shortest confidence interval on an average compared to all other methods. Methods II and III provide intervals with negative lower limits. This seems unrealistic because the direct average of area under corn is positive and large for all the 12 counties (the crude confidence intervals ($\chi_i \pm \tau_0.025 S_i$) do not contain zero for any of the areas either). Note that Method II does not have any theoretical support on its confidence intervals. Methods II and III produce wider confidence intervals when the sampling variance is high. For example, the sample size for both Franklin county and Pocahontas county is three, but sampling standard deviations are 5.704 and 43.406. Although the confidence interval under Method I is comparable, they are wide apart for Methods II and III. This is because although these methods consider the uncertainty in sampling variance estimates, the smoothing did not use the information from direct survey estimates, resulted the underlying sampling variance estimates remain highly variable (due to small sample size). In effect, the variance of the variance estimator (of the point estimates) is bigger compared to that in method I. This is further confirmed by the fact that the intuitive standard deviations of the “smoothed” small area estimates (one fourth of the interval) are smaller and less variable under method I compared to the others. Another noticeable aspect of our method is that the interval widths are similar for counties with same sample size. This could be an indication of obtaining equi-efficient estimators for equivalent sample sizes.

**Model selection:** For choosing the best fitting model, we used the Bayesian Information Criteria (BIC) which takes into account both the likelihood as well as the complexity of the fitted models. We calculated BICs for the models used in Methods I and III (Hwang et al. 2009). These two models have the same numbers of parameters with a difference in only the way the parameters are estimated. The model BIC for Method I is 210.025 and that for Method III is 227.372. This indicates superiority of our model. We could not compute the BIC for Wang and Fuller (2003) since they did not use any explicit likelihood.

### Table 7

Results of the corn data analysis. Here CI and LCI represent the confidence interval and the length of the confidence interval, respectively

<table>
<thead>
<tr>
<th>County</th>
<th>$\hat{\theta}_i$</th>
<th>CI</th>
<th>LCI</th>
<th>$\hat{\theta}_i$</th>
<th>CI</th>
<th>LCI</th>
</tr>
</thead>
<tbody>
<tr>
<td>Franklin</td>
<td>131.8106</td>
<td>104.085, 159.372</td>
<td>55.287</td>
<td>155.4338</td>
<td>124.151, 193.094</td>
<td>68.943</td>
</tr>
<tr>
<td>Pocahontas</td>
<td>108.7305</td>
<td>80.900, 136.436</td>
<td>55.536</td>
<td>102.3682</td>
<td>-38.973, 244.019</td>
<td>282.993</td>
</tr>
<tr>
<td>Winnebago</td>
<td>109.0559</td>
<td>81.430, 136.646</td>
<td>55.216</td>
<td>115.9093</td>
<td>-53.768, 279.314</td>
<td>333.083</td>
</tr>
<tr>
<td>Wright</td>
<td>131.6113</td>
<td>103.736, 159.564</td>
<td>55.828</td>
<td>131.0674</td>
<td>8.330, 280.263</td>
<td>271.932</td>
</tr>
<tr>
<td>Webster</td>
<td>113.1484</td>
<td>92.805, 133.348</td>
<td>40.543</td>
<td>109.4795</td>
<td>32.514, 202.675</td>
<td>170.161</td>
</tr>
<tr>
<td>Hancock</td>
<td>129.4279</td>
<td>111.781, 147.193</td>
<td>35.412</td>
<td>124.1028</td>
<td>56.750, 162.013</td>
<td>105.262</td>
</tr>
<tr>
<td>Kossuth</td>
<td>121.0071</td>
<td>103.451, 138.626</td>
<td>35.175</td>
<td>116.7147</td>
<td>-53.768, 279.314</td>
<td>333.083</td>
</tr>
<tr>
<td>Hardin</td>
<td>130.2520</td>
<td>112.373, 148.114</td>
<td>35.741</td>
<td>137.7983</td>
<td>51.734, 188.373</td>
<td>136.638</td>
</tr>
<tr>
<td>Franklin</td>
<td>158.4677</td>
<td>128.564, 188.370</td>
<td>59.805</td>
<td>157.7383</td>
<td>146.999, 168.477</td>
<td>21.478</td>
</tr>
<tr>
<td>Pocahontas</td>
<td>100.1276</td>
<td>-44.039, 244.295</td>
<td>288.334</td>
<td>101.1661</td>
<td>19.444, 182.887</td>
<td>163.442</td>
</tr>
<tr>
<td>Winnebago</td>
<td>114.1473</td>
<td>0.065, 228.228</td>
<td>228.163</td>
<td>113.7746</td>
<td>56.263, 171.286</td>
<td>115.022</td>
</tr>
<tr>
<td>Wright</td>
<td>140.3717</td>
<td>-24.119, 304.862</td>
<td>328.982</td>
<td>143.2244</td>
<td>41.559, 244.889</td>
<td>203.330</td>
</tr>
<tr>
<td>Webster</td>
<td>115.7865</td>
<td>50.297, 181.275</td>
<td>130.978</td>
<td>115.2244</td>
<td>75.124, 193.202</td>
<td>136.638</td>
</tr>
<tr>
<td>Hancock</td>
<td>111.3087</td>
<td>66.213, 156.403</td>
<td>90.189</td>
<td>113.1766</td>
<td>89.520, 135.127</td>
<td>45.607</td>
</tr>
<tr>
<td>Kossuth</td>
<td>110.9585</td>
<td>74.366, 147.550</td>
<td>73.184</td>
<td>112.3239</td>
<td>89.520, 135.127</td>
<td>45.607</td>
</tr>
</tbody>
</table>
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Figure 1 Corn hectares estimation. The horizontal line for each county displays the confidence interval of $\hat{\theta}_i$, with $\hat{\theta}_i$ marked by the circle, for (I) Proposed method, (II) Wang and Fuller (2003), (III) Hwang et al. (2009) and (IV) Qiu and Hwang (2007).

Figure 2 Boxplot of estimates of corn hectares for each county. (I) to (IV) are the 4 methods corresponding to Figure 1.

7. Conclusion

In this paper, joint area level modeling of means and variances is developed for small area estimation. The resulting small area estimators are shown to be more efficient than the traditional estimators obtained using Fay-Herriot models which only shrink the means. Although our model is same as one considered in Hwang et al. (2009), our method of estimation is different in two ways: In the determination of the tuning parameter $k$ and the use of $\pi(\sigma_i^2 \mid X_i, S_i^2, Z_i)$ (which depends additionally on $X_i$), instead of $\pi(\sigma_i^2 \mid S_i^2, Z_i)$, for constructing the conditional distribution of the small area parameters $\theta_i$. We demonstrated robustness properties of the model when the assumption that $\sigma_i^2$ arise from a inverse Gamma distribution is violated. The borrowing of $X_i$ information when estimating $\sigma_i^2$ as well as the robustness with respect to prior elicitation demonstrate the superiority of our proposed method. The parameter values chosen in the simulation study are different than in the real data analysis. The real data analysis given here is merely for illustration purposes. Our main aim was to
develop the methodology of mean-variance modeling and contrast with some closely related methods to show its effectiveness. For this reason, we chose parameter settings in the simulation to be the same as in the well-known small area estimation article Wang and Fuller (2003).

Obtaining improved sampling variance estimators is a byproduct of the proposed approach. We have provided an innovative estimation technique which is theoretically justified and user friendly. Computationally, the method is much simpler compared to other competitive methods such as Bayesian MCMC procedures or bootstrap resampling methods. We need sampling from posterior distribution only once during the model parameter estimation, and the sampled values can be used subsequently for all other purposes. The software is available from the authors upon request.

Acknowledgements

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Appendix

A. Derivation of the conditional distributions

From Equation (1) and (2), the conditional joint distribution of \( \{X_i, S_i^2, \theta_i, \sigma_i^2\} \), \( \pi(X_i, S_i^2, \theta_i, \sigma_i^2 \mid Z_i, B) \), is

\[
\pi(X_i, S_i^2, \theta_i, \sigma_i^2 \mid Z_i, B) = \frac{1}{\sqrt{2\pi\sigma_i^2}} \exp\left\{-\frac{(X_i - \theta_i)^2}{2\sigma_i^2}\right\} \frac{1}{\Gamma\left(\frac{n_i-1}{2}\right)} \frac{1}{2^{(n_i-1)/2}} \times \left\{(n_i-1)\frac{S_i^2}{\sigma_i^2}\right\}^{\frac{n_i-1}{2}} \times \left\{(n_i-1)\frac{1}{\sigma_i^2}\right\} \times \frac{1}{\Gamma(a)b^a} \frac{1}{\sigma_i^2} \exp\left\{-\frac{1}{b\sigma_i^2}\right\} \times \frac{1}{\Gamma(a)b^a} \frac{1}{\sigma_i^2} \exp\left\{-\frac{1}{b\sigma_i^2}\right\} \\
\times \exp\left\{-\frac{(\theta_i - Z_i^T \beta)^2}{2\tau^2}\right\} - \frac{(X_i - \theta_i)^2}{2\sigma_i^2} + \frac{(n_i-1)S_i^2}{2} + \frac{1}{b\sigma_i^2} \right\} \times \left\{1\right\} \frac{1}{\Gamma(a)b^a} \frac{1}{\sigma_i^2} \exp\left\{-\frac{1}{b\sigma_i^2}\right\}.
\]

Therefore the conditional distribution of \( \sigma_i^2 \) and \( \theta_i \) given the data and \( B \) are

\[
\pi(\sigma_i^2 \mid X_i, S_i^2, Z_i, B) = \int \pi(X_i, S_i^2, \theta_i, \sigma_i^2 \mid Z_i, B)d\theta_i \propto \frac{1}{(\sigma_i^2 + \tau^2)^{2\frac{n_i}{2}+a+1}} \exp\left\{-\frac{(X_i - \theta_i)^2}{2(\sigma_i^2 + \tau^2)} - \frac{1}{2}\frac{(n_i-1)S_i^2}{\sigma_i^2} + \frac{1}{b\sigma_i^2}\right\},
\]

\[
\pi(\theta_i \mid X_i, S_i^2, Z_i, B) = \int \pi(X_i, S_i^2, \theta_i, \sigma_i^2 \mid Z_i, B)d\sigma_i^2 \propto \exp\left\{-\frac{(\theta_i - Z_i^T \beta)^2}{2\tau^2}\right\} \psi_i^{\frac{n_i}{2}+a},
\]

where \( \psi_i \) is defined in Equation (4).

B. Details of the EM algorithm

The maximization of \( Q(B \mid B^{(t-1)}) \) is done by setting the partial derivatives with respect to \( B \) to be zero, that is,

\[
\frac{\partial Q(B \mid B^{(t-1)})}{\partial B} = 0.
\]

(B.1)

From the expression of \( Q(B \mid B^{(t-1)}) \) in the text, we give explicit expressions for the partial derivatives with respect to each component of \( B \). The partial derivative corresponding to \( \beta \) is

\[
\frac{\partial Q(B \mid B^{(t-1)})}{\partial \beta} = \sum_{i=1}^{n} \left\{Z_i \left(\frac{\theta_i - Z_i^T \beta}{\tau^2}\right) \exp\left\{-\frac{(\theta_i - Z_i^T \beta)^2}{2\tau^2}\right\} \psi_i^{\frac{n_i}{2}+a}\right\} d\theta_i = \sum_{i=1}^{n} E\left\{Z_i \left(\frac{\theta_i - Z_i^T \beta}{\tau^2}\right)\right\} d\theta_i
\]

where the expectation is with respect to the conditional distribution of \( \theta_i, \pi(\theta_i \mid X_i, S_i^2, B) \). The expression of the partial derivative corresponding to \( \tau^2 \) is:

\[
\frac{\partial Q(B \mid B^{(t-1)})}{\partial \tau^2} = \frac{1}{2}\sum_{i=1}^{n} \int \frac{(\theta_i - Z_i^T \beta)^2}{2\tau^2} \exp\left\{-\frac{(\theta_i - Z_i^T \beta)^2}{2\tau^2}\right\} \psi_i^{\frac{n_i}{2}+a} d\theta_i
\]

\[
= - \frac{n}{2\tau^2} + \sum_{i=1}^{n} \left\{\frac{(\theta_i - Z_i^T \beta)^2}{2(\tau^2)}\right\} \psi_i^{\frac{n_i}{2}+a} d\theta_i
\]

\[
= - \frac{n}{2\tau^2} + \sum_{i=1}^{n} E\left\{\frac{(\theta_i - Z_i^T \beta)^2}{2(\tau^2)}\right\}.
\]
Similarly for \(a\) and \(b\), we get the solutions by setting \(S_a = 0\) and \(S_b = 0\) where \(S_a\) and \(S_b\) are, respectively, the partial derivatives of \(Q(B \mid B^{(a-1)})\) with respect to \(a\) and \(b\) with expressions given in the main text. These equations are solved using the Newton-Raphson method which requires the matrix of second derivatives with respect to \(a\) and \(b\). These are given by the following expressions:

\[
S_{aa} = \sum_{i=1}^{n} \log'' \left\{ \Gamma \left( \frac{n_i}{2} + a \right) \right\} \\
- \log'' \{ \Gamma(a) \} + \text{Var} \{ \log(b) \}
\]

\[
S_{ab} = \sum_{i=1}^{n} \left[ -\frac{1}{b^2} \right] \left[ \frac{1}{b^3} \right] \left[ E \left( \frac{1}{\psi_i} \right) \right] \left( \frac{n_i}{2} + a \right) \frac{1}{b^2}
\]

\[
S_{bb} = \sum_{i=1}^{n} \left[ -\frac{1}{b^2} \right] \left[ \frac{1}{b^3} \right] \left[ E \left( \frac{1}{\psi_i} \right) \right] \left( \frac{n_i + 2}{b} \right) \frac{1}{b^4}
\]

\[
S_{ab} \text{ and } S_{bb} \text{ are } \text{given by (B.2).}
\]

\[
B_a \text{ and } B_b \text{ denote these quantities evaluated at the values of } a \text{ and } b \text{ at the } (u-1) \text{ iteration. Once the Newton Raphson procedure converges, the value of } a \text{ and } b \text{ at the } \tau \text{ step of the EM algorithm is set as } a^{(\tau)} = a^{(\infty)} \text{ and } b^{(\tau)} = b^{(\infty)}.
\]

\[
C. \text{ An alternative small area model formulation}
\]

It is possible to reduce the width of the confidence interval \(C(B)\) based on an alternative hierarchical model for small area estimation which has some mathematical elegance. The constant term \(n_i + 2a\) in (19) becomes \(n_i + 2a\) in this alternative model formulation. The model is given by

\[
X_i \mid \theta_i, \sigma_i^2 \sim N(\theta_i, \sigma_i^2), \quad (C.1)
\]

\[
\theta_i \mid \sigma_i^2 \sim N(\beta, \sigma_i^2), \quad (C.2)
\]

\[
(n_i - 1)S_i^2 \mid \sigma_i^2 \sim \chi^2_{n_i - 1}, \quad (C.3)
\]

\[
\sigma_i^2 \sim \text{Inverse - Gamma}(a, b), \quad (C.4)
\]

independently for \(i = 1, 2, ..., n\). Note that in the above formulation, it is assumed that the conditional variance of \(\theta_i\) is proportional to \(\sigma_i^2\) whereas the marginal variance is constant (by integrating out \(\sigma_i^2\) using (C.4)). In (1) and (2), the variance of \(\theta_i\) is a constant, \(\tau^2\), independent of \(\sigma_i^2\), and there is no conditional structure for \(\theta_i\) depending on \(\sigma_i^2\). The set of all unknown parameters in the current hierarchical model is \(B = (a, b, \beta, \lambda)\). The inference procedure for this model is given subsequently. The model essentially assumes that the true small area effects are not identically distributed even after eliminating the known variations.

\[
C.1 \text{ Inference methodology}
\]

By re-parameterizing the variance as in (C.2), some analytical simplifications are obtained in the derivation of the posteriors of \(\theta_i\) and \(\sigma_i^2\) given \(X_i, S_i^2\) and \(B\). We have

\[
\pi(\sigma_i^2 \mid X_i, S_i^2, B) = IG \left( \frac{n_i - 1}{2} \sigma_i^2 = \frac{(X_i - Z_i \beta)^2}{2(1 + \lambda)} + \frac{1}{b} \right)
\]

where \(IG(a,b)\) stands for the inverse Gamma distribution with shape and scale parameters \(a\) and \(b\), respectively. Given \(B\) and \(\sigma_i^2\), the conditional distribution of \(\theta_i\) is

\[
\pi(\theta_i \mid X_i, \sigma_i^2, B) = \text{Normal} \left( Z_i \beta, \frac{\lambda \sigma_i^2}{1 + \lambda} \right)
\]

Integrating out \(\sigma_i^2\), one obtains the conditional distribution of \(\theta_i\) given \(X_i, S_i^2\) and \(B\),

\[
\pi(\theta_i \mid X_i, S_i^2, B) = \int_0^{\infty} \pi(\theta_i \mid X_i, \sigma_i^2, B) \pi(\sigma_i^2 \mid X_i, S_i^2, B) \, d\sigma_i^2
\]

\[
= \int_0^{\infty} \pi(\theta_i \mid X_i, \sigma_i^2, B) \pi(\sigma_i^2 \mid X_i, S_i^2, B) \, d\sigma_i^2
\]

\[
= \int_0^{\infty} \frac{1}{2\lambda} \left( \frac{1}{1 + \lambda} \right) \frac{\delta^2}{2} \right)^{(n_i - 2a + 1)/2} \, d\sigma_i^2
\]

where \(\delta^2 = (n_i - 1)S_i^2 + (X_i - Z_i \beta)^2 / (1 + \lambda) / 2b\). We can rewrite (C.5) as

\[
\pi(\theta_i \mid X_i, S_i^2, B) = \frac{\Gamma((n_i + 1)/2 + a)}{\Gamma((n_i + 2)/2 + a)} \frac{\Gamma((n_i + 2)/2 + a)}{\Gamma((n_i + 2)/2 + a)} \frac{1}{\lambda} \frac{\delta^2}{2} \right)^{(n_i - 2a + 1)/2} \, d\sigma_i^2
\]

which can be seen to be a scaled t-distribution with \(n_i + 2a\) degrees of freedom and scale parameter \(\delta^2 / \lambda(1 + \lambda)\) with \(\delta^2 = \delta^2 / (n_i + 2a)\). Hence,

\[
E(\sigma_i^{-1} \mid X_i, S_i^2, B) = \frac{\Gamma((n_i + 1)/2 + a)\delta^2 / 2^{-(n_i + 1)/2 + a}}{\Gamma((n_i + 2)/2 + a)\delta^2 / 2^{-(n_i + 2)/2 + a}}
\]

\[
= \frac{\Gamma((n_i + 1)/2 + a)}{\Gamma((n_i + 2)/2 + a)} \frac{\sqrt{2}}{\delta^2 / (n_i + 2a)}.
\]
In this context, choosing
\[ k = k(B) = \left( 1 + \frac{t_{\alpha/2}^2}{n_i - 1} \right)^{-\frac{(n_i + 2\alpha + 1)/2}{2}} \left( \frac{1 + \lambda}{\lambda} \right)^{-\frac{1}{2}} \sqrt{2\pi}, \]
the confidence interval in (8) simplifies to
\[ C_i(B) = \left\{ \theta_i : \left| \frac{\theta_i - \mu_i}{\lambda} \left( \frac{1 + \lambda}{n_i - 1} \right) \right| \leq t_{\alpha/2} \right\}. \tag{C.6} \]
Using the similar arguments as before and noting that \((n_i + 2\alpha)\delta^2 \geq (n_i - 1)S_i^2\), we have \(P\{C_i(B)\} \geq P(D_i) = 1 - \alpha\), where \(D_i\) is the confidence interval in (20). When \(B\) is unknown, we replace \(B\) by its marginal maximum likelihood estimate \(\hat{B}\). It is expected that the pooling technique will result in an error small enough so that \(P\{C_i(\hat{B})\} \approx P\{C_i(B)\} \geq 1 - \alpha\).

**References**


